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High-density genetic linkage maps with over 2000 sequence-anchored DArT markers for genetic dissection in *Eucalyptus grandis* x *E. urophylla* hybrids

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Supplemental Figure S3. Alignment of the *Eucalyptus grandis* x *E. urophylla* consensus linkage maps with the draft (JGI-DOE V1.0) assembly of *E. grandis* genome.

Left: Consensus genetic map in cM derived from *E. grandis*, *E. urophylla* and their F1 hybrid. Right: Scaffolds from the draft genome assembly (V1.0) of *E. grandis* corresponding to the linkage group. The linkage map alignment was visualized graphically with MapChart.

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<th>LG1</th>
<th>Scaffold 1</th>
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