Phylogenetic relationships and taxonomic issues in *Gastridium* (Poaceae) inferred from plastid and nuclear DNA sequence analysis.
Scoppola et al. 2020

Supplementary Material

Figure S5a. *trn*L-*trn*F haplotype network of the investigated dataset. Coloured ellipses circumscribe the four investigated *Gastridium* taxa. Taxa colours as in Fig.1. Different line thickness indicates the relative number of mutations. *Ce: Cynosurus echinatus, Ae: Alopecurus aequalis, Gf: Gaudinia fragilis, Bm: Briza minor, Ac: Agrostis castellana, Pm: Polypogon monspeliensis, Tn: Triplachne nitens, Gv: Gastridium ventricosum, Gp: G. phleoides, Gs: G. scabrum, Gl: G. lainzii.*

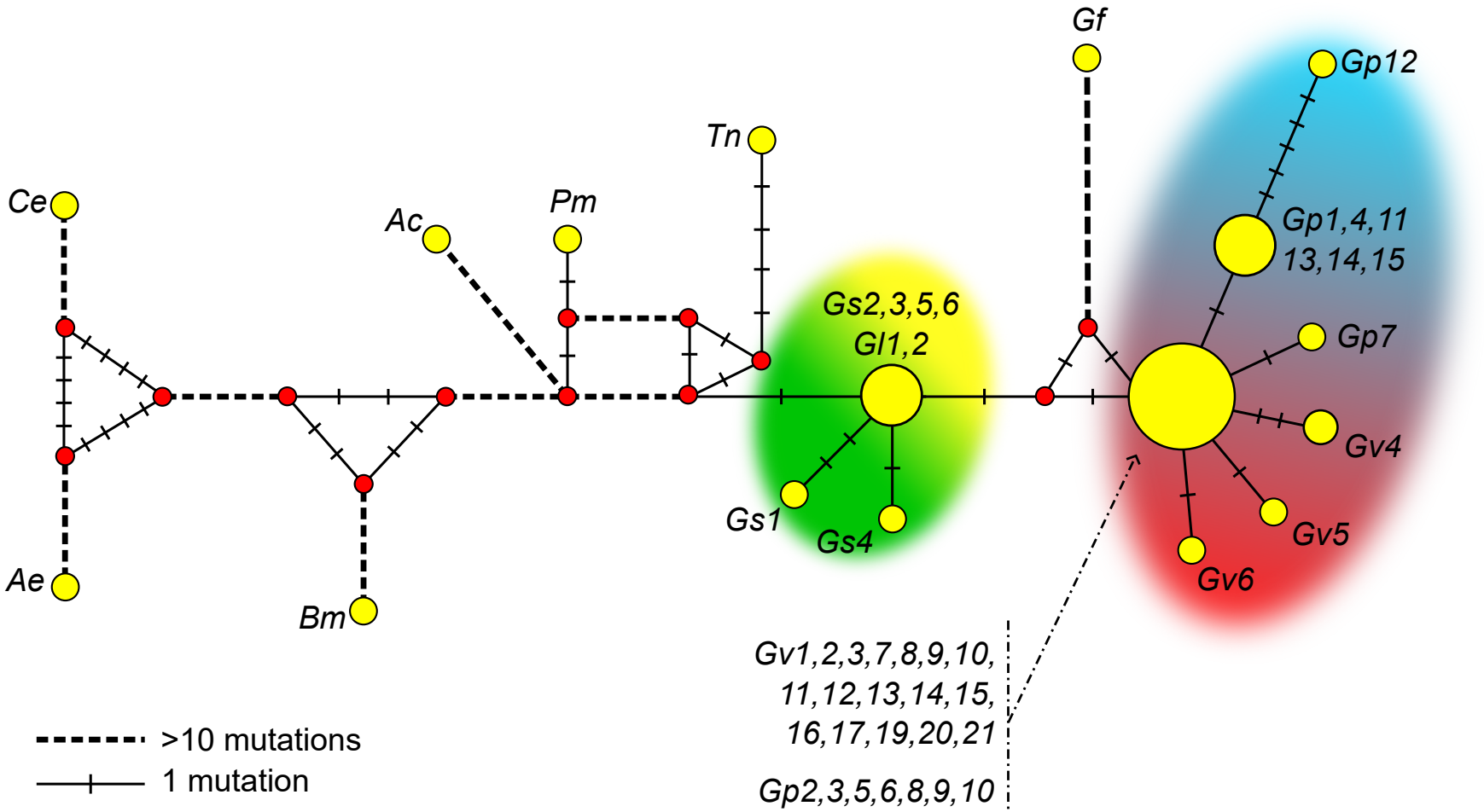
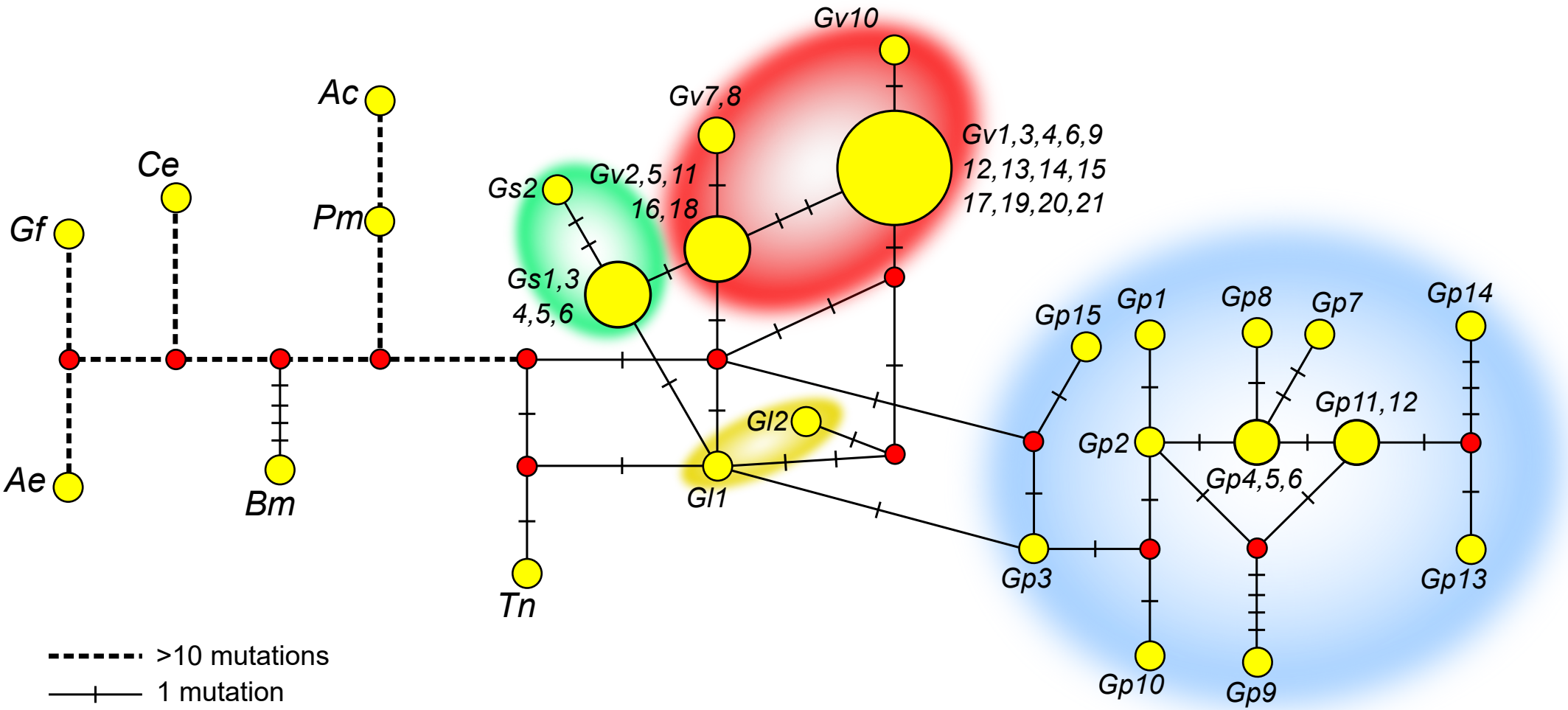
Figure S5b. *trn*H-*psb*A haplotype network of the investigated dataset. Coloured ellipses circumscribe the four investigated *Gastridium* taxa.Taxa colours as in Fig.1. Different line thickness indicates the relative number of mutations. *Ce: Cynosurus echinatus, Ae: Alopecurus aequalis, Gf: Gaudinia fragilis, Bm: Briza minor, Ac: Agrostis castellana, Pm: Polypogon monspeliensis, Tn: Triplachne nitens, Gv: Gastridium ventricosum, Gp: G. phleoides, Gs: G. scabrum, Gl: G. lainzii.*