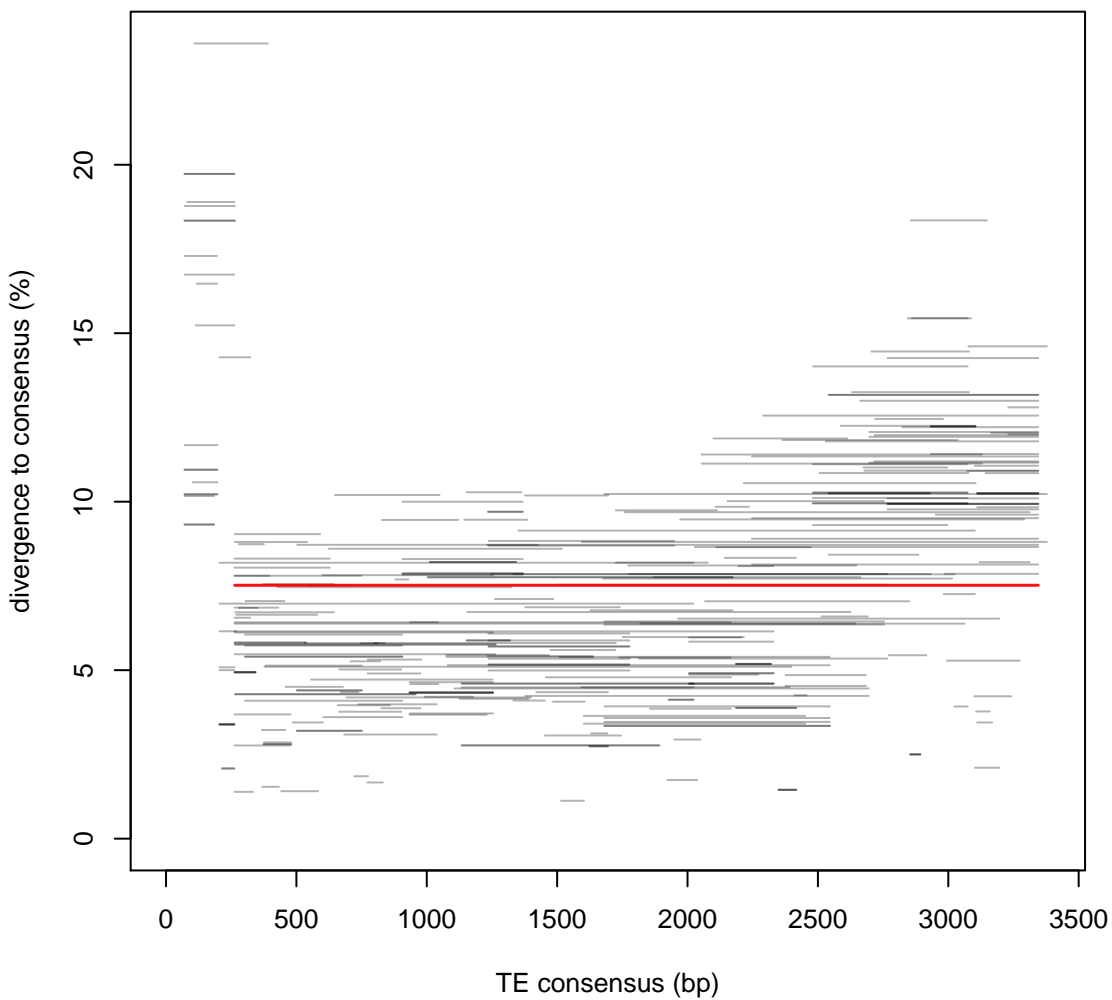
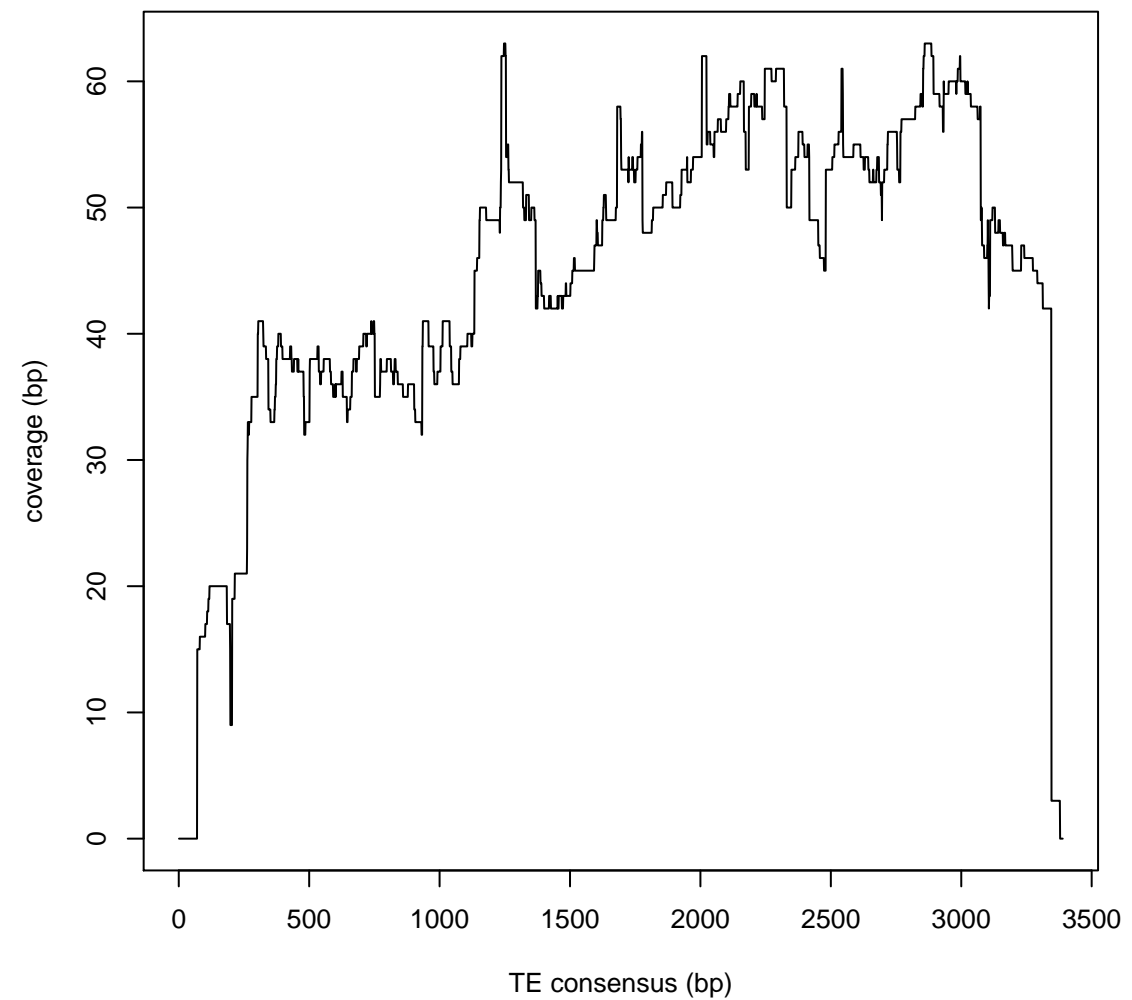


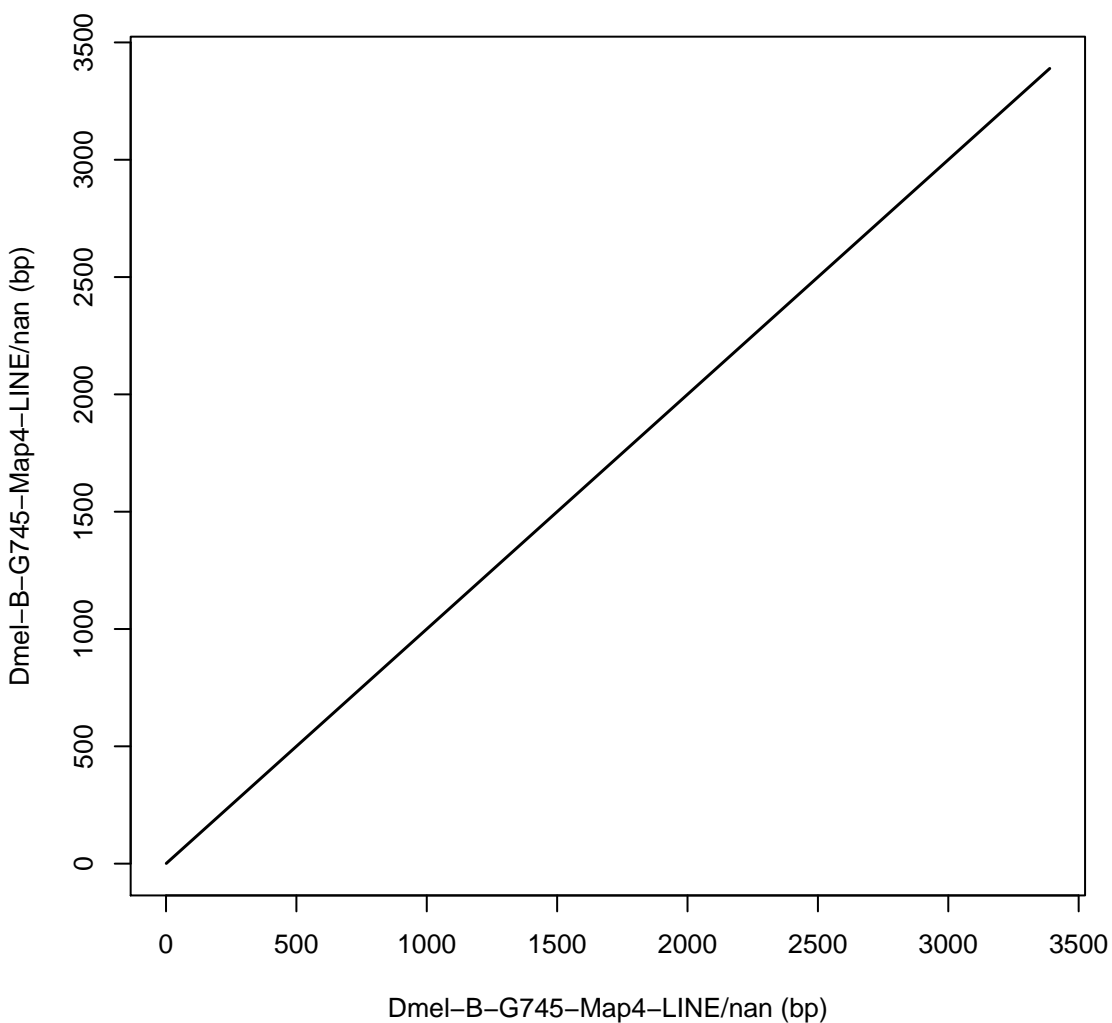
TE: Dmel-B-G745-Map4-LINE/nan
consensus size: 3389bp; fragments: 312; full length: 1 (>=3050.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

