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# Synthesis and evaluation of *galacto*-noeurostegine and its 2-deoxy analogue as glycosidase inhibitors

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# Quantum mechanical calculation setups

#### Noeurostegine, benzyl protected analogue:

Conformational search: Open and closed forms of the molecule were run through a conformational search using the MCMM method implemented in the Schrödinger Suite. The search was performed using 10,000 steps in the implicit CHCl<sub>3</sub> model with no cutoff. Searches were performed using both the OPLS-2005 and MMFF force fields. Unique conformations (defined by a 0.5 Å RMSD cutoff using all heavy atoms) within 10 kJ/mol of the lowest energy structure were kept for QM geometry optimisations.

Geometry optimisation: The conformations obtained from the conformational search were DFT optimised using the pcseg-1 basis set and the wb97xd functional. The IEFPCM method was used to model the presence of CHCl<sub>3</sub>. Conformations within 10 kJ/mol of the lowest energy structure were kept for MP2 single point calculations.

MP2 single point calculations: MP2 calculations were performed using the aug'-cc-pVDZ basis set and the wb97xd functional. The IEFPCM method was used to model the presence of CHCl<sub>3</sub>.

#### Noeurostegine:

Conformational search: Charged and neutral configurations of the open and closed forms of the molecule were run through a conformational search using the MCMM method implemented in the Schrödinger Suite. The search was performed using 10,000 steps in the implicit water model with no cutoff. Searches were performed using both the OPLS-2005 and MMFF force fields. Unique conformations (defined by a 0.2 Å RMSD cutoff using all heavy atoms) within 35 kJ/mol of the lowest energy structure were kept for QM geometry optimisations.

Geometry optimisation: The conformations obtained from the conformational search were DFT optimised using the pcseg-1 basis set and the wb97xd functional. The IEFPCM method was used to model the presence of water. All conformations were kept for MP2 single point calculations.

MP2 single point calculations: MP2 calculations were performed using the aug'-cc-pVTZ basis set and the wb97xd functional. The IEFPCM method was used to model the presence of water.

# Galacto-noeurostegine, benzyl protected analogue (28):

Conformational search: Open and closed forms of the molecule were run through a conformational search using the MCMM method implemented in the Schrödinger Suite. The search was performed using 10,000 steps in the implicit CHCl<sub>3</sub> model with no cutoff. Searches were performed using both the OPLS-2005 and MMFF force fields. Unique conformations (defined by a 0.5 Å RMSD cutoff using all heavy atoms) within 35 kJ/mol of the lowest energy structure were kept for QM geometry optimisations.

Geometry optimisation: The conformations obtained from the conformational search were DFT optimised using the pcseg-1 basis set and the wb97xd functional. The IEFPCM method was used to model the presence of CHCl<sub>3</sub>. Conformations within 20 kJ/mol of the lowest energy structure were kept for MP2 single point calculations.

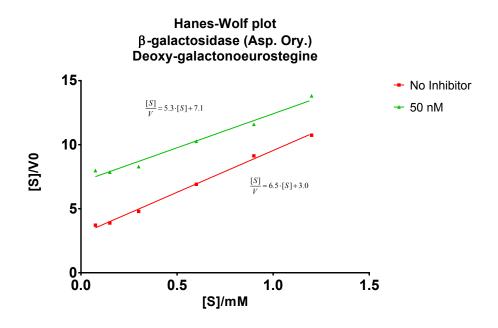
MP2 single point calculations: MP2 calculations were performed using the aug'-cc-pVDZ basis set and the wb97xd functional. The IEFPCM method was used to model the presence of CHCl<sub>3</sub>.

