Supporting Information

sp²-Iminosugars Targeting Human Lysosomal β-Hexosaminidase as Pharmacological Chaperone Candidates for Late-Onset Tay-Sachs Disease

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Compound	Commercial β -hexosaminidases (K_i , μ M)		
	Human	Bovine	Jack bean
	placenta	kidney	
DGJNAc			
thioureas			
1	4.8±0.2	3.5±0.2	3.0±0.1
2	3.0±0.1	2.2±0.1	7.2±0.2
3	16.0±0.5	6.3±0.2	5.1±0.2
4	0.24±0.02	0.14±0.01	0.065 ± 0.003
DGJNAc			
isothioureas			
5	0.055±0.002	0.024 ± 0.002	0.14 ± 0.01
6	0.026±0.001	0.015±0.001	0.048 ± 0.002
7	0.026±0.001	0.020±0.001	0.088 ± 0.003
8	0.044 ± 0.002	0.046 ± 0.002	0.13±0.01

Table S1. K_i or IC₅₀ values for sp²-iminosugars **1-8** against commercial β -*N*-acetylhexosaminidases.

^a K_i data are presented as the mean \pm SD (n = 3). Inhibition was competitive in all cases. No inhibition was observed for any compound at 1 mM concentration on baker's yeast isomaltase, *Aspergillus niger* amyloglucosidase, green coffee bean α -galactosidase, Jack bean α -mannosidase, and Helix pomatia β -mannosidase.

Cell-based recovery of HexA in TSD-GS fibroblast



Figure S1. Immunoblot of cell lysates for chaperoning effects of compound **6** in TSD patient fibroblast. The expression levels of HexA were determined in the control and TSD-GS fibroblast by Western blotting. Tubulin was used as a loading control.



Figure S2. ¹H and ¹³C NMR spectra (300 MHz, 75.5 MHz, CD₃OD) of **1**.





NHCOCH₃



Figure S5. ¹H and ¹³C NMR spectra (300 MHz, 75.5 MHz, CD₃OD) of 4











Figure S10. (A) Binding of **NAG-thiazoline** (green sticks) to HexA (tan sticks) from the crystallographic complex (PDB ID 2GK1). (B) Overlay of crystallographic **NAG-thiazoline** (green sticks) and docked **NAG-thiazoline** (magenta sticks) bound to HexA. Hydrogen atoms from ligands are not shown for clarity. Hydrogen bonds are represents as black dashed lines and sulfur- π interactions in red dashed lines.



Figure S11. Immunoblot assay using anti-OGA to test the purity of hOGA. The major band at approximately 130 kD correspond to the dimeric form of the enzyme. Lower molecular mass bands arise from truncated forms of the enzyme.

References

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