

Table S1. Physicochemical properties of tests solutes

Fragment	Name	Cas Number	log D (pH 7.4)	Charge	H-Bond Donor Count (PubChem)	H-Bond Acceptor Count (PubChem)
1	Benzylamine	100-46-9	-0,56	cationic	1	1
41	5-Phenyl-2-furoic acid	52938-97-3	-0,35	anionic	1	3
54	4-(Aminomethyl)pyridine	3731-53-1	-1,3	cationic	1	2
57	Histamine	51-45-6	-2,71	cationic	2	2
62	2-Phenylimidazole	670-96-2	1,64	neutral	1	1
66	1,1,4,7,10,10-Hexamethyltriethylenetetramine	4856-97-7	0,57	neutral	2	2
70	4-(2-Keto-1-benzimidazolyl)piperidine	20662-53-7	-1,13	cationic	2	2
74	2,2'-(Ethane-1,2-diyl)dianiline	34124-14-6	1,99	neutral	2	2
88	Thieno[3,2-b]pyridin-7-ol	107818-20-2	-0,34	neutral	1	3
93	4,4'-Dihydroxybiphenyl	92-88-6	2,26	neutral	2	2
117	2-Amino-3-benzoyloxy pyridine	24016-03-3	1,53	neutral	1	3
125	1-Methyl-2-pyrrolicarboxylic acid	6973-60-0	-1,86	anionic	1	2
139	2-Phenyl-2-imidazoline	936-49-2	-0,59	cationic	1	1
150	Salicylic acid	69-72-7	-0,77	anionic	2	3
153	2,4-Dihydroxybenzophenone	131-56-6	3,16	anionic	2	3
159	3-Furoic acid	488-93-7	-1,97	anionic	1	3
161	Sulfacetamide	144-80-9	-2,17	anionic	2	4
168	3-Cyano-6-methyl-2(1H)-pyridinone	4241-27-4	-0,29	neutral	1	2
169	Sulfabenzamide	127-71-9	-0,62	anionic	2	4
195	1-Benzoylpiperidine	776-75-0	1,95	neutral	0	1
209	Biphenyl-4-carboxylic acid	92-92-2	0,67	anionic	1	2
266	Benzhydrylamine	91-00-9	1,65	cationic	1	1
271	1,2,3,4-Tetrahydro-9-aminoacridine	321-64-2	0,89	cationic	1	2
286	(1R,2S)-(-)-2-Amino-1,2-diphenylethanol	23190-16-1	-0,08	cationic	2	2
288	2,3-Diaminopyridine	452-58-4	0,12	cationic	2	3
294	1-(1H-Pyrrol-2-yl)ethanone	1072-83-9	0,99	neutral	1	1
297	5-Amino-1-phenylpyrazole-4-carboxamide	50427-77-5	0,83	neutral	2	3
298	1-Phenyl-1,2,3,4-tetrahydropyrrolo[1,2-a]pyrazine	112758-89-1	2,53	cationic	1	1
302	4-Aminobiphenyl	92-67-1	2,71	neutral	1	1
336	trans-Ferulic acid	537-98-4	-1,38	anionic	2	4
346	2-Thiouracil	141-90-2	-3,6	neutral	2	2
366	3-Amino-2-chloropyridine	6298-19-7	0,97	neutral	1	2
368	4-Aminopyridine	504-24-5	-2,41	cationic	1	2
378	2,6-Dichloropyridine	2402-78-0	2,1	neutral	0	1
393	1-(4-Sulfophenyl)-3-methyl-5-pyrazolone	89-36-1	-4,31	anionic	1	5
411	Ethyl 5-amino-1-phenyl-4-pyrazolecarboxylate	16078-71-0	2,79	neutral	1	4
412	2-(carboxymethylthio)-4-methylpyrimidine	46118-95-0	-3,15	anionic	1	5
445	4-Aminohippuric acid	61-78-9	-4	anionic	3	4
449	Orotic acid	65-86-1	-4,75	anionic	3	4
725	Adenine	73-24-5	-2,06	neutral	2	4
771	4-Chlorophenyl methyl sulfone	98-57-7	1,45	neutral	0	2

