

Electronic supplementary material: Supplementary Table 2

Title: Genetic dissection of growth, wood basic density and gene expression in interspecific backcrosses of *Eucalyptus grandis* and *E. urophylla*

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Authors: Anand R.K. Kullan, Maria M van Dyk, Charles A. Hefer, Nicoletta Jones, Arnulf Kanzler, Alexander A. Myburg*

Affiliation and e-mail address of corresponding author:

Department of Genetics, Forestry and Agricultural Biotechnology Institute (FABI), University of Pretoria, Pretoria, 0002, South Africa

zander.myburg@fabi.up.ac.za

Supplementary Table 2a. Summary of the framework linkage maps constructed for QTL analysis in the *E. urophylla* backcross family.

	<i>E. urophylla</i> backcross parent map			<i>F</i> ₁ hybrid parent map (<i>E. urophylla</i> backcross family)		
Linkage group	Number of markers	Linkage group length (cM)	Mean distance between markers (cM)	Number of markers	Linkage group length (cM)	Mean distance between markers (cM)
1	8	82.4	10.3	11	91.3	8.3
2	8	38.6	4.8	12	101.0	8.4
3	13	100.4	7.7	12	107.7	8.9
4	8	72.0	9.0	12	60.1	5.0
5	10	88.9	8.8	13	95.2	7.3
6	14	100.6	7.1	14	124.3	8.8
7	10	79.5	7.9	9	74.0	8.2
8	13	112.4	8.6	13	108.4	8.3
9	8	95.4	11.9	12	74.0	6.1
10	10	97.1	9.7	12	91.9	7.6
11	10	87.6	8.7	10	90.7	9.0
Total	112	954.9	8.5	130	1018.0	7.8

BC, backcross; QTL, Quantitative trait locus.

Supplementary Table 2b. Markers used for QTL analysis, their positions and additive effects in the *E. urophylla* backcross family.

E. urophylla backcross parent					F1 hybrid of E. urophylla BC family				
Marker Name	Linkage group (LG)	Map Position (cM)	Additive effect (SD, DBH)	Additive effect (SD, Density)	Marker Name	Linkage group (LG)	Map Position (cM)	Additive effect (SD, DBH) ^a	Additive effect (SD, Density) ^a
ePt_638907	LG1	0.00	0.05	0.06	ePt_503701	LG1	0.00	0.02	0.01
ePt_640386	LG1	2.73	0.06	0.04	ePt_503224	LG1	10.54	0.01	0.00
ePt_571510	LG1	16.49	0.08	0.06	ePt_573788	LG1	17.87	0.06	0.05
ePt_636534	LG1	46.12	0.13	0.03	ePt_572856	LG1	30.50	-0.04	-0.10
ePt_571936	LG1	50.44	0.17	0.00	ePt_600232	LG1	36.17	0.05	-0.07
ePt_568625	LG1	70.49	0.12	0.07	ePt_642824	LG1	46.92	0.22	-0.03
ePt_641800	LG1	78.78	0.15	0.05	ePt_599740	LG1	56.39	0.16	-0.11
ePt_636825	LG1	82.36	0.14	0.03	ePt_565915	LG1	65.47	0.30	-0.06
ePt_637636	LG2	0.00	0.05	0.06	ePt_600664	LG1	81.49	-0.10	-0.06
ePt_563419	LG2	4.62	0.07	0.08	ePt_639849	LG1	87.91	0.02	-0.06
ePt_565691	LG2	14.27	0.14	0.12	ePt_563014	LG1	91.34	0.01	-0.06
ePt_574308	LG2	22.37	0.12	0.06	ePt_505021	LG2	0.00	0.19	-0.15
ePt_637223	LG2	26.94	0.23	0.11	ePt_574374	LG2	7.92	0.13	-0.04
ePt_575592	LG2	29.16	0.27	0.09	ePt_567712	LG2	16.07	0.09	-0.04
ePt_575031	LG2	36.01	0.21	0.14	ePt_569091	LG2	17.86	0.09	-0.05
ePt_566116	LG2	38.56	0.20	0.12	ePt_639347	LG2	36.63	-0.14	-0.02
ePt_564331	LG3	0.00	0.08	0.05	ePt_504902	LG2	42.22	-0.09	-0.02
ePt_503896	LG3	4.05	0.03	0.06	ePt_573048	LG2	53.26	-0.10	-0.12
ePt_599238	LG3	9.08	0.10	0.07	ePt_568036	LG2	68.07	-0.14	-0.07
ePt_566346	LG3	15.38	0.11	0.03	ePt_567298	LG2	76.97	-0.03	-0.22
ePt_643823	LG3	20.67	0.12	0.03	ePt_503852	LG2	92.48	0.09	-0.33
ePt_641822	LG3	26.72	0.22	0.02	ePt_565848	LG2	95.27	0.07	-0.30
ePt_504319	LG3	38.47	0.23	0.07	ePt_641866	LG2	101.00	0.02	-0.39
ePt_637193	LG3	45.45	0.11	0.11	ePt_644293	LG3	0.00	0.12	-0.13

