The discovery of high affinity and metabolically stable allosteric cyclin-dependent kinase 2 (CDK2) inhibitors from screening through lead optimization

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#### <u>Abstract</u>

Despite the status of cyclin-dependent kinase 2 (CDK2) as a validated target for both anticancer and contraceptive indications, a CDK2 inhibitor with exquisite selectivity has been historically challenging, largely due to the structural similarity of the ATP-binding site where most kinase inhibitors bind. We previously discovered an allosteric pocket in CDK2 with potential to bind a compound with desirable selectivity. Using high-throughput and virtual screening methods, we discovered and structurally confirmed an anthranilic acid scaffold that binds this pocket with high affinity. We previously reported that these allosteric CDK2 inhibitors demonstrate a negative cooperative relationship with cyclin binding, are selective for CDK2 over the structurally similar kinase CDK1 and show potential as a non-hormonal contraceptive agent. In this work, we describe our screening and lead optimization efforts that led to the discovery of compounds in this series like EF-4-177 with nanomolar affinity for CDK2. EF-4-177 is metabolically stable with a desirably long ½ life and adequate tissue distribution in mice, demonstrating the potential of this series as a therapeutic. This work details the discovery of the highest affinity allosteric CDK inhibitors reported and shows promise for further development of this series to yield an efficacious and selective allosteric CDK2 inhibitor.

**Introduction** 

Cyclin-dependent kinase 2 (CDK2) is part of the larger CDK family of kinases that regulate the mitotic cell cycle. It is activated by the binding of E-type cyclins at the G1-S transition and by A-type cyclins in the S phase and involves an additional phosphorylation event. Despite its established role in the cell cycle, surprisingly CDK2 knock-out mice are healthy but sterile, as compensatory mechanisms exist in its absence for mitosis but not meiosis.<sup>1-2</sup> However, in certain cancer contexts where E-type and A-type cyclins are overexpressed, CDK2 is hyperactivated, and preclinical animal models show that CDK2 inhibition for these cancers inhibit tumor growth.<sup>3-5</sup> Additionally, because CDK2 is necessary for meiosis and not mitosis, it is a validated target for non-hormonal contraceptive development.<sup>6</sup> While E- and A-type cyclins also play a role in meiosis, it appears that a cyclin-independent function of CDK2 involving the binding of Speedy1 is crucial in the pachytene stage of prophase I of spermatocytes.<sup>7</sup> Overall, developing a selective CDK2 inhibitor could be a great benefit for both cancer patients and individuals desiring safe and efficacious non-hormonal male contraception.

While the biochemical role of CDK2 has been studied for decades, an inhibitor with exquisite selectivity for CDK2 has been challenging and none have gained FDA approval.<sup>8-9</sup> Efforts to inhibit CDK2 have included the pan-CDK inhibitors such as dinaciclib, roscovitine, and flavopiridol, all of which have reached clinical trials but inhibit a multitude of CDK enzymes.<sup>10</sup> Most developed CDK2 inhibitors bind in a type I fashion, meaning they bind the ATP-binding site in CDK2's active conformation, a conformation highly similar to the active states of other similar kinases like cyclin-dependent kinase 1 (CDK1).<sup>9, 11</sup> This leads to undesirable off-target effects, especially considering the necessity of CDK1 function.<sup>12-13</sup> As a result, only one type I CDK2

inhibitor exists to our knowledge that has 100+-fold selectivity over other CDK family members.<sup>14-</sup><sup>15</sup> However, a different approach can be pursued to achieve this desired selectivity.

Targeting the inactive state of CDK2 is a promising avenue to develop selective inhibitors. For example, previous work has shown the structural divergence of inactive CDK2 and CDK1 in contrast to their active states.<sup>11</sup> One way to achieve the desirable selectivity is by targeting allosteric pockets distal to the ATP-binding site, i.e., developing type III inhibitors. Allosteric pockets are often unique to a particular kinase and can be exploited not only for their selectivity advantage, but also hold promise to overcome therapeutic resistance to type I ligands.<sup>9</sup> We previously discovered an unrecognized allosteric pocket within CDK2 that bound the commercial dye 8-anilino-1-naphthalene sulfonic acid (**ANS**) with moderate affinity.<sup>16</sup> We further developed an assay that exploited the environmentally sensitive nature of **ANS** fluorescence to develop a high-throughput assay to discover more drug-like chemical matter that bound to this allosteric pocket.<sup>17</sup> Despite this discovery, the only molecules reported to bind near the **ANS** pocket have been fragments confirmed by X-ray crystallography<sup>18-19</sup> or putative ligands whose allosteric binding mode were not empirically confirmed by such structural studies and whose ability to directly inhibit CDK2 is either weak or unknown.<sup>20-22</sup>

In contrast, we recently reported an anthranilic acid scaffold that binds this site with high affinity and demonstrates a negative cooperative relationship with cyclin binding to stabilize CDK2 in its inactive state. In addition, we confirmed this scaffold has exquisite selectivity for CDK2 over structurally similar CDK1 and shows promise as a contraceptive agent by recapitulating the  $Cdk2^{-/-}$  and  $Spdya^{-/-}$  phenotypes when added to explanted mouse testes.<sup>23</sup> Furthermore, the promise of disrupting the CDK2-cyclin A interaction for anticancer purposes has recently been demonstrated by the complex natural product homoharringtonine (HHT), although

confirmation of HHT's binding pose by empirical means such as X-ray crystallography and its selectivity over other kinases have yet to be explored.<sup>24</sup> In this paper, we will report our screening and lead optimization efforts that have resulted in a series of nanomolar affinity inhibitors that were crystallographically confirmed to bind the allosteric pocket of CDK2, and possess some of the desirable physicochemical and pharmacokinetic properties that demonstrate their potential as future therapeutics.

#### **Results**

To discover new chemical matter that binds the allosteric site of CDK2, we developed a fluorescent binding assay based on the occupancy of the type III site by ANS.<sup>17, 25</sup> The high affinity ATP site ligand staurosporine does not substantially affect ANS binding and can be used to differentiate between inhibitors of ANS binding directly at the allosteric site versus those binding to the active site and those cooperatively displacing ANS via conformational change.<sup>17</sup> The ANS analog *p*-Cl-ANS was used as the ligand for the assay instead of ANS due to its improved fluorescent properties and presumed similar binding to CDK2 (structures in Fig. 1).<sup>25</sup> In this assay, compound displacement of *p*-Cl-ANS from the hydrophobic interior of the allosteric site into the aqueous solvent results in a decrease in fluorescence yield due to the environmentally sensitive nature of ANS and its analogs.



#### Fig. 1. Structures of allosteric CDK2 ligands.

We conducted both a high-throughput screen (HTS) using the *p*-Cl-ANS displacement assay and a virtual high-throughput screen (vHTS) using the crystal structure of ANS bound to CDK2. In the HTS, we selected ~20k compounds from our in-house compound collection for screening, partly based on their similarity to ANS, particularly due to the presence of a negatively charged carboxylic or sulfonic acid. Hits producing > 50% loss of *p*-Cl-ANS fluorescence at the 30 µM screening concentration were cherry-picked from DMSO stocks and tested in doseresponse in the *p*-Cl-ANS assay in the absence and presence of staurosporine to distinguish between inhibitors acting at the ATP and allosteric sites. Hits that produced IC<sub>50</sub> values  $< 50 \mu M$ with little change in the absence and presence of staurosporine ("staurosporine shift" < 2-fold) were selected for repurchase. Ten repurchased hits decreased the fluorescent signal of *p*-Cl-ANS in a concentration-dependent manner in the absence and presence of staurosporine (SI Fig. 1, SI Fig. 2B, SI Table 1). All compounds completely inhibited *p*-Cl-ANS binding with IC<sub>50</sub> values ranging from 2-34  $\mu$ M in the absence of staurosporine. Staurosporine produced < 2-fold reduction in potency for 9 hits but reduced the potency of the ATP-competitive c-Met inhibitor PHA-665752 and the reference CDK2 ATP site inhibitor SU9516 by 2.8- and 234- fold, respectively (SI Fig. 1, SI Fig. 2A, SI Table 1). Although the IC<sub>50</sub> ratio for PHA-665752 was surprisingly low, these results indicate that the "staurosporine shift" is useful to detect ATP site inhibitors that allosterically inhibit the binding of *p*-Cl-ANS to the type III allosteric site.

Scheme 1. Anthranilic acid analogs were synthesized using nucleophilic aromatic substitution  $(S_nAr)$  followed by hydrolysis (method A) or by one-step Ullmann coupling (method B)



Simultaneously, a vHTS using ~720k structures from our in-house collection was conducted. Of the 260 vHTS hits selected for purchase, 230 were commercially available and tested for inhibition of *p*-Cl-ANS binding at 30  $\mu$ M, but only 7 hits produced > 25% inhibition. Of these, three compounds produced concentration-dependent inhibition of the fluorescent signal production of *p*-Cl-ANS with IC<sub>50</sub> values < 100  $\mu$ M, and of these the RAS inhibitor salirasib was the most potent (IC<sub>50</sub> = 3.1  $\mu$ M; Fig. 1; SI Fig. 2B, SI Table 1). Salirasib only partially inhibited binding, possibly due to limited solubility at the higher concentrations tested.

To confirm that these hits were not false positives and the result of fluorescence interference, we needed an orthogonal non-fluorescent assay. We chose to use HSQC NMR of N15 labeled CDK2 to accomplish this. We first noticed that ligands that bind the ATP-binding site, such as non-hydrolysable ATP, dinaciclib, and roscovitine, showed a similar set of global chemical shift perturbations upon binding, as described previously.<sup>25</sup> Furthermore, at saturating concentrations of roscovitine, we discovered that titration of **ANS** into CDK2 caused additional distinct global chemical shift perturbations (SI Fig. 3A). From this, we were able to discern an

NMR "fingerprint" of compounds that bound the ATP-binding site or the allosteric site of interest. Of the 11 hit compounds tested, three did not show any further chemical shift perturbations and were assumed to be either ATP-binding site ligands or non-binders, five compounds showed indeterminate binding due to observed protein aggregation in the assay, and three compounds (NPPB, salirasib, and Sigma S660124) demonstrated binding to the allosteric site (SI Fig. 3B-D). Of these three compounds, two (NPPB and salirasib) were ortho-substituted benzoic acids, yielding our lead scaffold (Fig. 1). Although the chloride channel inhibitor NPPB<sup>26</sup> was the least potent hit evaluated (IC<sub>50</sub> = 34  $\mu$ M; SI Fig. 2B, SI Table 1), it was selected for lead optimization due to its chemical tractability, low "staurosporine shift," and allosteric NMR "fingerprint." Importantly, we were able to confirm the binding of **NPPB** to both the allosteric pocket and ATPsite by X-ray crystallography (Fig. 2A, PDB ID 7RWE). The carboxylate on the phenyl ring engaged with the catalytic lysine (Lys33), which is the same lysine that interacts with the sulfonate of ANS. CDK2 adopted a previously undiscovered conformation with the activation loop folded inwards onto the kinase, like the inactive conformation seen in crystal structures of monomeric CDK2 (PDB ID 1HCL) but less compact to accommodate the small molecule.



