# Development of new 5-HT<sub>6</sub>R ligand series illustrates the search for selective transmembrane proteins ligands

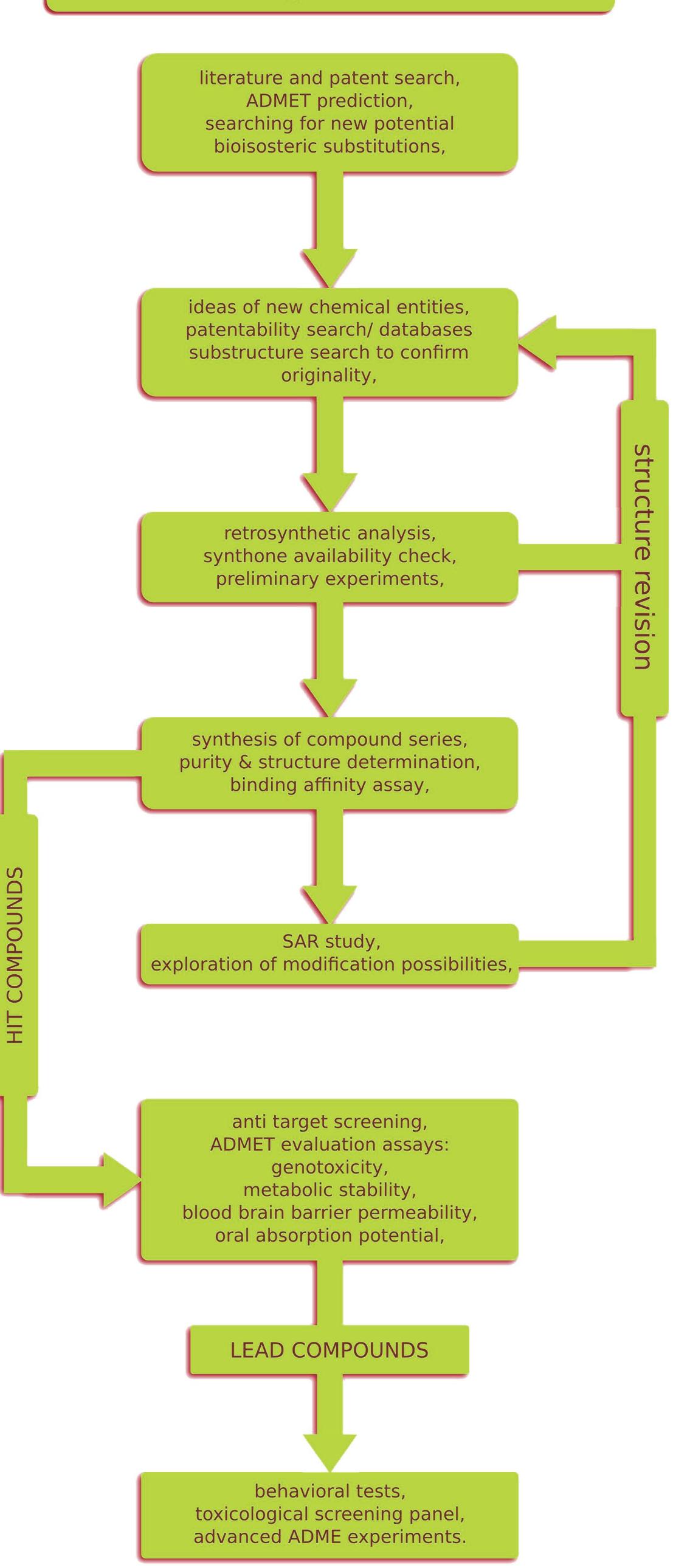
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## How does our drug candidate search work



#### Refferences:

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#### 5-HT<sub>6</sub> receptor - an emerging drug target for Alzheimer's disease

5-HT6R discovered in the early 90's has proven to be an important target for neurodegenerational disorders. 5-HT<sub>6</sub>R blockade induces acetylcholine release which might restore function in deteriorated cholinergic system of Alzheimer's Disease (AD) patients.<sup>2</sup> The receptor is expressed almost exclusively in brain which implicates low probability of peripheral side effects. 5-HT<sub>6</sub> antagonists administration can signifficantly improve cognitive performance in AD patients.<sup>3</sup> Two 5-HT<sub>6</sub> antagonists (Lu AE58054, SB-742457) have completed phase II clinical trials as an augmentation therapy for Alzheimer's disease.

### 5-HT<sub>6</sub>receptor ligand series development

Two independent series of selective 5-HT<sub>6</sub>R ligands consisting of 33 and 37 compounds respectively have been developed.

