



Supplémentar



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Supplementary Table 1 : Genotyping data of the EST-SSRs on *Moringa oleifera* genotypes

rs	alleles	G1	G2	G3	G4	G5	G6	G7	G8	G9	G10	G11	G12	G13	G14	G15	G16	G17	G18	G19	G20	G21	G22	G23	G24
MO13	A/G	R	R	A	G	A	A	R	G	R	R	G	R	R	R	R	R	R	R	R	R	G	R	R	
MO14	A/G	R	R	G	R	R	R	A	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	
MO15	A/G	R	A	R	A	R	A	G	R	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	
MO16	A/G	R	R	G	R	R	R	R	A	A	A	R	R	R	R	R	R	R	A	A	A	A	A	A	
MO17	A/G	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	
MO18	A/G	R	R	R	R	R	R	G	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	
MO19	A/G	R	A	A	R	R	R	R	A	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	
MO20	A/G	R	A	R	A	A	R	R	A	R	A	R	G	R	G	G	R	G	R	R	R	R	G	A	
MO21	A/G	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	
MO22	A/G	R	R	R	R	R	R	R	R	R	R	R	G	R	R	R	R	R	R	R	R	R	R	R	
MO23	A/G	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	
MO24	A/G	R	R	R	G	A	R	R	R	R	R	G	R	R	R	A	G	R	R	R	R	G	R	R	
MO25	A/G	R	R	R	R	R	R	R	R	R	R	A	R	R	R	R	R	R	R	R	R	R	R	R	
MO26	A/G	R	R	R	R	R	G	G	R	G	R	A	R	R	R	R	R	R	R	R	R	R	R	R	
MO27	A/G	G	R	R	R	R	A	R	A	R	A	R	A	R	R	R	R	R	R	R	R	R	R	R	
MO28	A/G	R	A	G	A	G	G	G	R	A	G	A	R	A	R	R	G	A	R	R	G	A	R	R	
MO29	A/G	R	A	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	
MO30	A/G	A	R	A	R	R	R	R	R	R	R	R	R	R	R	G	G	G	R	G	G	G	G	A	
MO31	A/G	G	A	A	A	R	R	A	R	R	A	A	A	R	R	R	R	R	G	A	R	A	R	A	
MO32	A/G	R	R	G	R	G	G	R	R	A	R	R	G	G	R	R	R	R	R	R	R	G	R	R	
MO33	A/G	A	R	G	A	A	A	A	A	R	R	R	R	G	G	G	G	G	R	R	G	R	R	R	
MO34	A/G	A	G	G	G	G	G	G	R	R	G	G	R	G	G	G	R	R	R	R	R	R	A	A	
MO35	A/G	A	R	A	R	R	R	R	R	R	A	G	G	R	R	G	R	A	G	R	R	G	R	R	
MO36	A/G	G	G	G	G	G	G	R	G	G	G	G	G	G	G	G	R	R	G	A	R	R	R	R	
MO37	A/G	R	R	G	R	A	R	R	R	R	R	R	R	R	R	R	R	R	G	R	G	R	R	R	
MO38	A/G	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	
MO39	A/G	R	A	R	G	G	R	R	A	A	R	R	R	R	R	R	R	R	R	A	G	R	R	R	
MO40	A/G	R	G	R	A	G	G	R	A	A	G	G	R	R	R	A	R	R	A	G	R	A	R	R	
MO41	A/G	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	G	R	G	R	G	
MO42	A/G	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	
MO43	A/G	R	A	R	G	A	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	A	
MO44	A/G	A	R	A	R	G	R	R	R	R	R	R	R	R	R	R	G	G	R	R	R	R	A	R	
MO45	A/G	R	R	R	R	R	R	R	R	R	R	G	R	R	R	R	R	R	R	R	R	R	R	G	
MO46	A/G	R	R	R	R	R	R	R	R	R	A	R	A	R	R	R	A	R	R	R	R	R	R	R	
MO47	A/G	R	A	R	R	R	R	R	R	R	G	R	R	R	R	R	R	R	G	R	R	R	R	R	
MO48	A/G	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	A	R	

*A- Parent 1 allele, G- Parent 2 allele, R- Heterozygote

Supplementary Table 2 : Marker association with traits/phenotypes obtained using GAPIT which is a R based software

Phenotypic trait	Marker associated	P-value	maf	nobs	Rsquare of Model without SNP	Rsquare of Model with SNP	FDR Adjusted P-values	effect
Total Phenols	MO15	0.03573	0.125	24	0.140982707	0.360433946	0.897624343	464.6535
FRAP Antioxidant activity mg/ 100 g (FW)	MO15	0.044616	0.125	24	0.154595288	0.35004726	0.541111501	597.0767
Antioxidant activity (DPPH)	MO43	0.042455	0.479167	24	0.234619244	0.415673741	0.58125149	-779.185
	MO17	0.048752	0.479167	24	0.234619244	0.404317985	0.58125149	-1498.18
	MO44	0.049791	0.479167	24	0.234619244	0.402606199	0.58125149	-804.938
H-Ca (%)	MO14	0.026793	0.5	24	0.282297423	0.489032556	0.711103098	-1.25293
I-Mg (%)	MO14	0.00502	0.5	24	0.255153806	0.629863155	0.18070865	-0.2995
	MO32	0.016255	0.3125	24	0.255153806	0.51408568	0.292591477	-0.17173
J-Fe (ppm)	MO44	0.02087	0.479167	24	0.148140159	0.418486233	0.751311657	80.89135
K-Mn (ppm)	MO44	0.012707	0.479167	24	-0.004897636	0.375359358	0.457452611	155.8797
	MO31	0.025774	0.3125	24	-0.004897636	0.289071245	0.463938949	129.215
L-Zn (ppm)	MO44	0.020141	0.479167	24	0.110294847	0.396432858	0.637277842	41.26446
	MO31	0.035404	0.3125	24	0.110294847	0.338490817	0.637277842	34.70698
M-Cu (ppm)	MO44	0.022929	0.479167	24	0.138232753	0.402111716	0.630725522	8.143296
	MO24	0.03504	0.458333	24	0.138232753	0.360253594	0.630725522	6.990332

