Table S1. Plant genotypes used for this study

Plant genotype	Description/reference
pATHB15::LhG4	Five thousand bp of the 5' upstream sequence of ATHB15 were amplified by PCR and cloned in front of the LhG4 sequence
10Op::ATHB8δmiR	A non-silent single-point mutation (G to D) was introduced in the miR165/166 complementary site of an ATHB8 cDNA and the cDNA was cloned behind the 10Op sequence
pLTP1::LhG4	Moore et al., 2006
pREV::LhG4	Moore et al., 2006
6Op::KAN1	Eshed et al., 2001
10Op::miR165	Alvarez et al., 2005
10Op::ATHB8δmiR	Alvarez et al., 2005
10Op::GFP	Alvarez et al., 2005
APL::GUS	Bonke et al., 2003
ATHB8::GUS	Baima et al., 1995
PIN1::GFP	Benkova et al., 2003
35S::PIN1	Benkova et al., 2003
DR5rev::GFP	Friml et al., 2003
35S::KAN1-GR	Hawker and Bowman, 2004
rev-9	Emery et al., 2003
rev-10d	Emery et al., 2003
phb-6 phv-5 rev-9	Emery et al., 2003
kan1-2 kan2-1 kan3-1 kan4-3	Izhaki and Bowman, 2007

Plants homozygous for the transactivation driver lines pATHB15::LhG4, pLTP1::LhG4 and pREV::LhG4 were crossed to plants homozygous for the various reporter constructs 60p::KAN1, 100p::miR165, 100p::ATHB88miR and 100p::GFP. pATHB15::LhG4 and 100p::miR165 were crossed to rev-9 mutants and segregating plants homozygous for pATHB15::LhG4 and rev-9 were crossed with segregating plants homozygous for 100p::miR165 and rev-9. APL::GUS and ATHB8::GUS plants were crossed with the pATHB15::LhG4 driver line and plants homozygous for pATHB15::LhG4 and either APL::GUS or ATHB8::GUS were crossed with homozygous reporter lines. Similarly, PlN1::GFP, 355::PlN1 and DR5rev::GFP were introduced into the pATHB15::LhG4 and the pOP::KAN1 lines. 355::KAN1-GR plants were crossed with 355::PlN1 plants and F3 plants homozygous for both gene constructs were selected.

References

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