

Table 4 Gene expression stability ranks of eight candidate reference genes in different sets of *C. speciosa* samples evaluated by RefFinder (RF) statistical algorithm

Gene	All samples		All treatments		Developmental stages		Tissues		MeJA treatments		CPPU treatments		GA ₃ treatments		PP ₃₃₃ treatments	
	Rank	Geomean	Rank	Geomean	Rank	Geomean	Rank	Geomean	Rank	Geomean	Rank	Geomean	Rank	Geomean	Rank	Geomean
<i>GAPDH</i>	3	2.21	5	3.46	2	1.68	1	1.19	2	2.11	2	2.11	2	3.08	6	4.79
<i>60S</i>	1	2.00	3	2.91	7	6.44	7	6.09	4	3.46	7	5.66	3	3.36	7	5.42
<i>ACTIN</i>	7	7.00	7	7.00	4	2.91	3	2.28	1	1.32	8	8.00	1	1.32	2	2.71
<i>TUA2</i>	8	8.00	8	8.00	8	8.00	4	4.47	8	8.00	5	4.12	7	6.16	5	4.45
<i>TUB1</i>	4	2.45	1	1.86	5	5.23	2	1.86	5	3.50	3	3.31	5	3.64	1	2.11
<i>TIF5</i>	2	2.06	4	3.16	3	2.63	6	5.96	6	5.63	4	3.83	4	3.46	3	2.78
<i>UBQ</i>	6	5.42	2	2.78	6	6.24	5	5.18	3	3.31	1	1.41	6	4.76	4	3.25
<i>EF2</i>	5	5.00	6	3.66	1	1.57	8	8.00	7	6.74	6	4.79	8	6.74	8	5.66