

## *Supporting Information*

### **Anomer-Specific Recognition and Dynamics in a Fucose-binding Lectin**

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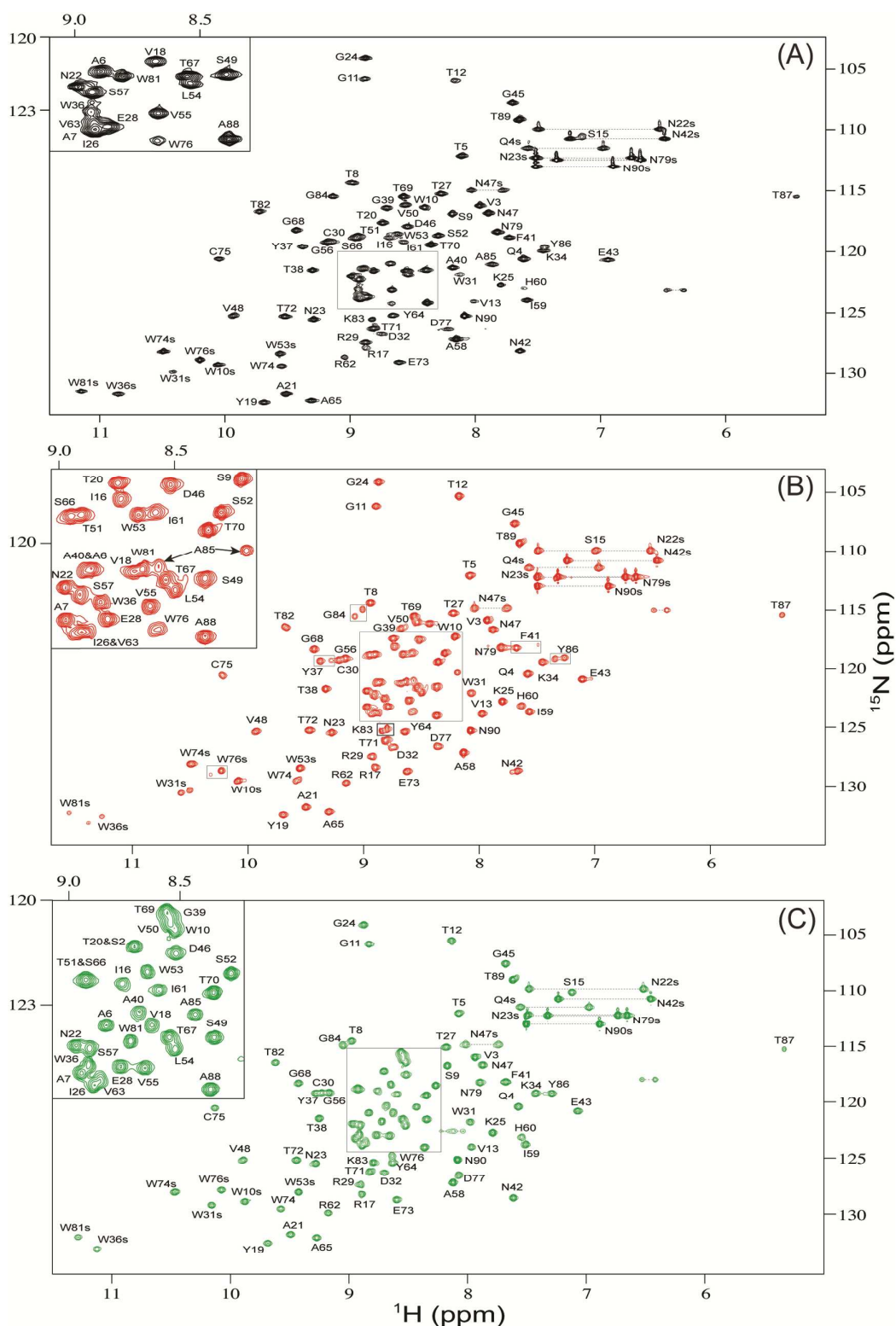
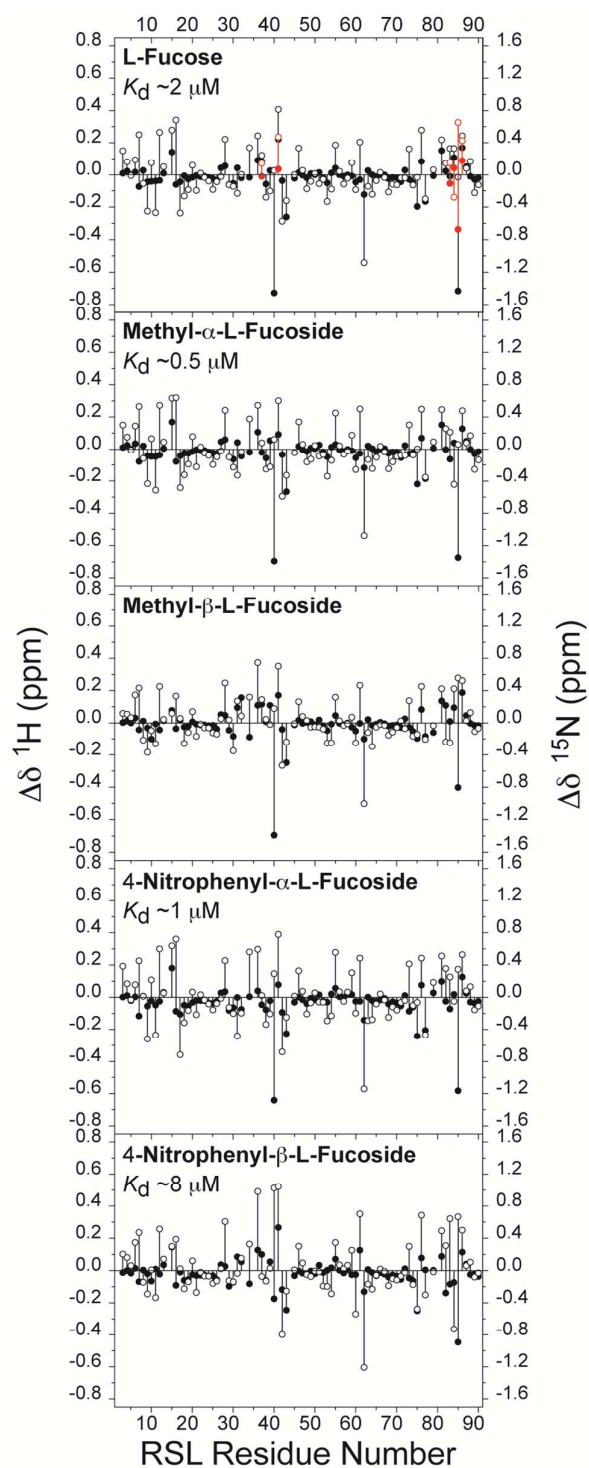
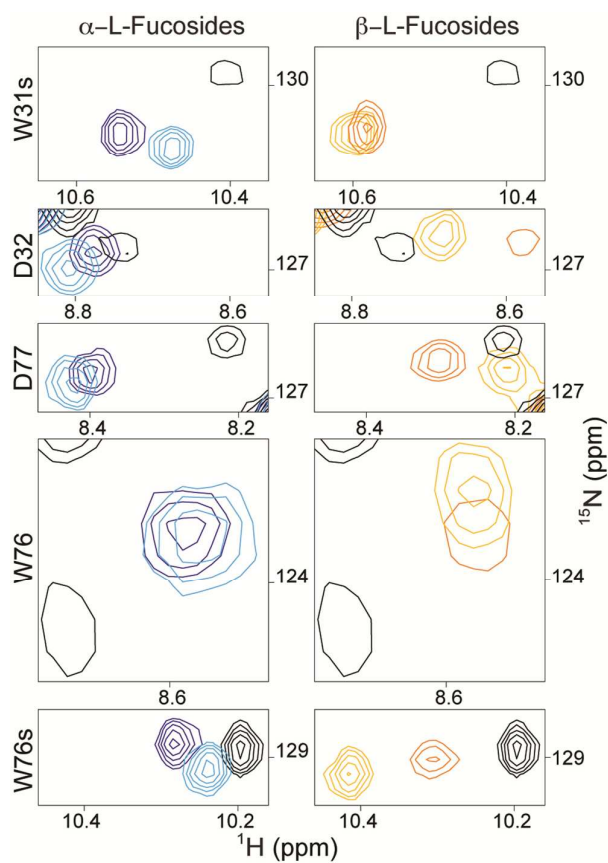


