

Expression analysis_qRT-PCR			ChIP analysis_qPCR
<i>P. persica</i> ID		Primer sequence 5'>3'	Primer sequence 5'>3' and relative position to predict TSS
PRUPE_4G204900	ppeUBQ	Fw: AAGGCTAAGATCCAAGACAAAGAG Rev: CCACGAAGACGAAGCACTAAG	
PRUPE_6G159200	ppeFLYind	Fw: GTGCGAATCAGTGACGACCCG Rev: GTCTTTGGAGGATGCGGATTTTCTC	FwA: CGGTTACTTTGGAATGTATTGTGAG (promoter) RevA: AACGAAACGGAGTGTAAGTGGAG (5'UTR)
			FwB: AAAGAAGCAGCAGCGCCAGG (5'UTR) RevB: ATTGGGTGTCAGGTGGGGCC (5'UTR)
			FwC: CGAATATTCTTCTCTTTGGCC (CDS) RevC: TCACCACCACCACCATCACCT (CDS)
			FwD: AGGTGATGGTGGTGGTGGTGA (CDS) RevD: GTCTTTGGAGGATGCGGATTTTCTC (CDS)
PRUPE_4G262200	ppePG22		Fw1: TTGTCCAGCGGAGCAACGG (CDS) Rev1: ATTTGATTACGTTCTGTGGAGTTTG (CDS) Fw2: CTGGTCAGGTAAGATATCATTATGC (CDS) Rev2: GAGCAAGCCCATGGAGGAAGG (CDS)