**Fig. 2. Binding of initially discovered allosteric CDK2 ligands.** A. Hit **NPPB** binds to both the ATP site and the allosteric pocket of CDK2 (PDB ID 7RWE). B. Anthranilic acid **TW-8-67-2** binds only the allosteric pocket with improved affinity in the *p*-Cl-ANS assay, unaffected by the presence of ATP-site inhibitor staurosporine.

Using NPPB as our hit to be optimized, we next prepared TW-8-67-2 using nucleophilic aromatic substitution ( $S_nAr$ ) followed by hydrolysis (Scheme 1, method A). Most anthranilic acid analogs were prepared in this manner but due to commercial availability and the desire to explore electron-rich aromatic systems, a one-step Ullmann coupling was employed to make some analogs (Scheme 1, method B). In our *p*-Cl-ANS assay, TW-8-67-2 has a 10-fold improved affinity for the allosteric site (*p*-Cl-ANS IC<sub>50</sub> =  $4.6 \pm 1.0 \mu$ M) compared to the initial hit compound NPPB. As described previously, the crystal structure of **TW-8-67-2** bound to CDK2 revealed an additional H-bond interaction made by the indole nitrogen in addition to the increased rigidification of the otherwise flexible alkyl linker.<sup>23</sup> Additionally, TW-8-67-2 only bound the allosteric pocket, unlike NPPB, which bound both the allosteric and ATP-binding site. The affinity of TW-8-67-2 was largely unchanged in the presence of staurosporine, the known high affinity ATP-binding site ligand (*p*-Cl-ANS IC<sub>50</sub> =  $5.7 \pm 1.3 \mu$ M), further confirming the allosteric nature of how TW-8-67-2 binds. To orthogonally assess the binding of these compounds, an isothermal titration calorimetry (ITC) assay was developed. For **TW-8-67-2**, the ITC  $K_D = 5.2 \mu M$ ,<sup>23</sup> comparable to the *p*-Cl-ANS  $IC_{50} = 4.6 \pm 1.0 \ \mu M$  (Fig. 2B).

Table 1. SAR of  $R^1$  and  $R^2$  positions of the phenyl ring of the anthranilic acid series

	R	<sub>1</sub> Н		2
Compound	R <sup>1</sup>	R <sup>2</sup>	R <sup>3</sup>	IC <sub>50</sub> (p-Cl-ANS, μM)
TW-8-67-2	Н	Н	NO <sub>2</sub>	4.6
1	Н	Н	Н	>300
2	NO <sub>2</sub>	Н	Н	38
3	CF3	Н	Н	>300
4	Me	Н	NO <sub>2</sub>	38
5	Н	NO <sub>2</sub>	Н	6.1
6	Н	CF <sub>3</sub>	Н	260
7	Н	Br	Н	13
8	Н	Br	NO <sub>2</sub>	2.8
9	Н	OMe	NO <sub>2</sub>	4.6
10	Н	NH <sub>2</sub>	NO <sub>2</sub>	42



To begin our structure-activity relationship (SAR) exploration, we first examined how tolerant the different positions on the phenyl ring were to changed substitution (Table 1), and how important substitution was at the R<sup>3</sup> position given that only one of the two preliminary screening hits contained a substituent at this position (Fig. 1). An analog without any substituents at the R<sup>1</sup>, R<sup>2</sup>, or R<sup>3</sup> positions (compound 1) did not show binding to the allosteric site in the *p*-Cl-ANS assay and showed weak binding by ITC. Nitro and trifluoromethyl substitutions at the R<sup>1</sup> position did not demonstrate significant binding to the allosteric site (compounds 2 and 3). However, a methyl group at the R<sup>1</sup> position combined with a nitro group at the R<sup>3</sup> position yielded compound 4 with moderate affinity to the allosteric site and 7-fold reduced affinity compared to the parent compound TW-8-67-2. Analogs with substitutions at the R<sup>2</sup> position of the phenyl ring (compounds 5-10) showed improved affinity to the allosteric site compared to analogs with similar substitutions at

the  $R^1$  position, but once again the affinity was strengthened by the presence of a nitro substituent at the  $R^3$  position (for example, compare compounds 7 and 8). Despite this, no analog with a substitution at the  $R^2$  position showed substantially improved affinity compared to parent compound **TW-8-67-2**. However, the presence of a substituent at the  $R^3$  position in general seemed to drive the binding substantially.

Table 2. SAR of the R<sup>3</sup> position of the phenyl ring of the anthranilic acid series

R <sup>1</sup> H						
R <sup>3</sup>	IC <sub>50</sub> ( <i>p</i> -Cl-ANS, μM)					
NO <sub>2</sub>	4.6					
CN	15					
	10					
	>300					
o	47					
O O S	>300					
O S S	>300					
Cl	8.0					
Br	4.0					
CF3	1.1					
OCF <sub>3</sub>	0.94					
OCF <sub>2</sub> H	8.9					
OCH <sub>3</sub>	45					
SF5	0.30					
	$\begin{array}{c} & \begin{array}{c} & \begin{array}{c} & \begin{array}{c} & \\ & \\ & \\ & \\ & \\ & \\ & \\ & \\ & \\ & $					

Next, we explored if replacement of the R<sup>3</sup> position substituent with groups other than nitro could improve the affinity towards CDK2 (Table 2). A nitrile substituent diminished the affinity

(compound 11), as did a methyl ester group more mildly (compound 12). However, the methyl amide, ketone, sulfone, and sulfonamide analogs (compounds 13-16) showed a substantial worsening in affinity. In contrast, large halogen atoms showed near equipotency to the nitro group (compounds 17 and 18). A trifluoromethyl group at the R<sup>3</sup> position (compound 19) improved the affinity 4-fold, and a trifluoromethoxy group at this position (compound 20) showed a similar improvement in affinity. Additionally, we crystallographically confirmed the tolerance of functional groups that replace the nitro group, such as the methyl ester, trifluoromethyl, and the trifluoromethoxy group (co-crystal structures of compounds 12 and 20 with CDK2 in SI Fig. 4-5; PDB IDs 7RXO and 7S85, respectively). Replacing the fluorine atoms in the trifluoromethoxy group with hydrogens, which decreases the size of the substituent and its electron-withdrawing potential, seemed to decrease the affinity substantially (compounds 21 and 22). Because bulky and electron-withdrawing groups seemed to be advantageous for binding, we hypothesized the larger pentafluorosulfanyl (SF<sub>5</sub>) group may further improve affinity.<sup>27</sup> Accordingly, the SF<sub>5</sub> analog 23 showed a 10+-fold improvement in affinity compared to TW-8-67-2. We therefore pursued both the CF<sub>3</sub> and SF<sub>5</sub> groups in future analogs.

**Table 3.** SAR of the indole ring of anthranilic acid series using a Topliss tree-like exploration of all the available synthetic sites



Compound	X	R <sup>2</sup>	R <sup>4</sup>	R <sup>5</sup>	R <sup>6</sup>	<b>R</b> <sup>7</sup>	IC <sub>50</sub> ( <i>p</i> -Cl-ANS, μM)
TW-8-67-2	NH	Η	Η	Н	Н	Н	4.6
24	NMe	Н	Н	Н	Н	Н	21
25	0	Н	Н	Н	Н	Н	32
26	NH	Me	Н	Н	Н	Н	16
27	NH	Н	Me	Н	Н	Н	5.0
28	NH	Н	Cl	Н	Н	Н	3.9
29	NH	Η	H	Me	Н	Н	21
30	NH	Н	Н	Cl	Н	Н	5.8
31	NH	Н	Н	Н	Н	Me	24
32	NH	Н	H	Н	Н	Cl	15
33	NH	Н	Н	Н	Me	Н	0.92
34	NH	Н	Н	Н	Cl	Н	0.44
35	NH	Н	H	Н	F	Н	0.37
36	NH	Н	Н	Н	Br	Н	0.25

We next turned to optimizing the indole ring in a similar manner as we had done with the phenyl ring (Table 3). We first set out to explore how important the NH was in the indole as a hydrogen bond donor by making methylated analog **24** and benzofuran analog **25**. In both cases, there was a 4-6-fold loss in affinity, corresponding to a loss of a hydrogen bond at this position as seen crystallographically (Fig. 2A). Next, a methyl group at the R<sup>2</sup> position (compound **26**) did not improve the binding of these compounds to the allosteric pocket. At the R<sup>4</sup>, R<sup>5</sup>, and R<sup>7</sup> positions, a similar lack of substantial improvement in affinity was observed as methyl and chloro groups were introduced at these positions in a Topliss tree-like fashion (Table 3, compounds **27**-**32**).<sup>28</sup> In particular, a methyl group at the R<sup>5</sup> position (compound **29**) worsened binding

substantially, as did methyl and chloro substituents at the R<sup>7</sup> position (compounds **31** and **32**). The R<sup>4</sup> position in the crystal structure faces back towards the linker and phenyl ring so it is unlikely that small hydrophobic groups at this position, like those explored (compounds **27** and **28**), would drastically affect the binding affinity towards the protein. However, substituents at the R<sup>6</sup> position substantially improved the affinity for CDK2. For example, a R<sup>6</sup> methyl group (compound **33**) improved the affinity 5-fold while a chloro substituent (compound **34**) improved the affinity 10-fold, with other halogens showing similarly large improvements in affinity (compounds **35** and **36**). This aligns with the observed unfilled space near this position in the crystal structure of **TW-8-67-2** (Fig. 3A). An overlay of **TW-8-67-2** and compound **34** bound to CDK2 reveals a near identical binding pose, despite a 10+-fold difference in affinity (Fig. 3B). The difference in affinity is ostensibly due to the space filling by the chlorine. Overall, halogens on the R<sup>6</sup> position on the indole were found to best improve the affinity of these compounds towards CDK2. Other small substituents at this position, like nitrile, also showed improved affinity towards CDK2, such as in **EF-4-084** and **EF-4-177** (Table 4).



