



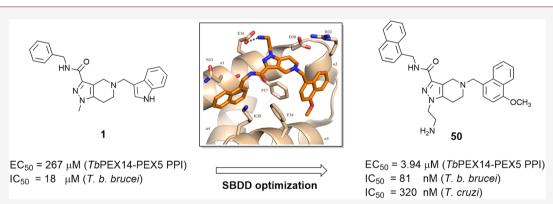
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Structure—Activity Relationship in Pyrazolo[4,3-c]pyridines, First Inhibitors of PEX14-PEX5 Protein-Protein Interaction with Trypanocidal Activity

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Supporting Information



ABSTRACT: Trypanosoma protists are pathogens leading to a spectrum of devastating infectious diseases. The range of available chemotherapeutics against Trypanosoma is limited, and the existing therapies are partially ineffective and cause serious adverse effects. Formation of the PEX14-PEX5 complex is essential for protein import into the parasites' glycosomes. This transport is critical for parasite metabolism and failure leads to mislocalization of glycosomal enzymes, with fatal consequences for the parasite. Hence, inhibiting the PEX14-PEX5 protein-protein interaction (PPI) is an attractive way to affect multiple metabolic pathways. Herein, we have used structure-guided computational screening and optimization to develop the first line of compounds that inhibit PEX14-PEX5 PPI. The optimization was driven by several X-ray structures, NMR binding data, and molecular dynamics simulations. Importantly, the developed compounds show significant cellular activity against Trypanosoma, including the human pathogen Trypanosoma brucei gambiense and Trypanosoma cruzi parasites.

INTRODUCTION

Human African trypanosomiasis (HAT, sleeping sickness) and Chagas disease (American trypanosomiasis) are among the

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HO₃S
$$\rightarrow$$
 SO₃H \rightarrow O \rightarrow HO₃S \rightarrow SO₃H \rightarrow SO₃

Figure 1. Structures of drugs used for the treatment of Trypanosoma-related diseases and clinical candidates.

most devastating and deadliest parasitic diseases. They occur predominantly in sub-Saharian Africa and Latin America, respectively, but Chagas disease has spread globally because of international human migration. Both qualify as neglected tropical diseases threatening millions of people in poor communities. The impact of *Trypanosoma* protists on livestock is also threatening the well-being of multiple communities.

HAT is caused by two protozoans: Trypanosoma brucei gambiense and Trypanosoma brucei rhodesiense (accounting for 98 and 2% of HAT cases, respectively), transmitted by the bite of an infected tsetse fly. The disease consists of two phases: the haemolymphatic stage (stage I) with trypanosomes being present in the blood and lymphatic system and the meningoencephalitic stage (stage II), when parasites have spread to the central nervous system. Without appropriate diagnosis and treatment, the disease can be fatal.^{3,6} No effective vaccine has so far been developed because the parasites employ several immune evasion strategies such as antigenic variation. Therefore, chemotherapy is the only therapeutic option currently available (Figure 1).^{7,8} Suramin and pentamidine are used in stage I of HAT, while melarsoprol, eflornithine, and nifurtimox-eflornithine combination therapy (NECT) are used in stage II of the disease. Unfortunately, these drugs have severe side effects and require continuous parenteral administration, which is not optimal, given that the patients live in rural African areas with very limited health resources.^{6,9} There is a better treatment option because recently a first all-oral drug fexinidazole completed clinical trials in humans and was approved for HAT treatment. 10-13 The efficacy of this monotherapy is comparable to intravenous NECT and the possibility of oral administration makes it easier to use in rural African realities. Nevertheless, cross-resistance with nifurtimox is likely to develop because of the same mechanism of action and no

convenient polytherapy is available to overcome this issue. ^{12,14–16} Another drug candidate, the benzoxaborole derivative SCYX-7158 (acoziborole) is currently in the advanced phase of clinical trials. ^{12,17} In spite of these promising developments, the Drugs for Neglected Diseases initiative (DNDi) still requires new candidates for stage-2 HAT treatment. ¹⁶

The situation is more dramatic for Chagas disease, caused by *Trypanosoma cruzi*. The parasite is spread by triatomine bugs ("kissing bugs"), by blood transfusions, congenitally, and even orally, by contaminated food.¹⁸ It is considered to be the parasitic disease with the greatest socioeconomic impact in Latin America and one of the leading causes of myocarditis. There are two stages of the disease. In the acute phase, the symptoms vary between the individuals and in most cases they decline spontaneously after 1–2 months. In the chronic phase, the patients may develop potentially fatal cardiac and digestive pathologies. Currently, there are two drugs for Chagas disease, benznidazole and nifurtimox (Figure 1). However, both have serious side effects and are ineffective in the chronic phase of the disease.^{4,19}

Undeniably, there is an urgent need for novel drugs against *Trypanosoma* infections, especially for Chagas disease, with high clinical efficiency, favorable pharmacological profiles, and ease of use. In an ideal setting, the pipeline would comprise several new agents that are suitable for combination therapy, which, in turn, would allow for better efficacy and minimal risk of resistance development.^{2,16} This can be achieved inter alia by finding promising targets and pathways essential for parasite survival. In the past, various molecular targets have been proposed and successfully validated.^{16,20,21} It is intriguing that although the trypanosomatids share similarities in genetic sequence and biochemical pathways,^{22,23} so far most drug discovery campaigns aimed either at *T. brucei* or *T. cruzi*, with

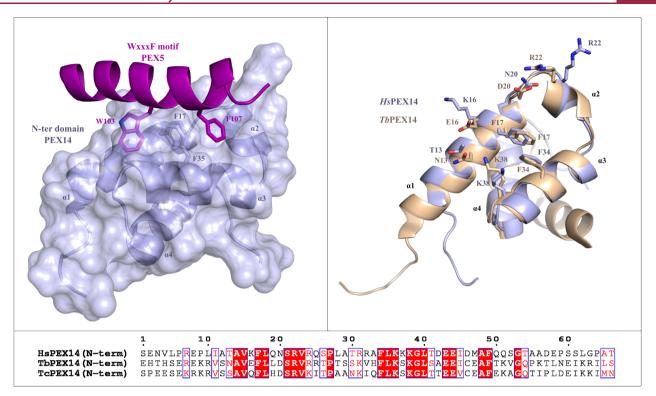


Figure 2. Structural basis for PEX14–PEX5 PPI interface targeting. Left: Solution NMR structure of the HsPEX14 NTD in complex with the diaromatic WXXXF pentapeptide motif of PEX5 (PDB code: 2W84³⁶). PEX14 is shown in light blue, whereas PEX5 is shown in magenta. Two hydrophobic cavities on the PEX14 surface accommodating Trp103 and Phe107 residues of PEX5 WXXXF are separated by two π-stacked phenylalanine side chains (Phe17 and Phe34). Right: Superposition of the solution NMR structures of HsPEX14 (light blue, PDB code: 2W84) and TbPEX14 (gold, PDB code: 5MMC³⁵) NTDs. The polar residues (represented as sticks) surrounding the binding pocket are significantly different in both species. Bottom: Sequence alignment of HsPEX14, TbPEX14, and TcPEX14.

fewer seeking for new, broad-spectrum agents which would affect both species. 24-27

Glycosomes are peroxisome-related organelles unique to kinetoplastids, including *Trypanosoma* spp. These organelles contain the enzymes required for glycolysis and for other intermediary metabolic pathways.²⁸ Hence, glycosomes are essential for parasite survival and they have long been considered attractive targets for development of new drugs against trypanosomatids.^{20,29,30}

As glycosomes lack DNA, glycosomal matrix proteins encoded in the nuclear genome have to be post-translationally transported from the cytosol to the organelle. Cargo proteins are imported by a cascade of interactions of specialized transport proteins, called peroxins (PEX#). The PEX14-PEX5 protein-protein interaction (PPI) plays a key role in this process. The PEX5 import receptor is capable of binding short peroxisomal targeting sequences (PTS) of the cargo. Next, the PEX5-cargo complex binds PEX14, which is localized at the glycosomal membrane and is required for cargo translocation into the matrix of the organelle. 30,31 Many of the glycosomal enzymes have been proposed and validated as potential drug targets.²⁰ However, the glycosomal transport system should be a more attractive drug target because it provides the opportunity to interfere with the function of many critical metabolic processes at once. Furthermore, mislocalization to the cytosol provides the glycosomal kinases access to the cytosolic ATP pool. Enzymes such as T. brucei hexokinase function at such a high rate that they deplete ATP faster than it can be replenished by substrate-level phosphorylation, which will be fatal to the trypanosomes.³²

A series of RNAi studies has previously shown the importance of PEX14 function for glucose metabolism in *Trypanosoma*. Recently, we reported a proof-of-concept study for targeting the PEX14–PEX5 PPI in *T. brucei* and *T. cruzi* parasites by small molecules. Using a structure-based drug discovery approach we developed a subset of prototypic pyrazolo [4,3-c] pyridine derivatives—the first inhibitors of this PPI and the first compounds that disrupt glycosomal import of matrix proteins and kill *T. brucei* and *T. cruzi* in vitro in nanomolar concentrations. Here, we present the details on the medicinal chemistry program that led to this group of active molecules along with the in-depth structure—activity relationship (SAR) study.

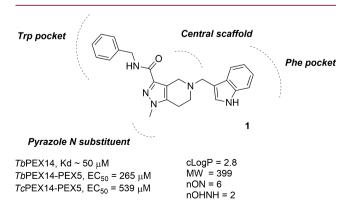


Figure 3. Structure, properties, and proposed derivatization sites of the pyrazolo [4,3-c] pyridine in silico hit 1.

Table 1. Initial Optimization of 1



				TbPEX14	TcPEX14	HsPEX14	T. brucei	HepG2	
#	\mathbf{R}_1	R ₂	R ₃		EC ₅₀ [μM] ^a				SIc
1	,£		[≯] CH ₃	265±37	539±75	223±30	18.5	>50	>2.7
2 ^d	£ \$	× NH	² CH₃	>1000	>1000	>1000	18.6 (13.7-25.2)	>50	>2.7
3	,5	× NH	×Ή	441±47	768±150	>1000		29.4 (20.8-41.6)	>5.7
4	,5-	* NH	∤ ∕∕он	421±19	828±127	>1000	21.0 (17.8-24.7)	>50	>2.4
5	2	A NH	<i>≱</i> _соон	>1000	>1000	>1000	>50	>50	nd
6	£	, st. NH	COOEt	550±144	>1000	>1000	16.7 (13.6-20.5)	>50	>3.0
7	,4	, st. NH	[≠] COOEt	>1000	>1000	>1000	3.05 (2.83-3.27)	29.7 (22.5 - 36.9)	>9.7
8	/CI	, the NH	X∕он	740±167	384±69	555±120	5.74 (4.57-7.22)	>20	>3.5
9	*** F	X NH	≯ ∕он	559±33	505±42	509±35	11.5 (6.75-19.7)	nd	nd
10	∤	× NH	≯ ✓OH	381±60	632±83	218±15	5.19 (3.04-7.34)	nd	nd
11	¢CH ₃	× NH	∕ ∕он	233±25	422±29	244±21	11.8 (3.9-35.7)	34.8 (31.3-38.7)	2.9
12	CF ₃	× NH	×∕он	879±140	625±94	427±54	4.25 (3.84-4.71)	nd	nd
13	*	* NH	[≠] CH ₃	51.3±2.1	99.5±25.7	51.1±4.1	2.96 (2.84 - 3.08)	14.4 (12.8-16.8)	4.9
14	, ž	X NH	×∕он	53.7±4.8	68.0±4.3	57.9±3.3	4.50 (3.87-5.23)	7.87 (7.28-8.44)	2.7
15	ž 🔾	ž C	 ≯∕он	>1000	>1000	>1000	2.15 (1.42-3.26)	>50	>23.2
16	£	₹ CI	<i></i> ≮∕∕oн	>1000	nd	nd		nd	nd
17	,\$	* CI	/ ∕ОН	>1000	nd	nd	nd	nd	nd
18	,5	[≯] COCH₃	×∕ОН	627±76	>1000	>1000	1.74 (0.82-3.60)	>50	>28.7
19	**		^{yk} ∕∕OH	>1000	nd	nd	nd	nd	nd
20	**************************************	V OCH₃	<i>у</i> с ОН	46.3±0.40	704±449	135±15	9.33 (8.43 - 10.3)	34.9 (28.3-41.5)	>3.6

 $[^]a$ EC $_{50}$ values were calculated as a Hill curve fit to 12 point titration. SD represents fitting error. nd = not determined. b EC $_{50}$ values are shown as mean (n = 4). Values in parentheses are 95% confidence intervals. c Selectivity index is calculated as HepG2 EC $_{50}$ [μ M]/T. brucei EC $_{50}$ [μ M]. d N-2 regioisomer of 1.

■ RESULTS AND DISCUSSION

Structural Basis for PEX14–PEX5 PPI Targeting. During the initial phase of the project, the structural basis for the PEX14–PEX5 PPI interaction was known (Figure 2). The N-terminal part of PEX5 contains several diaromatic WXXX(F/Y) motifs that address the respective hydrophobic Trp and Phe/Tyr pockets on the surface of the N-terminal helical domain (NTD) of PEX14. These two relatively shallow cavities are separated by two π -stacked phenyl rings of PEX14 (Phe17 and Phe34, the explanation of

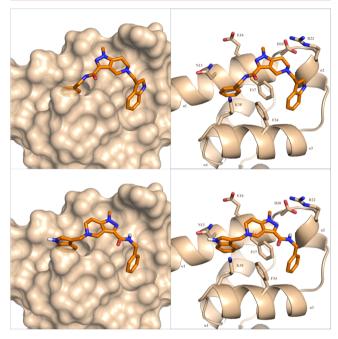


Figure 4. Docking poses of **1** to the TbPEX14—PEX5 PPI interface. Top: Phenyl residue of **1** occupies the Trp pocket while the indole is positioned in the Phe pocket on the PEX14 surface. Bottom: The "reversed" binding mode of **1** to the TbPEX14—PEX5 PPI interface. Surface (left) and cartoon (right) representation of TbPEX14 (gold).

residue numbering among different species is given in Table S1) that protrude to the solvent. In addition to the hydrophobic and aromatic binding of Trp and Phe/Tyr of WXXX(F/Y) motifs in PEX5 with their respective hotspots on the PEX14 surface, the positively charged Lys38 and Arg22 side chains of PEX14 plausibly facilitate the protein—protein complex formation by cation— π interactions. Other hydrophilic residues that might be of an importance for TbPEX14 ligand design are Asn13, Glu16, and Asp20. These three residues are different in *T. cruzi* and human sequences, and thus they might account for the ligands' cross-species selectivity (Figure 2, right).

The PEX14–PEX5 protein–protein interface poses several challenges for medicinal chemistry that are typical for the PPI target class.³⁹ First, it is hydrophobic in nature, with only few water exposed polar amino acid side chains in the spatial vicinity. Furthermore, the interface is relatively flat, with only two shallow cavities in the highly solvent-exposed surface. The Phe hotspot appeared as particularly difficult to address, being a "shelf" rather than a classical PPI pocket.

We created a pharmacophore model that mimicked the native binding of the PEX5 WXXX(F/Y) motif to TbPEX14. Two pharmacophoric elements were set as hydrophobic,

aromatic rings addressing the respective Trp and Phe pockets. Further, we anticipated that the presence of "protruding" Phe17 and Phe34 side-chains can be an exploitable feature for ligand design and we defined the third pharmacophore as another connecting aromatic ring that would allow for

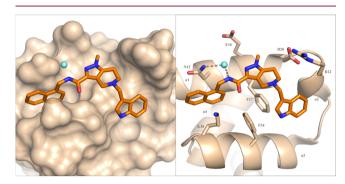


Figure 5. Crystal structure of TbPEX14 in complex with **13** (PDB code: 5L87³⁵). Left: surface representation of TbPEX14 (gold). A highly conserved water molecule is represented as a sphere. Right: cartoon representation of TbPEX14; relevant residues are shown as sticks. A water-mediated hydrogen bond between the PEX14 N13 side chain and the amide nitrogen of **13** is presented (gray dashed line).

additional π -stacking. The in silico screening cascade using a TbPEX14 homology model prepared from the NMR structure of HsPEX14 NTD (PDB 2W84)³⁶ identified several compounds with convincing docking poses. We tested these hits for binding to ¹⁵N-labelled TbPEX14 NTD using the ¹H–¹⁵N HSQC NMR chemical shift perturbations (CSP) assay. A pyrazolo[4,3-c]pyridine derivative 1 was the most potent hit, with a $K_{\rm D}$ of 163 μ M in the NMR CSP analysis and was resynthesized. The compound disrupted the interaction of the PEX5-derived peptide (ALSENWAQEFLA) with both TbPEX14 and TcPEX14, having an EC₅₀ of 265 and 539 μ M, respectively, in the AlphaScreen⁴³ assays (Figure 3 and Table 1). Furthermore, 1 was characterized by a proper drug-likeness profile and was chemically tractable, rendering this compound an attractive starting point for medicinal chemistry optimization.

The docking results suggested that the phenyl residue of 1 addressed the Trp pocket on the TbPEX14 surface, while the indole moiety filled the Phe hotspot (Figure 4, top). Such placement of aromatic residues is in fact a mirror image of the native binding mode of PEX5 WXXX(F/Y) fragments to PEX14. However, we have previously shown that the Trp pocket of PEX14 can effectively accommodate the Phe residue of the F/YXXXF motif in its PPI with another peroxin, PEX19.³⁶ Another hallmark of the docking-derived binding pose of 1 to PEX14 is that the pyrazolo [4,3-c] pyridine central scaffold of the hit compound lays over the Phe17 and Phe34 residues, forming the favorable π – π interactions. We have also identified lower-scored poses, in which the projection of the aromatic phenyl and indole residues of 1 in TbPEX14 cavities was reversed, that is, directly mimicking the native placement of the aromatic rings in the PEX14-PEX5 PPI (Figure 4, bottom). Although this inverted binding pose seemed less plausible because of the number of possible unfavorable interactions with the hydrophilic residues present in the vicinity of the hotspots, it could not be completely disregarded

Table 2. Secondary Optimization of the Residue Addressing the Phe Pocket



ThPEX14 TcPEX14 HsPEX14 T. brucei HenG2

14	3.3
14	3.3
21	3.3
22	10.14)
24 Section 166±15 141 (2.82-11.70) (18.0-1)	nd
25 18±12 176 (5.66-6.56) (9.9-1) (9.56-6.56) (9.9-1) (9.56-6.56) (9.9-1) (1.9-	27.4) 4.0
25	7.5)
26 10	3.2
27	8.42)
28 HN 20.3±2.6 43.6±4.4 35.5±1.6 (4.29-6.84) (8.64- 29 14.8±1.9 104±40 39.7±5.0 (3.29-3.85) >50 30 454±143 >1000 >1000 7.95 9.11 31 454±143 >1000 >1000 20 >50 32 454±143 142±49 60.8±4.7 nd nd 33 45CH ₃ 19.0±4.4 38.0±11.8 32.4±14.8 (2.78-5.51) nd 34 45CH ₃ 19.0±4.4 38.0±11.8 32.4±14.8 (2.78-5.51) nd 34 45CH ₃ 19.0±4.4 38.0±11.8 32.4±14.8 (2.78-5.51) nd	2.7
29	1.7
31	>14.0
32 CH ₃ 38.3±1.7 142±49 60.8±4.7 nd nd nd 33 CH ₃ 19.0±4.4 38.0±11.8 32.4±14.8 (2.78-5.51) nd 34 CH ₃ 122±27 168±11.1 437±137 8.40 29.2 (7.53-9.36) (25.9±10.15) (7.53-9.36) (25.9±10.15) (7.53-9.36) (25.9±10.15) (25.9±10.1	9.24)
OCH ₃ 33 19.0±4.4 38.0±11.8 32.4±14.8 (2.78-5.51) nd 34 122±27 168±11.1 437±137 (7.53-9.36) (25.9±10)	nd
33 19.0±4.4 38.0±11.8 32.4±14.8 (2.78-5.51) nd	nd
34 122±27 168±11.1 437±137 (7.53-9.36) (25.9-5	nd
	3.5
35 16.2±5.9 21.0±2.2 16.4±2.3 5.70 (4.81-6.76) >50	>8.8
36 200±25 >1000 503±55 13.1 38.2 (11.3-14.9) (30.6-	2.9
37 75.1±2.9 387±178 181±19 7.26 (6.07-8.45) nd	nd
38 7.62 nd nd (6.68-8.72) nd	nd
39 56.2±5.6 126±51 149±30 3.29 nd (3.16-3.44)	nd
40 51.7±4.5 369±201 113 6.22 (5.90-6.52) nd	nd
41 53.2±7.0 229±44 86.1±8.6 3.98 14.1 (12.3-	3.5
42 OCF ₃ 95.0±14.8 280±54 116±15 5.64 (5.08-6.21) nd	nd
43 COCH ₃ 50.9±4.8 287±64 138±13 10.4 40.8 (8.3-12.9) (34.7-	3.9
44.7±5.4 354±173 95.5±9.8 4.29 nd (3.36-5.22)	nd
45 137±63 259±160 422±342 nd 13.8 (12.3-	nd 15.4)

Table 2. continued

 $^a\mathrm{EC}_{50}$ values were calculated as a Hill curve fit to 12 point titration. SD represents fitting error. nd = not determined. $^b\mathrm{EC}_{50}$ values are shown as mean (n=4). Values in parentheses are 95% confidence intervals. $^c\mathrm{Selectivity}$ index is calculated as HepG2 EC_{50} [$\mu\mathrm{M}$]/T. brucei EC_{50} [$\mu\mathrm{M}$].

because the overall shape of the PEX14—PEX5 PPI interface is highly symmetrical.

Initial Optimization Trials. For optimization purposes we divided the initial hit 1 into four regions: (1) the pyrazole N substituent, (2) the residue addressing the Trp pocket, (3) the residue addressing the Phe pocket, and (4) the central scaffold (Figure 3).

According to the preferred docking pose of 1, the methyl group on pyrazole N-1 atom points to the solvent-exposed region of the interface, suggesting that substituents of different sizes should be tolerable at this position. Despite the proximity of Asn13 (Ser13 in TcPEX14 and Thr13 in HsPEX14) and Glu16 (Gln16 in TcPEX14 and Lys16 in HsPEX14), we expected that modifications addressing this region of the interface may affect the ligand's physicochemical properties (e.g., solubility) rather than significantly influence the affinity to the target. We have found that the binding to PEX14 depends on the position of the substituent in the pyrazole ring. Contrary to the active hit 1, the N-2 regioisomer 2 showed modest inhibitory activity in the AlphaScreen assay (Table 1) and did not bind TbPEX14 in a 1H-15N HSQC NMR experiment. This was inconsistent with the anticipated binding mode of 1 to TbPEX14 because the distance to the proximal Asn13 and Glu16 should allow the methyl group to be tolerated at the N-2 position of the pyrazole. Compound 3 lacking substituents at the N-1 atom of the pyrazole had a slightly decreased ability to disrupt PEX5 interaction with both T. brucei and T. cruzi PEX14, and it was inactive against the human protein. Likewise, the hydroxyethyl derivative 4 showed no superior activity compared to 1. Carboxylate derivatives 5-7 proved inactive. In conclusion, the initial SAR of the solventexposed region could not be fully rationalized on the basis of molecular docking. We have chosen the hydroxyethyl residue in this position because of the superior solubility of the compounds and sought further structural modifications that would yield more potent compounds suitable for cocrystallization trials with TbPEX14.

Because the binding of the PEX5 WXXX(F/Y) motifs to PEX14 is hydrophobic and aromatic in nature, we assumed that optimizing the residues addressing the Trp and Phe hotspots would likely improve ligand affinity. Structural data show that the Trp pocket is quite shallow (Figures 2 and 4). As a result, the orientation of the indole ring in the native PEX14-PEX5 PPI is vertical with respect to the surface of the interface, and the imidazole component of the fused system slightly points out from the hotspot cavity, being surrounded by Lys38 and Asn13 (Ser13 in TcPEX14 and Thr13 in HsPEX14). On the other hand, this pocket has an elongated shape and the docking pose of 1 suggested that there is space that could be filled by a residue larger than the unsubstituted phenyl. Hence, we first modified the substitution pattern on the phenyl ring (Table 1). We synthesized derivatives 8-12having various small lipophilic substituents in meta- and paraposition, none of which showed superior activity or species selectivity, when compared to 1. In contrast to the flat SAR of

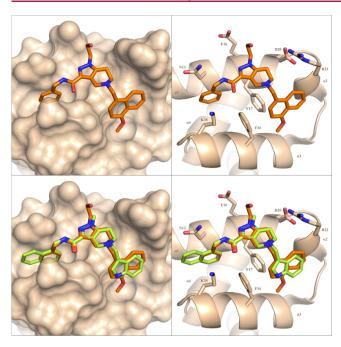


Figure 6. Top: Crystal structure of TbPEX14 in complex with **20** (PDB code: 5L8A³⁵). Bottom: overlay of the binding poses of **13** (purple) and **20** (green). Surface (left) and cartoon (right) representation of TbPEX14 (gold).

the substituted phenyls, derivatives 13 and 14 bearing naphthalene moieties, were more effective in disrupting the TbPEX14–PEX5 PPI, with AlphaScreen-derived EC_{50} values

five and eightfold lower than the respective parent compounds 1 and 4. We have also observed the increased ability of these derivatives to interrupt the Tc and HsPEX14-PEX5 complex formation.

As already mentioned, the Phe hotspot on the PE14–PEX5 PPI interface appeared more difficult to target than the Trp pocket because this cavity is shallow and more exposed to solvent. The docking pose of 1 in TbPEX14 showed that this region of the interface is well filled by the indole moiety of the inhibitor. In fact, replacement of the bicyclic aromatic system of 1 with smaller phenyls yielded only very weak compounds 15–18 (Table 1). Interestingly, inhibitor 19 bearing a bulky naphthalene was inactive, while its 4-methoxy derivative 20 proved 10-fold more potent in disrupting the TbPEX14–PEX5 PPI than 4.

Secondary Optimization—Phe Pocket. The above SAR was in large parts difficult to rationalize without additional structural data. Fortunately, at this stage of the project we were able to obtain a high-resolution X-ray structure of TbPEX14 bound to 13 (PDB code: 5L87,³⁵ Figure 5) that guided further optimization. Overall, the crystallographic data were consistent with the binding mode of 1 to TbPEX14 as predicted by docking. According to the TbPEX14-13 complex structure, the double aromatic naphthalene system effectively filled the Trp hotspot. However, unlike the vertically oriented Trp side chain in the native PEX5 WXXX(F/Y)—PEX14 PPI, the orientation of the naphthyl group in the Trp pocket is horizontal. A very important feature of the X-ray structure that could not have been predicted on the basis of docking is the presence of one structural water that mediated binding of 13 to the TbPEX14

Table 3. Secondary Optimization around the Trp Pocket

	_	TbPEX14	TcPEX14	HsPEX14	T. brucei	HepG2	GT:
#	compound structure	EC ₅₀ [μM] ^a	EC ₅₀ [μM] ^a	EC ₅₀ [μM] ^a	EC ₅₀ [μM] ^b	EC ₅₀ [μM] ^b	SIc
46	N N OCH3	49.2±19.5	463±203	>1000	6.42 (5.61-7.20)	>50	>7.8
47	OCH3	103±17	448±408	>1000	5.61 (5.13-6.11)	>50	>8.9
48	HN N OCH3	174±17	176±9	nd	nd	29.6 (24.4-36.2)	nd
49	Q, O HN N N OCH ₃	>1000	208±33	>1000	5.94 (3.63-9.73)	16.4 (13.6-19.7)	2.8

 $[^]a$ EC₅₀ values were calculated as a Hill curve fit to 12 point titration. SD represents fitting error. nd = not determined. b EC₅₀ values are shown as mean (n = 4). Values in parentheses are 95% confidence intervals. c Selectivity index is calculated as HepG2 EC₅₀ [μ M]/T. brucei EC₅₀ [μ M].

Table 4. Secondary Optimization of the Pyrrole N-Substituent: Amines

	_	TbPEX14	TcPEX14	HsPEX14	T. brucei	HepG2	
#	R	EC ₅₀ [μM] ^a		EC ₅₀ [μM] ^a	EC ₅₀ [μM] ^b	EC ₅₀ [μM] ^b	SIc
29	∕ ∕он	14.8±1.9	104±40	39.7±5.0	3.56 (3.29-3.85)	>50	>14.0
50	× NH ₂	3.94±0.40	22.1±2.3	19.3±2.3	0.081 (0.075-0.087)	3.74 (3.04-4.44)	46.1
51	[₹] NH	1.86±0.09	31.9±1.7	9.31±1.07	0.099 (0.095-0.103)	7.36 (6.33-8.39)	74.3
52	× NH	1.80±0.10	25.3±1.44	7.89±1.14	0.070 (0.051-0.096)	nd	nd
53	× NH	1.92±0.19	19.9±1.5	9.55±0.93	0.292 (0.275-0.310)	>50	>100
54	**\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\	3.2±0.48	21.5±1.7	8.69±0.84	0.236 (0.223 - 0.248)	8.84 (6.53-10.75)	37.5
55	× N N N	2.9±0.2	10.7±1.3	7.25±0.56	2.46 (2.22-2.79)	7.54 (5.30-10.35)	3.1
56	p. N. N.	34.4±11.2	247±113	94.5±21.4	3.60 (3.23-4.20)	nd	nd
57	*	88.1±13.1	162±43	166±35	3.58 (3.19-4.02)	42.0 (32.9-59.9)	11.7
58	× NH OH	17.9±1.17	70.8±7.6	33.5±5.0	5.73 (4.90-6.70)	4.92 (3.60-6.25)	<1
59	N COOH	4.05±0.60	41.9±3.2	nd	4.74 (4.14-5.38)	nd	nd
60	N COOMe	57.6±6.5	333±34	nd	4.68 (4.36-5.03)	nd	nd
61	» OH	7.78±1.6	27.5±3.4	9.31±1.07	0.503 (0.486 - 0.531)	6.50 (6.06-7.16)	12.9

 a EC₅₀ values were calculated as a Hill curve fit to 12 point titration. SD represents fitting error. nd = not determined. b EC₅₀ values are shown as mean (n = 4). Values in parentheses are 95% confidence intervals. c Selectivity index is calculated as HepG2 EC₅₀ [μ M]/T. brucei EC₅₀ [μ M].

Asn13 side chain. This has a pronounced influence on the ligand binding mode. Most of all, the ligand is shifted toward Asn13 and Glu16 residues with respect to its orientation speculated on the basis of docking. As a consequence, the indole residue of 13 is shifted out toward the upper rim of the Phe pocket rather than being deeply inserted into this cavity as we postulated previously. Thus, the filling of the hotspot by the indole can be considered shallow and suboptimal. On the other hand, the aromatic residue of 13 is closer to the Arg22 side chain of PEX14, which, in turn, can stabilize the ligand—protein complex by forming cation— π interactions. The presence of the abovementioned thermodynamically stable water molecule may also explain the loss of affinity of 2, in which the methyl group attached to the N-2 pyrazole nitrogen

atom interferes with the solvation shell of the protein—ligand complex.

Driven by the obtained high-resolution structural information we decided to systematically exploit the SAR of the Phe hotspot, while keeping the Trp pocket occupied by the naphthyl (Table 2). The overall optimization strategy was thus focused on preserving the orientation of the aromatic portion of the substituents addressing the Phe hotspot close to its rim, while simultaneously extending the ligand deeper toward its bottom. First, we synthesized compounds 21–24 bearing various small substituents in the N-1 position of the indole system, of which only the methyl and ethyl derivatives 21 and 22, respectively, showed potency comparable to 14 in the AlphaScreen assay. On the other hand, compounds 23 and 24, with larger, more polar pendant residues, were slightly less

active in disrupting the PEX14–PEX5 PPI. As seen in the TbPEX14-13 complex structure, the Phe pocket of the interface is filled tightly by the indole residue of the ligand. Hence, there is no possibility of rotation along the $N-CH_2$ bond that links the indole residue to the central scaffold, which would be required to accommodate large indole N-1 substituents. Furthermore, as observed in the TbPEX14-13 complex structure, the ligand binds to the interface in its low-energy conformation. Hence, a considerable entropic penalty would have to be paid for the N-1 substituted ligand adaptation to the hotspot.

While indole N-1 substituents are in a steric conflict with the amino acid side chain forming the Phe pocket and there is no possibility to compensate this by the adjustment of ligand's orientation, the analysis of the TbPEX14-13 complex structure showed that there is enough space in the hotspot to accommodate substituents in the C-7 position. Therefore, in the next optimization round we synthesized compounds 25-27 having small residues attached to this carbon atom (Table 2). Gratifyingly, these proved superior to 14 in disrupting the TbPEX14-PEX5 complex formation, each being 2 times more potent than the parent compound in the AlphaScreen assay. As of Tc and HsPEX14-PEX5 PPI inhibition, the results varied. Compound 25 had a comparable activity to the parent 14, while its N-1 methyl derivative 26 was 2 times more active. On the other hand, 27 bearing a methyl ester function on C-7 of indole was twofold less potent in inhibiting the *T. cruzi* protein. We have also synthesized 28, in which indole was attached to the central scaffold through a C-2 carbon atom, allowing for a deeper insertion of the aromatic residue into the pocket (Table

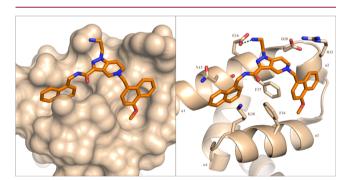


Figure 7. High resolution crystal structure of N-terminal domain of TbPEX14 in complex with compound **50** (PDB code: 5N8V³⁵). Surface (left) and cartoon (right) representation of TbPEX14 (gold).

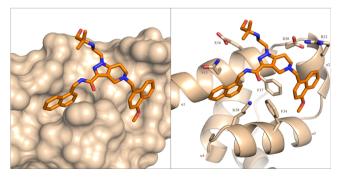


Figure 8. High resolution crystal structure of N-terminal domain of TbPEX14 in the complex with compound **61** (1.2 Å, PDB code: 6SPT). Surface (left) and cartoon (right) representation of TbPEX14 (gold).

2). As a result, 28 was more active than its C-3 regioisomer 14 and roughly equipotent to the isomeric methoxy indole derivative 26.

The different activities of the undecorated indole derivative 14 and 26 resulting from the unequally deep occupation of the Phe pocket recalled the observations made previously for naphthalenes 19 and 20 during the initial optimization trials. With the aim to improve binding of the ligands and to further investigate SAR in the Phe pocket, we solved the crystal structure of the methoxynaphthalene derivative 20 in complex with TbPEX14 (PDB code: 5L8A, 35 Figure 6). Overall, the binding mode of 20 to the TbPEX14-PEX5 PPI interface closely resembled the one observed for 13 in the TbPEX14-13 complex. Thus, the phenyl ring of the inhibitor was buried in the Trp pocket, while the naphthalene system filled the upper part of the Phe hotspot, with the methoxy residue reaching the bottom part of the cavity. Importantly, the orientation of the pyrazolo [4,3-c] pyridine central scaffold in 13 and 20 was identical, which was a clear indication that a more potent PEX14-PEX5 PPI inhibitor can be obtained by merging those two ligands. Indeed, the hybrid molecule 29 showed a superior activity in the AlphaScreen TbPEX14-PEX5 PPI inhibition assay, when compared to both parent compounds (Table 2). The methoxy group present at the C-4 atom of the naphthalene in 29 was crucial for the high-affinity interaction with TbPEX14 because derivatives 30 and 31 lacking this residue proved significantly less active. Next, we synthesized 32, in which one of the fused benzenes of the naphthalene system was saturated. The compound was 3 times less potent than the parent 29, most likely because of its inability to form cation $-\pi$ and π – π interactions with the Arg22 and Phe17 residues, respectively. Next, we investigated compounds 33-35 having small substituents other than methoxy at C-4 carbon of the naphthalene. Switching to a larger methylthiol group in 33 resulted in a 3 times stronger inhibition of the TcPEX14-PEX5 complex formation when compared to 29. Sulfoxide 34, an expected metabolite of 33, was significantly less potent. The dimethylamino derivative 35 was as active as the parent compound 29 in inhibiting the TbPEX14-PEX5 PPI and significantly more potent against Tc and HsPEX14.