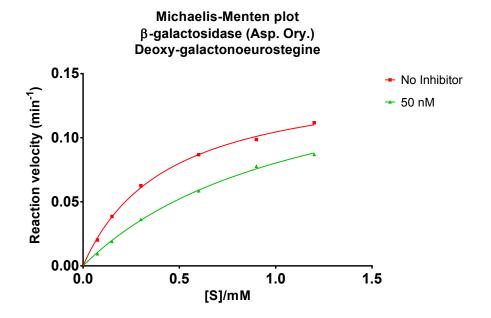
#### Galacto-noeurostegine (5):

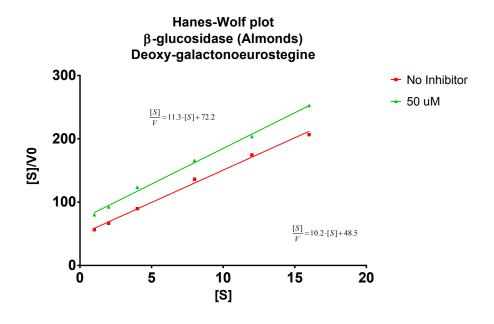
Conformational search: Charged and neutral configurations of the open and closed forms of the molecule were run through a conformational search using the MCMM method implemented in the Schrödinger Suite. The search was performed using 10,000 steps in the implicit water model with no cutoff. Searches were performed using both the OPLS-2005 and MMFF force fields. Unique conformations (defined by a 0.2 Å RMSD cutoff using all heavy atoms) within 50 kJ/mol of the lowest energy structure were kept for QM geometry optimisations.

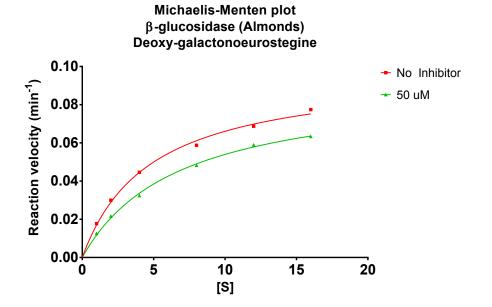
Geometry optimisation: The conformations obtained from the conformational search were DFT optimised using the pcseg-1 basis set and the wb97xd functional. The IEFPCM Method was used to model the presence of water. Conformations within 20 kJ/mol of the lowest energy structure were kept for MP2 single point calculations.

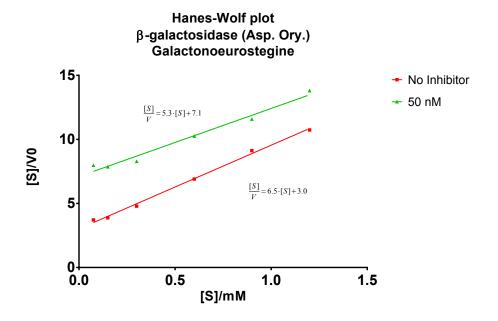
MP2 single point calculations: MP2 calculations were performed using the aug'-cc-pVTZ basis set and the wb97xd functional. The IEFPCM method was used to model the presence of water.