**Fig. 3. Empty space off the indole.** A. The crystal structure of **TW-8-67-2** bound to CDK2 reveals an unoccupied pocket off the R<sup>6</sup> position of the indole, as highlighted by the yellow arrow. B. Overlay of crystal structures of CDK2 bound to **TW-8-67-2** (pink; CDK2 in cyan; PDB ID 7RWF) and compound 34 (yellow; CDK2 in green; PDB ID 7S4T).

With potency elements discovered on both the phenyl and indole rings, we next combined these elements to yield compounds with higher affinity. Because some high affinities were approaching the sensitivity limit of the *p*-Cl-ANS assay, ITC was used to measure and rank the relative affinities of compounds with  $K_D < 100$  nM (Table 4, SI Fig. 6, or previously reported<sup>23</sup>). By combining some of the potency elements found from both the phenyl and indole ring SAR, ITC revealed that high affinity compounds **EF-3-006**, **EF-4-084**, and **EF-4-177** had a  $K_D < 50$  nM. Combining the SF<sub>5</sub> group at the R<sup>3</sup> position of the benzene ring with the nitrile at the R<sup>6</sup> position of the indole ring (**EF-4-177**) leads to the highest affinity compound by ITC of the series ( $K_D =$  7.4 nM) and is at the sensitivity limit of the instrument. This demonstrates the interchangeability of these features to modify the physicochemical properties of lead compounds.

NH NH H R <sup>6</sup>					
Compound	R <sup>3</sup>	R <sup>6</sup>	IC <sub>50</sub> (p-Cl-ANS, nM)	K <sub>D</sub> (ITC, nM)	
TW-8-67-2	NO <sub>2</sub>	Н	4,600	5,200	
19	CF <sub>3</sub>	Н	1,100	2,300	
EF-3-006	CF <sub>3</sub>	Br	160	18	
EF-4-084	CF <sub>3</sub>	CN	100	23	
23	SF <sub>5</sub>	Н	300	260	
EF-4-177	SF <sub>5</sub>	CN	87	7.4	

 Table 4. Combining features from the phenyl ring SAR and indole ring SAR yield high affinity allosteric inhibitors of CDK2

We have previously shown that this series, in particular **EF-3-006**, binds CDK2 in a manner that is negatively cooperative with cyclin binding. **EF-3-006** was selective for CDK2 over structurally similar CDK1, showing the promise of targeting the allosteric pocket to achieve selectivity.<sup>23</sup> Lastly, in a cyclin-independent contraceptive context, **EF-3-006** recapitulated the desired phenotype of  $Cdk2^{-/-}$  and  $Spdya^{-/-}$  spermatocytes, showing the therapeutic potential of this series.<sup>23</sup> From these promising results, we next explored the physicochemical and pharmacokinetic properties of these compounds for future study in animals and ultimately humans.

The metabolic stability of lead compound **EF-3-006** had a short ½ life in both human and mouse microsomes. However, higher affinity compound **EF-4-177** demonstrated a ½ life that was 8+-fold longer in humans and 7+-fold longer in mice (Fig. 4A) compared to **EF-3-006**. These long ½ lives were desirable given the therapeutic indications of anticancer and contraceptive purposes we are targeting. Additionally, oral administration of **EF-4-177** in mice demonstrated a corresponding long exposure in both plasma and more specifically, testis for contraceptive uses (Fig. 4B). Coupled with observed low clearance, the overall stability of **EF-4-177** demonstrates

potential for multiple therapeutic uses that require selective targeting of CDK2 and the potential of this series as oral medications.



**Fig. 4. Metabolic profiles of high affinity allosteric CDK2 inhibitors.** A. Metabolic stability in human and mouse microsomes for **EF-3-006** and **EF-4-177** (data provided by Pharmaron). B. **EF-4-177** shows a long ½ life in mice and distribution into the testis.

#### **Discussion and Conclusion**

We have measured and confirmed by crystallography some of the highest affinity allosteric kinase inhibitors known to date, with affinities in the double and single digit nanomolar range as measured by ITC. To discover the new anthranilic acid scaffold, we employed a combination of HTS and vHTS, and confirmed allosteric binding by crystallography and by HSQC NMR changes. Apart from the **ANS** scaffold, this newly developed anthranilic acid series is only the second scaffold to be structurally confirmed by crystallography to bind an allosteric pocket of CDK2 in a type III fashion. The anthranilic acid scaffold has the additional advantage of replacing the sulfonate in **ANS** that engages with the catalytic Lys33 with a more drug-like carboxylate. The improvement in affinity from **ANS** to **EF-4-177** is 5,000-fold. This makes **EF-4-177** the highest

affinity allosteric CDK inhibitor to date, spurring promise for other efficacious and selective inhibitors towards other CDKs involved in disease processes.

While our lead compounds are certainly more lipophilic compared to our starting compound **TW-8-67-2**, the lipophilic ligand efficiencies ( $pK_D - \log D_{7.4}$ ) of the compounds improved from 4.58 for **TW-8-67-2** to 5.62 for **EF-4-177**. While **EF-4-177** replaces the nitro group in **TW-8-67-2** with a more hydrophobic SF<sub>5</sub> group and has an additional nitrile at the R<sup>6</sup> position of the indole, the improvement in affinity is substantial at 600+-fold. Crystal structures of these compounds reveal these allosteric ligands stabilize CDK2 in an inactive conformation of the kinase incompatible with cyclin binding.

We have previously described the mechanism of action and exquisite selectivity of this series for CDK2 over other closely related kinases.<sup>23</sup> While lead compound **EF-3-006** showed promise in cellular studies as a contraceptive agent, its short ½ life was undesirable considering its therapeutic indication. However, by replacing the CF<sub>3</sub> group with a larger SF<sub>5</sub> group on the benzene ring and substituting a nitrile for a bromine on the indole, the metabolic stability improved substantially both in microsome incubation studies as well as in mouse pharmacokinetic studies for **EF-4-177**. The distribution of this compound into the testis further demonstrates the promise of this agent or similar compounds as contraceptives, and its general stability in plasma hints as the potential for this work to yield an efficacious yet selective CDK2 inhibitor for use in animals or even humans.

#### **Experimental Section**

The expression and purification of CDK2 for the *p*-Cl-ANS displacement assay, crystallography, and ITC have been described previously.<sup>23</sup> The liver microsome studies were completed by Pharmaron.

#### *p*-Cl-ANS Displacement Assay

The *p*-Cl-ANS fluorescent binding assay was conducted as described previously.<sup>25</sup> Compounds in DMSO were added to 384-well black clear bottom plates (Greiner 781091) using the Echo 550 (Beckman). SU9516 (50 µM final) and DMSO were used as controls defining 100% and 0% inhibition of *p*-Cl-ANS binding, respectively. For HTS, ~21,000 compounds (final 30  $\mu$ M) were tested in the presence of staurosporine (DMSO 0.3% final). 20 µL of assay buffer (150 mM NaCl, 50 mM HEPES, pH 7.5 containing 0.01% Triton and 2 mM DTT) was added to each well, the plate was shaken for 1 min, centrifuged at 2000 rpm for 5 min intervals until air bubbles were eliminated and then read on a CLARIOstar plate reader (BMG; Ex 388 nm, Em 455 nm) to obtain background fluorescence measurements for correction of intrinsic compound fluorescence. Absorbance was then measured at both the excitation and emission wavelengths for correction of the inner filter effect. Next, 5 µL of buffer containing *p*-Cl-ANS (synthesis reported previously;<sup>29</sup> 15 µM final) and staurosporine (Enzo Life Sciences; final 5 µM) was added. Lastly, 5 µL of CDK2 (final 0.5  $\mu$ M) in buffer was added (final volume of 30  $\mu$ L) and the plate was shaken, centrifuged, incubated for 2 h at rt, and read as described above. The data were analyzed by correcting the raw fluorescence data using the pre-read fluorescence and absorbance data to minimize test compound fluorescence interference and inner filter effect as described previously.<sup>25</sup> For HTS, hit confirmation was conducted with cherry-picked compounds from DMSO stocks (10 mM; DMSO 1% final) in 8-point dose response in duplicate in the absence and presence of staurosporine.

Selected hits were repurchased and tested in the same manner (purchasing information in SI). IC<sub>50</sub> values were calculated using the four-parameter logistic equation using Prism 6.0 (GraphPad). For all compounds in Tables 1-4, a method with 8-point dose response in duplicate, analogous to that done for HTS hit confirmation, was employed. The *p*-Cl-ANS IC<sub>50</sub> values in those tables are reflective of single (n=1) independent experiments. For SI Table 1, the number of independent replicates is noted next to each compound in the table.

#### Virtual High Throughput Screening

Two sets of vHTS were performed against the most interior allosteric ANS site in CDK2 (PDB ID 3PXQ): a first using an in-house GPHR collection with 400k structures and the second using an in-house GPHR collection with 320k structures. Protein preparation, ligand preparation, and docking screening were performed using Protein Preparation Wizard, LigPrep, and Glide in Maestro (Schrödinger, LLC), respectively. Using HTVS in Glide, the top 12.5% scored compounds were advanced to SP docking, where the subsequent top 10% scored compounds were advanced to XP docking, and then the subsequent top 40% scored compounds were advanced to MM-GBSA scoring. MM-GBSA calculations were performed using Prime (Schrödinger, LLC). The top 10% scored compounds from the MM-GBSA analysis were selected for purchase (154 compounds from the first GPHR collection and 106 compounds from the second) and a commercially available subset of 230 compounds were tested in the *p*-Cl-ANS displacement assay as described above.

