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Background

- Lysosomal storage disorders (LSDs) are complex metabolic disorders characterised by an abnormal build-up of toxic materials within lysosomes as a result of lysosomal enzyme defects and deficiencies¹. Due to the complex nature of these disorders, there is currently no available treatment for the majority of LSDs.
- Acid ceramidase (ACase) is a lysosomal hydrolase that hydrolyses ceramide into sphingosine and a free fatty acid². However, ACase can also hydrolyse accumulating glycosphingolipids into lyso-glycosphingolipids (Fig. 1), a biomarker of LSDs³.
- Studies have shown that inhibition or genetic removal of ACase prevents the formation of lyso-glycosphingolipids in Gaucher, Fabry⁴ and Krabbe disease⁵, thus ameliorating lyso-glycosphingolipid induced pathology.
- ACase therefore makes a promising therapeutic target as a potential treatment for LSDs.

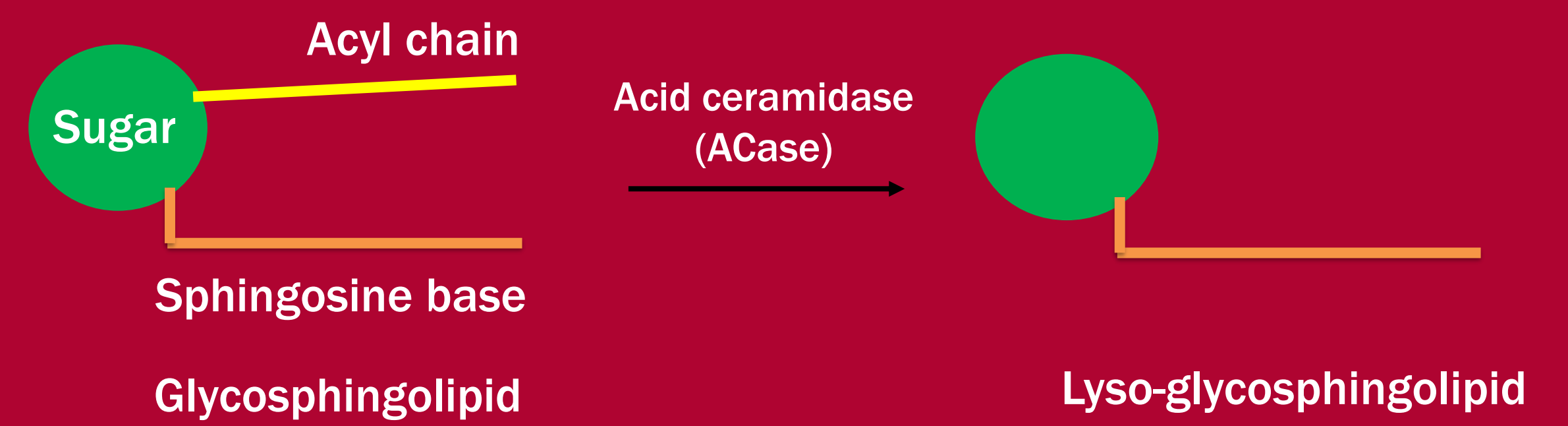


Figure 1. Simplified overview of the ACase mechanism in LSDs. In LSDs, glycosphingolipids accumulate as a result of lysosomal enzyme defect or deficiency. ACase deacylates accumulating glycosphingolipids into lyso-glycosphingolipids, a biomarker of LSDs.

Why a non-covalent inhibitor?

