

Supplemental Table 3. Oligonucleotide Sequences:

ACT2 (AT3G18780)	5'-TCCCTCAGCACATTCCTGCAGAT-3', 5'-AACGATTCCTGGACCTGCCTCATC-3'
LEC1 (At1g21970)	5'-ATCCATCTCTGAATTGAACTTGGA-3' (exon1/exon2), 5'-GGCATGTATTGGTCTTGCTCA-3'
Lea (At1g52690)	5'-GAGGCAAGGCTCAGGAGAA-3' (exon1/exon2), 5'-GCAGCTTGAGACGTCTTGTCTT-3'
Lea (At5g06760)	5'-GAGGAAAAGGCGGAGAAGAT-3' (exon1/exon2), 5'-TTGTCCAGTGGTCGAGTGAGT-3'
Hsp17.4 (At3g46230)	5'-ACGTGGCAGCGTTCACAA-3', 5'-CTTTACTTCCTCCACCTTTGCAT-3'
AtHSP90.1 (At5g52640)	5'-GACGATCAGCTTGAATACTTGGA-3' (exon3,exon4), 5'-ACTTCTTCAACCTCACCTTCGTT-3'
AtHSP70 (At2g32120)	5'-GGTGACAGGAAATGGCAGAA-3' (exon1/exon2), 5'-GTGAACTTGAGAACCGTTCCAA-3'
HSP26.5 (At1g52560)	5'-TCCCACCCTAAATGAGTTCTT-3' (exon1/exon2), 5'-GAACCTCGTATCTGAGCTTGTA-3'
HSP23.5-M (At5g51440)	5'-CTTCTCACATATACTCGATCCGTTT-3' (exon1/exon2), 5'-TAGATGCAACGCGTCGTCTT-3'
ARF4 (At5g60450)	5'-TGTTGGTGCGATGGGATGA-3' (exon8/exon9), 5'-GGAGCAGCAGCTGGACTGAA-3'
ARF3 (At2g33860)	5'-TGGTAAGGTGGGACGACATT-3' (exon8/exon9), 5'-CGGAAGAGGGTGATGATGATA-3'
TAS3 (At3g17185)	5'-TCTCCAAGGCATTAAGGAA-3' (exon1/exon2), 5'-CAAGGTCAAGAAAAGAGAATAAT-3'
PHB (At2g34710)	5'-TCCTAAGCCAAGTTCTCTAAG-3', 5'-TGAGCAAGAATCTTCGATG-3'
PHB –	5'-ACATCAACAGCAAAACCCAAAT-3' 5'-GACCAGGCTTCATCCCAATCA-3'
REV (At5g60690)	5'- GAATAGTCCTGCTGGATTGCTCT-3'

5'- GGATCTCTGCAATCTTCATAGGTT-3'

AthB15 (At1g52150) 5'-CCAAATCCTTCACTCCCAGCTA-3'
5'-CCAGGCTTCATTCCAGGCAT-3'

PHV (At1g30490) 5'-GCAAGGCTACAGGAACTGCTGT-3'
5' - TTGTAGGAGCATAAATCTGAGTGTT-3'

AthB8 (At4g32880) 5'-GTCCTGCTGGATTATTGTCCATT-3'
5'-CAAGCACGAGCAGCGATT-3'

SFC (At5g13300) 5'-GACTCTCCCATGTTCCGCAA-3',
5'-CTCGCCTAGTCCTTCAGTGTATTT-3' (exon2/exon3)

BPS1 (At1g01550) 5'-CTTCTCTCGTCTTTCCTGG-3'
5'-CCGCGTTCTTCGAAGATA-3'

BME3 (At3g54810) 5'-CCCCACTCCTTACTCCTTTTT-3',
5'-GTCTCTCTAGCTCTGTTGTTCTTA-3' (exon1/exon2)

MYB4 (At1g22640) 5'-TCCCGGCAATAAGCGACCATTTG-3'
5'-TCACCGAGGAAGAAGACGAACTCA-3'

REV – 5'-ATCTGTGGTCACAACCTCC-3'
5'-GTTTGTGAGAGGTCGCTA-3'

AthB8 – 5'-GCTACCACAGATACTAGC-3'
5'-TCGCAAGGTCTAATGAGG-3'

AthB15 – 5'-TCAAAGGCAACTGGAACC-3'
5'-CACCCAAAGTACTTGAC-3'

The three primers listed above were taken from Li et al. (2005)

PHV – 5'-AGACCTTGGCGGAGTTCCTTTG-3'
5'-GGTATCGTAGCTGTTTCACGCAAC-3'

DCL1 (At1g01040) 5'-GATCCATTCCCTAAGCGAAGTTTCAGAG-3'
5'-CTATGGATCTTTATGTTGCTCGGGC-3'

The two primers listed above were taken from from Vaucheret et al. (2004).

ARF8 (At5g37020) – 5'-AGATGTTTGCTATCGAAGGGTGTGTTG-3'
5'-CTTCTCCTTGGTGATGACCCATGG-3'

ARF17 (At1g77850) – 5'-AGCACCTGATCCAAGTCCTTCTATG-3'
5'-GAAATCCTCCCCAGCTATTCACCA-3'

AGO1 (At1g48410) – 5'-AAGGAGGTCGAGGAGGGTATGG-3'
5'-CCTACTGTTCTGGCTCAGCAATTTG-3'

The three primers listed above were taken from Vazquez et al. (2004).

To genotype *tdt-1*, we amplified genomic DNA using primers that spanned the 50 bp deletion. To genotype *tdt-2*, we used the SALK T-DNA left border-specific oligonucleotide LBB1 and TDT-specific primers listed below:

tdt-1 (1) 5': GTGCTATTCTCCTGGTGAATG
tdt-1 (2) 5': CACCTCATTACTTGCTGGCT

tdt-2 (1) 5': TCAATCAGAGGAAGCCAAAG
tdt-2 (2) 5': CAGTACTGGACTCGACACTT
LBB1 5': GCGTGGACCGCTTGCTGCAACT

To genotype *vcs-7*, we used the SALK T-DNA left-border-specific oligonucleotide LBB1, and VCS-specific primers listed below:

5': GCGACTAACTTGCATCCTCAG
5': CGCAGTACATGTTTACCAATTG
LBB1 5': GCGTGGACCGCTTGCTGCAACT

RACE

5' RACE Adapter
(5'-CUGAUGGCGAUGAAUGAACACUGCGUUUGCUGGCUUUGAUGAA-3')

5' RACE Outer Primer 5'-GCTGATGGCGATGAATGAACACTG-3'

At5g51440 (HSP23.5-M) gene specific primer
5'-GCATCTGGCTCAAGCTTCTCGTTGGAGTAA-3'

At3g46230 (HSP17.4) gene specific primer
5'-CTCACTGCTCCTCTCACCGCTTATCTGAA-3'