#### **Crystallization**

Purified CDK2 was buffer exchanged into 100 mM Na/K phosphate, pH 6.2, 2 mM DTT using PD10 columns (GE Life Sciences) and concentrated to 9 mg/mL. Crystallization was performed at 19 °C in hanging droplets by vapor diffusion, and crystals grew from 5% to 7.5%

(v/v) PEG3350, 50 mM HEPES, pH 7.5 and 4.5 mg/mL CDK2. Each structure had only 1 crystal each. Most cocrystals were generated by in-diffusion of ligand-free crystals with inhibitors as detailed in SI Table 1.

#### Data Collection and Refinement

X-ray diffraction data of CDK2 liganded with **NPPB**, **20**, and **34** were collected on a Rigaku Micro-Max 007-HF X-ray generator, equipped with a CCD Saturn 944 system located in the Chemical Biology Core of the Moffitt Cancer Center. Data of CDK2 liganded with **12** were collected at the GM/CA beamline (23ID-B with Dectris PILATUS 3-6M detector, 23ID-D with Dectris EIGER 16M detector) at Argonne National Laboratory. Collection wavelength was 1.54178 Å for **NPPB**, **20**, and **34** and was 1.03318 Å for **12**. All data were collected at 93.2 K and processed with XDS and Aimless of the CCP4 suite.<sup>30</sup> Initial phasing of all structures was done by molecular replacement using PDB deposited structure 4KD1 as the model in PHASER.<sup>31</sup> The inhibitor geometrical restraints were obtained with Phenix Elbow.<sup>32</sup> All structures were refined with Phenix,<sup>33</sup> and model building was performed with Coot.<sup>34</sup> Data collection and refinement statistics are shown in SI Table 2.

#### HSQC NMR

The method of expression and purification of N15 labeled CDK2 have been described previously.<sup>25</sup> NMR samples were prepared with 50 μM CDK2 and 500 μM roscovitine, plus 1 mM of allosteric compound. Trosy-HSQC spectra were collected on a Bruker Avance III 850 MHz spectrometer, using 2048 points in the proton dimension and 96 points in the indirect dimension. Data were analyzed with NMRPipe<sup>35</sup> and SPARKY.<sup>36</sup>

#### Isothermal Titration Calorimetry (ITC)

The ITC method for CDK2 has been described previously.<sup>23</sup> The inhibitor concentration for **23** and **EF-4-177** in the syringe was 75  $\mu$ M, and the CDK2 concentration in the sample cell was 5  $\mu$ M. As before, a one-site binding model was used to fit the ITC data.

#### Pharmacokinetic Studies in Mice

All animal procedures in this study were approved by the University of Minnesota Institutional Animal Care and Use Committee. Animals were allowed to acclimate in their cages after arrival for at least 72 h and were allowed access to rodent chow and water *ad libitum*. The inlife portion of the study consisted of dosing 12 male CD-1 mice (6-8 weeks, Envigo, Indianapolis, IN) PO with 10 mg/kg of **EF-4-177**. At specified time points (30 min, 90 min, 2 h and 4 h) sets of 3 animals were bled, euthanized and testes harvested. Resulting plasma and tissue were frozen at  $\leq$  -20 °C until processing and analysis by LC/MS/MS. Processing of testes consisted of weighing samples and adding 2x volume of cold H<sub>2</sub>O. The tissue mixtures were then homogenized using a Polytron PT 2500E (Kinematica AG, Luzern, Switzerland) with a 5 mm probe (PT-DA 05/2 EC-E85, Kinematica AG). Homogenates were stored at  $\leq$  -20 °C until analysis.

#### LC/MS/MS Quantitation of Animal Samples

LC/MS/MS analysis was performed using a Quattro Ultima triple quadrupole mass spectrometer (Waters, Milford, MA) coupled with a Waters Acquity Ultra Performance Liquid Chromatography system by operating in electrospray in the negative ion mode. Mass-to-charge ratio (m/z) transitions for the analyte (**EF-4-177**, MW = 431.4 Da) and internal standard (MW = 485.3 Da) were determined to be 429.84 > 126.76 and 484.64 > 126.81, respectively. For liquid chromatographic separation, an isocratic elution (30% H<sub>2</sub>O/0.1% formic acid, 70% acetonitrile/0.1% formic acid) at was performed using a Synergi Polar-RP column (75 × 2 mm, 4 µm; Phenomenex, Torrance, CA) with a flow rate of 0.5 mL/min. The total run time was 4.0 min. Standard curve working solutions were prepared by serial dilution of stock solutions (made from 1 mg/mL master stock in DMSO) with 100% acetonitrile to obtain working concentrations at 0.21, 0.62, 1.86, 5.56, 16.67, 50, 150 and 450 ng/ $\mu$ l. Calibration standards were prepared by spiking 50  $\mu$ L of blank with freshly prepared working solutions to achieve standards with concentrations of 0.23, 0.7, 2.1, 6.2, 18.6, 55.6, 166.7, 500, 1500 and 4500 ng/mL. Quality control samples were prepared by spiking 50 µL of blank with freshly prepared working solutions of 0.21, 0.62, 5.56 and 450 ng/µL to obtain the limit of quantitation, low quality control, medium quality control, and high-quality control. Plasma and tissue samples (50  $\mu$ L) were placed in a 1.5 mL Eppendorf tube, 100 µL 100% acetonitrile was added with a spiked amount of internal standard, and the mixture was vortexed. Samples were left to stand at 4 °C for at least 30 min and then microcentrifuged at 16,000 x g for 5 min and the supernatant was recovered. Supernatants (100  $\mu$ L) were transferred to a clean tube and evaporated to dryness under a nitrogen stream at 25 °C for approximately 10-15 min. Samples were reconstituted with 100 µL of HPLC elution buffer (70% H<sub>2</sub>O/0.1% Formic Acid, 30% acetonitrile/0.1% Formic Acid). Processed samples were then transferred to 12 x 32 mm glass vials with 400  $\mu$ L inserts and loaded into 96-vial auto-sampler plates from where 7.5  $\mu$ L was injected into the LC/MS/MS system. Sample analysis was performed using MassLynx Software (v4.1, Waters, Milford, MA). Data are expressed as the mean  $\pm$  standard deviation from the 3 replicate animals.

#### Compound synthesis

All tested compounds were determined to have at least 95% purity as determined by qHNMR. The synthesis of compound *p*-Cl-ANS was published previously,<sup>29</sup> as were the syntheses of **TW-8-67-2**, **19**, **EF-3-006**, and **EF-4-084**.<sup>23</sup> Most anthranilic acids were prepared using S<sub>n</sub>Ar followed by hydrolysis using commercial methyl esters and tryptamine derivatives. Compounds

**13**, **23**, and **EF-4-177** required additional synthesis to produce the desired methyl ester. The synthesis to produce the desired tryptamine derivative for **EF-4-177** has been described previously.<sup>37</sup> NMR spectra and purity data of previously uncharacterized compounds can be found in the Supporting Information.

## General method of S<sub>n</sub>Ar followed by hydrolysis to yield anthranilic acids compounds 1-11, 17-18, 24-36

Commercial electron-deficient methyl 2-chloro- or 2-fluorobenzoic esters (0.30 mmol) were dissolved in DMF (5 mL) together with commercial tryptamine derivatives (1.1 equiv, in their neutral form or as HCl salts) and sodium bicarbonate (3-4 equiv). Potassium carbonate was often an acceptable substitute for NaHCO<sub>3</sub> and was sometimes used instead. The reaction temperature employed varied from room temperature to 90 °C (room temperature for 2, 4, 8, 24-36; 70 °C for 3, 5-7, 9-11, 17-18; 90 °C for 1) and was dependent on the degree of electrondeficiency of the benzene ring and whether the substitution patterns around the benzene ring stabilize the Meisenheimer complex formed as a reactive intermediate in S<sub>n</sub>Ar. The solution was stirred for 12 h and then 1 M HCl (aq) was added until pH < 7 and extracted two times into DCM. The DCM solvent was removed via rotary evaporation and the crude product was dissolved in MeOH to proceed to the hydrolysis step without purification. Water was added to the MeOH mixture until a 1:1 ratio of solvents was reached. A solution of 5 M NaOH (aq, 1 mL) was added, and the hydrolysis proceeded at 50 °C for 12 h and monitored by TLC for completion. The mixture was cooled to room temperature and the solvent was removed by rotary evaporation. 1 M HCl (aq) was added until pH < 4 and the solution was extracted two times into EtOAc. The EtOAc layer was dried with MgSO<sub>4</sub> and loaded onto Celite<sup>®</sup>. Subsequent purification was accomplished by silica gel flash column chromatography with a gradient of EtOAc in hexanes of 0-100%, eluting around 60% EtOAc as the free carboxylic acid. The solvent was removed by rotary evaporation to furnish the final product.

#### Ullmann coupling to yield compounds 12, 14-16, 20-22

The respective 2-bromobenzoic acid (0.9 mmol), tryptamine (1.5 equiv), potassium phosphate (2 equiv), copper (I) iodide (0.1 equiv), [1,1'-binaphthalene]-2,2'-diol (0.2 equiv) were added to anhydrous DMF (8 mL) and stirred at rt for 24 h under nitrogen. The reaction was then filtered through Celite<sup>®</sup>, whereby 1M HCl (aq) was added dropwise until pH < 4. The product was extracted three times into EtOAc. The EtOAc layer was then dried with MgSO<sub>4</sub>, filtered, and concentrated under reduced pressure. The remaining residue was purified by silica flash column chromatography with a gradient of EtOAc in hexanes of 0-100% over 7 min, eluting around 60% EtOAc as the free carboxylic acid. The solvent was removed by rotary evaporation to furnish the final product.

