

Selection and Validation of Novel RT-qPCR Reference Genes under Hormonal Stimuli and in Different Tissues of *Santalum album*

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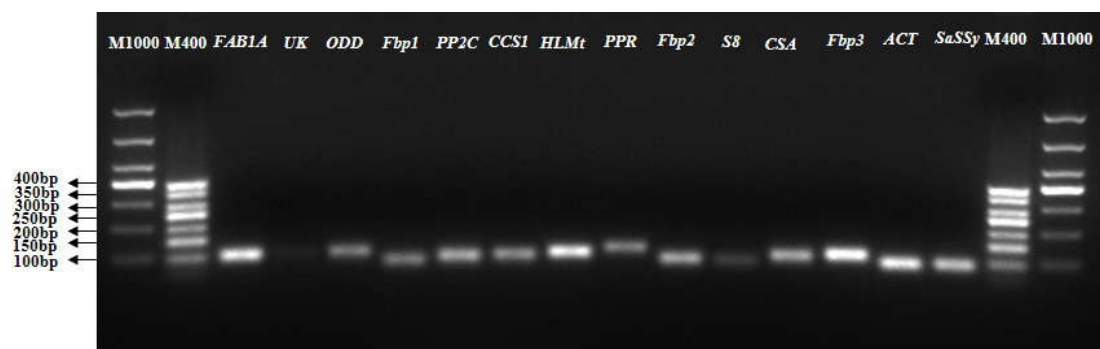
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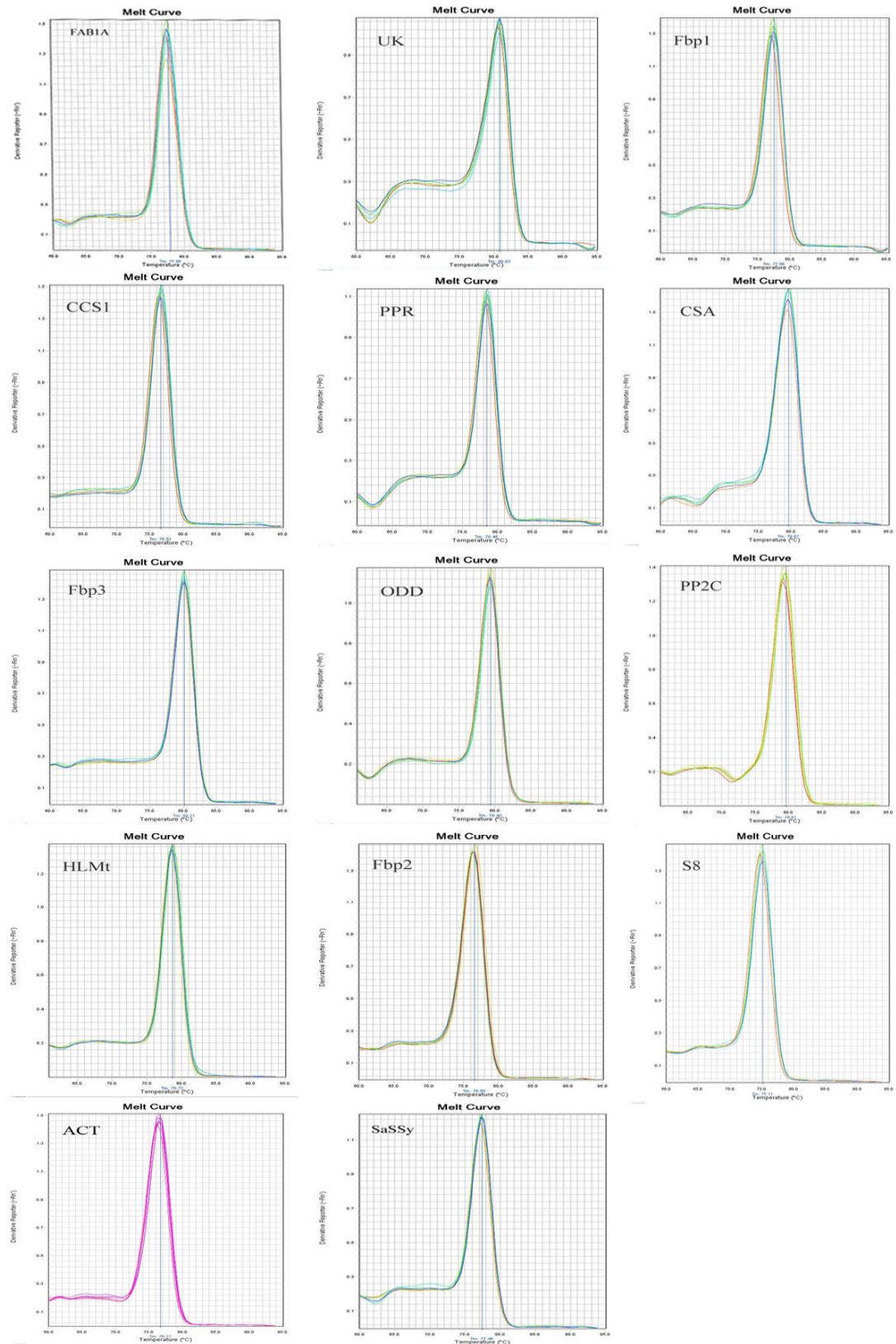
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Supplementary Table S1. Novel candidate reference genes selected based on RNA-seq data.