ePt_572258	LG3	59.13	0.03	0.06	ePt_568037	LG3	5.44	0.04	-0.07
ePt_644491	LG3	70.44	0.08	0.20	ePt_572721	LG3	13.68	0.07	-0.12
ePt_642546	LG3	79.28	0.08	0.19	ePt_568116	LG3	33.73	-0.04	0.07
ePt_600566	LG3	92.96	0.01	0.20	ePt_644403	LG3	45.94	-0.03	0.03
ePt_566961	LG3	100.42	0.03	0.17	ePt_640977	LG3	53.33	-0.02	-0.08
ePt_567857	LG4	0.00	0.35	0.10	ePt_640786	LG3	59.57	0.02	-0.13
ePt_567945	LG4	3.07	0.29	0.16	ePt_567607	LG3	67.42	0.13	-0.31
ePt_569322	LG4	7.23	0.25	0.19	ePt_571789	LG3	76.02	0.17	-0.35
ePt_642860	LG4	30.64	0.07	0.12	ePt_566415	LG3	87.80	0.23	-0.36
ePt_637285	LG4	37.23	0.11	0.02	ePt_636665	LG3	93.39	0.17	-0.33
ePt_599283	LG4	58.90	0.05	0.21	ePt_570151	LG3	107.70	0.15	-0.38
ePt_570780	LG4	65.23	0.06	0.24	ePt_570493	LG4	0.00	0.07	0.04
ePt_636880	LG4	72.03	0.07	0.19	ePt_600655	LG4	3.79	0.10	0.04
ePt_643707	LG5	0.00	0.08	0.12	ePt_503766	LG4	4.62	0.11	0.01
ePt_639138	LG5	6.23	0.15	0.14	ePt_643984	LG4	6.27	0.12	0.10
ePt_571353	LG5	18.81	0.01	0.04	ePt_636515	LG4	14.28	0.18	0.06
ePt_572057	LG5	35.54	0.05	0.02	ePt_571769	LG4	22.42	0.19	-0.01
ePt_636498	LG5	48.88	0.08	0.05	ePt_573193	LG4	25.58	0.19	-0.06
ePt_573321	LG5	55.79	0.01	0.06	ePt_599702	LG4	29.14	0.18	0.51
ePt_636480	LG5	66.81	0.01	0.15	ePt_573837	LG4	34.75	0.15	0.56
ePt_643842	LG5	78.12	0.03	0.12	ePt_638305	LG4	39.77	0.19	0.62
ePt_600653	LG5	85.68	0.08	0.18	ePt_503674	LG4	57.27	0.16	-0.17
ePt_503808	LG5	88.87	0.07	0.20	ePt_570682	LG4	60.09	0.19	-0.13
ePt_568290	LG6	0.00	0.03	0.17	ePt_570054	LG5	0.00	-0.03	0.04
ePt_575429	LG6	1.12	0.04	0.16	ePt_643259	LG5	9.46	0.09	0.01
ePt_636902	LG6	5.86	0.07	0.11	ePt_567769	LG5	17.14	0.08	-0.05
ePt_505086	LG6	14.44	0.02	0.01	ePt_644484	LG5	19.99	0.07	-0.06
ePt_636956	LG6	17.66	0.02	0.05	ePt_570274	LG5	31.29	0.08	-0.12
ePt_644163	LG6	31.49	0.27	0.04	ePt_575685	LG5	36.70	0.07	-0.11
ePt_638183	LG6	43.44	0.24	0.06	ePt_639908	LG5	45.52	0.15	-0.04