Finally, we synthesized a series of phenyl derivatives 36–45 having small lipophilic substituents in para- or (and) metapositions (Table 2). By this, we wanted to verify whether a single aromatic system can outcompete, or at least be as active as, the bulkier and highly lipophilic indoles and naphthalenes. By changing the substitution patterns on the phenyl, we have observed that a gain of potency can be achieved by expanding the ligand toward the bottom of the pocket, similar to what had been observed for the bicyclic aromatic systems in 14 and 21–35. Several phenyl derivatives were equipotent to 14 in disrupting the TbPEX14–PEX5 complex formation; however, none of them could outcompete the more potent 25–28 or the C-4-substituted naphthalenes 29 and 33–35.

Secondary Optimization—around Trp Pocket. To further expand the SAR information in PEX14–PEX5 PPI targeting we turned our attention to the region of the interface near the entrance to the Trp hotspot. The crystallographic structures of TbPEX14 in complex with 13 and 20 show that the amide moiety that projects the aromatic residue of the ligand into the Trp pocket is in a low-energy conformation that is coplanar with the pyrazole ring. Hence, we posed the question whether an entropic gain from freezing this favorable conformation can be achieved and we synthesized 46 and 47,

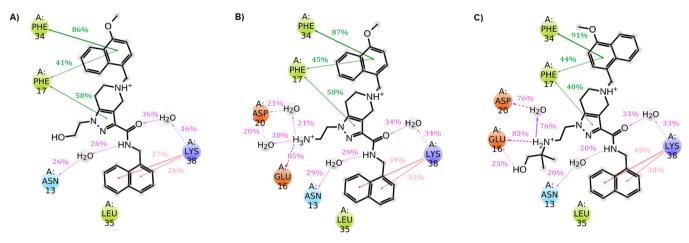


Figure 9. Ligand interaction diagrams of the final MDs simulations snapshots. The MDs were performed for TbPEX14-**29** (A), TbPEX14-**50** (B), and TbPEX14-**61** (C) complexes. Blue lines represent H-bonding interactions, red cation— π , and green π — π interactions. Introduction of alkylamine residues at N-1 pyrazolo position leads to formation of direct and water-mediated interaction patterns significantly improving compound affinity.

tricyclic derivatives of 29 (Table 3). Both compounds were less potent than the parent inhibitor in the AlphaScreen assay. As previously mentioned, the TbPEX14-13 complex structure shows an important interaction of the amide NH proton with Asn13 of the protein, mediated through a water molecule. Hence, it is possible that the gain in entropy in 46 and 47 from freezing the molecules in their low-energy conformations cannot compensate the loss in the binding affinity caused by the inability of forming the mentioned important connection through water. Intriguingly, this structural change proved deleterious for the ability of the inhibitors to bind Tc and HsPEX14 but did not abolish its interaction with the T. brucei protein. As a result, 46 and 47 are highly selective in inhibiting the latter PPI. To further explore the importance of the amide linkage in this region of the ligand, we synthesized 48, a derivative of 29 in which carbonyl oxygen was removed (Table 3). The obtained inhibitor projecting the naphthalene into the Trp pocket through an alkylamino linker was significantly weaker in inhibiting the PEX14-PEX5 PPI than the parent molecule. We attributed this to the repulsive interactions with Lys38, weaker hydrogen bond-accepting properties of the amine, and to the increased conformational freedom of the linker when compared to the amide. With the aim to explore the possible polar interactions with Asn13 and Lys38 surrounding the entrance to the Trp pocket, we synthesized sulfonamide 49 (Table 3). The compound failed to inhibit the TbPEX14-PEX5 PPI, likely because of the inability of the relatively narrow Trp pocket to accommodate the sulfonamide moiety.

Secondary Optimization—Amines in N-1 Pyrazolo Substituent. Our initial optimization trials showed that the substitution pattern at N-1 of pyrazole does not significantly alter binding of the ligands to the PEX14–PEX5 PPI interface. However, a closer inspection of the TbPEX14-20 complex structure revealed that the pendant hydroxyalkyl chain can adopt a conformation in which the hydroxyl lies in a spatial proximity of the carboxylic amino acid residues, Glu16, Asp20, in the solvent-exposed region. Therefore, we posed a question if a switch to an amine can lead to a further improvement in the binding affinity originating from additional electrostatic contacts. Indeed, a simple –OH to–NH₂ transition resulted in primary amine 50 which was sixfold more active in disrupting

the TbPEX14–PEX5 PPI and fivefold more potent against TcPEX14, respectively, than the parent **29** (Table 4). Encouraged by this result, we pursued a series of derivatives **51–61** with various alkylamine residues at the N-1 pyrazolo position. Secondary and tertiary amines were tolerated. An important observation was that binding to PEX14 correlates positively with the basicity of the substituents (e.g., **50** vs **58**, **54** vs **57**, **55** vs **56**, and **59** vs **60**) rather than with their hydrogen bond donor or acceptor capabilities (e.g., **50** vs **58**).

We solved high-resolution crystal structures of 50 and 61 in the complex with the TbPEX14 N-terminal domain (PDB codes: 5N8V³⁵ and 6SPT, Figures 7 and 8, respectively). The TbPEX14-50 complex structure provides a snapshot of the pendant amino group interacting with Glu16 (Figure 7). However, this interaction is not seen in the TbPEX14-61 complex structure (Figure 8), which reflects the flexibility of the hydroxyalkyl chain. As X-ray crystallography may not always precisely reveal the flexible ligand-protein interactions because of the crystal symmetry-related contacts, we analyzed the binding of the amino groups in 50 and 61 and hydroxyl of 29 with the carboxylic residues of Glu16 and Asp20 of TbPEX14 by performing molecular dynamics (MD) simulations (Figures 9 and S2—see Supporting Information). The computations clearly show the high occurrence of interactions of the amino groups with both residues. In contrast, the MD simulation did not show significant interaction of the pendant hydroxyethyl group of derivative 29. Here, the occurrence of the interaction during the MD run is much lower.

The negatively charged Glu16 residue is present in the TbPEX14 amino acid sequence, but not in TcPEX14 (Gln16) or HsPEX14 (Lys16). Furthermore, in the HsPEX14 sequence Asp20 is replaced by the neutral Asn20. In this context, it is intriguing that the ligands bearing positive charge (i.e., amines 50–55, 59, and 61) show enhanced inhibitory activity in the latter two PPIs. On the other hand, it has been shown that the binding affinity and thermodynamics in solvent-exposed protein cavities, such as those present on the PEX14 surface, is not only affected by the direct interactions of ligand atoms with protein residues, but can also depend significantly on the alterations of the water envelope around the ligand—protein complex.⁴⁴ Indeed, both TbPEX14-50 and TbPEX14-61 complex X-ray structures document the involvement of the

Table 5. Secondary Optimization—Central Scaffold

"		TbPEX14	TcPEX14	HsPEX14	T. brucei IC50	HepG2	CITE
#	Compound structure	EC ₅₀ [μM] ^a	EC ₅₀ [μM] ^a	EC ₅₀ [μM] ^a	[<i>µ</i> M] ^b	IC50 [μM]b	SIc
29	HN OCH3	14.8±1.9	104±40	39.7±5.0	3.56 (3.29-3.85)	>50	>14.0
62	HN OCH3	17.8±1.8	54.4±3.9	34.0±3.9	10.1 (8.5-13.4)	>50	>5.0
63	HN O OMe	8.17±0.69	30.0±1.8	15.3±0.5	6.01 (4.65-7.77)	nd	nd
50	HN O OCH3	3.94±0.40	22.1±2.3	19.3±2.3	0.081	3.74 (3.02-4.44)	46.2
64	H ₂ N OMe	1.22±0.22	15.8±1.2	4.38±0.71	0.225 (0.207-0.244)	3.67 (3.52-3.82)	16.3
65	HN OME	0.67±0.15	11.7±0.4	3.48±0.11	0.549 (0.513 - 0.587)	3.82 (3.60-4.04)	7.0

 $^{a}\text{EC}_{50}$ values were calculated as a Hill curve fit to 12 point titration. SD represents fitting error. nd = not determined. $^{b}\text{EC}_{50}$ values are shown as mean (n = 4). Values in parentheses are 95% confidence intervals. $^{c}\text{Selectivity}$ index is calculated as HepG2 EC₅₀ $[\mu M]/T$. brucei EC₅₀ $[\mu M]$.

flexible alkylamines in the water network around the ligand. Similar engagement can be expected in Tc and HsPEX14, which might explain the enhancement in inhibitory activity of the ligands against the PPIs of those proteins with PEX5.

Overall, the sum of the effects described above provided a significant improvement of the activity of the ligands against all three tested PEX14 proteins. This enhancement was most pronounced in the case of TbPEX14–PEX5 PPI, which resulted in the higher selectivity of compounds toward this PPI.

Secondary Optimization—Central Scaffold. Finally, we decided to obtain some SAR information of the central bicyclic scaffold. Quaternization of the tertiary amine in **29** resulted in a permanently charged derivative **62**, which did not show

superior binding to PEX14 (Table 5). Decreasing the size of the aliphatic ring of **29** by one methylene unit produced the pyrrolo[3,4-c]pyrazole **63**, which was twofold more active than the parent compound in inhibiting the TbPEX14–PEX5 PPI, and threefold more potent against TcPEX14 and HsPEX14. The amino derivative of alcohol **63**, compound **64**, showed the expected enhancement in PEX14 binding, with respect to its six-membered ring homologue **50**. Finally, the amino alcohol **65** was the most potent PEX14 binder, representing a first inhibitor of TbPEX14–PEX5 PPI with submicromolar EC₅₀ value in the AlphaScreen assay.

Cellular Activity of the PEX14–PEX5 PPI Inhibitors. The synthesized compounds were evaluated for their trypanocidal effect against the *T. brucei brucei* bloodstream

Table 6. In Vitro Trypanocidal Activity of Compounds Against T. b. rhodesiense STIB900 and T. Cruzi

#	T. b. rhodesiense IC_{50} $[\mu M]^a$	T. cruzi IC_{50} $[\mu M]^a$	L6 IC ₅₀ [μΜ] ^a	SI ^b
1	1.52 ± 0.67	9.16 ± 0.87	16.8 ± 0.2	1.8
2	6.00 ± 0.06	16.8 ± 0.9	34.5 ± 3.9	2.1
4	6.38 ± 0.40	21.5 ± 0.2	43.2 ± 3.2	2.0
14	2.14 ± 0.18	5.22 ± 0.01	5.93 ± 0.16	1.1
29	4.42 ± 0.16	3.83 ± 1.94	12.7 ± 2.6	3.3
33	5.54 ± 0.28	4.35 ± 0.09	17.6 ± 0.70	4.0
50	0.012 ± 0.004	0.320 ± 0.074	1.80 ± 0.06	5.6
52	0.009 ± 0.001	3.16 ± 1.36	1.66 ± 0.01	<1
58	1.98 ± 0.57	7.22 ± 1.30	5.84 ± 0.70	<1
61	0.029 ± 0.001	2.20 ± 0.28	1.82 ± 0.10	<1
63	1.89 ± 0.28	1.52 ± 0.32	9.19 ± 0.42	6.0
64	0.075 ± 0.008	0.627 ± 0.290	1.96 ± 0.04	3.1
65	0.179 ± 0.048	2.28 ± 0.12	1.96 ± 0.10	<1

 $[^]a\mathrm{EC}_{50}$ values are shown as mean \pm deviation from the mean (n=2). $^b\mathrm{Selectivity}$ index is calculated as L6 EC_{50} [$\mu\mathrm{M}$]/T. cruzi EC_{50} [$\mu\mathrm{M}$].

form (Lister 427, MITat 1.2) in the resazurin-based cell survival assay. T. b. brucei is the causative agent of animal trypanosomiasis and was used as a model organism for the primary screening of trypanocidal activities of the inhibitors. The obtained EC_{50} values are presented in Tables 1–5. Besides several outliers, the trypanocidal activity of the tested derivatives correlates well with the results of the AlphaScreen-based TbPEX14-PEX5 PPI inhibition assay. We have also observed a remarkable enhancement of the antiparasitic activity in compounds bearing an aminoalkyl chain attached to the N-1 position of the pyrazole, resulting in submicromolar EC50 values in the resazurin-based assay (Tables 4 and 5). We attribute this to the greater inhibitory potency of these compounds against the TbPEX14-PEX5 PPI. Compounds 50-52 were most active from the series, having EC₅₀ values below 100 nM (Table 4).

Additionally, the investigated PEX14–PEX5 PPI inhibitors were assayed for their cytotoxicity in the human-derived HepG2 cell line (Tables 1–5). In general, the cytotoxicity of some of the compounds followed their inhibitory activity against HsPEX14–PEX5 PPI. This was most pronounced for many of the indole derivatives (e.g., 2–8, 14, and 21–28). On the other hand, some other active HsPEX14 ligands did not follow this trend (e.g., naphthalenes 20, 29, and 35). Overall, the selectivity index for trypanocidal activity against *T. b. brucei*

improved along with increased efficacy of compounds in disrupting PEX14—PEX5 PPI. This was most pronounced in the alkylamine derivatives (Tables 4 and 5), many of which displayed favorable selectivity profiles (e.g., 50, 51, 53, 54, 61, 64, and 65).

The PEX14—PEX5 PPI inhibitors showing high trypanocidal efficacy against *T. b. brucei* were further evaluated activity against *T. b. rhodesiense* STIB900, a human pathogen derived from an African (United Republic of Tanzania) patient (Table 6). In general, the results followed the SAR pattern observed in the primary *T. b. brucei* screening. The *T. b. rhodesiense* STIB900 strain was even more susceptible to the PEX14—PEX5 PPI inhibitors, which resulted in more preferable selectivity indices. The high activity of the compounds against the human pathogen may have practical implications and merit further development of the PEX14—PEX5 PPI inhibitors as potential treatments against HAT.

Unlike T. brucei, T. cruzi parasites do not use glycolysis as the main energy source for the cell. However, they still use glycosomes to compartmentalize other essential metabolic processes. Hence, we wanted to check whether impairing the function of the organelle by PEX14-PEX5 PPI inhibition is also lethal to T. cruzi. When assayed for their trypanocidal activity against the amastigote forms of the T. cruzi Tulahuen strain C2C4 grown in rat L-6 myoblasts (Table 6), the tested PEX14-PEX5 PPI inhibitors demonstrated varied efficiency. The IC₅₀ values were higher than those obtained for *T. brucei* and thus the selectivity indices determined with uninfected L-6 cells were narrower. The aminoalkyl derivatives 50, 61, and 64 displayed highest potency, showing submicromolar trypanocidal activities and fair therapeutic indices. A general observation was that the correlation between the ability to break the PEX14-PEX5 PPI and trypanocidal efficacy is less clear in T. cruzi than in T. brucei. Presumably, this reflects the characteristics of the assay, in which the T. cruzi parasites are located within myoblasts and therefore the compounds have to pass multiple cellular membranes and environments to reach the target.

Chemistry. The synthetic pathway used to access the in silico hit 1 and its C-2 regioisomer 2 is shown in Scheme 1. The key pyrazolo[4,3-c]pyridine 66 was obtained by adopting a literature method⁴⁵ to a one-pot process, without isolating the intermediate mixed-Claisen condensation product. The subsequent N-methylation of the pyrazole ring of 66 resulted in two chromatographically separable regioisomers 67 and 68, in a 60:40 proportion. The 1,5,7-triazabicyclo[4.4.0]dec-5-ene

Scheme 1. Synthetic Route to the in Silico Hit 1 and Its C-2 Regioisomer 2^a

"Reagents and conditions: (a) LHMDS, THF, -78 °C, then (COOEt)₂, THF, -78 °C to rt; (b) hydrazine hydrate, THF, EtOH, AcOH, reflux; (c) Cs₂CO₃, DMF, rt, then MeI, DMF, rt; (d) benzylamine, TBD, THF, 60 °C; (e) 4 M HCl, 1,4-dioxane, rt; (f) TEA, THF, rt, then AcOH, indole-3-carboxaldehyde, NaBH(OAc)₃, THF, rt.

Scheme 2. Synthetic Routes Employed for the Initial Optimization Trials (Table 1)^a

"Reagents and conditions: (a) benzylamine, TBD, THF, 60 °C; (b) 4 M HCl, 1,4-dioxane, rt; (c) TEA, THF, rt, then AcOH, indole-3-carboxaldehyde, NaBH(OAc)₃, THF, rt; (d) Cs₂CO₃, DMF, rt, then ethyl 2-bromoacetate, DMF, rt (for 71) or ethyl chloroformate, DIPEA, 0 °C to rt (for 72); (e) KOH, EtOH/H₂O, rt, then NH₄Cl, EtOH/H₂O, rt; (f) LHMDS, THF, -78 °C, then (COOEt)₂, THF, -78 °C to rt; (g) 2-hydrazinoethanol, THF, EtOH, AcOH, reflux; (h) amine, TBD, THF, 60 °C; (i) TEA, THF, rt, then AcOH, aldehyde, NaBH(OAc)₃, THF, rt.

Scheme 3. Synthetic Routes Employed for the Optimization of the Residue Addressing the Phe Pocket (Table 2)

^aReagents and conditions: (a) 4 M HCl, 1,4-dioxane, rt; (b) TEA, THF, rt, then AcOH, aldehyde, NaBH(OAc)₃, THF, rt (for 14, 21–26, 28, and 30–45) or 7-methoxycarbonyl-1*H*-indole, HCHO_{aq}, ZnCl₂, EtOH, rt (for 27).

(TBD)-catalyzed aminolysis⁴⁶ of the respective esters **67** and **68**, followed by Boc-deprotection and reductive amination steps, gave the corresponding regioisomeric compounds **1** and **2**.

Highlighted in Scheme 2 are the synthetic routes employed for initial optimization trials. Compound 66 was used as a common intermediate to synthesize the derivatives having various substituents in the N-1 position of the pyrazolo[4,3-c]pyridine system. Aminolysis of 66 resulted in 70, which was subjected to Boc-deprotection and reductive amination steps

to give the N-1 unsubstituted derivative 3. Conversion of 70 to 71 and 72 by N-alkylation and N-acylation reactions, respectively, followed by the deprotection/reductive amination sequence, resulted in corresponding derivatives 6 and 7. Ester 6 was saponified to give the carboxylate 5. Attempts to obtain the hydroxyethyl derivatives 73 and 74 by the direct N-alkylation of 66 with 2-bromoethanol resulted in low conversions and afforded complex mixtures that were difficult to purify. Hence, we turned to a more practical route employing one pot mixed-Claisen condensation and pyrazole

Scheme 4. Synthetic Routes Employed for the Optimization around the Trp Pocket (Table 3)^a

"Reagents and conditions: (a) 1-naphthalenemethylamine, TBD, THF, 60 °C; (b) 4 M HCl, 1,4-dioxane, rt; (c) TEA, THF, rt, then AcOH, 4-methoxy-1-naphthaldehyde, NaBH(OAc)₃, THF, rt; (d) MsCl, DIPEA, DMAP, DCM, 0 °C to rt; (e) 'BuOK, THF, 0 °C to rt; (f) LiAlH₄, THF, rt, then reflux; (g) CDI, DCM, rt; (h) NH $_{3aq}$, EtOH, reflux; (i) TEA, THF, rt, then AcOH, 1-naphthalenecarboxaldehyde, NaBH(OAc)₃, THF, rt (for 48), or 1-naphthalenesulfonyl chloride, TEA, DMAP, DCM, 0 °C to rt (for 49).

Scheme 5. Synthetic Routes Employed for the Secondary Optimization of the Pyrazole N-Substituent (Table 4)^a

"Reagents and conditions: (a) MeNH₂, DCM/EtOH, reflux (for **51**), imidazole, Cs₂CO₃, DMF, 60 °C (for **55**), pyrazole, Cs₂CO₃, DMF, 60 °C (for **56**), morpholine, Cs₂CO₃, DMF, 60 °C (for **57**), methyl glycinate hydrochloride, K₂CO₃, MeCN, reflux (for **59**); (b) NaOH, MeOH/H₂O, rt; (c) acetone, AcOH, NaBH(OAc)₃, THF, rt (for **52**), HCHO_{aq}, AcOH, NaBH(OAc)₃, THF, rt (for **54**) or Ac₂O (neat), 80 °C (for **58**); (d) *tert*-butyl 3-((methylsulfonyl)oxy)azetidine-1-carboxylate, Cs₂CO₃, DMF, 90 °C; (e) 4 M HCl, 1,4-dioxane, rt, then K₂CO₃, rt.

formation that gave the mixture of regioisomers 73 and 74 in a 75:25 proportion. These regioisomers were readily separable by recrystallization and chromatography. Deprotection and reductive amination of intermediate 73 afforded compound 75 that served as a starting material for subsequent aminolyses,

leading to derivatives 4 and 8–12 bearing various phenyl substituents. Finally, reaction of ester 73 with benzylamine gave intermediate 76 that was used in the subsequent Bocdeprotection and reductive amination steps that led to

Scheme 6. Synthetic Routes Employed for the Optimization of the Central Scaffold (Table 5)^a

"Reagents and conditions: (a) MeI, CHCl₃, rt; (b) LDA, THF, -78 °C, then (COOEt)₂, THF, -78 °C to rt; (c) 2-hydroxyethylhydrazine, THF, EtOH, AcOH, reflux; (d) AcOH, reflux; (e) K₂CO₃, EtOH, 60 °C; (f) 1-naphthalenemetylamine, TBD, THF, 60 °C; (g) 1,4-cyclohexadiene, 10% Pd/C, MeOH, reflux; (h) HCl, Et₂O, DCM, 0 °C to rt; (i) 4-methoxy-1-naphthaldehyde, NaBH(OAc)₃, AcOH, THF, rt; (j) TsCl, TEA, DMAP, DCM, 0 °C to rt; (k) NaN₃, DMF, 70 °C; (l) PPh₃, THF, rt, then H₂O, THF, 50 °C; (m) 2-amino-2-methyl-1-propanol, K₂CO₃, CH₃CN, reflux.

derivatives 15-19 with different aromatic substituents addressing the Phe pocket of PEX14.

The synthetic route used for the secondary optimization trials of the residue addressing the Phe pocket is shown in Scheme 3. Briefly, ester 73 was converted to amide 77 by the previously described method.³⁵ Intermediate 77 was then Bocdeprotected and the resulting amine salt 78 was used to generate the desired derivatives 14, 21–28, and 30–45 either by reductive aminations or by the ZnCl₂-mediated Mannich reaction.⁴⁷

The synthesis of compounds 46–49, having modifications within the region projecting aromatic residues to the Trp pocket of PEX14, is depicted in Scheme 4. Intermediate 80 was obtained from compound 74 by an aminolysis/Bocdeprotection/reductive amination sequence. The hydroxyl group of 80 was then converted to mesylate and the subsequent treatment with a strong base induced the ring closure leading to compound 46. To obtain the sterically constrained derivative 47, intermediate 66 was first converted

to compound 82 in a manner similar to synthesis of compound 80 from 74. The amide 82 was then reduced with LiAlH₄ and the resulting amine 83 was treated with 1,1'-carbonyldiimidazole (CDI) to afford tricyclic compound 47. Derivatives 48 and 49 were accessed by reductive amination or acylation, respectively, of the common intermediate amine 86 obtained from compound 73 by a sequence comprising aminolysis/Bocdeprotection/reductive amination/LiAlH₄ reduction.

Scheme 5 shows the synthesis of pyrazolo[4,3-c]pyridine derivatives bearing various alkylamino substituents on the N-1 atom. Compounds 51, 55–57, and 59 were obtained from the tosylate 87 by the N-alkylation reactions of the appropriate aliphatic amines or heterocyclic bases. Glycinate ester 59 was saponified to give the corresponding carboxylic acid 60. Compounds 52, 54, and 58 were obtained by reductive amination or acetylation of the primary amine 50. The azetidine derivative 53 was synthesized from compound 82 by alkylation of the pyrazole nitrogen atom with *tert*-butyl 3-

((methylsulfonyl)oxy)azetidine-1-carboxylate, followed by removal of the Boc protecting group.

Synthetic routes to compounds with modifications within the central heterobicyclic core are shown in Scheme 6. The quaternary ammonium salt 62 was obtained by treatment of 29 with excess of MeI. To synthesize derivatives with pyrazolo-[4,3-c]pyrrole heterocycle as the central scaffold we attempted to apply the one-pot procedure used for construction of the homologous (2-hydroxyethyl)-1,4,6,7-tetrahydro-5H-pyrazolo-[4,3-c] pyridines 73 and 74. This resulted in formation of mixtures containing cyclized regioisomeric products and openchain hydrazones, as judged by liquid chromatography-mass spectrometry analyses. An additional step comprising prolonged refluxing in acetic acid was required to drive the cyclocondensation process to completion. Acetylation of the hydroxyl group in products 89 and 90 as well as partial lactonization of the N-2 regioisomer 89 to compound 90 was observed at the same time. Acetyl removal with K2CO3 in intermediate 89 afforded the free alcohol 92 that gave the final compounds 63, 64, and 65 in similar synthetic sequences to those used previously for synthesis of derivatives 29, 50, and $61.^{35}$

CONCLUSIONS

Glycosomal enzymes have long been considered attractive molecular targets for combating parasitic diseases, such as HAT and Chagas disease. Targeting the PEX14-PEX5 PPI leads to mislocalization of multiple glycosomal enzymes, which has a dual effect: the absence of the enzymes in the lumen of the glycosome compromises glycosome biogenesis and function, while the presence of the glycosomal kinases in the cytosol dissipates ATP levels. To verify that this is indeed a useful target for molecular intervention, we have used a structure-based drug design approach to develop the first small-molecule agents that inhibit the PEX14-PEX5 PPI. We have performed a systematic optimization of different sites in the initial in silico pyrazolo [4,3-c] pyridine hit 1 and found compounds that disrupt PEX14-PEX5 PPI, kill T. brucei and T. cruzi parasites in submicromolar concentrations, and display fair therapeutic windows between trypanocidal activity and cytotoxicity for the mammalian cells. Hence, the pyrazolo [4,3c]pyridine series of compounds represent an attractive lead series for treatment of HAT and Chagas disease and can also serve as tools for studying biochemical processes related to glycosomes and peroxisomes.

There have been successful medicinal chemistry campaigns that developed new PPI inhibitors as preclinical and clinical candidates. Still, this target class is considered difficult to be targeted by classical small molecules and successful drug development may require involvement of an alternative chemical matter that goes far beyond the established criteria of drug-likeness.³⁹ A good example of such compound is Venetoclax, a Bcl-2-Bak/Bax PPI inhibitor that displays violations of the classical drug-likeness principles (e.g., large MW, high log D, and poor aqueous solubility) but has quite recently been approved for treatment of chronic lymphocytic leukemia.48 The presented PEX14-PEX5 PPI inhibitors display suboptimal pharmacochemical properties³⁵ because the binding of the ligands to PEX14 hotspots is mostly lipophilicity-driven. However, we also show that polar interactions with charged amino-acids in the spatial proximity of the PPI hotspots can be addressed and illustrate this by providing detailed structure activity data. Importantly, our

results indicate that addressing specific water-mediated interactions is very important for design of compounds that bind the highly solvent-exposed PEX14 surface, and may be useful to improve the overall physicochemical profile of the obtained inhibitors, which is another subject of our ongoing efforts.

■ EXPERIMENTAL SECTION

Chemistry. Unless otherwise noted, all reagents were obtained from commercial sources and used without further purification. Technical grade solvents for chromatography and aqueous workup were distilled prior to use. Dry DCM, DMF, and THF were purchased from Acros. Manual flash column chromatography (FC) was performed using Merck silica gel 60 (particle size: 0.040-0.063 mm). Automated preparative chromatography was performed on a Grace Reveleris Prep purification system using linear gradient elution and Buchi Reveleris Silica 40 µm cartridges for normal phase (SiO₂) and Buchi Reveleris C-18 40 µm cartridges for reverse-phase (RP-C18) separations. Analytical thin layer chromatography (TLC) was performed on Merck silica coated plates (silica gel 60 F 254). Compounds were detected by ultraviolet (UV) irradiation at 254 or 366 nm. The final compounds were ≥95% pure, as determined by high-performance liquid chromatography-mass spectrometry analyses performed on a Dionex UltiMate 3000 HPLC system coupled with a Thermo Finnigan LCQ ultrafleet mass spectrometer, using the following methods: (A) Waters X-Bridge C18 (4.6 \times 30 mm, 3.5 μ m) column; gradient: 5-95% of acetonitrile +0.1% formic acid v/v in water +0.1% formic acid v/v over 5 min period; flow rate: 1.1 mL/ $\,$ min; UV detection at 214 and 280 nm; (B) Thermo Scientific Accucore aQ (2.1 \times 50 mm, 2.6 μ m) column; gradient: 5–95% of acetonitrile +0.1% formic acid v/v in water +0.1% formic acid v/v over 5 min period; flow rate: 0.9 mL/min; UV detection at 214 and 280 nm. High-resolution mass spectrometry (HRMS) measurements were performed on a Thermo Finnigan LTQ FT apparatus using an electrospray ionization (ESI) detector. NMR spectra were recorded on a Bruker AV250, Bruker AVHD300, Bruker AV360, Bruker AVHD400 or Bruker AV500C (equipped with a QNP CryoProbe) spectrometer. NMR peaks are reported as follows: chemical shift (δ) in parts per million (ppm) relative to residual nondeuterated solvent as an internal standard (CHCl₃: δ H = 7.26, δ C = 77.2; DMSO: δ H = 2.50, δ C = 39.5 ppm), multiplicity (s = singlet, d = doublet, t = triplet, q = quartet, dd = doublet of doublets, dt = doublet of triplets, m = multiplet and bs = broad signal), coupling constant (in Hz) and integration. Compounds 1, 13, 20, 29, 50, 61, 77, and 87 were synthesized as described previously.

5-((1H-Indol-3-yl)methyl)-N-benzyl-2-methyl-4,5,6,7-tetrahydro-2H-pyrazolo[4,3-c]pyridine-3-carboxamide (2). Intermediate 69 (100 mg, 0.27 mmol 1.0 equiv) was dissolved in a 4 M solution of HCl in 1,4-dioxane (0.5 mL). The mixture was stirred for 4 h at rt and concentrated in vacuo. The crude solid material was suspended in dry THF (1 mL) and TEA (27 mg, 0.27 mmol, 1.0 equiv) was added. The mixture was stirred for 0.5 h followed by addition of indole-3carboxaldehyde (39 mg, 0.27 mmol, 1.0 equiv), AcOH (16 mg, 0.27 mmol, 1.0 equiv), and NaBH(OAc)₃ (86 mg, 0.40 mmol, 1.5 equiv). After stirring for 12 h, the mixture was quenched with saturated aqueous solution of NaHCO₃ (2 mL) and extracted with EtOAc (2 × 2 mL). The combined organic extracts were washed with water (1 mL), brine (1 mL), dried over anhydrous Na₂SO₄, filtered, and concentrated in vacuo. The residue was purified by FC (EtOAc/ MeOH/TEA 99:1 to 94:5:1) to yield 65 mg (60%) of the title compound 2 as a pale yellow solid. ¹H NMR (500 MHz, CDCl₃): δ 8.30 (br s, 1H), 7.69 (d, J = 7.9 Hz, 1H), 7.37 (d, J = 8.2 Hz, 1H), 7.34-7.30 (m, 3H), 7.22 (ddd, J = 7.1, 3.1, 1.8 Hz, 2H), 7.13-7.07(m, 2H), 5.79 (t, J = 5.2 Hz, 1H), 4.53 (d, J = 5.7 Hz, 2H), 4.10 (s, 3H), 3.93 (s, 2H), 3.65 (s, 2H), 2.88 (t, J = 5.8 Hz, 2H), 2.79 (t, J = 5.7 Hz, 2H). 13 C NMR (126 MHz, CDCl₃): δ 160.2, 145.8, 137.7, 136.2, 130.9, 128.8, 127.8, 127.7, 127.6, 123.7, 122.2, 119.7, 119.2, 115.0, 112.3, 111.2, 52.4, 50.1, 49.5, 43.5, 39.1, 23.4. ESI HRMS (m/z): calcd for $C_{24}H_{26}N_5O$ [M + H]⁺, 400.21319; found, 400.21345.

5-((1H-Indol-3-yl)methyl)-N-benzyl-4,5,6,7-tetrahydro-1Hpyrazolo[4,3-c]pyridine-3-carboxamide (3). Compound 3 was synthesized by employing the procedure described for compound 2, using intermediate 70 (200 mg, 0.56 mmol, 1.0 equiv), 4 M solution of HCl in 1,4-dioxane (1 mL), TEA (56 mg, 0.56 mmol, 1.0 equiv), indole-3-carboxaldehyde (81 mg, 0.56 mmol, 1.0 equiv), AcOH (34 mg, 0.56 mmol, 1.0 equiv) and NaBH(OAc)₃ (119 mg, 0.84 mmol, 1.5 equiv), and dry THF (2 mL). Purification by FC (hexane/EtOAc 1:1 to 0:1, then AcOEt/TEA 99:1) gave 2.15 g (89%) of the title compound 3 as a white solid. ¹H NMR (500 MHz, DMSO- d_6): δ 12.86 (br s, 1H), 10.94 (br s, 1H), 8.52 (t, J = 6.5 Hz, 1H), 7.63 (d, J = 7.9 Hz, 1H), 7.35 (d, J = 8.1 Hz, 1H), 7.35–7.24 (m, 5H), 7.24– 7.16 (m, 1H), 7.06 (t, J = 7.5 Hz, 1H), 6.95 (t, J = 7.5 Hz, 1H), 4.35 (d, I = 6.5 Hz, 2H), 3.83 (s, 2H), 3.35 (s, 1H), 2.72 (t, I = 5.6 Hz, 2.22 (t, I = 5.6 (t, I = 5.6 (t, I = 5.6 (t, I = 5.6 (t, I = 5.62H), 2.66 (t, J = 5.6 Hz, 2H). ¹³C NMR (126 MHz, DMSO- d_6): δ 163.0 (br s), 141.5 (br s), 140.5 (br s), 139.0 (br s), 136.9, 128.6, 127.9, 127.7, 127.0, 124.9, 121.4, 119.6, 118.8, 115.9 (br s), 111.8, 111.8, 53.1, 49.7, 49.6, 42.1, 22.0 (br s). ESI HRMS (m/z): calcd for $C_{23}H_{24}N_5O [M + H]^+$, 386.19809; found, 386.19743.

5-((1H-Indol-3-yl)methyl)-N-benzyl-1-(2-hydroxyethyl)-4,5,6,7tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (4). Compound 4 was synthesized by employing the procedure described for compound 69, using intermediate 75 (120 mg, 0.40 mmol, 1.0 equiv), benzylamine (51 mg, 0.40 mmol, 1.2 equiv), TBD (15 mg, 0.11 mmol, 0.3 equiv), and dry THF (1 mL). Purification by FC (EtOAc/ MeOH/TEA 99:0:1 to 93:6:1) gave 90 mg (64%) of the title compound 4 as a white solid. ¹H NMR (500 MHz, DMSO- d_6): δ 10.95 (s, 1H), 8.44 (t, J = 6.0 Hz, 1H), 7.63 (d, J = 7.8 Hz, 1H), 7.36 (d, J = 8.1 Hz, 1H), 7.33-7.24 (m, 5H), 7.24-7.18 (m, 1H), 7.07 (t, 1H)J = 7.4 Hz, 1H), 6.97 (t, J = 7.4 Hz, 1H), 4.91 (t, J = 5.5 Hz, 1H), 4.36 (d, J = 6.3 Hz, 2H), 4.05 (t, J = 5.3 Hz, 2H), 3.84 (br s, 2H), 3.72 (d, J = 4.8 Hz, 2H), 3.58 (br s, 2H), 2.73 (br s, 4H). 13 C NMR (101 MHz, DMSO- d_6): δ 162.6, 140.50, 140.47, 139.9, 136.3, 128.6, 127.9, 127.8, 127.0, 125.0, 121.4, 119.6, 118.8, 116.5, 111.8, 60.6, 53.0, 52.0, 49.6, 49.5, 42.1, 22.2. ESI HRMS (m/z): calcd for $C_{25}H_{28}N_5O_2$ [M + H]⁺, 430.22375; found, 430.22358.

