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Supporting Information

Carbohydrate Receptors Combining both a Macrocyclic Building Block and Flexible Side Arms as Recognition Units: Binding Properties of Compounds with CH₂OH Groups as Side Arms

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1 <u>Description of the binding studies</u>

1.1 ¹H NMR titrations

¹H NMR titrations were carried out in CDCl₃ at 25°C (chloroform was stored over activated molecular sieves and deacidified). Dilution experiments show that the receptors do not self-aggregate in the used concentration range.

Stock solutions in CDCl₃ were prepared for the receptor and sugar. These solutions and the corresponding solvent were combined in a manner so that the concentration of the receptor was kept constant and that of the sugar varied. For each titration 15-20 samples were prepared and the ¹H NMR spectra were recorded. For each receptor-sugar system at least two ¹H NMR titrations were carried out.

<u>1.2</u> Microcalorimetric titrations.

Isothermal titration calorimetry (ITC) measurements were performed in CDCl₃ and watercontaining CDCl₃ at 298 K using a power compensation calorimeter VP-ITC (Malvern). A solution of octyl- β -D-glucopyranoside was injected stepwise (8 μ L × 30 times) into a dilute solution of receptor (appr. 0.3 mM) in CDCl₃ or water-containing CDCl₃ (0.035 mol/L and 0.07 mol/L H₂O in CDCl₃). The measured heat rate was recorded as a function of time and converted into enthalpies by integration of the appropriate reaction peaks using NanoAnalyze software provided by TA Instruments. Dilution effects were corrected by subtracting the results of a blank experiment. For each receptor-sugar system at least three titrations were carried out.

It should be noted that initial microcalorimetric studies were performed using microcalorimeter TAM (Thermal Activity Monitor, Thermometrics); however, the required long measurement time (up to 20 h compared with 2 h in the case of VP-ITC) was less favorable for the investigation of the binding processes in chloroform.

1.3 Binding studies in two-phase systems: Liquid-liquid extraction (CDCl₃/H₂O)

To study the extraction of monosaccharides from water into organic phase, aqueous sugar solutions (1 M) were equilibrated with the corresponding receptor in chloroform (1 mM solution) at 25 °C. Solutions of receptor in chloroform were then shaken vigorously with aqueous carbohydrate, the phases were separated, and the organic phase was passed through hydrophobic filter paper to remove residual aqueous solution (similar to the procedure used by Davis et al., see references below). The chloroform was evaporated and the residue analysed by ¹H NMR, first in CDCl₃ and afterwards in DMSO-d₆ (after the evaporation of CDCl₃).

T. Velasco, G. Lecollinet, T. Ryan, A. P. Davis, Org. Biomol. Chem. 2004, 2, 645-647

T. J. Ryan, G. Lecollinet, T. Velasco, A. P. Davis, Proc. Natl. Acad. Sci. U.S.A. 2002, 99, 4863-4866

- 2 Microcalorimetric titrations in water-containing CDCl₃.
- **2.1** Titration of **13** with **24** in water-containing $CDCl_3$ (0.035 mol/L H_2O)



Figure S1. ITC thermogram (left) and titration curve-fitting (right) of **13** with **24** in watercontaining CDCl₃ (0.035 mol/L H₂O). Titration mode: addition of **24** ($c_{syringe} = 3.09$ mM) into **13** ($c_{cell} = 0.30$ mM) at 298 K in 30 steps.





time (minutes)

[sugar]/[receptor]

Figure S2. ITC thermogram (left) and titration curve-fitting (right) of 13 with 24 in watercontaining CDCl₃ (0.07 mol/L H₂O). Titration mode: addition of **24** ($c_{syringe} = 3.02$ mM) into **13** ($c_{cell} = 0.30$ mM) at 298 K in 30 steps.

2.3 Titration of **12** with **24** in water-containing $CDCl_3$ (0.035 mol/L H₂O)



Figure S3. ITC thermogram (left) and titration curve-fitting (right) of **12** with **24** in in watercontaining CDCl₃ (0.035 mol/L H₂O). Titration mode: addition of **24** ($c_{syringe} = 3.01$ mM) into **12** ($c_{cell} = 0.30$ mM) at 298 K in 30 steps.

2.4 Titration of **13** with **24** in water-containing $CDCl_3$ (0.035 mol/L H_2O)



Figure S4. ITC thermogram (left) and titration curve-fitting (right) of of **12** with **24** in in watercontaining CDCl₃ (0.07 mol/L H₂O). Titration mode: addition of **24** ($c_{syringe} = 3.01$ mM) into **12** ($c_{cell} = 0.30$ mM) at 298 K in 30 steps.

3 Molecular modelling calculations (examples)

Program MacroModel V.8.5, OPLS 2001 force field, MCMM, 50000 steps.



Figure S5. Energy-minimized structure of the 1:1 complex formed between 13 and octyl- β -glucoside 24. Shown are three structures with lowest energy: (a) octyl chain of 24 between the bridges, (b) octyl chain next to the bridges.



Figure S6. Examples of energy-minimized structures of compound **12** (Color code: N, blue; O, red; C, gray). The possible intramolecular OH····O, NH···O and NH··· π interactions are shown as broken lines.



Figure S7. (a) Energy-minimized structure of 13 with marked intramolecular OH···O and NH···O hydrogen bonds indicated by the calculations. (b) Energy-minimized structure of the1:1 complex formed between 13 and octyl β -glucoside 24.

4 ROESY experiments for 13-24 and 14-24



b)



Figure S8. Partial ROESY spectrum of receptor **13** (a) and **14** (b) (1 mM) with β -glucoside **24** (1 mM) showing intermolecular connections between carbohydrate and receptor (mixing time = 165 ms)

5. 1 H and 13 C NMR spectra of compounds 12-14, 17 and 18.



5.1. ¹H and ¹³C NMR spectra of compound **12**.

Figure S9. ¹H NMR spectrum of compound 12 in CDCl₃.



Figure S10. ¹³C NMR spectrum of compound 12 in CDCl₃.

5.2. ¹H and ¹³C NMR spectra of compound **13**.



Figure S11a. ¹H NMR spectrum of compound 13 in CDCl₃ (0.3 mM).



Figure S11b. ¹H NMR spectrum of compound 13 in CDCl₃ (\approx 1 mM).



re S12. ¹³C NMR spectrum of compound 13 in CDCl₃.

5.3	¹ H and ¹³ C NMF	spectra of	compound 14.
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Figure S13. ¹H NMR spectrum of compound 14 in CDCl₃.



Figure S14. ¹³C NMR spectrum of compound 14 in CDCl₃.



5.4 ¹H and ¹³C NMR spectra of compound 17.

Figure S15. ¹H NMR spectrum of 17 in CDCl₃.



Figure S16. ¹³C NMR spectrum of 17 in CDCl₃.







Figure S18. ¹³C NMR spectrum of 18 in CDCl₃/CD₃OD 50:1 (ν/ν).