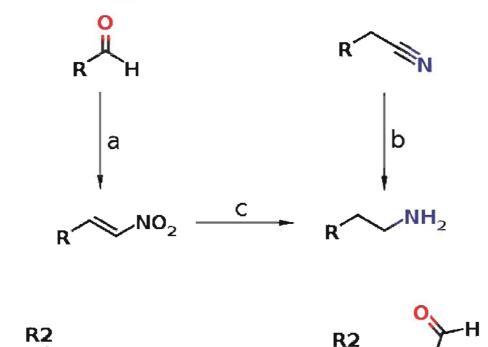
Pilot compound of serie 1, AH-122 emerged from analysis of pharmacological profile of an ultrapotent 5-HT<sub>2A</sub>R agonist, CIMBI-5<sup>4,5,6</sup> (Fig. 1). Replacement of the benzylic fragment of CIMBI-5 with an 3-indolemethyl scaffold resulted in a steep drop of 5-HT<sub>2A</sub> activity, while 5-HT<sub>6</sub>R affinity was

enhanced. To reveal the optimal substitution pattern for 5-HT<sub>6</sub>R binding, 33 compounds have been synthetised till date. This was achieved thanks to the very concise exploiting H3C synthetic protocol commercialy aviable arylacetonitriles developed in ourlab (Fig. 2). It was found that proper substitution of the

H<sub>3</sub>C~O AH-122 Cimbi-5

Fig. 1: non-classical bioisosteric substitution yields a hit compound

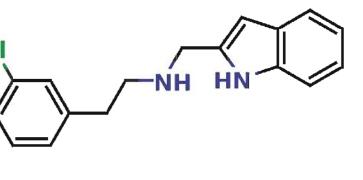
phenylethylamine backbone is crucial for 5-HT<sub>6</sub>R activity (Tab. 1). To our HNdelight, two additional serotonin receptor hit compounds werefound in radioligand binding assays (Fig. 3). A virtual combinatorial library consisting of 3364 structures was generated using Jchem Reactor tool.



Virtual docking protocol (5-HT<sub>6</sub>R, 5-HT<sub>2A</sub>R, 5-HT<sub>2C</sub>R homology models used) was performed using Glide software. Our plan is to investigate binding mode(s) of this serie by docking results of CI analyzing determined compounds with binding affinities. Next, rigid, constrained conformationally analogues will be synthetised to bring additional evidence. The synthesis and evaluation of

hit compound AH-246 derivatives Fig. 3: may bring an insight into 5-HT<sub>1A</sub>R R1 binding mechanisms.

AH-226  $5-HT_{2A}R K_i=20 nM$ 



AH-246  $5-HT_{1A}R K_i=19 nM$ 

5-HT<sub>2A</sub>R and 5-HT<sub>1A</sub>R hit compounds - byproducts of the research

Br; R3 = H, Me

Serie 2, opened with hit compound AH-42 (10nM at 5-HT6R), consists of 37 compounds with 5-HT6R binding affinities in the range of 2 - 455 nM. The selectivity over related targets (5-HT1A, 5-HT2A, 5-HT7 and D2 **Further** receptors) is excellent. revealed experiments good metabolic stability, lack of (Ames test) genotoxicity and selectivity over wide range of targets (two compounds screened for H1, H3, M1, M5, 5-HT3, alpha1, alpha3,). Functional assay confirmed that the compounds act as antagonists of 5-New compounds HT6R. synthetised in order to improve ADME properties. Lead structures will go through a panel of additional invitro and in-vivo experiments to evaluate drug-likeness.

ID	R1 N R2		K <sub>i</sub> [nM]				
	R1	R2	5-HT <sub>1A</sub>	5-HT <sub>2A</sub>	5-HT <sub>6</sub>	5-HT <sub>7</sub>	$\mathbf{D}_2$
CIMBI-5	0		3552	1 (0,04 *)	41 (73 *)	>10000	_
AH-122	· C	O H	>10000	300	39	>10000	1522
AH-120	0	₩ PH	4214	1	87	6224	1488
AH-125			2721	1	46	3875	938
AH-189	0	₩H	-	64	543	>10000	5223
AH-190		O NH	-	1238	310	7556	3572
AH-195		NH NH	398	_	534	-	% <b>-</b> -
AH-196	· Co		1895	424	318	4257	_
AH-185		O NH	>10000	610	1179	5338	-
AH-184		₩H	-	2394	38	5888	7115
AH-186		NH NH	-	3180	39	4182	2071
AH-199		• NH	5730	-	703	>10000	-
AH-228		Br	3795	1099	246	4166	-
AH-226	₩H	₩H	1531	20	172	2861	-
AH-236	cı	NH NH	173	3422	1605	1559	-
AH-246	CI	NH	19	2006	475	923	-

Tab. 1: Binding affinities of selected compounds from serie 1

data from ChEMBL database