2-(5-((1H-Indol-3-yl)methyl)-3-(benzylcarbamoyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridin-1-yl)acetic Acid (5). Compound 6 (100 mg, 0.21 mmol, 1.0 equiv) was dissolved in EtOH (4 mL), followed by addition of aqueous solution of KOH (59 mg, 1.05 mmol, 5.0 equiv in 0.5 mL of water). The solution was stirred for 0.5 h at rt and saturated aqueous solution of NH₄Cl (0.5 mL) was added. The mixture was concentrated in vacuo and the residue was refluxed with water (3 mL). The resulting solid was filtered and dried in vacuo to give 63 mg (67%) of the title compound 5 as a white solid. ¹H NMR (360 MHz, DMSO- d_6): δ 11.16 (br s, 1H), 8.54 (t, J = 6.4 Hz, 1H), 7.69 (d, J = 7.9 Hz, 1H), 7.39 (d, J = 8.2 Hz, 2H), 7.33–7.15 (m, SH), 7.10 (t, J = 7.5 Hz, 1H), 7.01 (t, J = 7.4 Hz, 1H), 4.75 (s, 2H), 4.33 (d, J = 6.4 Hz, 2H), 4.17 (s, 2H), 3.85 (s, 2H), 3.03 (s, 2H), 2.73 (t, J = 5.8 Hz, 2H). ¹³C NMR (91 MHz, DMSO- d_6): δ 169.9, 162.3, 140.4, 140.3, 139.1 (br s), 136.7, 128.6, 128.0, 127.7, 127.0, 126.6 (br s), 121.7, 119.3, 114.7 (br s), 112.0, 108.4 (br s), 52.5 (br s), 51.6 (br s), 48.8 (br s), 48.4 (br s), 42.2, 20.9 (br s). ESI HRMS (m/z): calcd for C₂₅H₂₆N₅O₃ [M + H]⁺, 444.20302; found, 444.20284.

Ethyl 2-(5-((1H-Indol-3-yl)methyl)-3-(benzylcarbamoyl)-4,5,6,7tetrahydro-1H-pyrazolo[4,3-c]pyridin-1-yl)acetate (6). Compound 6 was synthesized by employing the procedure described for compound 2, using intermediate 71 (701 mg, 1.59 mmol, 1.0 equiv), 4 M solution of HCl in 1,4-dioxane (5 mL), TEA (161 mg, 1.59 mmol, 1.0 equiv), indole-3-carboxaldehyde (231 mg, 1.59 mmol, 1.0 equiv), AcOH (95 mg, 1.59 mmol, 1.0 equiv), NaBH(OAc)₃ (506 mg, 2.39 mmol, 1.5 equiv), and dry THF (10 mL). Purification by FC (EtOAc/MeOH/TEA 99:0:1 to 94:5:1) gave 533 mg (71%) of the title compound 6 as a pale yellow solid. ¹H NMR (360 MHz, CDCl₃): δ 8.48 (br s, 1H), 7.76 (d, J = 7.8 Hz, 1H), 7.40–7.29 (m, 5H), 7.22– 7.16 (m, 1H), 7.16-7.08 (m, 4H), 4.73 (s, 2H), 4.59 (d, J = 6.0 Hz,2H), 4.22 (q, J = 7.1 Hz, 2H), 3.98 (s, 2H), 3.96 (s, 2H), 2.86 (t, J =5.7 Hz, 2H), 2.61 (t, J = 5.6 Hz, 2H), 1.27 (t, J = 7.1 Hz, 3H). ¹³C NMR (91 MHz, CDCl₃): δ 167.2, 162.4, 141.4, 140.0, 138.4, 136.4, 128.6, 127.9, 127.9, 127.3, 123.9, 121.9, 119.5, 119.4, 118.1, 112.6,

111.1, 62.0, 52.4, 50.8, 49.6, 48.5, 42.9, 22.1, 14.1. ESI HRMS (m/z): calcd for $C_{27}H_{30}N_5O_3$ [M + H] $^+$, 472.23432; found, 472.23447.

Ethyl 5-((1H-Indol-3-yl)methyl)-3-(benzylcarbamoyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-1-carboxylate (7). Compound 7 was synthesized by employing the procedure described for compound 2, using intermediate 72 (545 mg, 1.27 mmol, 1.0 equiv), 4 M solution of HCl in 1,4-dioxane (5 mL), TEA (128 mg, 1.27 mmol, 1.0 equiv), indole-3-carboxaldehyde (184 mg, 1.27 mmol, 1.0 equiv), AcOH (76 mg, 1.27 mmol, 1.0 equiv), NaBH(OAc)₃ (404 mg, 1.91 mmol, 1.5 equiv), and dry THF (10 mL). Purification by FC (EtOAc/MeOH 99:1 to 95:5) gave 510 mg (88%) of the title compound 7 as a white solid. ^{1}H NMR (250 MHz, CDCl₃): δ 8.41 (br s, 1H), 7.81-7.66 (m, 1H), 7.42-7.27 (m, 7H), 7.24-7.07 (m, 3H), 4.57 (d, J = 6.1 Hz, 2H), 4.47 (q, J = 7.1 Hz, 2H), 4.05-3.86(m, 4H), 3.02 (t, J = 5.8 Hz, 2H), 2.85 (t, J = 5.7 Hz, 2H), 1.42 (t, J =7.1 Hz, 3H). 13 C NMR (63 MHz, CDCl₃): δ 161.5, 149.6, 145.2, 143.0, 137.9, 136.3, 128.7, 127.9, 127.8, 127.5, 124.1, 122.0, 120.4, 119.6, 119.4, 111.8, 111.1, 64.7, 52.2, 49.0, 48.6, 43.0, 25.5, 14.2. ESI HRMS (m/z): calcd for $C_{27}H_{30}N_5O_3$ [M + H]⁺, 458.21867; found, 458.21897.

5-((1H-Indol-3-vl)methyl)-N-(4-chlorobenzyl)-1-(2-hydroxyethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (8). Compound 8 was synthesized by employing the procedure described for compound 69, using intermediate 75 (120 mg, 0.33 mmol, 1.0 equiv), 4-chlorobenzylamine (57 mg, 0.40 mmol, 1.2 equiv), TBD (15 mg, 0.11 mmol, 0.3 equiv), and dry THF (1 mL). Purification by FC (EtOAc/MeOH/TEA 99:0:1 to 93:6:1) gave 63 mg (41%) of the title compound 8 as a white solid. ¹H NMR (360 MHz, DMSO- d_6): δ 10.92 (s, 1H), 8.50 (t, J = 6.3 Hz, 1H), 7.62 (d, J= 7.9 Hz, 1H), 7.39-7.30 (m, 3H), 7.30-7.22 (m, 3H), 7.06 (t, I = 1.00 (m, 3H)7.5 Hz, 1H), 6.95 (t, J = 7.5 Hz, 1H), 4.90 (br s, 1H), 4.32 (d, J = 6.3Hz, 2H), 4.04 (t, J = 5.4 Hz, 2H), 3.81 (s, 2H), 3.72 (t, J = 5.1 Hz, 2H), 3.55 (s, 2H), 2.71 (br s, 4H). 13 C NMR (91 MHz, DMSO- d_6): δ 162.2, 139.9, 139.5, 139.1, 136.4, 131.1, 129.1 (2× C), 128.1 (2× C), 127.4, 124.4, 120.9, 119.1, 118.3, 116.2, 111.3 (2× C), 60.1, 52.5, 51.5, 49.2, 49.1, 41.0, 21.8. ESI HRMS (m/z): calcd for $C_{25}H_{27}ClN_5O_2$ [M + H]⁺, 464.18478; found, 464.18476.

5-((1H-Indol-3-yl)methyl)-N-(4-fluorobenzyl)-1-(2-hydroxyethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (**9**). Compound 9 was synthesized by employing the procedure described for compound 69, using intermediate 75 (120 mg, 0.33 mmol, 1.0 equiv), 4-fluorobenzylamine (50 mg, 0.40 mmol, 1.2 equiv), TBD (15 mg, 0.11 mmol, 0.3 equiv), and dry THF (1 mL). Purification by FC (EtOAc/MeOH/TEA 99:0:1 to 93:6:1) gave 60 mg (41%) of the title compound 9 as a white solid. ¹H NMR (500 MHz, CDCl₃): δ 10.91 (s, 1H), 8.45 (t, J = 6.2 Hz, 1H), 7.62 (d, J = 7.7 Hz, 1H), 7.35 (d, J =7.7 Hz, 1H), 7.30 (t, J = 7.1 Hz, 2H), 7.25 (s, 1H), 7.10 (t, J = 8.8 Hz, 2H), 7.05 (t, J = 8.0 Hz, 1H), 6.95 (t, J = 7.5 Hz, 1H), 4.89 (br s, 1H), 4.32 (d, J = 5.6 Hz, 2H), 4.04 (t, J = 5.0 Hz, 2H), 3.82 (s, 2H), 3.71 (d, J = 4.7 Hz, 2H), 3.71 (d, J = 4.7 Hz, 2H), 3.55 (s, 2H), 2.71(br s, 4H). 13 C NMR (91 MHz, CDCl₃): δ 162.1, 161.0 (d, $^{1}J_{CF}$ = 240.0 Hz), 139.0, 139.5, 136.4, 136.2 (d, $J_{CF} = 3.0 \text{ Hz}$), 129.3, 129.2, 127.4, 124.4, 120.9, 119.1, 118.3, 116.2, 114.9, 114.7, 111.3 (2× C), 60.1, 52.5, 51.5, 49.2, 49.1, 41.0, 21.8. ESI HRMS (m/z): calcd for $C_{25}H_{27}FN_5O_2$ [M + H]⁺, 448.21433; found, 448.21425.

5-((1H-Indol-3-yl)methyl)-N-(3-chlorobenzyl)-1-(2-hydroxyethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (10). Compound 10 was synthesized by employing the procedure described for compound 69, using intermediate 75 (120 mg, 0.33 mmol, 1.0 equiv), 3-chlorobenzylamine (57 mg, 0.40 mmol, 1.2 equiv), TBD (15 mg, 0.11 mmol, 0.3 equiv), and dry THF (1 mL). Purification by FC (EtOAc/MeOH/TEA 99:0:1 to 93:6:1) gave 92 mg (60%) of the title compound 10 as a white solid. ¹H NMR (500 MHz, DMSO- d_6): δ 10.91 (s, 1H), 8.53 (t, J = 6.4 Hz, 1H), 7.62 (d, J = 7.8 Hz, 1H), 7.37-7.19 (m, 6H), 7.06 (t, J = 7.5 Hz, 1H), 6.95 (t, J = 7.5 Hz, 1H), 4.89 (br s, 1H), 4.34 (d, J = 6.4 Hz, 2H), 4.05 (t, J = 5.5 Hz, 2H), 3.82 (s, 2H), 3.72 (br s, 2H), 3.55 (s, 2H), 2.71 (br s, 4H). ¹³C NMR (91 MHz, DMSO- d_6): δ 162.2, 142.7, 139.8, 139.6, 136.4, 132.8, 130.1, 127.1, 126.5, 126.0, 124.4, 120.9, 119.1, 118.3, 116.2, 111.3 (2× C), 60.1, 52.5, 51.5, 49.2, 49.1, 41.2, 21.8. ESI

HRMS (m/z): calcd for $C_{25}H_{27}ClN_5O_2$ [M + H]⁺, 464.18478; found, 464.18481.

5-((1H-Indol-3-yl)methyl)-N-(3-methylbenzyl)-1-(2-hydroxyethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (11). Compound 11 was synthesized by employing the procedure described for compound 69, using intermediate 75 (120 mg, 0.33 mmol, 1.0 equiv), 3-methylbenzylamine (48 mg, 0.40 mmol, 1.2 equiv), TBD (15 mg, 0.11 mmol, 0.3 equiv), and dry THF (1 mL). Purification by FC (EtOAc/MeOH/TEA 99:0:1 to 93:6:1) gave 76 mg (52%) of the title compound 11 as a white solid. ¹H NMR (500 MHz, DMSO- d_6): δ 10.92 (s, 1H), 8.36 (t, J = 6.2 Hz, 1H), 7.62 (d, J= 7.8 Hz, 1H), 7.35 (d, J = 8.1 Hz, 1H), 7.25 (s, 1H), 7.17 (t, J = 7.5Hz, 1H), 7.10-6.99 (m, 4H), 6.95 (t, J = 7.1 Hz, 1H), 4.88 (t, J = 5.5Hz, 1H), 4.31 (d, J = 6.3 Hz, 2H), 4.04 (t, J = 5.5 Hz, 2H), 3.83 (br s, 2H), 3.71 (q, J = 5.5 Hz, 2H), 3.50 (br s, 2H), 2.71 (br s, 4H), 2.26 (s, 3H). ¹³C NMR (91 MHz, CDCl₃): δ 162.5, 141.0, 139.7, 138.5, 136.4, 128.7, 128.7, 128.3, 128.0, 125.0, 124.4, 122.1, 119.7, 119.4, 117.4, 111.9, 111.4 (2× C), 61.4, 52.5, 51.2, 49.8, 48.9, 43.0, 22.4, 21.5. ESI HRMS (m/z): calcd for $C_{26}H_{30}N_5O_2$ [M + H]⁺, 444.23940; found, 444.23924.

5-((1H-Indol-3-vl)methyl)-N-(3-trifluoroethylbenzyl)-1-(2-hydroxyethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (12). Compound 12 was synthesized by employing the procedure described for compound 69, using intermediate 75 (120 mg, 0.33 mmol, 1.0 equiv), 3-methylbenzylamine (70 mg, 0.40 mmol, 1.2 equiv), TBD (15 mg, 0.11 mmol, 0.3 equiv), and dry THF (1 mL). Purification by FC (EtOAc/MeOH/TEA 99:0:1 to 93:6:1) gave 68 mg (41%) of the title compound 12 as a white solid. ¹H NMR (500 MHz, DMSO- d_6): δ 10.91 (s, 1H), 8.61 (t, J = 6.3 Hz, 1H), 7.67-7.47 (m, 5H), 7.35 (d, J = 8.1 Hz, 1H), 7.25 (d, 2.0 Hz, 1H), 7.06 (t, J = 7.3 Hz, 1H), 6.95 (t, J = 7.3 Hz, 1H), 4.89 (t, J = 9.2 Hz, 1H), 4.42 (d, J = 6.2 Hz, 2H), 4.05 (t, J = 5.4 Hz, 2H), 3.81 (s, 2H), 3.72 (d, J = 4.9 Hz, 2H), 3.55 (s, 2H), 2.71 (br s, 4H). 13 C NMR (91) MHz, DMSO-d₆): δ 162.3, 141.6, 139.8, 139.6, 136.4, 131.5, 129.2, 128.9 (d, J_{CF} = 31.6 Hz), 128.3 (q, J_{CF} = 151 Hz), 127.4, 124.4, 123.8 $(d, J_{CF} = 3.8 \text{ Hz}), 123.3 (d, J_{CF} = 3.3 \text{ Hz}), 120.9, 119.1, 118.3, 116.2,$ 111.3 (2× C), 60.1, 52.5, 51.5, 49.2, 49.1, 41.3, 21.8. ESI HRMS (*m*/ z): calcd for $C_{26}H_{27}F_3N_5O_2$ [M + H]⁺, 498.21114; found, 498.21120.

5-((1H-Indol-3-yl)methyl)-1-(2-hydroxyethyl)-N-(naphthalen-1ylmethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (14). Compound 78 (120 mg, 0.31 mmol, 1.0 equiv), was suspended in dry THF (5 mL), followed by addition of TEA (31 mg, 0.31 mmol, 1.0 equiv). The mixture was stirred for 0.5 h and indole-3carboxaldehyde (45 mg, 0.31 mmol, 1.0 equiv), AcOH (19 mg, 0.31 mmol, 1.0 equiv), and NaBH(OAc)₃ (100 mg, 0.46 mmol, 1.5 equiv) were added. After stirring for 12 h, the mixture was quenched with saturated aqueous solution of NaHCO3 (5 mL) and extracted with EtOAc (2×5 mL). The combined organic extracts were washed with water (5 mL), brine (5 mL), dried over anhydrous Na2SO4, filtered, and concentrated in vacuo. The residue was purified by automated preparative chromatography (RP-C18, linear gradient from water/ MeOH 90:10 to 15:85) to yield 81 mg (54%) of the title compound 14 as a pale yellow solid. ¹H NMR (400 MHz, CDCl₃): δ 8.82 (br s, 1H), 8.07-8.03 (m, 1H), 7.86-7.82 (m, 1H), 7.78 (d, J = 8.1 Hz, 1H), 7.70 (d, J = 7.8 Hz, 1H), 7.50–7.44 (m, 2H), 7.44 (dd, J = 6.9Hz, J = 1.5 Hz, 1H), 7.41-7.35 (m, 1H), 7.28 (d, J = 7.9 Hz, 1H), 7.19 (t, J = 5.8 Hz, 1H), 7.16 - 7.07 (m, 2H), 7.03 (d, J = 4.6 Hz, 1H), $5.02 \text{ (d, } J = 5.6 \text{ Hz, } 2\text{H}), 3.92 \text{ (s, } 2\text{H}), 3.89 \text{ (s, } 2\text{H}), 3.80 \text{ (t, } J = 4.9 \text{ (s, } 2\text{H}), 3.80 \text{ (t,$ Hz, 2H), 3.68 (t, J = 4.9 Hz, 2H), 2.75 (t, J = 5.9 Hz, 2H), 2.61 (t, J =5.9 Hz). 13 C NMR (101 MHz, CDCl₃): δ 162.4, 140.8, 139.8, 136.4, 140.0, 133.8, 131.6, 128.8, 128.6, 128.0, 126.8, 126.7, 126.0, 125.5, 124.3, 123.6, 122.0, 119.6, 119.3, 117.5, 111.6, 111.4, 61.2, 52.4, 51.1, 49.8, 48.7, 41.0, 22.4. ESI HRMS (m/z): calcd for $C_{29}H_{30}N_5O_2$ [M + H]+, 480.23940; found, 480.23940.

N,5-Dibenzyl-1-(2-hydroxyethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (15). The crude amine hydrochloride liberated from intermediate 76 (305 mg, 0.91 mmol, 1.0 equiv) was suspended in dry THF (10 mL), followed by addition of TEA (92 mg, 0.91 mmol, 1.0 equiv). The mixture was stirred for 0.5 h and benzaldehyde (96 mg, 0.91 mmol, 1.0 equiv), AcOH (55

mg, 0.91 mmol, 1.0 equiv), and NaBH(OAc)₃ (288 mg, 1.36 mmol, 1.5 equiv) were added. After stirring for 12 h, the mixture was quenched with saturated aqueous solution of NaHCO₃ (10 mL) and extracted with EtOAc (2 × 10 mL). The combined organic extracts were washed with water (5 mL), brine (5 mL), dried over anhydrous Na₂SO₄, filtered, and concentrated in vacuo. The residue was purified by FC (EtOAc/MeOH/TEA 99:1 to 94:5:1) to yield 184 mg (52%) of the title compound 15 as a pale yellow solid. ¹H NMR (400 MHz, CDCl₃): δ 7.43–7.30 (m, 8H), 7.30–7.23 (m, 2H), 7.20 (t, J = 6.0, 1H), 4.56 (d, J = 6.1 Hz, 1H), 4.01 (t, J = 4.8, 1H), 3.92–3.84 (m, 4H), 3.76 (s, 2H), 3,20 (br s, 1H) 2.79–2.67 (m, 4H); ¹³C NMR (101 MHz, CDCl₃): δ 162.4, 140.8, 139.5, 138.5, 137.9, 129.2, 128.6, 128.4, 127.8, 127.4, 127.3, 117.2, 61.9, 61.2, 51.1, 49.9, 48.7, 42.8, 22.2. ESI HRMS (m/z): calcd for C₂₃H₂₇N₄O₂ [M + H]⁺, 391.21285; found, 391.21275.

N-Benzyl-5-(3-chlorobenzyl)-1-(2-hydroxyethyl)-4.5.6.7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (16). Compound 16 was synthesized by employing the procedure described for compound 15, using the crude amine hydrochloride liberated from intermediate 76 (105 mg, 0.31 mmol, 1.0 equiv), TEA (31 mg, 0.31 mmol, 1.0 equiv), 3-chlorobenzaldehyde (44 mg, 0.31 mmol, 1.0 equiv), AcOH (19 mg, 0.31 mmol, 1.0 equiv), NaBH(OAc)₃ (98 mg, 0.46 mmol, 1.5 equiv), and dry THF (2 mL). Purification by FC (EtOAc/MeOH/ TEA 99:0:1 to 95:4:1) gave 89 mg (68%) of the title compound 16 as a white solid. ¹H NMR (500 MHz, CDCl₃): δ 7.38 (br s, 1H), 7.32 (d, J = 4.4 Hz, 4H), 7.28-7.25 (m, 1H), 7.24 (br s, 3H), 7.11 (t, J = 4.4 Hz, 4Hz)5.8 Hz, 1H), 4.55 (d, I = 5.8 Hz, 2H), 4.04 (t, I = 5.6 Hz, 2H), 3.93 (t, J = 5.6 Hz, 2H), 3.83 (s, 2H), 3.70 (s, 2H), 2.72 (d, J = 5.0 Hz,2H), 2.70 (d, J = 5.0 Hz, 2H). ¹³C NMR (91 MHz, CDCl₃): δ 162.5, 141.0, 140.7, 139.6, 138.6, 134.4, 129.8, 129.0, 128.8 (2× C), 128.0 (2× C), 127.5, 127.5, 127.2, 117.4, 61.5, 61.4, 51.1, 50.0, 49.0, 43.0, 22.4. ESI HRMS (m/z): calcd for $C_{23}H_{26}ClN_4O_2$ [M + H]⁺, 425.17388; found, 425.17420.

N-Benzyl-5-(4-chlorobenzyl)-1-(2-hydroxyethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (17). Compound 17 was synthesized by employing the procedure described for compound 15, using the crude amine hydrochloride liberated from intermediate 76 (105 mg, 0.31 mmol, 1.0 equiv), TEA (31 mg, 0.31 mmol, 1.0 equiv), 4-chlorobenzaldehyde (44 mg, 0.31 mmol, 1.0 equiv), AcOH (19 mg, 0.31 mmol, 1.0 equiv), NaBH(OAc)₃ (98 mg, 0.46 mmol, 1.5 equiv), and dry THF (5 mL). Purification by FC (EtOAc/MeOH/ TEA 99:0:1 to 95:4:1) gave 110 mg (84%) of the title compound 17 as a white solid. ¹H NMR (500 MHz, CDCl₃): δ 7.31 (d, J = 4.4 Hz, 4H), 7.29 (s, 4H), 7.27–7.24 (m, 1H), 7.13 (t, J = 6.1 Hz, 1H), 4.55 (d, J = 6.1 Hz, 2H), 4.02 (t, J = 4.9 Hz, 2H), 3.91 (t, J = 4.9 Hz, 2H),3.81 (s, 2H), 3.70 (s, 2H), 3.35 (br s, 1H), 2.72 (d, J = 5.2 Hz, 2H), 2.69 (d, I = 5.2 Hz, 2H). ¹³C NMR (91 MHz, CDCl₃): δ 162.5, 141.0, 139.6, 138.6, 136.7, 133.1, 130.5 (2× C), 128.8 (2× C), 128.7 (2× C), 128.0 (2× C), 127.5, 117.3, 61.4, 61.2, 51.1, 49.9, 48.9, 43.0, 22.4. ESI HRMS (m/z): calcd for $C_{23}H_{26}CIN_4O_2$ [M + H]⁺, 425.17388; found, 425.17406.

N-Benzyl-5-(4-methoxybenzyl)-1-(2-hydroxyethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (18). Compound 18 was synthesized by employing the procedure described for compound 15, using the crude amine hydrochloride liberated from intermediate 76 (105 mg, 0.31 mmol, 1.0 equiv), TEA (31 mg, 0.31 mmol, 1.0 equiv), 4-methoxybenzaldehyde (42 mg, 0.31 mmol, 1.0 equiv), AcOH (19 mg, 0.31 mmol, 1.0 equiv), NaBH(OAc)₃ (98 mg, 0.46 mmol, 1.5 equiv), and dry THF (5 mL). Purification by FC (EtOAc/MeOH/TEA 99:0:1 to 95:4:1) gave 107 mg (82%) of the title compound 18 as a white solid. ¹H NMR (500 MHz, CDCl₃): δ 7.31 (d, J = 4.3 Hz, 4H), 7.27 (d, J = 6.7 Hz, 3H), 7.13 (t, J = 5.8 Hz, 1H), 6.85 (d, J = 8.4 Hz, 2H), 4.54 (d, J = 5.8 Hz, 2H), 4.00 (t, J = 4.7Hz, 2H), 3.89 (t, J = 5.0 Hz, 2H), 3.81 (s, 2H), 3.79 (s, 3H), 3.68 (s, 2H), 3.09 (br s, 1H), 2.71 (d, J = 5.0 Hz, 2H), 2.67 (d, J = 5.0 Hz, 2H). 13 C NMR (91 MHz, CDCl₃): δ 162.5, 159.0, 141.0, 139.7, 138.6, 130.5 (2× C), 130.1, 128.8 (2× C), 127.9 (2× C), 127.5 (2× C), 117.5, 113.9 (2× C), 61.4, 55.4, 51.1, 49.8, 48.7, 42.9, 22.4. ESI HRMS (m/z): calcd for $C_{24}H_{29}N_4O_3$ [M + H]⁺, 421.22342; found, 421.22375.

N-Benzyl-1-(2-hydroxyethyl)-5-(naphthalen-1-ylmethyl)-4,5,6,7tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (19). Compound 19 was synthesized by employing the procedure described for compound 15, using the crude amine hydrochloride liberated from intermediate 76 (52 mg, 0.16 mmol, 1.0 equiv), TEA (16 mg, 0.16 mmol, 1.0 equiv), 1-naphthaldehyde (25 mg, 0.16 mmol, 1.0 equiv), AcOH (9 mg, 0.16 mmol, 1.0 equiv), NaBH(OAc)₃ (51 mg, 0.24 mmol, 1.5 equiv), and dry THF (2 mL). Purification by FC (EtOAc/ MeOH/TEA 99:0:1 to 95:4:1) gave 32 mg (45%) of the title compound 19 as a pale yellow solid. 1 H NMR (500 MHz, CDCl₃): δ 8.28 (dd, I = 6.7 Hz, 2.7 Hz, 1H), 7.84 (dd, I = 6.3 Hz, 3.1 Hz, 1H), 7.79 (d, J = 8.1 Hz, 1H), 7.49 - 7.46 (m, 3H), 7.42 (dd, J = 8.1 Hz, J = 8.6.9 Hz, 1H), 7.32 (d, J = 4.4 Hz, 4H), 7.28 - 7.24 (m, 1H), 7.14 (t, J = 4.4 Hz) 6.1 Hz, 1H), 4.56 (d, I = 6.1 Hz, 2H), 4.14 (s, 2H), 4.00 (t, I = 5.6Hz, 2H), 3.96 (s, 2H), 3.90 (t, J = 5.6 Hz, 2H), 2.78 (t, J = 5.6 Hz, 2H), 2.61 (t, J = 5.6 Hz, 2H), 2.02 (s, 1H). ¹³C NMR (91 MHz. CDCl₃): δ 162.5, 141.0, 139.9, 138.6, 134.1, 134.0, 132.8, 128.8 (2× C), 128.5, 128.3, 128.0 (2 \times C), 127.7, 127.5, 126.0, 125.8, 125.3, 124.8, 117.6, 61.4, 60.1, 51.0, 50.4, 48.8, 43.0, 22.4. ESI HRMS (m/

z): calcd for $C_{27}H_{29}N_4O_2~[M+H]^+$, 441.22850; found, 441.22828. 1-(2-Hydroxyethyl)-5-((1-methyl-1H-indol-3-yl)methyl)-N-(naphthalen-1-ylmethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (21). Compound 21 was synthesized by employing the procedure described for compound 14, using intermediate 78 (60 mg, 0.16 mmol, 1.0 equiv), TEA (16 mg, 0.16 mmol, 1.0 equiv), 1methyl-1H-indole-3-carbaldehyde (24 mg, 0.16 mmol, 1.0 equiv), AcOH (10 mg, 0.16 mmol, 1.0 equiv), NaBH(OAc)₃ (50 mg, 0.23 mmol, 1.5 equiv), and dry THF (2 mL). Purification by automated preparative chromatography (RP-C18, linear gradient from water/ MeOH 90:10 to 15:85) gave 60 mg (76%) of the title compound 21 as a pale yellow solid. ¹H NMR (500 MHz, CDCl₃): δ 8.07 (d, J = 7.9Hz, 1H), 7.85 (dd, J = 7.0 Hz, 1.7 Hz, 1H), 7.79 (d, J = 8.1 Hz, 1H), 7.72 (d, I = 7.9 Hz, 1H), 7.55–7.47 (m, 2H), 7.46 (d, I = 6.8 Hz, 1H), 7.40 (t, J = 6.8 Hz, 1H), 7.30 (d, J = 8.1 Hz, 1H), 7.23 (t, J = 7.6Hz, 1H), 7.15-7.09 (m, 2H), 7.08 (s, 1H), 4.99 (d, J = 5.7 Hz, 2H), 3.96 (s, 2H), 3.94 (s, 2H), 3.88 (t, J = 5.3 Hz, 2H), 3.78-3.73 (m, 5H), 2.79 (t, J = 5.6 Hz, 2H), 2.61 (t, J = 5.6 Hz, 2H). ¹³C NMR (75 MHz, CDCl₃): δ 162.3, 140.9, 139.7, 137.2, 133.9, 133.8, 131.6, 128.9, 128.8, 128.6, 128.5, 126.8, 126.7, 126.0, 125.5, 123.8, 121.7, 119.6, 119.2, 117.6, 110.7, 109.3, 61.3, 52.4, 51.0, 49.8, 48.8, 41.0, 32.3, 22.4. ESI HRMS (m/z): calcd for $C_{30}H_{32}N_5O_2$ $[M + H]^+$, 494.25505; found, 494.25478.

1-(2-Hydroxyethyl)-5-((1-ethyl-1H-indol-3-yl)methyl)-N-(naphthalen-1-ylmethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (22). Compound 22 was synthesized by employing the procedure described for compound 14, using intermediate 78 (60 mg, 0.16 mmol, 1.0 equiv), TEA (16 mg, 0.16 mmol, 1.0 equiv), 1ethyl-1H-indole-3-carbaldehyde (26 mg, 0.16 mmol, 1.0 equiv), AcOH (10 mg, 0.16 mmol, 1.0 equiv), NaBH(OAc)₃ (50 mg, 0.23 mmol, 1.5 equiv), and dry THF (2 mL). Purification by automated preparative chromatography (RP-C18, linear gradient from water/ MeOH 90:10 to 15:85) gave 25 mg (31%) of the title compound 22 as a pale yellow solid. ¹H NMR (400 MHz, CDCl₃): δ 8.07 (dd, I =8.0 Hz, 1.3 Hz, 1H), 7.85 (dd, J = 7.2 Hz, 2.2 Hz, 1H), 7.79 (d, J =8.0 Hz, 1H), 7.71 (d, J = 8.0 Hz, 1H), 7.55–7.43 (m, 4H), 7.41 (d, J= 8.0 Hz, 1H), 7.34 (d, I = 8.2 Hz, 1H), 7.24 - 7.18 (m, 2H), 7.15 -7.10 (m, 2H), 4.99 (d, J = 5.7 Hz, 2H), 4.15 (q, J = 7.3 Hz, 2H), 4.04 (m, 2H)(s, 2H), 4.02 (s, 2H), 3.90 (t, J = 4.4 Hz, 2H), 3.78 (t, J = 5.4 Hz, 2H)3H), 2.86 (t, J = 5.7 Hz, 2H), 2.69 (t, J = 5.7 Hz, 2H), 1.46 (t, J = 7.3Hz, 3H). 13 C NMR (101 MHz, CDCl₃): δ 162.2, 141.0, 139.4, 136.2, 134.0, 133.8, 131.6, 128.8, 128.7, 128.6, 127.7, 126.8, 126.7, 126.0, 125.5, 123.8, 121.7, 119.5, 116.6, 109.6, 109.3, 61.3, 53.6, 52.1, 51.4, 49.4, 48.4, 41.1, 22.0, 15.6. ESI HRMS (m/z): calcd for $C_{31}H_{34}N_5O_2$ [M + H]⁺, 508.27070; found, 508.27076.

1-(2-Hydroxyethyl)-5-((1-acetyl-1H-indol-3-yl)methyl)-N-(naphthalen-1-ylmethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (23). Compound 23 was synthesized by employing the procedure described for compound 14, using intermediate 78 (60 mg, 0.16 mmol, 1.0 equiv), TEA (16 mg, 0.16 mmol, 1.0 equiv), 1-acetyl-1H-indole-3-carbaldehyde (29 mg, 0.16 mmol, 1.0 equiv),

AcOH (10 mg, 0.16 mmol, 1.0 equiv), NaBH(OAc)₃ (50 mg, 0.23 mmol, 1.5 equiv), and dry THF (2 mL). Purification by FC (EtOAc/ MeOH/TEA 99:0:1 to 95:4:1) gave 53 mg (63%) of the title compound 23 as a pale yellow solid. ¹H NMR (300 MHz, DMSO d_{δ}): δ 8.43 (t, I = 6.0 Hz, 1H), 8.32 (d, I = 8.1 Hz, 1H), 8.24–8.09 (m, 1H), 7.93 (dd, J = 6.4 Hz, J = 3.1 Hz, 1H), 7.87–7.76 (m, 2H), 7.75 (d, J = 7.4 Hz, 1H), 7.58-7.49 (m, 2H), 7.48-7.40 (m, 2H), 7.32 (t, J = 7.1 Hz, 1H), 7.24 (t, J = 7.0 Hz, 1H), 4.89 (t, J = 5.4 Hz, 1H), 4.83 (d, J = 6.0 Hz, 2H), 4.04 (t, J = 5.4 Hz, 2H), 3.83 (s, 2H), 3.71 (q, J = 5.4 Hz, 2H), 3.65 (s, 2H), 2.75 (br s, 4H), 2.64 (s, 3H).¹³C NMR (75 MHz, DMSO-*d*₆): δ 169.3, 162.1, 140.0, 139.5, 135.4, 134.9, 133.2, 130.8, 130.2, 128.5, 127.3, 126.1, 125.7, 125.4 (2× C), 125.3, 124.7, 123.5, 123.1, 120.1, 118.7, 115.9 (2× C), 60.1, 52.0, 51.5, 49.6, 49.2, 23.9, 21.8. 1 C-atom overlaps with DMSO-d₆ signal. ESI HRMS (m/z): calcd for $C_{31}H_{32}N_5O_3$ [M + H]⁺, 522.24997; found, 522.24997.

