

## Supporting Information

# Molecular Recognition of Methyl $\alpha$ -D-Mannopyranoside by Antifreeze (Glyco)Proteins

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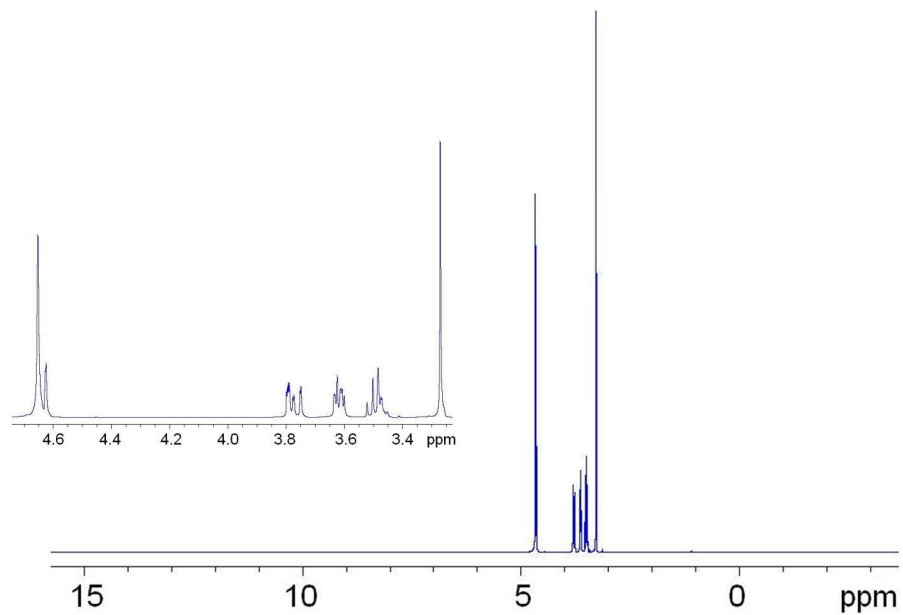


Figure S1.  $^1\text{H}$  NMR spectrum of MDM in  $\text{D}_2\text{O}$  (500 MHz).

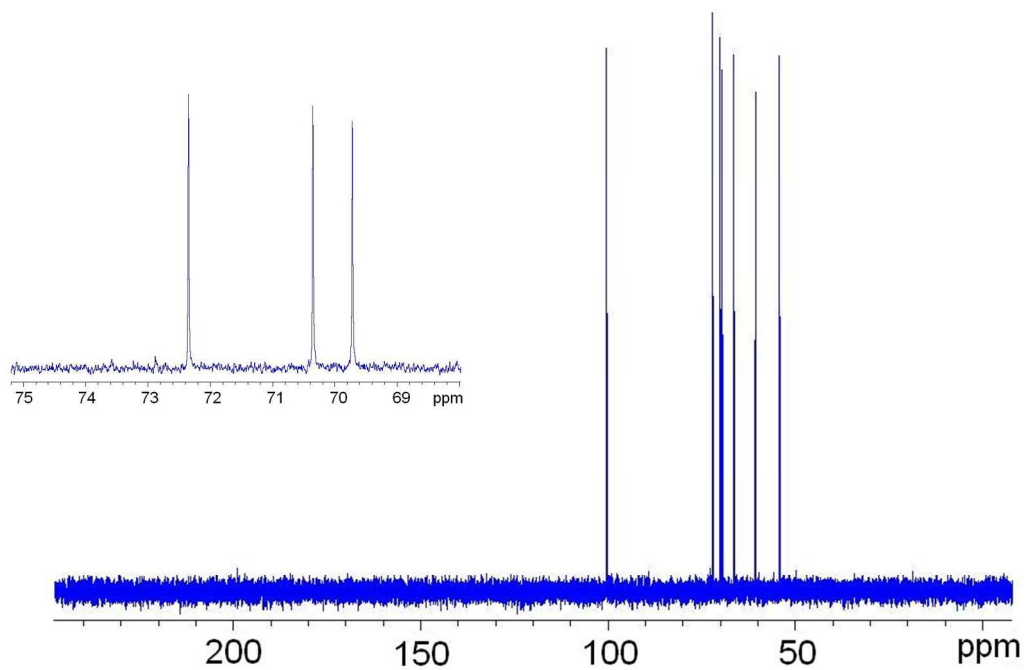


Figure S2.  $^{13}\text{C}$  NMR spectrum of MDM in  $\text{D}_2\text{O}$  (125.8 MHz).

