First in Class Dual Non-ATP Competitive Glycogen Synthase Kinase 3b **/ Histone Deacetylase Inhibitors as a Potential Therapeutic to Treat Alzheimer's Disease**

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Compounds	$%$ inhibition at 1 µM ^a			
	$GSK-3\beta$	HDAC2	HDAC ₆	
1	44.78±22.43	36.24 ± 13.40	36.99 ± 5.59	
$\mathbf{2}$	47.31 ± 4.57	21.95 ± 2.43	37.23 ± 2.51	
3	54.64 ± 6.07	55.93 ± 5.02	78.43±0.74	
$\boldsymbol{4}$	95.33 ± 0.61	88.43 ± 0.49	89.39 ± 0.82	
5	87.81 ± 2.90	86.65 ± 2.80	87.68 ± 0.23	
6	91.93 ± 3.02	85.70 ± 3.87	88.36 ± 0.29	
7	35.65±3.96	63.07 ± 2.16	44.84±3.08	
8	94.45 ± 0.73	88.47±1.25	86.80 ± 0.40	
Tideglusib	n.d. ^b	63.68 ± 5.00	64.34±2.07	
Vorinostat	n.a. ^c	$t.i^d$	$\frac{1}{2}$ i. $\frac{d}{dx}$	
Nexturastat A	37.60%	n.d. ^b	n.d. ^b	

1. Table SI1. Effects of compounds **1-8**, tideglusib, vorinostat and nexturastat A, on GSK-3β, HDAC2, and HDAC6 activity.

a. % of activity inhibition at 1 μM of compounds are reported as a mean value of at least three determinations; b. not determined; c. not active up to 50 μ M; d. t.i.: total inhibition.

Figure SI1. Protein surface and electrostatic potential map comparison between HDAC2 and HDAC6. HDAC2 structure in complex with SAHA (PDB ID: 4LXZ) and HDAC6 structure in complex with TSA (PDB ID: 5EDU).

3.

Figure SI2. HDAC2 and HDAC6 Molecular Interaction Field (MIF) pocket analysis. A) HDAC2 in complex with SAHA (PDB ID: 4LXZ); B) HDAC6 in complex with TSA (PDB ID: 5EDU). Red and blue contours identify respectively regions in which interactions with protein H-bond donors and acceptors are favorable. Potential protein hydrophobic interaction regions are highlighted

by yellow contours.

Figure SI3. Contact map of compound **5** in HDAC2 and HDAC6. Percentage of MD frames for each residue in which the center of mass is located within 5.0 A of the ligand.

Figure SI4. Solvent Accessible Surface Area (SASA) analysis of compound 5 in HDAC2 and HDAC6. SASA hydrophobic, hydrophilic and total values along the 50ns MD simulation.

Figure SI5. Protein RMSD analysis of compound **5** in HDAC2 and HDAC6. RMSD values of the protein backbone along the 50ns MD simulation

7.

Donor-acceptor	H-bond occupancy	Donor-acceptor	H-bond occupancy (%)
	(%)	610HIP(HE2) - $5(01)$	82.1
308 TYR(HH) - 5(O1)	98.9	782TYR(HH) – $5(02)$	81.5
145HIP(HE2) - 5(O2)	68.5	$5(H5) - 611HID(NE2)$	40.6
$5(H5) - 146HID(NE2)$	52.1	568SER(HG) - 5(O3)	9.6

Figure SI6. Donor-acceptor pairs in HDAC2 and HDAC6 complexes with compound **5**. Pairs identified along the 50ns MD simulation in which the percentage of frames satisfying H-bond distance and angle criteria were higher than 5%.

Figure SI7. SH-SY5Y cell viability was assessed through MTT assay after 24 h of treatment with: vehicle (0.1% DMSO) or increasing concentrations (0.1 μM, 1 μM, 10 μM, 25 μM, 50 μM, and 100 μ M) of compound 4 (n = 10), tideglusib (n = 5), nexturastat A (n = 5), or vorinostat (n = 5). The graph shows the dose-dependent effect of treatments as a percentage of the vehicle-treated condition (0 μM). Values are given as means $±$ SEM. *p < 0.05, **p < 0.01, ***p < 0.001. Turkey's test after twoway ANOVA.

Figure SI8. A) Effect of treatment with Nexturastat A and Tideglusib on GSK-3β phosphorylation, histone H3 acetylation, and alpha tubulin acetylation in SH-SY5Y cells. Western blot analysis of phospho-GSK3β (Ser9; P-GSK-3β), acetylated H3 (H3Ac) and alpha tubulin (TubAc) levels in protein extracts from SH-SY5Y cells treated with vehicle (0.1% DMSO) or different concentrations (0.1 μM, 1 μM and 10 μM) Nexturastat A or Tideglusib for 24 h. Immunoblots are examples from two biological replicates of each experimental condition. B-D) Graphs show P-GSK-3B protein levels normalized to corresponding total protein levels (B), acetylated H3 levels normalized to total

H3 levels (C), and acetylated alpha tubulin (TubAc) levels normalized to GAPDH levels in cells treated as in A (C). Data in B-D are expressed as a percentage of vehicle-treated cells. Values are represented as means \pm SEM. *p < 0.05, ** p < 0.01, *** p < 0.001. Fisher's LSD test after two-way ANOVA.

10.

a)

Vorinostat

Tideglusib

Figure SI9. Effect of a) Vorinostat, and b) Tideglusib on murine N9 cells following induction into M1 activation state by LPS (100 ng/mL) treatment. iNOS, TREM2, and TGFβ2 expression were analyzed using Western blot after a 24 h treatment with LPS and increasing concentrations $(0.1, 1, \text{ and } 10 \,\mu\text{M})$. For the indirect determination of Nitric Oxide (NO) release, nitrites derived from N9-cultured media were quantified through Griess reaction. Results are presented as means \pm SE from three independent experiments. Statistical significance was calculated using two-way ANOVA (Dunnett's post-hoc comparison test): # $P \le 0.05$, ## $P \le 0.01$, ### $P \le 0.001$, compared to LPS-activated microglia; * P $<$ 0.05, ** P $<$ 0.01, *** P $<$ 0.001, compared to control.

Table SI2. Assessment of compound **4** and parent compounds tideglusib and vorinostat with respect to some Molecular-Structural and Physicochemical properties.

 $\frac{1}{a}$. Obtained from http://www.swissadme.ch/, with the exception of b. that was obtained from

ChemDraw 21.0.0.

11.

12. Representative copies of 1 H-NMR and 13C-NMR spectra.

Compound 2. 1 H-NMR (400 MHz) and 13C-NMR (100 MHz) in DMSO-d.

Compound 3. 1 H-NMR (400 MHz) and 13C-NMR (100 MHz) in DMSO-d

Compound 5. 1 H-NMR (400 MHz) and 13C-NMR (100 MHz) in DMSO-d

AS319_H1 AS 319 C

Compound 6. 1 H-NMR (400 MHz) and 13C-NMR (100 MHz) in DMSO-d

13.

Table SI3. List of antibodies used.

