

Gene name	All samples			Fruit developmental series	Different	Different storage	Different regulator	Different
	No subgroups	2 subgroups	5 subgroups		genotypes	time series	treatments	tissues
<i>18S rRNA</i>	0.140	0.098	0.056	0.101	0.051	0.087	0.162	0.140
<i>ACT</i>	0.048	0.057	0.020	0.023	0.023	0.023	0.025	0.060
<i>CYP2</i>	0.045	0.028	0.015	0.028	0.030	0.028	0.058	0.015
<i>GAPDH</i>	0.047	0.064	0.018	0.036	0.021	0.023	0.007	0.068
<i>PLA2</i>	0.055	0.061	0.021	0.054	0.053	0.056	0.035	0.056
<i>RPII</i>	0.037	0.017	0.010	0.005	0.002	0.003	0.024	0.033
<i>TEF2</i>	0.019	0.027	0.007	0.020	0.005	0.020	0.006	0.010
<i>TUA</i>	0.042	0.044	0.015	0.024	0.026	0.009	0.060	0.034
<i>TUB</i>	0.043	0.047	0.012	0.008	0.002	0.014	0.052	0.050
<i>UBQ10</i>	0.027	0.039	0.013	0.026	0.033	0.031	0.007	0.031