

Figure S9. Normalized relative quantity (NRQ, compared to *Ae. tauschii* (D), set as 1, see text) of Gypsy LTR retrotransposons in the following species: *T. urartu* (A^u), *Ae. speltoides* (S), *Ae. tauschii* (D) (Yaakov et al., 2013); *P. spicata* (St), *Th. bessarabicum* (J^b), *D. villosum* (V). The area of the circle is equal to the NRQ of each transposable element.

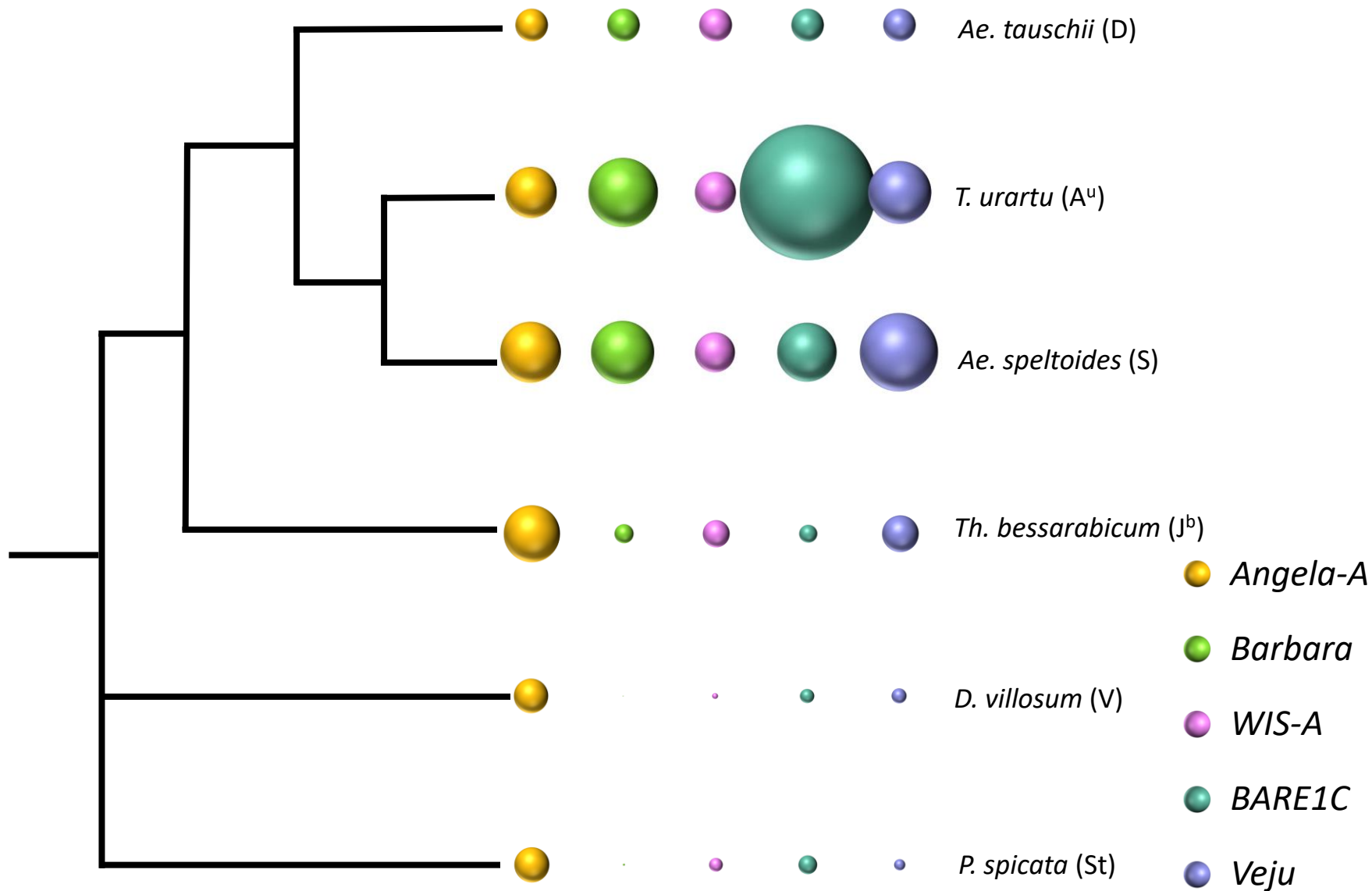


Figure S10. Normalized relative quantity (NRQ, compared to *Ae. tauschii* (D), set as 1, see text) of *Copia* LTR retrotransposons in the following species: *T. urartu* (A^u), *Ae. speltoides* (S), *Ae. tauschii* (D) (Yaakov et al., 2013); *P. spicata* (St), *Th. bessarabicum* (J^b), *D. villosum* (V). The area of the circle is equal to the NRQ of each transposable element.

