



eFP Browser by B. Vinegar, drawn by J. Alls and N. Provart. Data from Gene Expression Map of Arabidopsis Development: Schmid et al., 2005, Nat. Gen. 37:501, and the Nambara lab for the imbibed and dry seed stages. Data are normalized by the GCOS method, TGT value of 100. Most tissues were sampled in triplicate.

Figure S1. Gene Expression Patterns of *PIN3*.

Tissue expression pattern of *PIN3* gene from publicly available microarray datasets visualized by the Arabidopsis eFP Browser (<http://bar.utoronto.ca/efp/cgi-bin/efpWeb.cgi?primaryGene=AT1G70940&modeInput=Absolute>).