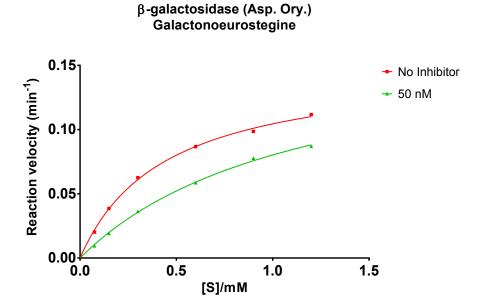






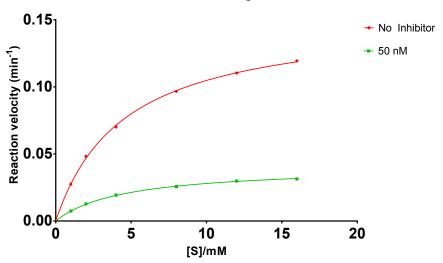


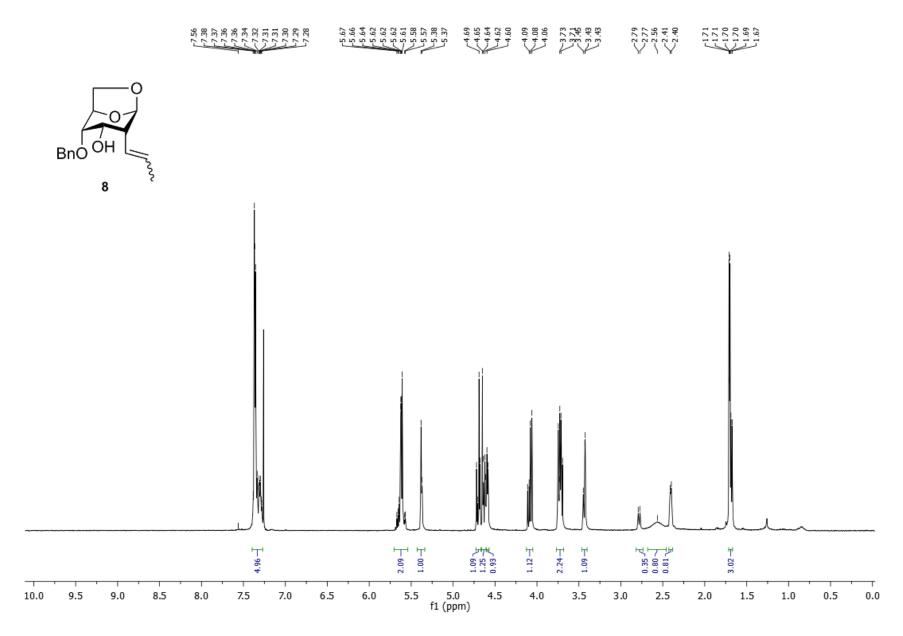


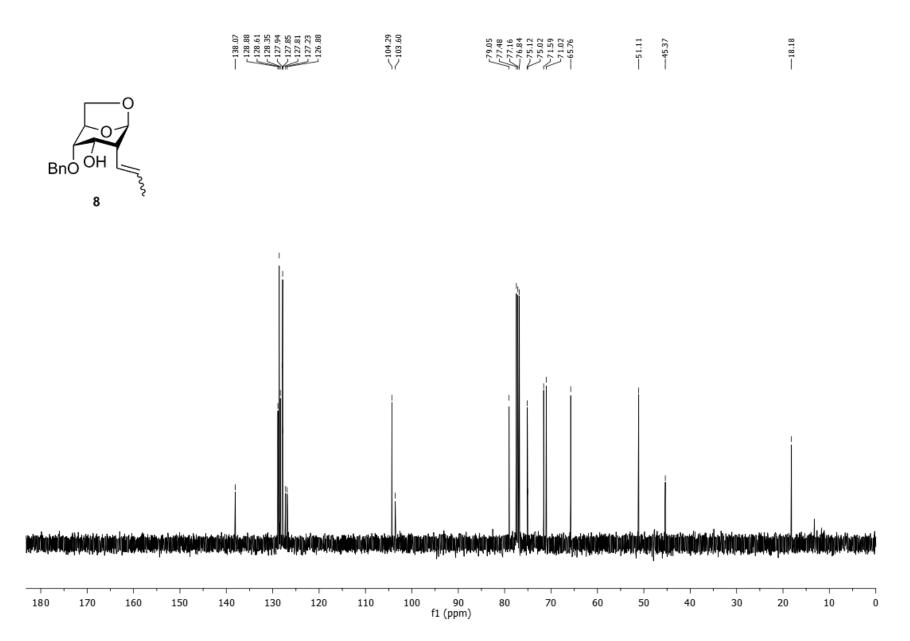


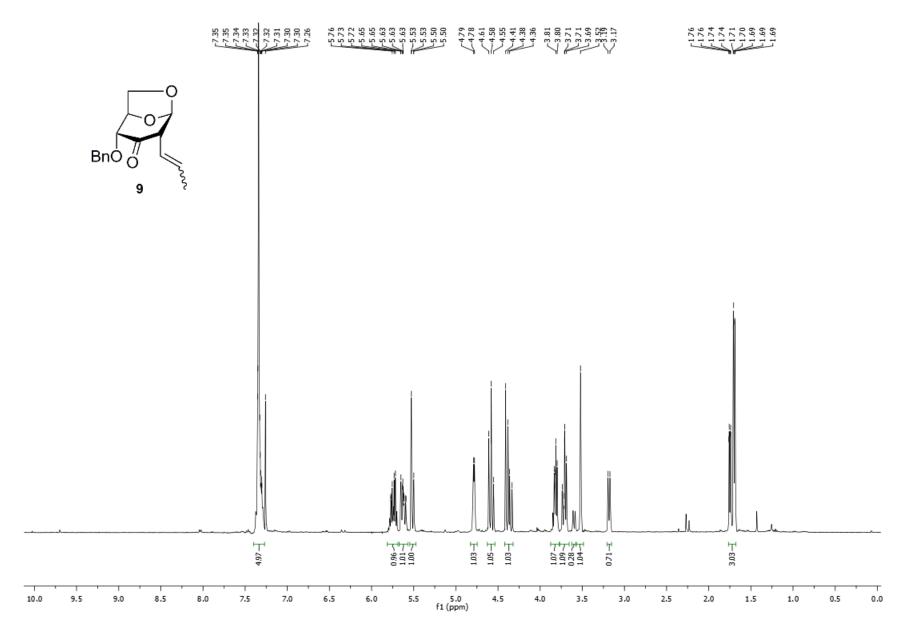
Michaelis-Menten plot

