Please see below for the correct Figs. 1 and 2

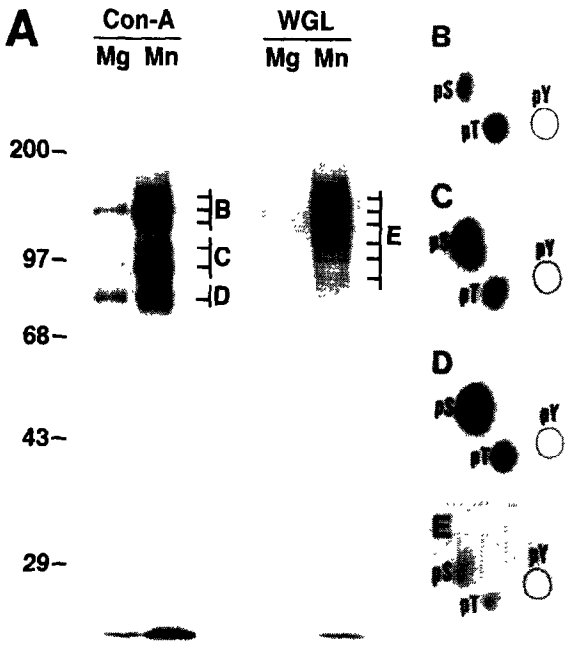

Fig. 1. Kinase activity of glycosylated proteins isolated from a membrane-enriched fraction of *Arabidopsis*. (A) Proteins bound to concanavalin-A agarose (Con-A) or wheat-germ-lectin agarose (WGL) were assayed for kinase activity in the presence of 10 mM $MgCl_2$ or $MnCl_2$, and subjected to SDS-PAGE. An autoradiogram of phosphorylated polypeptides is shown. Migration positions of molecular mass markers are indicated in kilodaltons. Tick marks indicate positions of polypeptides which show greater phosphorylation in the presence of Mn^{2+} than Mg^{2+}. Phosphoamino acid analysis was performed on the regions marked B, C, D, and E in panel (A), and the results are shown in panels (B), (C), (D), and (E), respectively. Positions of nonradioactive phosphoserine (pS), phosphothreonine (pT), and phosphotyrosine (pY) references are indicated.

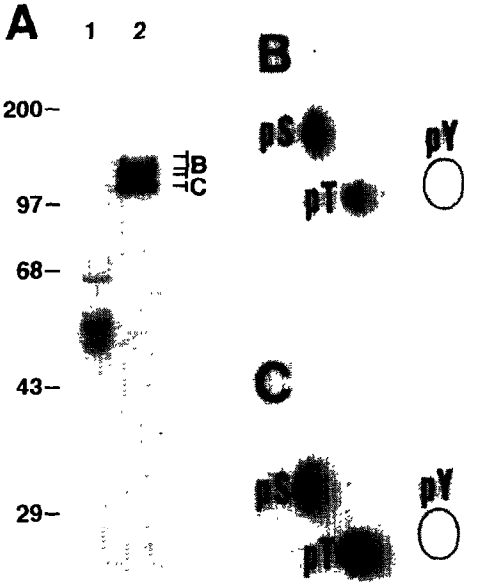

Fig. 2. Blot renaturation of protein kinase. (A) 10 μl Triton X-100 solubilized membranes (lane 1) or concanavalin-A purified proteins from 100 μl solubilized membranes (lane 2) were analyzed for kinase activity following blot renaturation. An autoradiogram is shown. Tick marks indicate positions of autophosphorylating polypeptides enriched for by binding to concanavalin-A. Phosphoamino acid analysis was performed on the regions marked B and C in panel (A) and the results are shown in panels (B) and (C), respectively.

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Further comparison of ubiquinol and cytochrome *c* terminal oxidases

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In the second column, bottom paragraph on p. 297 the sentence:

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should read:

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