

S2 Fig. AtPEPR2 determines root growth inhibition by AtPep1.

Allele distribution of *pepr1* and *pepr2* in a segregating F2 population of a *pepr1 pepr2* cross with *promoter::YFP_N* lines selected for loss of root inhibition on 0.5x MS containing 100 nM AtPep1. Numbers represent individual plants genotyped by PCR for the respective *pepr* T-DNA insertion.

	<i>pepr1</i>	<i>pepr2</i>
WT	14	-
heterozygous	47	6
homozygous	38	93