TE: DmeI-B-R672-Map3_s_2_s_3-CLASSI/LINE/nan
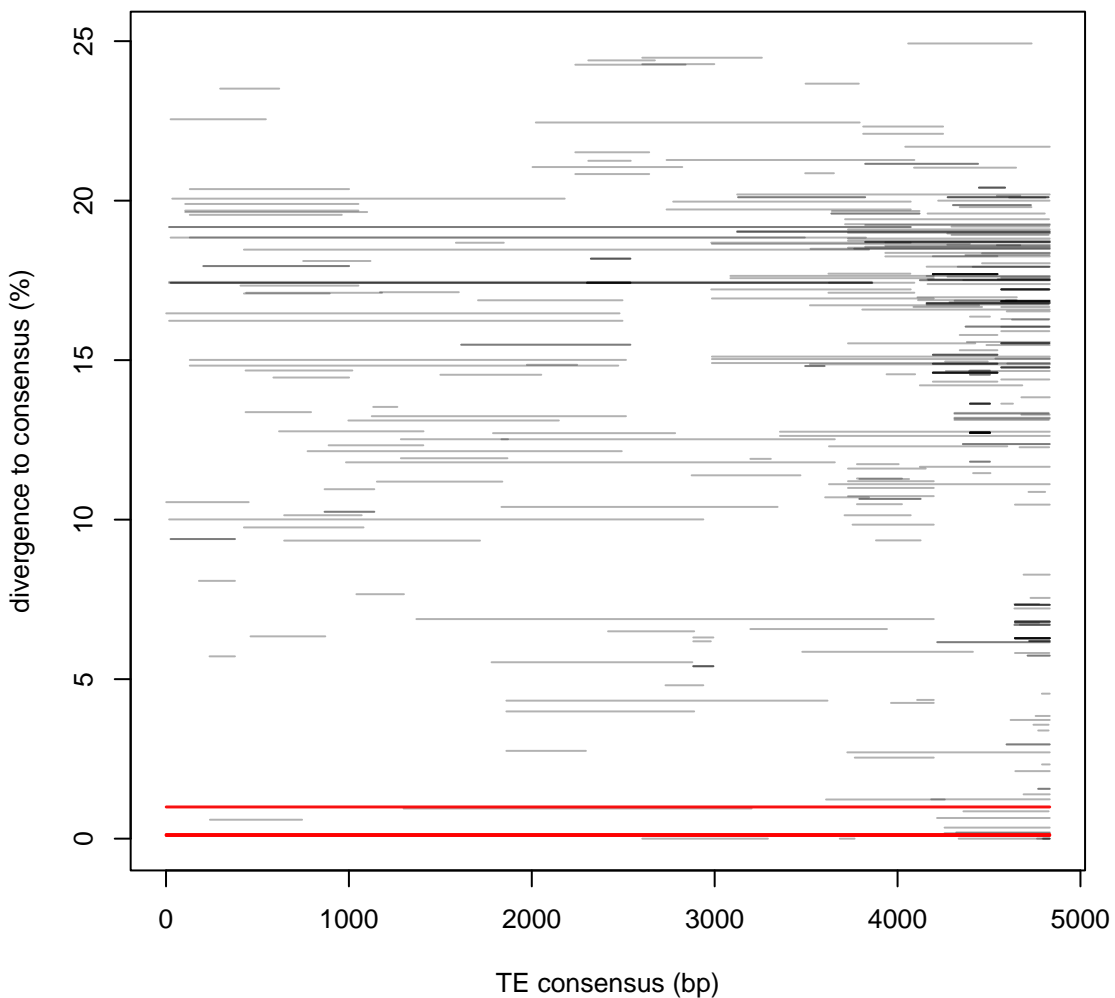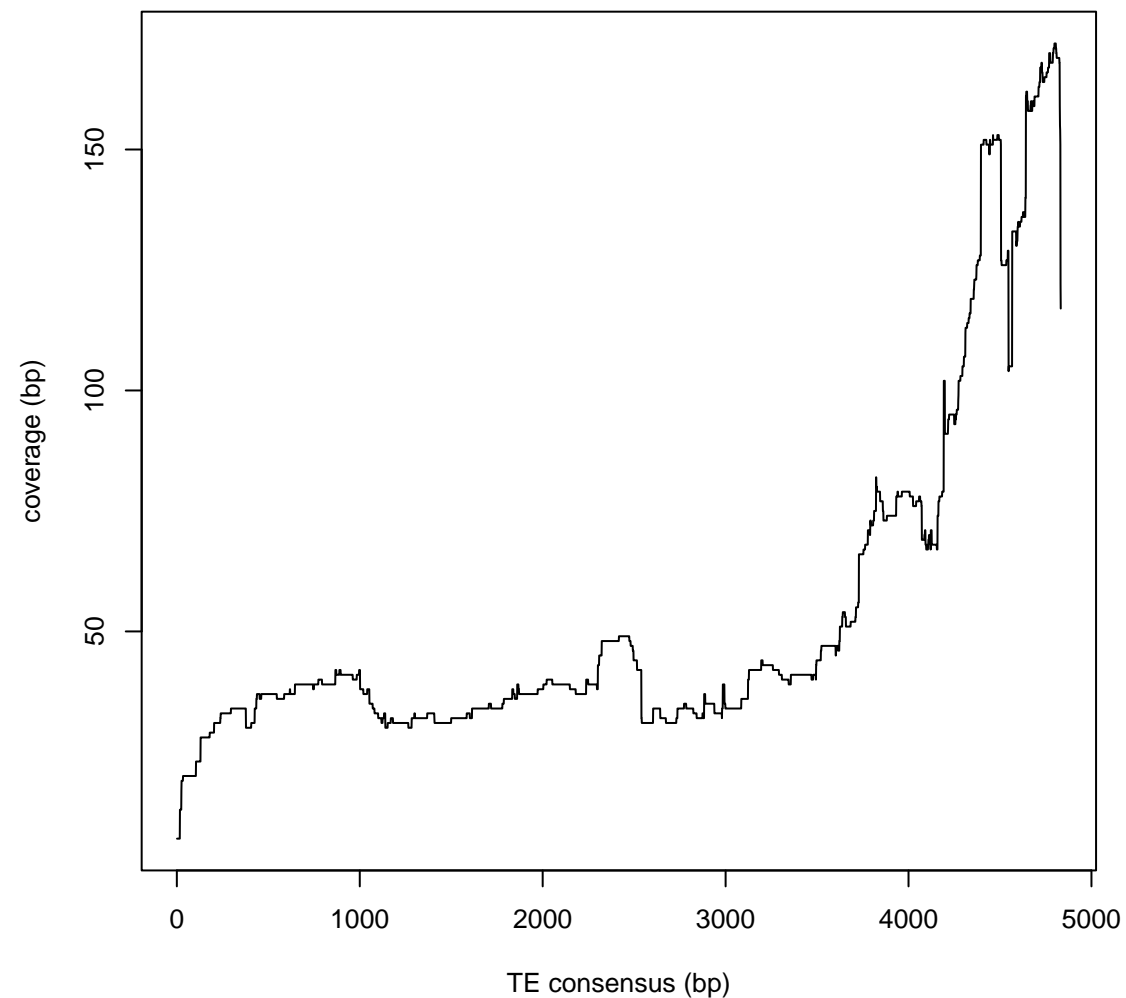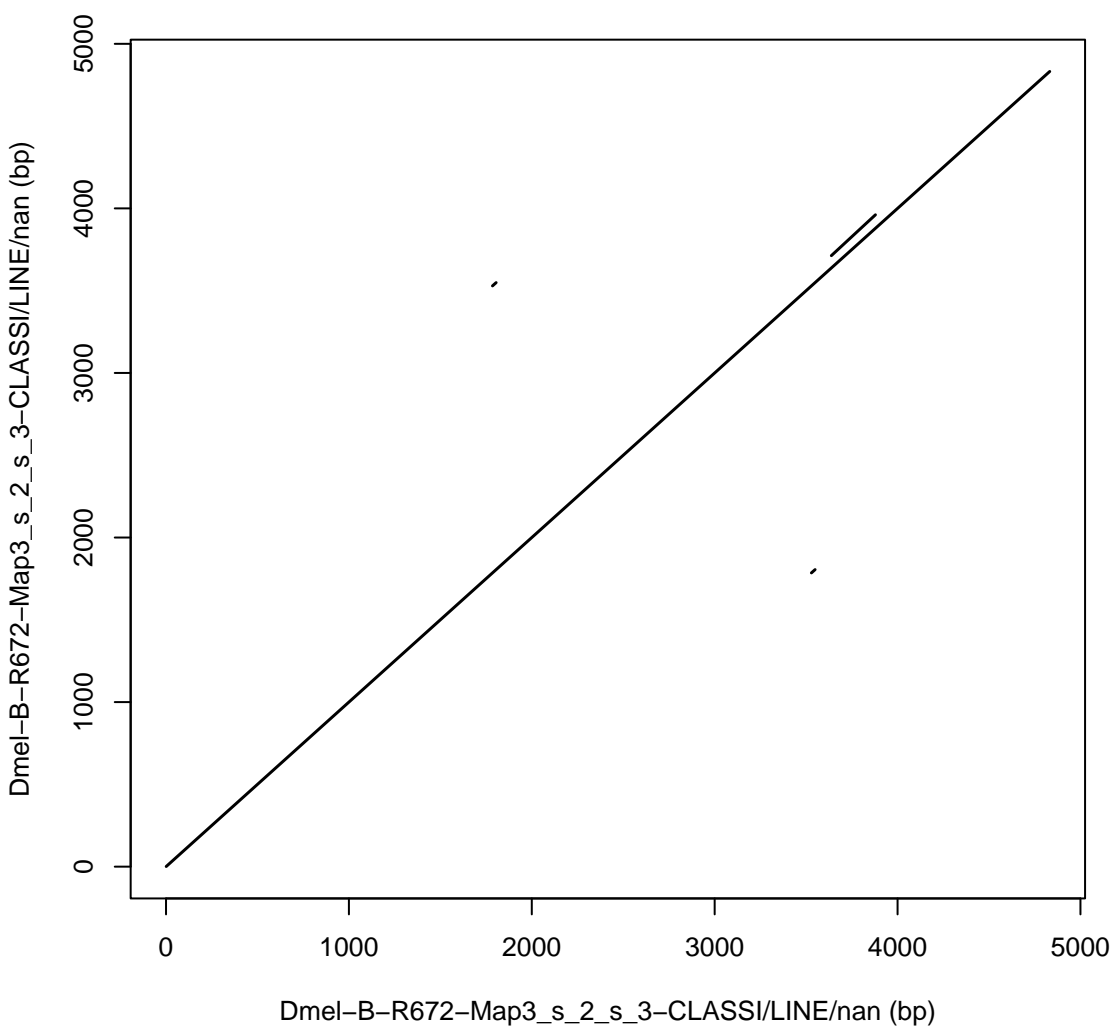**consensus size: 4832bp; fragments: 390; full length: 5 (>=4348.8bp)**

divergence to consensus (%)

TE consensus (bp)

TE consensus genomic coverage

coverage (bp)

TE consensus (bp)

TE consensus self dotplot (blastn)

Dmel-B-R672-Map3_s_2_s_3-CLASSI/LINE/nan (bp)

Dmel-B-R672-Map3_s_2_s_3-CLASSI/LINE/nan (bp)

TE consensus structure and protein hits

HETA_gag LINE/I-Jockey

Dmel-B-R672-Map3_s_2_s_3-CLASSI/LINE/nan (bp)