ePt_637303	LG6	50.76	0.28	0.10	ePt_569419	LG5	50.88	0.21	-0.05
ePt_569463	LG6	63.45	0.08	0.02	ePt_568347	LG5	57.71	0.15	-0.01
ePt_641907	LG6	67.82	0.16	0.02	ePt_575678	LG5	64.98	0.11	0.02
ePt_639634	LG6	82.84	0.14	0.16	ePt_574584	LG5	78.05	0.09	0.04
ePt_640477	LG6	89.20	0.21	0.16	ePt_573934	LG5	80.80	0.11	0.00
ePt_565994	LG6	103.14	0.23	0.10	ePt_643170	LG5	95.23	0.04	0.01
ePt_566534	LG6	110.56	0.28	0.10	ePt_565718	LG6	0.00	-0.17	-0.37
ePt_566646	LG7	0.00	0.06	0.05	ePt_641950	LG6	23.00	-0.19	-0.47
ePt_571137	LG7	2.67	0.07	0.20	ePt_563588	LG6	27.73	0.18	-0.44
ePt_573687	LG7	10.17	0.12	0.29	ePt_599939	LG6	37.06	0.36	-0.03
ePt_637829	LG7	10.61	0.11	0.31	ePt_504039	LG6	45.05	0.47	-0.15
ePt_563468	LG7	19.98	0.11	0.37	ePt_566935	LG6	59.85	0.25	-0.20
ePt_637575	LG7	35.70	0.16	0.04	ePt_641192	LG6	73.55	0.09	-0.18
ePt_566192	LG7	62.52	0.05	0.15	ePt_643704	LG6	76.43	0.10	-0.19
ePt_571262	LG7	70.61	0.06	0.18	ePt_504218	LG6	85.57	-0.17	-0.32
ePt_644456	LG7	72.78	0.09	0.19	ePt_643314	LG6	101.90	-0.10	-0.21
ePt_566482	LG7	79.47	0.03	0.27	ePt_575722	LG6	107.12	-0.22	-0.18
ePt_639224	LG8	0.00	0.09	0.07	ePt_638034	LG6	114.40	0.12	-0.07
ePt_504097	LG8	8.17	0.04	0.01	ePt_568058	LG6	118.37	-0.28	-0.02
ePt_565436	LG8	11.79	0.04	0.03	ePt_574570	LG6	124.32	-0.18	-0.01
ePt_575292	LG8	12.50	0.05	0.19	ePt_642265	LG7	0.00	-0.23	-0.14
ePt_574247	LG8	36.24	0.04	0.37	ePt_568515	LG7	5.52	-0.23	-0.06
ePt_639354	LG8	44.53	0.02	0.35	ePt_571000	LG7	23.29	-0.14	0.12
ePt_637292	LG8	53.21	0.07	0.00	ePt_575149	LG7	34.99	-0.07	0.01
ePt_564161	LG8	67.69	0.01	0.01	ePt_643010	LG7	43.49	-0.10	-0.04
ePt_503998	LG8	73.25	0.01	0.02	ePt_574160	LG7	46.16	-0.07	-0.02
ePt_637028	LG8	90.80	0.12	0.07	ePt_639371	LG7	56.16	-0.09	-0.05
ePt_574691	LG8	93.35	0.13	0.07	ePt_568436	LG7	64.60	-0.11	-0.08
ePt_570192	LG8	107.51	0.11	0.00	ePt_564451	LG7	73.97	-0.08	-0.03
ePt_644171	LG8	112.35	0.17	0.06	ePt_504398	LG8	0.00	0.05	0.26