1-(2-Hydroxyethyl)-5-((1-methanesulfonyl-1H-indol-3-yl)methyl)-N-(naphthalen-1-ylmethyl)-4,5,6,7-tetrahydro-1Hpyrazolo[4,3-c]pyridine-3-carboxamide (24). Compound 24 was synthesized by employing the procedure described for compound 14, using intermediate 78 (60 mg, 0.16 mmol, 1.0 equiv), TEA (16 mg, 0.16 mmol, 1.0 equiv), 1-methanesulfonyl-1H-indole-3-carbaldehyde (32 mg, 0.16 mmol, 1.0 equiv), AcOH (10 mg, 0.16 mmol, 1.0 equiv), NaBH(OAc)₃ (50 mg, 0.23 mmol, 1.5 equiv), and dry THF (2 mL). Purification by FC (EtOAc/MeOH 99:1 to 95:5) gave 48 mg (41%) of the title compound 24 as a pale yellow solid. ¹H NMR (400 MHz, CDCl₃): δ 8.07 (dd, J = 7.6 Hz, 1.5 Hz, 1H), 7.90 (d, J = 8.2 Hz, 1H), 7.86 (dd, J = 7.0 Hz, 2.0 Hz, 1H), 7.79 (t, J = 8.4 Hz, 2H), 7.56-7.34(m, 6H), 7.30 (t, J = 7.6 Hz, 1H), 7.10 (t, J = 5.6 Hz, 1H), 5.00 (d, J)= 5.6 Hz, 2H), 3.98 - 3.94 (m, 3H), 3.91 (s, 2H), 3.85 (d, J = 5.4 Hz,2H), 3.11 (s, 3H), 2.81 (t, J = 5.6 Hz, 2H), 2.67 (t, J = 5.6 Hz, 2H). 2protons could not be detected. 13 C NMR (101 MHz, CDCl₃): δ 162.3, 140.9, 139.5, 135.6, 134.0, 133.8, 131.6, 130.8, 128.8, 128.6, 126.8, 126.7, 126.1, 125.5, 125.3, 124.9, 123.7, 123.6, 120.8, 119.3, 117.1, 113.2, 61.4, 52.3, 51.1, 50.1, 49.0, 41.1, 40.8, 22.3. ESI HRMS (m/z): calcd for $C_{30}H_{32}N_5O_4S$ [M + H]⁺, 558.21695; found,

1-(2-Hydroxyethyl)-5-((7-methoxy-1H-indol-3-yl)methyl)-N-(naphthalen-1-ylmethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (25). Compound 25 was synthesized by employing the procedure described for compound 14, using intermediate 78 (80 mg, 0.21 mmol, 1.0 equiv), TEA (21 mg, 0.21 mmol, 1.0 equiv), 7-methoxyindole-3-carboxaldehyde (33 mg, 0.21 mmol, 1.0 equiv), AcOH (13 mg, 0.21 mmol, 1.0 equiv), NaBH(OAc)₃ (67 mg, 0.32 mmol, 1.5 equiv), and dry THF (3 mL). Purification by FC (EtOAc/MeOH/TEA 99:0:1 to 95:4:1) gave 61 mg (57%) of the title compound 25 as a pale pink solid. ¹H NMR (500 MHz, CDCl₃): δ 8.75 (s, 1H), 8.07–8.05 (m, 1H), 7.86–7.84 (m, 1H), 7.79 (d, J = 8.1 Hz, 1H), 7.49-7.45 (m, 3H), 7.41-7.38 (m, 1H), 7.79 (d, J = 8.1 Hz, 1H), 7.49-7.45 (m, 3H), 7.41-7.38 (m, 1H), 7.79 (d, J = 8.1 Hz, 1H), 7.49-7.45 (m, 3H), 7.41-7.38 (m, 1H), 7.49-7.45 (m, 2H), 7.41-7.38 (m, 2H),1H), 7.33 (d, J = 8.0 Hz, 1H), 7.14 (t, J = 5.7 Hz, 1H), 7.10 (d, J = 2.2Hz, 1H), 7.04 (t, J = 7.9 Hz, 1H), 6.64 (d, J = 7.7 Hz, 1H), 5.00 (d, J= 5.7 Hz, 1H), 3.95 (s, 2H), 3.94 (s, 5H), 3.81 (t, J = 5.2 Hz, 2H),3.70 (t, J = 5.1 Hz, 2H), 2.73 (t, J = 5.7 Hz, 2H), 2.53 (t, J = 5.6 Hz, 2H). ¹³C NMR (126 MHz, CDCl₃): δ 162.3, 146.3, 140.8, 139.8, 133.9, 133.8, 131.6, 129.4, 128.8, 128.6, 126.9, 126.8, 126.7, 126.1, 125.5, 123.8, 123.7, 120.0 117.7, 112.6, 112.1, 101.8, 61.2, 55.4, 52.5, 51.0, 49.9, 48.6, 41.1, 22.4. ESI HRMS (m/z): calcd for $C_{30}H_{32}N_5O_3$ [M + H]⁺, 510.24997; found, 510.24997.

1-(2-Hydroxyethyl)-5-((7-methoxy-1-methyl-1H-indol-3-yl)-methyl)-N-(naphthalen-1-ylmethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (26). Compound 26 was synthesized by employing the procedure described for compound 14, using intermediate 78 (60 mg, 0.16 mmol, 1.0 equiv), TEA (16 mg, 0.16 mmol, 1.0 equiv), 7-methoxy-1-methyl-1H-indole-3-carbalde-hyde (24 mg, 0.16 mmol, 1.0 equiv), AcOH (10 mg, 0.16 mmol, 1.0 equiv), NaBH(OAc) $_3$ (50 mg, 0.23 mmol, 1.5 equiv), and dry THF (2 mL). Purification by automated preparative chromatography (RP-C18, linear gradient from water/MeOH 90:10 to 15:85) gave 26 mg (31%) of the title compound 26 as a pale yellow solid. 1 H NMR (500 MHz, CDCl $_3$): δ 8.06 (d, J = 8.1 Hz, 1H), 7.85 (dd, J = 7.4 Hz, 1.4

Hz, 1H), 7.79 (d, J = 8.1 Hz, 1H), 7.56–7.44 (m, 3H), 7.40 (t, J = 7.6 Hz, 1H), 7.25 (d, J = 6.5 Hz, 1H), 7.13 (t, J = 5.6 Hz, 1H), 7.01 (s, 1H), 7.00 (t, J = 8.1 Hz, 1H), 6.61 (d, J = 7.7 Hz, 1H), 4.99 (d, J = 5.6 Hz, 2H), 4.03–4.01 (m, 7H), 3.91–3.89 (m, 6H), 3.78 (d, J = 4.5 Hz, 2H), 2.85 (t, J = 4.9 Hz, 2H), 2.70 (t, J = 5.3 Hz, 2H). ¹³C NMR (126 MHz, CDCl₃): δ 162.2, 148.0, 140.9, 139.3, 133.9, 133.7, 131.6, 130.8, 130.7, 128.8, 128.6, 126.8, 126.7 (2× C), 126.0, 125.5, 123.7, 120.1, 116.4, 112.0, 108.7, 102.5, 61.3, 55.5, 51.9, 49.2, 48.4, 41.1, 36.7, 21.9. ESI HRMS (m/z): calcd for C₃₁H₃₄N₅O₃ [M + H]⁺, 524.26562; found, 524.26560.

Methyl 3-((1-(2-Hydroxyethyl)-3-((naphthalen-1-ylmethyl)carbamoyl)-1,4,6,7-tetrahydro-5H-pyrazolo[4,3-c]pyridin-5-yl)methyl)-1H-indole-7-carboxylate (27). Compound 78 (80 mg, 0.21 mmol, 1.0 equiv), methyl 1H-indole-7-carboxylate (40 mg, 0.21 mmol, 1.0 equiv), and TEA (21 mg, 0.21 mmol, 1.0 equiv) were dissolved in MeOH (2 mL). 37% aqueous solution of formaldehyde (19 mg, 0.21 mmol, 1.0 equiv) and ZnCl₂ (43 mg, 0.32 mmol, 1.5 equiv) were then added and the resulting mixture was stirred overnight at rt. The reaction mixture was concentrated in vacuo. Purification of the residue by FC (EtOAc/MeOH/TEA 99:0:1 to 93:6:1) gave 50 mg (44%) of the title compound 27 as a white solid. ¹H NMR (500 MHz, CDCl₃): δ 9.76 (s, 1H), 8.07 (d, J = 8.0 Hz, 1H), 7.98 (d, J = 7.7 Hz, 1H), 7.88 (d, J = 7.5 Hz, 1H), 7.85 (d, J =7.2 Hz, 1H), 7.79 (d, J = 8.0 Hz, 1H), 7.52–7.46 (m, 3H), 7.40 (t, J =7.5 Hz, 1H), 7.31 (br s, 1H), 7.14 (t, J = 7.7 Hz, 1H), 7.07 (t, J = 5.5Hz, 2H), 3.99-3.93 (m, 10H), 3.82 (t, J = 4.9 Hz, 2H), 2.79 (t, J =5.6 Hz, 2H), 2.64 (t, J = 5.6 Hz, 2H). ¹³C NMR (91 MHz, CDCl₃): δ 168.0, 162.3, 141.0, 139.7, 136.4, 134.0, 133.8, 131.6, 129.1, 128.8, 128.6, 126.8, 126.7, 126.1, 125.5, 125.4, 125.1, 124.6, 123.8, 119.1, 117.5, 112.6, 112.5, 61.4, 52.5, 52.0, 51.1, 50.0, 48.8, 41.1, 22.4. ESI HRMS (m/z): calcd for $C_{31}H_{32}N_5O_4$ [M + H]⁺, 538.24488; found, 538.24484.

1-(2-Hvdroxvethyl)-5-((1H-indol-2-vl)methyl)-N-(naphthalen-1ylmethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (28). Compound 28 was synthesized by employing the procedure described for compound 14, using intermediate 78 (80 mg, 0.21 mmol, 1.0 equiv), TEA (21 mg, 0.21 mmol, 1.0 equiv), indole-2carboxaldehyde (30 mg, 0.21 mmol, 1.0 equiv), AcOH (13 mg, 0.21 mmol, 1.0 equiv), NaBH(OAc)₃ (67 mg, 0.32 mmol, 1.5 equiv), and dry THF (3 mL). Purification by FC (EtOAc/MeOH/TEA 99:0:1 to 93:6:1) gave 55 mg (53%) of the title compound 28 as a pale pink solid. ¹H NMR (500 MHz, CDCl₃): δ 8.82 (s, 1H), 8.09–8.07 (m, 1H), 7.87-7.85 (m, 1H), 7.79 (d, J = 8.2 Hz, 1H), 7.57 (d, J = 7.8Hz, 1H), 7.49-7.45 (m, 3H), 7.39-7.34 (m, 2H), 7.17-7.13 (m, 1H), 7.10-7.07 (m, 2H), 6.40 (s, 1H), 4.99 (d, J = 5.8 Hz, 2H), 3.91(s, 2H), 3.89–3.87 (m, 4H), 3.78 (t, J = 4.8 Hz, 2H), 2.65 (t, J = 5.6 Hz, 2H), 2.49 (t, J = 5.6 Hz, 2H). ¹³C NMR (126 MHz, CDCl₃): δ 162.1, 140.8, 139.6, 136.4, 135.8, 134.0, 133.8, 131.6, 128.8, 128.7, 128.5, 127.0, 126.7, 126.1, 125.5, 123.8, 121.7, 120.3, 119.7, 117.3, 111.0, 101.9, 61.2, 54.7, 51.2, 50.4, 48.5, 41.1, 22.2. ESI HRMS (m/ z): calcd for $C_{29}H_{30}N_5O_2$ [M + H]⁺, 480.23940; found, 480.23912.

1-(2-Hydroxyethyl)-N,5-bis(naphthalen-1-ylmethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (30). Compound 30 was synthesized by employing the procedure described for compound 14, using intermediate 78 (120 mg, 0.31 mmol, 1.0 equiv), TEA (31 mg, 0.31 mmol, 1.0 equiv), 1-naphthaldehyde (48 mg, 0.31 mmol, 1.0 equiv), AcOH (19 mg, 0.31 mmol, 1.0 equiv), NaBH(OAc)₃ (100 mg, 0.46 mmol, 1.5 equiv), and dry THF (5 mL). Purification by FC (EtOAc/MeOH/TEA 99:0:1 to 95:4:1) gave 60 mg (39%) of the title compound 30 as a pale yellow solid. ¹H NMR (500 MHz, CDCl₃): δ 8.30 (dd, J = 7.4 Hz, 2.1 Hz, 1H), 8.08 (d, J = 8.2 Hz, 1H), 7.87-7.83 (m, 2H), 7.80 (d, J = 8.2 Hz, 2H), 7.54-7.46(m, 6H), 7.42 (ddd, J = 8.0, 7.0, 5.6 Hz, 2H), 7.09 (t, J = 5.7 Hz, 1H), 5.01 (d, J = 5.7 Hz, 2H), 4.15 (s, 2H), 3.99 (s, 2H), 3.91 (t, J = 4.4Hz, 2H), 3.82 (t, J = 5.1 Hz, 2H), 2.77 (t, J = 5.7 Hz, 2H), 2.57 (t, J =5.7 Hz, 2H). ¹³C NMR (126 MHz, CDCl₃): δ 162.3, 140.9, 139.9, 134.3, 134.0, 133.9, 133.8, 132.7, 131.6, 128.8, 128.6, 128.5, 128.2, 127.6, 126.8, 126.7, 126.1, 126.0, 125.8, 125.5, 125.4, 124.9, 123.8, 117.7, 61.4, 60.2, 50.9, 50.5, 48.8, 41.1, 22.4. ESI HRMS (m/z): calcd for $C_{31}H_{31}N_4O_2$ [M + H]⁺, 491.24415; found, 491.24404.

1-(2-Hydroxyethyl)-5-((4-hydroxynaphthalen-1-yl)methyl)-N-(naphthalen-1-ylmethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (31). Compound 31 was synthesized by employing the procedure described for compound 14, using intermediate 78 (60 mg, 0.16 mmol, 1.0 equiv), TEA (16 mg, 0.16 mmol, 1.0 equiv), 4-hydroxy-1-naphthaldehyde (27 mg, 0.16 mmol, 1.0 equiv), AcOH (10 mg, 0.16 mmol, 1.0 equiv), NaBH(OAc)₃ (50 mg, 0.23 mmol, 1.5 equiv), and dry THF (2 mL). Purification by FC (EtOAc/MeOH/TEA 99:0:1 to 95:4:1) gave 40 mg (49%) of the title compound 31 as a white solid. ¹H NMR (500 MHz, DMSO- d_6): δ 10.09 (s, 1H), 8.41 (t, J = 5.5 Hz, 1H), 8.26–8.09 (m, 3H), 7.96– 7.93 (m, 1H), 7.83 (dd, I = 7.1, 2.0 Hz, 1H), 7.60–7.37 (m, 6H), 7.25 (d, J = 7.5 Hz, 1H), 6.82 (d, J = 7.5 Hz, 1H), 4.89 (t, J = 4.9 Hz, 1H),4.83 (d, J = 6.1 Hz, 2H), 4.03 (t, J = 4.9 Hz, 2H), 3.95 (s, 2H), 3.77-3.65 (m, 2H), 3.58 (s, 2H), 2.76 (br s, 2H), 2.70 (br s, 2H). ¹³C NMR (101 MHz, DMSO- d_6): δ 162.5, 153.4, 140.4, 140.0, 135.4, 133.7, 133.7, 131.3, 128.9, 128.6, 127.8, 126.6, 126.3, 126.2, 125.8, 125.8, 125.4, 125.3, 124.8, 124.7, 124.0, 122.7, 116.4, 107.4, 60.5, 60.1, 52.0, 49.8, 49.7, 22.28. ESI HRMS (m/z): calcd for $C_{31}H_{31}N_4O_3$ [M + H]⁺, 507.23907; found, 507.23902.

5-((4-Hydroxy-5,6,7,8-tetrahydronaphthalen-1-yl)methyl)-1-(2hydroxyethyl)-N-(naphthalen-1-ylmethyl)-4,5,6,7-tetrahydro-1Hpyrazolo[4,3-c]pyridine-3-carboxamide (32). Compound 32 was synthesized by employing the procedure described for compound 14, using intermediate 78 (360 mg, 0.93 mmol, 1.0 equiv), TEA (93 mg, 0.93 mmol, 1.0 equiv), 4-methoxy-5,6,7,8-tetrahydronaphthalene-1carbaldehyde (176 mg, 0.93 mmol, 1.0 equiv), AcOH (57 mg, 0.93 mmol, 1.0 equiv), NaBH(OAc)₃ (300 mg, 1.38 mmol, 1.5 equiv), and dry THF (10 mL). Purification by FC (EtOAc/MeOH/TEA 99:0:1 to 95:4:1) gave 265 mg (56%) of the title compound 32 as a pale white solid. ¹H NMR (400 MHz, CDCl₃): δ 8.08 (d, J = 7.9 Hz, 1H), 7.86 (dd, J = 7.3 Hz, 2.0 Hz, 1H), 7.80 (d, J = 8.1 Hz, 1H), 7.55 - 7.46(m, 3H), 7.43-7.40 (m, 1H), 6.64 (d, I = 8.3 Hz, 1H), 5.01 (d, I = 85.7 Hz, 2H), 3.93 (t, J = 5.1 Hz, 2H), 3.86 - 3.81 (m, 4H), 3.81 (s, 3H), 3.62 (s, 2H), 2.82 (br s, 2H), 2.71 (t, J = 5.5 Hz, 2H), 2.66 (br s, 2H), 2.60 (t, J = 5.5 Hz, 2H), 1.75 (t, J = 2.9 Hz, 4H). ¹³C NMR (101 MHz, CDCl₃): δ 162.3, 156.8, 140.9, 139.9, 138.1, 134.0, 133.8, 131.7, 128.8, 128.6, 128.2, 127.8, 126.8, 126.7, 126.4, 126.0, 125.5, 123.8, 117.9, 106.3, 61.4, 59.5, 55.3, 50.9, 50.3, 48.6, 41.1, 26.4, 23.7, 22.9, 22.5 (2× C). ESI HRMS (m/z): calcd for $C_{32}H_{37}N_4O_3$ [M + H]+, 525.28602; found, 525.28590.

1-(2-Hydroxyethyl)-5-((4-(methylthio)naphthalen-1-yl)methyl)-N-(naphthalen-1-ylmethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (33). Compound 33 was synthesized by employing the procedure described for compound 14, using intermediate 78 (60 mg, 0.16 mmol, 1.0 equiv), TEA (16 mg, 0.16 mmol, 1.0 equiv), 4-methylthio-1-naphthaldehyde (32 mg, 0.16 mmol, 1.0 equiv), AcOH (10 mg, 0.16 mmol, 1.0 equiv), NaBH(OAc)₃ (50 mg, 0.23 mmol, 1.5 equiv), and dry THF (2 mL). Purification by FC (EtOAc/MeOH/TEA 99:0:1 to 95:4:1) gave 37 mg (43%) of the title compound 33 as a white solid. ¹H NMR (400 MHz, CDCl₃): δ 8.34–8.26 (m, 2H), 8.07 (dd, J = 7.8 Hz, 1.2 Hz, 1H), 7.86 (dd, J = 7.2 Hz, 2.2 Hz, 1H), 7.79 (d, J = 8.1 Hz, 1H), 7.56-7.38 (m, 7H), 7.32 (d, J = 7.5 Hz, 1H), 7.09 (t, J = 5.7 Hz, 1H), 5.00 (d, J = 5.7 Hz, 2H), 4.12 (s, 2H), 3.97 (br s, 2H), 3.90 (dd, J = 5.7 Hz, 2H)5.6 Hz, 4.0 Hz, 2H), 3.81 (dd, J = 5.6 Hz, 4.0 Hz, 2H), 2.78 (t, J = 5.6 HzHz, 2H), 2.60-2.56 (m, 2H), 2.58 (s, 3H). 13 C NMR (101 MHz, CDCl₃): δ 162.3, 140.9, 139.8, 136.1, 134.0, 133.8, 132.8, 132.1, 131.6, 128.8, 128.6, 127.7, 126.8, 126.7, 126.4, 126.1, 126.0, 125.5, 124.8, 123.8, 123.0, 117.4, 61.4, 60.0, 51.0, 50.4, 48.9, 41.1, 22.4, 16.3. ESI HRMS (m/z): calcd for $C_{32}H_{33}N_4O_2S$ [M + H]⁺, 537.23187;

1-(2-Hydroxyethyl)-5-((4-(methylsulfinyl))naphthalen-1-yl)-methyl)-N-(naphthalen-1-ylmethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (34). Compound 34 was synthesized by employing the procedure described for compound 14, using intermediate 78 (80 mg, 0.21 mmol, 1.0 equiv), TEA (21 mg, 0.21 mmol, 1.0 equiv), 4-(methylsulfinyl)-1-naphthaldehyde (46 mg, 0.21 mmol, 1.0 equiv), AcOH (13 mg, 0.21 mmol, 1.0 equiv), NaBH(OAc)₃ (67 mg, 0.32 mmol, 1.5 equiv), and dry THF (3 mL). Purification by FC (EtOAc/MeOH/TEA 99:0:1 to 95:4:1) gave 56

mg (48%) of the title compound 34 as a white solid. 1 H NMR (500 MHz, CDCl₃): δ 8.40–8.38 (m, 1H), 8.11 (d, J = 7.4 Hz, 1H), 8.06 (d, J = 8.1 Hz, 1H), 7.94–7.90 (m, 1H), 7.84 (dd, J = 7.4 Hz, 1.9 Hz, 1H), 7.78 (d, J = 8.1 Hz, 1H), 7.72 (d, J = 7.0 Hz, 1H), 7.60–7.54 (m, 2H), 7.53–7.44 (m, 3H), 7.39 (dd, J = 8.1 Hz, 7.1 Hz, 1H), 7.17 (t, J = 5.6 Hz, 2H), 4.99 (d, J = 5.6 Hz, 2H), 4.20 (s, 2H), 4.02 (q, J = 14.9 Hz, 2H), 3.93 (t, J = 4.9 Hz, 2H), 3.83 (t, J = 4.9 Hz, 2H), 2.83 (s, 3H), 2.81 (br s, 2H), 2.65 (br s, 2H). 13 C NMR (75 MHz, CDCl₃): δ 162.3, 141.5, 140.9, 139.6, 133.9, 133.7, 132.7, 131.6, 129.1, 128.8, 128.6, 127.4, 127.4, 127.2, 126.9, 126.8, 126.7, 126.0, 126.0, 125.5, 123.7, 121.9, 121.8, 117.1, 61.3, 59.8, 51.1, 50.4, 49.0, 43.0, 41.0, 22.3. ESI HRMS (m/z): calcd for $C_{32}H_{33}N_4O_3S$ [M + H] $^+$, 553.22679: found, 553.22666.

5-((4-(Dimethylamino)naphthalen-1-yl)methyl)-1-(2-hydroxyethyl)-N-(naphthalen-1-ylmethyl)-4,5,6,7-tetrahydro-1Hpyrazolo[4,3-c]pyridine-3-carboxamide (35). Compound 78 (120 mg, 0.31 mmol, 1.0 equiv) was suspended in dry 1,2-dichloroethane (3 mL), followed by addition of TEA (31 mg, 0.31 mmol, 1.0 equiv). The mixture was stirred for 0.5 h and 4-dimethylaminonaphthaldehyde (62 mg, 0.31 mmol, 1.0 equiv), AcOH (19 mg, 0.31 mmol, 1.0 equiv), and NaBH(OAc)₃ (100 mg, 0.46 mmol, 1.5 equiv) were added. After stirring for 12 h, the mixture was quenched with saturated aqueous solution of NaHCO3 (5 mL) and extracted with DCM (2×5 mL). The combined organic extracts were washed with water (5 mL), brine (5 mL), dried over anhydrous Na2SO4, filtered, and concentrated in vacuo. The residue was purified by FC (EtOAc/ MeOH/TEA 99:0:1 to 93:6:1) to yield 57 mg (34%) of the title compound 35 as a white solid. ¹H NMR (400 MHz, CDCl₃): δ 8.28– 8.26 (m, 2H), 8.08 (dd, J = 8.1 Hz, 1.1 Hz, 1H), 7.86 (dd, J = 7.5 Hz, 2.1 Hz, 1H), 7.79 (d, I = 8.1 Hz, 1H), 7.54–7.45 (m, 5H), 7.40 (d, I= 8.1 Hz, 1H), 7.39 (t, J = 7.5 Hz, 1H), 7.10 (t, J = 5.7 Hz, 1H), 7.01(d, J = 7.5 Hz, 1H), 5.01 (d, J = 5.7 Hz, 2H), 4.08 (s, 2H), 3.97 (s, 2H), 3.97 (s, 2Hz)2H), 3.89 (dd, J = 5.6 Hz, J = 4.0 Hz, 2H), 3.80 (t, J = 5.2 Hz, 2H), 2.90 (s, 6H), 2.78 (t, J = 5.6 Hz, 2H), 2.58 (t, J = 5.6 Hz, 2H). ¹³C NMR (126 MHz, CDCl₃): δ 162.3, 151.0, 140.9, 139.9, 134.0, 133.9, 133.8, 131.6, 129.3, 128.8, 128.6, 127.9, 126.8, 126.7, 126.0 (2× C), 125.9, 125.5, 125.3, 125.0, 124.7 (2× C), 123.8, 117.7, 113.4, 61.3, 60.2, 51.0, 50.4, 45.4 (2× C), 41.1, 22.4. ESI HRMS (m/z): calcd for $C_{33}H_{36}N_5O_2$ [M + H]⁺, 534.28635; found, 534.28633.

5-Benzyl-1-(2-hydroxyethyl)-N-(naphthalen-1-ylmethyl)-4,5,6,7tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (36). Compound 36 was synthesized by employing the procedure described for compound 14, using intermediate 78 (80 mg, 0.21 mmol, 1.0 equiv), TEA (21 mg, 0.21 mmol, 1.0 equiv), benzaldehyde (22 mg, 0.21 mmol, 1.0 equiv), AcOH (13 mg, 0.21 mmol, 1.0 equiv), NaBH(OAc)₃ (67 mg, 0.32 mmol, 1.5 equiv), and dry THF (3 mL). Purification by FC (EtOAc/MeOH/TEA 99:0:1 to 95:4:1) gave 57 mg (62%) of the title compound 36 as a white solid. ¹H NMR (500 MHz, CDCl₃): δ 8.07 (d, J = 7.9 Hz, 1H), 7.87 (dd, J = 7.4 Hz, 1.9 Hz, 1H), 7.81 (d, I = 8.1 Hz, 1H), 7.54–7.47 (m, 3H), 7.44–7.40 (m, 1H), 7.39-7.37 (m, 2H), 7.33 (t, J = 7.4 Hz, 2H), 7.28 (dt, J =8.2 Hz, 1.8 Hz, 1H), 7.05 (t, J = 7.2 Hz, 1H), 5.01 (d, J = 5.7 Hz, 2H), 3.96 (t, J = 5.3 Hz, 2H), 3.87 (s, 2H), 3.85 (t, J = 5.6 Hz, 2H), 3.76 (s, 2H), 2.73 (t, J = 5.6 Hz, 2H), 2.66 (t, J = 5.4 Hz, 2H). ¹³C NMR (126 MHz, CDCl₃): δ 162.2, 141.0, 139.7, 138.2, 134.0, 133.8, 131.6, 129.3 (2× C), 128.8, 128.7, 128.5 (2× C), 127.4, 126.8, 125.7, 126.1, 125.5, 123.8, 117.6, 62.1, 61.4, 51.0, 50.1, 48.8, 41.1, 22.4. ESI HRMS (m/z): calcd for $C_{27}H_{29}N_4O_2$ [M + H]⁺, 441.22850; found, 441.22.865.

1-(2-Hydroxyethyl)-5-(4-methylbenzyl)-N-(naphthalen-1-yl-methyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (37). Compound 37 was synthesized by employing the procedure described for compound 14, using intermediate 78 (80 mg, 0.21 mmol, 1.0 equiv), TEA (21 mg, 0.21 mmol, 1.0 equiv), 4-methylbenzaldehyde (24 mg, 0.21 mmol, 1.0 equiv), AcOH (13 mg, 0.21 mmol, 1.0 equiv), NaBH(OAc)₃ (67 mg, 0.32 mmol, 1.5 equiv), and dry THF (3 mL). Purification by FC (EtOAc/MeOH/TEA 99:0:1 to 95:4:1) gave 54 mg (57%) of the title compound 37 as a white solid. ¹H NMR (300 MHz, CDCl₃): δ 8.11–8.04 (m, 1H), 7.88–7.85 (m, 1H), 7.81 (d, J = 8.0 Hz, 1H), 7.57–7.37 (m, 4H),

7.26 (d, J = 7.9 Hz, 1H), 7.14 (d, J = 7.8 Hz, 1H), 7.04 (t, J = 5.7 Hz, 1H), 5.02 (d, J = 5.7 Hz, 2H), 3.99–3.93 (m, 2H), 3.90–3.82 (m, 4H), 3.73 (s, 2H), 2.73 (t, J = 5.1 Hz, 2H), 2.66 (t, J = 5.0 Hz, 2H), 2.35 (s, 3H). ¹³C NMR (91 MHz, CDCl₃): δ 162.2, 141.0, 139.6, 137.1, 134.6, 134.0, 133.8, 131.6, 129.3 (2× C), 129.3 (2× C), 128.8, 128.6, 126.8, 126.7, 126.1, 125.5, 123.8, 117.3, 61.7, 61.4, 51.1, 50.0, 48.7, 41.1, 22.3, 21.3. ESI HRMS (m/z): calcd for C₂₈H₃₁N₄O₂ [M + H]⁺, 455.24415; found, 455.24392.

1-(2-Hvdroxyethyl)-5-(4-ethylbenzyl)-N-(naphthalen-1-ylmethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (38). Compound 38 was synthesized by employing the procedure described for compound 14, using intermediate 78 (80 mg, 0.21 mmol, 1.0 equiv), TEA (21 mg, 0.21 mmol, 1.0 equiv), 4ethylbenzaldehyde (28 mg, 0.21 mmol, 1.0 equiv), AcOH (13 mg, 0.21 mmol, 1.0 equiv), NaBH(OAc)₃ (67 mg, 0.32 mmol, 1.5 equiv), and dry THF (3 mL). Purification by FC (EtOAc/MeOH/TEA 99:0:1 to 95:4:1) gave 55 mg (56%) of the title compound 38 as a pale yellow solid. ¹H NMR (360 MHz, CDCl₃): δ 8.07 (d, J = 8.2 Hz, 1 1H), 2 7.86 (dd, 2 J = 6.7 Hz, 1.9 Hz, 1H), 2 7.80 (d, 2 J = 8.0 Hz, 1H), 7.57-7.37 (m, 4H), 7.29 (d, J = 7.9 Hz, 2H), 7.17 (d, J = 7.9 Hz, 2H), 7.07 (t, J = 5.3 Hz, 1H), 5.01 (t, J = 5.6 Hz, 2H), 3.94 (t, J = 4.8Hz, 2H), 3.89 (s, 2H), 3.84 (t, I = 4.5 Hz, 2H), 3.74 (s, 2H), 2.89 (br s, 1H), 2.73 (t, J = 5.3 Hz, 2H), 2.68-2.62 (m, 4H), 1.24 (t, J = 7.6Hz, 3H). 13 C NMR (91 MHz, CDCl₃): δ 162.2, 143.4, 141.0, 139.7, 135.3, 134.0, 133.8, 131.7, 129.3 (2× C), 128.8, 128.6, 128.0 (2× C), 126.8, 126.7, 126.1, 125.5, 123.8, 117.6, 61.7, 61.4, 51.0, 50.0, 48.7, 41.1, 28.7, 22.4, 15.7. ESI HRMS (m/z): calcd for $C_{29}H_{33}N_4O_2$ [M + H]+, 469.25980; found, 469.25967.

1-(2-Hydroxyethyl)-5-(4-isopropylbenzyl)-N-(naphthalen-1-ylmethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (39). Compound 39 was synthesized by employing the procedure described for compound 14, using intermediate 78 (80 mg, 0.21 mmol, 1.0 equiv), TEA (21 mg, 0.21 mmol, 1.0 equiv), 4isopropylbenzaldehyde (31 mg, 0.21 mmol, 1.0 equiv), AcOH (13 mg, 0.21 mmol, 1.0 equiv), NaBH(OAc)₃ (67 mg, 0.32 mmol, 1.5 equiv), and dry THF (3 mL). Purification by FC (EtOAc/MeOH/ TEA 99:0:1 to 95:4:1) gave 67 mg (65%) of the title compound 39 as a white solid. 1 H NMR (250 MHz, CDCl₃): δ 8.09–8.05 (m, 1H), 7.87-7.78 (m, 2H), 7.54-7.37 (m, 4H), 7.30 (d, J = 8.0 Hz, 2H), 7.19 (d, J = 8.0 Hz, 2H), 7.11 (t, J = 5.4 Hz, 1H), 4.99 (d, J = 5.7 Hz, 1Hz, 1Hz)2H), 3.87 (br s, 4H), 3.79 (t, J = 5.2 Hz, 2H), 2.91 (sept, J = 6.9 Hz, 1H), 2.72 (t, J = 4.9 Hz, 2H), 2.65 (t, J = 4.9 Hz, 2H), 1.26 (d, J = 6.9Hz, 6H). 13 C NMR (63 MHz, CDCl₃): δ 162.2, 148.0, 141.0, 139.6, 135.3, 134.0, 133.8, 131.6, 129.3 (2× C), 128.8, 128.6, 126.8, 126.7, 126.6 (2× C), 126.0, 125.5, 123.8, 117.5, 61.7, 61.3, 51.0, 50.0, 48.8, 41.1, 33.9, 24.2 (2× C), 22.4. ESI HRMS (m/z): calcd for $C_{30}H_{35}N_4O_2$ [M + H]⁺, 483.27545; found, 483.27524.

1-(2-Hydroxyethyl)-5-(4-ethoxybenzyl)-N-(naphthalen-1-ylmethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (40). Compound 40 was synthesized by employing the procedure described for compound 14, using intermediate 78 (80 mg, 0.21 mmol, 1.0 equiv), TEA (21 mg, 0.21 mmol, 1.0 equiv), 4ethoxybenzaldehyde (33 mg, 0.21 mmol, 1.0 equiv), AcOH (13 mg, 0.21 mmol, 1.0 equiv), NaBH(OAc)₃ (67 mg, 0.32 mmol, 1.5 equiv), and dry THF (3 mL). Purification by FC (EtOAc/MeOH/TEA 99:0:1 to 95:4:1) gave 62 mg (63%) of the title compound 40 as a white solid. ¹H NMR (500 MHz, CDCl₃): δ 8.07 (d, J = 7.9 Hz, 1H), 7.87 (dd, J = 7.4 Hz, 2.0 Hz, 1H), 7.81 (d, J = 8.1 Hz, 1H), 7.55-7.45(m, 3H), 7.42 (dd, J = 8.1 Hz, J = 7.0 Hz, 1H), 7.27 (d, J = 8.9 Hz, 2H), 7.04 (t, J = 5.6 Hz, 1H), 6.85 (d, J = 8.7 Hz, 2H), 5.01 (d, J = 5.7Hz, 2H), 4.03 (q, J = 7.0 Hz. 2H), 3.95 (t, J = 5.3 Hz, 2H), 3.86-3.84(m, 4H), 3.69 (s, 3H), 2.72 (t, J = 5.5 Hz, 2H), 2.65 (t, J = 5.3 Hz, 2H), 1.41 (t, J = 7.0 Hz, 3H). ¹³C NMR (126 MHz, CDCl₃): δ 162.2, 158.3, 140.9, 139.7, 134.0, 133.8, 131.6, 130.5 (2×), 130.1, 128.1, 128.7, 126.8, 126.7, 126.1, 125.5, 123.8, 117.6, 114.4 (2×), 63.5, 61.5, 61.4, 51.0, 50.0, 48.6, 41.1, 22.4, 15.0. ESI HRMS (m/z): calcd for $C_{29}H_{33}N_4O_3$ [M + H]⁺, 485.25472; found, 485.25447.

