**TE: Dmel−B−R1202−Map4−CLASSI/LTR/Gypsy**
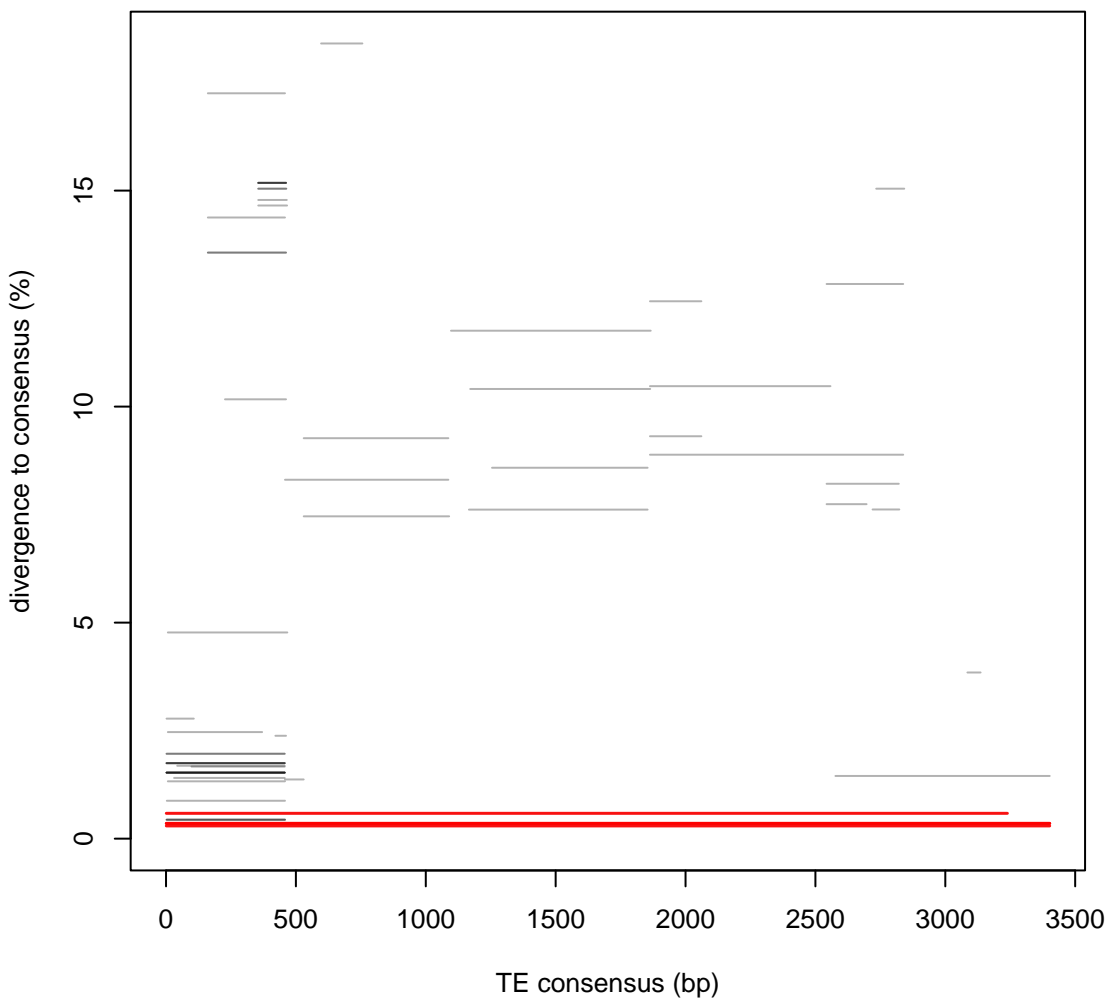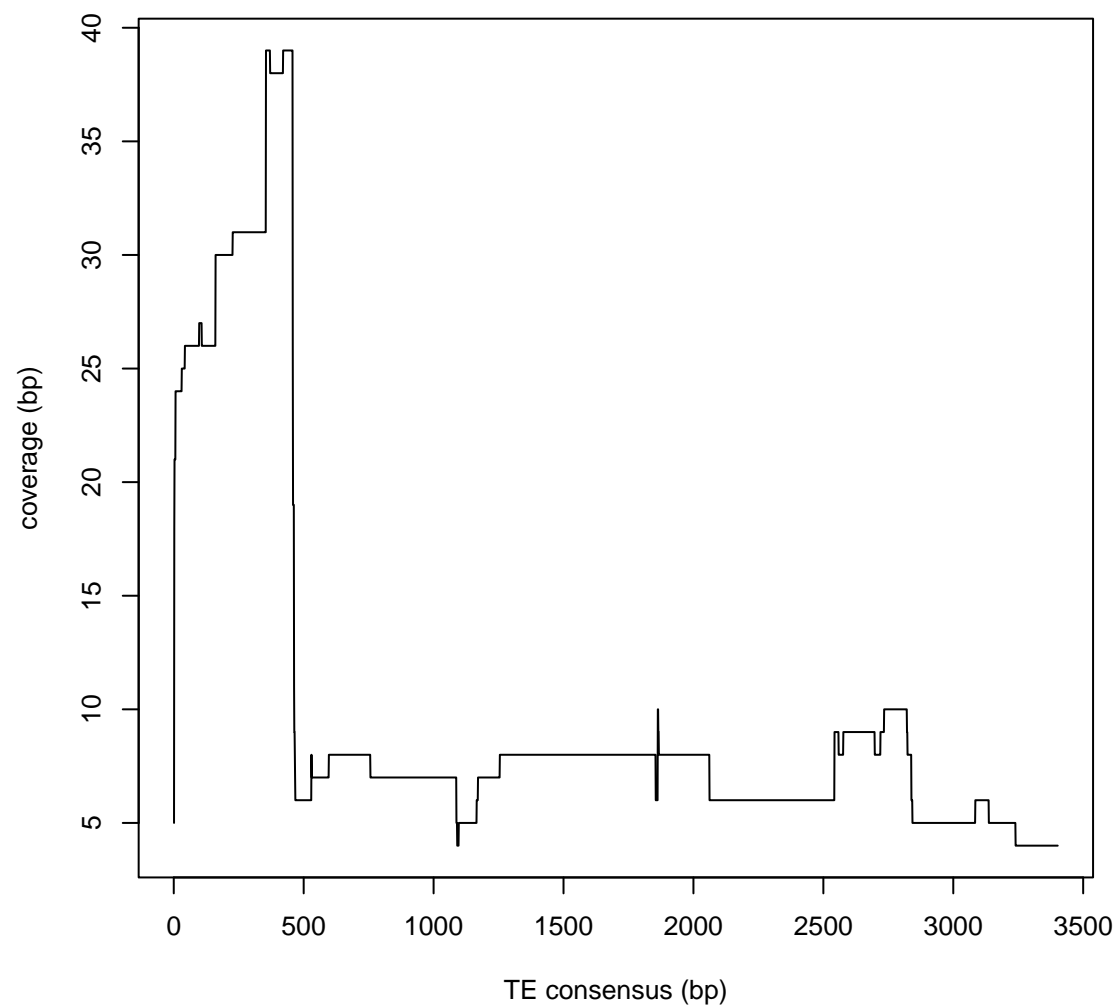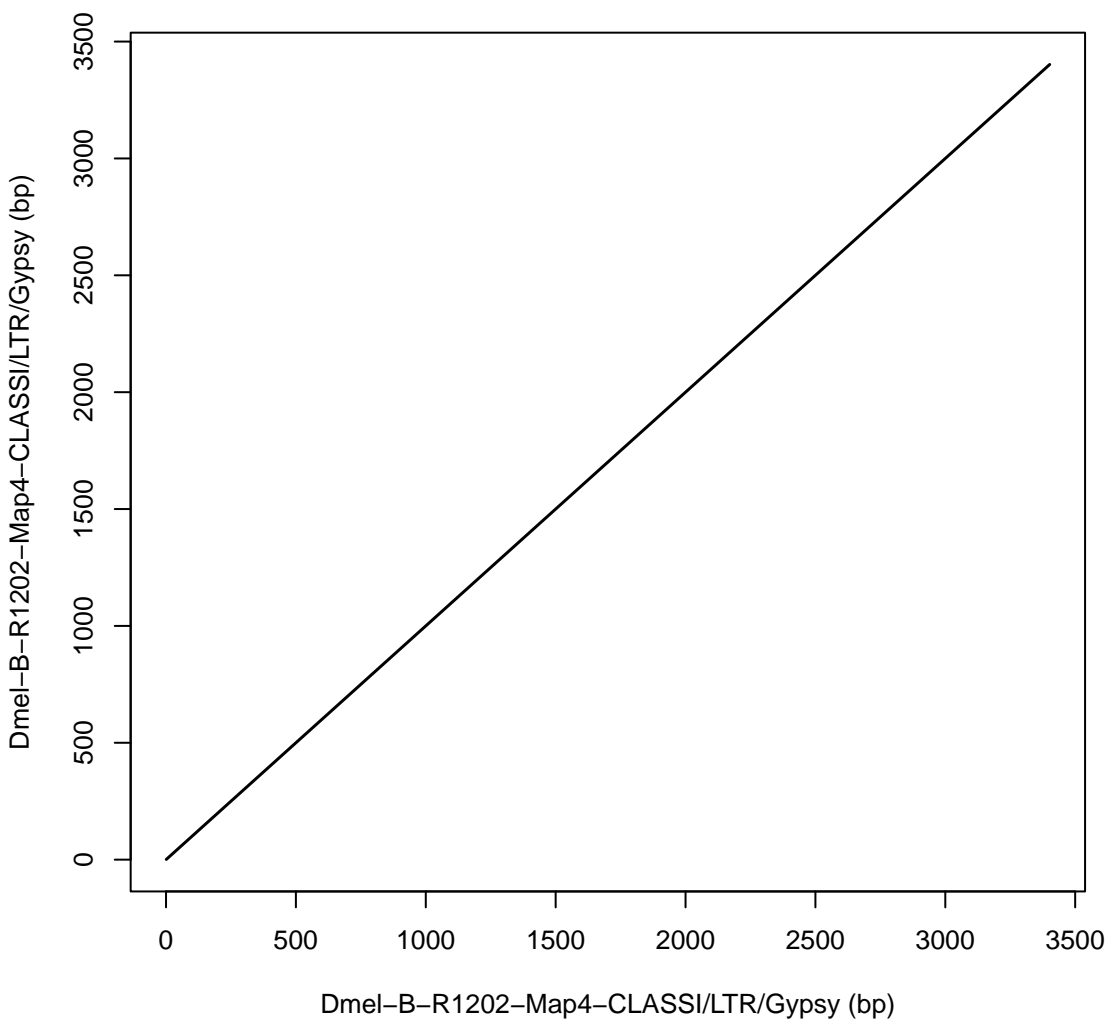**consensus size: 3402bp; fragments: 61; full length: 4 (>=3061.8bp)**

**TE consensus genomic coverage**

**TE consensus self dotplot (blastn)**

**TE consensus structure and protein hits**

YOYO_pol LTR/Gypsy

Gypsy−44_DWil_env LTR/Gypsy

Gypsy4_env LTR/Gypsy