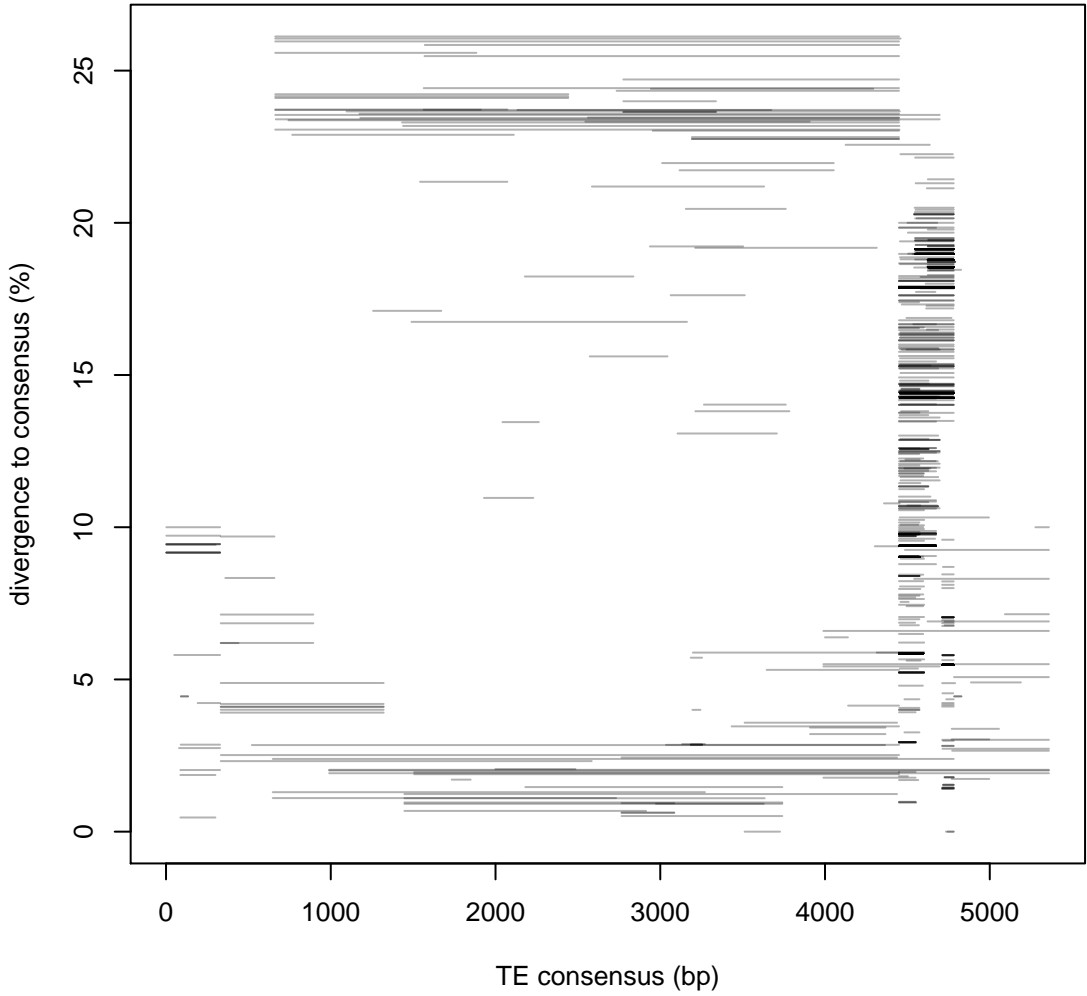
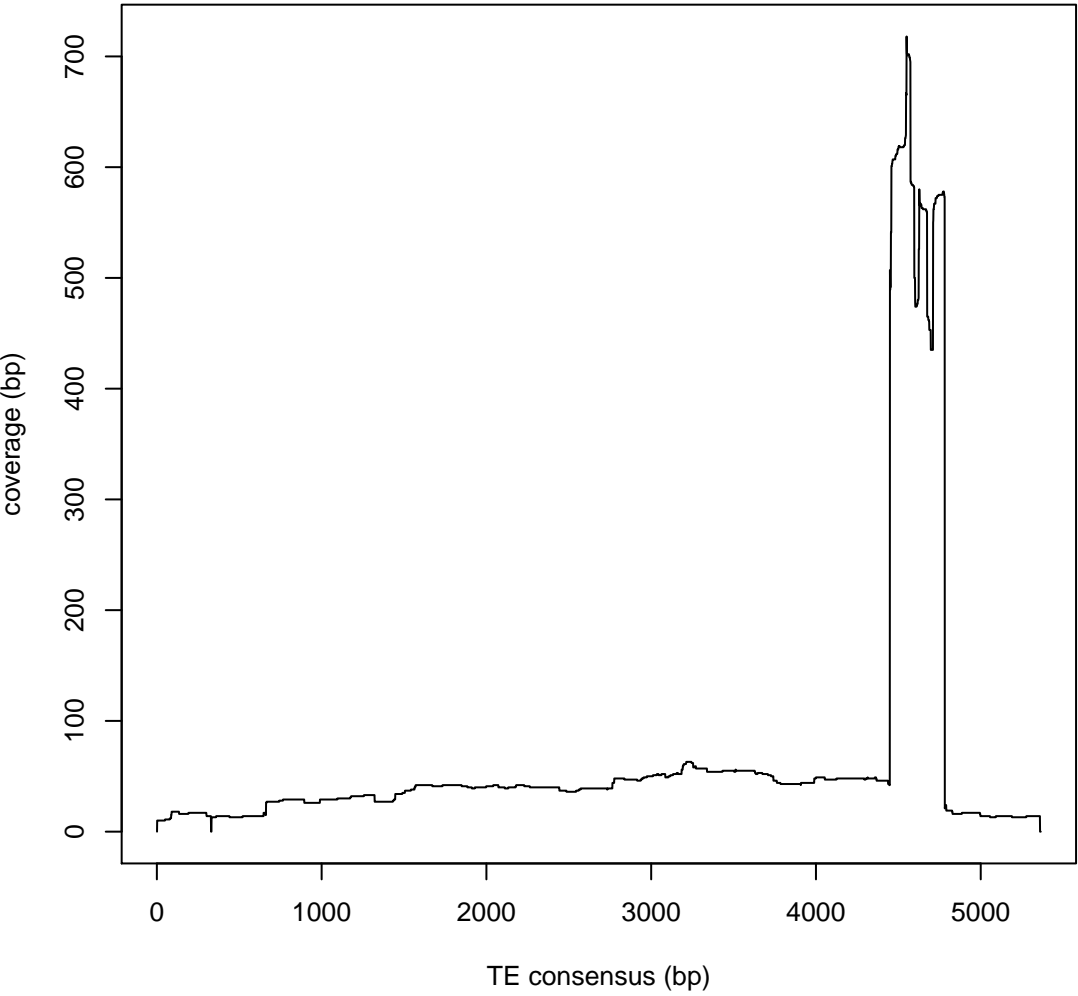


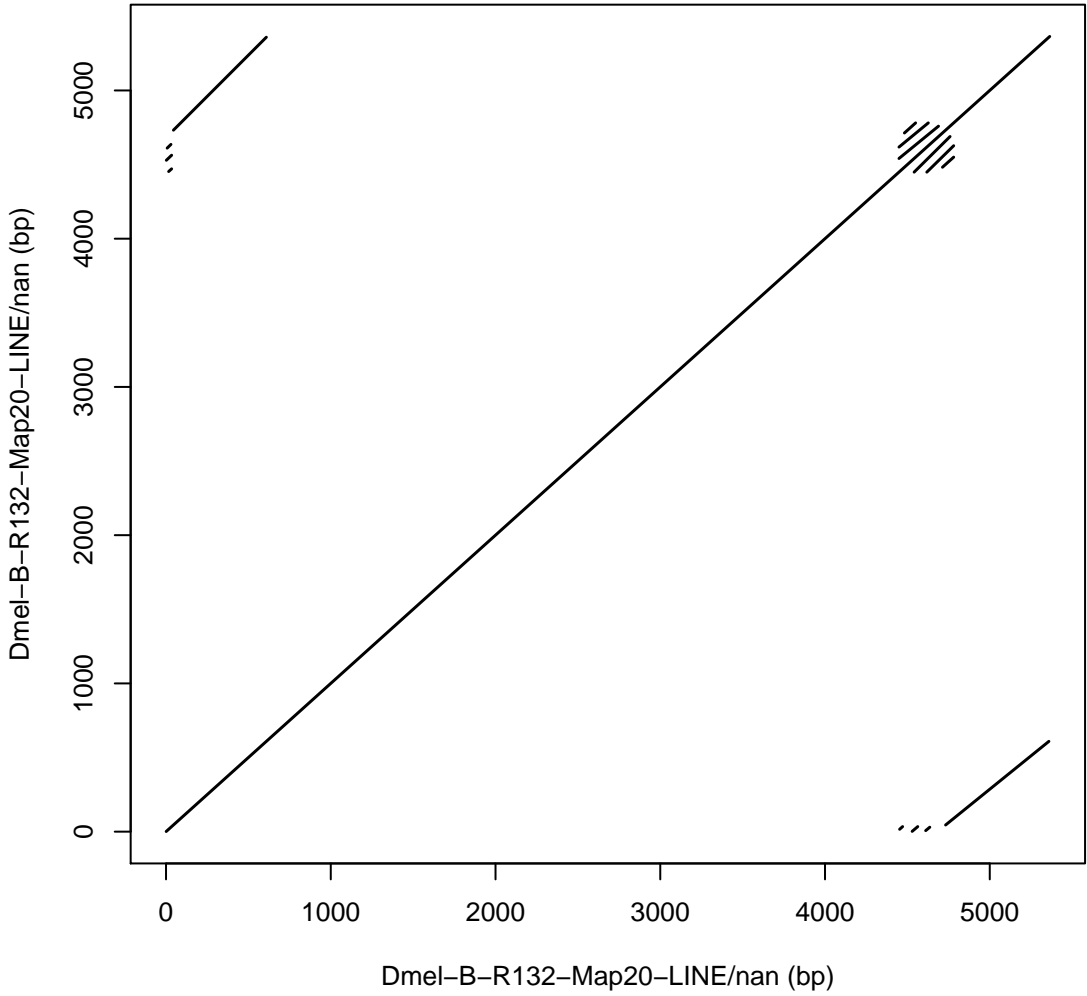
TE: Dmel-B-R132-Map20-LINE/nan  
consensus size: 5364bp; fragments: 1106; full length: 0 (>=4827.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