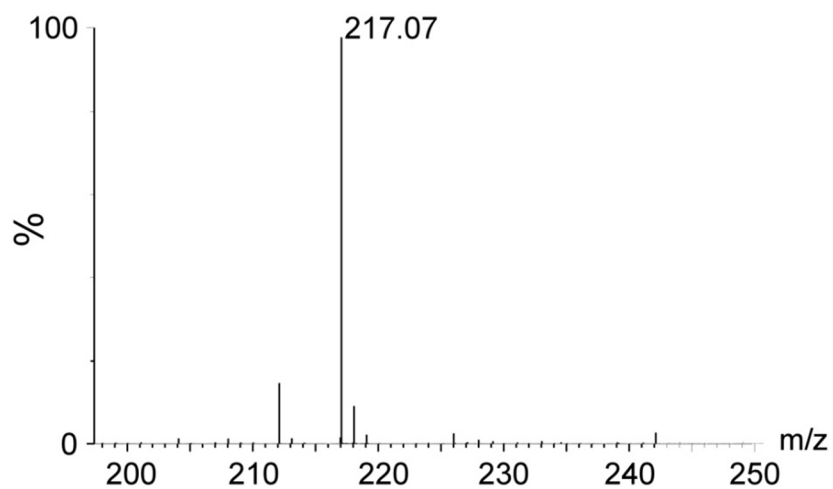


Figure S3. LC-MS spectrum of MDM. Calculated mass for  $C_7H_{14}O_6Na$  is 217.07 and observed mass is 217.07.

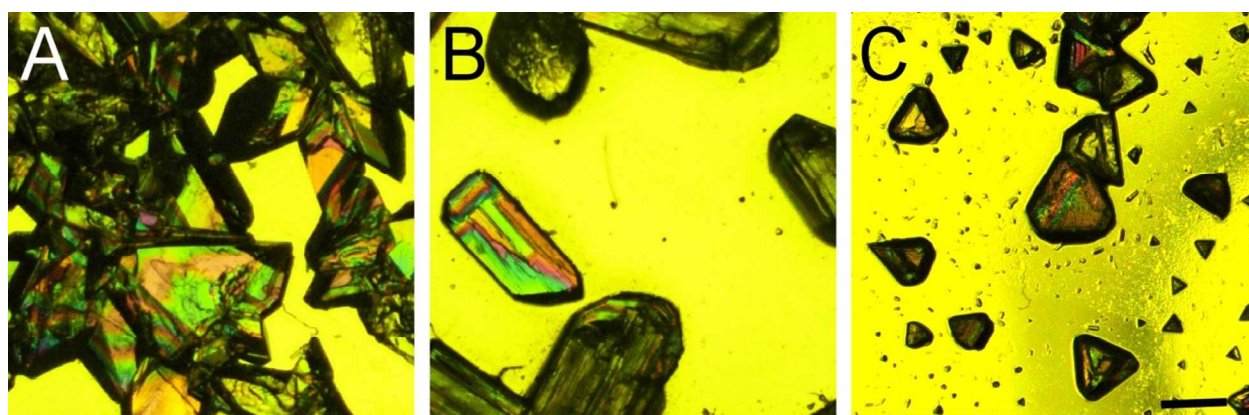


Figure S4. Optical micrographs of the MDM crystals obtained in the presence of the seed MDM crystals and (A) AFGP8, (B) AFGP1-5, and (C) DAFP1. The length of the scale bar, 1 mm, is the same in all the panels.

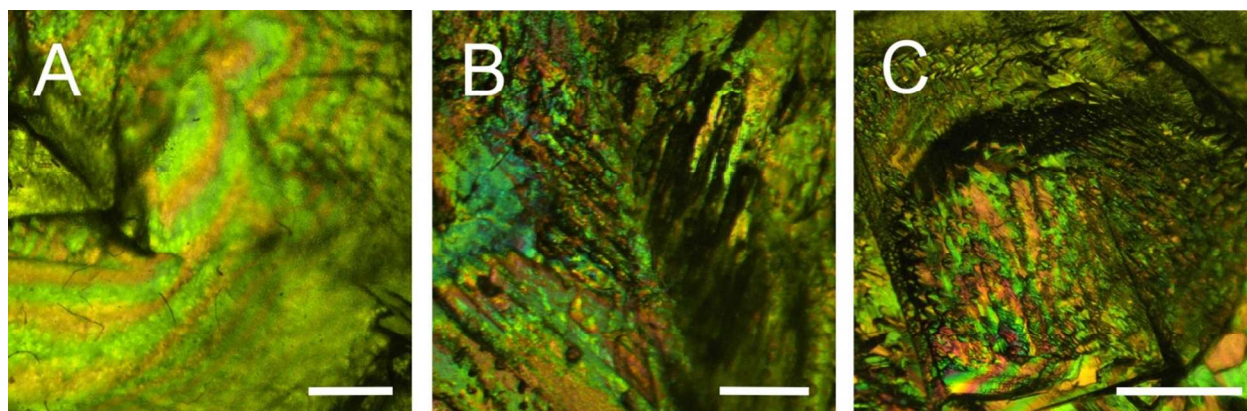


Figure S5. Optical micrographs of the MDM crystals obtained in the presence of the controls: (A) Gal1- $\beta$ -3GalNAc, (B) denatured DAFP1, (C) LCA. The length of the scale bar is 1 mm.