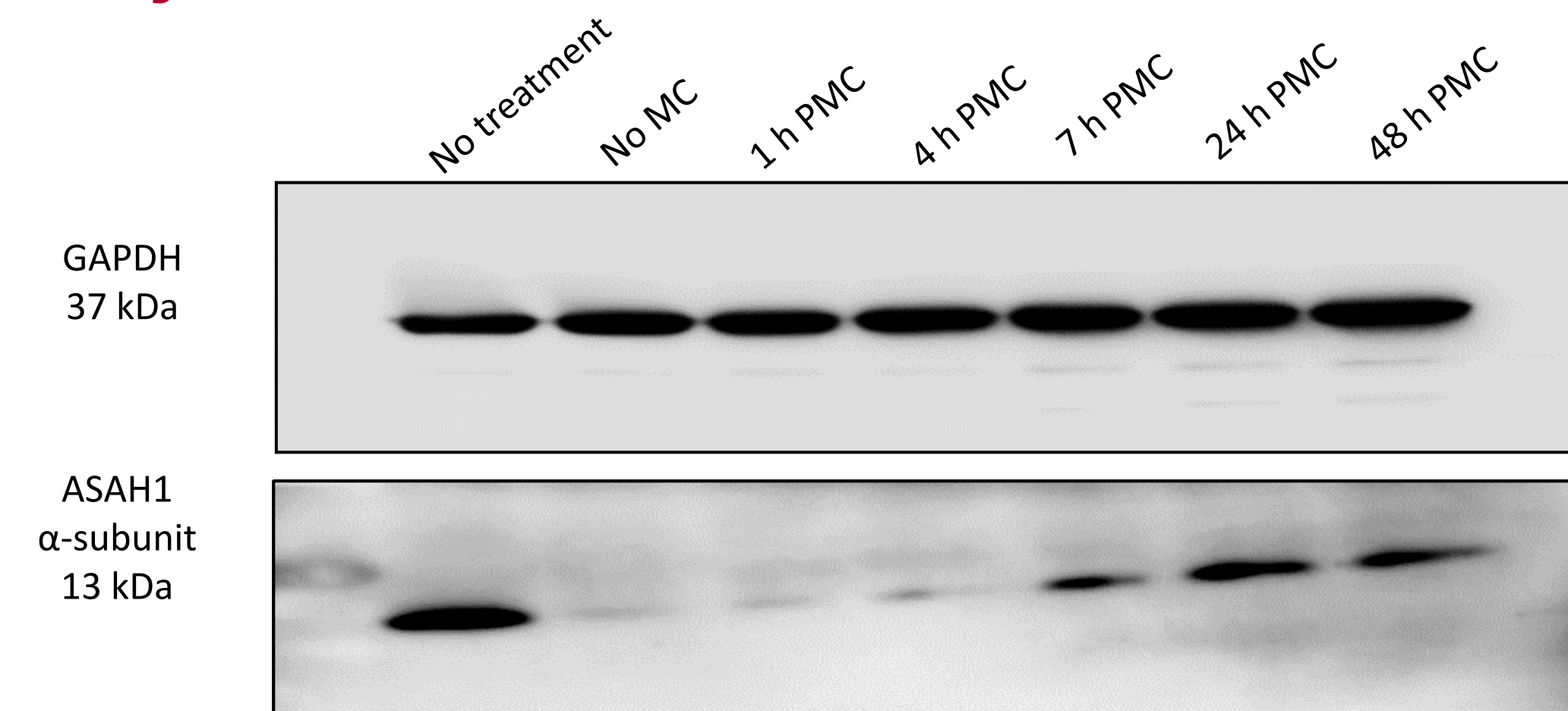


Figure 2. ACase has a turnover time of 24 h. Cells were treated with 15 μ M tamoxifen for 24 h. After 24 h, the media was removed and replaced with fresh media. Cells were collected 1, 4, 7, 24 and 48 h post-media change (MC). Post-media change (PMC).

- Over-inhibition of ACase causes the development of Farber disease, an aggressive LSD.
- Our data shows that ACase has a relatively slow turnover time of 24 h (Fig. 2), corroborating our aim to find a non-covalent inhibitor of ACase.