Fragment	Name	Cas Number	log D (pH 7.4)	Global charge	H-Bond Donor Count (PubChem)	H-Bond Acceptor Count (PubChem)
1	Benzylamine	100-46-9	-0,56	1	1	1
41	5-Phenyl-2-furoic acid	52938-97-3	-0,35	-1	1	3
54	4-(Aminomethyl)pyridine	3731-53-1	-1,3	1	1	2
57	Histamine	51-45-6	-2,71	1,8	2	2
62	2-Phenylimidazole	670-96-2	1,64	0,3	1	1
66	1,1,4,7,10,10-Hexamethyltriethylenetetramine	4856-97-7	0,57	0	2	2
70	4-(2-Keto-1-benzimidazolyl)piperidine	20662-53-7	-1,13	1	2	2
74	2,2'-(Ethane-1,2-diyl)dianiline	34124-14-6	1,99	0	2	2
88	Thieno[3,2-b]pyridin-7-ol	107818-20-2	-0,34	0	1	3
93	4,4'-Dihydroxybiphenyl	92-88-6	2,26	0	2	2
117	2-Amino-3-benzoyloxy pyridine	24016-03-3	1,53	0	1	3
125	1-Methyl-2-pyrrolicarboxylic acid	6973-60-0	-1,86	-1	1	2
139	2-Phenyl-2-imidazoline	936-49-2	-0,59	1	1	1
150	Salicylic acid	69-72-7	-0,77	-1	2	3
153	2,4-Dihydroxybenzophenone	131-56-6	3,16	-0,5	2	3
159	3-Furoic acid	488-93-7	-1,97	-1	1	3
161	Sulfacetamide	144-80-9	-2,17	-1	2	4
168	3-Cyano-6-methyl-2(1H)-pyridinone	4241-27-4	-0,29	-0,3	1	2
169	Sulfabenzamide	127-71-9	-0,62	-1	2	4
195	1-Benzoylpiperidine	776-75-0	1,95	0	0	1
209	Biphenyl-4-carboxylic acid	92-92-2	0,67	-1	1	2
266	Benzhydrylamine	91-00-9	1,65	1	1	1
271	1,2,3,4-Tetrahydro-9-aminoacridine	321-64-2	0,89	1	1	2
286	(1R,2S)-(-)-2-Amino-1,2-diphenylethanol	23190-16-1	-0,08	1	2	2
288	2,3-Diaminopyridine	452-58-4	0,12	0,5	2	3
294	1-(1H-Pyrrol-2-yl)ethanone	1072-83-9	0,99	0	1	1
297	5-Amino-1-phenylpyrazole-4-carboxamide	50427-77-5	0,83	0	2	3
298	1-Phenyl-1,2,3,4-tetrahydropyrrolo[1,2-a]pyrazine	112758-89-1	2,53	0,7	1	1
302	4-Aminobiphenyl	92-67-1	2,71	0	1	1
336	trans-Ferulic acid	537-98-4	-1,38	-1	2	4
346	2-Thiouracil	141-90-2	-3,6	-0,2	2	2
366	3-Amino-2-chloropyridine	6298-19-7	0,97	0	1	2
368	4-Aminopyridine	504-24-5	-2,41	1	1	2
378	2,6-Dichloropyridine	2402-78-0	2,1	0	0	1
393	1-(4-Sulfophenyl)-3-methyl-5-pyrazolone	89-36-1	-4,31	-1	1	5
411	Ethyl 5-amino-1-phenyl-4-pyrazolecarboxylate	16078-71-0	2,79	0	1	4
412	2-(carboxymethylthio)-4-methylpyrimidine	46118-95-0	-3,15	-1	1	5
445	4-Aminohippuric acid	61-78-9	-4	-1	3	4
449	Orotic acid	65-86-1	-4,75	-1	3	4
725	Adenine	73-24-5	-2,06	0	2	4
771	4-Chlorophenyl methyl sulfone	98-57-7	1,45	0	0	2

- Nano-FAC data processing for column characterization (protein specific interactions (active binding sites, affinity (dissociation constant)) and non-specific interactions)

Nano-FAC experiments were used to determine the quantity of ligand capture at different ligand concentrations (q_{captured}). In staircase method, increasing concentrations of ligand were continuously injected into the nanocolumn (Figure S1 A). This allows to determine the cumulative captured quantities ($q_{\text{captured cumul.}}$) of ligand at each concentration step (n).

