

Supplementary Material

Identification of the bisabolol synthase in the endangered Candeia tree (*Eremanthus erythropappus* (DC) McLeisch)

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Supplementary table 1. Main components of Candeia tissues, at three development stages.

RT	RI	Compound	Sapling (<1 year)				5 year old tree						10 year old tree								Identification method				
			Leaf	Stem	Root	Twig	Limb	Leaf	New root	Root	Inflorescence	Seeds	Twig	Limb wood	Limb Bark	New leaf	Old leaf	New root	Old root	Old root core		Trunk core	Trunk bark		
11.80	1336	Elemene	1.20	-	-	-	-	-	0.23	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	RI, MS
11.91	1342	Presilphiperfol-7-ene	-	-	0.38	-	-	-	3.28	14.98	-	-	-	-	-	-	-	-	-	-	-	-	-	-	RI, MS
12.07	1348	α -Cubebene	-	-	0.14	-	-	-	0.70	-	-	-	-	-	-	0.22	0.37	-	-	-	-	-	-	-	RI, MS
12.13	1350	Silphin-1-ene	-	-	0.14	-	-	-	1.17	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	RI, MS
12.44	1370	Cyclosativene	-	-	0.24	-	-	-	2.70	-	-	-	-	-	-	0.45	-	-	-	-	-	-	-	-	RI, MS
12.40	1379	α -Copaene	30.61	-	1.46	24.17	-	16.17	2.23	-	15.37	-	1.54	-	-	23.41	23.40	-	-	-	-	-	-	-	RI, MS
12.60	1389	β -Elemene	4.65	-	1.27	-	-	1.92	1.64	-	1.51	6.98	-	0.12	-	1.79	2.46	0.25	-	0.23	-	-	-	-	RI, MS
12.72	1388	α -Isocomene	-	-	7.93	-	-	-	0.70	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	RI, MS
12.79	1399	sesquithujene	-	-	0.28	-	-	-	0.47	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	RI, MS
12.91	1402	cyperene	-	-	9.87	-	-	-	10.20	14.48	-	21.85	-	-	-	-	-	0.12	-	-	-	-	-	-	RI, MS
12.93	1413	α -Gurjunene	1.80	-	-	-	-	-	-	-	2.41	-	-	-	-	2.02	1.97	-	-	-	-	-	-	-	RI, MS
13.10	1420	(E)- β -Caryophyllene	28.03	68.90	7.40	34.49	-	28.19	4.30	-	27.14	-	1.34	-	-	23.53	24.63	0.12	-	-	-	-	-	-	RI, MS, ST