Project aim

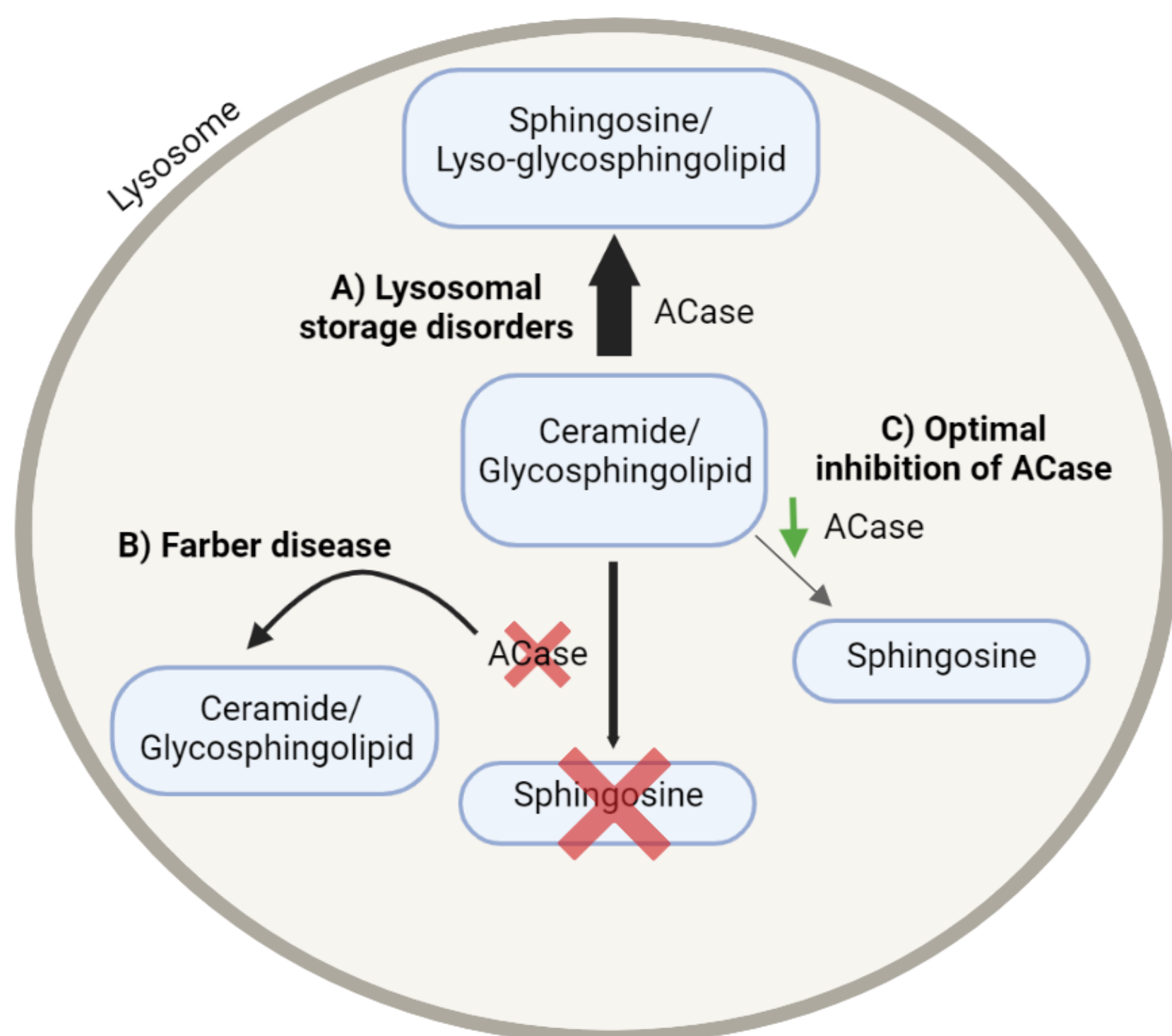


Figure 3. Simplified overview of the overall project aim. Our aim is inhibit ACase enough to alleviate lyso-glycosphingolipid induced pathology, but not enough to induce Farber disease. Image created using BioRender.