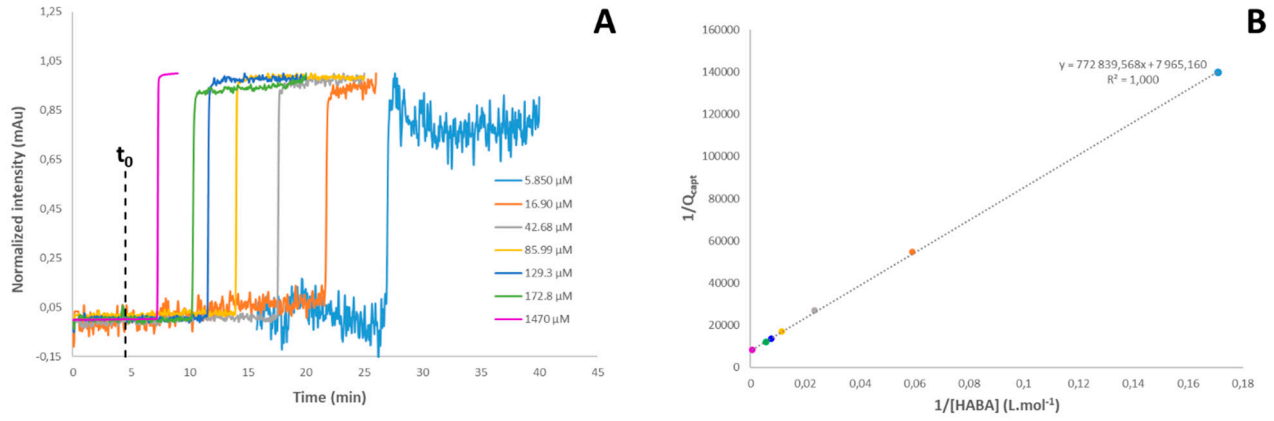


Figure S1. Frontal affinity chromatograms of 4'-Hydroxyazobenzene-2-carboxylic acid (HABA) on streptavidin-functionalized poly(DHPMA-co-MBA) monolith. A) Breakthrough curves of HABA at different concentrations in phosphate buffer (67mM, pH 7.4); UV detection at 347 nm; short injection of DMSO 0.1 % in water was used to mark the dead time. B) Plot of the reciprocal of the quantity of the specific captured quantity of HABA minus the non-specific captured quantity of HABA ($1/(\text{cumul}Q_{\text{capt}} - (K_{\text{NS}} \times [\text{HABA}]))$) (μmol^{-1}) versus the reciprocal of HABA concentration ($1/[\text{HABA}]$) (L.mol^{-1})

If only specific interactions are observed, the plot of $1/q_{\text{Cumul captured}}(n)$ vs $1/[L]_n$ is linear. A deviation from linearity in the high concentration region of the plot indicates the presence of a non-specific interaction contribution. For unspecific interactions (if any), $q_{\text{Cumul captured}}(n)$ is the sum of the amounts bound by specific interactions $\frac{B_{\text{act}} \times [L]_n}{K_d + [L]_n}$ and the amount of ligand captured by non-specific interactions $K_{\text{non-specific}} \times [L]_n$, leading to equation S1:

$$q_{\text{Cumul captured}}(n) = \frac{B_{\text{act}} \times [L]_n}{K_d + [L]_n} + K_{\text{non-specific}} \times [L]_n \quad (\text{Equation S1})$$

Assuming $K_{\text{non-specific}}$ is constant, equation S1 can be written as follows:

$$\frac{1}{q_{\text{captured}} - K_{\text{non-specific}} \times [L]_n} = \frac{K_d}{B_{\text{act}}} \times \frac{1}{[L]_n} + \frac{1}{B_{\text{act}}} \quad (\text{Equation S2})$$

The excel solver function is used to compute the $K_{\text{non-specific}}$ factor for which the best r^2 value is reached.

In case of presence of non-specific interaction, to determine the number of active site (B_{act}) and dissociation constant properly the plot of $1/[q_{\text{cumul Capt}}(n) - (K_{\text{NS}} \times [L]_n)]$ vs $1/[L]_n$ was used (figure S1B right).