

Table 3 Gene expression stability ranks of eight candidate reference genes in different sets of *C. speciosa* samples evaluated by *BestKeeper* statistical algorithm

Gene	All samples		All treatments		Developmental stage:		Tissues		MeJA treatments		CPPU treatments		GA ₃ treatments		PP ₃₃₃ treatments	
	Rank	SD	Rank	SD	Rank	SD	Rank	SD	Rank	SD	Rank	SD	Rank	SD	Rank	SD
<i>GAPDH</i>	6	1.13	6	1.46	1	0.11	2	0.18	5	0.84	5	1.04	5	0.57	7	0.89
<i>60S</i>	1	0.82	1	0.84	5	0.75	7	0.64	4	0.76	3	0.78	1	0.44	4	0.54
<i>ACTIN</i>	7	1.73	7	1.87	2	0.24	3	0.36	3	0.63	8	1.92	3	0.48	5	0.77
<i>TUA2</i>	8	1.84	8	1.88	8	1.41	4	0.42	8	1.14	6	1.24	8	0.76	8	1.22
<i>TUB1</i>	3	0.95	3	0.88	6	0.82	1	0.09	1	0.52	2	0.50	7	0.70	5	0.77
<i>TIF5</i>	2	0.86	5	0.97	4	0.47	5	0.50	7	1.06	1	0.31	4	0.50	3	0.43
<i>UBQ</i>	4	0.96	4	0.89	7	0.91	6	0.57	2	0.61	4	0.83	1	0.44	1	0.09
<i>EF2</i>	5	1.02	1	0.84	3	0.45	8	1.03	6	0.89	7	1.30	6	0.69	2	0.42