

Table S1 **Amino acid sequences of epitope tags.** The amino acid sequence of each tandem affinity epitope tag are shown with the Biotin Ligase Recognition Peptide (BLRP) in bold type, the L to R mutation in the mutated BLRP tag in red, the 3C protease cleavage site underlined and the Flag, Myc or HA tag in large, non-bold text. \* indicates a stop codon.

Epitope Tag	Amino Acid Sequence
<b>Carboxy-terminal 3xFlag</b>	KLGTDYKDDDDKDYKDDDDKDYKDDDDKGS <u>LEVLFQGP</u> LEGS <b>MAGGLNDIFEAQ</b> <b>RIEWHEDTGG</b> *
<b>Carboxy-terminal 3xFlag-BLRP</b>	KLGTDYKDDDDKDYKDDDDKDYKDDDDKGS <u>LEVLFQGP</u> LEGS <b>MAGGLNDIFEAQ</b> <b>KIEWHEDTGG</b> *
<b>Carboxy-terminal 9xMyc</b>	GRAGTGRSRTSGEQKLISEEDLN <b>GEQKLISEEDLN</b> GEQKLISEEDLN <b>GS</b> SRGEQKLIS EEDLN <b>GEQKLISEEDLN</b> GEQKLISEEDLN <b>GS</b> SRGEQKLISEEDLN <b>GEQKLISEEDL</b> NGEQKLISEEDLN <b>GS</b> SGS <u>LEVLFQGP</u> LEGS <b>MAGGLNDIFEAQ</b> <b>RIEWHEDTGG</b> *
<b>Carboxy-terminal 3xMyc-BLRP</b>	GRAGTGRSRTSGEQKLISEEDLN <b>GEQKLISEEDLN</b> GEQKLISEEDLN <b>GS</b> SGS <u>LEVLF</u> <u>QGP</u> LEGS <b>MAGGLNDIFEAQ</b> <b>KIEWHEDTGG</b> *
<b>Amino-terminal BLRP-3xHA</b>	RYH <b>MAGGLNDIFEAQ</b> <b>KIEWHEDTGG</b> SSIPG <u>LEVLFQGP</u> LEMGYPDVDPDYAGMG YPYDVPDYAGMGYPYDVPDYAGMv

Table S2 DNA sequences of primers and quantitative PCR probes.

Locus	Primer numbers	Primer sequence 5' to 3'
<i>NRPD1</i>	3912	CACCGGTGTCTCACATTCCAAAGTCCCC
	3913	CGGGTTTTTCGGAGAAACCACC
<i>RDR2</i>	7853	CACCAGCAGATAGTGAGTTCTTGG
	7854	AATGGATACAAGTCCACTTG
<i>CLSY1</i>	7857	CACCGTACCTTTTCCATCCTTCTCTG
	7859	GCCACCTGTTGAAGCTTT
<i>SHH1</i>	7927	CACCCAGAGAAAGAGATAAGCAAAGATGG
	7856	CTCTTCAGGGCGGGCGGCA
<i>SHH1 (At1g15215) (A)</i>	7971	CGCTCGATAATGGCTGCT
	7900	TCCCAACAAAGGTAGCATTTG
<i>SHH1 (At1g15215) (B)</i>	7928	CGTGTACGGTTTTCTGGGT CTCTTCAGGGCGGGCGGCA
	7856	
<i>SHH1 (At1g15215) (C)</i>	7971	CGCTCGATAATGGCTGCT
	7856	CTCTTCAGGGCGGGCGGCA
<i>UBQ</i>	3483	GATCTTTGCCGAAAACAATTGGAGG
	3484	CGACTTGTCATTAGAAAGAAAGAGAT
<i>AtSN1</i>	6699	ACTTAATTAGCACTCAAATTAACAAAATAAGT
	6700	TTTAAACATAAGAAGAAGTTCCTTTTTTCATCTAC
<i>FWA</i>	2004	GGTTTTATATTAATATTAAGAGTTATGGGTGGAAGTTT
	2005	CAAATACTTTACACATAAACRAAAAACAAACAAATCRAA
<i>MEA-ISR</i>	3733	CGCGAACGACTATTGCTAAA
	3734	TGAAATCTAACCGGATTTTGG
	M16	TGACCACGGTTAAATGGCGGG-HEX
<i>IGN5</i>	6606	TCCCGAGAAGAGTAGAACAAATGCTAAAA
	6607	CTGAGGTATTCCATAGCCCCTGATCC
	M36	TTGGGCCGAATAACAGCAAGTCC-HEX
<i>ACTIN</i>	2452	TCGTGGTGGTGAGTTTGTTAC
	2453	CAGCATCATCACAAGCATCC
	M17	TTTTCCCTAGTTGAGATGGGAATT-HEX