TE: Dmel−B−R1039−Map9−TRIM/nan
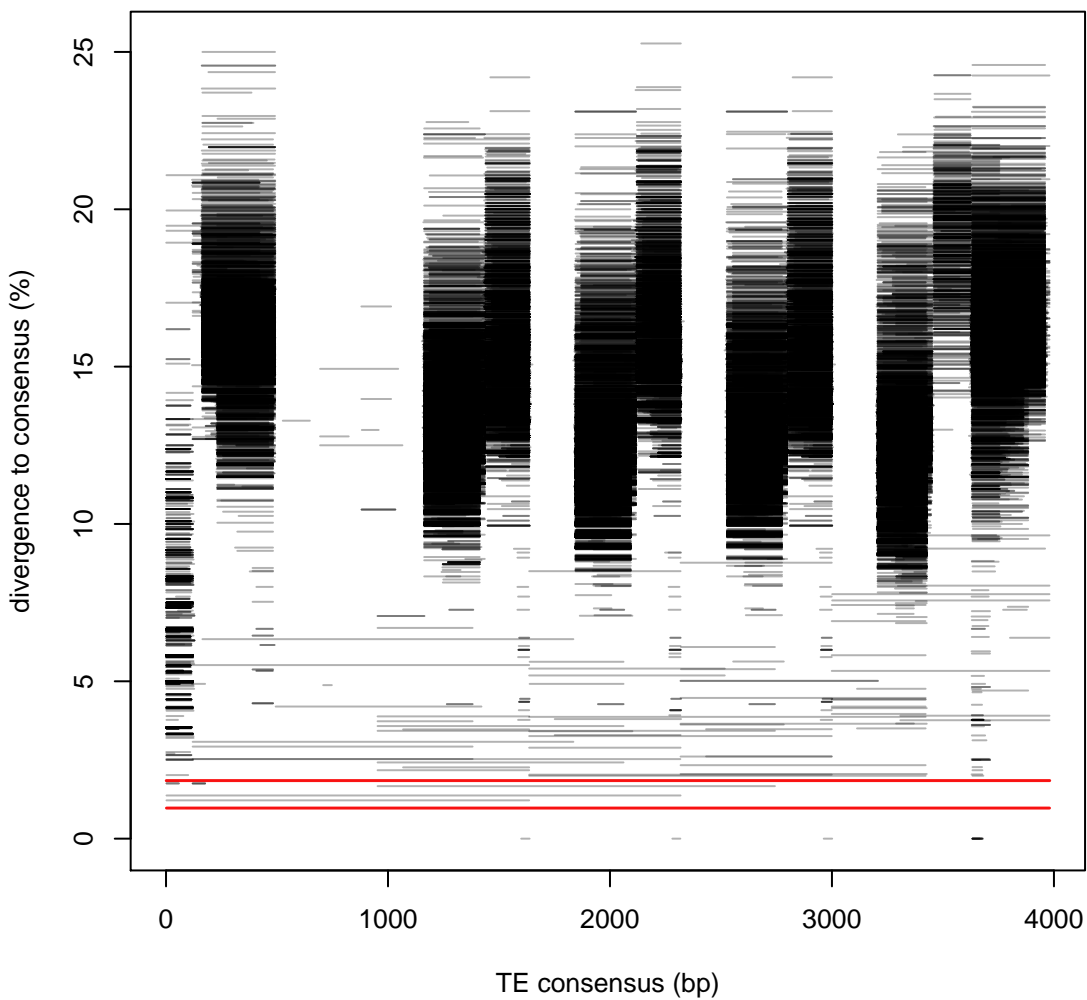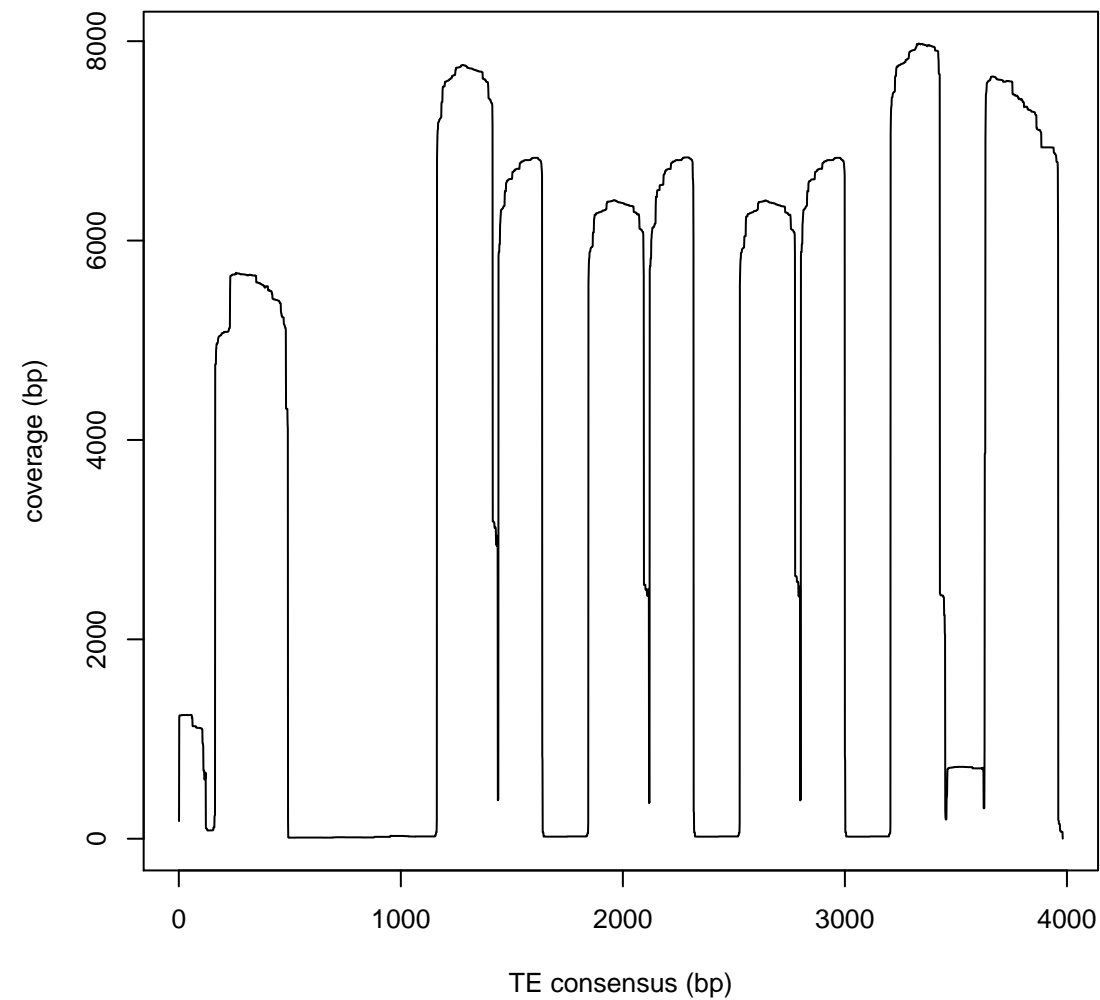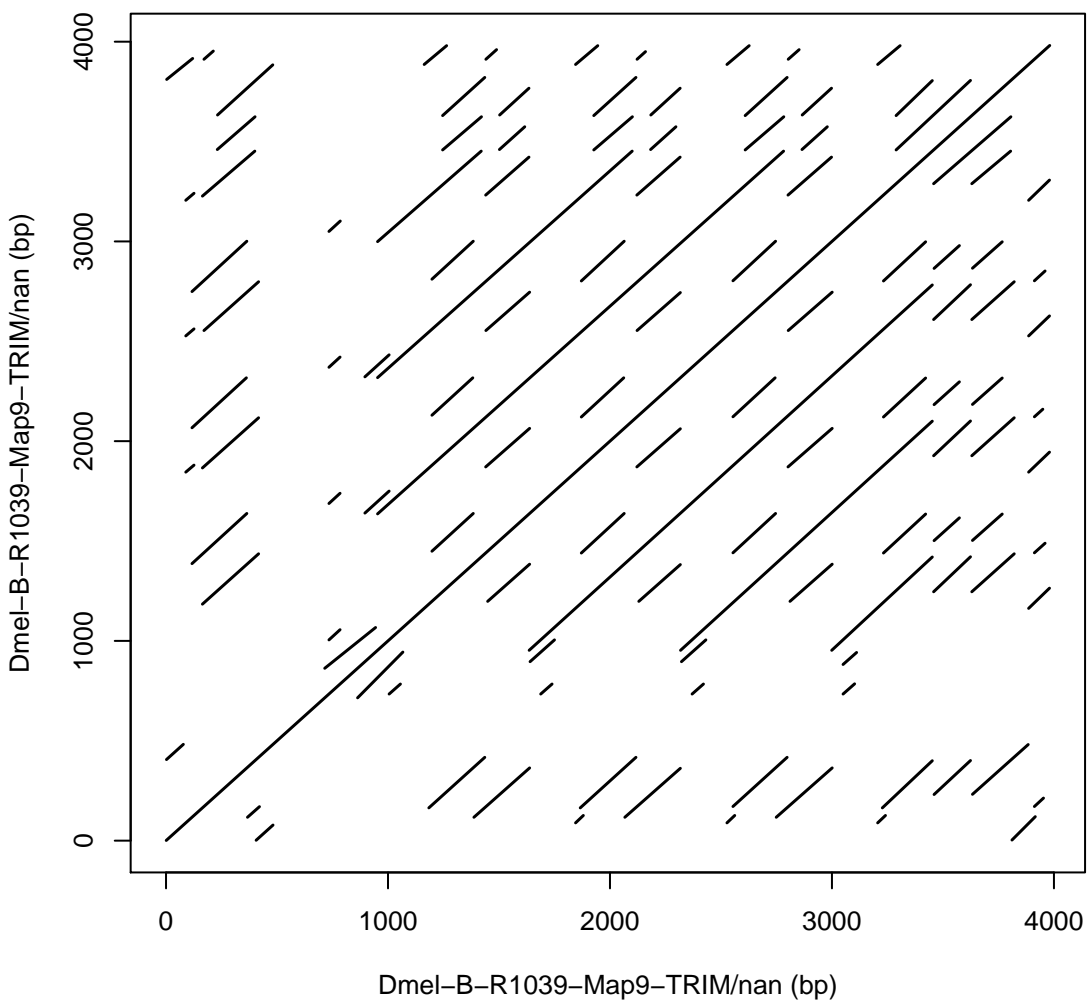consensus size: 3981bp; fragments: 63450; full length: 2 (>=3582.9bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

Gypsy−1_LG_env LTR/Gypsy