1-(2-Hydroxyethyl)-5-(4-chlorobenzyl)-N-(naphthalen-1-ylmethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (41). Compound 41 was synthesized by employing the procedure described for compound 14, using intermediate 78 (80 mg, 0.21

mmol, 1.0 equiv), TEA (21 mg, 0.21 mmol, 1.0 equiv), 4chlorobenzaldehyde (29 mg, 0.21 mmol, 1.0 equiv), AcOH (13 mg, 0.21 mmol, 1.0 equiv), NaBH(OAc)₃ (67 mg, 0.32 mmol, 1.5 equiv), and dry THF (3 mL). Purification by FC (EtOAc/MeOH/TEA 99:0:1 to 95:4:1) gave 50 mg (50%) of the title compound 41 as a pale yellow solid. ¹H NMR (500 MHz, DMSO- d_6): δ 8.44 (t, J = 5.8Hz, 1H), 8.19 (d, J = 7.6 Hz, 1H), 7.93 (d, J = 7.1 Hz, 1H), 7.82 (t, J= 4.6 Hz, 1H, 7.55-7.51 (m, 2H), 7.45 (d, I = 4.6 Hz, 2H), 7.37(virt. q, $J = J \approx 8.3$ Hz, 4H), 4.90 (t, J = 5.2 Hz, 1H), 4.84 (d, J = 5.8Hz, 2H), 4.04 (t, J = 5.2 Hz, 2H), 3.71 (d, J = 5.2 Hz, 2H), 3.66 (s, 2H), 3.54 (s, 2H), 2.73-2.70 (m, 4H). ¹³C NMR (126 MHz, DMSO d_6): δ 162.1, 140.3, 139.4, 137.9, 135.0,133.3, 131.5, 130.8, 130.4 (2× C), 128.5, 128.2 (2 \times C), 127.3, 126.2, 125.7, 125.4 (2 \times C), 123.5, 115.8, 60.2 (2× C), 51.6, 49.2, 49.2, 21.8. 1 CH₂ group overlaps with the DMSO signal. ESI HRMS (m/z): calcd for $C_{27}H_{28}ClN_4O_2$ [M + H]+, 475.18953; found, 475.18939.

1-(2-Hvdroxvethvl)-N-(naphthalen-1-vlmethvl)-5-(4-(trifluoromethoxy)benzyl)-4,5,6,7-tetrahydro-1H-pyrazólo[4,3-c]pyridine-3-carboxamide (42). Compound 42 was synthesized by employing the procedure described for compound 14, using intermediate 78 (80 mg, 0.21 mmol, 1.0 equiv), TEA (21 mg, 0.21 mmol, 1.0 equiv), 4-trifluoromethyoxybenzaldehyde (37 mg, 0.21 mmol, 1.0 equiv), AcOH (13 mg, 0.21 mmol, 1.0 equiv), NaBH(OAc)₃ (67 mg, 0.32 mmol, 1.5 equiv), and dry THF (3 mL). Purification by FC (EtOAc/MeOH/TEA 99:0:1 to 95:4:1) gave 70 mg (64%) of the title compound 42 as a pale yellow solid. ¹H NMR (500 MHz, CDCl₂): δ 8.08 (dd, I = 8.0 Hz, 1.1 Hz, 1H), 7.87 (dd, J = 7.3 Hz, 2.0 Hz, 1H), 7.81 (d, J = 8.0 Hz, 1H), 7.55-7.47 (m, J = 8.0 Hz, 1Hz), 7.55-7.47 (m, J = 8.0 Hz), 7.55-7.47 (m, J = 8.0 Hz),3H), 7.44 (d, J = 8.0 Hz, 1H), 7.40 (d, J = 8.7 Hz, 2H), 7.18 (d, J =8.0 Hz, 2H), 7.02 (t, J = 5.7 Hz, 1H), 5.02 (d, J = 5.7 Hz, 2H), 4.00 (d, J = 5.3 Hz, 2H), 3.92 - 3.89 (m, 2H), 3.88 (s, 2H), 3.76 (s, 2H),2.74 (t, J = 5.5 Hz, 2H), 2.68 (t, J = 5.5 Hz, 2H). ¹³C NMR (91 MHz, CDCl₃): δ 162.2, 148.5, 141.0, 139.6, 137.3, 134.0, 133.8, 131.7, 130.3 (2× C), 128.8, 128.7, 126.8, 126.7, 126.1, 125.5, 123.8, 121.0 (2× C), 117.5, 61.4, 61.2, 51.0, 50.1, 49.0, 41.1, 22.5. CF₃ signal could not be detected. ESI HRMS (m/z): calcd for $C_{28}H_{28}F_3N_4O_3$ [M + H]+, 525.21080; found, 525.21064.

5-(3,4-Dimethoxybenzyl)-1-(2-hydroxyethyl)-N-(naphthalen-1ylmethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (43). Compound 43 was synthesized by employing the procedure described for compound 14, using intermediate 78 (80 mg, 0.21 mmol, 1.0 equiv), TEA (21 mg, 0.21 mmol, 1.0 equiv), 3,4dimethoxybenzaldehyde (37 mg, 0.21 mmol, 1.0 equiv), AcOH (13 mg, 0.21 mmol, 1.0 equiv), NaBH(OAc)₃ (67 mg, 0.32 mmol, 1.5 equiv), and dry THF (3 mL). Purification by FC (EtOAc/MeOH/ TEA 99:0:1 to 95:4:1) gave 34 mg (32%) of the title compound 43 as a white solid. ¹H NMR (300 MHz, CDCl₃): δ 8.07 (dd, I = 6.6 Hz, 2.4 Hz, 1H), 7.86 (dd, J = 6.3 Hz, 3.1 Hz, 1H), 7.80 (d, J = 7.9 Hz, 1H), 7.58-7.35 (m, 4H), 7.10 (t, J = 5.6 Hz, 1H), 6.94 (d, 1.6 Hz, 1H), 6.88 (dd, J = 8.1 Hz, 1.6 Hz, 1H), 6.81 (d, J = 8.1 Hz, 1H), 5.00 (d, J = 5.6 Hz, 2H), 3.96 - 3.88 (m, 3H), 3.87 (s, 3H), 3.87 (s, 3H),3.85-3.79 (m, 3H), 3.69 (s, 2H), 2.69 (d, J = 4.8 Hz, 2H), 2.64 (d, J = 4.8 Hz, 2H). 13 C NMR (75 MHz, CDCl₃): δ 162.2, 149.1, 148.4, 141.0, 139.7, 134.0, 133.8, 131.6, 131.0, 128.8, 128.6, 126.8, 126.7, 126.0, 125.5, 123.8, 121.4, 117.6, 112.3, 111.0, 61.8, 61.3, 56.1, 51.0, 50.1, 48.6, 41.1, 22.4. ESI HRMS (m/z): calcd for $C_{29}H_{33}N_4O_4$ [M + H]⁺, 501.24963; found, 501.24950.

5-(3,4-Dichlorobenzyl)-1-(2-hydroxyethyl)-N-(naphthalen-1-ylmethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (44). Compound 44 was synthesized by employing the procedure described for compound 14, using intermediate 78 (80 mg, 0.21 mmol, 1.0 equiv), TEA (21 mg, 0.21 mmol, 1.0 equiv), 3,4dichlorobenzaldehyde (36 mg, 0.21 mmol, 1.0 equiv), AcOH (13 mg, 0.21 mmol, 1.0 equiv), NaBH(OAc)₃ (67 mg, 0.32 mmol, 1.5 equiv), and dry THF (3 mL). Purification by FC (EtOAc/MeOH/TEA 99:0:1 to 95:4:1) gave 55 mg (50%) of the title compound 44 as a pale yellow solid. ¹H NMR (500 MHz, CDCl₃): δ 8.06 (d, J = 7.8 Hz, 1H), 7.86 (dd, J = 6.9 Hz, 2.2 Hz, 1H), 7.80 (d, J = 8.1 Hz, 1H), 7.53-7.45 (m, 5H), 7.41 (d, J = 8.0 Hz, 1H), 7.38 (d, J = 8.1 Hz, 1H), 7.20 (dd, J = 8.1 Hz, 1.8 Hz, 1H), 7.10 (t, J = 5.7 Hz, 1H), 4.99

(d, J = 5.7 Hz), 3.94 (t, J = 4.5 Hz, 2H), 3.85 - 3.83 (m, 4H), 3.68 (s, 4H)2H), 2.71 (t, J = 5.4 Hz), 2.65 (t, J = 5.4 Hz, 2H), 2.16 (s, 1H). ¹³C NMR (126 MHz, CDCl₃): δ 162.2, 140.0, 139.5, 139.0, 134.0, 133.8, 132.6, 131.6, 131.2, 130.8, 130.5, 128.8, 128.6, 128.3, 126.8, 126.7, 126.1, 125.5, 123.7, 117.3, 61.3, 60.8, 51.0, 50.0, 49.0, 41.1, 22.4. ESI HRMS (m/z): calcd for $C_{27}H_{27}Cl_2N_4O_2$ [M + H]⁺, 509.15056; found, 509.15037.

5-((2,3-Dihydro-1H-inden-2-yl)methyl)-1-(2-hydroxyethyl)-N-(naphthalen-1-ylmethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (45). Compound 45 was synthesized by employing the procedure described for compound 14, using intermediate 78 (80 mg, 0.21 mmol, 1.0 equiv), TEA (21 mg, 0.21 mmol, 1.0 equiv), 2,3-dihydro-1H-indene-2-carbaldehyde (31 mg, 0.21 mmol, 1.0 equiv), AcOH (13 mg, 0.21 mmol, 1.0 equiv), NaBH(OAc)₃ (67 mg, 0.32 mmol, 1.5 equiv), and dry THF (3 mL). Purification by FC (EtOAc/MeOH/TEA 99:0:1 to 95:4:1) gave 71 mg (70%) of the title compound 45 as a white solid. ¹H NMR (300 MHz, CDCl₃): δ 8.14–8.05 (m, 1H), 7.87 (dd, J = 7.0 Hz, 2.4 Hz, 1H), 7.81 (d, J = 8.0 Hz, 1H), 7.58-7.47 (m, 3H), 7.44 (d, J = 8.0Hz, 1H), 7.23-7.17 (m, 2H), 7.16-7.11 (m, 2H), 7.05 (t, J = 5.7 Hz, 1H), 5.03 (d, I = 5.7 Hz, 2H), 4.04–3.95 (m, 2H), 3.92–3.85 (m, 4H), 3.12 (d, J = 7.3 Hz, 1H), 3.07 (d, J = 7.6 Hz, 1H), 2.90-2.76(m, 5H), 2.71 (t, J = 6.5 Hz, 3H), 2.64 (d, J = 7.3 Hz, 2H). ¹³C NMR (75 MHz, CDCl₃): δ 162.2, 143.0 (2× C), 140.9, 139.5, 133.9, 133.8, 131.6, 128.8, 128.6, 126.8, 126.7, 126.3 (2× C), 126.0, 125.5, 124.7 $(2 \times C)$, 123.8, 117.1, 62.8, 61.2, 51.1, 50.1, 49.7, 41.0, 37.9 $(2 \times C)$, 37.1, 22.1. ESI HRMS (m/z): calcd for $C_{30}H_{33}N_4O_2$ $[M + H]^+$, 481.25980; found, 481.25949.

2-((4-Methoxynaphthalen-1-yl)methyl)-9-(naphthalen-1-ylmethyl)-1,2,3,4,8,9-hexahydropyrido[4',3':3,4]pyrazolo[1,5-a]pyrazin-10(7H)-one (46). To a stirred, cooled (ice-salt bath) solution of compound 80 (75 mg, 0.14 mol, 1.0 equiv), DIPEA (57 mg, 0.44 mmol, 3.0 equiv), and DMAP (1 mg, 5 mol %) in dry DCM (2 mL) was added methanesulfonly chloride (37 mg, 0.28 mmol, 2.0 equiv). The reaction mixture was allowed to reach rt and stirred for 1 h. The solution was washed with water (0.5 mL), brine (0.5 mL), dried over anhydrous Na2SO4, filtered, and concentrated in vacuo. The residue was redissolved in dry toluene (2 × 1 mL) and evaporated in vacuo. The obtained brown oil was dissolved in dry THF (0.5 mL). The solution was cooled (ice-salt bath) and ^tBuOK (31 mg, 0.28 mmol, 2.0 equiv) was added in portions. The reaction mixture was allowed to warm to rt, stirred overnight followed by quenching with saturated aqueous solution of NH₄Cl (1 mL). The mixture was extracted with AcOEt (2 × 1 mL) and the combined organic extracts were washed with brine, dried over anhydrous Na₂SO₄, filtered, and evaporated in vacuo. Purification of the residue by FC (hexane/EtOAc 1:1 to 0:1) gave 40 mg (57%) of the title compound 46 as a beige solid. ¹H NMR (500 MHz, CDCl₃): δ 8.29 (dd, J = 8.5 Hz, 1.2 Hz, 2H), 8.08 (dd, J = 6.7 Hz, 2.2 Hz, 1H), 7.88 (dd, J = 6.7 Hz, 2.9 Hz, 1H), 7.85 (dd, J =7.5 Hz, 1.9 Hz, 1H), 7.56-7.45 (m, 5H), 7.45 (d, J = 7.0 Hz, 1H), 7.39 (d, J = 7.7 Hz, 1H), 6.76 (d, J = 7.7 Hz, 1H), 5.16 (s, 2H), 4.08(s, 2H), 4.07 (d, J = 6.2 Hz, 2H), 4.01 (s, 3H), 3.97 (s, 2H), 3.55-3.52 (m, 2H), 2.84 (t, J = 5.7 Hz, 2H), 2.74 (t, J = 5.7 Hz, 2H). ¹³C NMR (126 MHz, CDCl₃): δ 158.5, 155.4, 148.3, 134.1, 133.5, 131.8, 131.8, 129.3, 128.9, 128.8, 128.1, 127.8, 127.1, 126.5, 126.4, 126.3, 126.1, 125.2, 125.1, 124.8, 123.9, 122.3, 119.1, 103.0, 60.5, 55.6, 50.0 $(2 \times C)$, 46.9, 46.1, 43.9, 24.0. ESI HRMS (m/z): calcd for $C_{32}H_{31}N_4O_2$ [M + H]⁺, 503.24415; found, 503.24455.

8-((4-Methoxynaphthalen-1-yl)methyl)-2-(naphthalen-1-ylmethyl)-1,2,6,7,8,9-hexahydro-3H-imidazo[1',5':1,5]pyrazolo[4,3c]pyridin-3-one (47). To a stirred, cooled (ice-salt bath) solution of compound 83 (100 mg, 0.22 mmol, 1.0 equiv) in dry DCM (5 mL) was added CDI (42 mg, 0.26 mmol, 1.2 equiv). The reaction mixture was allowed to reach rt, stirred overnight, and evaporated in vacuo. Purification by automated preparative chromatography (SiO₂, linear gradient hexane/EtOAc 1:0 to 0:1) gave 35 mg (33%) of the title compound 47 as a white solid. ¹H NMR (400 MHz, CDCl₃): δ 8.31– 8.23 (m, 1H), 8.21-8.12 (m, 2H), 7.92-7.92 (m, 2H), 7.59-7.48 (m, 2H), 7.48-7.39 (m, 4H), 7.31 (d, J = 8.5 Hz, 1H), 6.72 (d, J =7.8 Hz, 1H), 5.12 (s, 2H), 4.03-3.95 (m, 5H), 3.90 (s, 2H), 3.34 (s,

2H), 2.90 (s, 4H). 13 C NMR (101 MHz, CDCl₃): δ 155.5, 149.4, 135.5 (br s), 134.0, 133.3, 131.4, 131.0, 129.5, 128.8, 127.8, 127.3, 126.5, 126.4, 125.9, 125.1, 124.4 (br s), 123.5, 122.3, 102.8, 76.7 (br s), 60.1, 55.5, 50.5, 47.8 (br s), 46.1, 42.8, 24.4 (br s). ESI HRMS (m/z): calcd for $C_{31}H_{29}N_4O_2$ [M + H] $^+$, 489.22850; found, 489.22856.

2-(5-((4-Methoxynaphthalen-1-yl)methyl)-3-(((naphthalen-1ylmethyl)amino)methyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridin-1-yl)ethan-1-ol (48). Compound 48 was synthesized by employing the procedure described for compound 14, using intermediate 86 (40 mg, 0.11 mmol, 1.0 equiv), 1-naphthaldehyde (17 mg, 0.11 mmol, 1.0 equiv), AcOH (7 mg, 0.11 mmol, 1.0 equiv), NaBH(OAc)₃ (34 mg, 0.16 mmol, 1.5 equiv), and dry THF (1 mL). Purification by FC (DCM/MeOH/7 M solution of NH₃ in MeOH 99:0:1 to 94:5:1) gave 17 mg (30%) of the title compound 48 as a white solid. ¹H NMR (500 MHz, DMSO- d_6): δ 8.21 (d, J = 8.5 Hz, 1H), 8.17 (d, J = 7.9 Hz, 1H), 8.12 (d, J = 7.4 Hz, 1H), 7.91 (dd, J =6.8 Hz, 2.3 Hz, 1H), 7.81 (d, J = 8.1 Hz, 1H), 7.54–7.37 (m, 7H), 7.33 (d, I = 7.8 Hz, 1H), 6.87 (d, I = 7.8 Hz, 1H), 4.79 (t, I = 5.4 Hz, 1H), 4.11 (s, 2H), 3.96 (s, 3H), 3.92 (br s, 4H), 3.67-3.60 (m, 4H), 3.41 (s, 2H), 2.73 (t, J = 5.7 Hz, 2H), 2.63 (t, J = 5.7 Hz, 2H). ¹³C NMR (126 MHz, DMSO- d_6): δ 154.5, 144.9, 137.7, 136.0, 133.4, 132.9, 131.5, 128.3, 127.7, 127.2, 126.2, 126.1, 125.9, 125.8, 125.6, 125.3, 125.2, 125.1, 125.0, 124.2, 121.7, 112.3, 103.4, 60.4, 59.6, 55.5, 50.7, 50.0, 49.5, 48.9, 45.4, 21.8. ESI HRMS (m/z): calcd for $C_{32}H_{35}N_4O_2$ [M + H]⁺, 507.27545; found, 507.27562.

N-((1-(2-Hydroxyethyl)-5-((4-methoxynaphthalen-1-yl)methyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridin-3-yl)methyl)naphthalene-1-sulfonamide (49). To a stirred, cooled (ice-salt bath) solution of compound 86 (40 mg, 0.11 mol, 1.0 equiv), TEA (16 mg, 0.16 mmol, 1.5 equiv), and DMAP (1 mg, 5 mol %) in dry DCM (1 mL) was added naphthalene-1-sulfonyl chloride (25 mg, 0.11 mmol, 1.0 equiv). The reaction mixture was allowed to reach rt and stirred overnight. The solution was washed with water (0.5 mL), brine (0.5 mL), dried over anhydrous Na2SO4, filtered, and concentrated in vacuo. Purification by FC (EtOAc/MeOH 99:1 to 95:5) gave 35 mg (57%) of the title compound 49 as a white solid. ¹H NMR (500 MHz, CDCl₃): δ 8.61 (d, J = 8.3 Hz, 1H), 8.30 (d, J = 7.9 Hz, 1H), 8.16 (d, J = 7.2 Hz, 1H), 8.12 (d, J = 8.3 Hz, 1H), 7.96 (d, J = 8.3 Hz, 1H), 7.86 (d, J = 7.6 Hz, 1H), 7.59 - 7.45 (m, 4H), 7.41 (t, J = 7.8 Hz, 1H),7.32 (d, J = 7.8 Hz, 1H), 6.76 (d, J = 7.8 Hz, 1H), 6.22 (br s, 1H), 4.01 (s, 2H), 3.91 (s, 2H), 3.84 (d, J = 4.9 Hz, 2H), 3.77 (t, J = 4.9Hz, 2H), 3.62 (t, I = 4.9 Hz, 2H), 3.29 (br s, 2H), 2.66 (t, I = 5.2 Hz, 2H), 2.46 (t, J = 5.5 Hz, 2H). ¹³C NMR (126 MHz, CDCl₃): δ 155.6, 142.2, 138.8, 134.6, 134.2, 134.1, 133.4, 129.6, 129.0, 128.3, 128.2, 128.1, 126.9, 126.7, 126.1, 125.2, 124.7, 124.5, 124.2, 122.5, 119.9, 112.9, 103.1, 61.3, 59.7, 55.7, 50.2, 49.1, 48.7, 39.6, 22.0. ESI HRMS (m/z): calcd for $C_{31}H_{33}N_4O_4S$ $[M + H]^+$, 557.22170; found, 557.22161.

5-((4-Methoxynaphthalen-1-yl)methyl)-1-(2-(methylamino)ethyl)-N-(naphthalen-1-ylmethyl)-4,5,6,7-tetrahydro-1H-pyrazolo-[4,3-c]pyridine-3-carboxamide (51). Compound 87 (120 mg, 0.18 mmol, 1.0 equiv) was dissolved in DCM (1 mL), followed by addition of 33% ethanolic solution of MeNH₂ (2 mL, 15.60 mmol, 86.6 equiv). The mixture was refluxed for 4 h, cooled to rt, and evaporated in vacuo. Purification by FC (DCM/MeOH/7 M solution of NH3 in MeOH 99:0:1 to 94:5:1) gave 44 mg (46%) of the title compound 51 as a pale yellow solid. ¹H NMR (400 MHz, DMSO- d_6): δ 8.42 (t, J =6.0 Hz, 1H), 8.24 (d, J = 7.9 Hz, 1H), 8.18 (d, J = 8.8 Hz, 1H), 7.93 (dd, I = 6.0 Hz, 3.2 Hz, 1H), 7.86 - 7.78 (m, 1H), 7.56 - 7.45 (m, 5H),7.43 (d, J = 5.3 Hz, 2H), 7.37 (d, J = 7.9 Hz, 1H), 6.90 (d, J = 7.9 Hz, 1H), 4.83 (d, J = 6.0 Hz, 2H), 4.02 (t, J = 6.0 Hz, 2H), 3.98 (s, 2H), 3.96 (s, 2H), 3.58 (s, 2H), 2.80 (d, J = 6.1 Hz, 2H), 2.77 (d, J = 6.1Hz, 2H), 2.67 (t, J = 5.1 Hz, 2H), 2.23 (s, 3H). ¹³C NMR (101 MHz, DMSO- d_6): δ 162.1, 154.5, 139.9, 139.0, 134.9, 133.2, 132.8, 130.8, 128.4, 127.8, 127.3, 126.2, 126.1, 126.0, 125.7, 125.4, 125.3, 125.2, 125.0, 124.9, 123.4, 121.7, 116.0, 103.4, 59.5, 55.4, 51.0, 49.3, 49.2, 48.5, 35.8, 28.1, 21.7. ESI HRMS (m/z): calcd for $C_{33}H_{36}N_5O_2$ [M + H]⁺, 534.28635; found, 534.28647.

1-(2-(Isopropylamino)ethyl)-5-((4-methoxynaphthalen-1-yl)-methyl)-N-(naphthalen-1-ylmethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (52). To a solution of

compound 50 (100 mg, 0.19 mmol) in dry THF (2 mL) was added acetone (11 mg, 0.19 mmol, 1.0 equiv), AcOH (11 mg, 0.19 mol, 1.0 equiv), and NaBH(OAc)₃ (60 mg, 0.28 mmol, 1.5 equiv). The reaction mixture was stirred overnight at rt, followed by addition of saturated aqueous solution of NaHCO₂ (1 mL). The mixture was extracted with EtOAc (2 × 2 mL) and the combined organic phase was washed with water (1 mL), brine (1 mL), dried over anhydrous Na2SO4, filtered, and concentrated in vacuo. Purification by FC (DCM/MeOH/7 M solution of NH₃ in MeOH 99:0:1 to 94:5:1) gave 85 mg (80%) of the title compound 52 as a white solid. ¹H NMR (500 MHz, DMSO- d_6): δ 8.44 (t, J = 6.2 Hz, 1H), 8.24 (d, J = 7.9 Hz, 1H), 8.17 (d, J = 7.9 Hz, 2H), 7.92 (dd, J = 6.5 Hz, 2.8 Hz, 1H), 7.84-7.77 (m, 1H), 7.57-7.46 (m, 4H), 7.45-7.40 (m, 2H), 7.37 (d, I = 7.9 Hz, 1H), 6.89 (d, I = 7.9 Hz, 1H), 4.82 (d, I = 6.1 Hz, 2H), 4.00 (t, J = 6.6 Hz, 2H), 3.98 (s, 2H), 3.96 (s, 3H), 3.58 (s, 2H), 2.82(t, J = 6.6 Hz, 2H), 2.77 (t, J = 5.4 Hz, 2H), 2.68 (t, J = 5.1 Hz, 2H),2.61 (sept, I = 6.2 Hz, 1H), 0.90 (d, I = 6.2 Hz, 6H). ¹³C NMR (126 MHz, DMSO- d_6): δ 162.1, 154.5, 139.9, 139.0, 135.0, 133.2, 132.9, 130.8, 128.5, 127.8, 127.3, 126.2, 126.2, 126.0, 125.7, 125.4, 125.2, 125.2, 125.0, 125.0, 123.5, 121.7, 116.0, 103.4, 59.5, 55.5, 49.4, 49.3, 49.3, 47.8, 46.5, 22.8 (2× C), 21.8. 1 C atom overlaps with DMSO- d_6 signal. ESI HRMS (m/z): calcd for $C_{35}H_{40}N_5O_2$ $[M + H]^+$, 562.31765; found, 562.31897.

1-(Azetidin-3-yl)-5-((4-methoxynaphthalen-1-yl)methyl)-N-(naphthalen-1-ylmethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (53). Compound 88 (240 mg, 0.38 mmol) was stirred in 4 M solution of HCl in 1,4-dioxane (2 mL) for 4 h at rt. The solution was concentrated in vacuo and the residue was partitioned between DCM (2 mL) and saturated aqueous solution of NaHCO₃ (1 mL). The phases were separated and the organic layer was dried over anhydrous Na2SO4, filtered, and evaporated in vacuo. Purification by FC (DCM/MeOH/7 M solution of NH3 in MeOH 99:0:1 to 93:6:1) gave 123 mg (61%) of the title compound 53 as a pale yellow solid. ¹H NMR (400 MHz, DMSO- d_6): δ 8.43 (t, J = 5.9Hz, 1H), 8.23 (d, J = 7.9 Hz, 1H), 8.17 (d, J = 8.1 Hz, 2H), 7.94-7.92 (m, 1H), 7.82 (t, J = 4.6 Hz, 1H), 7.56 - 7.41 (m, 6H), 7.37 (d, J= 7.9 Hz, 2H), 6.90 (d, J = 7.9 Hz, 1H), 5.07 (p, J = 7.3 Hz, 1H), 4.85 (p, J = 7.3 Hz, 2H), 3.85 (p, J = 7.3 Hz, 3.85 (p, J = 7.3 Hz), $3.85 \text{ (p, } J = 7.3 \text{$ (d, J = 5.9 Hz, 2H), 3.98 (s, 3H), 3.96 (s, 4H), 3.63 (t, J = 7.3 Hz,2H), 3.57 (s, 2H), 2.76 (t, J = 5.3 Hz, 2H), 2.65 (t, J = 5.3 Hz, 2H). 13 C NMR (101 MHz, DMSO- d_6): δ 162.0, 154.5, 140.1, 138.6, 134.9, 133.2, 132.8, 130.8, 128.5, 127.7, 127.3, 126.2, 126.1, 125.9, 125.7, 125.4, 125.4, 125.2, 125.0, 124.9, 123.4, 121.6, 116.4, 103.4, 59.4, 55.5, 52.7 (2× C), 51.4, 49.1 (2× C), 21.5. 1 signal overlaps with DMSO- d_6 . ESI HRMS (m/z): calcd for $C_{33}H_{34}N_5O_2$ $[M + H]^+$, 532.27070; found, 532.27141.

1-(2-(Dimethylamino)ethyl)-5-((4-methoxynaphthalen-1-yl)methyl)-N-(naphthalen-1-ylmethyl)-4,5,6,7-tetrahydro-1Hpyrazolo[4,3-c]pyridine-3-carboxamide (54). To a solution of compound 50 (80 mg, 0.15 mmo) in dry THF (1 mL) was added 37% aqueous solution of formaldehyde (36 mg, 0.46 mmol, 3.0 equiv) and NaBH(OAc)₃ (131 mg, 0.62 mmol, 4.0 equiv). The reaction mixture was stirred overnight at rt, followed by addition of saturated aqueous solution of NaHCO₃ (1 mL). The mixture was extracted with EtOAc (2 × 1 mL) and the combined organic phase was washed with water (1 mL), brine (1 mL), dried over anhydrous Na₂SO₄, filtered, and concentrated in vacuo. Purification by FC (DCM/ MeOH/7 M solution of NH₃ in MeOH 99:0:1 to 94:5:1) gave 68 mg (83%) of the title compound 54 as a white solid. ¹H NMR (500 MHz, DMSO- d_6): δ 8.43 (t, J = 6.2 Hz, 1H), 8.23 (d, J = 7.8 Hz, 1H), 8.17 (d, J = 7.8 Hz, 2H), 7.93 (dd, J = 5.9 Hz, 2.4 Hz, 1H), 7.81 (dd, J =6.9 Hz, 2.4 Hz, 1H), 7.56–7.44 (m, 4H), 7.44–7.40 (m, 2H), 7.37 (d, J = 7.8 Hz, 1H, 6.90 (d, J = 7.8 Hz, 1H), 4.82 (d, J = 6.2 Hz, 2H),4.05 (t, J = 6.7 Hz, 2H), 3.98 (s, 2H), 3.96 (s, 3H), 3.57 (s, 2H), 2.77 (t, J = 5.4 Hz, 2H), 2.68 (t, J = 5.4 Hz, 2H), 2.56 (t, J = 6.7 Hz, 2H),2.12 (s, 6H). ¹³C NMR (126 MHz, DMSO- d_6): δ 162.1, 154.4, 139.9, 139.0, 135.0, 133.3, 132.9, 130.8, 128.5, 127.9, 127.3, 126.3, 126.2, 126.0, 125.7, 125.4, 125.3, 125.2, 125.1, 125.0, 123.5, 121.7, 116.1, 103.4, 59.5, 58.4, 55.5, 49.4, 49.3, 47.1 (2× C), 45.3, 21.7. 1 C atom overlaps with DMSO- d_6 signal. ESI HRMS (m/z): calcd for $C_{34}H_{38}N_5O_2$ [M + H]⁺, 548.30200; found, 548.30223.

1-(2-(1H-Imidazole-1-yl)ethyl)-5-((4-methoxynaphthalen-1-yl)methyl)-N-(naphthalen-1-ylmethyl)-4,5,6,7-tetrahydro-1Hpyrazolo[4,3-c]pyridine-3-carboxamide (55). A mixture of compound 87 (132 mg, 0.20 mmol, 1 equiv), imidazole (21 mg, 0.30 mmol, 1.5 equiv), and Cs₂CO₃ (98 mg, 0.30 mmol, 1.5 equiv) in dry DMF (2 mL) was stirred for 2 h at 60 °C. The reaction mixture was cooled and partitioned between EtOAc (15 mL) and water (40 mL). The aqueous phase was extracted with EtOAc (2 \times 10 mL). The combined organic layers were washed with water (2 × 10 mL), brine (2 × 10 mL), dried over anhydrous Na₂SO₄, filtered, and evaporated in vacuo. Purification by FC (DCM/MeOH/7 M solution of NH3 in MeOH 99:0:1 to 94:5:1) gave 77 mg (67%) of the title compound 55 as a white solid. ¹H NMR (500 MHz, CDCl₃): δ 8.27 (d, I = 8.0 Hz, 1H), 8.20 (d, J = 8.2 Hz, 1H), 8.10 (d, J = 8.2 Hz, 1H), 7.88 (d, J =8.0 Hz, 1H), 7.82 (d, J = 8.2 Hz, 1H), 7.55 (dd, J = 6.8 Hz, 1.1 Hz, 1H), 7.53-7.48 (m, 3H), 7.48-7.42 (m, 2H), 7.34 (d, J = 7.8 Hz, 1H), 7.14 (s, 1H), 7.07 (t, J = 5.6 Hz, 1H), 6.93 (s, 1H), 6.73 (d, J =7.8 Hz, 1H), 6.43 (s, 1H), 5.03 (d, J = 5.6 Hz, 2H), 4.21 (t, J = 5.2Hz, 2H), 4.04 (s, 2H), 4.02 (t, J = 5.2 Hz, 2H), 3.98 (s, 3H), 3.92 (s, 2H), 2.05 (t, J = 5.4 Hz, 2H), 1.98 (t, J = 5.4 Hz, 2H). ¹³C NMR (126 MHz, CDCl₃): δ 162.2, 155.4, 141.6, 140.3, 137.0, 133.9, 133.7, 133.4, 131.6, 129.8, 128.8, 128.7, 128.1, 126.9, 126.8, 126.5, 126.1, 126.0, 125.5, 125.5, 125.1, 124.7, 123.7, 122.3, 119.1, 117.2, 103.0, 59.6, 55.6, 50.0, 49.8, 48.4, 46.6, 41.1, 21.2. ESI HRMS (m/z): calcd for $C_{35}H_{35}N_6O_2$ [M + H]⁺, 571.28160; found, 571.28301.

1-(2-(1H-Pyrazol-1-yl)ethyl)-5-((4-methoxynaphthalen-1-yl)methyl)-N-(naphthalen-1-ylmethyl)-4,5,6,7-tetrahydro-1Hpyrazolo[4,3-c]pyridine-3-carboxamide (56). Compound 56 was synthesized by employing the procedure described for compound 55, using intermediate 87 (132 mg, 0.20 mmol, 1 equiv), pyrazole (21 mg, 0.30 mmol, 1.5 equiv), Cs₂CO₃ (98 mg, 0.30 mmol, 1.5 equiv), and dry DMF (2 mL). Purification by FC (DCM/MeOH/7 M solution of NH₃ in MeOH 99:0:1 to 94:5:1) gave 66 mg (57%) of the title compound **56** as a white solid. ¹H NMR (500 MHz, CDCl₃): δ 8.28 (d, J = 7.8 Hz, 1H), 8.21 (d, J = 7.8 Hz, 1H), 8.12 (d, J = 8.2 Hz, 1H), 7.89 (d, J = 7.8 Hz, 1H), 7.84 (d, J = 8.2 Hz, 1H), 7.60-7.42 (m, 7H), 7.32 (d, J = 6.1 Hz, 1H), 7.08 (t, J = 5.1 Hz, 1H), 6.77 (d, 1.5 Hz, 1H), 6.73 (d, J = 7.8 Hz, 1H), 6.06 (br s, 1H), 5.05 (d, J = 5.5Hz, 2H), 4.42 (t, J = 5.5 Hz, 2H), 4.22 (t, J = 5.0 Hz, 2H), 4.00 (s, 5H), 3.89 (s, 2H), 2.58 (br s, 2H), 1.98 (br s, 2H). ¹³C NMR (126 MHz, CDCl₃): δ 162.4, 155.4, 141.5, 140.5, 140.4, 134.0, 133.8, 133.5, 131.7, 130.3, 128.8, 128.7, 127.8, 126.9, 126.7, 126.4, 126.1 (2× C), 126.1, 125.6, 125.1, 124.8, 123.8, 122.3, 117.0, 105.9, 103.0, 59.7, 55.6, 51.9, 50.0, 49.2, 48.5, 41.1, 21.2. ESI HRMS (m/z): calcd for $C_{35}H_{35}N_6O_2$ [M + H]⁺, 571.28160; found, 571.28301.