13.45	1437	Sesquisabinene-A	0.68	-	3.90	-	-	-	2.26	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	RI, MS
13.50	1440	Humulene	3.94	-	1.49	-	-	4.27	2.27	-	3.87	-	-	-	-	3.36	2.83	-	-	-	-	-	-	-	RI, MS
13.88	1480	Germacrene D	10.72	31.10	1.99	-	-	25.53	2.11	-	23.44	-	1.77	-	-	20.17	19.11	-	-	-	-	-	-	-	RI, MS
14.03	1485	β -Selinene	-	-	13.63	-	-	-	1.88	-	-	-	-	-	-	-	-	-	-	0.58	-	-	-	-	RI, MS
14.06	1494	Bicyclogermacrene	4.41	-	1.92	-	-	12.71	-	-	11.50	-	-	-	-	10.64	9.85	-	-	0.58	-	-	-	-	RI, MS
14.19	1496	α -Bisabolene	-	-	29.83	-	-	-	-	-	-	-	-	-	-	-	-	-	-	0.58	-	-	-	-	RI, MS
14.30	1513	Amorphene (γ -cadinene)	4.52	-	3.22	-	-	11.21	4.40	-	9.46	20.27	-	-	-	11.27	9.85	-	-	-	-	-	-	-	RI, MS
15.13	1570	Caryophyllene oxide	1.07	-	3.97	-	-	-	-	-	1.08	-	-	-	-	-	-	-	-	-	-	-	-	-	RI, MS
15.30	1592	Presilphiperfolanol	-	-	1.93	-	-	-	24.14	31.11	-	-	-	-	-	-	-	-	-	-	-	-	-	-	RI, MS
15.70	1610	Cubenol	-	-	0.46	-	-	-	8.44	-	1.08	-	-	-	-	-	-	-	-	-	-	-	-	-	RI, MS
15.80	1636	Cadinol	2.97	-	6.55	-	-	-	-	-	-	42.79	-	-	-	1.57	3.08	-	-	-	-	-	-	-	RI, MS
15.90	1652	α -Cadinol	2.81	-	-	-	-	-	3.75	-	3.20	8.11	-	-	-	1.62	2.46	-	-	0.60	0.60	1.97	-	-	RI, MS
15.96	1744	Bisabololoxide	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	0.62	0.60	1.97	-	-	-	-	RI, MS
16.27	1683	(-)- α -bisabolol sesquiterpene lactone (Vanillosmin)	-	-	-	26.11	36.59	-	0.59	5.71	-	-	32.31	76.92	0.43	-	-	37.45	44.18	39.35	30.23	-	-	-	RI, MS, ST
19.05		Diterpene I	2.58	-	0.31	-	-	-	-	-	-	-	-	-	49.79	-	-	5.26	7.88	35.88	46.51	35.50	-	-	MS
19.17		Diterpene I	-	-	0.99	15.23	23.65	-	2.34	17.55	-	-	1.54	-	1.42	-	-	-	0.17	-	-	-	-	-	MS
19.68		Diterpene II	-	-	0.66	-	39.76	-	20.16	16.17	-	-	61.54	22.96	48.36	-	-	56.18	46.57	20.83	23.26	64.50	-	-	MS

