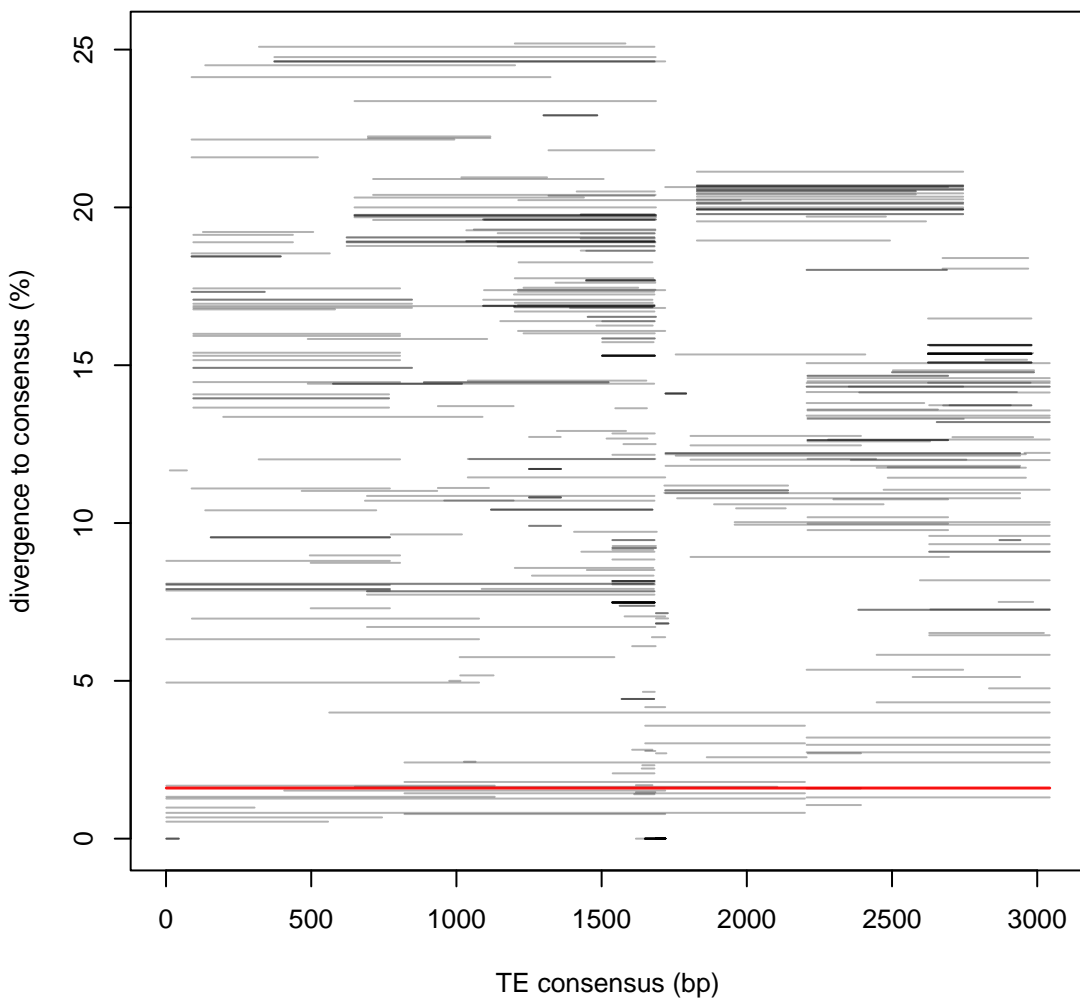
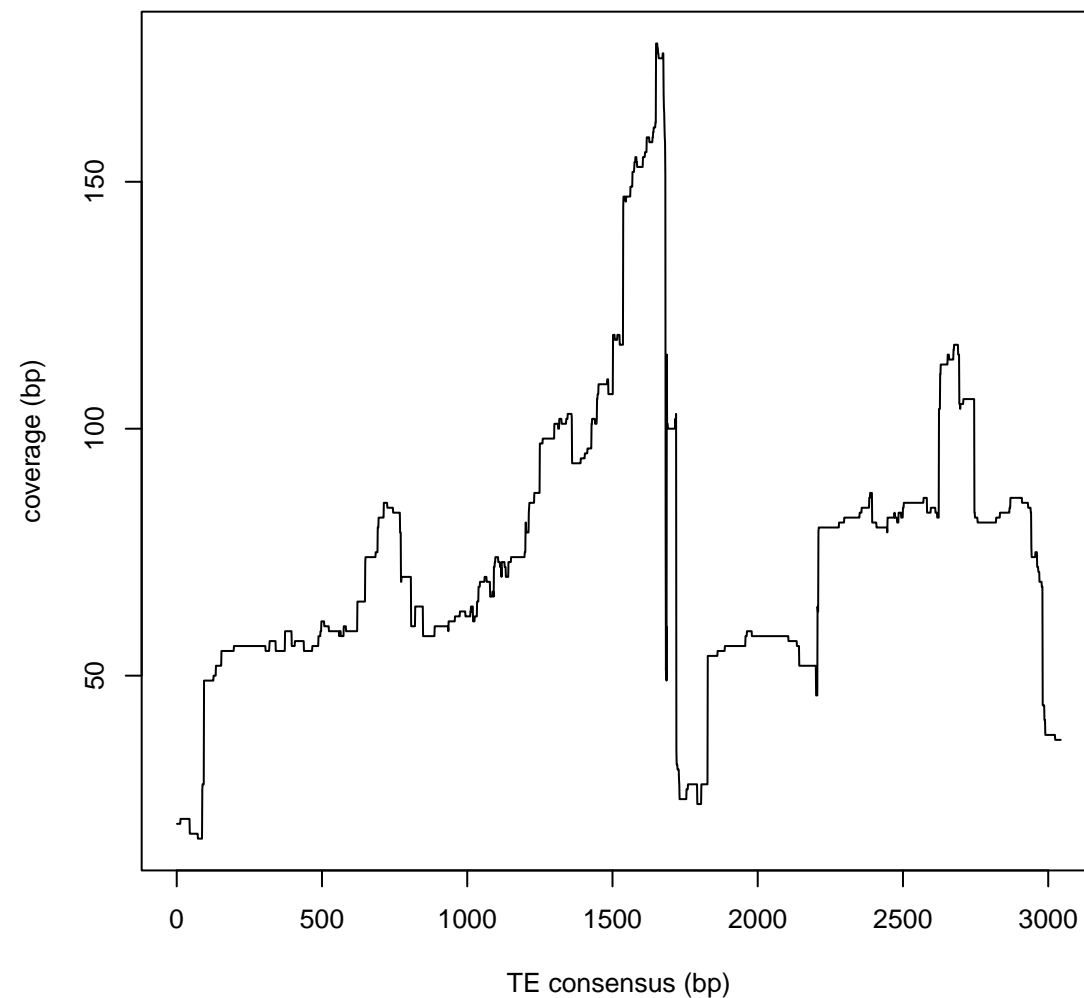


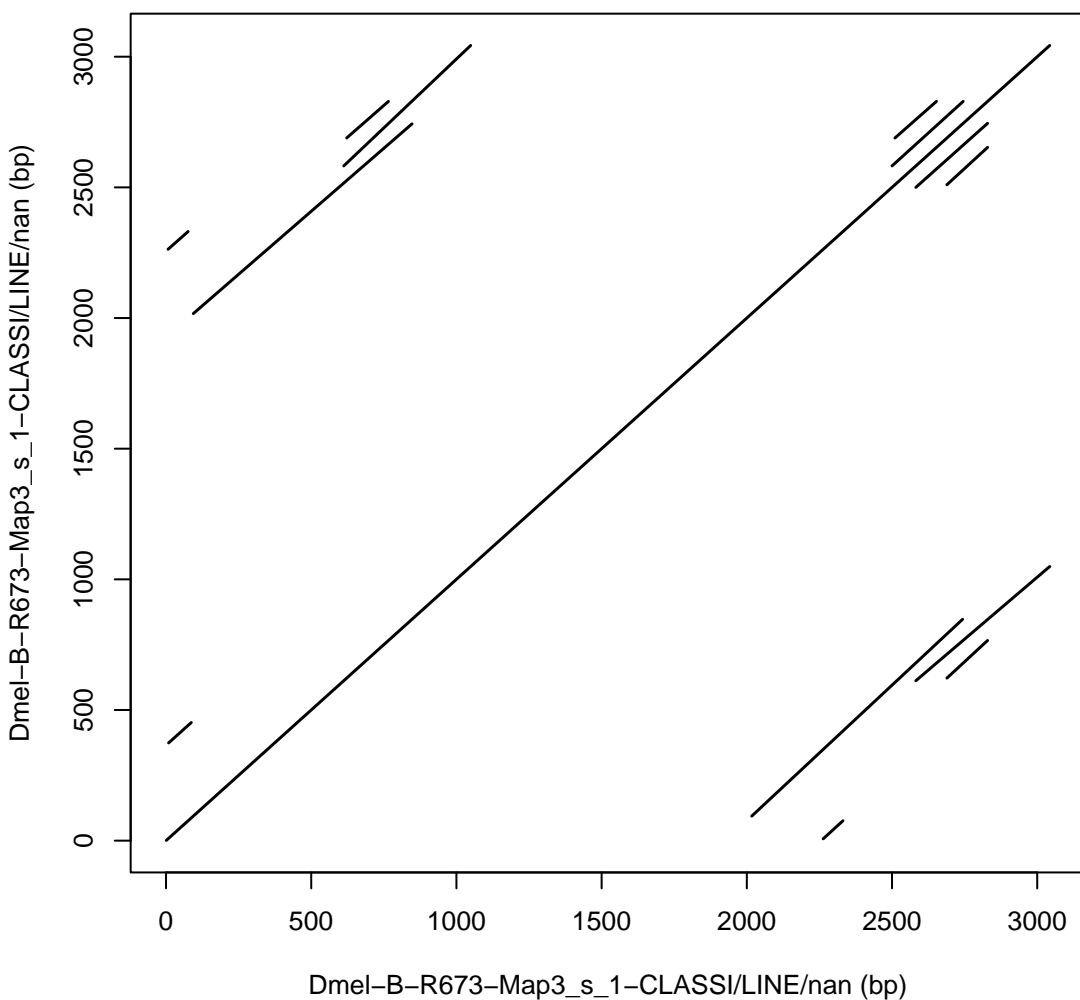
TE: Dmel-B-R673-Map3_s_1-CLASSI/LINE/nan
consensus size: 3043bp; fragments: 502; full length: 1 (>=2738.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

