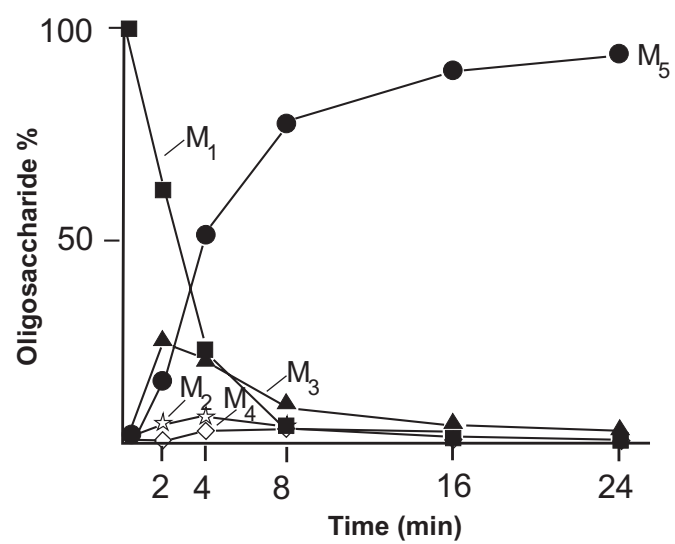
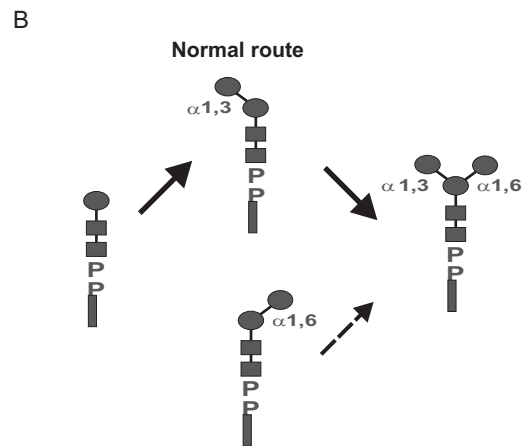
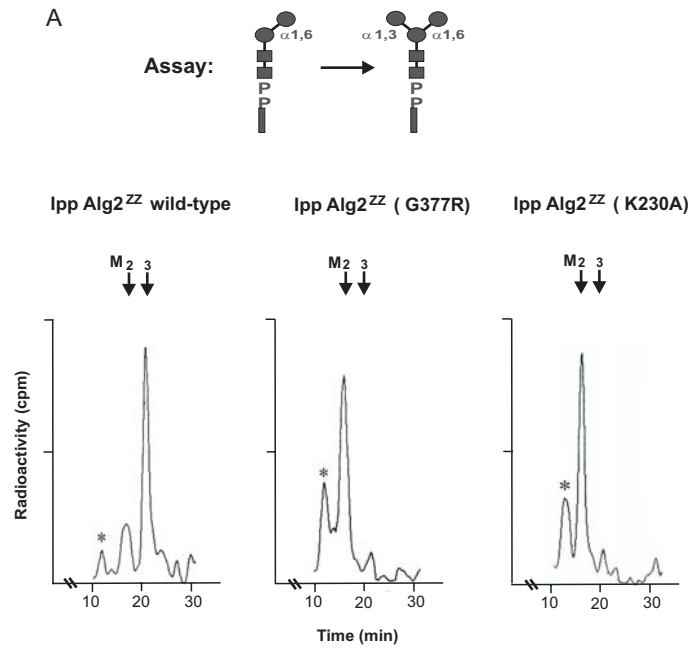


Supplemental Fig. 1



Supplemental Fig. 2



Supplemental Fig. 3

SUPPLEMENTAL FIGURES

SUPPLEMENTAL FIGURE 1. Alg2 has three potential N-glycosylation sites, but is not N-glycosylated. A, membranes were isolated from cells expressing Alg2 C-terminally tagged with Protein A (ZZ). Membranes were incubated without (-) (lanes 1 and 2), or with (+) endoH for 3 h (lane 3) at the temperatures indicated. Alg2 was then analyzed by Western blot. B, as a control, to monitor glycan release, the soluble fraction was also incubated under the same conditions, and carboxypeptidase Y (CPY) was analyzed (lanes 4-6) with anti-CPY antiserum.

SUPPLEMENTAL FIGURE 2. Time dependence of lipid-linked oligosaccharide formation. Solubilized enzyme was incubated for the times indicated using assay 1, as described in Experimental Procedures. Glycolipid fraction was extracted and analyzed by thin layer chromatography on silica gel60-coated plates (Merck), developed in chloroform/methanol/water 65:25:4 (v/v/v). Radioactivity was monitored and quantified by phosphorimaging.

SUPPLEMENTAL FIGURE 3. Elongation of Man(1,6)Man-GlcNAc₂-PP-Dol is defective in *alg2-1* (G377R) and *alg2*(K230A). A, Man(1,6)Man-GlcNAc₂-PP-Dol, obtained by limited digestion of [³H]Man₅GlcNAc₂-PP-Dol with jack bean mannosidase, according to (21), was used as acceptor and incubated with immunoprecipitated Alg2^{ZZ} as enzyme source under the conditions of assay I. Whereas wild-type Alg2 was able to elongate to Man₃GlcNAc₂-PP-Dol (left image), Alg2 with a G377R (middle image) and K230A (right image) mutation was inactive. The *asterisks* indicate Man₁GlcNAc₂-PP-Dol impurity of the acceptor substrate. B, pathway of Man₃GlcNAc₂-PP-Dol formation. The normal route employed is from α1,3- to α1,6-linked mannose. *In vitro*, Alg2 seems to have a somewhat relaxed specificity.