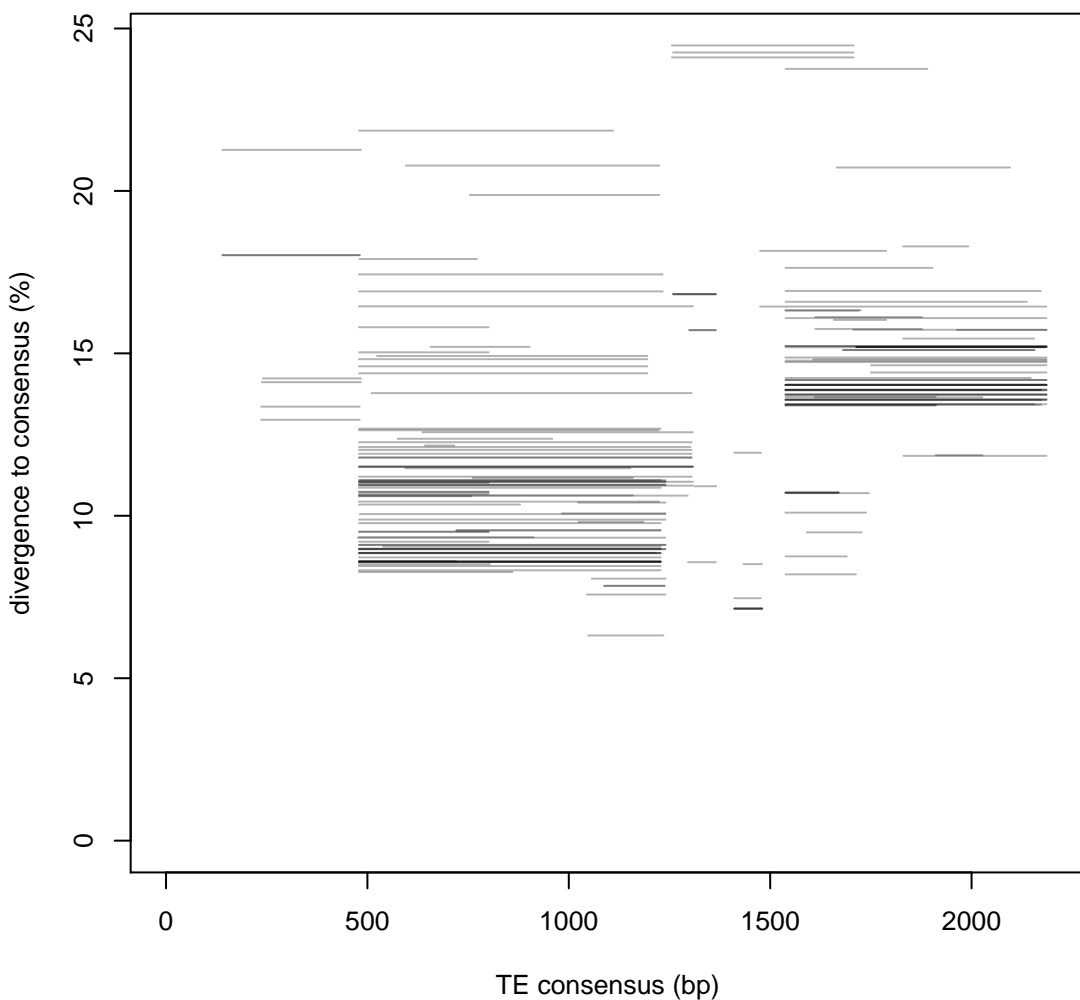
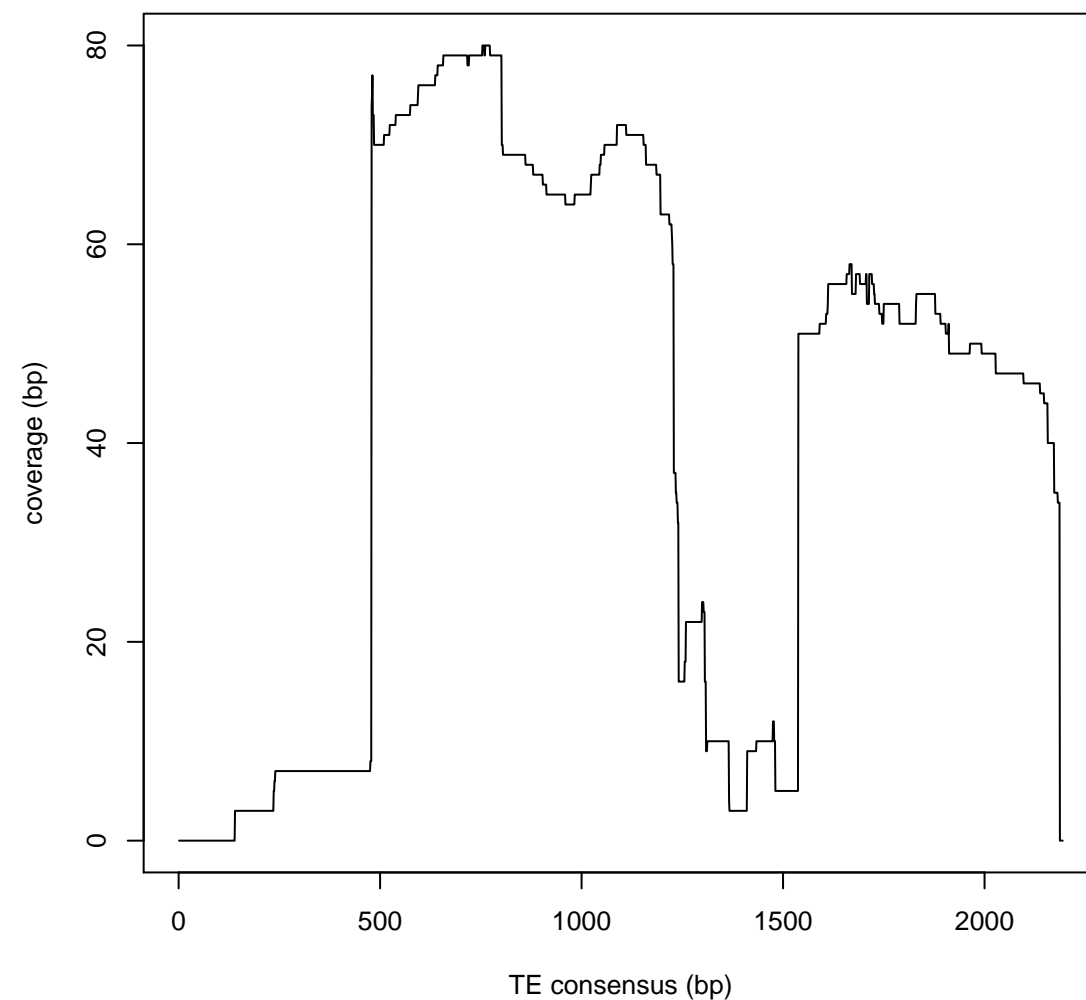


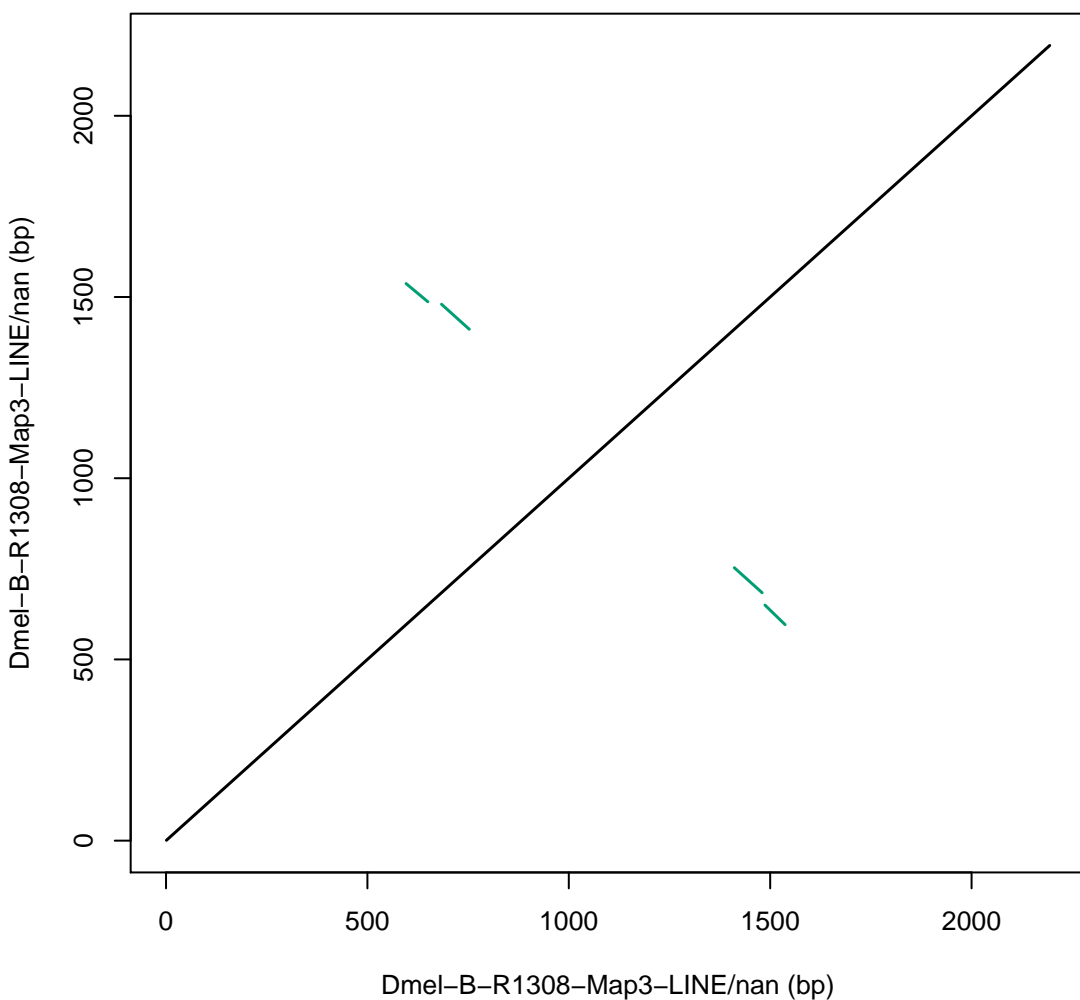
TE: Dmel-B-R1308-Map3-LINE/nan
consensus size: 2194bp; fragments: 184; full length: 0 (>=1974.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