#### Procedure to yield compound 13

Compound **13** was made via amide coupling of methylammonium chloride (1.1 equiv) with the commercial 4-fluoro-3-(methoxycarbonyl)benzoic acid (0.5 mmol) using hexafluorophosphate azabenzotriazole tetramethyl uronium (1 equiv) and triethylamine (3 equiv) in DMF (5 mL) at room temperature, stirred for 1 h. Then 1 M HCl (aq) was added until pH < 7 and the intermediate was extracted two times into EtOAc, dried over MgSO4, and loaded onto Celite<sup>®</sup>. Purification of the intermediate methyl ester was accomplished by silica gel flash column chromatography with a gradient of EtOAc in hexanes of 0-100% over 7 min, eluting at 70% EtOAc. Next, the methyl ester and tryptamine (1.1 equiv) were dissolved in DMF (5 mL) and NaHCO<sub>3</sub> (3 equiv) was added. The reaction mixture was stirred for 12 h at 70 °C in an S<sub>n</sub>Ar step as described above for the "General method of S<sub>n</sub>Ar followed by hydrolysis to yield anthranilic acids." After extraction into EtOAc from 1 M HCl (aq), a similar hydrolysis step was performed to yield compound **13**, which was worked up and purified as the free carboxylic acid via flash column chromatography in 0-30% MeOH in DCM in 5 min, eluting at 25% MeOH.

# Preparation of methyl 2-fluoro-5-(pentafluoro- $\lambda^6$ -sulfaneyl)benzoate for the synthesis of compounds 23 and EF-4-177

To 2-Fluoro-5-(pentafluoro- $\lambda^6$ -sulfaneyl)benzoic acid (0.50 mmol) was added methyl iodide (5 equiv) in DMF (5 mL) under anhydrous conditions together with K<sub>2</sub>CO<sub>3</sub> (4 equiv) and was stirred at 40 °C for 12 h. The reaction was extracted into DCM twice and dried over MgSO<sub>4</sub>. Subsequent purification was accomplished by silica gel flash column chromatography using a gradient of EtOAc in hexanes of 0-100% over 7 min, eluting around 30% EtOAc in 71% yield as an oil. Next, the general S<sub>n</sub>Ar/hydrolysis method was employed to yield **23** and **EF-4-177**.

**2-((2-(1***H***-Indol-3-yl)ethyl)amino)benzoic Acid (compound 1)**. 7% yield. Colorless solid. m.p. 158-161 °C. <sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>) δ 10.86 (s, 1H), 7.77 (d, *J* = 8.0 Hz, 1H), 7.58 (d, *J* = 7.9 Hz, 1H), 7.35 (d, *J* = 8.0 Hz, 2H), 7.23 (s, 1H), 7.08 (t, *J* = 7.5 Hz, 1H), 6.98 (t, *J* = 7.4 Hz, 1H), 6.77 (d, *J* = 8.5 Hz, 1H), 6.53 (t, *J* = 7.5 Hz, 1H), 3.46 (t, *J* = 7.1 Hz, 2H), 3.03 (t, *J* = 7.0 Hz, 2H).

**2-((2-(1***H***-Indol-3-yl)ethyl)amino)-3-nitrobenzoic Acid (compound 2)**. 47% yield. Yellow solid. m.p. 75-77 °C. <sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>) δ 10.87 (s, 1H), 8.70 (s, 1H), 8.05 (d, *J* = 7.6 Hz, 1H), 7.96 (d, *J* = 8.1 Hz, 1H), 7.54 (d, *J* = 7.9 Hz, 1H), 7.34 (d, *J* = 8.0 Hz, 1H), 7.16 (s, 1H), 7.08 (t, *J* = 7.5 Hz, 1H), 6.98 (t, *J* = 7.4 Hz, 1H), 6.73 (t, *J* = 7.9 Hz, 1H), 3.16 – 3.09 (m, 2H), 3.00 (t, *J* = 6.8 Hz, 2H).

**2-((2-(1***H***-Indol-3-yl)ethyl)amino)-3-(trifluoromethyl)benzoic Acid (compound 3)**. 52% yield. Yellow solid. m.p. 86-89 °C. <sup>1</sup>H NMR (400 MHz, acetone-d<sub>6</sub>)  $\delta$  10.05 (s, 1H), 8.17 (d, *J* = 9.8 Hz, 1H), 7.77 (d, *J* = 9.8 Hz, 1H), 7.59 (d, *J* = 8.2 Hz, 1H), 7.37 (d, *J* = 8.1 Hz, 1H), 7.22 (s, 1H), 7.09 (t, *J* = 7.6 Hz, 1H), 7.01 (t, *J* = 6.9 Hz, 1H), 6.89 (t, *J* = 8.2 Hz, 1H), 3.61 (t, *J* = 7.9 Hz, 2H), 3.09 – 3.14 (m, 2H). <sup>19</sup>F NMR (376 MHz, acetone-d<sub>6</sub>)  $\delta$  120.3. <sup>13</sup>C NMR (101 MHz, acetone-d<sub>6</sub>)  $\delta$  169.7, 151.2, 137.8, 137.0, 134.12, 134.06, 128.4. 123.8, 123.6, 122.2, 119.5, 119.2, 118.3, 117.3, 112.6, 112.2, 49.6, 27.4.

**2-((2-(1***H***-Indol-3-yl)ethyl)amino)-3-methyl-5-nitrobenzoic Acid (compound 4)**. 81% yield. Yellow solid. m.p. 210-211 °C. <sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>) δ 13.40 (s, 1H), 10.86 (s, 1H), 8.47 (d, *J* = 2.9 Hz, 1H), 8.01 (d, *J* = 2.9 Hz, 1H), 7.56 (d, *J* = 7.9 Hz, 1H), 7.34 (d, *J* = 8.0 Hz, 1H), 7.18 (s, 1H), 7.07 (t, *J* = 7.6 Hz, 1H), 6.98 (t, *J* = 7.4 Hz, 1H), 3.82 – 3.77 (m, 2H), 3.00 (t, *J* = 7.0 Hz, 2H), 2.43 (s, 3H).

**2-((2-(1***H***-Indol-3-yl)ethyl)amino)-4-nitrobenzoic Acid (compound 5)**. 68% yield. Yellow solid. m.p. 200-202 °C. <sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>) δ 10.88 (s, 1H), 8.59 (s, 1H), 8.00 (d, *J* = 8.6 Hz, 1H), 7.61 (d, *J* = 7.8 Hz, 1H), 7.43 (s, 1H), 7.35 (d, *J* = 8.1 Hz, 1H), 7.27 (d, *J* = 8.8 Hz, 1H), 7.24 (s, 1H), 7.08 (t, *J* = 7.5 Hz, 1H), 6.99 (t, *J* = 7.4 Hz, 1H), 3.54 (t, *J* = 7.2 Hz, 2H), 3.06 (t, *J* = 7.0 Hz, 2H). <sup>13</sup>C NMR (101 MHz, DMSO-d<sub>6</sub>) δ 150.7, 150.6, 136.3, 133.1, 127.0, 123.1, 121.0, 118.3, 118.2, 111.4, 111.3, 107.5, 104.7, 43.0, 24.4.

**2-((2-(1***H***-Indol-3-yl)ethyl)amino)-4-(trifluoromethyl)benzoic Acid (compound 6**). 42% yield. Yellow solid. m.p. 191-196 °C. <sup>1</sup>H NMR (400 MHz, acetone-d<sub>6</sub>) δ 11.27 (s, 1H), 10.06 (s, 1H), 8.21 (s, 1H), 8.07 (d, *J* = 8.2 Hz, 1H), 7.65 (d, *J* = 7.9 Hz, 1H), 7.39 (d, *J* = 8.1 Hz, 1H), 7.29 (s, 1H), 7.13 – 7.06 (m, 2H), 7.03 (t, *J* = 8.0 Hz, 1H), 6.84 (d, *J* = 8.2 Hz, 1H), 3.61 – 3.69 (m, 2H), 3.15 – 3.21 (m, 2H). <sup>19</sup>F NMR (376 MHz, acetone-d<sub>6</sub>) δ 113.2. <sup>13</sup>C NMR (101 MHz, acetone-d<sub>6</sub>) δ 169.6, 152.1, 137.8, 134.0, 128.4, 126.4, 123.8, 122.2, 119.5, 119.2, 113.2, 112.9, 112.2, 110.63, 110.60, 108.65, 108.61, 44.1, 25.6.

**2-((2-(1***H***-Indol-3-yl)ethyl)amino)-4-bromobenzoic Acid (compound 7)**. 66% yield. Colorless solid. m.p. 201-202 °C. <sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>) δ 12.77 (s, 1H), 10.87 (s, 1H), 8.01 (s, 1H), 7.67 (d, *J* = 8.4 Hz, 1H), 7.58 (d, *J* = 7.9 Hz, 1H), 7.35 (d, *J* = 8.1 Hz, 1H), 7.23 (s, 1H), 7.08 (t, *J* = 7.8 Hz, 1H), 6.98 (t, *J* = 7.4 Hz, 1H), 6.93 (s, 1H), 6.70 (dd, *J* = 8.4, 1.9 Hz, 1H), 3.48 (t, *J* = 7.0 Hz, 2H), 3.03 (t, *J* = 6.9 Hz, 2H). <sup>13</sup>C NMR (101 MHz, DMSO-d<sub>6</sub>) δ 169.4, 151.5, 136.3, 133.4, 128.6, 127.0, 123.1, 121.0, 118.3, 116.8, 113.4, 111.4, 111.2, 109.0, 42.7, 24.4.

**2-((2-(1***H***-Indol-3-yl)ethyl)amino)-4-bromo-5-nitrobenzoic Acid (compound 8)**. 56% yield. Yellow solid. m.p. 218-220 °C. <sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>) δ 13.55 (s, 1H), 10.89 (s, 1H), 8.70 (s, 1H), 8.54 (s, 1H), 7.59 (d, *J* = 7.9 Hz, 1H), 7.35 (d, *J* = 8.1 Hz, 1H), 7.24 (s, 1H), 7.15 (s, 1H), 7.08 (t, *J* = 7.5 Hz, 1H), 6.99 (t, *J* = 7.5 Hz, 1H), 3.65 (q, *J* = 6.4 Hz, 2H), 3.05 (t, *J* = 6.9 Hz, 2H).

**2-((2-(1***H***-Indol-3-yl)ethyl)amino)-4-methoxy-5-nitrobenzoic Acid (compound 9)**. 21% yield. Yellow solid. m.p. 220-222 °C. <sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>) δ 13.07 (s, 1H), 10.87 (s, 1H), 8.76 (s, 1H), 8.51 (s, 1H), 7.60 (d, *J* = 7.8 Hz, 1H), 7.34 (d, *J* = 8.1 Hz, 1H), 7.22 (s, 1H), 7.07 (t, *J* = 7.5 Hz, 1H), 6.98 (t, *J* = 7.4 Hz, 1H), 6.21 (s, 1H), 3.84 (s, 3H), 3.64 (q, *J* = 6.5 Hz, 2H), 3.07 (t, *J* = 6.8 Hz, 2H).

