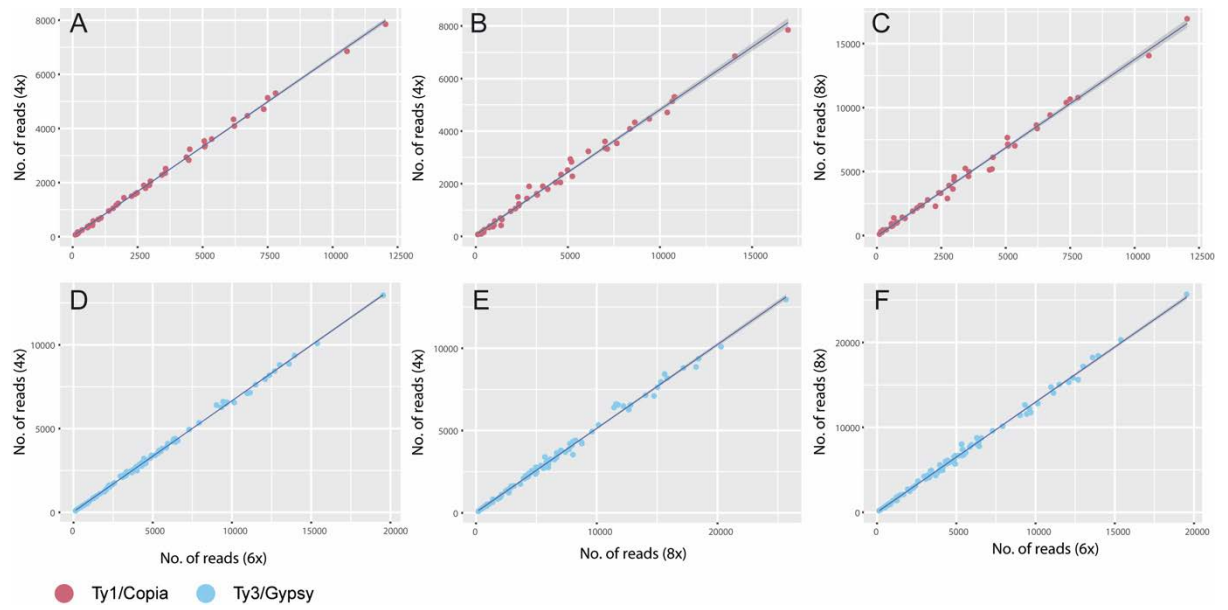


Supplementary Figure S1. Genomic composition of all *Senecio doricum* accessions across cytotypes (4x, 6x and 8x). Estimates of the genomic abundances (in Mb/1C) of different repeats are indicated and colored by repeat class [GP = Genome proportion (%), LTR = Long Terminal Repeat].



Supplementary Figure S2. Pairwise scatterplot comparisons of the number of reads included in repeat clusters from each cytotype. The slope indicates the genome size ratio between each cytotype. (A–C) Ty1/Copia. (D–F) Ty3/Gypsy.