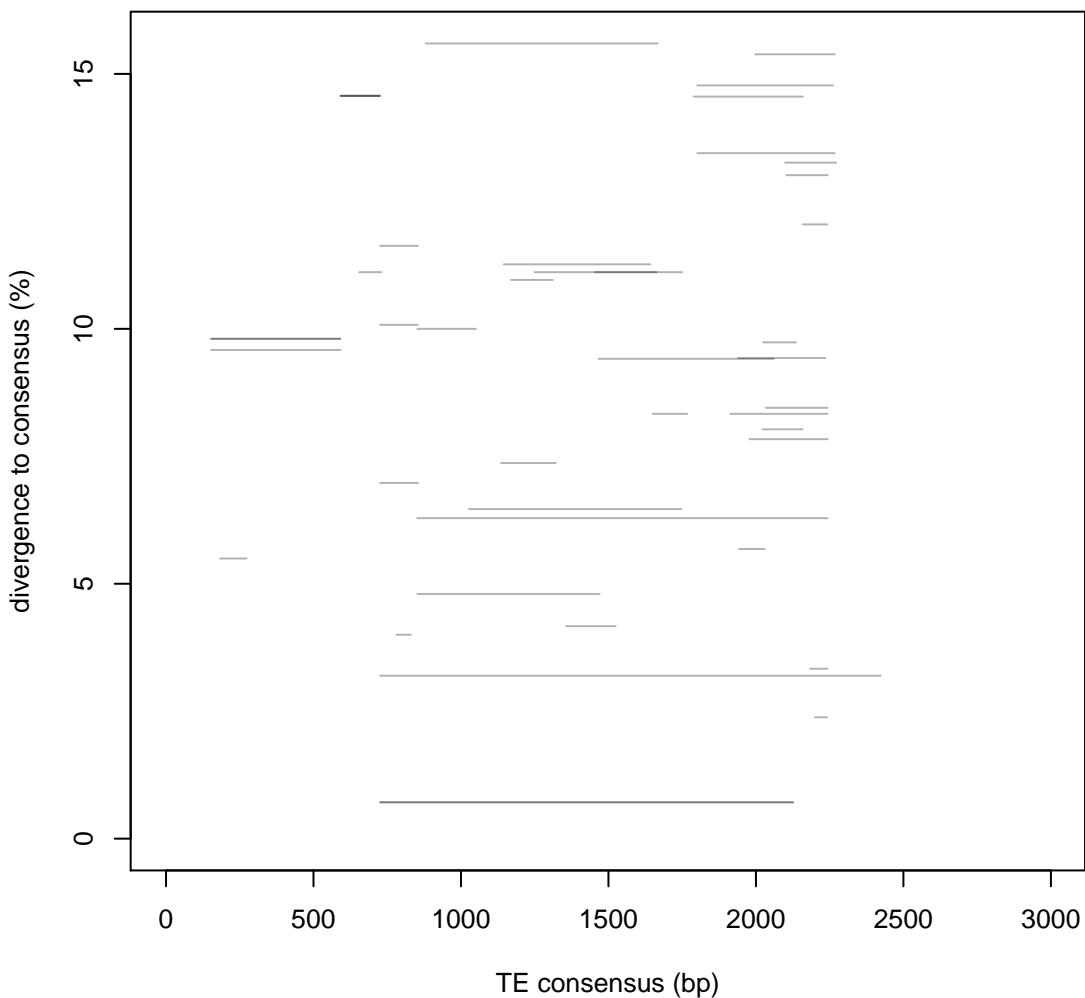
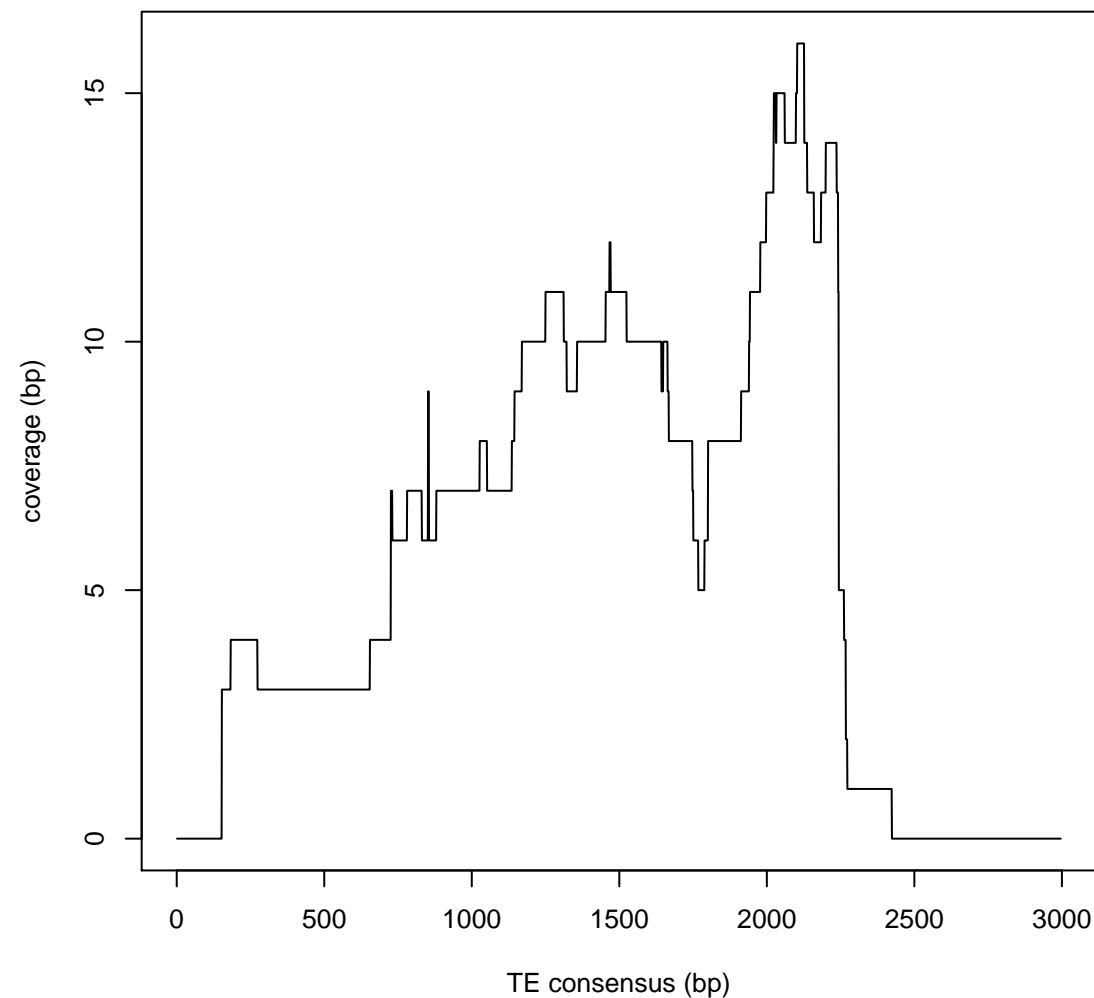


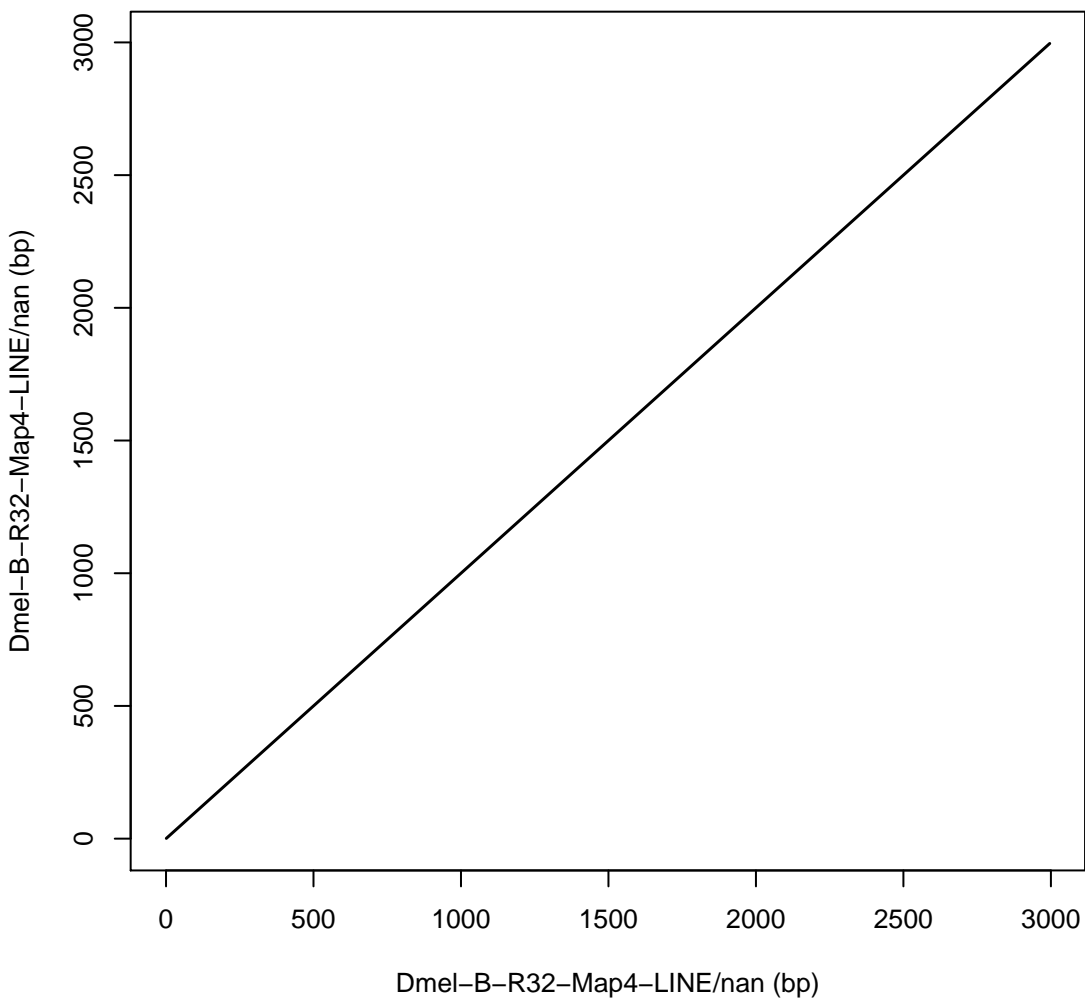
TE: Dmel-B-R32-Map4-LINE/nan
consensus size: 2996bp; fragments: 44; full length: 0 (>=2696.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