Gene name	FPKM root1	FPKM root2	FPKM root3	FPKM heartwood1	FPKM heartwood2	FPKM heartwood3	FPKM heartwood4	FPKM sapwood1	FPKM sapwood2	FPKM sapwood3	FPKM sapwood4	FPKM transition zone1	FPKM transition zone2	FPKM transition zone3	FPKM transition zone4	FPKM leaves	MV	SD	CVs (%)
<i>FAB1A</i>	22.11	23.86	26.51	31.26	26.25	27.18	26.48	29.63	25.69	22.02	25.42	28.84	27.10	24.15	27.06	23.87	26.09	2.54	9.75
<i>UK</i>	2.35	3.31	2.97	2.50	2.67	2.90	2.63	2.82	2.64	2.66	2.32	2.48	2.38	2.89	2.60	2.52	2.67	0.26	9.84
<i>ODD</i>	7.04	9.33	7.93	7.83	6.10	7.82	7.91	7.07	7.60	8.01	8.12	6.47	7.52	7.18	6.67	8.42	7.56	0.80	10.57
<i>Fbp1</i>	4.89	4.53	4.64	3.96	3.97	3.43	3.91	3.76	4.09	3.55	3.73	3.59	3.51	3.73	3.62	3.67	3.91	0.43	11.01
<i>PP2C</i>	8.59	8.41	8.62	8.78	8.16	8.96	8.49	8.36	7.28	6.59	8.53	7.81	8.13	7.77	8.63	11.04	8.38	0.93	11.1
<i>CCS1</i>	5.26	6.15	5.75	4.48	4.66	4.74	5.95	5.42	4.70	4.89	5.76	4.95	4.89	5.21	6.51	5.09	5.28	0.59	11.25
<i>HLMt</i>	9.27	8.26	10.94	8.41	8.62	9.01	7.57	7.31	9.04	9.18	7.61	9.30	8.74	7.85	8.41	10.45	8.75	0.99	11.3
<i>PPR</i>	4.90	4.40	5.12	4.59	5.99	5.85	5.06	5.05	6.11	6.00	5.85	5.48	6.86	5.62	5.81	5.26	5.50	0.64	11.57
<i>Fbp2</i>	9.14	8.40	7.39	8.56	7.91	6.89	8.42	10.70	8.32	8.34	8.01	7.41	8.46	6.70	7.58	8.71	8.18	0.95	11.58
<i>S8</i>	4.28	4.74	3.22	4.16	3.91	3.94	3.89	4.29	4.58	4.18	4.57	4.01	3.66	5.22	3.93	3.57	4.13	0.49	11.81
<i>CSA</i>	16.63	15.18	15.70	17.82	15.83	13.15	12.99	15.36	16.48	14.15	14.62	19.54	14.08	13.04	14.14	14.39	15.19	1.80	11.82
<i>Fbp3</i>	4.15	4.38	5.47	5.47	4.85	4.50	4.68	5.21	4.31	4.72	5.08	6.47	4.87	5.16	4.44	5.51	4.95	0.59	11.95



Supplementary figure S1. 2% agarose gel electrophoresis of 13 candidate reference genes



Supplementary figure S2. Melting curve of selected genes