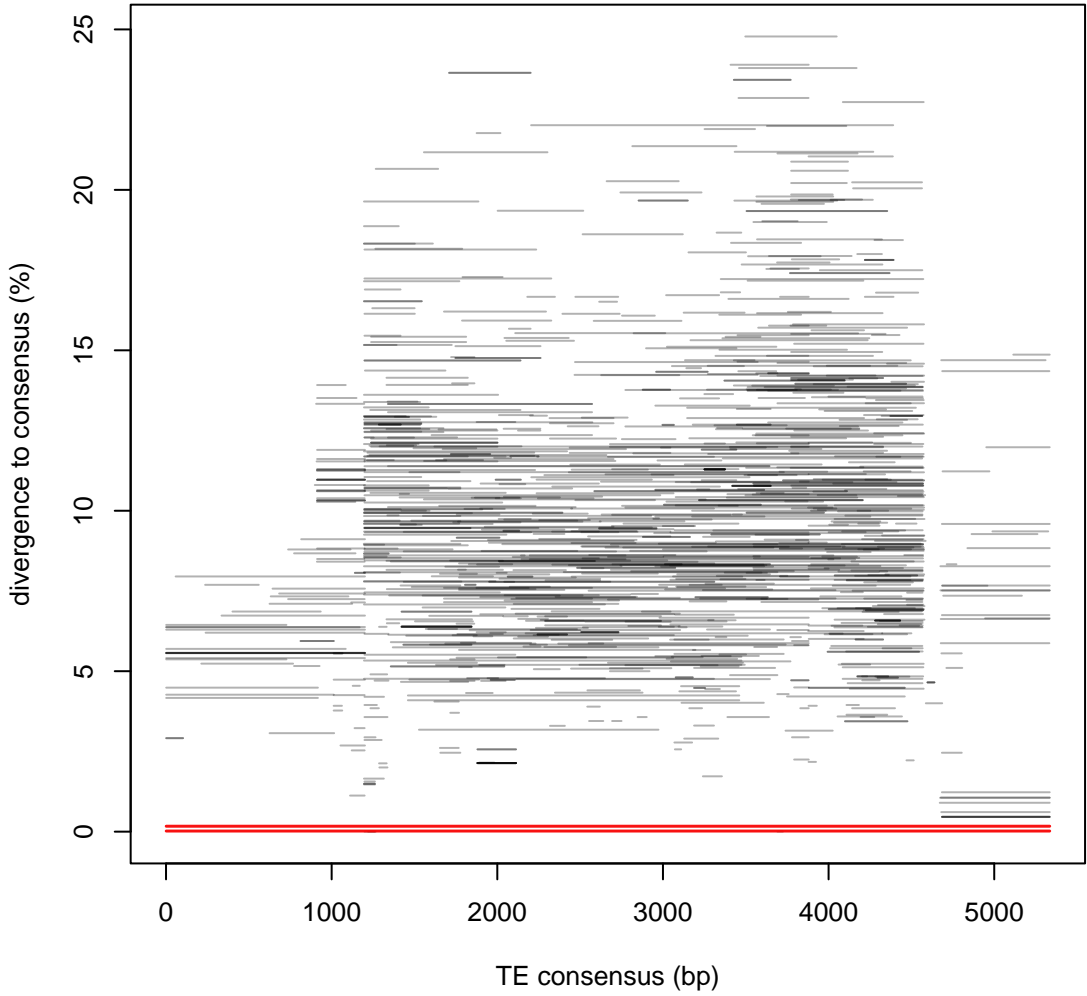
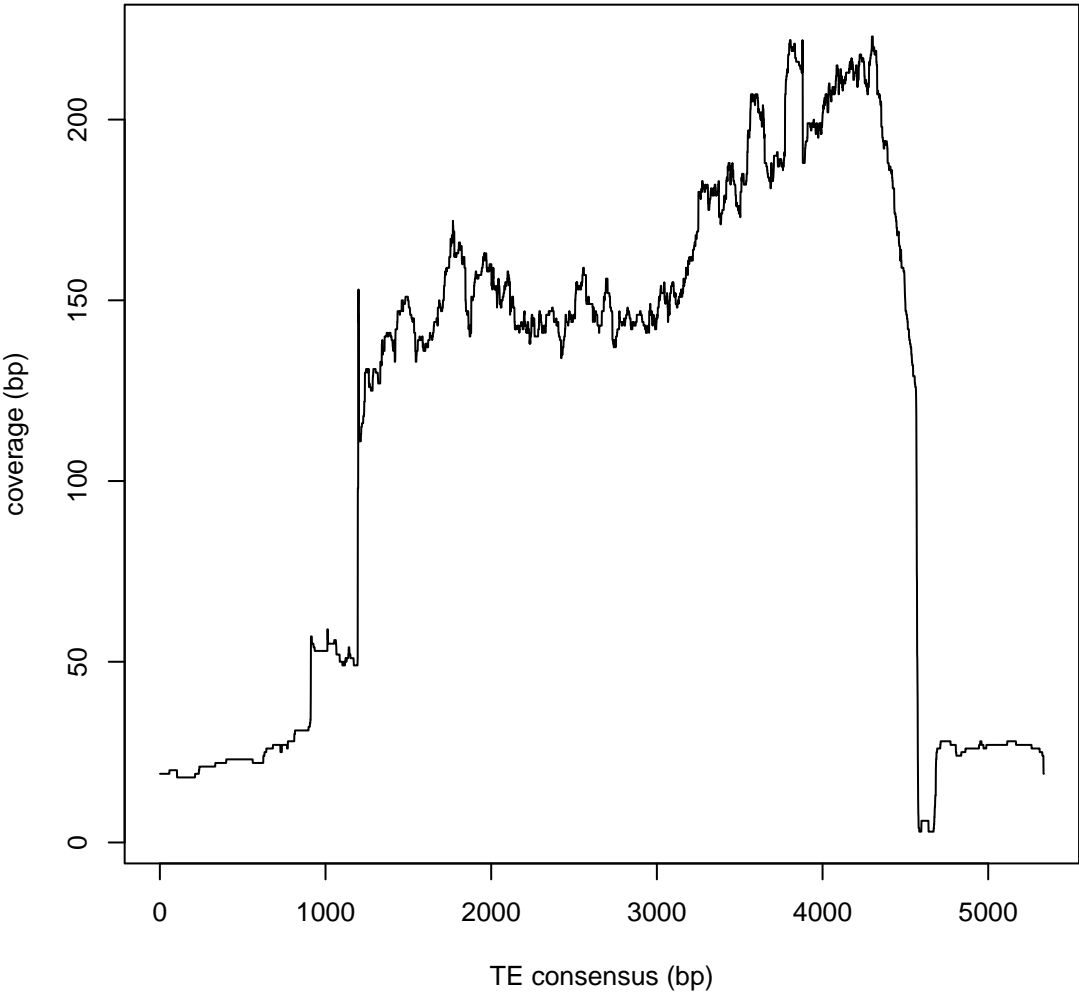


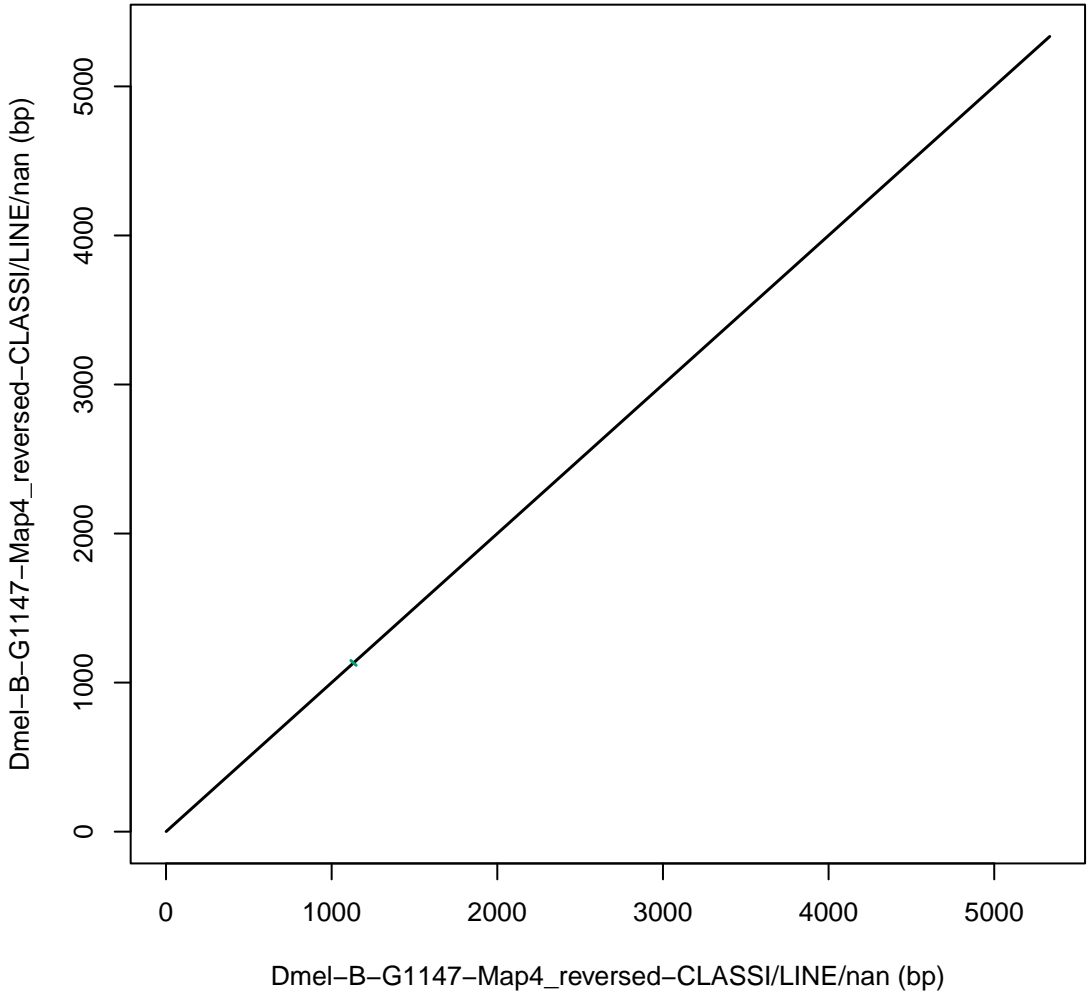
TE: Dmel-B-G1147-Map4_reversed-CLASSI/LINE/nan
consensus size: 5335bp; fragments: 1097; full length: 2 (>=4801.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