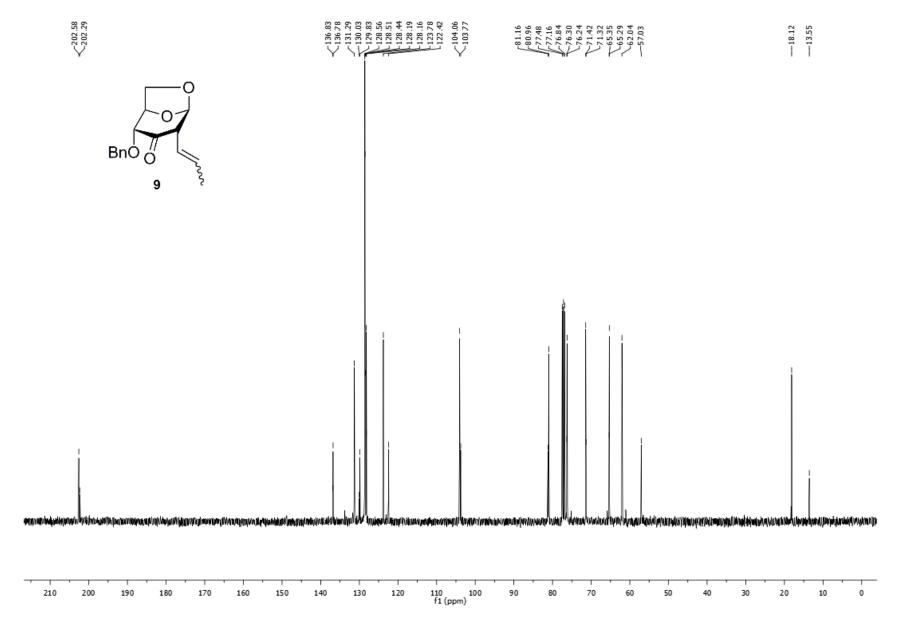
# Non-competitive - 30 min preincubation Michaelis-Menten plot β-glucosidase (Almonds) Galactonoeurostegine











# 0.91 0.91 0.30 10.0 9.5 9.0 8.5 8.0 7.5 7.0 6.5 5.5 5.0 f1 (ppm) 4.0 3.5 3.0 2.5 2.0 1.5 1.0 0.5

