Table S2 DEGs in KEGG enrichment in 'Plant hormone signal transduction', 'Starch and sucrose metabolism', 'MAPK signaling pathway'.

Grafting combination	KEGG description	Gene ID	Gene description
	MAPK	LOC103424105	ethylene receptor 2-like [ Malus domestica (apple)]
F-I-S Vs I-F-R	signaling pathway - plant	LOC103436283	basic form of pathogenesis-related protein 1-like [ Malus domestica (apple) ]
	Starch and	LOC103438567	probable trehalose-phosphate phosphatase D [ Malus domestica (apple) ]
	sucrose metabolism	LOC103453816	endoglucanase 24-like [ Malus domestica (apple) ]
	Plant hormone signal	LOC103424105	ethylene receptor 2-like [ Malus domestica (apple)]
		LOC103445717	auxin-induced protein AUX28 [ Malus domestica (apple) ]
		LOC103436283	basic form of pathogenesis-related protein 1-like [ Malus domestica (apple) ]
	MAPK	LOC103404780	transcription factor MYC2-like [ Malus domestica (apple) ]
F-I-R Vs	signaling pathway - plant	LOC103412151	serine/threonine-protein kinase OXI1-like [ Malus domestica (apple) ]
I-I-R	Plant hormone	LOC103444275	auxin-responsive protein SAUR32-like [ Malus domestica (apple) ]
	signal transduction	LOC103404780	transcription factor MYC2-like [ Malus domestica (apple) ]
	MAPK	LOC103436283	basic form of pathogenesis-related protein 1-like [ Malus domestica (apple) ]
I-F-S Vs I-I-S	signaling	LOC103436302	pathogenesis-related protein 1-like [ Malus domestica (apple) ]
	pathway - plant	LOC103434045	pathogenesis-related protein 1-like [ Malus domestica (apple) ]
	Plant hormone signal transduction	LOC103436283	basic form of pathogenesis-related protein 1-like [ Malus domestica (apple) ]
		LOC103434165	auxin-responsive protein IAA3-like [ Malus domestica (apple) ]
		LOC103436302	pathogenesis-related protein 1-like [ Malus domestica (apple) ]
		LOC103445478	probable xyloglucan endotransglucosylase/hydrolase protein 23 [ <i>Malus domestica</i> (apple) ]
		LOC103432477	gibberellin receptor GID1B-like [ Malus domestica (apple) ]
		LOC103410576	auxin-responsive protein IAA1-like [ Malus domestica (apple)]
		LOC103434045	pathogenesis-related protein 1-like [ Malus domestica (apple) ]
	Starch and sucrose metabolism	LOC103400901	sucrose synthase [ Malus domestica (apple) ]
		LOC103450387	hexokinase-2, chloroplastic [ Malus domestica (apple) ]
		LOC103444250	sucrose synthase 5 [ Malus domestica (apple) ]
		LOC103431945	alpha,alpha-trehalose-phosphate synthase [UDP-forming] 1-like [ Malus domestica (apple) ]
		LOC103436320	beta-glucosidase 12-like [ Malus domestica (apple) ]
		LOC103444513	probable sucrose-phosphate synthase 4 [ Malus domestica (apple) ]
		LOC103440507	glucose-6-phosphate isomerase, cytosolic 2B-like [ Malus domestica (apple) ]