Conclusions

- Compound A provides a suitable and promising starting point to find a compound that will inhibit ACase as a mechanism to treat LSDs.

Future directions

- Establish crystallisation conditions for ACase to aid drug design.
- Determine which LSDs are likely to benefit from ACase inhibition.

Results

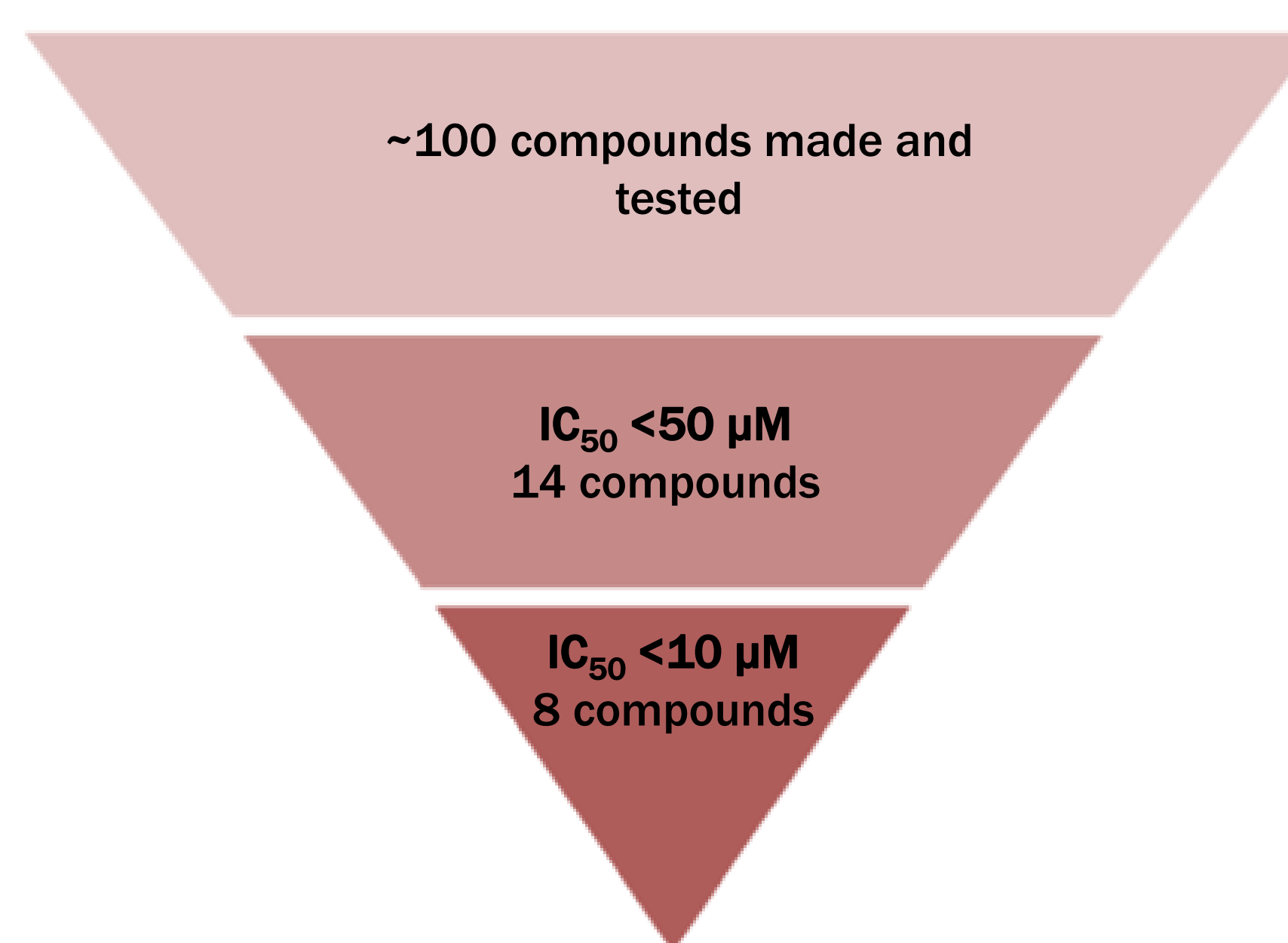


Figure 4. Brief overview of the number of compounds tested, and the number of compounds that have an IC_{50} less than 50 μ M and 10 μ M.