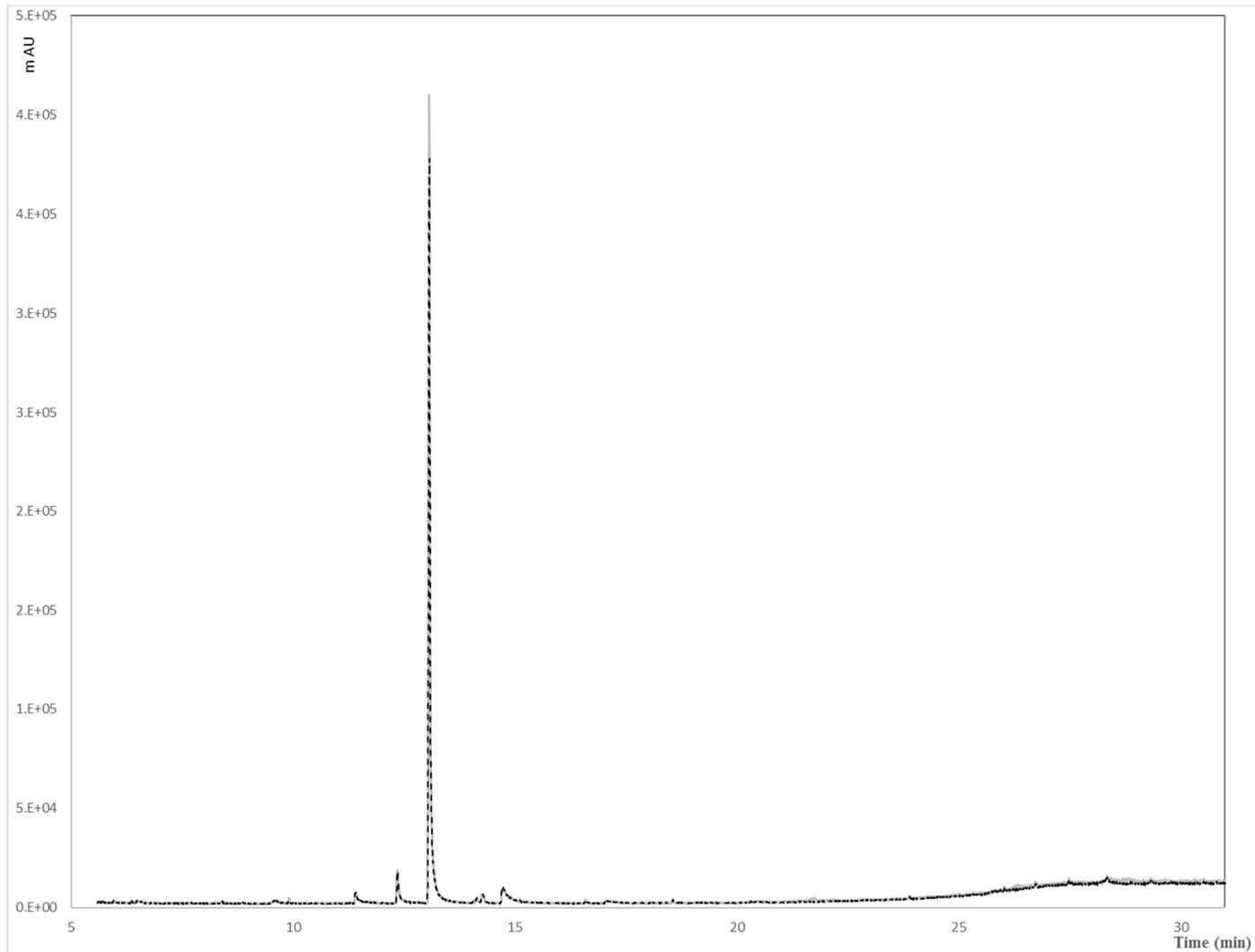
Supplementary table 2. Primers use during this study

