

Table 1 Gene expression stability ranks of eight candidate reference genes in different sets of *C. speciosa* samples evaluated by delta Ct (ΔCT) statistical algorithm

Gene	All samples		All treatments		Developmental stages		Tissues		MeJA treatment		CPPU treatment		GA ₃ treatment		PP ₃₃₃ treatment	
	Rank	ΔCT	Rank	ΔCT	Rank	ΔCT	Rank	ΔCT	Rank	ΔCT	Rank	ΔCT	Rank	ΔCT	Rank	ΔCT
<i>GAPDH</i>	1	1.23	2	1.16	1	1.03	1	0.63	2	0.86	2	0.88	2	0.63	5	1.02
<i>60S</i>	4	1.32	3	1.18	7	1.44	7	0.88	3	0.88	7	1.38	4	0.71	6	1.08
<i>ACTIN</i>	7	1.78	7	1.54	3	1.04	3	0.69	1	0.78	8	1.83	1	0.61	3	0.93
<i>TUA2</i>	8	1.92	8	1.65	8	2.15	4	0.8	8	1.78	3	1.02	6	0.82	7	1.09
<i>TUB1</i>	2	1.25	1	1.13	5	1.33	2	0.68	5	1.02	3	1.02	5	0.73	1	0.89
<i>TIF5</i>	3	1.28	4	1.2	4	1.06	6	0.87	6	1.07	6	1.23	3	0.69	2	0.9
<i>UBQ</i>	6	1.56	4	1.2	6	1.35	5	0.86	4	0.94	1	0.83	8	1.21	4	0.98
<i>EF2</i>	5	1.47	6	1.39	1	1.03	8	1.34	7	1.34	5	1.09	7	1.17	8	1.15