- A fluorescent-based biochemical assay⁶ has been established in-house to measure ACase activity.
- Approximately 100 compounds have been made, and their effectiveness as ACase inhibitors have been tested (Fig. 4).
- 14 and 8 compounds can inhibit ACase activity with an IC_{50} of <50 μ M and <10 μ M respectively.
- Compound A is our most promising compound with an IC_{50} of 1.4 μ M (Fig. 5A).

Compound A: biochemical assays

- With an IC_{50} of 1.4 μ M, compound A is our lead compound (Fig. 5A).
- Exploratory enzyme kinetic experiments have shown that compound A is a non-competitive inhibitor of ACase (Fig. 5B).

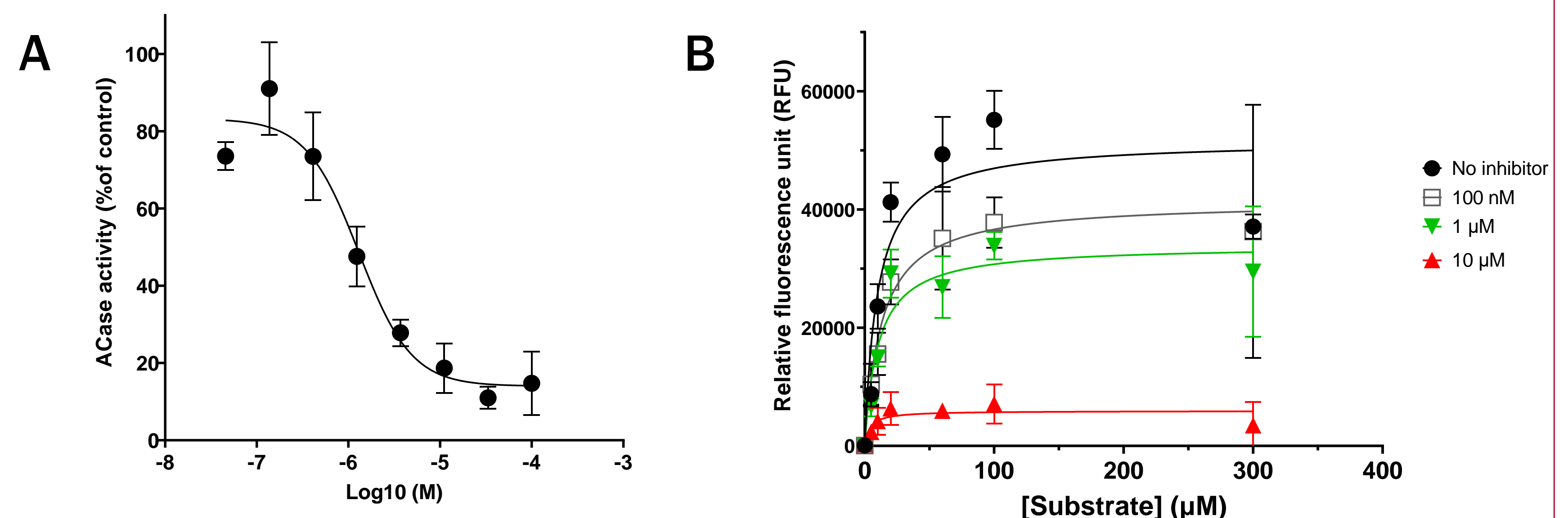


Figure 5. Compound A is a non-competitive inhibitor of ACase with an IC_{50} value of 1.4 μ M. (A) Effects of compound A on ACase activity. (B) Michaelis-Menten analysis of ACase in the presence of 10 μ M, 1 μ M and 100 nM of Compound A. Results are presented as mean \pm SD ($n = 3$).

