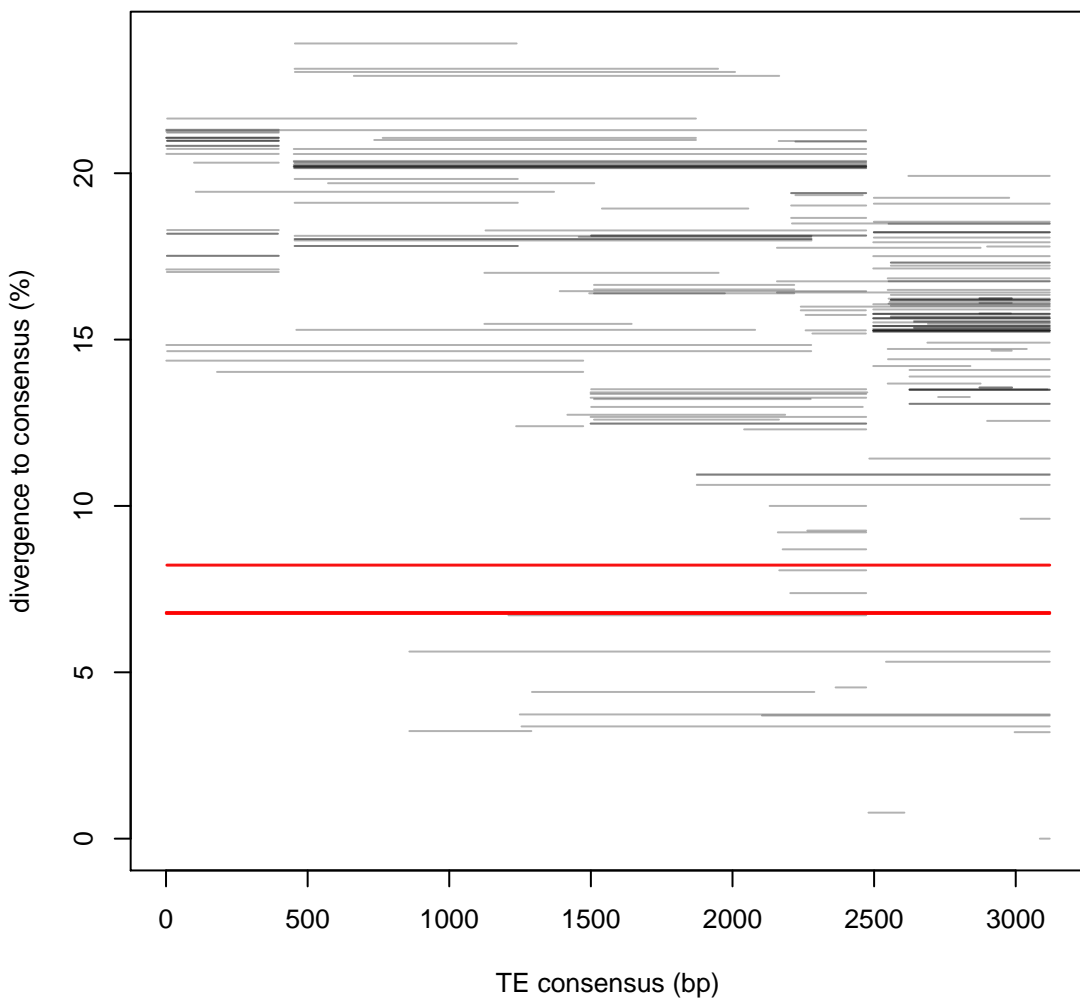
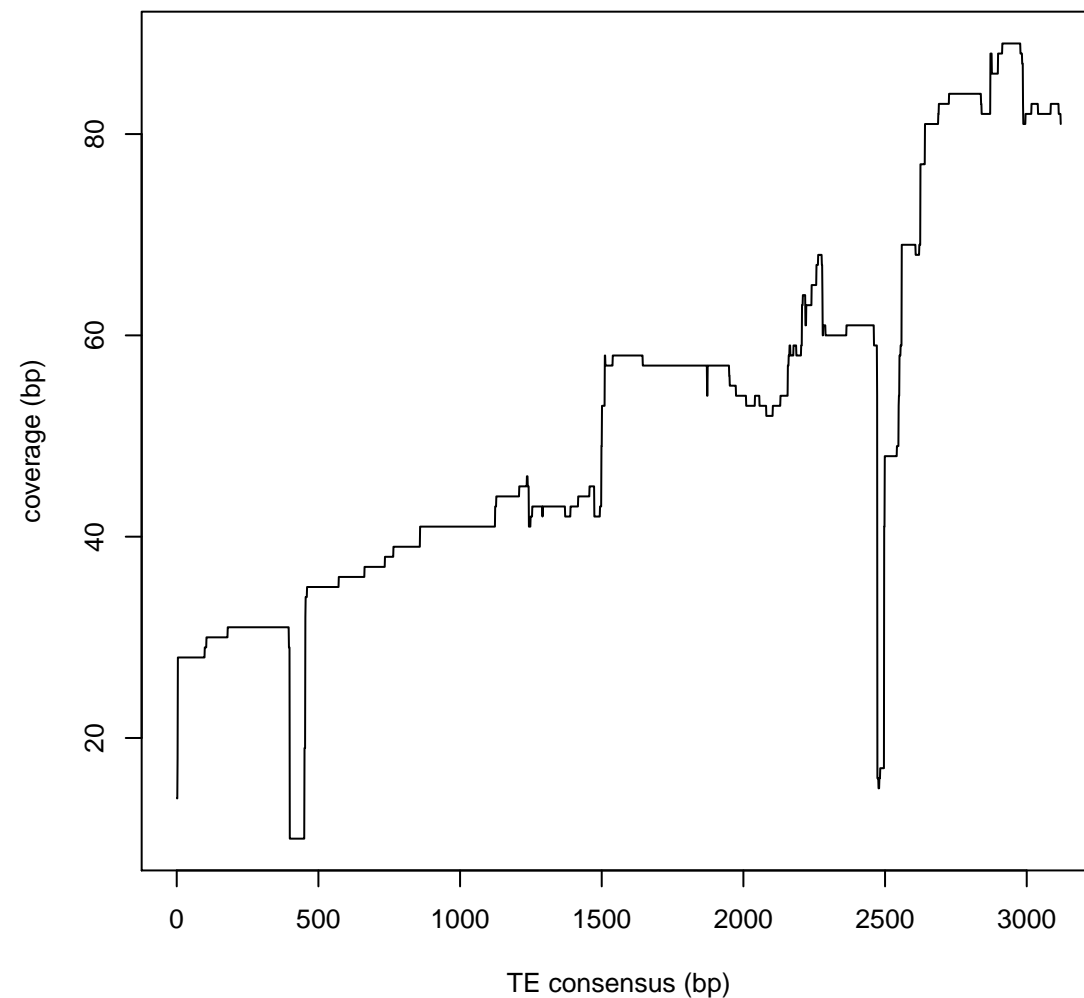


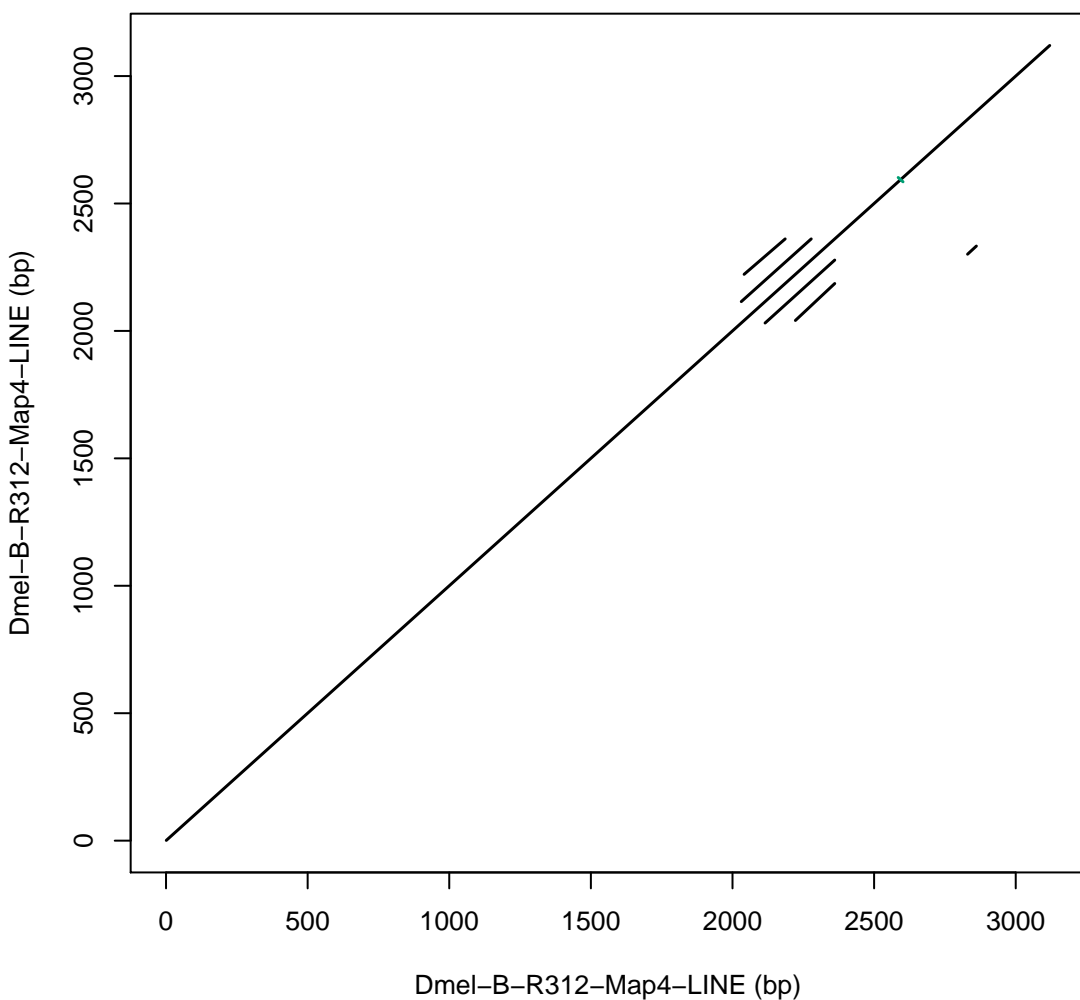
TE: Dmel-B-R312-Map4-LINE
consensus size: 3120bp; fragments: 200; full length: 3 (>=2808bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