5-((4-Methoxynaphthalen-1-yl)methyl)-1-(2-morpholinoethyl)-N-(naphthalen-1-ylmethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (57). Compound 57 was synthesized by employing the procedure described for compound 55, using intermediate 87 (132 mg, 0.20 mmol, 1 equiv), morpholine (26 mg, 0.30 mmol, 1.5 equiv), Cs₂CO₃ (98 mg, 0.30 mmol, 1.5 equiv), and dry DMF (2 mL). Purification by FC (DCM/MeOH/7 M solution of NH₂ in MeOH 99:0:1 to 98:1:1) gave 41 mg (35%) of the title compound 57 as a white solid. ¹H NMR (500 MHz, CDCl₃): δ 8.33-8.27 (m, 2H), 8.13 (d, J = 8.1 Hz, 1H), 7.90 (dd, J = 7.7, 1.7Hz, 1H), 7.84 (d, J = 8.2 Hz, 1H), 7.61-7.43 (m, 6H), 7.40 (d, J =7.8 Hz, 1H), 7.07 (t, J = 5.7 Hz, 1H), 6.78 (d, J = 7.8 Hz, 1H), 5.06 (d, J = 5.7 Hz, 2H), 4.10 (s, 2H), 4.03 (s, 3H), 4.02-3.93 (m, 4H),3.64 (t, I = 4.7 Hz, 4H), 2.81 (t, I = 5.7 Hz, 2H), 2.71–2.61 (m, 4H), 2.40 (t, J = 4.7 Hz, 4H). ¹³C NMR (91 MHz, CDCl₃): δ 162.5, 155.3, 140.5, 139.2, 133.9, 133.4, 131.6, 129.8, 128.7, 128.4, 127.7, 126.6, 126.6, 126.4, 126.2, 126.0, 125.9, 125.4, 125.0, 124.7, 123.7, 122.2, 117.5, 102.9, 66.9, 60.0, 58.0, 55.5, 53.8, 50.3, 48.6, 47.0, 40.9, 22.5. ESI HRMS (m/z): calcd for $C_{36}H_{40}N_5O_3$ [M + H]⁺, 590.31257; found, 590.31338.

1-(2-Acetamidoethyl)-5-((4-methoxynaphthalen-1-yl)methyl)-N-(naphthalen-1-ylmethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]-pyridine-3-carboxamide (58). A solution of compound 50 (60 mg, 0.11 mmol, 1.0 equiv) in acetic anhydride (100 mg, 0.99 mmol, 9.0 equiv) was heated at 80 °C for 1 h. The mixture was evaporated in vacuo and the residue was purified by FC (EtOAc/MeOH 99:1 to

95:5) to give 45 mg (73%) of the title compound **58** as a white solid.

¹H NMR (400 MHz, CDCl₃): δ 8.28 (dd, J = 6.6 Hz, 2.0 Hz, 1H), 8.23 (dd, J = 7.1 Hz, 2.0 Hz, 1H), 8.09 (dd, J = 8.7 Hz, 1.3 Hz, 1H), 7.86 (dd, J = 7.1 Hz, 2.3 Hz, 1H), 7.80 (d, J = 8.1 Hz, 1H), 7.55–7.38 (m, 6H), 7.33 (d, J = 7.8 Hz, 1H), 7.08 (t, J = 5.6 Hz, 1H), 6.73 (d, J = 7.8 Hz, 1H), 5.82 (t, J = 5.6 Hz, 1H), 5.01 (d, J = 5.6 Hz, 2H), 4.05 (s, 2H), 4.00 (s, 3H), 3.95–3.88 (m, 4H), 3.48 (q, J = 5.8 Hz, 2H), 2.72 (t, J = 5.5 Hz, 2H), 2.52 (t, J = 5.5 Hz, 2H), 1.75 (s, 3H).

NMR (101 MHz, CDCl₃): δ 170.6, 162.4, 155.4, 141.0, 139.8, 134.0, 133.9, 133.5, 131.6, 128.9, 128.6, 127.9, 126.9, 126.7, 126.4, 126.1, 126.1, 125.6, 125.1, 124.7, 123.7, 122.4, 117.8, 103.0, 60.2, 55.6, 50.2, 48.9, 48.0, 41.1, 39.2, 23.1, 22.4. ESI HRMS (m/z): calcd for $C_{34}H_{36}N_5O_3$ [M + H]⁺, 562.28127; found, 562.28196.

(2-(5-((4-methoxynaphthalen-1-yl)methyl)-3-((naphthalen-1ylmethyl)carbamoyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridin-1-yl)ethyl)glycine (59). The solution of compound 59 (37 mg, 0.06 mmol, 1.0 equiv) and NaOH (12 mg, 0.30 mmol, 5.0 equiv) in MeOH (1 mL)/water (0.1 mL) was stirred at rt for 2 h. The reaction mixture was evaporated in vacuo and the residue was partitioned between DCM (1 mL) and saturated aqueous solution of NH₄Cl. The aqueous phase was washed with DCM (2 × 0.5 mL) and the combined organic phase was dried over anhydrous Na₂SO₄, filtered, and concentrated in vacuo. Purification by automated preparative chromatography (RP-C18, linear gradient from water/MeOH 90:10 to 15:85) gave 20 mg (58%) of the title compound 59 as a white solid. ¹H NMR (500 MHz, DMSO- d_6): δ 8.68 (t, J = 6.2 Hz, 1H), 8.23 (d, J = 8.3 Hz, 1H), 8.18 (dd, J = 12.8 Hz, 8.1 Hz, 2H), 7.92 (d, J= 7.1 Hz, 1H), 7.80 (d, J = 5.0 Hz, 1H), 7.56–7.45 (m, 5H), 7.45– 7.41 (m, 2H), 7.38 (d, J = 7.8 Hz, 1H), 6.90 (d, J = 7.8 Hz, 1H), 4.83 (d, J = 6.2 Hz, 2H), 4.18 (t, J = 6.2 Hz, 2H), 3.99 (s, 2H), 3.96 (s, 2H)3H), 3.58 (s, 2H), 3.25 (s, 2H), 3.17 (t, J = 6.2 Hz, 2H), 2.79 (t, J = 4.3 Hz, 2H), 2.69 (t, J = 4.3 Hz, 2H). 13 C NMR (101 MHz, DMSO d_6): δ 169.1, 161.9, 154.5, 140.2, 139.3, 134.8, 133.2, 132.8, 130.7, 130.5, 128.4, 127.8, 127.2, 126.2, 126.2, 125.9, 125.7, 125.2, 125.1, 125.0, 124.9, 123.5, 121.7, 116.1, 103.4, 59.3, 55.5, 49.7, 49.1 (2× C), 46.8, 46.0, 21.4. 1 C atom overlaps with DMSO-d₆ signal. ESI HRMS (m/z): calcd for $C_{34}H_{36}N_5O_4$ [M + H]⁺, 578.27618; found, 578.27710.

Methyl (2-(5-((4-Methoxynaphthalen-1-yl)methyl)-3-((naphthalen-1-ylmethyl)carbamoyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridin-1-yl)ethyl)glycinate (60). A mixture of compound 87 (70 mg, 0.10 mmol, 1.0 equiv), methyl glycinate hydrochloride (50 mg, 0.40 mmol, 4.0 equiv), and K₂CO₃ (110 mg, 0.80 mmol, 8.0 equiv) in MeCN (2 mL) was stirred at reflux overnight. The reaction mixture was cooled, filtered, and evaporated in vacuo. The residue was partitioned between AcOEt (2 mL) and water (0.5 mL). The layers were separated and the organic phase was washed with brine (1 mL), dried over anhydrous Na2SO4, filtered, and concentrated in vacuo. Purification by FC (EtOAc/MeOH 99:1 to 92:8) gave 88 mg (57%) of the title compound 60 as a white solid. ¹H NMR (500 MHz, DMSO- d_6): δ 8.40 (t, J = 6.2 Hz, 1H), 8.23 (d, J = 8.5 Hz, 1H), 8.17 (dd, J = 8.7 Hz, 1.3 Hz, 2H), 7.94-7.91 (m, 1H), 7.82 (dd, J = 7.1)Hz, 2.3 Hz, 1H), 7.56-7.45 (m, 4H), 7.44-7.40 (m, 2H), 7.38 (d, J =7.8 Hz, 1H), 6.91 (d, J = 7.8 Hz, 1H), 4.82 (d, J = 6.2 Hz, 2H), 4.03 (t, J = 6.2 Hz, 2H), 3.99 (s, 2H), 3.97 (s, 3H), 3.57 (s, 2H), 3.56 (s, 2H), 3.56 (s, 2H), 3.56 (s, 2H), 3.97 (s, 2H), 3.99 (s, 2H), 3.97 (s, 3H), 3.57 (s, 2H), 3.56 (s, 2H), 3.97 (s, 2H), 3.98 (s, 2H), 3.97 (s, 3H), 3.57 (s, 2H), 3.98 (s, 2H), 3.97 (s, 3H), 3.57 (s, 2H), 3.56 (s, 2H), 3.98 (s, 2H), 3.93H), 3.29 (s, 2H), 2.88 (t, J = 6.3 Hz, 2H), 2,78 (t, J = 5.8 Hz, 2H), 2.69 (t, J = 5.8 Hz, 2H). ¹³C NMR (126 MHz, DMSO- d_6): δ 172.6, 162.1, 154.5, 139.9, 139.1, 134.9, 133.2, 132.8, 130.8, 128.5, 127.8, 127.3, 126.3, 126.2, 126.0, 125.7, 125.4, 125.2 (2× C), 125.1, 125.0, 123.5, 121.7, 116.0, 103.4, 59.5, 55.5, 51.3, 49.7, 49.3, 49.3, 48.9, 48.2, 21.7. 1 C atom overlaps with DMSO- d_6 signal. ESI HRMS (m/z): calcd for $C_{35}H_{38}N_5O_4$ [M + H]⁺, 592.29183; found, 592.29275.

1-(2-Hydroxyethyl)-5-((4-methoxynaphthalen-1-yl)methyl)-5-methyl-3-((naphthalen-1-ylmethyl)carbamoyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridin-5-ium lodide (62). To a cooled (ice-salt bath) solution of compound 29 (100 mg, 0.19 mmol, 1.0 equiv) in acetone (0.5 mL) was added MeI (41 mg, 0.288 mmol, 1.5 equiv). The reaction mixture was allowed to reach rt and stirred overnight. The resulting precipitate was filtered off, washed with EtOAc, and further purified by automated preparative chromatography (RP-C18,

linear gradient from water/MeOH 90:10 to 40:60) to give 63 mg (50%) of the title compound 62 as a white solid. ¹H NMR (500 MHz, DMSO- d_6): δ 8.85 (d, J = 6.2 Hz, 1H), 8.32 (d, J = 8.5 Hz, 1H), 8.27 (dd, J = 8.5 Hz, 1.5 Hz, 1H), 8.24 (dd, J = 7.9 Hz, 1.5 Hz, 1H), 7.96 (dd, J = 7.3 Hz, 1.8 Hz, 1H), 7.85 (dd, J = 6.8 Hz, 2.7 Hz, 1H), 7.74 (d, J = 8.1 Hz, 1H), 7.62 - 7.53 (m, 4H), 7.51 - 7.45 (m, 2H), 7.12 (d, s)J = 8.1 Hz, 1H), 5.14 (q, 13.6 Hz, 2H), 4.98 (t, J = 5.6 Hz, 1H), 4.87 (d, J = 6.2 Hz, 2H), $4.\overline{73}$ (q, J = 15.2 Hz, 2H), 4.16 (t, J = 5.2 Hz, 2H), 4.04 (s, 3H), 3.85-3.80 (m, 2H), 3.76 (q, J = 5.4 Hz, 2H), 3.31(dt, 17.0 Hz, J = 5.0 Hz, 1H), 3.12 (dt, 17.0 Hz, J = 7.5 Hz, 1H), 2.96 (br s. 3H). 13 C NMR (75 MHz, DMSO- d_6): δ 161.3, 156.9, 141.1, 135.9, 134.7, 134.7, 133.9, 133.3, 130.8, 128.5, 127.7, 127.4, 126.2, 125.7, 125.7, 125.5, 125.4, 125.2, 123.8, 123.5, 122.3, 115.5, 109.1, 104.0, 62.6, 62.6, 60.0, 56.7, 56.6, 56.0, 52.3, 45.8, 17.7. ESI HRMS (m/z): calcd for $C_{33}H_{35}N_4O_3$ [M + H]⁺, 535.27037; found, 535.27027

1-(2-Hvdroxvethvl)-5-((4-methoxvnaphthalen-1-vl)methvl)-N-(naphthálen-1-ylméthyl)-1,4,5,6-tetrahydropyrrolo[3,4-c]pyrazole-3-carboxamide (63). Compound 63 was synthesized by employing the procedure described for compound 2, using intermediate 94 (226 mg, 0.61 mmol, 1.0 equiv), TEA (62 mg, 0.61 mmol, 1.0 equiv), 4methoxy-1-naphthaldehyde (114 mg, 0.61 mmol, 1.0 equiv), AcOH (37 mg, 0.61 mmol, 1.0 equiv), NaBH(OAc)₃ (195 mg, 0.92 mmol, 1.5 equiv), and dry THF (5 mL). Purification by FC (EtOAc/ MeOH/TEA 99:0:1 to 94:5:1) gave 211 mg (68%) of the title compound 63 as a white solid. ¹H NMR (300 MHz, CDCl₃): δ 8.38– 8.29 (m, 1H), 8.25 (dd, J = 7.6, 1.5 Hz, 1H), 8.12-8.02 (m, 1H), 7.92-7.84 (m, 1H), 7.81 (d, J = 7.8 Hz, 1H), 7.61-7.35 (m, 7H), 7.06 (t, J = 5.7 Hz, 1H), 6.75 (d, J = 7.8 Hz, 1H), 5.01 (d, J = 5.7 Hz, 2H), 4.30 (s, 2H), 4.12-4.06 (s, 2H), 4.02 (s, 3H), 3.88 (td, J = 5.5, 4.9, 2.2 Hz, 2H), 3.84-3.72 (m, 4H), 2.55 (br s, 1H). ¹³C NMR (75 MHz, CDCl₃): δ 161.4, 155.5, 149.0, 139.1, 133.8, 133.6, 133.0, 131.5, 128.7, 128.5, 127.2, 126.8, 126.7, 126.6, 126.2, 126.0, 125.9, 125.4, 125.2, 125.0, 124.1, 123.6, 122.4, 102.8, 61.2, 58.2, 55.5, 53.1, 52.4, 50.7, 41.1. ESI HRMS (m/z): calcd for $C_{31}H_{31}N_4O_3$ $[M + H]^+$, 507.23907; found, 507.23903.

1-(2-Aminoethyl)-5-((4-methoxynaphthalen-1-yl)methyl)-N-(naphthalen-1-ylmethyl)-1,4,5,6-tetrahydropyrrolo[3,4-c]pyrazole-3-carboxamide (64). To a solution of compound 96 (64 mg, 0.12 mmol, 1.0 equiv) in THF (0.5 mL) was added PPh₃ (94 mg, 0.36 mmol, 3 equiv). The reaction mixture was stirred for further 1 h, followed by addition of water (65 mg, 3.60 mmol, 10.0 equiv). The solution was then heated to 50 °C for 4 h and concentrated in vacuo. Purification by FC (DCM/MeOH/7 M solution of NH₃ in MeOH 99:0:1 to 98:1:1) gave 39 mg (64%) of the title compound 64 as a white solid. ¹H NMR (400 MHz, DMSO- d_6): δ 8.47 (t, J = 6.1 Hz, 1H), 8.27 (d, I = 8.0 Hz, 1H), 8.22–8.15 (m, 2H), 7.93 (dd, I = 7.1Hz, 1.9 Hz, 1H), 7.84-7.79 (m, 1H), 7.59-7.48 (m, 5H), 7.44 (dd, J = 5.8 Hz, 2.0 H, 3H), 6.91 (d, J = 8.0 Hz, 1H), 4.85 (d, J = 6.1 Hz, 1Hz)2H), 4.24 (s, 2H), 3.97 (s, 3H), 3.94 (t, I = 6.0 Hz, 2H), 3.81 (s, 2H), 3.79 (s, 2H), 2.85 (t, *J* = 5.8 Hz, 2H). ¹³C NMR (101 MHz, DMSO d_6): δ 161.3, 154.5, 148.4, 138.4, 134.9, 133.2, 132.5, 130.8, 128.4, 127.3, 127.1, 126.6, 126.3, 126.1, 125.7, 125.4, 125.3, 125.2, 125.1, 124.6, 123.6, 123.5, 121.7, 103.4, 57.5, 55.5, 53.8, 51.9, 50.4, 41.6. 1 C atom overlaps with DMSO- d_6 signal ESI HRMS (m/z): calcd for $C_{31}H_{32}N_5O_2[M+H]^+$, 506.25505; found, 506.25515.

1-(2-((1-Hydroxy-2-methylpropan-2-yl)amino)ethyl)-5-((4-methoxynaphthalen-1-yl)methyl)-N-(naphthalen-1-ylmethyl)-1,4,5,6-tetrahydropyrrolo[3,4-c]pyrazole-3-carboxamide (65). A mixture of compound 95 (57 mg, 0.09 mmol, 1.0 equiv), 2-amino-2-methyl-1-propanol (40 mg, 0.45 mmol, 5 equiv), and K_2CO_3 (30 mg, 0.22 mmol, 2.5 equiv) in CH₃CN (0.5 mL) was stirred at reflux for 8 h. The reaction mixture was cooled to rt and concentrated in vacuo. The residue was purified by FC (DCM/MeOH/7 M NH₃ in MeOH 99:0:1 to 94:5:1) to yield 40 mg (77%) of 65 as a white solid. 1 H NMR (400 MHz, DMSO- d_6): δ 8.45 (t, J = 6.1 Hz, 1H), 8.27 (d, J = 7.9 Hz, 1H), 8.18 (d, J = 8.3 Hz, 2H), 7.93 (dd, J = 6.8 Hz, 2.0 Hz, 1H), 7.82 (dd, J = 6.1 Hz, 3.1 Hz, 1H), 7.58–7.49 (m, 4H), 7.46–7.41 (m, 3H), 6.91 (d, J = 7.9 Hz, 1H), 4.84 (d, J = 6.1 Hz, 2H), 4.47 (br s, 1H), 4.24 (s, 2H), 4.00–3.97 (m, 5H), 3.82 (s, 2H), 3.78 (s,

2H), 3.09 (s, 2H), 2.77 (t, J=6.2 Hz, 2H), 0.84 (s, 6H). 13 C NMR (101 MHz, DMSO- d_6): δ 161.3, 154.5, 148.4, 138.2, 134.8, 133.2, 132.5, 130.8, 128.4, 127.3, 127.1, 126.6, 126.2, 126.1, 125.7, 125.4, 125.3, 125.2, 125.1, 124.6, 123.6, 123.5, 121.7, 103.4, 68.0, 57.4, 55.5, 53.3, 52.3, 51.8, 50.6, 41.6, 23.6 (2× C). 1 C atom overlaps with DMSO- d_6 signal ESI HRMS (m/z): calcd for $C_{35}H_{40}N_5O_3$ [M + H]⁺, 578.31257; found, 578.31258.

5-(tert-Butyl) 3-Ethyl 1,4,6,7-tetrahydro-5H-pyrazolo[4,3-c]pyridine-3,5-dicarboxylate (66). To a stirred solution of LHMDS (25.0 mL of 1 M solution in THF, 25.0 mmol, 1.0 equiv) in dry THF (50 mL) at -78 °C was added a solution of 1-boc-4-piperidinone (5.00 g, 25.0 mmol, 1.0 equiv) in dry THF (20 mL), dropwise. The reaction mixture was stirred for further 30 min at -78 °C, followed by a dropwise addition of a diethyl oxalate (3.92 g, 25.0 mmol, 1.0 equiv). The mixture was then warmed to rt and stirred for further 30 min. Subsequently, EtOH (40 mL), AcOH (10 mL), and hydrazine monohydrate (1.50 mL of 80% aqueous solution, 25 mmol, 1.0 equiv) were added. A slightly exothermic reaction was observed. The reaction mixture was then heated to reflux for 1 h, cooled to rt, and concentrated in vacuo. The residue was partitioned between EtOAc (100 mL) and saturated aqueous solution of NaHCO₃ (100 mL). The aqueous phase was extracted with EtOAc (2 × 25 mL). The combined organic extracts were sequentially washed with water (25 mL), 1 M aqueous solution of citric acid (50 mL), water (25 mL), and brine (25 mL), dried over anhydrous Na₂SO₄, filtered, and concentrated in vacuo. The residue was purified by FC (hexane/ EtOAc 2:1 to 0:1) to yield 4.35 g (59%) of the target compound 66 as a white solid. ¹H NMR (500 MHz, DMSO- d_6): δ 13.70 (br s, 0.3 H), 13.30 (br s, 0.7H), 4.49 (br s, 2H), 4.27 (br s, 2H), 3.59 (t, J = 5.8Hz, 2H), 2.67 (bst, J = 5.8 Hz, 2H), 1.42 (s, 9H), 1.29 (t, J = 7.1 Hz, 3H). ¹³C NMR (126 MHz, DMSO- d_6): δ 162.7 (br s), 159.5 (br s), 154.6, 147.6 (br s), 138.8 (br s), 128.5 (br s), 117.4 (br s), 115.9 (br s), 79.6, 61.0 (br s), 60.4 (br s), 41.6 (br s), 41.0 (br s), 28.5, 23.5 (br s), 21.4 (br s), 14.6 (br s). ESI HRMS (m/z): calcd for $C_{14}H_{22}N_3O_4$ [M + H]⁺, 296.16048; found, 296.16027.

5-(tert-Butyl) 3-Ethyl 1-Methyl-1,4,6,7-tetrahydro-5H-pyrazolo-[4,3-c]pyridine-3,5-dicarboxylate (67) and 5-(tert-Butyl) 3-Ethyl 2-Methyl-1,4,6,7-tetrahydro-5H-pyrazolo[4,3-c]pyridine-3,5-dicarboxylate (68). Intermediate 66 (2.30 g, 7.80 mmol, 1.0 equiv) was dissolved in dry DMF (20 mL). $Cs_2\tilde{C}O_3$ (2.54 g, 7.80 mmol, 1.0 equiv) was added followed by MeI (3.33 g, 23.4 mmol, 3.0 equiv). The mixture was stirred at rt overnight and partitioned between water (100 mL) and EtOAc (50 mL). The layers were separated and the aqueous phase was extracted with EtOAc (2 × 15 mL). The combined organic extracts were washed with water $(3 \times 25 \text{ mL})$ saturated aqueous solution of NH₄Cl (25 mL), dried over anhydrous Na2SO4, filtered, and concentrated in vacuo. The residue was purified by FC (hexane/EtOAc 3:1 to 0:1) to yield 1.38 g (57%) of slower eluting 67 as a pale yellow solid and 0.85 g (35%) of faster eluting N-2 regioisomer 68 as yellow oil that solidified upon storing in the cold. 67: ¹H NMR (250 MHz, CDCl₃): δ 4.60 (s, 2H), 4.38 (q, J = 7.1 Hz, 2H), 3.81 (s, 3H), 3.71 (t, J = 5.7 Hz, 2H), 2.67 (t, J = 5.8 Hz, 2H), 1.47 (s, 9H), 1.38 (t, J = 7.1 Hz, 3H). ¹³C NMR (63 MHz, CDCl₃): δ 159.6, 155.4, 144.9, 136.9, 116.6, 85.2, 60.0, 41.5, 40.0, 36.4, 28.4, 21.8, 14.4. ESI HRMS (m/z): calcd for $C_{15}H_{24}N_3O_4$ $[M + H]^+$, 310.17613; found, 310.17650. **68** 1 H NMR (250 MHz, CDCl₃): δ 4.59 (s, 2H), 4.33 (q, J = 7.1 Hz, 2H), 4.12 (s, 3H), 3.67 (t, J = 5.8 Hz, 2H), 2.72 (t, J = 5.8 Hz, 2H), 1.47 (s, 9H), 1.37 (t, J = 7.1 Hz, 3H). 13 C NMR (63 MHz, CDCl₃): δ 159.9, 155.0, 145.7 (br s), 127.7 (br s), 119.0, 79.9, 60.9, 41.7 (br s), 40.4 (br s), 35.5, 27.4, 22.9 (br s), 14.4. ESI HRMS (m/z) calcd for $C_{15}H_{24}N_3O_4$ [M + H]⁺, 310.17613; found, 310.17625.

tert-Butyl 3-(Benzylcarbamoyl)-2-methyl-2,4,6,7-tetrahydro-5H-pyrazolo[4,3-c]pyridine-5-carboxylate (69). A solution of compound 68 (309 mg, 1.00 mmol, 1.0 equiv), benzylamine (128 mg, 1.00 mmol, 1.2 equiv), and TBD (44 mg, 0.30 mmol, 0.3 equiv) in dry THF (2 mL) was stirred at 60 °C for 12 h. The mixture was concentrated in vacuo and the residue was purified by FC (hexane/EtOAc 2:1 to 1:2) to yield 295 mg (80%) of the title compound 69 as a pale yellow solid. $^1\mathrm{H}$ NMR (500 MHz, CDCl3): δ 7.44–7.30 (m,

5H), 5.96 (br s, 0.6H), 5.79 (br s, 0.4H), 4.68–4.46 (m, 4H), 4.12 (s, 3H), 3.67 (m, t, J=5.5 Hz, 2H), 2.75 (t, J=5.7 Hz, 2H), 1.48 (s, 9H). 13 C NMR (91 MHz, CDCl₃): δ 160.0, 155.0 (br s), 145.2 (br s), 137.6, 131.0, 128.9, 127.8, 127.7, 113.3, 80.4, 43.7, 41.9 (br s), 41.0 (br s), 39.2, 28.4, 23.5 (br s). ESI HRMS (m/z): calcd for $C_{20}H_{27}N_4O_3$ [M + H]⁺, 371.20777; found, 371.20809.

tert-Butyl 3-(Benzylcarbamoyl)-1,4,6,7-tetrahydro-5H-pyrazolo-[4,3-c]pyridine-5-carboxylate (70). Compound 70 was synthesized by employing the procedure described for compound 69, using intermediate 66 (2.00 g, 6.78 mmol, 1.0 equiv), benzylamine (0.87 g, 8.14 mmol, 1.2 equiv), TBD (0.34 g, 2.44 mmol, 0.3 equiv), and dry THF (10 mL). Purification by FC (hexane/EtOAc 1:1 to 0:1) gave 2.15 g (89%) of the title compound 70 as a white solid. ¹H NMR (500 MHz, DMSO- d_6): δ 13.04 (br s, 1H), 8.65 (br s, 1H), 7.35–7.27 (m, 4H), 7.23 (td, J = 6.0, 2.7 Hz, 1H), 4.50 (br s, 2H), 4.40 (d, J = 6.4 Hz, 2H), 3.59 (t, J = 5.7 Hz, 2H), 2.68 (t, J = 5.7 Hz, 2H), 1.42 (s, 9H). ¹³C NMR (126 MHz, DMSO- d_6): δ 162.8 (br s), 154.6 (br s), 141.5 (br s), 140.5 (br s), 138.7 (br s), 128.7, 127.7, 127.1, 114.2 (br s), 79.5 (br s), 42.1 (br s), 41.7 (br s), 41.1 (br s), 28.5, 21.6 (br s). ESI HRMS (m/z): calcd for $C_{19}H_{25}N_3O_4$ [M + H]⁺, 357.19212; found, 357.19220.

tert-Butyl 3-(Benzylcarbamoyl)-1-(2-ethoxy-2-oxoethyl)-1,4,6,7tetrahydro-5H-pyrazolo[4,3-c]pyridine-5-carboxylate (71). Compound 71 was synthesized by employing the procedure described for compounds 67 and 68, using intermediate 66 (1.50 g, 4.21 mmol, 1.0 equiv), Cs₂CO₃ (1.37 g, 4.21 mmol, 1.0 equiv), ethyl 2bromoacetate (703 mg, 4.21 mmol, 1.0 equiv), and dry DMF (20 mL). Purification by FC (hexane/EtOAc 3:1 to 1:1) gave 1.65 g (89%) of the title 71 as a white solid. ¹H NMR (500 MHz, CDCl₃): δ 7.40-7.31 (m, 4H), 7.32-7.32 (m, 1H), 7.11 (br s, 1H), 4.78 (s, 2H), 4.74 (s, 2H), 4.60 (d, J = 6.0 Hz, 2H), 4.25 (q, J = 7.1 Hz, 2H), 3.74 (t, J = 5.8 Hz, 2H), 2.65 (t, J = 5.7 Hz, 2H), 1.50 (s, 9H), 1.30 (t, I = 7.1 Hz, 3H). ¹³C NMR (91 MHz, CDCl₃): δ 167.1, 162.0 (br s), 141.4 (br s), 139.9 (br s), 138.3 (br s), 128.7, 127.9, 127.4, 116.6 (br s), 80.2 (br s), 62.2, 50.7, 42.9, 41.3 (br s), 39.7 (br s), 28.5, 21.8 (br s), 14.1. ESI HRMS (m/z): calcd for $C_{23}H_{31}N_4O_5$ $[M + H]^+$, 443.22890; found, 443.22895.

5-(tert-Butyl) 1-Ethyl 3-(benzylcarbamoyl)-6,7-dihydro-1Hpyrazolo[4,3-c]pyridine-1,5(4H)-dicarboxylate (72). To a stirred, cooled (ice-salt bath) solution of intermediate 66 (1.00 g, 2.81 mmol, 1.0 equiv) and DIPEA (0.72 g, 5.62 mmol, 2.0 equiv) in dry THF (10 mL) was added ethyl chloroformate (0.31 g, 2.81 mmol, 1.0 equiv), dropwise. The mixture was then warmed to rt overnight and concentrated in vacuo. The residue was purified by FC (hexane/ EtOAc 3:1 to 2:1) to yield 1.09 g (91%) of the target compound 72 as a white solid. ¹H NMR (360 MHz, CDCl₃): δ 7.44–7.29 (m, 5H), 4.71 (s, 2H), 4.61 (d, J = 6.1 Hz, 2H), 4.51 (q, J = 7.2 Hz, 2H), 3.71(t, J = 5.8 Hz, 2H), 3.06 (t, J = 5.8 Hz, 2H), 1.50 (s, 9H), 1.48-1.43(m, 4H). 13 C NMR (91 MHz, CDCl₃): δ 161.2 (br s), 154.9 (br s), 149.5, 145.0 (br s), 142.9 (br s), 137.8, 128.7, 128.0, 127.6, 119.3 (br s), 80.3 (br s), 64.9, 43.1, 41.0 (br s), 39.8 (br s), 28.4, 25.2, 14.2. ESI HRMS (m/z): calcd for $C_{22}H_{29}N_4O_5$ [M + H]⁺, 429.21325; found, 429,21322.

5-(tert-Butyl) 3-Ethyl 1-(2-Hydroxyethyl)-1,4,6,7-tetrahydro-5Hpyrazolo[4,3-c]pyridine-3,5-dicarboxylate (73) and 5-(tert-Butyl) 3-Ethyl 2-(2-Hydroxyethyl)-2,4,6,7-tetrahydro-5H-pyrazolo[4,3-c]pyridine-3,5-dicarboxylate (74). Compounds 73 and 74 were synthesized by employing the procedure described for compound 66, using a solution of LHMDS (37.5 mL of 1 M solution in THF, 37.5 mmol, 1.0 equiv) in dry THF (75 mL), solution of 1-boc-4piperidinone (7.50 g, 37.5 mmol, 1.0 equiv) in dry THF (50 mL), diethyl oxalate (5.88 g, 37.5 mmol, 1.0 equiv), and 2-hydroxyethylhydrazine (2.85 g, 37.5 mmol, 1.0 equiv). Recrystallization of the reaction mixture from hexane/DCM gave 6.38 g (50%) of the title compound 73 as a white solid. The filtrate was concentrated in vacuo and the residue was purified by FC (hexane/EtOAc 5:1 to 0:1, then EtOAc/MeOH 95:5) to give further 1.21 g (10%) of 73 (total yield: 7.59 g, 60%) and 2.00 g (16%) of the N-2 regioisomer 74 as yellow oil. 73 ¹H NMR (360 MHz, CDCl₃): δ 4.61 (s, 2H), 4.39 (q, J = 7.1Hz, 2H), 4.22-4.13 (m, 2H), 4.02 (q, J = 5.0 Hz, 2H), 3.70 (t, J = 5.2

Hz, 2H), 2.82–2.62 (m, 3H), 1.49 (s, 9H), 1.37 (t, J = 7.1 Hz, 3H). 13 C NMR (91 MHz, CDCl₃): δ 162.4, 155.0 (br s), 139.4 (br s), 138.5 (br s), 117.3, 80.2, 61.4, 60.8, 51.4, 41.5 (br s), 39.8 (br s), 28.4, 22.0, 14.4. ESI HRMS (m/z): calcd for C₁₆H₂₆N₃O₅ [M + H]⁺, 340.18670; found, 340.18670. 74 1 H NMR (360 MHz, CDCl₃): δ 4.70 (t, J = 4.7, 2H), 4.63 (br s, 2H), 4.35 (q, J = 7.1 Hz, 2H), 4.00 (t, J = 4.9 Hz, 2H), 3.70 (br s, 2H), 3.00 (br s, 1H), 2.76 (t, J = 6.2 Hz, 1H), 1.50 (s, 9H), 1.40 (t, J = 7.1 Hz, 3H). 13 C NMR (91 MHz, CDCl₃): δ 160.2, 155.0, 146.6 (br s), 127.9 (br s), 119.3, 80.0, 62.2, 61.2, 53.4, 53.1, 42.0 (br s), 28.4, 23.4 (br s), 14.3 (br s). ESI HRMS (m/z): calcd for C₁₆H₂₆N₃O₅ [M + H]⁺, 340.18670; found, 340.18669.

Ethyl 5-((1H-Indol-3-yl)methyl)-1-(2-hydroxyethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxylate (75). Compound 75 was synthesized by employing the procedure described for compound 2, using intermediate 73 (1.30 g, 3.83 mmol, 1.0 equiv), 4 M solution of HCl in 1,4-dioxane (15 mL), TEA (0.39 g, 3.83 mmol, 1.0 equiv), indole-3-carboxaldehyde (0.55 g, 3.83 mmol, 1.0 equiv), AcOH (0.23 g, 3.83 mmol, 1.0 equiv), NaBH(OAc)₃ (1.22 g, 5.74 mmol, 1.5 equiv), and dry THF (20 mL). Purification by FC (EtOAc/MeOH/TEA 99:0:1 to 94:5:1) gave 1.02 g (72%) of the title compound 75 as a yellow solid. ¹H NMR (300 MHz, CDCl₃): δ 8.64 (br s, 1H), 7.71 (d, I = 7.8 Hz, 1H), 7.36 (dt, I = 8.0, 1.1 Hz, 1H), 7.30-7.07 (m, 3H), 4.34 (q, J = 7.2 Hz, 2H), 4.12 (t, J = 5.0 Hz, 2H), 4.00 (s, 2H), 3.94 (t, J = 5.0 Hz, 2H), 3.84 (s, 2H), 3.46 (br s, 2H),2.87 (t, I = 5.8 Hz, 2H), 2.72 (t, I = 5.9 Hz, 2H), 1.32 (t, I = 7.2 Hz, 3H). 13 C NMR (75 MHz, CDCl₃): δ 162.5, 139.4, 138.5, 136.2, 127.9, 124.4, 122.1, 119.7, 119.1, 118.3, 111.4, 111.3, 61.3, 60.7, 52.0, 51.5, 49.6, 48.7, 22.0, 14.3. ESI HRMS (m/z): calcd for $C_{20}H_{25}N_4O_3$ $[M + H]^+$, 369.19212; found, 369.19203.

tert-Butyl 3-(Benzylcarbamoyl)-1-(2-hydroxyethyl)-1,4,6,7-tetra-hydro-5H-pyrazolo[4,3-c]pyridine-5-carboxylate (76). Compound 76 was synthesized by employing the procedure described for compound 69, using intermediate 73 (2.30 mg, 8.14 mmol, 1.2 equiv), benzylamine (0.86 g, 6.78 mmol, 1.2 equiv), TBD (0.28 g, 2.03 mmol, 0.3 equiv), and dry THF (5 mL). Purification by FC (hexane/EtOAc 1:1 to 0:1) gave 1.94 g (72%) of the title compound 76 as a white solid. 1 H NMR (360 MHz, CDCl₃): δ 7.37–7.21 (m, 5H), 7.12 (br s, 1H), 4.70 (s, 2H), 4.59 (d, J = 6.1 Hz, 2H), 4.13–4.07 (m, 2H), 4.05–3.92 (m, 2H), 3.72 (t, J = 5.7 Hz, 2H), 2.81 (t, J = 5.8 Hz, 1H), 2.72 (t, J = 5.8 Hz, 2H), 1.49 (s, 9H). 13 C NMR (91 MHz, CDCl₃): δ 162.1, 155.1, 140.9, 139.6, 138.4, 128.7, 127.9, 127.4, 116.1, 80.2, 61.4, 51.1, 42.9, 41.3, 40.0, 28.5, 22.1. ESI HRMS (m/z): calcd for $C_{21}H_{29}N_4O_4$ [M + H] $^+$, 401.21833; found, 401.21830.