Compound A: biophysical assay

- A biophysical assay was performed to help our understanding of the interactions between ACase and our compounds at a molecular level.
- Using the WAVE, we found that Compound A binds non-covalently to ACase (Fig. 6).
- Further assay optimisation is needed to obtain association and dissociation rates of Compound A and ACase.

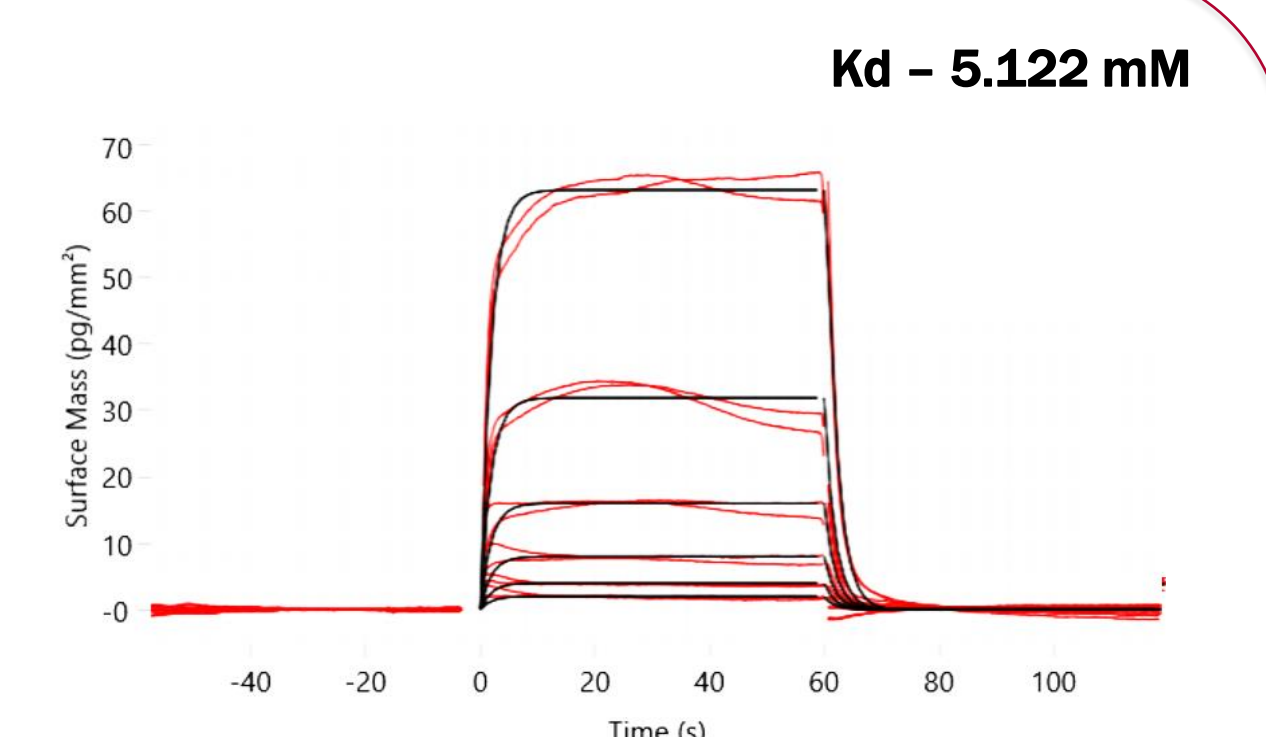


Figure 6. Compound A binds non-covalently to ACase and has quick on/off rates with a K_d of 5.122 nM.

References

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