

Table S1. Primer and their main features. Table reports primer name, sequence, amplicon length, name/gene symbol and identifier (ITAG annotation) used in the experimental procedures.

Primer	Sequence (5'→3')	Amplicon length (bp)	Name/Gene symbol	Accession Number
P11F1 Rw	CGCGCGCCATGGGAACCTTCATATGATATCA			
Del Rv	CGCGCGCTCGAGTTACTAAAGATCTTCTCGTACTAT AATTTTCTCAC	560	ProSys ₍₁₋₁₇₈₎	AAA34184.1
GST Fw	AGATGAGACATTTGAAGGCCCT			
GST Rv	TACCACCGCTCCACCTTAT	106	GST	Solyc09g011500.2.1
EF Fw Rt	CTCCATTGGGTCGTTTTGCT			
EF Rv Rt	GGTCACCTTGGCACCAGTTG	101	elongation Factor 1 alpha	Solyc06g005060.2
bZIP Fw	CGGCGTGTGGAAGATGAGAT			
bZIP Rv	CCTCAAGGGCCTCTGTCACTC	145	bZIP	Solyc01g090270.2.1
PPR2 Fw	ATACCCAGAACGACCCGGTA			
PPR2 Rv	CACCACCAGTCCCATTCTC	103	PPR2	Solyc01g110460.2.1
MLO3 Fw	CGGCACATTTAGCACACAG			
MLO3 Rv	GAGTAAGAAGAGCACGGCGA	136	MLO3	Solyc06g010030.2.1
Crocetin Fw	GGAGTACCTGTCGTGGCTTT			
Crocetin Rv	ACTCCACTCTTCCACACATCT	83	Crocetin	Solyc12g098590.1.1
SRP Fw	CTTAGTGGACACAGCGAGCA			
SRP Rv	GATCTCCATGCCAAAACCGC	123	Stress-related protein	Solyc09g074930.2.1
DSP Fw	TCATCAGCCTCATCACCTCC			
DSP Rv	TGGAGTCAGACGGTGTGAA	148	Dual specificity phosphatase 1	Solyc05g054700.2.1
Lap Fw	ATCTCAGGTTTCCTGGTGGAAGGA			
Lap Rv	AGTTGCTATGGCAGAGGCAGAG	99	leucine amino peptidase	Solyc00g187050.2
Mate Fw	ACCCATCAATGACACCCAAG			
Mate Rv	GGCATGTGGTATGGGATGTT	154	mate efflux protein	BI933305
PG Fw	TCAGCCCTTTTCCTTGTCGA	125	Polygalacturonase	Solyc08g082170.2.1
PG Rv	TGCGACCACCATCATTTCCA			
PcSys Fw	TCTGAATTTGTCTCCCCTTAGAA			
PcSys Rv	AGCCAAAAGAAAGGAAGCAATAC	119	Endogenous Prosystemin	Solyc05g051750.2.1
Sys Fw	GGGAGGGTGCACTAGAAATA			
Sys Rv	TTGTCGAAACCGATGATACG	108	ProSystemin	Solyc05g051750.2.1
Sysdel Fw	GAAGGAGATGATGATGCACAA			
Sysdel Rv	CACCCTCCCCTTCCATCT	208	ProSystemin	Solyc05g051750.2.1