Q-PCR_EeBOS	Fw	CGGGAAATTCGATTCTGGAG
Q-PCR_EeBOS	Rv	CAAGCACCCAGGCATATT
Q-PCR_EeEF1	Fw	GGCTGATTGTGCTGTTCTTATC
Q-PCR_EeEF1	Rv	ACCCAAGAGTGAAAGCAAG
Chrysolaeana EF	Fw	CTGGAGGAATTGTACAAG
Chrysolaeana EF	Rv	GCTTGACCCCAAGAGTGA
EeBOS_NotI	Rv	<u>ATATGCGGCCGCTCAAACA</u> ACTAAAGGGTGAACAACGAGC
EeBOS_Sall	Fw	<u>ATATGTCGACCGATCAAATCATGTCA</u> ACTGCTTTTCCA
RACE primer		CAGTGTAGTGCATTATCAGTGCGAGTG

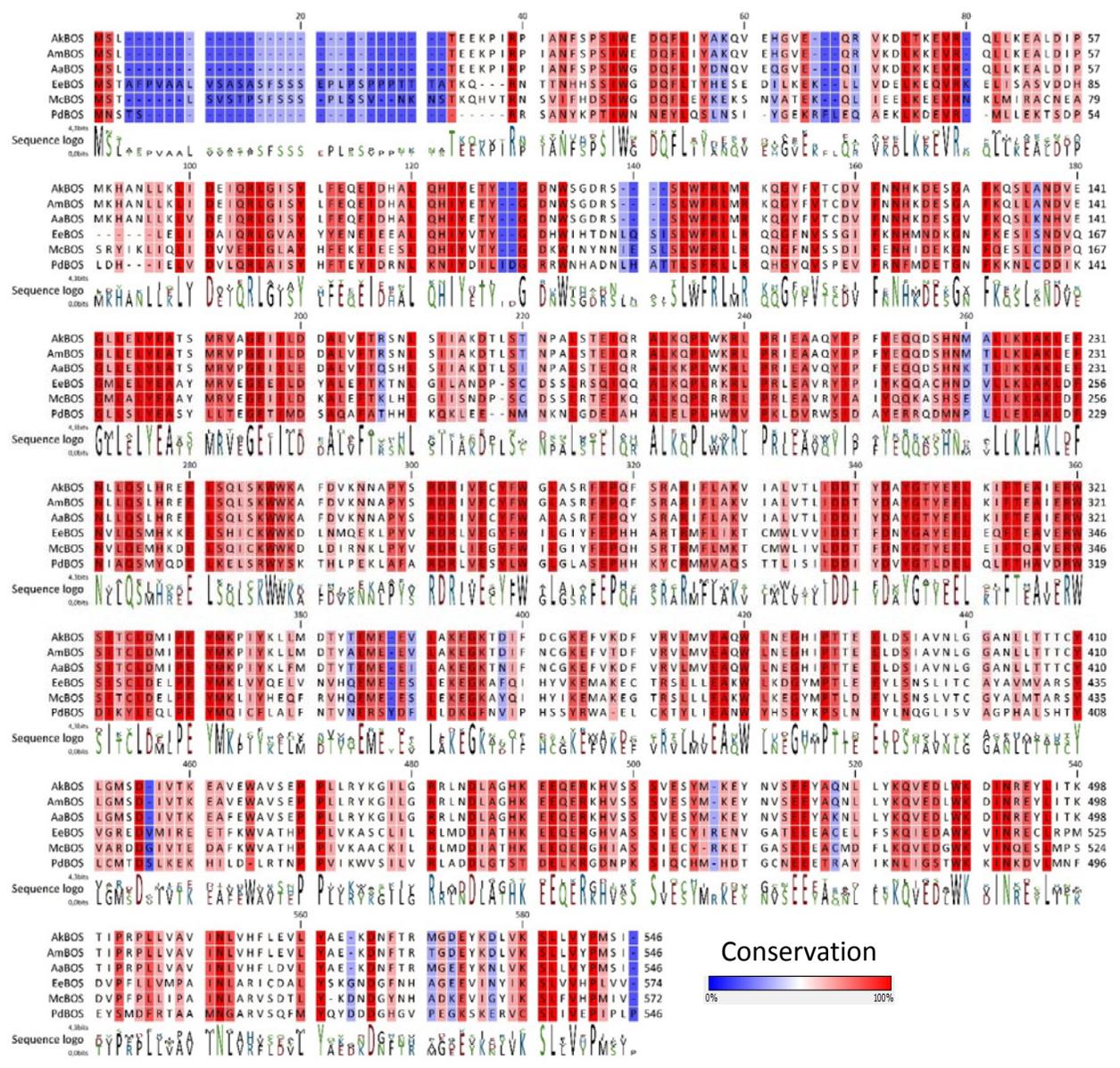
Supplementary table 3. Amino acid identity (%) of six Bisabolol synthase

		1	2	3	4	5	6
PdBOS	1	-	32.2	32.02	31.83	35.27	35.2
AmBOS	2	32.2	-	98.53	94.14	48.05	48.35
AkBOS	3	32.02	98.53	-	94.14	48.24	48.53
AaBOS	4	31.83	94.14	94.14	-	48.05	48.53
EeBOS	5	35.27	48.05	48.24	48.05	-	70.85
MrBOS	6	35.2	48.35	48.53	48.53	70.85	-