ePt_572414	LG9	0.00	0.17	0.05	ePt_571985	LG8	11.03	0.09	0.21
ePt_571185	LG9	13.53	0.13	0.13	ePt_565753	LG8	16.28	0.16	0.14
ePt_571912	LG9	25.42	0.26	0.16	ePt_599669	LG8	20.98	0.17	0.05
ePt_566194	LG9	30.85	0.40	0.20	ePt_572210	LG8	29.04	0.15	0.03
ePt_563442	LG9	46.07	0.16	0.21	ePt_643850	LG8	48.00	0.04	-0.27
ePt_642787	LG9	72.00	0.04	0.04	ePt_503845	LG8	53.61	-0.01	-0.33
ePt_573318	LG9	74.05	0.04	0.08	ePt_640365	LG8	59.81	0.03	-0.38
ePt_599981	LG9	95.36	0.13	0.13	ePt_640629	LG8	67.70	0.06	-0.33
ePt_641100	LG10	0.00	0.40	0.16	ePt_575022	LG8	79.29	0.16	-0.37
ePt_570364	LG10	3.51	0.43	0.21	ePt_640149	LG8	88.85	0.20	-0.27
ePt_640576	LG10	20.93	0.06	0.12	ePt_572120	LG8	97.61	0.22	-0.22
ePt_575378	LG10	32.04	0.03	0.12	ePt_573365	LG8	108.47	0.10	-0.22
ePt_568620	LG10	46.40	0.03	0.09	ePt_575684	LG9	0.00	-0.04	-0.27
ePt_644096	LG10	61.93	0.04	0.05	ePt_641785	LG9	7.11	0.05	-0.21
ePt_563364	LG10	73.27	0.03	0.14	ePt_567963	LG9	9.11	0.07	-0.18
ePt_642097	LG10	83.94	0.22	0.19	ePt_572213	LG9	12.91	0.13	-0.18
ePt_643528	LG10	88.39	0.18	0.22	ePt_567473	LG9	26.26	0.23	0.15
ePt_638623	LG10	97.06	0.18	0.23	ePt_644133	LG9	35.99	0.19	0.34
ePt_599825	LG11	0.00	0.04	0.02	ePt_639625	LG9	40.55	0.11	0.42
ePt_599720	LG11	3.36	0.06	0.10	ePt_640713	LG9	42.30	0.12	0.45
ePt_566717	LG11	13.92	0.04	0.11	ePt_599692	LG9	52.17	0.09	0.59
ePt_564887	LG11	17.94	0.01	0.14	ePt_600528	LG9	61.79	0.04	0.68
ePt_641324	LG11	22.85	0.02	0.13	ePt_639912	LG9	70.95	0.12	0.64
ePt_640702	LG11	25.43	0.07	0.09	ePt_504705	LG9	73.95	0.09	0.65
ePt_643222	LG11	37.19	0.07	0.21	ePt_565743	LG10	0.00	0.12	0.42
ePt_00411	LG11	50.32	0.10	0.25	ePt_637643	LG10	6.53	0.04	0.40
ePt_571164	LG11	72.78	0.07	0.09	ePt_504458	LG10	28.94	0.07	0.20
ePt_503841	LG11	87.60	0.24	0.05	ePt_643963	LG10	36.64	0.09	0.24
					ePt_568851	LG10	42.77	0.11	0.23
					ePt_644176	LG10	45.86	0.05	0.17

ePt_600200	LG10	50.88	-0.03	0.57
ePt_566772	LG10	62.32	-0.11	0.64
ePt_638929	LG10	66.87	-0.11	0.62
ePt_574725	LG10	77.32	-0.04	0.16
ePt_575312	LG10	87.79	-0.13	0.23
ePt_640871	LG10	91.92	-0.18	0.28
ePt_571374	LG11	0.00	0.11	0.07
ePt_638531	LG11	2.23	0.08	0.11
ePt_567666	LG11	11.06	0.13	0.14
ePt_575643	LG11	25.68	0.01	0.27
ePt_504138	LG11	29.34	-0.02	0.26
ePt_504803	LG11	46.11	-0.07	0.15
ePt_570513	LG11	52.30	-0.09	0.14
ePt_600156	LG11	60.11	-0.11	0.10
ePt_573069	LG11	69.61	-0.10	0.07
ePt_503383	LG11	90.66	0.07	0.11

SD, Standard deviation. QTL effects at marker positions based on composite interval mapping (Model 6, QTL Cartographer v2.5).

^a Marker data in the F1 hybrid maps were recoded so that positive and negative additive effect values on all linkage groups are associated with the effect of replacement of the *E. grandis* allele with the *E. urophylla* allele in backcross progeny. Linkage phases were arbitrary from one linkage group to the next for the backcross parents and the directions of the effect are therefore not indicated for these parents.