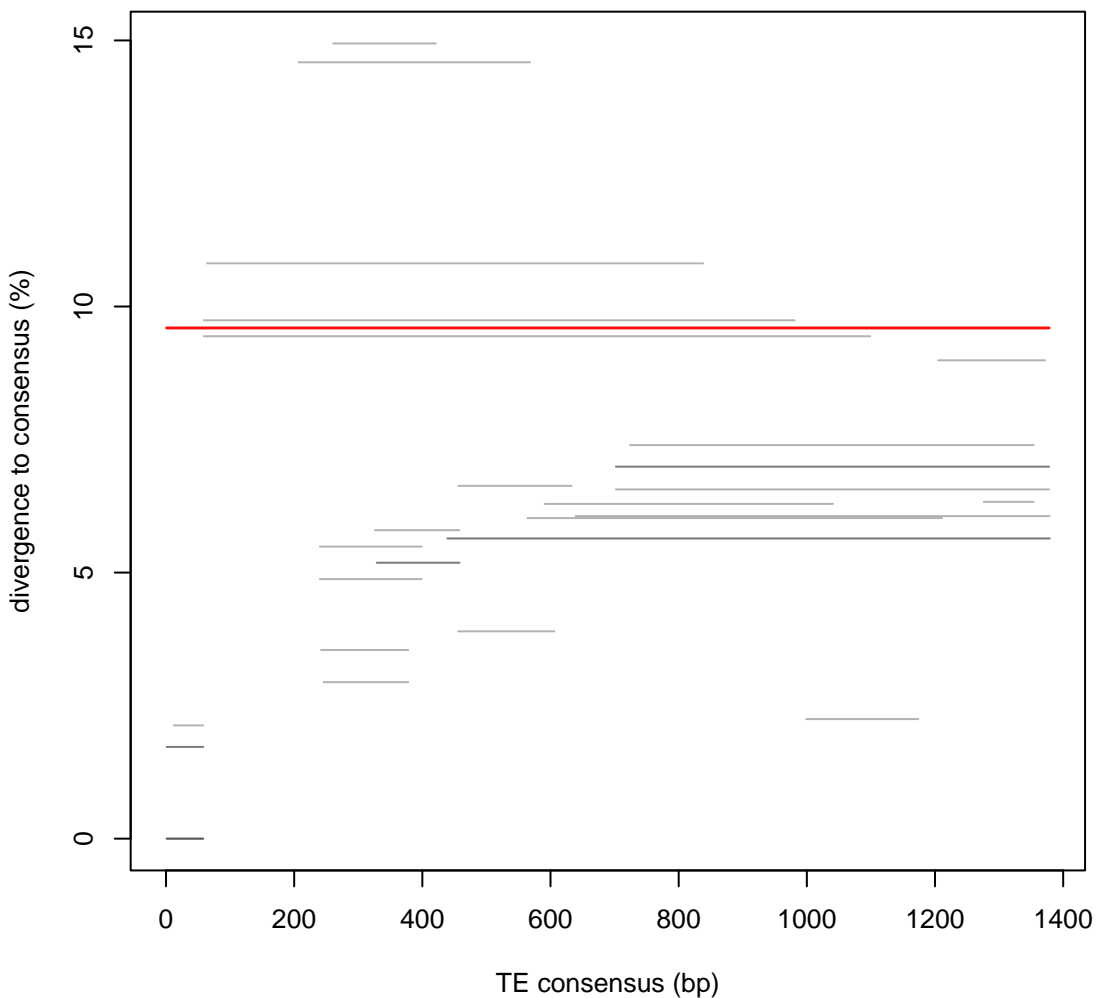
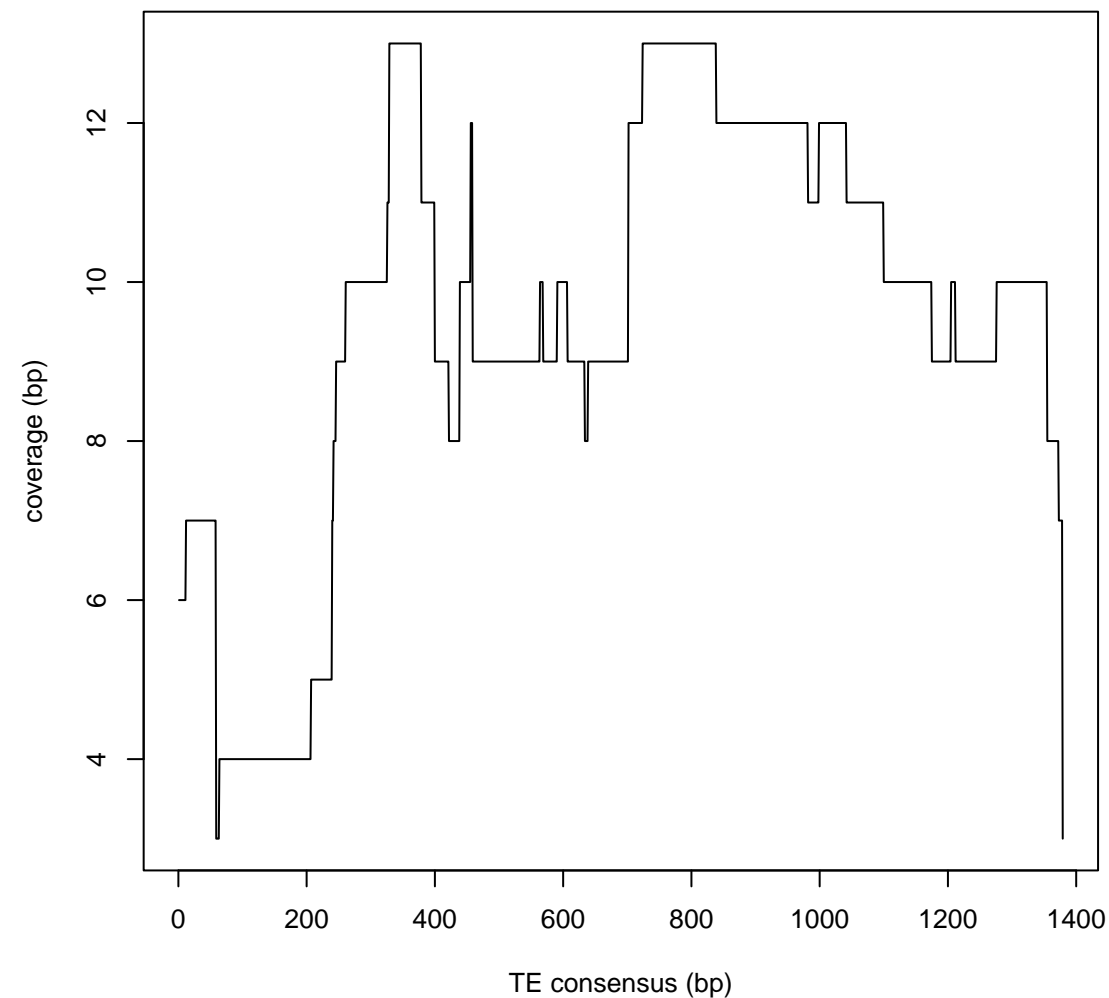


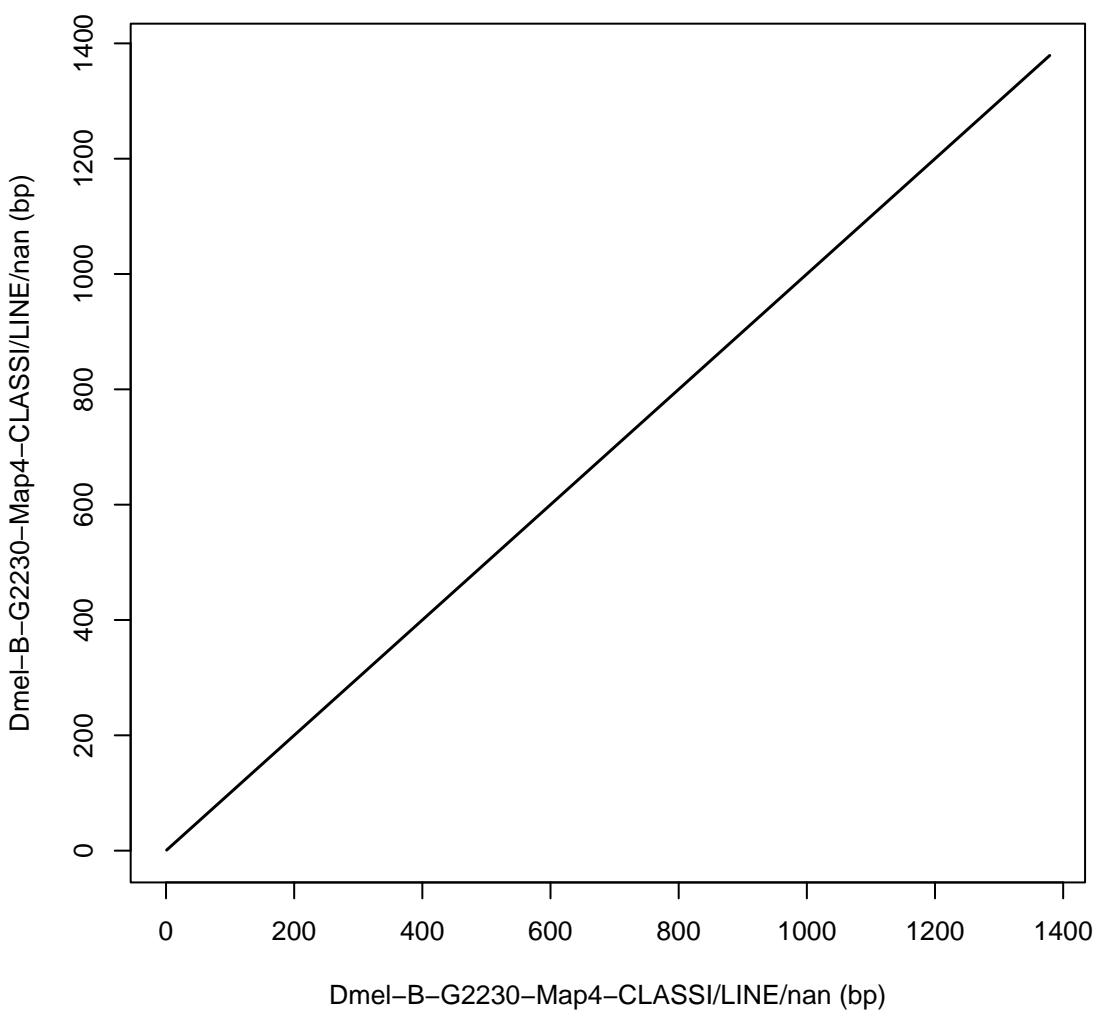
TE: Dmel-B-G2230-Map4-CLASSI/LINE/nan
consensus size: 1379bp; fragments: 33; full length: 1 (>=1241.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