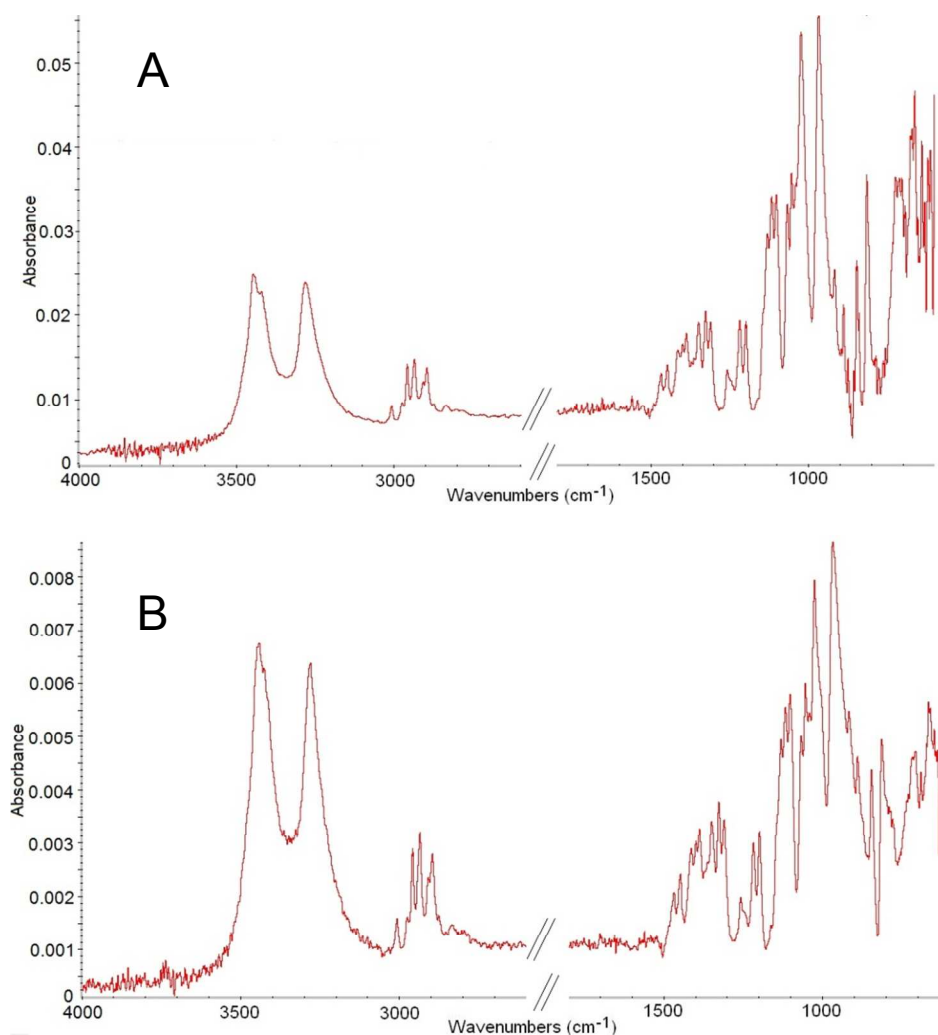


Figure S6. Representative ATR-FTIR spectra of the MDM crystals in the absence and presence of AF(G)Ps. The ATR-FTIR spectra of MDM crystals in the absence of AF(G)Ps (A) and in the presence of DAFP1 (B).

**Table S1. Sample results of methyl  $\alpha$ -D-mannopyranoside (MDM) crystal growth in the presence of additives.**

Sample <sup>a</sup>	Polypeptide concentration ( $\mu$ M)	Induction time (day) <sup>b</sup>	Complete time (day) <sup>b</sup>	Twin defects (%) <sup>c</sup>
MDM	0	12	14	> 75%
MDM/denatured DAFP1	170.0	12	14	> 75%
MDM + LCA	275.4	12	14	> 75%
MDM + Gal1- $\beta$ -3GalNAc	408.0	12	14	> 75%
MDM + DAFP1	14.0	12	17	< 5%
MDM + DAFP1	0.15	12	16	< 5%
MDM + AFGP1-5	1.89	12	15	< 15%
MDM + AFGP1-5	1.00	12	14.5	< 15%
MDM + AFGP8	170.0	12	14	< 40%
MDM + AFGP8	112.2	12	14	< 40%

<sup>a</sup> Each sample contained 510 mM MDM on day 1. Results of each MDM alone and in the presence of each of the three control compounds, the denatured DAFP1, *Lens culinaris* lectin (LCA), and 2-acetamido-2-deoxy-O-( $\beta$ -D-galactopyranosyl)-D-galactose (Gal1- $\beta$ -3GalNAc) are listed for comparison. <sup>b</sup> The day the first solid was observed. Complete means no more weight loss from the vial. Time errors are less than 8 hours. <sup>c</sup> The identity and quality of single crystals were examined using polarized microscope and single crystal x-ray diffraction. The estimated percentages of crystals with twin defects are listed.