



b)

Representative isoform	Corresponding dot in figure	Name	Sequence
Isotig01227		Tubulin Fwd	TGCAGAGGGCTGTTTGCATGA
Isotig01227		Tubulin Rev	ACCCACGTACCAGTGAACAAAAG
Isotig02972	a	Art v 2 Fwd	GGCTGATCAAAGGAAGGACTGCAA
Isotig02972	a	Art v 2 Rev	TGCACCGCGTATGAACCATTGA
Isotig03100	b	Amb a 10 Fwd	GACGAGCTATTCGCCATCCTGAAA
Isotig03100	b	Amb a 10 Rev	GCTAATGAAACCATCGCCATCTGC
Contig32361	c	Amba 8.2 Fwd	GAATTCGATGCAGCTGGTACCCTT
Contig32361	c	Amb a 8.2 Rev	GGGCTCTTCGTAATGCCAAACAC
Isotig03734	d	Clathrin Fwd	AGGCGTTCATTCGCGCTTACTA
Isotig03734	d	Clathrin Rev	ATCCTGGAGATCCTGCAACCAAAC
Isotig02293	e	02293 Fwd	AGCTCACTTGGATGCTCACCTTCT
Isotig02293	e	02293 Rev	CCATGGGCAAGGTGGAAGTTGTA

Figure S6. Comparison of high and low expressed transcripts, qRT-PCR vs. RPKM. **a)** Fold-change of tubulin normalized expression data are plotted against high and low RPKM-values of transcripts from control plants (black) (n=3). RPKM-values normalized to tubulin are shown in grey. Pearson-correlation $r = 0.984$. **b)** Corresponding isoforms with primer sequences are indicated.