The free amine hydrochloride used for synthesis of derivatives 15-19 was obtained by treatment of compound 76 with 4 M solution of HCl in 1,4-dioxane at rt for 4 h, followed by evaporation of the solvent in vacuo.

1-(2-Hydroxyethyl)-N-(naphthalen-1-ylmethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (78). To a stirred solution of compound 77 (2.10 g, 0.47 mmol) in DCM (50 mL) was added slowly 4 M solution of HCl in 1,4-dioxane (20 mL). The mixture was stirred for 3 h at rt, followed by evaporation of the solvent in vacuo to give 1.80 g (quantitative) of the title compound 78 as a pale yellow solid. 1H NMR (300 MHz, DMSO- d_6): δ 9.53 (br s, 2H), 8.73 (t, J = 6.1 Hz, 1H), 8.22 (d, J = 8.7 Hz, 1H), 7.95 (d, J = 8.7 Hz 8.9 Hz, 1H), 7.84 (dd, J = 6.0, 3.4 Hz, 1H), 7.55 (p, J = 7.2, Hz, 2H), 7.49-7.40 (m, 2H), 4.89 (d, J = 6.1 Hz, 2H), 4.25 (br s, 2H), 4.13 (t, J = 5.2 Hz, 2H), 3.85 (br s, 2H), 3.73 (d, J = 10.4 Hz, 2H), 3.60 (br s, 1H), 3.36 (br s, 2H), 3.02 (t, J = 6.0 Hz, 2H). ¹³C NMR (75 MHz, DMSO- d_6): δ 162.0, 141.2, 137.8, 135.2, 133.7, 131.3, 129.0, 127.9, 126.7, 126.2, 125.9, 125.8, 124.0, 110.7, 60.5, 52.3, 19.2. ESI HRMS (m/z): calcd for $C_{20}H_{23}N_4O_2$ [M + H]⁺, 351.18155; found, 351.18146.

tert-Butyl 2-(2-Hydroxyethyl)-3-((naphthalen-1-ylmethyl)-carbamoyl)-2,4,6,7-tetrahydro-5H-pyrazolo[4,3-c]pyridine-5-carboxylate (79). Compound 79 was synthesized employing the procedure described for compound 69, using intermediate 74 (600 mg, 1.77 mmol, 1.0 equiv), 1-naphthalenemethylamine (333 mg, 2.12

mmol, 1.2 equiv), TBD (74 mg, 0.53 mmol, 0.3 equiv), and dry THF (5 mL). Purification by FC (EtOAc) gave 670 mg (84%) of the title compound 79 as a pale yellow solid. 1 H NMR (500 MHz, CDCl₃): δ 8.07 (d, J = 8.4 Hz, 1H), 7.91 (d, J = 8.1 Hz, 1H), 7.86 (d, J = 8.2 Hz, 1H), 7.59 (ddd, J = 8.4, 6.8, 1.4 Hz, 1H), 7.57–7.54 (m, 1H), 7.54–7.49 (m, 1H), 7.46 (dd, J = 8.2, 6.9 Hz, 1H), 6.73 (br s, 0.5H), 6.53 (br s, 0.5H), 5.07 (d, J = 5.5 Hz, 2H), 4.64–4.33 (m, 4H), 3.99 (t, J = 4.9 Hz, 2H), 3.65 (t, J = 5.7 Hz, 2H), 2.72 (t, J = 5.9 Hz, 2H), 1.54–1.29 (m, 9H). 13 C NMR (91 MHz, CDCl₃): δ 160.0, 154.8, 134.0, 132.6, 132.4, 131.2, 129.0, 126.9, 126.8, 126.1, 125.4, 123.2, 80.3, 62.3, 52.8, 42.0, 40.6, 28.3, 23.4. ESI HRMS (m/z): calcd for $C_{25}H_{31}N_4O_4$ [M + H] $^+$, 451.23398; found, 451.23398.

2-(2-Hydroxyethyl)-5-((4-methoxynaphthalen-1-yl)methyl)-N-(naphthalen-1-ylmethyl)-4,5,6,7-tetrahydro-2H-pyrazolo[4,3-c]pyridine-3-carboxamide (80). Compound 80 was synthesized by employing the procedure described for compound 2, using intermediate 79 (279 mg, 0.62 mmol, 1.0 equiv), 4 M solution of HCl in 1,4-dioxane (5 mL), TEA (63 mg, 0.62 mmol, 1.0 equiv), 4methoxy-1-naphthaldehyde (115 mg, 0.62 mmol, 1.0 equiv), AcOH (37 mg, 0.62 mmol, 1.0 equiv), NaBH(OAc)₃ (197 mg, 0.93 mmol, 1.5 equiv), and dry THF (5 mL). Purification by FC (EtOAc/ MeOH/TEA 99:0:1 to 94:5:1) gave 131 mg (40%) of the title compound 80 as a white solid. ¹H NMR (300 MHz, DMSO- d_6): δ 8.74 (t, J = 5.6 Hz, 1H), 8.21-8.15 (m, 2H), 8.13-8.07 (m, 1H), 7.99-7.92 (m, 1H), 7.86 (t, J = 4.8 Hz, 1H), 7.54-7.47 (m, 4H), 7.41 (d, J = 5.4 Hz, 2H), 7.28 (d, J = 7.8 Hz, 1H), 6.84 (d, J = 7.8 Hz, 1H), 5.05 (t, J = 5.1 Hz, 1H), 4.87 (d, J = 5.6 Hz, 2H), 4.28 (t, J = 5.8Hz, 2H), 3.97 (s, 3H), 3.89 (s, 2H), 3.66 (q, J = 5.6 Hz, 2H), 3.55 (s, 2H), 2.73 (t, I = 5.6 Hz, 2H), 2.58 (t, I = 5.2 Hz, 2H). ¹³C NMR (75 MHz, DMSO- d_6): δ 160.1, 154.5, 145.2, 134.0, 133.3, 132.8, 132.7, 130.7, 128.5, 127.6, 127.6, 126.2, 126.2, 125.8, 125.8, 125.3, 125.3, 125.2, 125.0, 124.9, 123.4, 121.7, 115.3, 103.3, 60.5, 59.5, 55.5, 52.3, 49.8, 49.0, 40.6, 23.3. ESI HRMS (m/z): calcd for $C_{32}H_{33}N_4O_3$ [M + H]+, 521.25472; found, 521.25468.

tert-Butyl 3-((Naphthalen-1-ylmethyl)carbamoyl)-1,4,6,7-tetrahydro-5H-pyrazolo[4,3-c]pyridine-5-carboxylate (81). Compound 81 was synthesized by employing the procedure described for compound 69, using intermediate 66 (720 g, 2.44 mmol, 1.0 equiv), 1-naphthalenemethylamine (460 mg, 2.93 mmol, 1.2 equiv), TBD (102 mg, 0.73 mmol, 0.3 equiv), and dry THF (10 mL). Purification by recrystallization from EtOAc/MeOH gave 610 mg (62%) of the title compound 81 as a white solid. ¹H NMR (500 MHz, DMSO- d_6): δ 13.02 (br s, 1H), 8.61 (br s, 1H), 8.23 (d, J = 8.4 Hz, 1H), 7.96 (d, J = 7.7 Hz, 1H), 7.86–7.82 (m, 1H), 7.62–7.52 (m, 2H), 7.47 (d, J = 5.5 Hz, 2H), 4.89 (d, J = 6.3 Hz, 2H), 4.52 (s, 2H), 3.60 (t, J = 5.8 Hz, 2H), 2.69 (t, J = 5.8 Hz, 2H), 1.42 (s, 9H). ¹³C NMR (91 MHz, DMSO- d_6): δ 162.8, 154.6, 138.8, 135.4, 133.7, 131.3, 128.9, 127.8, 126.6, 126.2, 125.9, 125.6, 123.9, 114.2, 79.5, 28.6. ESI HRMS (m/z): calcd for $C_{23}H_{27}N_4O_3$ [M + H] $^+$, 407.20777; found, 407.20791.

5-((4-Methoxynaphthalen-1-yl)methyl)-N-(naphthalen-1-ylmethyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (82). Compound 82 was synthesized by employing the procedure described for compound 2, using intermediate 81 (379 mg, 0.93 mmol, 1.0 equiv), 4 M solution of HCl in 1,4-dioxane (5 mL), TEA (94 mg, 0.93 mmol, 1.0 equiv), 4-methoxy-1naphthaldehyde (172 mg, 0.93 mmol, 1.0 equiv), AcOH (56 mg, 0.62 mmol, 1.0 equiv), NaBH(OAc)₃ (296 mg, 1.40 mmol, 1.5 equiv), and dry THF (7 mL). Purification by recrystallization from EtOAc/MeOH gave 245 mg (55%) of the title compound 82 as a white solid. ¹H NMR (400 MHz, DMSO- d_6): δ 12.90 (s, 1H), 8.51 (t, J = 6.1 Hz, 1H), 8.25 (d, J = 7.7 Hz, 1H), 8.21–8.14 (m, 2H), 7.93 (dd, J = 6.2, 3.0 Hz, 1H), 7.81 (d, J = 6.6 Hz, 1H), 7.58-7.35 (m, J = 6.6 Hz, 1H)7H), 6.91 (d, J = 7.9 Hz, 1H), 4.82 (d, J = 6.0, 2H), 4.00 (s, 2H), 3.97 (s, 3H), 3.59 (s, 2H), 2.77 (t, J = 5.8 Hz, 2H), 2.68 (s, J = 5.8 Hz, 2H). 13 C NMR (101 MHz, DMSO- d_6): δ 162.9, 155.0, 141.5, 139.0, 135.4, 133.7, 133.3, 131.2, 128.9, 128.2, 127.7, 126.7, 126.6, 126.5, 126.1, 125.8, 125.7, 125.6, 125.5, 125.4, 123.9, 122.1, 115.6, 103.9, 60.0, 56.0, 49.8, 46.0, 22.0. ESI HRMS (m/z): calcd for $C_{30}H_{29}N_4O_2$ [M + H]⁺, 477.22850; found, 477.22853.

1-(5-((4-Methoxynaphthalen-1-yl)methyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridin-3-yl)-N-(naphthalen-1-ylmethyl)methanamine (83). To a stirred, cooled (ice-salt bath) solution of compound 82 (600 mg, 1.26 mmol, 1.0 equiv) in dry THF (15 mL) was added LiAlH₄ (3.15 mL of a 2 M solution in THF, 6.30 mmol, 5.0 equiv). The reaction mixture was allowed to reach rt followed by refluxing overnight. The solution was cooled (ice-salt bath) and water (0.5 mL) was added slowly. The mixture was treated with 15% aqueous solution of NaOH (0.5 mL) and water (1.5 mL), followed by MgSO₄. Next, DCM (5 mL) and MeOH (5 mL) were added, followed by filtration of the mixture and evaporation in vacuo. Purification by FC (DCM/MeOH/7 M solution of NH₃ in MeOH 94:5:1) gave 380 mg (65%) of the title compound 83 as a white solid. ¹H NMR (400 MHz, DMSO- d_6): δ 12.21 (br s, 0.5H), 12.13 (br s, 0.5H), 8.23 (d, I = 7.8 Hz, 1H), 8.18 (dd, I = 8.2, 1.7 Hz, 1H), 8.10(d, I = 8.0 Hz, 1H), 7.97 - 7.86 (m, 1H), 7.80 (d, I = 8.0 Hz, 1H),7.58-7.40 (m, 4H), 7.39 (t, J = 7.5 Hz, 1H), 7.34 (d, J = 8.0 Hz, 1H), 6.87 (d, J = 7.8 Hz, 1H), 4.07 (s, 2H), 3.96 (s, 3H), 3.93 (s, 2H), 3.66(s, 2H), 3.44 (s, 2H), 2.73 (t, J = 5.8 Hz, 2H), 2.59 (t, J = 5.7 Hz, 2H),2.37 (br s, 1H). ¹³C NMR (126 MHz, DMSO- d_6): δ 154.9, 146.4 (br s), 136.5 (br s), 133.8, 133.3, 131.9, 128.8, 128.2, 127.6, 126.6, 126.6, 126.2, 126.1, 126.0, 125.8, 125.7, 125.5, 125.4, 124.6, 122.1, 112.1 (br s), 103.8, 61.2, 60.3 (br s), 56.0, 50.5, 49.4, 45.9 (br s), 43.0 (br s), 29.7. ESI HRMS (m/z): calcd for $C_{30}H_{31}N_4O [M + H]^+$, 463.24924; found, 463.25043.

tert-Butyl 3-Carbamoyl-1-(2-hydroxyethyl)-1,4,6,7-tetrahydro-5H-pyrazolo[4,3-c]pyridine-5-carboxylate (84). The solution of compound 73 (1.50 g, 4.42 mmol) in 7 M methanolic solution of NH₃ (20 mL) was refluxed in a sealed pressure tube for 2 days. The mixture was cooled and evaporated in vacuo. The residue was heated with diethyl ether and the resulting solid was filtered and dried in vacuo to give 1.36 g (quantitative) of the title compound 84 as a white solid. 1 H NMR (400 MHz, DMSO- d_6): δ 7.34 (s, 1H), 7.16 (s, 1H), 4.91 (t, J = 5.4 Hz, 1H), 4.48 (s, 2H), 4.06 (t, J = 5.4 Hz, 2H), 3.72 (q, J = 5.4 Hz, 2H), 3.59 (t, J = 5.7 Hz, 2H), 2.72 (t, J = 5.7 Hz, 2H), 1.42 (s, 9H). 13 C NMR (101 MHz, DMSO- d_6): δ 164.4, 154.6, 140.8, 139.7, 115.0, 79.5, 60.6, 52.0, 41.7, 41.1, 28.6, 22.0.

1-(2-Hydroxyethyl)-5-((4-methoxynaphthalen-1-yl)methyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridine-3-carboxamide (**85**). Compound 85 was synthesized by employing the procedure described for compound 2, using intermediate 84 (728 mg, 2.35 mmol, 1.0 equiv), 4 M solution of HCl in 1,4-dioxane (5 mL), TEA (237 mg, 2.35 mmol, 1.0 equiv), 4-methoxy-1-naphthtaldehyde (437 mg, 2.35 mmol, 1.0 equiv), AcOH (141 mg, 2.35 mmol, 1.0 equiv), NaBH(OAc)₃ (747 mg, 3.52 mmol, 1.5 equiv), and dry THF (15 mL). Purification by FC (DCM/MeOH 99:1 to 95:5) gave 557 mg (62%) of the title compound 85 as a white solid. ¹H NMR (400 MHz, CDCl₃): δ 8.30 (dd, J = 8.2, 1.6 Hz, 1H), 8.23 (dd, J = 7.8, 2.0 Hz, 1H), 7.56-7.42 (m, 2H), 7.37 (d, J = 7.8 Hz, 1H), 6.76 (d, J = 7.8Hz, 1H), 6.62 (br s, 1H), 5.45 (br s, 1H), 4.09-4.02 (m, 4H), 4.02 (s, 3H), 3.95 (dd, *J* = 5.7, 4.0 Hz, 2H), 3.88 (s, 2H), 2.91 (br s, 1H), 2.78 (t, J = 5.7 Hz, 2H), 2.63 (t, J = 5.8 Hz, 2H). ¹³C NMR (101 MHz, $CDCl_3$): δ 164.5, 155.3, 140.2, 139.9, 133.4, 127.7, 126.4, 126.0, 125.0, 124.6, 122.3, 117.8, 102.9, 76.7, 61.3, 59.8, 55.5, 51.0, 50.1, 48.5, 22.3. ESI HRMS (m/z): calcd for $C_{21}H_{25}N_4O_3$ $[M + H]^+$, 381.19212; found, 381.19212.

2-(3-(Aminomethyl)-5-((4-methoxynaphthalen-1-yl)methyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3-c]pyridin-1-yl)ethan-1-ol (86). To a stirred, cooled (ice-salt bath) solution of compound 85 (545 mg, 1.43 mmol, 1.0 equiv) in dry THF (10 mL) was added LiAlH₄ (2.89 mL of a 2 M solution in THF, 5.74 mmol, 4.0 equiv). The reaction mixture was allowed to reach rt followed by refluxing overnight. The solution was cooled (ice-salt bath) and water (0.5 mL) was added slowly. The mixture was treated with 15% aqueous solution of NaOH (0.5 mL) and water (1.5 mL), followed by MgSO₄. The solids were filtered off and the resulting solution was evaporated in vacuo. Purification by automated preparative chromatography (RP-C18, linear gradient from water/MeOH 90:10 to 15:85) gave 220 mg (42%) of the title compound 86 as a white solid. ¹H NMR (400 MHz, DMSO- d_6): δ 8.24 (d, J = 8.1 Hz, 1H), 8.18 (d, J = 7.9 Hz, 1H),

7.61–7.43 (m, 2H), 7.38 (d, J = 7.8 Hz, 1H), 6.91 (d, J = 7.8 Hz, 1H), 4.77 (br s, 1H), 3.97 (s, 3H), 3.96–3.93 (m, 2H), 3.89 (t, J = 5.8 Hz, 2H), 3.63 (t, J = 5.8 Hz, 2H), 3.52 (s, 2H), 3.46 (s, 2H), 2.71 (t, J = 5.8 Hz, 2H), 2.60 (t, J = 5.8 Hz, 2H), 1.86 (br s, 2H). 13 C NMR (101 MHz, DMSO- d_6): δ 155.0, 148.2, 138.0, 133.4, 128.2, 126.7, 126.6, 125.7, 125.5, 125.4, 122.1, 112.0, 103.9, 60.8, 60.0, 56.0, 51.1, 49.8, 49.3, 39.0, 22.2. ESI HRMS (m/z): calcd for $C_{21}H_{27}N_4O_2$ [M + H]⁺, 367.21285; found, 367.21289.

tert-Butyl 3-(5-((4-methoxynaphthalen-1-yl)methyl)-3-((naphthalen-1-ylmethyl)carbamoyl)-4,5,6,7-tetrahydro-1H-pyrazolo[4,3c]pyridin-1-yl)azetidine-1-carboxylate (88). A mixture of compound 82 (270 mg, 0.57 mmol, 1.0 equiv), tert-butyl 3-((methylsulfonyl)oxy)azetidine-1-carboxylate (231 mg, 0.86 mmol, 1.5 equiv) and Cs₂CO₃ (336 mg, 1.14 mmol, 2.0 equiv) in dry DMF (4 mL) was stirred at 95 °C overnight. The mixture was cooled and partitioned between water (30 mL) and EtOAc (5 mL). The aqueous layer was extracted with EtOAc (3 × 5 mL) and the combined organic phase was washed with water $(3 \times 5 \text{ mL})$, brine $(3 \times 5 \text{ mL})$, dried over anhydrous Na2SO4, filtered, and evaporated in vacuo. Purification by FC (hexane/EtOAc 2:1 to 0:1) gave 300 mg (83%) of the title compound 88 as colorless oil. ¹H NMR (400 MHz, CDCl₂): δ 8.31 (dd, J = 7.7, 2.1 Hz, 1H), 8.28-8.23 (m, 1H), 8.18-8.11 (m, 1H),7.91 (dd, J = 7.6, 2.0 Hz, 1H), 7.85 (d, J = 8.2 Hz, 1H), 7.61–7.44 (m, 6H), 7.38 (d, J = 7.8 Hz, 1H), 7.18 (t, J = 5.7 Hz, 1H), 6.77 (d, J)= 7.9 Hz, 1H), 5.09 (d, J = 5.7 Hz, 2H), 4.82 (tt, J = 8.0, 5.6 Hz, 1H), 4.36-4.28 (m, 2H), 4.23 (t, J = 8.5 Hz, 2H), 4.09 (s, 2H), 4.03 (s, 3H), 3.98 (s, 2H), 2.79 (t, J = 5.7 Hz, 2H), 2.57 (t, J = 5.7 Hz, 2H), 1.41 (s, 9H). 13 C NMR (101 MHz, CDCl₃): δ 162.2, 156.1, 155.4, 141.2, 139.2, 133.9, 133.8, 133.4, 131.6, 128.7, 128.5, 127.7, 126.7, 126.6, 126.4, 126.1, 126.03, 125.9, 125.5, 125.0, 124.7, 123.7, 122.3, 118.2, 102.9, 80.0, 60.1, 55.9 (br s), 55.5, 50.2, 48.5, 46.8, 40.9, 28.3, 22.3. ESI HRMS (m/z): calcd for $C_{38}H_{42}N_5O_4$ [M + H]⁺, 632.32313; found, 632,32395.

Ethyl 1-(2-Acetoxyethyl)-5-benzyl-1,4,5,6-tetrahydropyrrolo[3,4c]pyrazole-3-carboxylate (89), Ethyl 2-(2-Acetoxyethyl)-5-benzyl-2,4,5,6-tetrahydropyrrolo[3,4-c]pyrazole-3-carboxylate (90) and 2-Benzyl-2,3,6,7-tetrahydropyrrolo[3',4':3,4]pyrazolo[5,1-c][1,4]oxazin-9(1H)-one (91). To a stirred solution of DIPA (8.84 g, 68.5 mmol, 1.2 equiv) in dry THF (100 mL) at -78 °C was added n-BuLi (25.1 mL of 2.5 M solution in hexane, 62.8 mmol, 1.1 equiv). After stirring at -78 °C for 45 min, a solution of 1-benzylpyrrolidin-3-one (10.0 g, 57.1 mmol, 1.0 equiv) in dry THF (50 mL) was added dropwise. The reaction mixture was stirred for further 30 min at -78 °C, followed by a dropwise addition of a solution of diethyl oxalate (8.34 mL, 57.1 mmol, 1.0 equiv) in dry THF (50 mL). The mixture was then warmed to rt and stirred for further 30 min. Subsequently, EtOH (40 mL), AcOH (40 mL), and hydroxyethylhydrazine (3.87 mL, 57.1 mmol, 1.0 equiv) were added. The reaction mixture was then heated to reflux for 1 h, cooled to rt, and concentrated in vacuo. The residue was dissolved in AcOH (40 mL) and refluxed for further 4 h after which TLC (EtOAc) showed a formation of a predominant spot $(R_f \approx 0.2)$. The solution was evaporated in vacuo and the resulting oil was partitioned between EtOAc (100 mL) and saturated aqueous solution of NaHCO₃ (70 mL). The aqueous phase was extracted with EtOAc (2 \times 25 mL). The combined organic extracts were washed with water (25 mL), brine (25 mL), dried over anhydrous Na2SO4, filtered, and concentrated in vacuo. The residue was purified by FC (hexane/EtOAc 1:1 to 0:1, then EtOAc/MeOH 99:1 to 95:5) which gave 5.82 g of oil. LC/MS (ESI+) analysis of this material showed the presence of both tetrahydropyrrolo [3,4-c]pyrazole $(m/z = 316 [M + H]^+)$ and the corresponding acyclic hydrazone species $(m/z = 334 [M + H]^{+})$. To achieve a full cyclization, the oil was again dissolved in AcOH (50 mL) and refluxed for 14 h. The solution was evaporated in vacuo and the resulting oil was partitioned between EtOAc (100 mL) and saturated aqueous solution of NaHCO₃ (70 mL). The aqueous phase was extracted with EtOAc (2 × 25 mL). The combined organic extracts were washed with water (25 mL), brine (25 mL), dried over anhydrous Na₂SO₄, filtered, and concentrated in vacuo. Purification of the residue by FC (hexane/EtOAc 3:1 to 0:1) gave the title acetylated derivatives 89

(3.51 g, 17%, brown oil) and 90 (0.84 g, 4%, yellow oil) along with the cyclic lactone 91 (0.60 g, 4%, brown solid). 89: ¹H NMR (400 MHz, CDCl₃): δ 7.45–7.34 (m, 4H), 7.34–7.29 (m, 1H), 4.43–4.34 (m, 4H), 4.34-4.29 (m, 2H), 4.04-3.98 (m, 4H), 3.87 (br s, 2H), 2.01 (s, 3H), 1.37 (t, J = 7.1 Hz, 3H). ¹³C NMR (101 MHz, CDCl₃): δ 170.3, 162.0, 148.1, 138.2, 136.7, 128.7, 128.6, 127.5, 126.8, 62.8, 60.9, 60.3, 52.6, 51.1, 50.6, 20.7, 14.4. ESI HRMS (m/z): calcd for $C_{19}H_{24}N_3O_4$ [M + H]⁺, 358.17613; found, 358.17639. **90**: 1H NMR (400 MHz, CDCl₃): δ 7.46–7.41 (m, 2H), 7.38 (ddd, J = 7.8, 6.8, 0.9 Hz, 2H), 7.34-7.29 (m, 1H), 4.79 (t, J = 5.4 Hz, 2H), 4.47-4.38 (m, 2H), 4.31 (q, J = 7.1 Hz, 2H), 4.01 (s, 2H), 3.95 (s, 2H), 3.87 (s, 2H), 2.02 (s, 3H), 1.33 (t, J = 7.1 Hz, 3H). ¹³C NMR (101 MHz, CDCl₂): δ 170.7, 159.6, 156.1 (br s), 138.4 (br s), 128.7, 128.5, 127.4, 126.0, 63.3, 61.0, 60.3, 52.5, 51.9, 50.4, 20.8, 14.3. ESI HRMS (m/z): calcd for $C_{19}H_{24}N_3O_4$ [M + H]⁺, 358.17613; found, 358.17630. 91: ¹H NMR (500 MHz, CDCl₃): δ 7.41 (d, J = 7.2 Hz, 2H), 7.37 (t, J =7.6 Hz, 2H), 7.34-7.27 (m, 1H), 4.74-4.65 (m, 2H), 4.47-4.41 (m, 2H), 4.01 (s, 2H), 3.99 (s, 2H), 3.89 (s, 2H). ¹³C NMR (101 MHz, CDCl₃): δ 158.1, 157.1, 138.2, 128.7, 128.6, 127.4, 127.2, 123.1, 66.3, 60.2, 51.7, 51.5, 46.0. ESI HRMS (m/z): calcd for $C_{15}H_{16}N_3O_2$ [M + H]+, 270.12370; found, 270.12378.

Ethyl 5-Benzyl-1-(2-hydroxyethyl)-1,4,5,6-tetrahydropyrrolo[3,4-c]pyrazole-3-carboxylate (92). A mixture of compound 90 (3.46, 9.69 mmol, 1.0 equiv), K_2CO_3 (1.34 g, 9.69 mmol, 1.0 equiv), and EtOH (15 mL) was stirred for 0.5 h at 60 °C. The reaction mixture was cooled, filtered, and evaporated in vacuo. Purification of the residue by FC (EtOAc/MeOH 99:1 to 95:5) gave 2.93 g (96%) of the title compound 92 as brown oil. ¹H NMR (400 MHz, CDCl₃): δ 7.43–7.34 (m, 4H), 7.34–7.30 (m, 1H), 4.36 (q, J = 7.2 Hz, 2H), 4.20–4.14 (m, 2H), 4.04–3.96 (m, 6H), 3.93 (t, J = 2.2 Hz, 2H), 2.88 (br s, 1H), 1.36 (t, J = 7.2 Hz, 3H). ¹³C NMR (101 MHz, CDCl₃): δ 162.0, 148.2, 137.9, 136.4, 128.8, 128.6, 127.6, 126.2, 76.7, 61.4, 60.9, 60.3, 53.7, 52.6, 51.2, 14.4. ESI HRMS (m/z): calcd for $C_{17}H_{22}N_3O_3$ [M + H]⁺, 316.16557; found, 316.16540.

5-Benzyl-1-(2-hydroxyethyl)-N-(naphthalen-1-ylmethyl)-1,4,5,6tetrahydropyrrolo[3,4-c]pyrazole-3-carboxamide (93). Compound 93 was synthesized by employing the procedure described for compound 69, using intermediate 92 (1.97 g, 5.68 mmol, 1.0 equiv), 1-naphthalenemethylamine (1.34 g, 8.52 mmol, 1.5 equiv), TBD (0.24 g, 1.70 mmol, 0.3 equiv), and dry THF (5 mL). Purification by FC (EtOAc) gave 1.14 mg (47%) of the title compound 93 as a beige solid. ¹H NMR (500 MHz, DMSO- d_6): δ 8.49 (t, J = 6.2 Hz, 1H), 8.21 (dd, I = 8.1, 1.7 Hz, 1H), 7.95 (dd, I = 7.6, 1.9 Hz, 1H), 7.84 (dt, J = 6.2, 3.5 Hz, 1H), 7.60–7.50 (m, 2H), 7.50–7.42 (m, 2H), 7.42– 7.32 (m, 4H), 7.30-7.24 (m, 1H), 4.92 (t, J = 5.4 Hz, 1H), 4.86 (d, J= 6.2 Hz, 2H), 4.06 (t, J = 5.2 Hz, 2H), 3.92 (s, 2H), 3.85 - 3.76 (m, 2H)4H), 3.68 (q, J = 5.3 Hz, 2H). ¹³C NMR (126 MHz, DMSO- d_6): δ 161.8, 149.2, 139.6, 138.8, 135.4, 133.7, 131.3, 129.0, 128.9, 128.8, 127.8, 127.4, 126.6, 126.2, 125.9, 125.8, 124.1, 124.0, 60.4, 59.90, 54.2, 52.5, 51.2. ESI HRMS (m/z): calcd for $C_{26}H_{27}N_4O_2$ $[M + H]^+$, 427.21285; found, 427.21280.

1-(2-Hvdroxyethyl)-N-(naphthalen-1-ylmethyl)-1,4,5,6tetrahydropyrrolo[3,4-c]pyrazole-3-carboxamide Hydrochloride (94). The mixture of compound 94 (1.10 g, 2.58 mmol, 1.0 equiv), 1,4-cyclohexadiene (2.07 g, 25.80 mmol, 10.0 equiv), 10% Pd/C (0.11 g), and MeOH (15 mL, degassed by bubbling argon through for 1 h) was stirred for 4 h at 70 °C in a sealed pressure tube. The mixture was then cooled, filtered through a pad of Celite, and evaporated in vacuo. The residue was dissolved in DCM (10 mL) and cooled (ice-salt bath), followed by treatment with 1 M solution of hydrochloric acid in Et₂O (5 mL). The solvents were evaporated in vacuo to give 0.91 g (quantitative) of the title compound 94 as a white solid. ¹H NMR (500 MHz, DMSO- d_6): δ 10.59 (s, 2H), 8.78 (t, J = 6.1 Hz, 1H), 8.22 (d, J = 8.2 Hz, 1H), 7.95 (d, J = 7.9 Hz, 1H), 7.84 (dd, I = 6.5, 3.1 Hz, 1H), 7.62–7.51 (m, 2H), 7.51–7.43 (m, 2H), 5.15 (t, J = 5.2 Hz, 1H), 4.88 (d, J = 6.1 Hz, 2H), 4.45 (s, 2H), 4.35 (s, 2H), 4.18 (t, J = 4.9 Hz, 2H), 3.72 (q, J = 5.0 Hz, 2H). ¹³C NMR (101 MHz, DMSO- d_6): δ 161.0, 145.9, 139.3, 135.1, 133.7, 131.3, 129.0, 127.9, 126.7, 126.2, 125.9, 124.0, 121.4, 60.3, 54.6, 44.9,

43.7. ESI HRMS (m/z): calcd for $C_{19}H_{21}N_4O_2$ [M + H]⁺, 337.16590; found, 337.16568.

2-(5-((4-Methoxynaphthalen-1-yl)methyl)-3-((naphthalen-1ylmethyl)carbamoyl)-5,6-dihydropyrrolo[3,4-c]pyrazol-1(4H)-yl)ethyl 4-Methylbenzenesulfonate (95). To a stirred, cooled (ice-salt bath) solution of 63 (193 mg, 0.38 mmol, 1 equiv), and DMAP (2 mg, 0.02 mmol, 5 mol %) in dry DCM (5 mL) was added 4toluenesulfonyl chloride (80 mg, 0.42 mmol, 1.1 equiv) and TEA (58 mg, 0.57 mmol, 1.5 equiv). The solution was allowed to warm up to rt then stirred overnight. The reaction mixture was washed with water (20 mL), brine (20 mL), dried over anhydrous Na₂SO₄, filtered, and concentrated in vacuo. Purification by FC (EtOAc/MeOH/TEA 99:0:1 to 94:5:1) gave 191 mg (76%) of the title compound 95 as a white solid. ¹H NMR (500 MHz, CDCl₃): δ 8.34 (dd, J = 8.4, 1.4 Hz, 1H), 8.28 (d, I = 8.3 Hz, 1H), 8.10 (d, I = 8.3 Hz, 1H), 7.90 (dd, I =8.0, 1.6 Hz, 1H), 7.84 (d, J = 8.1 Hz, 1H), 7.63-7.49 (m, 7H), 7.46 (dd, J = 8.1, 7.0 Hz, 1H), 7.40 (d, J = 7.8 Hz, 1H), 7.12 (d, J = 8.0 Hz, 1H), 7.12 (d, J = 8.2H), 6.89 (t, J = 5.8 Hz, 1H), 6.79 (d, J = 7.8 Hz, 1H), 5.02 (d, J = 5.8Hz, 2H), 4.33-4.24 (m, 4H), 4.06 (t, J = 5.0 Hz, 2H), 4.05 (s, 3H), 3.99 (t, J = 2.4 Hz, 2H), 3.76 (s, 2H), 2.34 (s, 3H). ¹³C NMR (75 MHz, CDCl₃): δ 162.5, 161.2, 155.5, 149.2, 145.3, 139.4, 133.9, 133.7, 133.02, 132.04, 131.51, 129.83, 128.76, 128.58, 127.64, 127.14, 126.76, 126.65, 126.02, 126.0, 125.4, 125.2, 124.2, 123.6, 122.4, 102.9, 67.8, 58.3, 55.6, 52.2, 50.6, 50.0, 41.1, 21.6. ESI HRMS (m/z): calcd for C₃₈H₃₇N₄O₅S [M + H]⁺, 661.24792; found, 661.24811.

