Table S1. Phosphorylation of LAZY4 proteins in plants or in *in vitro* kinase assays, related to Figures 5 and 6

Peptides	Up-regulated upon gravistimulation	Phosphorylation sites	IP-MS				in vitro kinase assay (MPK3)				Included in		
			R1	R2	R3	R4	KR1	KR2	KR3	KR4	A5/D5	A9/D9	A13/D13
QEITHRPSISSASSHHPR	V	T18		*	√	V	V	V	√	√	V	V	V
		S22		*	V	√	√	√	V	√	√	√	V
		S24		٠	V	√	√	√	√	√	√	V	√
		S25		*	√	√	√	√	√	√	√	√	V
		S27		٠	V	*	√	√	√	V	√	1	√
EEFNDWPHGLLAIGTFGNK/ EEFNDWPHGLLAIGTFGNKK		T47					V	√	√	V			V
KQTPQTLDQEVIQEETVSNLHV EGR		T54					V	√	√	V			V
FLNCPSSLEVDR/ FLNCPSSLEVDRR	٧	S139	V	V	V	√	√	√	√	√		√	V
		S140	√	√	√	√	√	√	√	√		√	√
ISNALCDEKEEDIER/ RISNALCDEKEEDIER		S148	√	V	√	√	√	√		4		1	√
TISVILGR		S164	√	V	V	√		√	√	4		1	√
MFVCTEGFSPVPRPILR		S208					√	√	V	4			V
QQLSLKNEEEEGR	√	S258	*		V	√		√	√	√			V

This table summarizes the mass spectrometry results of the IP-MS experiment shown in Figures 5A and 5B (four replicates: R1 to R4), and *in vitro* kinase assays shown in Figure 5E (four replicates: KR1 to KR4). The phosphorylation sites with explicit MS/MS spectra are labeled "√", and possible phosphorylation sites are labeled "*". In R1 to R4, MS1 area intensities of phosphorylated peptides were normalized by the total area intensity of phosphorylated and non-phosphorylated peptides. If max normalized MS1 intensity of a phosphorylated peptide in samples of gravistimulation was 5-fold greater than the control under vertical condition in at least two experimental replicates, the phosphorylation of the peptide is recorded as "up-regulated upon gravistimulation". For LAZY4-A5/D5, A9/D9, and A13/D13 mutagenesis, selected amino acids were mutated to Ala (A) or Asp (D): A5/D5, the 5 phosphorylated amino acids identified in a phosphorylated peptide in domain I; A9/D9, the 9 total phosphorylated amino acids identified in domain I and III; A13/D13, all the 13 phosphorylated amino acids identified in plants and in *in vitro* kinase assays.