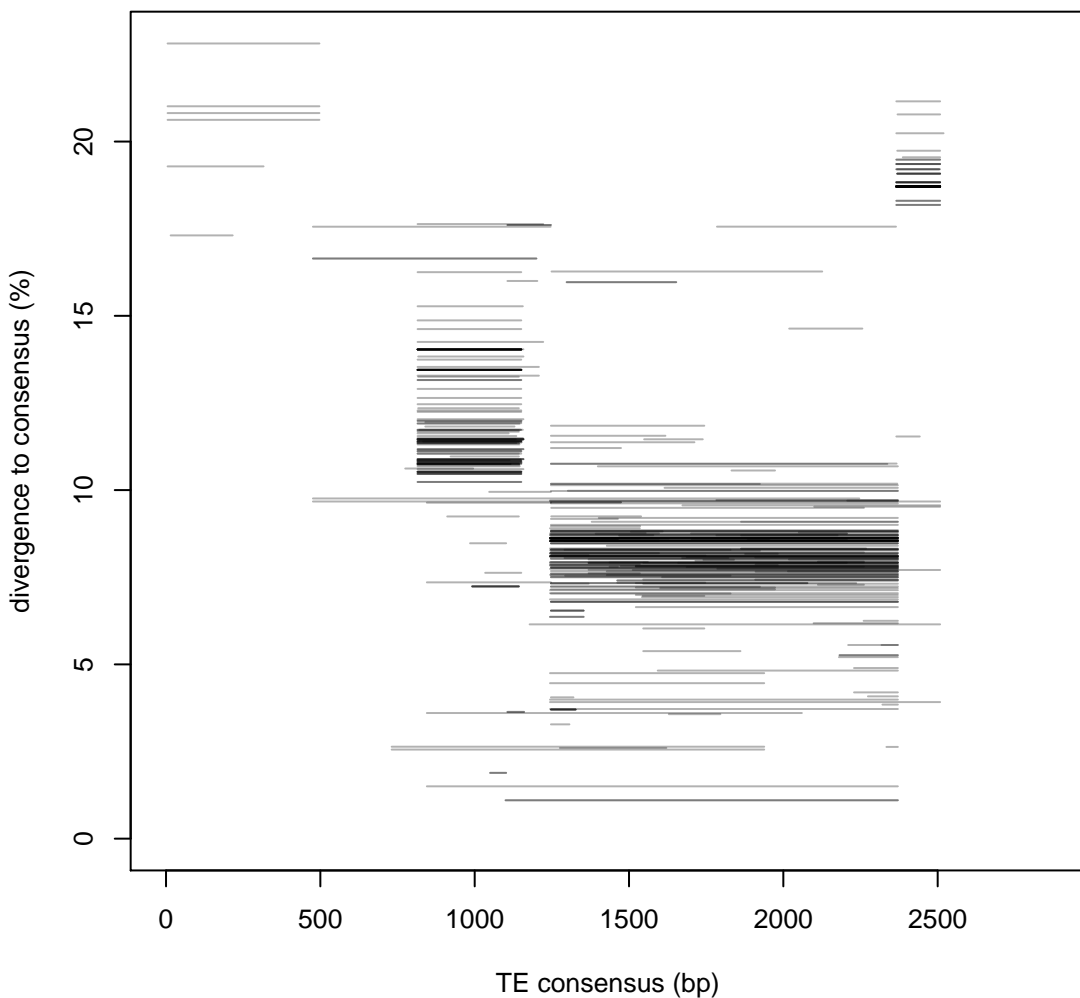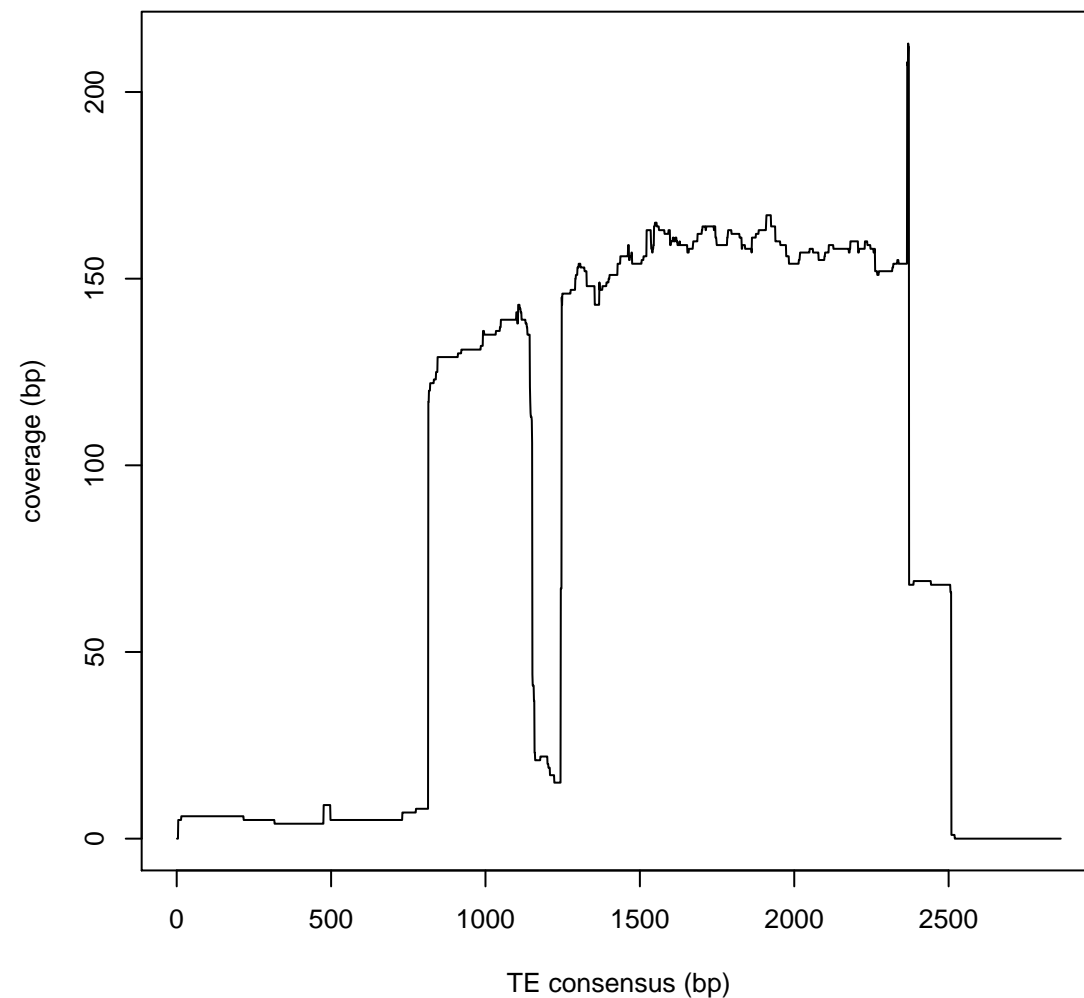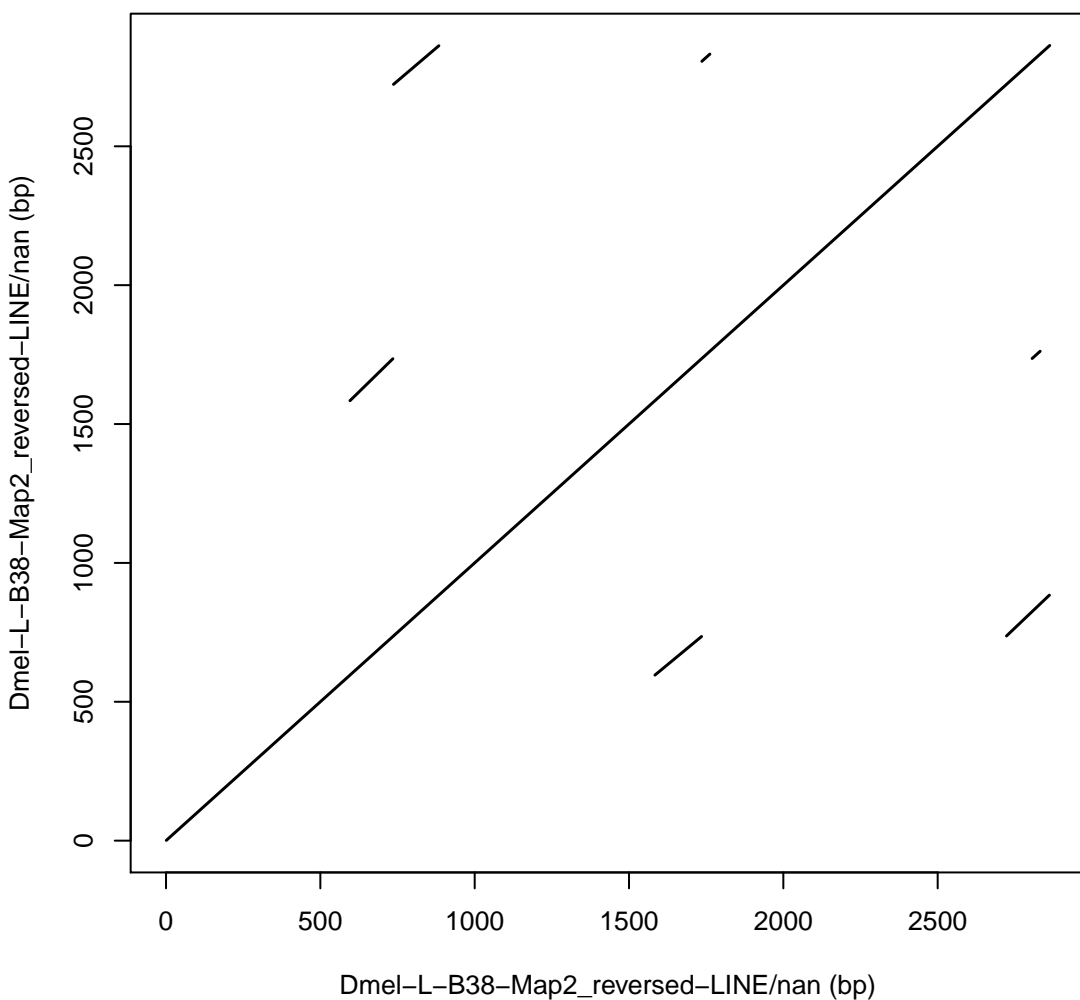**TE: Dmel−L−B38−Map2_reversed−LINE/nan**
consensus size: 2863bp; fragments: 448; full length: 0 (>=2576.7bp)

TE consensus (bp)

divergence to consensus (%)

TE consensus genomic coverage

coverage (bp)

TE consensus (bp)

TE consensus self dotplot (blastn)

Dmel−L−B38−Map2_reversed−LINE/nan (bp)

Dmel−L−B38−Map2_reversed−LINE/nan (bp)

TE consensus structure and protein hits

R1_DSe_gag LINE/R1
R1_DM_pol LINE/R1
R1_DM_pol LINE/R1

Dmel−L−B38−Map2_reversed−LINE/nan (bp)