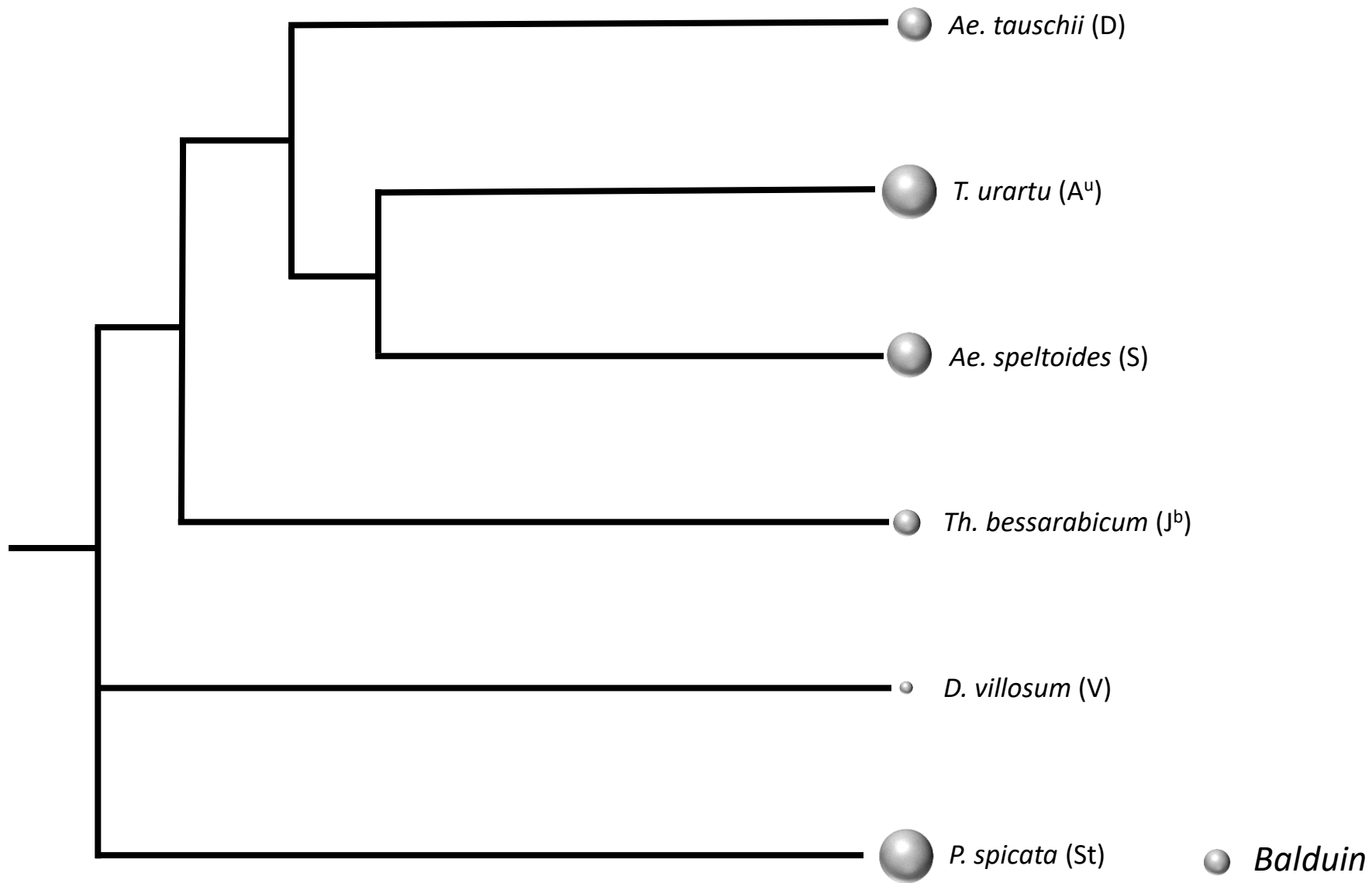


Figure S11. Normalized relative quantity (NRQ, compared to *Ae. tauschii* (D), set as 1, see text) of DNA transposon *Balduin* in the following species: *T. urartu* (A^u), *Ae. speltooides* (S), *Ae. tauschii* (D) (Yaakov et al., 2013); *P. spicata* (St), *Th. bessarabicum* (J^b), *D. villosum* (V). The area of the circle is equal to the NRQ of each transposable element.