**2-((2-(1***H***-Indol-3-yl)ethyl)amino)-4-amino-5-nitrobenzoic Acid (compound 10)**. 69% yield. Yellow solid. m.p. 221-223 °C. <sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>) δ 12.81 (s, 1H), 10.89 (s, 1H), 8.61 (s, 1H), 8.39 (s, 1H), 7.61 (d, *J* = 7.8 Hz, 1H), 7.54 (s, 2H), 7.35 (d, *J* = 8.1 Hz, 1H), 7.24 (s, 1H), 7.08 (t, *J* = 7.5 Hz, 1H), 6.99 (t, *J* = 7.4 Hz, 1H), 6.05 (s, 1H), 3.45 (q, *J* = 6.6 Hz, 2H), 3.06 (t, *J* = 7.1 Hz, 2H). <sup>13</sup>C NMR (101 MHz, DMSO-d<sub>6</sub>) δ 168.4, 153.5, 149.9, 136.3, 132.6, 127.0, 123.1, 122.2, 121.0, 118.34, 118.31, 111.4, 111.1, 103.0, 93.2, 43.0, 24.0.

**2-((2-(1***H***-Indol-3-yl)ethyl)amino)-5-cyanobenzoic Acid (compound 11)**. 24% yield. Light yellow solid. m.p. 247 °C. <sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>)  $\delta$  13.12 (s, 1H), 10.87 (s, 1H), 8.53 (s, 1H), 8.07 (s, 1H), 7.68 (d, *J* = 11.1 Hz, 1H), 7.58 (d, *J* = 7.8 Hz, 1H), 7.34 (d, *J* = 8.1 Hz, 1H), 7.23 (d, *J* = 2.3 Hz, 1H), 7.07 (t, *J* = 8.2 Hz, 1H), 6.98 (t, *J* = 7.5 Hz, 1H), 6.91 (d, *J* = 9.0 Hz, 1H), 3.63 – 3.53 (m, 2H), 3.04 (t, *J* = 7.0 Hz, 2H). <sup>13</sup>C NMR (101 MHz, DMSO-d<sub>6</sub>)  $\delta$  168.5, 152.8, 136.9, 136.5, 136.3, 127.0, 123.2, 121.0, 119.6, 118.30, 118.28, 112.3, 111.4, 111.0, 110.2, 95.0, 42.6, 24.3.

**2-((2-(1***H***-Indol-3-yl)ethyl)amino)-5-(methoxycarbonyl)benzoic Acid (compound 12)**. 56% yield. Colorless solid. m.p. 233-234 °C. <sup>1</sup>H NMR (400 MHz, acetone-d<sub>6</sub>) δ 10.09 (s, 1H), 8.63 (s, 1H), 8.55 (s, 1H), 7.98 (d, *J* = 9.0 Hz, 1H), 7.66 (d, *J* = 7.8 Hz, 1H), 7.40 (d, *J* = 7.8 Hz, 1H), 7.28 (s, 1H), 7.12 (t, *J* = 7.6 Hz, 1H), 7.04 (t, *J* = 8.0 Hz, 1H), 6.91 (d, *J* = 9.0 Hz, 1H), 3.83 (s, 3H), 3.70 – 3.63 (m, 2H), 3.18 (t, *J* = 7.1 Hz, 2H). <sup>13</sup>C NMR (101 MHz, acetone-d<sub>6</sub>) δ 170.0, 166.8, 155.1, 137.8, 136.1, 135.4, 128.4, 123.8, 122.2, 119.5, 119.3, 116.5, 112.8, 112.2, 111.9, 109.8, 51.8, 44.1, 25.6.

**2-((2-(1***H***-Indol-3-yl)ethyl)amino)-5-(methylcarbamoyl)benzoic Acid (compound 13)**. 43% yield across three steps. Colorless solid. m.p. 245-248 °C. <sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>) δ 10.90 (s, 1H), 8.55 (s, 1H), 8.36 (d, *J* = 2.3 Hz, 1H), 8.19 – 8.13 (m, 1H), 7.82 (dd, *J* = 8.8, 2.4 Hz, 1H), 7.58 (d, *J* = 7.8 Hz, 1H), 7.35 (d, *J* = 8.1 Hz, 1H), 7.24 (s, 1H), 7.08 (t, *J* = 7.5 Hz, 1H), 6.98 (t, *J* = 7.5 Hz, 1H), 6.77 (d, *J* = 8.9 Hz, 1H), 3.51 (t, *J* = 7.1 Hz, 2H), 3.04 (t, *J* = 7.0 Hz, 2H), 2.73 (d, *J* = 4.3 Hz, 3H). <sup>13</sup>C NMR (101 MHz, DMSO-d<sub>6</sub>) δ 166.1, 152.2, 136.2, 132.4, 131.6, 127.0, 123.1, 120.9, 119.7, 118.3, 111.4, 110.2, 42.8, 26.1, 24.6.

**2-((2-(1***H***-Indol-3-yl)ethyl)amino)-5-acetylbenzoic Acid (compound 14)**. 54% yield. Light yellow solid. m.p. 228-229 °C. <sup>1</sup>H NMR (400 MHz, acetone-d<sub>6</sub>) δ 10.09 (s, 1H), 8.60 (s, 1H), 8.00 (d, *J* = 9.0 Hz, 1H), 7.66 (d, *J* = 8.9 Hz, 1H), 7.40 (d, *J* = 7.5 Hz, 1H), 7.28 (s, 1H), 7.12 (t, *J* = 7.6 Hz, 1H), 7.04 (t, *J* = 7.4 Hz, 1H), 6.92 (d, *J* = 9.0 Hz, 1H), 3.68 (q, *J* = 6.7 Hz, 2H), 3.15 –3.21 (m, 2H), 2.48 (s, 3H). <sup>13</sup>C NMR (101 MHz, acetone-d<sub>6</sub>) δ 195.1, 170.2, 155.0, 137.8, 135.3, 134.6, 128.4, 125.1, 123.8, 122.2, 119.5, 119.3, 112.8, 111.8, 109.6, 44.1, 25.9, 25.7.

**2-((2-(1***H***-Indol-3-yl)ethyl)amino)-5-(methylsulfonyl)benzoic Acid (compound 15)**. 58% yield. Colorless solid. m.p. 224-226 °C. <sup>1</sup>H NMR (400 MHz, acetone-d<sub>6</sub>)  $\delta$  11.38 (s, 1H), 10.08 (s, 1H), 8.61 (s, 1H), 8.45 (s, 1H), 7.84 (d, *J* = 9.2 Hz, 1H), 7.66 (d, *J* = 8.1 Hz, 1H), 7.41 (d, *J* = 8.1 Hz, 1H), 7.29 (s, 1H), 7.12 (t, *J* = 6.9 Hz, 1H), 7.07 – 6.99 (m, 2H), 3.74 – 3.64 (m, 2H), 3.19 (t, *J* = 7.0 Hz, 2H), 3.05 (s, 3H). <sup>13</sup>C NMR (101 MHz, acetone-d<sub>6</sub>)  $\delta$  169.5, 155.0, 137.8, 133.9, 133.4, 128.4, 126.8, 123.8, 122.3, 119.6, 119.3, 112.7, 112.6, 112.2, 109.7, 44.9, 44.1, 25.6.

**2-((2-(1***H***-Indol-3-yl)ethyl)amino)-5-(***N***-methylsulfamoyl)benzoic Acid (compound 16). 48% yield. Colorless solid. m.p. 233-237 °C. <sup>1</sup>H NMR (400 MHz, acetone-d<sub>6</sub>) δ 10.09 (s, 1H), 8.52 (s, 1H), 8.38 (s, 1H), 7.77 (d,** *J* **= 11.3 Hz, 1H), 7.66 (d,** *J* **= 8.2 Hz, 1H), 7.40 (d,** *J* **= 7.2 Hz, 1H), 7.29 (s, 1H), 7.11 (t,** *J* **= 7.6 Hz, 1H), 7.03 (t,** *J* **= 6.9 Hz, 1H), 6.99 (d,** *J* **= 9.0 Hz, 1H), 6.11 (s, 1H), 3.72 – 3.63 (m, 2H), 3.19 (t,** *J* **= 7.0 Hz, 2H), 2.55 (d,** *J* **= 5.3 Hz, 3H). <sup>13</sup>C NMR (101 MHz, acetone-d<sub>6</sub>) δ 169.6, 154.4, 137.8, 133.9, 133.1, 128.4, 125.0, 123.8, 123.7, 122.2, 119.5, 119.3, 112.8, 112.3, 112.2, 109.6, 44.1, 25.6.** 

**2-((2-(1***H***-Indol-3-yl)ethyl)amino)-5-chlorobenzoic Acid (compound 17)**. 19% yield. Colorless solid. m.p. 206-210 °C. <sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>) δ 10.86 (s, 1H), 7.70 (s, 1H), 7.58 (d, *J* = 7.8 Hz, 1H), 7.41 – 7.31 (m, 2H), 7.22 (s, 1H), 7.07 (t, *J* = 7.5 Hz, 1H), 6.98 (t, *J* = 7.4 Hz, 1H), 6.83 (d, *J* = 9.1 Hz, 1H), 3.47 (t, *J* = 7.0 Hz, 2H), 3.02 (t, *J* = 6.9 Hz, 2H).

**2-((2-(1***H***-Indol-3-yl)ethyl)amino)-5-bromobenzoic Acid (compound 18)**. 27% yield. Colorless solid. m.p. 204-209 °C. <sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>) δ 10.86 (s, 1H), 7.83 (d, *J* = 2.5 Hz, 1H), 7.57 (d, *J* = 7.8 Hz, 1H), 7.48 (dd, *J* = 9.0, 2.6 Hz, 1H), 7.34 (d, *J* = 8.1 Hz, 1H), 7.21 (s, 1H), 7.07 (t, *J* = 6.9 Hz, 1H), 6.98 (t, *J* = 8.0 Hz, 1H), 6.78 (d, *J* = 9.1 Hz, 1H), 3.47 (t, *J* = 7.0 Hz, 2H), 3.02 (t, *J* = 6.9 Hz, 2H).