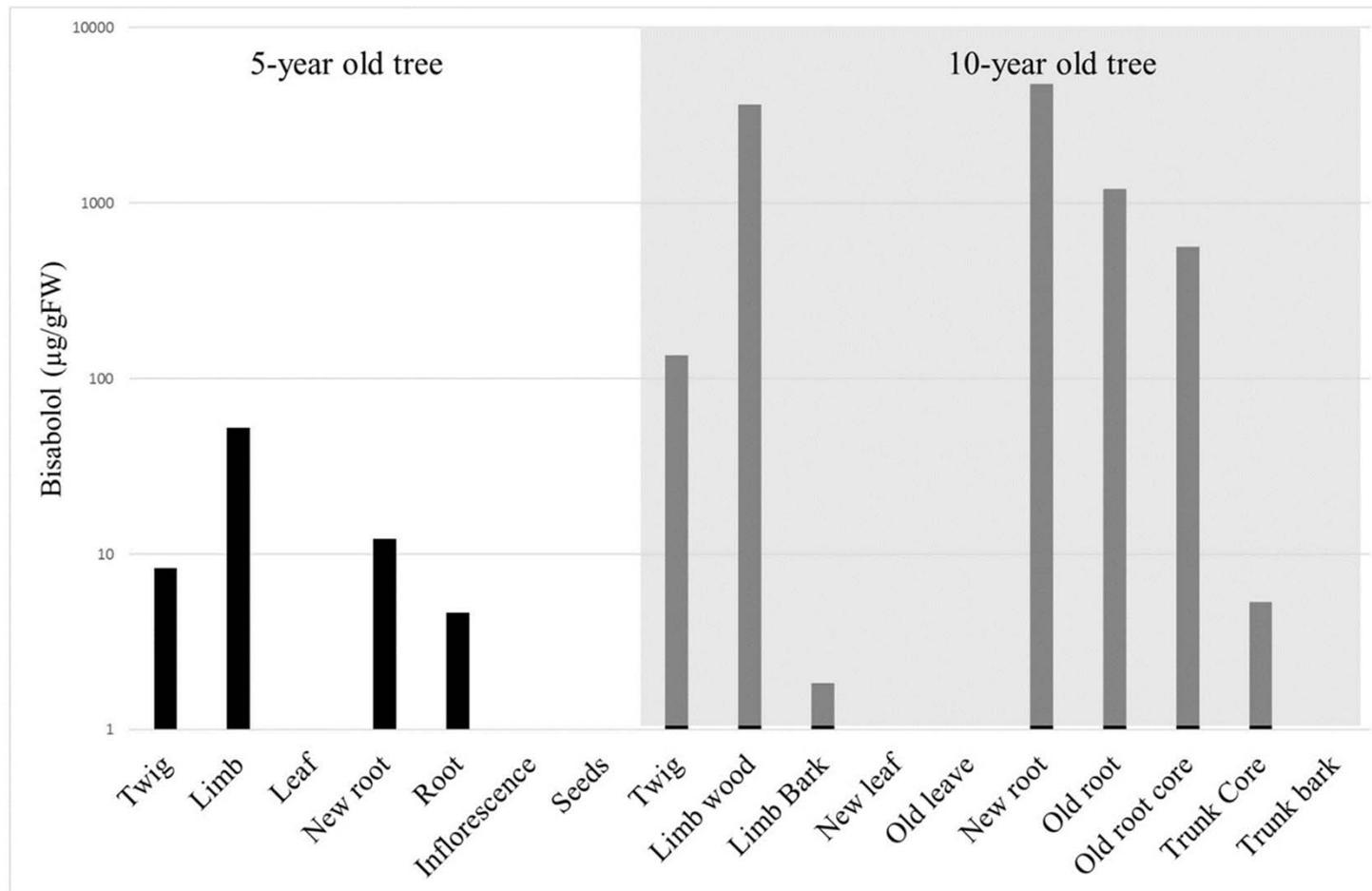
AkBOS = *Artemisia kurramensis* bisabolol synthase (BAW34955.1); AaBOS= *Artemisia annua* bisabolol synthase (AFV40969.1); AmBOS= *Artemisia maritima* bisabolol synthase (BAW34954.1); McBOS= *Matricaria recutita* bisabolol synthase (AIG92846.1); PdBOS= *Phyla dulcis* bisabolol synthase (AFR23372.1)