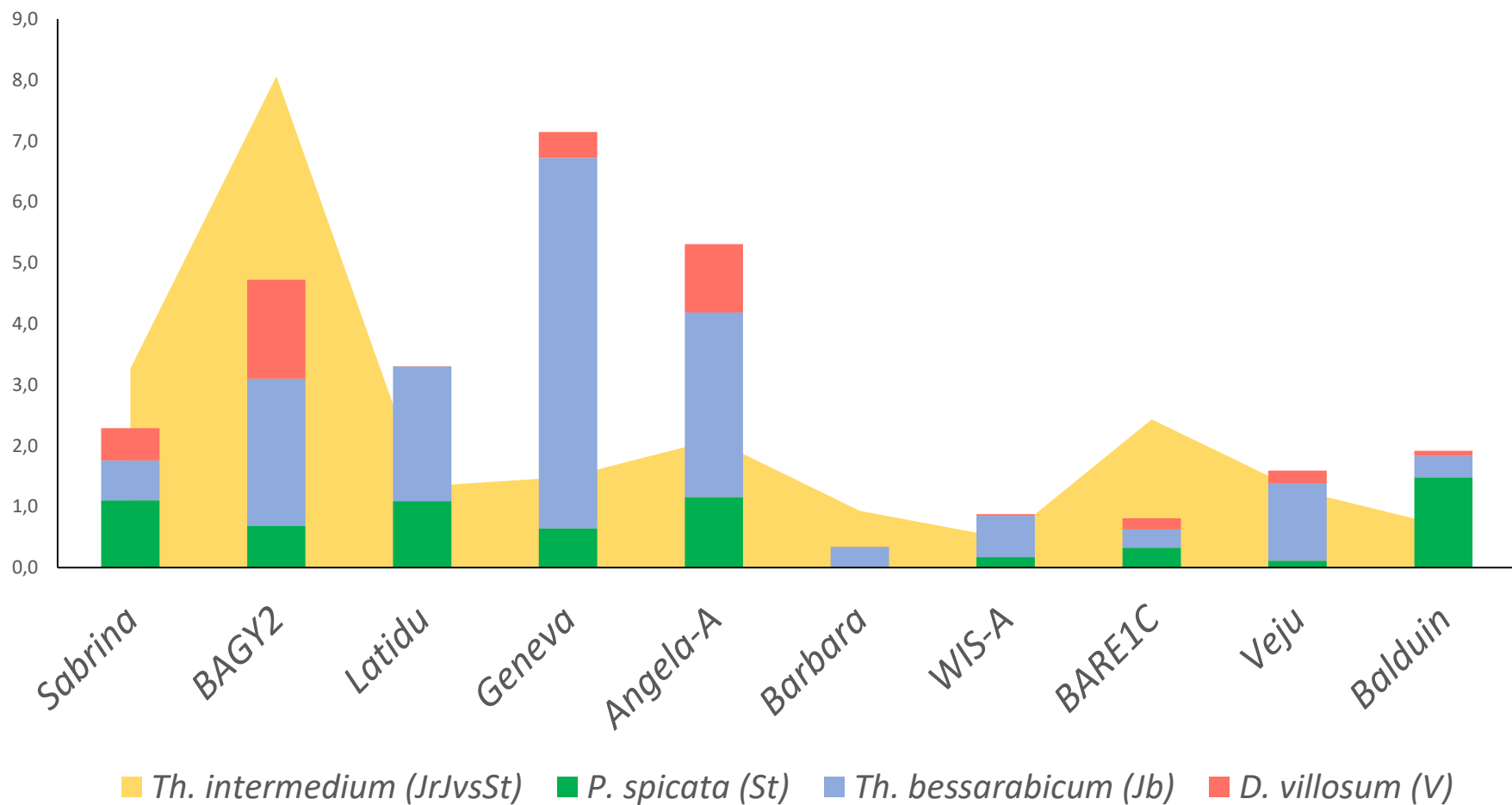


Figure S12. Normalized relative quantity (compared to *Ae. tauschii* (D), set as 1, see text) of the studied mobile elements in the following species: *P. spicata* (St), *Th. bessarabicum* (J^b), *D. villosum* (V), *Th. intermedium* (J^rJ^{vs}St, per whole allopolyploid genome).