**2-((2-(1***H***-Indol-3-yl)ethyl)amino)-5-(trifluoromethoxy)benzoic Acid (compound 20)**. 78% yield. Colorless solid. m.p. 177-179 °C. <sup>1</sup>H NMR (400 MHz, acetone-*d*<sub>6</sub>)  $\delta$  11.22 (s, 1H), 10.05 (s, 1H), 8.07 (s, 1H), 7.78 (s, 1H), 7.64 (s, 1H), 7.38 (t, *J* = 9.0 Hz, 2H), 7.27 (s, 1H), 7.11 (t, *J* = 7.6 Hz, 1H), 7.02 (t, *J* = 7.4 Hz, 1H), 6.93 (d, *J* = 9.3 Hz, 1H), 3.58 – 3.64 (m, 2H), 3.16 (t, *J* = 7.1 Hz, 2H). <sup>19</sup>F NMR (376 MHz, acetone-*d*<sub>6</sub>)  $\delta$  118.1. <sup>13</sup>C NMR (101 MHz, acetone-*d*<sub>6</sub>)  $\delta$  169.3, 151.3, 137.8, 129.1, 128.4, 125.2, 123.7, 122.2, 110.1, 119.5, 119.3, 113.4, 113.0, 112.2, 110.2, 44.3, 25.7.

**2-((2-(1***H***-Indol-3-yl)ethyl)amino)-5-(difluoromethoxy)benzoic Acid (compound 21).** 69% yield. Colorless solid. m.p. 179-180 °C. <sup>1</sup>H NMR (400 MHz, acetone-d<sub>6</sub>) δ 11.11 (s, 1H), 10.06 (s, 1H), 7.97 (s, 1H), 7.71 (s, 1H), 7.65 (d, *J* = 7.0 Hz, 1H), 7.40 (d, *J* = 8.9 Hz, 1H), 7.31 – 7.24 (m, 2H), 7.11 (t, *J* = 6.9 Hz, 1H), 7.03 (t, *J* = 6.9 Hz, 1H), 7.01 – 6.60 (m, 2H), 3.59 (t, *J* = 7.2 Hz, 2H), 3.15 (t, *J* = 7.1 Hz, 2H). <sup>19</sup>F NMR (376 MHz, acetone-d<sub>6</sub>) δ 95.7, 95.5. <sup>13</sup>C NMR (101 MHz, acetone-d<sub>6</sub>) δ 169.6, 150.5, 140.2, 137.8, 128.5, 124.0, 123.7, 122.2, 119.5, 119.3, 118.0, 113.3, 113.1, 112.2, 110.3, 44.3, 25.8.

**2-((2-(1***H***-Indol-3-yl)ethyl)amino)-5-methoxybenzoic Acid (compound 22)**. 89% yield. Yellow solid. m.p. 170-173 °C. <sup>1</sup>H NMR (400 MHz, acetone-d<sub>6</sub>) δ 10.05 (s, 1H), 7.64 (d, *J* = 7.8 Hz, 1H), 7.45 (s, 1H), 7.39 (d, *J* = 7.5 Hz, 1H), 7.25 (s, 1H), 7.13 – 7.07 (m, 2H), 7.03 (d, *J* = 7.9 Hz, 1H), 6.83 (d, *J* = 9.0 Hz, 1H), 3.74 (s, 3H), 3.54 (t, *J* = 7.2 Hz, 2H), 3.12 (t, *J* = 7.2 Hz, 2H).

<sup>13</sup>C NMR (101 MHz, acetone-d<sub>6</sub>) δ 170.1, 150.1, 147.6, 137.8, 128.5, 123.9, 123.6, 122.1, 119.4, 119.3, 115.7, 113.6, 113.4, 112.2, 110.4, 56.0, 44.6, 26.0.

**2-((2-(1***H***-Indol-3-yl)ethyl)amino)-5-(pentafluoro-λ<sup>6</sup>-sulfaneyl)benzoic Acid (compound 23)**. 40% yield over three steps. Colorless solid. m.p. 237-238 °C. <sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>) δ 13.31 (s, 1H), 10.89 (s, 1H), 8.40 (s, 1H), 8.15 (d, J = 2.9 Hz, 1H), 7.80 (dd, J = 9.5, 2.9 Hz, 1H), 7.59 (d, J = 7.8 Hz, 1H), 7.35 (d, J = 8.1 Hz, 1H), 7.24 (s, 1H), 7.08 (t, J = 6.9 Hz, 1H), 6.98 (t, J = 8.0 Hz, 1H), 6.92 (d, J = 9.5 Hz, 1H), 3.56 (t, J = 7.1 Hz, 2H), 3.05 (t, J = 7.0 Hz, 2H). <sup>19</sup>F NMR (376 MHz, DMSO-d<sub>6</sub>) δ 91.2, 90.8, 90.4, 90.0, 66.2, 65.8. <sup>13</sup>C NMR (101 MHz, DMSO-d<sub>6</sub>) δ 168.53, 152.08, 136.25, 131.26, 129.29, 126.99, 123.15, 120.98, 118.29, 111.36, 111.27, 111.01, 108.46, 42.74, 24.28.

**2-((2-(1-Methyl-1***H***-indol-3-yl)ethyl)amino)-5-nitrobenzoic Acid (compound 24)**. 44% yield. Yellow solid. m.p. 220-223 °C. <sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>) δ 13.45 (s, 1H), 8.87 (s, 1H), 8.63 (d, *J* = 2.8 Hz, 1H), 8.18 (dd, *J* = 9.4, 2.9 Hz, 1H), 7.61 (d, *J* = 7.9 Hz, 1H), 7.40 (d, *J* = 8.2 Hz, 1H), 7.24 (s, 1H), 7.15 (t, *J* = 7.1 Hz, 1H), 7.02 (t, *J* = 8.0 Hz, 1H), 6.96 (d, *J* = 9.5 Hz, 1H), 3.75 (s, 3H), 3.64 (q, *J* = 6.7 Hz, 2H), 3.06 (t, *J* = 7.0 Hz, 2H). <sup>13</sup>C NMR (101 MHz, DMSO-d<sub>6</sub>) δ 168.6, 154.5, 136.7, 134.5, 129.4, 128.6, 127.6, 127.3, 121.2, 118.6, 118.4, 111.6, 110.2, 109.6, 109.3, 43.1, 32.3, 24.1.

**2-((2-(Benzofuran-3-yl)ethyl)amino)-5-nitrobenzoic Acid (compound 25)**. 18% yield. Yellow solid. m.p. 205-207 °C. <sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>) δ 13.48 (s, 1H), 8.91 (s, 1H), 8.63 (d, *J* = 2.8 Hz, 1H), 8.18 (dd, *J* = 9.4, 2.9 Hz, 1H), 7.90 (s, 1H), 7.74 (d, *J* = 6.5 Hz, 1H), 7.56 (d, *J* = 8.2 Hz, 1H), 7.32 (td, *J* = 7.2, 1.4 Hz, 1H), 7.26 (td, *J* = 7.4, 1.1 Hz, 1H), 7.00 (d, *J* = 9.5 Hz, 1H), 3.72 (q, *J* = 6.7 Hz, 2H), 3.05 (t, *J* = 6.7 Hz, 2H). <sup>13</sup>C NMR (101 MHz, DMSO-d<sub>6</sub>) δ 168.6, 154.6, 154.5, 142.8, 134.7, 129.4, 128.6, 127.5, 124.4, 122.5, 119.9, 116.9, 111.7, 111.3, 41.7, 22.5.

**2-((2-(2-Methyl-1***H***-indol-3-yl)ethyl)amino)-5-nitrobenzoic Acid (compound 26)**. 79% yield. Yellow solid. m.p. 119-120 °C. <sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>) δ 13.44 (s, 1H), 10.76 (s, 1H), 8.78 (s, 1H), 8.61 (d, *J* = 2.9 Hz, 1H), 8.14 (dd, *J* = 9.4, 2.9 Hz, 1H), 7.48 (d, *J* = 7.8 Hz, 1H), 7.23 (d, *J* = 7.8 Hz, 1H), 6.99 (t, *J* = 6.8 Hz, 1H), 6.93 (t, *J* = 6.8 Hz, 1H), 6.88 (d, *J* = 9.5 Hz, 1H), 3.61 - 3.52 (m, 2H), 3.01 (t, *J* = 6.7 Hz, 2H), 2.34 (s, 3H). <sup>13</sup>C NMR (101 MHz, DMSO-d<sub>6</sub>) δ 168.6, 154.5, 135.3, 134.4, 132.8, 129.3, 128.6, 128.0, 120.1, 118.2, 117.3, 111.6, 110.4, 106.5, 42.9, 23.2, 11.2.

**2-((2-(4-Methyl-1***H***-indol-3-yl)ethyl)amino)-5-nitrobenzoic Acid (compound 27)**. 93% yield. Yellow solid. m.p. 253-256 °C. <sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>) δ 13.45 (s, 1H), 10.91 – 10.86 (m, 1H), 8.90 (s, 1H), 8.64 (d, *J* = 2.8 Hz, 1H), 8.17 (dd, *J* = 9.5, 2.9 Hz, 1H), 7.21 – 7.14 (m, 2H), 7.05 – 6.89 (m, 2H), 6.72 (d, *J* = 7.0 Hz, 1H), 3.65 (q, *J* = 6.6 Hz, 2H), 3.23 (t, *J* = 6.9 Hz, 2H), 2.65 (s, 3H), 2.62 (s, 2H). <sup>13</sup>C NMR (101 MHz, DMSO-d<sub>6</sub>) δ 168.6, 154.5, 136.7, 134.5, 129.4, 128.6, 125.4, 123.4, 121.0, 120.0, 111.8, 111.6, 109.4, 44.1, 26.0, 20.1.

**2-((2-(4-Chloro-1***H***-indol-3-yl)ethyl)amino)-5-nitrobenzoic Acid (compound 28)**. 68% yield. Yellow solid. m.p. 170-174 °C. <sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>) δ 11.29 (s, 1H), 8.67 (s, 1H), 8.00 (dd, *J* = 9.2, 2.6 Hz, 1H), 7.37 – 7.31 (m, 2H), 7.09 – 6.98 (m, 2H), 6.72 (d, *J* = 9.5 Hz, 1H), 3.57 (q, *J* = 6.6 Hz, 2H), 3.25 (t, *J* = 7.2 Hz, 2H).