Figure S1. Assigned  $^1\text{H}$ - $^{15}\text{N}$  HSQC spectra of RSL in (A) sugar-free, (B) L-fucose-bound and (C) D-mannose-bound forms. Split resonances due to  $\alpha$  or  $\beta$ -L-fucose are boxed; Y37, F41, W76s, K83 and G84. Sample conditions were 0.25 mM protein in 20 mM potassium phosphate, 50 mM NaCl, pH 6.0 at 303 K.



**Figure S2.** Plot of chemical shift perturbations for RSL backbone resonances in the presence of L-fucose, methyl- $\alpha$ -L-fucoside, methyl- $\beta$ -L-fucoside, nitrophenyl- $\alpha$ -L-fucoside or nitrophenyl- $\beta$ -L-fucoside. In the L-fucose complex some resonances were split in two (red data points) due to the presence of the  $\alpha$ - and  $\beta$ -monomers.



**Figure S3.** Regions from the overlaid  $^1\text{H}$ - $^{15}\text{N}$  HSQC spectra of pure RSL (black contours) and RSL bound to methyl- $\alpha$ -L-fucoside (navy), methyl- $\beta$ -L-fucoside (dark orange), nitrophenyl- $\alpha$ -L-fucoside (light blue) or nitrophenyl- $\beta$ -L-fucoside (light orange). See Figure 5 in the main text.



**Figure S4.** The RSL monomer (PDB 2bt9) coloured by  $R_{2ex}$  values (see scale bar) in the sugar-free (upper), D-mannose-bound (middle) and L-fucose-bound (lower panel) forms. The residues for which  $R_{2ex}$  data could not be obtained are grey. The intra-monomeric binding site residues are shown as sticks, with the backbone nitrogen shown as spheres. Methyl- $\alpha$ -L-fucoside is shown as sticks.