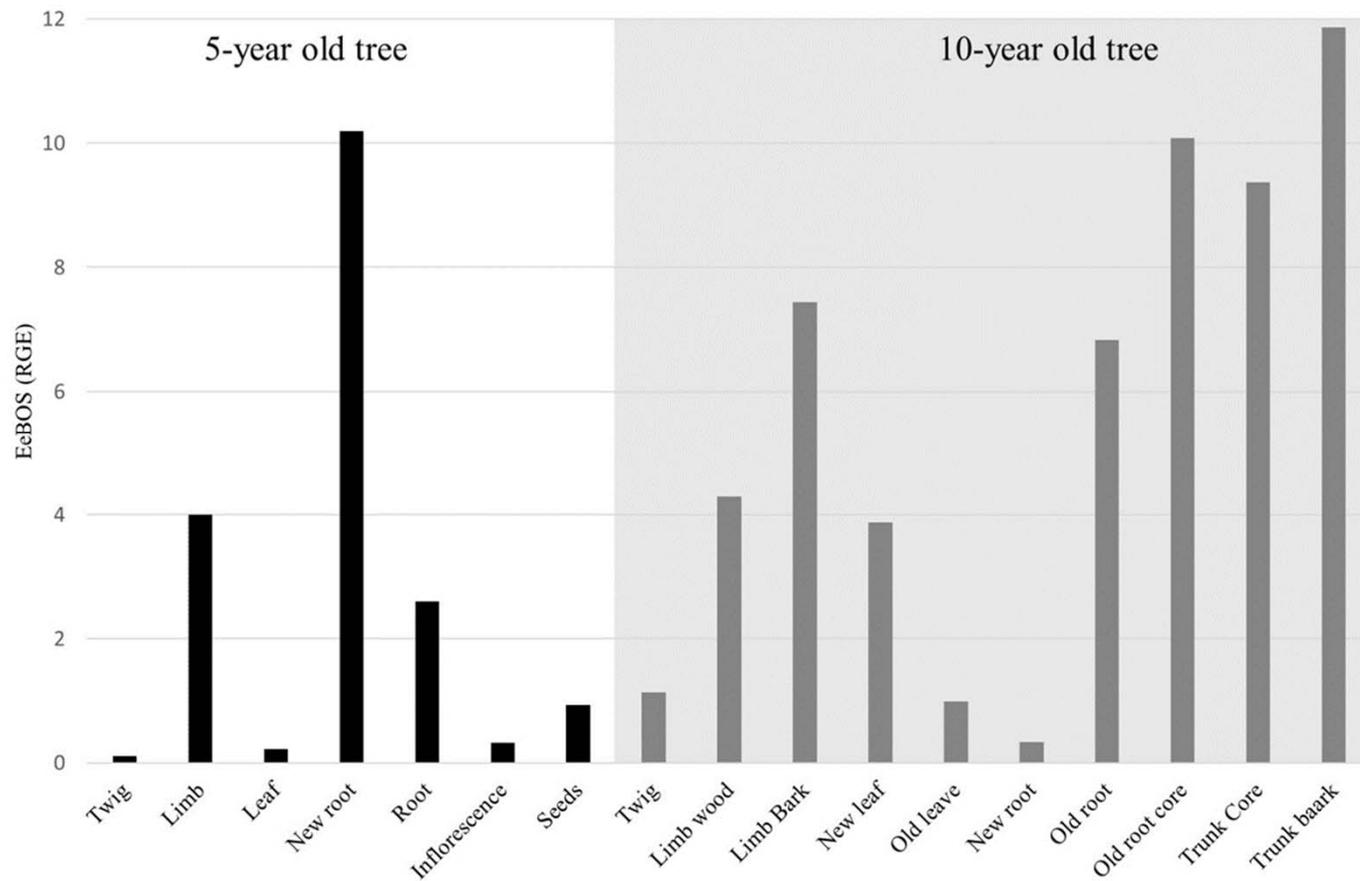
Supplementary figure 1. Detection of *in vitro* EeBOS products with GPP as a substrate. In vitro enzymatic assay with cell extracts expressing (black dotted line) or not (grey solid line) EeBOS were compared in the presence of GPP. The results were analysed by GC-MS.



Supplementary figure 2. Protein alignment between six bisabolol synthase. The conservation of amino acids in the protein are depicted with a range from low to highly conserved (from blue to red, respectively). AkBOS = *Artemisia kurramensis* bisabolol synthase (BAW34955.1); AaBOS= *Artemisia annua* bisabolol synthase (AFV40969.1); AmBOS= *Artemisia maritima* bisabolol synthase (BAW34954.1); McBOS= *Matricaria Recutita* bisabolol synthase (AIG92846.1); PdBOS= *Phyla dulcis* bisabolol synthase (AFR23372.1)



Supplementary figure 3. Bisabolol content in different tissues. (-)- α -Bisabolol was extracted with dichloromethane and quantified by GC-MS for different tissues originating from the 5 and 10-year old trees. Each bar represents a single analysis. The grayed out area of the bar diagram represent sample originating from the 10-year old tree. Each bar represents a single analysis.



Supplementary figure 4. Relative gene expression of EeBOS in different tissues. The reference gene used was EeEF1. The grayed out area of the bar diagram represent sample originating from the 10-year old tree. Each bar represents a single analysis.