**2-((2-(5-Methyl-1***H***-indol-3-yl)ethyl)amino)-5-nitrobenzoic Acid (compound 29)**. 33% yield. Yellow solid. m.p. 247-251 °C. <sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>) δ 13.42 (s, 1H), 10.74 (s, 1H), 8.87 (s, 1H), 8.63 (d, *J* = 2.9 Hz, 1H), 8.17 (dd, *J* = 9.5, 2.8 Hz, 1H), 7.35 (s, 1H), 7.23 (d, *J* = 8.2 Hz, 1H), 7.19 (s, 1H), 6.96 (d, *J* = 9.5 Hz, 1H), 6.90 (d, *J* = 9.1 Hz, 1H), 3.63 (q, *J* = 6.6 Hz, 2H), 3.03 (t, *J* = 6.8 Hz, 2H), 2.37 (s, 3H). **2-((2-(5-Chloro-1***H***-indol-3-yl)ethyl)amino)-5-nitrobenzoic Acid (compound 30)**. 37% yield. Yellow solid. m.p. 272-275 °C. <sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>) δ 13.42 (s, 1H), 11.10 (s, 1H), 8.88 (s, 1H), 8.63 (d, *J* = 2.9 Hz, 1H), 8.16 (dd, *J* = 9.5, 2.9 Hz, 1H), 7.66 (s, 1H), 7.36 (d, *J* = 8.6 Hz, 1H), 7.34 (s, 1H), 7.07 (dd, *J* = 8.6, 2.1 Hz, 1H), 6.96 (d, *J* = 9.6 Hz, 1H), 3.66 – 3.60 (m, 2H), 3.04 (t, *J* = 7.1 Hz, 2H).

**2-((2-(7-Methyl-1***H***-indol-3-yl)ethyl)amino)-5-nitrobenzoic Acid (compound 31)**. 47% yield. Yellow solid. m.p. 253-257 °C. <sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>) δ 13.42 (s, 1H), 10.86 (s, 1H), 8.86 (s, 1H), 8.63 (d, *J* = 2.8 Hz, 1H), 8.18 (dd, *J* = 9.5, 2.9 Hz, 1H), 7.42 (d, *J* = 7.2 Hz, 1H), 7.24 (s, 1H), 6.96 (d, *J* = 9.6 Hz, 1H), 6.92 – 6.86 (m, 2H), 3.67 – 3.61 (m, 2H), 3.06 (t, *J* = 6.9 Hz, 2H), 2.44 (s, 3H).

**2-((2-(7-Chloro-1***H***-indol-3-yl)ethyl)amino)-5-nitrobenzoic Acid (compound 32)**. 37% yield. Yellow solid. m.p. 275-279 °C. <sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>) δ 13.43 (s, 1H), 11.28 (s, 1H), 8.86 (s, 1H), 8.63 (d, *J* = 2.8 Hz, 1H), 8.18 (dd, *J* = 9.5, 2.8 Hz, 1H), 7.59 (d, *J* = 7.8 Hz, 1H), 7.34 (s, 1H), 7.17 (d, *J* = 7.5 Hz, 1H), 7.04 – 6.94 (m, 2H), 3.66 (q, *J* = 6.7 Hz, 2H), 3.08 (t, *J* = 6.9 Hz, 2H). <sup>13</sup>C NMR (101 MHz, DMSO-d<sub>6</sub>) δ 168.6, 154.5, 134.5, 133.0, 129.4, 129.0, 128.6, 124.7, 120.6, 119.4, 117.5, 115.8, 112.4, 111.7, 42.9, 24.2.

**2-((2-(6-Methyl-1***H***-indol-3-yl)ethyl)amino)-5-nitrobenzoic Acid (compound 33)**. 48% yield. Yellow solid. m.p. 222-224 °C. <sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>) δ 13.42 (s, 1H), 10.72 (s, 1H), 8.85 (s, 1H), 8.63 (d, *J* = 2.9 Hz, 1H), 8.18 (dd, *J* = 9.4, 2.9 Hz, 1H), 7.47 (d, *J* = 8.1 Hz, 1H), 7.17 - 7.11 (m, 2H), 6.95 (d, *J* = 9.5 Hz, 1H), 6.82 (dd, *J* = 8.1, 1.5 Hz, 1H), 3.63 (q, *J* = 6.7 Hz, 2H), 3.03 (t, *J* = 7.0 Hz, 2H), 2.38 (s, 3H).

**2-((2-(6-Chloro-1***H***-indol-3-yl)ethyl)amino)-5-nitrobenzoic Acid (compound 34)**. 31% yield. Yellow solid. m.p. 249-251 °C. <sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>) δ 13.43 (s, 1H), 11.05 (s, 1H), 8.85 (s, 1H), 8.62 (s, 1H), 8.17 (d, *J* = 7.3 Hz, 1H), 7.61 (d, *J* = 8.6 Hz, 1H), 7.39 (s, 1H), 7.30 (s, 1H), 7.03 – 6.92 (m, 2H), 3.64 (q, *J* = 6.6 Hz, 2H), 3.05 (t, *J* = 7.1 Hz, 2H). <sup>13</sup>C NMR (101 MHz, DMSO-d<sub>6</sub>) δ 168.6, 154.5, 136.6, 134.5, 129.4, 128.6, 125.9, 125.8, 124.5, 119.8, 118.7, 111.7, 111.3, 111.0, 109.1, 42.9, 24.1.

**2-((2-(6-Fluoro-1***H***-indol-3-yl)ethyl)amino)-5-nitrobenzoic Acid (compound 35)**. 42% yield. Yellow solid. m.p. 249-250 °C. <sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>) δ 10.96 (s, 1H), 8.85 (t, *J* = 5.2 Hz, 1H), 8.62 (d, *J* = 2.9 Hz, 1H), 8.16 (dd, *J* = 9.5, 3.0 Hz, 1H), 7.58 (dd, *J* = 8.8, 5.4 Hz, 1H), 7.24 (s, 1H), 7.11 (dd, *J* = 10.1, 2.4 Hz, 1H), 6.94 (d, *J* = 9.5 Hz, 1H), 6.83 (td, *J* = 9.2, 2.4 Hz, 1H), 3.63 (q, *J* = 6.6 Hz, 2H), 3.04 (t, *J* = 7.0 Hz, 2H). <sup>19</sup>F NMR (376 MHz, DMSO-d<sub>6</sub>) δ -122.3 (td, *J* = 9.7, 5.5 Hz).

**2-((2-(6-Bromo-1***H***-indol-3-yl)ethyl)amino)-5-nitrobenzoic Acid (compound 36)**. 91% yield. Yellow solid. m.p. 253-257 °C. <sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>) δ 13.43 (s, 1H), 11.05 (s, 1H), 8.85 (s, 1H), 8.63 (d, *J* = 2.8 Hz, 1H), 8.17 (dd, *J* = 9.4, 2.9 Hz, 1H), 7.57 (d, *J* = 8.5 Hz, 1H), 7.53 (s, 1H), 7.33 – 7.26 (m, 1H), 7.11 (dd, *J* = 8.4, 1.8 Hz, 1H), 6.96 (d, *J* = 9.6 Hz, 1H), 3.64 (q, *J* = 6.6 Hz, 2H), 3.05 (t, *J* = 7.0 Hz, 2H).

**2-((2-(6-Cyano-1***H***-indol-3-yl)ethyl)amino)-5-(pentafluoro-λ<sup>6</sup>-sulfaneyl)benzoic Acid (EF-4-177)**. 9.8% yield over five steps.<sup>37</sup> Colorless solid. m.p. 210-212 °C. <sup>1</sup>H NMR (400 MHz, DMSOd<sub>6</sub>) δ 13.31 (s, 1H), 11.50 (d, J = 2.6 Hz, 1H), 8.39 (s, 1H), 8.15 (d, J = 2.8 Hz, 1H), 7.85 (s, 1H), 7.84 – 7.75 (m, 2H), 7.57 (s, 1H), 7.31 (d, J = 8.3 Hz, 1H), 6.92 (d, J = 9.5 Hz, 1H), 3.58 (q, J =6.2 Hz, 2H), 3.07 (t, J = 7.0 Hz, 2H). <sup>19</sup>F NMR (376 MHz, DMSO-d<sub>6</sub>) δ 91.5, 91.1, 90.7, 90.3, 89.9, 66.2, 65.7. <sup>13</sup>C NMR (101 MHz, DMSO-d<sub>6</sub>) δ 168.5, 152.1, 139.0, 134.9, 131.2, 130.1, 129.3, 128.0, 121.0, 120.7, 119.6, 116.3, 112.4, 111.3, 108.5, 102.3, 42.7, 23.9.

#### Supporting Information

Representative *p*-Cl-ANS data for screening hits, HSQC NMR data for allosterically bound ligands, four additional crystal structures, ITC data, crystallography conditions table, NMR spectra and qHNMR purity data of newly described compounds.

#### PDB ID Codes

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#### Abbreviations Used

ANS, 8-anilino-1-naphthalene sulfonic acid; CDK1, cyclin-dependent kinase 1; CDK2, cyclindependent kinase 2; CL<sub>int</sub>, clearance; D<sub>7.4</sub>, distribution constant at pH 7.4; Em, emission wavelength; EtOAc, ethyl acetate; Ex, excitation wavelength; HEPES, 4-(2-hydroxyethyl)-1piperazineethanesulfonic acid; ITC, isothermal titration calorimetry; K<sub>D</sub>, dissociation equilibrium constant; MeOH, methanol; MM-GBSA, molecular mechanics with generalized Born and surface area solvation; NPPB, 5-nitro-2-(3-phenylpropylamino)benzoic acid; *p*-Cl-ANS, 8-(4chloroanilino)-1-naphthalene sulfonic acid; qHNMR, quantitative <sup>1</sup>H NMR; S<sub>n</sub>Ar, nucleophilic aromatic substitution; SP, standard precision; vHTS, virtual high-throughput screen; XP, extra precision <u>References</u>

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### Table of Contents Graphic