1-(2-Azidoethyl)-5-((4-methoxynaphthalen-1-yl)methyl)-N-(naphthalen-1-ylmethyl)-1,4,5,6-tetrahydropyrrolo[3,4-c]pyrazole-3-carboxamide (96). A mixture of compound 95 (114 mg, 0.17 mmol) and NaN₃ (56 mg, 0.86 mmol, 5.0 equiv) in dry DMF (2 mL) was vigorously stirred at 70 °C for 4 h. The reaction mixture was partitioned between EtOAc (5 mL) and water (10 mL) and the aqueous layer was extracted with EtOAc (2×2 mL). The combined organic extracts were washed with water (2 \times 2 mL), brine (2 \times 2 mL), dried over anhydrous Na2SO4, filtered, and concentrated in vacuo. Purification by FC (hexane/EtOAc 1:1 to 0:1) gave 81 mg (78%) of the title compound 96 as a white solid. ¹H NMR (400 MHz, CDCl₃): δ 8.36–8.29 (m, 1H), 8.27 (d, I = 8.3 Hz, 1H), 8.11 (ddd, I= 7.8, 0.8 Hz, 1H), 7.90 (dd, J = 8.0, 1.5 Hz, 1H), 7.84 (dt, J = 8.2, 1.1 Hz, 1H), 7.60-7.49 (m, 5H), 7.48-7.41 (m, 2H), 7.01 (t, J = 5.7 Hz, 1H), 6.78 (d, I = 7.8 Hz, 1H), 5.05 (d, I = 5.6 Hz, 2H), 4.36 (s, 2H), 4.15 (s, 2H), 4.04 (s, 3H), 3.98 (dd, J = 6.2, 5.1 Hz, 2H), 3.85 (s, 2H), 3.63 (dd, J = 6.2, 5.1 Hz, 2H). ¹³C NMR (101 MHz, CDCl₃): δ 161.4, 155.5, 149.0, 139.5, 133.9, 133.7, 133.0, 131.6, 128.7, 128.5, 127.1, 126.7, 126.7, 126.6, 126.5, 126.0, 126.0, 125.4, 125.2, 124.2, 123.7, 122.4, 102.9, 58.4, 55.5, 52.4, 50.6, 50.4, 50.2, 41.1. ESI HRMS (m/z): calcd for $C_{31}H_{30}N_7O_2$ [M + H]⁺, 532.24555; found, 532,24615

Protein Expression and Purification. Human, *T. cruzi*, and *T. brucei* N-His-PEX14 (19–84) proteins were cloned into a pETM-11 (EMBL) vector and expressed in a standard *Escherichia coli* BL21 system using an autoinduction medium. ⁴⁹ The proteins were initially purified by affinity chromatography using a NiNTA column. Following dialysis to TEV cleavage buffer (50 mM Tris pH 8.0, 1 mM EDTA, 5 mM β-mercaptoethanol), the TEV protease was added and the reaction was allowed to proceed overnight at room temperature. This step was omitted when the proteins were prepared for AlphaScreen assay that requires presence of a His-tag. Next, the proteins were purified by size exclusion chromatography into either PBS buffer or crystallization buffer (5 mM Tris pH 8.0, 50 mM NaCl, 5 mM β-mercaptoethanol).

AlphaScreen—Based Competition Assays. To determine the compounds' effective concentration (EC₅₀) values against PEX14—PEX5 PPI, an AlphaScreen-based assay was developed according to a PerkinElmer manual. The assay mixture was composed of 3 nM N-His-PEX14 (21–84) and 10 nM of biotinylated PEX5-derived peptide (ALSENWAQEFLA) in a PBS buffer supplemented with 5 mg/mL of BSA and 0.01% (v/v) Tween-80. The assay employed 5 μ g/mL of streptavidin donor beads and 5 μ g/mL of nickel chelate acceptor beads (PerkinElmer). Compound was added to the assay mixture as a DMSO solution. DMSO concentration was kept constant

at 5% (v/v). This concentration was shown to have no effect on the assay readout. The competition curves were measured using a serial dilution of the inhibitor (12 points) while keeping the concentrations of all other assay components constant. The inhibitor EC_{50} was calculated from the Hill sigmoidal fit of the experimental data with asymptotes fixed at maximal assay signal (no inhibitor added) and 0, respectively, using OriginLab OriginPro 9.0. 50

Crystallization and X-ray Structure Solution. The purified protein was mixed with 10-fold molar excess of ligand (50 mM solution in DMSO) in storage buffer (10 mM Tris pH 8.0, 100 mM NaCl and 5 mM β -mercaptoethanol) with 3% DMSO and the mixture was incubated for 1 h at room temperature. The excess ligand and DMSO were removed by washing the complex at 4 °C on a 10 kDacutoff Centricon concentrator with storage buffer. Prior to crystallization the complex was concentrated to about 30 mg/mL. Initial crystallization trials were set up using commercial kits and TTP LabTech Mosquito automated crystallization workstation. Crystals suitable for diffraction testing appeared under different conditions, after few days. They were transferred into a cryoprotectant solution containing the harvesting solution and 25% (v/v) glycerol and cryocooled in liquid nitrogen. Diffraction data were collected at the European Synchrotron Radiation Facility (ESRF, Grenoble, France) beamline ID29. The best diffracting crystal was obtained at room temperature in 0.2 M MgCl₂, 0.1 M Tris pH 8.0, and 20% (w/v) PEG 6000. The experimental data were processed using XDS and XSCALE software. 51 Crystals of TbPEX14-ligand complex diffracted up to 1.2 Å resolution and belonged to the orthorhombic P3₂21 space group. The Matthews coefficient analysis suggested the presence of one molecule in the asymmetric unit. 52 The structure was solved by molecular replacement using Phaser 53 with the TbPEX14 structure (PDB code: 5AON⁵⁴) as a search model. The analysis of the electron density calculated with $(F_0 - F_c)$ and $(2F_0 - F_c)$ coefficients allowed to build the initial model and to unambiguously place the ligand using COOT.55 The starting model was refined by iterations of manual and automated refinement using Refmac5.56 Throughout the refinement, 5% of the reflections were used for the crossvalidation analysis, 57 and the behavior of R_{free} was employed to monitor the refinement strategy. Water molecules were added using Arp/Warp⁵⁸ and subsequently manually inspected. Hydrogens were generated and refined using Refmac5. Detailed information on data collection and refinement statistics are reported in Table S2—see Supporting Information.

MDs Simulations. The PDB files of the three different ligandprotein complexes (TbPEX14-50: 5N8V35 and TbPEX14-61: 6SPT, TbPEX14-29 complex structure was generated in Maestro v11.1⁵⁹ by replacing the nitrogen of the pendant amine of 50 in 5N8V with the oxygen, assuming the same consensus-binding mode) were prepared adding missing sidechains and hydrogens using YASARA Biosciences YASARA Structure's 60 clean built-in command. Structures were then imported into Schrödinger Maestro version 2018.3⁵⁹ and further refined using the Maestro v11.1 "Protein Preparation Wizard".61 Protonation states were calculated using PROPKA 62,63 at pH 7.0 \pm 2.0 and minimization of hydrogen positions with restrained backbone was performed using OPLS_2005 FF⁶⁴ in order to optimize the hydrogen bonding network. All the systems were then prepared for simulation using Maestro v11.1 "System Builder" GUI using a TIP4P solvent model^{65,66} (crystallographic water molecules were deleted) in an automatically generated cubic cell with periodic boundary conditions. In addition to the solvated complex, Na+ and Cl- ions corresponding to a 150 mM buffer were placed in the cell in order to set the total net charge to zero. MDs simulations were run using Schrödinger Maestro Desmond Molecular Dynamics Package version on a NVIDIA 1060 graphics processing unit using the Desmond graphical user interface for a total simulation time of 50 ns to ensure system convergence, with xyz coordinates recording interval every 50 ps (1000 snapshots in total) and 1.2 ps for potential energy calculations of the ensemble. Ensemble class was set to NPT at a temperature of 300 K and pressure of 1.01 bar, force cutoff radius was set to 9.0 Å and each solvated model was relaxed with the Desmond default relaxation protocol before starting the actual simulation.

In Vitro Trypanocidal Activity of Compounds Against T. b. brucei. T. b. brucei bloodstream form (Lister 427, MITat 1.2) parasites were grown in a HMI-11 medium⁶⁸ containing 10% fetal bovine serum (FBS) at 37 °C with 5% CO₂. Antitrypanosomal activities of the compounds were tested using resazurin-based 96-well plate assay. Twofold serial dilutions of each compound (10 wells in each row) were prepared in 96-well plates in HMI-11 medium (100 μ L/well, quadruplicates). As controls, each row included a well without compound and a well with medium alone. 100 μ L of parasite cultures $(4 \times 10^3/\text{mL})$ were inoculated in all wells, except in the well with media alone. Final concentration of parasites was $2 \times 10^3 / \text{mL}$. The plates were incubated for 66 h. Resazurin (25 μ L of 0.1 mg/mL in Hanks Balanced Salt Solution) was added to all wells and the plates were further incubated until 72 h timepoint. Reduction of resazurin by living cells was quantified by measuring the fluorescence with a Synergy H1 microplate reader (excitation 530 nm, emission 585 nm). After subtracting the background fluorescence of the well with media alone, percent survival values were calculated by setting the fluorescence of the wells without compound to "100% survival". Nonlinear regression graphs were plotted in GraphPad Software GraphPad Prism 6.04⁶⁹ to yield sigmoidal dose-response curves and half-maximal effective concentration (EC₅₀) values were determined.

Cytotoxicity of Compounds against HepG2 Cells. HepG2 (Hepatocyte) cells were seeded in 96-well plates (5000 cells/well in rows B–H) and grown overnight at 37 °C in humidified incubator with 5% CO₂. Compounds were tested in triplicate from 100 to 3.125 μ M (twofold serial dilutions, from row H to row C). Row A contained medium alone and served as a negative control. Row B contained cells alone without inhibitors and served as a positive control. Hygromycin B (InvivoGen) was used as a positive control for cytotoxicity. After incubation for 66 h, 25 μ L of 0.1 mg/mL resazurin (dissolved in Hanks Balanced Salt Solution HBSS, Sigma) was added to all wells. Plates were further incubated for 6 h. Fluorescence was measured and data processed as described above for the *T. b. brucei* cytotoxicity assay.

In Vitro Trypanocidal Activity of Compounds Against T. b. rhodesiense STIB900. This stock was isolated in 1982 from a human patient in Tanzania and after several mouse passages cloned and adapted to axenic culture conditions.⁷⁰ Minimum essential medium (50 μ L) supplemented with 25 mM HEPES, 1 g/L additional glucose, 1% MEM nonessential amino acids (100×), 0.2 mM β -mercaptoethanol, 1 mM Na-pyruvate, and 15% heat-inactivated horse serum was added to each well of a 96-well microtiter plate. Serial drug dilutions of eleven threefold dilution steps covering a range from 100 to 0.002 μ g/mL were prepared. Next, 4 × 10³ bloodstream forms of T. b. rhodesiense STIB 900 in 50 µL was added to each well and the plate incubated at 37 °C under a 5% CO₂ atmosphere for 70 h. Alamar Blue (resazurin, 10 μL of 12.5 mg solution in 100 mL double-distilled water) was then added to each well and incubation was continued for a further 2-4 h.71 Subsequently, the plates were read with a Molecular Devices SpectraMax Gemini XS microplate fluorometer, using an excitation wave length of 536 nm and an emission wave length of 588 nm. The IC₅₀ values were calculated by linear regression⁷² from the sigmoidal dose inhibition curves using Molecular Devices SoftmaxPro software.

In Vitro Trypanocidal Activity of Compounds Against *T. Cruzi*. Rat skeletal myoblasts (L-6 cells) were seeded in 96-well microtitre plates at 2000 cells/well in 100 μ L RPMI 1640 medium with 10% FBS and 2 mM L-glutamine. After 24 h, the medium was removed and replaced with 100 μ L per well of 5000 trypomastigote forms of *T. cruzi* Tulahuen strain C2C4 expressing the β -galactosidase (Lac Z) gene. That a h, the medium was removed from the wells and replaced by 100 μ L fresh medium with or without a serial drug dilution of eleven threefold dilution steps covering a range from 100 to 0.002 μ g/mL. After 96 h of incubation the plates were inspected under an inverted microscope to assure growth of the controls and sterility. Then the substrate CPRG/Nonidet (50 μ L) was added to all wells. A color reaction developed within 2–6 h and was read photometrically at 540 nm. Data were analyzed with Molecular

Devices Softmax Pro software, which calculated the IC_{50} values by linear regression⁷² from the sigmoidal dose inhibition curves.

ASSOCIATED CONTENT

Supporting Information

The Supporting Information is available free of charge at https://pubs.acs.org/doi/10.1021/acs.jmedchem.9b01876.

Crystal structure data collection and refinement statistics for TbPEX14-61 complex structure; details on MDs simulations performed for TbPEX14-50, TbPEX14-61, and TbPEX14-20 complexes; and ¹H and ¹³C NMR spectra (PDF)

Molecular formula strings (CSV)

PDB File of the TbPEX14 model used for docking (PDB)

Accession Codes

The coordinates of the crystal structures have been deposited to the RCSB Protein Data Bank under the following accession codes: 6SPT (TbPEX14-61).

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Author Contributions

The manuscript was written through contributions of all the authors. All the authors have given approval to the final version of the manuscript.

Notes

The authors declare no competing financial interest.

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ABBREVIATIONS

PPI, protein—protein interaction; HAT, human African trypanosomiasis; NTD, N-terminal domain; NECT, nifurtimox—eflornithine combination therapy; DNDi, Drugs for Neglected Diseases initiative; PEX, peroxin; PTS, peroxisome targeting sequences; CSP, chemical shift perturbation; TBD, 1,5,7-triazabicyclo [4.4.0] dec-5-ene; TEA, triethylamine; DIPEA, N,N-diisopropyl-N-ethylamine; CDI, 1,1'-carbonyldimidazole; FC, flash column chromatography; dd, doublet of doublets; dt, doublet of triplets; bs, broad signal; SI, selectivity index; FBS, fetal bovine serum; TbPEX14, T. brucei PEX14 N-terminal domain; HsPEX14, human PEX14 N-terminal domain; TcPEX14, T. cruzi PEX14 N-terminal domain

REFERENCES

- (1) Molyneux, D. H.; Savioli, L.; Engels, D. Neglected tropical diseases: progress towards addressing the chronic pandemic. *Lancet* **2017**, 389, 312–325.
- (2) WHO. Research Priorities for Chagas Disease, Human African Trypanosomiasis and Leishmaniasis. World Health Organization Technical Report Series; WHO, 2012; (975), Vol. v-xii, pp 1-100.
- (3) Trypanosomiasis, Human African Trypanosomiasis (Sleeping Sickness), Fact Sheet. http://www.who.int/mediacentre/factsheets/fs259/en/ (accessed April 10, 2018).
- (4) Chagas Disease (American Trypanosomiasis), Fact Sheet. http://www.who.int/mediacentre/factsheets/fs340/en/ (accessed April 10, 2018).
- (5) Giordani, F.; Morrison, L. J.; Rowan, T. G.; De Koning, H. P.; Barrett, M. P. The animal trypanosomiases and their chemotherapy: a review. *Parasitology* **2016**, *143*, 1862–1889.
- (6) Büscher, P.; Cecchi, G.; Jamonneau, V.; Priotto, G. Human African trypanosomiasis. *Lancet* **2017**, *390*, 2397–2409.
- (7) Magez, S.; Caljon, G.; Tran, T.; Stijlemans, B.; Radwanska, M. Current status of vaccination against African trypanosomiasis. *Parasitology* **2010**, *137*, 2017–2027.
- (8) Stijlemans, B.; Radwanska, M.; De Trez, C.; Magez, S. African trypanosomes undermine humoral responses and vaccine development: link with inflammatory responses? *Front. Immunol.* **2017**, *8*, 1–14
- (9) Kennedy, P. G. Clinical features, diagnosis, and treatment of human African trypanosomiasis (sleeping sickness). *Lancet Neurol.* **2013**, *12*, 186–194.
- (10) Mesu, V. K. B. K.; Kalonji, W. M.; Bardonneau, C.; Mordt, O. V.; Blesson, S.; Simon, F.; Delhomme, S.; Bernhard, S.; Kuziena, W.; Lubaki, J.-P. F.; Vuvu, S. L.; Ngima, P. N.; Mbembo, H. M.; Ilunga, M.; Bonama, A. K.; Heradi, J. A.; Solomo, J. L. L.; Mandula, G.; Badibabi, L. K.; Dama, F. R.; Lukula, P. K.; Tete, D. N.; Lumbala, C.; Scherrer, B.; Strub-Wourgaft, N.; Tarral, A. Oral fexinidazole for latestage African Trypanosoma brucei gambiense trypanosomiasis: a pivotal multicentre, randomised, non-inferiority trial. *Lancet* 2018, 391, 144–154.
- (11) Pollastri, M. P. Fexinidazole: a new drug for African sleeping sickness on the horizon. *Trends Parasitol.* **2018**, 34, 178–179.
- (12) Patterson, S.; Wyllie, S. Nitro drugs for the treatment of trypanosomatid diseases: past, present, and future prospects. *Trends Parasitol.* **2014**, 30, 289–298.
- (13) Deeks, E. D. Fexinidazole: First Global Approval. *Drugs* **2019**, 79, 215–220.
- (14) Wyllie, S.; Foth, B. J.; Kelner, A.; Sokolova, A. Y.; Berriman, M.; Fairlamb, A. H. Nitroheterocyclic drug resistance mechanisms in Trypanosoma brucei. *J. Antimicrob. Chemother.* **2016**, *71*, 625–634.
- (15) Sokolova, A. Y.; Wyllie, S.; Patterson, S.; Oza, S. L.; Read, K. D.; Fairlamb, A. H. Cross-resistance to nitro drugs and implications

- for treatment of human African trypanosomiasis. Antimicrob. Agents Chemother. 2010, 54, 2893–2900.
- (16) Field, M. C.; Horn, D.; Fairlamb, A. H.; Ferguson, M. A. J.; Gray, D. W.; Read, K. D.; De Rycker, M.; Torrie, L. S.; Wyatt, P. G.; Wyllie, S.; Gilbert, I. H. Anti-trypanosomatid drug discovery: an ongoing challenge and a continuing need. *Nat. Rev. Microbiol.* **2017**, *15*, 217–231.
- (17) DNDi Announces Successful Completion of SCYX-7158 Phase I Study for Treatment of Sleeping Sickness. https://www.news-medical.net/news/20150909/DNDi-announces-successful-completion-of-SCYX-7158-Phase-I-study-for-treatment-of-sleeping-sickness.aspx (accessed April 10, 2018).
- (18) Alarcón de Noya, B.; Díaz-Bello, Z.; Colmenares, C.; Ruiz-Guevara, R.; Mauriello, L.; Zavala-Jaspe, R.; Suarez, J. A.; Abate, T.; Naranjo, L.; Paiva, M.; Rivas, L.; Castro, J.; Márques, J.; Mendoza, I.; Acquatella, H.; Torres, J.; Noya, O. Large urban outbreak of orally acquired acute Chagas disease at a school in Caracas, Venezuela. *J. Infect. Dis.* **2010**, *201*, 1308–1315.
- (19) Pérez-Molina, J. A.; Molina, I. Chagas disease. Lancet 2018, 391, 82-94.
- (20) Barros-Alvarez, X.; Gualdron-Lopez, M.; Acosta, H.; Caceres, A. J.; Graminha, M. A. S.; Michels, P. A. M.; Concepcion, J. L.; Quinones, W. Glycosomal targets for anti-trypanosomatid drug discovery. *Curr. Med. Chem.* **2014**, *21*, 1679–1706.
- (21) Smirlis, D.; Soares, M. B. P. Selection of Molecular Targets for Drug Development Against Trypanosomatids. In *Proteins and Proteomics of Leishmania and Trypanosoma*; Santos, A. L. S., Branquinha, M. H., d'Avila-Levy, C. M., Kneipp, L. F., Sodré, C. L., Eds.; Springer Netherlands: Dordrecht, 2014; pp 43–76.
- (22) El-Sayed, N. M.; Myler, P. J.; Blandin, G.; Berriman, M.; Crabtree, J.; Aggarwal, G.; Caler, E.; Renauld, H.; Worthey, E. A.; Hertz-Fowler, C.; Ghedin, E.; Peacock, C.; Bartholomeu, D. C.; Haas, B. J.; Tran, A. N.; Wortman, J. R.; Alsmark, U. C.; Angiuoli, S.; Anupama, A.; Badger, J.; Bringaud, F.; Cadag, E.; Carlton, J. M.; Cerqueira, G. C.; Creasy, T.; Delcher, A. L.; Djikeng, A.; Embley, T. M.; Hauser, C.; Ivens, A. C.; Kummerfeld, S. K.; Pereira-Leal, J. B.; Nilsson, D.; Peterson, J.; Salzberg, S. L.; Shallom, J.; Silva, J. C.; Sundaram, J.; Westenberger, S.; White, O.; Melville, S. E.; Donelson, J. E.; Andersson, B.; Stuart, K. D.; Hall, N. Comparative genomics of trypanosomatid parasitic protozoa. *Science* 2005, 309, 404–409.
- (23) Stuart, K.; Brun, R.; Croft, S.; Fairlamb, A.; Gürtler, R. E.; McKerrow, J.; Reed, S.; Tarleton, R. Kinetoplastids: related protozoan pathogens, different diseases. *J. Clin. Invest.* **2008**, *118*, 1301–1310.
- (24) Khare, S.; Nagle, A. S.; Biggart, A.; Lai, Y. H.; Liang, F.; Davis, L. C.; Barnes, S. W.; Mathison, C. J. N.; Myburgh, E.; Gao, M.-Y.; Gillespie, J. R.; Liu, X.; Tan, J. L.; Stinson, M.; Rivera, I. C.; Ballard, J.; Yeh, V.; Groessl, T.; Federe, G.; Koh, H. X. Y.; Venable, J. D.; Bursulaya, B.; Shapiro, M.; Mishra, P. K.; Spraggon, G.; Brock, A.; Mottram, J. C.; Buckner, F. S.; Rao, S. P. S.; Wen, B. G.; Walker, J. R.; Tuntland, T.; Molteni, V.; Glynne, R. J.; Supek, F. Proteasome inhibition for treatment of leishmaniasis, Chagas disease and sleeping sickness. *Nature* 2016, 537, 229–233.
- (25) Russell, S.; Rahmani, R.; Jones, A. J.; Newson, H. L.; Neilde, K.; Cotillo, I.; Rahmani Khajouei, M.; Ferrins, L.; Qureishi, S.; Nguyen, N.; Martinez-Martinez, M. S.; Weaver, D. F.; Kaiser, M.; Riley, J.; Thomas, J.; De Rycker, M.; Read, K. D.; Flematti, G. R.; Ryan, E.; Tanghe, S.; Rodriguez, A.; Charman, S. A.; Kessler, A.; Avery, V. M.; Baell, J. B.; Piggott, M. J. Hit-to-lead optimization of a novel class of potent, broad-spectrum trypanosomacides. *J. Med. Chem.* **2016**, *59*, 9686–9720.
- (26) Fraser, A. L.; Menzies, S. K.; King, E. F. B.; Tulloch, L. B.; Gould, E. R.; Zacharova, M. K.; Smith, T. K.; Florence, G. J. Design and synthesis of broad spectrum trypanosomatid selective inhibitors. *ACS Infect. Dis.* **2018**, *4*, 560–567.
- (27) Di Pietro, O.; Vicente-García, E.; Taylor, M. C.; Berenguer, D.; Viayna, E.; Lanzoni, A.; Sola, I.; Sayago, H.; Riera, C.; Fisa, R.; Clos, M. V.; Pérez, B.; Kelly, J. M.; Lavilla, R.; Muñoz-Torrero, D. Multicomponent reaction-based synthesis and biological evaluation of

- tricyclic heterofused quinolines with multi-trypanosomatid activity. Eur. J. Med. Chem. 2015, 105, 120-137.
- (28) Haanstra, J. R.; González-Marcano, E. B.; Gualdrón-López, M.; Michels, P. A. M. Biogenesis, maintenance and dynamics of glycosomes in trypanosomatid parasites. *Biochim. Biophys. Acta* **2016**, *1863*, 1038–1048.
- (29) Gualdrón-lópez, M.; Brennand, A.; Avilán, L.; Michels, P. A. M. Translocation of solutes and proteins across the glycosomal membrane of trypanosomes; possibilities and limitations for targeting with trypanocidal drugs. *Parasitology* **2013**, *140*, 1–20.
- (30) Schliebs, W. Sleeping sickness: PEX and drugs. *Biochim. Biophys. Acta* **2006**, 1763, 4–5.
- (31) Moyersoen, J.; Choe, J.; Kumar, A.; Voncken, F. G. J.; Hol, W. G. J.; Michels, P. A. M. Characterization of Trypanosoma brucei PEX14 and its role in the import of glycosomal matrix proteins. *Eur. J. Biochem.* **2003**, *270*, 2059–2067.
- (32) Haanstra, J. R.; Van Tuijl, A.; Kessler, P.; Reijnders, W.; Michels, P. A. M.; Westerhoff, H. V.; Parsons, M.; Bakker, B. M. Compartmentation prevents a lethal turbo-explosion of glycolysis in trypanosomes. *Proc. Natl. Acad. Sci. U.S.A.* **2008**, *105*, 17718–17723.
- (33) Kessler, P. S.; Parsons, M. Probing the role of compartmentation of glycolysis in procyclic form Trypanosoma brucei: RNA interference studies of PEX14, hexokinase, and phosphofructokinase. *J. Biol. Chem.* **2005**, 280, 9030–9036.
- (34) Furuya, T.; Kessler, P.; Jardim, A.; Schnaufer, A.; Crudder, C.; Parsons, M. Glucose is toxic to glycosome-deficient trypanosomes. *Proc. Natl. Acad. Sci. U.S.A.* **2002**, *99*, 14177–14182.
- (35) Dawidowski, M.; Emmanouilidis, L.; Kalel, V. C.; Tripsianes, K.; Schorpp, K.; Hadian, K.; Kaiser, M.; Mäser, P.; Kolonko, M.; Tanghe, S.; Rodriguez, A.; Schliebs, W.; Erdmann, R.; Sattler, M.; Popowicz, G. M. Inhibitors of PEX14 disrupt protein import into glycosomes and kill Trypanosoma parasites. *Science* **2017**, 355, 1416–1420.
- (36) Neufeld, C.; Filipp, F. V.; Simon, B.; Neuhaus, A.; Schüller, N.; David, C.; Kooshapur, H.; Madl, T.; Erdmann, R.; Schliebs, W.; Wilmanns, M.; Sattler, M. Structural basis for competitive interactions of Pex14 with the import receptors Pex5 and Pex19. *EMBO J.* **2009**, 28, 745–754.
- (37) Su, J.-R.; Takeda, K.; Tamura, S.; Fujiki, Y.; Miki, K. Crystal structure of the conserved N-terminal domain of the peroxisomal matrix protein import receptor, Pex14p. *Proc. Natl. Acad. Sci. U.S.A.* **2009**, *106*, 417–421.
- (38) Neuhaus, A.; Kooshapur, H.; Wolf, J.; Meyer, N. H.; Madl, T.; Saidowsky, J.; Hambruch, E.; Lazam, A.; Jung, M.; Sattler, M.; Schliebs, W.; Erdmann, R. A novel Pex14 protein-interacting site of human Pex5 is critical for matrix protein import into peroxisomes. *J. Biol. Chem.* **2014**, 289, 437–448.
- (39) Scott, D. E.; Bayly, A. R.; Abell, C.; Skidmore, J. Small molecules, big targets: drug discovery faces the protein–protein interaction challenge. *Nat. Rev. Drug Discovery* **2016**, *15*, 533.
- (40) Rew, Y.; Sun, D. Discovery of a small molecule MDM2 inhibitor (AMG 232) for treating cancer. *J. Med. Chem.* **2014**, *57*, 6332–6341.
- (41) Zhao, Y.; Aguilar, A.; Bernard, D.; Wang, S. Small-molecule inhibitors of the MDM2–p53 protein–protein interaction (MDM2 inhibitors) in clinical trials for cancer treatment. *J. Med. Chem.* **2015**, 58, 1038–1052.
- (42) Jennings, L. E.; Schiedel, M.; Hewings, D. S.; Picaud, S.; Laurin, C. M. C.; Bruno, P. A.; Bluck, J. P.; Scorah, A. R.; See, L.; Reynolds, J. K.; Moroglu, M.; Mistry, I. N.; Hicks, A.; Guzanov, P.; Clayton, J.; Evans, C. N. G.; Stazi, G.; Biggin, P. C.; Mapp, A. K.; Hammond, E. M.; Humphreys, P. G.; Filippakopoulos, P.; Conway, S. J. BET bromodomain ligands: probing the WPF shelf to improve BRD4 bromodomain affinity and metabolic stability. *Bioorg. Med. Chem.* 2018, 26, 2937–2957.
- (43) Yasgar, A.; Jadhav, A.; Simeonov, A.; Coussens, N. P. AlphaScreen-based assays: ultra-high-throughput screening for small-molecule inhibitors of challenging enzymes and protein-protein interactions. *Methods Mol. Biol.* **2016**, *1439*, 77–98.

- (44) Cramer, J.; Krimmer, S. G.; Heine, A.; Klebe, G. Paying the price of desolvation in solvent-exposed protein pockets: impact of distal solubilizing groups on affinity and binding thermodynamics in a series of thermolysin inhibitors. *J. Med. Chem.* **2017**, *60*, 5791–5799.
- (45) Hartman, G. D. K.; Kuduk, S. Derivatives and methods of treating Hepatitis B infections. U.S. Patent 2,016,185,777 A1, 2016.
- (46) Sabot, C.; Kumar, K. A.; Meunier, S.; Mioskowski, C. A convenient aminolysis of esters catalyzed by 1,5,7-triazabicyclo[4.4.0]-dec-5-ene (TBD) under solvent-free conditions. *Tetrahedron Lett.* **2007**, *48*, 3863–3866.
- (47) Dai, H. G.; Li, J. T.; Li, T. S. Efficient and practical synthesis of mannich bases related to gramine mediated by zinc chloride. *Synth. Commun.* **2006**, *36*, 1829–1835.
- (48) DeGoey, D. A.; Chen, H.-J.; Cox, P. B.; Wendt, M. D. Beyond the rule of 5: lessons learned from AbbVie's drugs and compound collection. *J. Med. Chem.* **2018**, *61*, 2636–2651.
- (49) Studier, F. W. Protein production by auto-induction in high density shaking cultures. *Protein Expression Purif.* **2005**, 41, 207–234.
- (50) OriginPro, 9.0; OriginLab Corporation: Northampton, MA, USA.
- (51) Kabsch, W. XDS. Acta Crystallogr., Sect. D: Biol. Crystallogr. **2010**, 66, 125–132.
- (52) Kantardjieff, K. A.; Rupp, B. Matthews coefficient probabilities: improved estimates for unit cell contents of proteins, DNA, and protein-nucleic acid complex crystals. *Protein Sci.* **2003**, *12*, 1865–1871.
- (53) McCoy, A. J.; Grosse-Kunstleve, R. W.; Adams, P. D.; Winn, M. D.; Storoni, L. C.; Read, R. J. Phaser crystallographic software. *J. Appl. Crystallogr.* **2007**, *40*, 658–674.
- (54) Watanabe, Y.; Kawaguchi, K.; Okuyama, N.; Sugawara, Y.; Obita, T.; Mizuguchi, M.; Morita, M.; Imanaka, T. Characterization of the interaction between Trypanosoma brucei Pex5p and its receptor Pex14p. FEBS Lett. 2016, 590, 242–250.
- (55) Emsley, P.; Lohkamp, B.; Scott, W. G.; Cowtan, K. Features and development of Coot. *Acta Crystallogr., Sect. D: Biol. Crystallogr.* **2010**, *66*, 486–501.
- (56) Murshudov, G. N.; Skubak, P.; Lebedev, A. A.; Pannu, N. S.; Steiner, R. A.; Nicholls, R. A.; Winn, M. D.; Long, F.; Vagin, A. A. REFMAC5 for the refinement of macromolecular crystal structures. *Acta Crystallogr., Sect. D: Biol. Crystallogr.* **2011**, *67*, 355–367.
- (57) Brünger, A. T. Free R value: a novel statistical quantity for assessing the accuracy of crystal structures. *Nature* **1992**, *355*, 472–475.
- (58) Morris, R. J.; Perrakis, A.; Lamzin, V. S. ARP/wARP and automatic interpretation of protein electron density maps. *Methods Enzymol.* **2003**, 374, 229–244.
- (59) Schrödinger Release 2018.3; Maestro Schrödinger, LLC: New York, NY, 2019, 2018.
- (60) Krieger, E.; Vriend, G. YASARA View—molecular graphics for all devices—from smartphones to workstations. *Bioinformatics* **2014**, 30, 2981–2982.
- (61) Madhavi Sastry, G.; Adzhigirey, M.; Day, T.; Annabhimoju, R.; Sherman, W. Protein and ligand preparation: parameters, protocols, and influence on virtual screening enrichments. *J. Comput.-Aided Mol. Des.* 2013, 27, 221–234.
- (62) Søndergaard, C. R.; Olsson, M. H. M.; Rostkowski, M.; Jensen, J. H. Improved treatment of ligands and coupling effects in empirical calculation and rationalization of pKa values. *J. Chem. Theory Comput.* **2011**, *7*, 2284–2295.
- (63) Olsson, M. H. M.; Søndergaard, C. R.; Rostkowski, M.; Jensen, J. H. PROPKA3: consistent treatment of internal and surface residues in empirical pKa predictions. *J. Chem. Theory Comput.* **2011**, *7*, 525–537.
- (64) Jorgensen, W. L.; Tirado-Rives, J. The OPLS [optimized potentials for liquid simulations] potential functions for proteins, energy minimizations for crystals of cyclic peptides and crambin. *J. Am. Chem. Soc.* **1988**, *110*, 1657–1666.

- (65) Jorgensen, W. L.; Chandrasekhar, J.; Madura, J. D.; Impey, R. W.; Klein, M. L. Comparison of simple potential functions for simulating liquid water. *J. Chem. Phys.* **1983**, *79*, 926–935.
- (66) Jorgensen, W. L.; Madura, J. D. Temperature and size dependence for Monte Carlo simulations of TIP4P water. *Mol. Phys.* **1985**, *56*, 1381–1392.
- (67) Bowers, K. J.; Chow, E.; Xu, H.; Dror, R. O.; Eastwood, M. P.; Gregersen, B. A.; Klepeis, J. L.; Kolossvary, I.; Moraes, M. A.; Sacerdoti, F. D.; Salmon, J. K.; Shan, Y.; Shaw, D. E. Scalable Algorithms for Molecular Dynamics Simulations on Commodity Clusters. In *Proceedings of the 2006 ACM/IEEE Conference on Supercomputing*; ACM: Tampa, Florida, 2006.
- (68) Hirumi, H.; Hirumi, K. Continuous cultivation of Trypanosoma brucei blood stream forms in a medium containing a low concentration of serum protein without feeder cell layers. *J. Parasitol.* **1989**, 75, 985–989.
- (69) GraphPad Prism, 6.04; GraphPad Software, Inc: San Diego, CA, USA.
- (70) Baltz, T.; Baltz, D.; Giroud, C.; Crockett, J. Cultivation in a semi-defined medium of animal infective forms of Trypanosoma brucei, T. equiperdum, T. evansi, T. rhodesiense and T. gambiense. *EMBO J.* **1985**, *4*, 1273–1277.
- (71) Räz, B.; Iten, M.; Grether-Bühler, Y.; Kaminsky, R.; Brun, R. The Alamar Blue assay to determine drug sensitivity of African trypanosomes (T.b. rhodesiense and T.b. gambiense) in vitro. *Acta Trop.* **1997**, *68*, 139–147.
- (72) Huber, W.; Koella, J. C. A comparison of three methods of estimating EC50 in studies of drug resistance of malaria parasites. *Acta Trop.* **1993**, *55*, 257–261.
- (73) Buckner, F. S.; Verlinde, C. L.; La Flamme, A. C.; Van Voorhis, W. C. Efficient technique for screening drugs for activity against Trypanosoma cruzi using parasites expressing beta-galactosidase. *Antimicrob. Agents Chemother.* **1996**, *40*, 2592–2597.