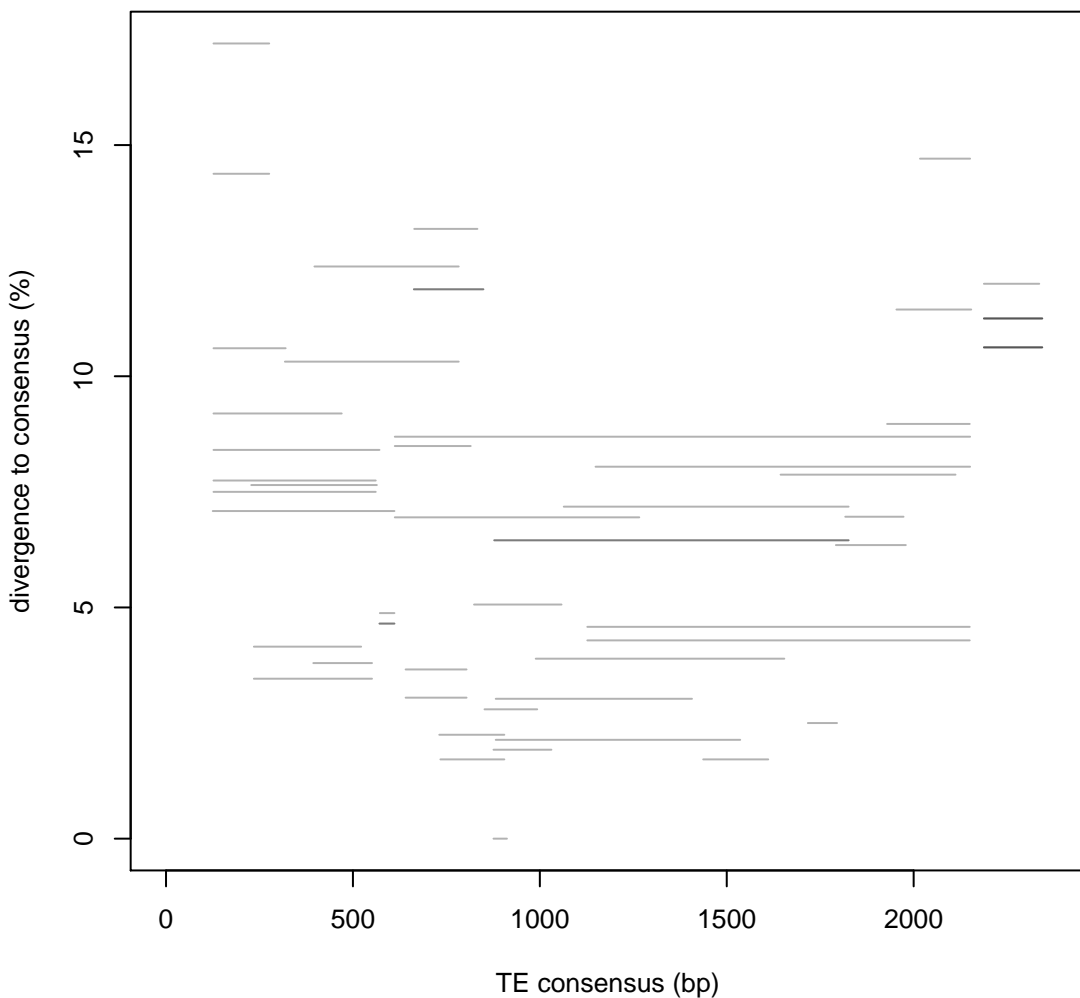
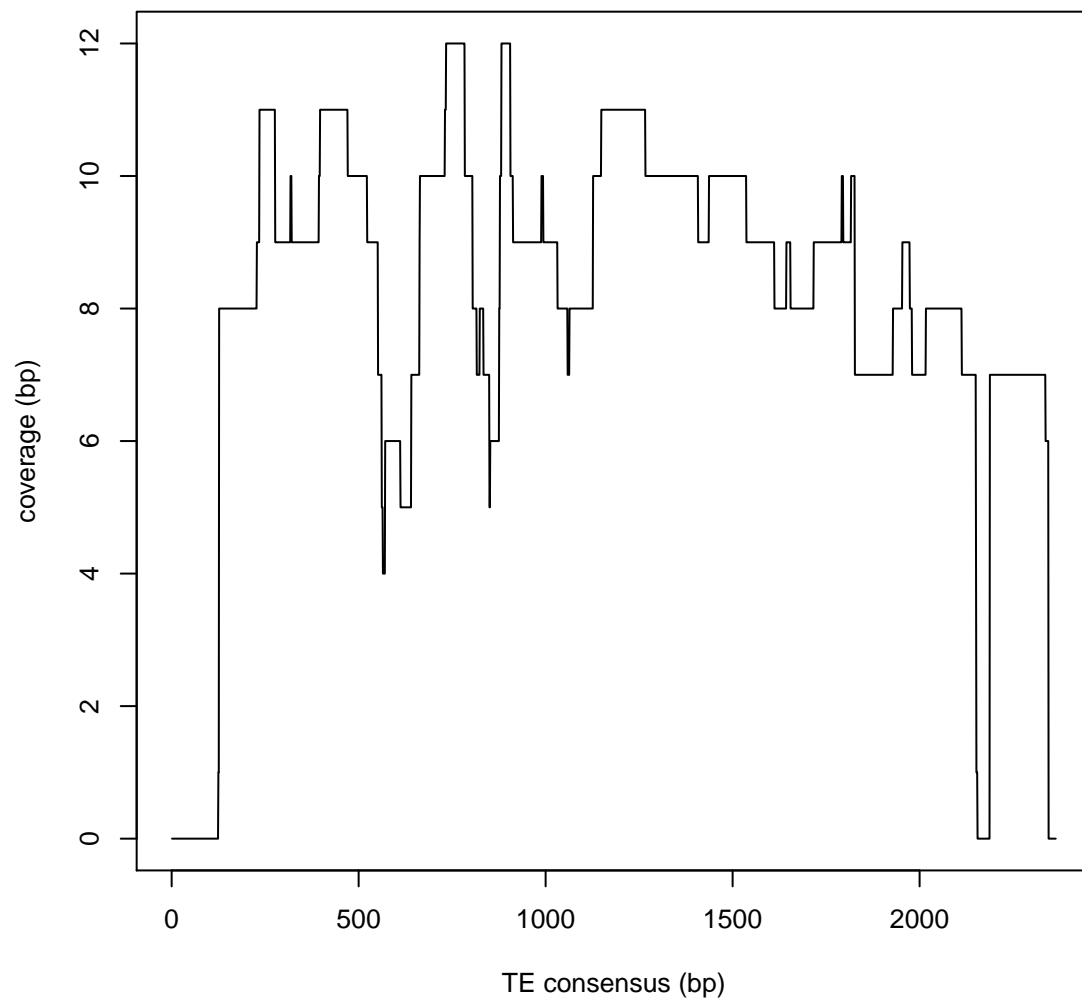


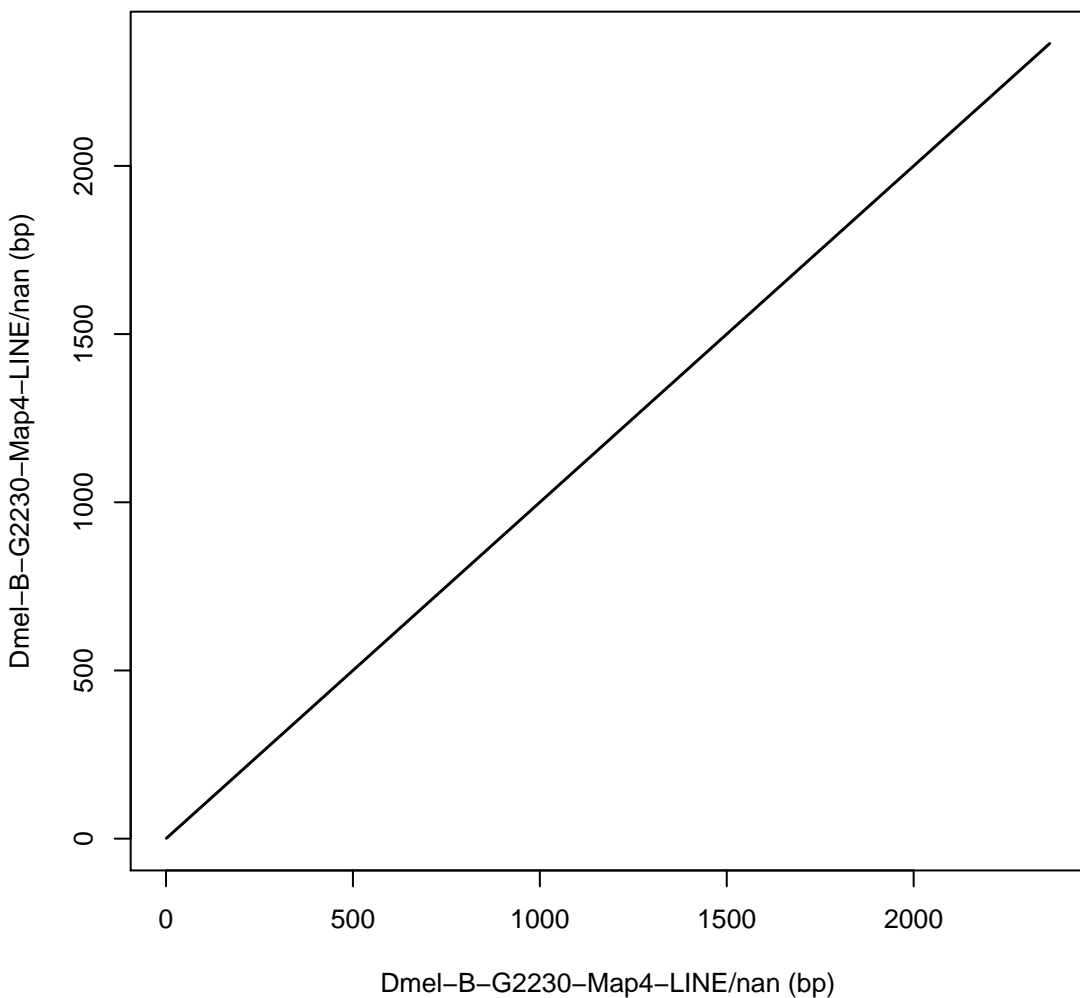
TE: Dmel-B-G2230-Map4-LINE/nan
consensus size: 2364bp; fragments: 55; full length: 0 (>=2127.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

