**Supporting Information Table S1. Information on source of RNA used for next-generation sequencing of 24 tropical tree species.** Normally, embryonic tissue was pooled from 2 to 4 individuals to provide sufficient material for RNA extraction.

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| **Species** | **Source of RNA** |
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| *Acacia mangium* | Root only |
| *Acacia senegal* | Coleoptile only |
| *Acrocarpus fraxinifolius* | Root only |
| *Adansonia digitata* | Root only |
| *Albizia lebbeck* | Root only |
| *Calliandra calothyrsus* | Root only |
| *Diospyros mespiliformis* | Root only |
| *Enterolobium cyclocarpum* | Root only |
| *Faidherbia albida* | Root only |
| *Gliricidia sepium* | Root only |
| *Jacaranda mimosifolia* | Root only |
| *Jatropha curcas* | Root only |
| *Leucaena diversifolia* | Root only |
| *Leucaena leucocephala* | Root and embryo |
| *Moringa stenopetala* | Root only |
| *Prunus africana* | Root and coleoptile |
| *Samanea saman* | Root and embryo |
| *Senna siamea* | Root only |
| *Sesbania macrantha* | Root only |
| *Sesbania sesban* | Root only |
| *Tephrosia candida* | Root and coleoptile |
| *Tipuana tipu* | Root and coleoptile |
| *Warburgia ugandensis* | Root only |
| *Ziziphus mauritiana* | Root only |
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