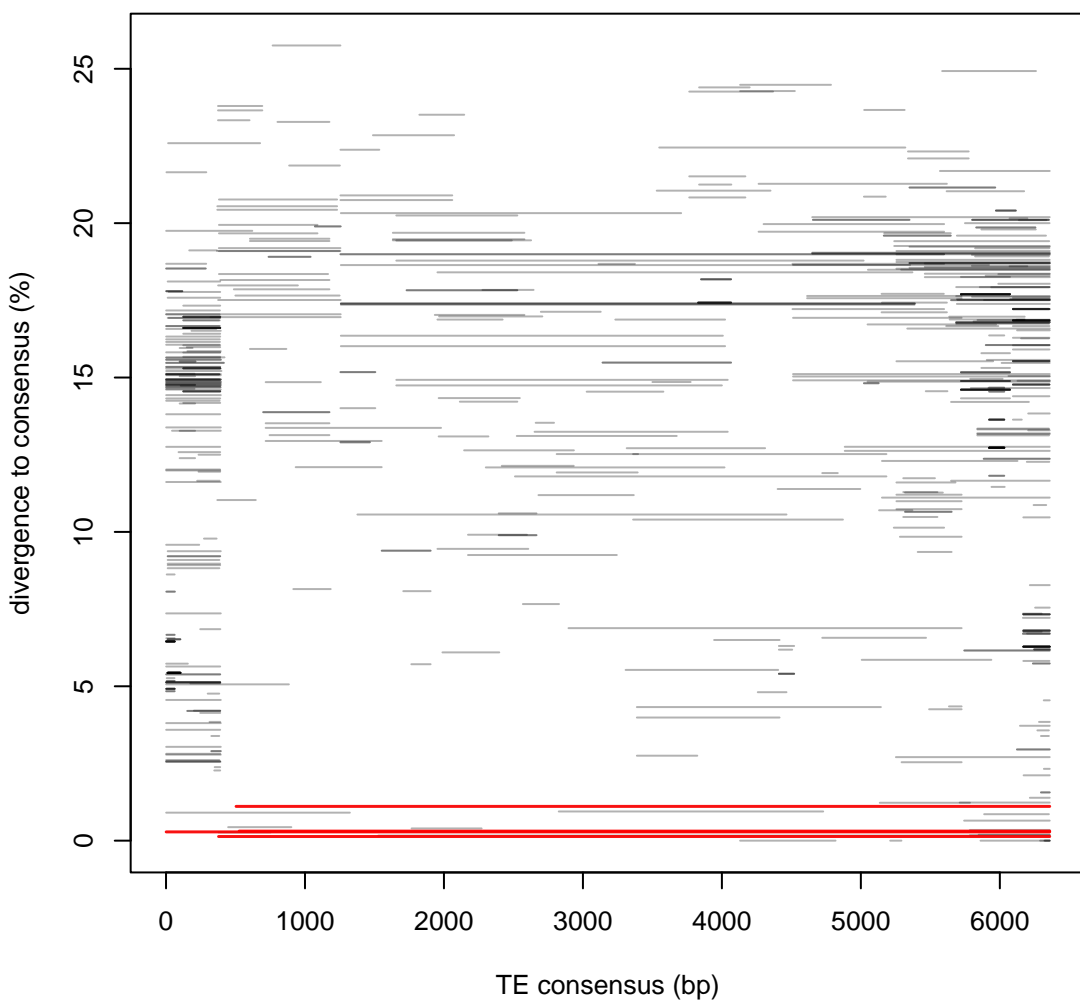
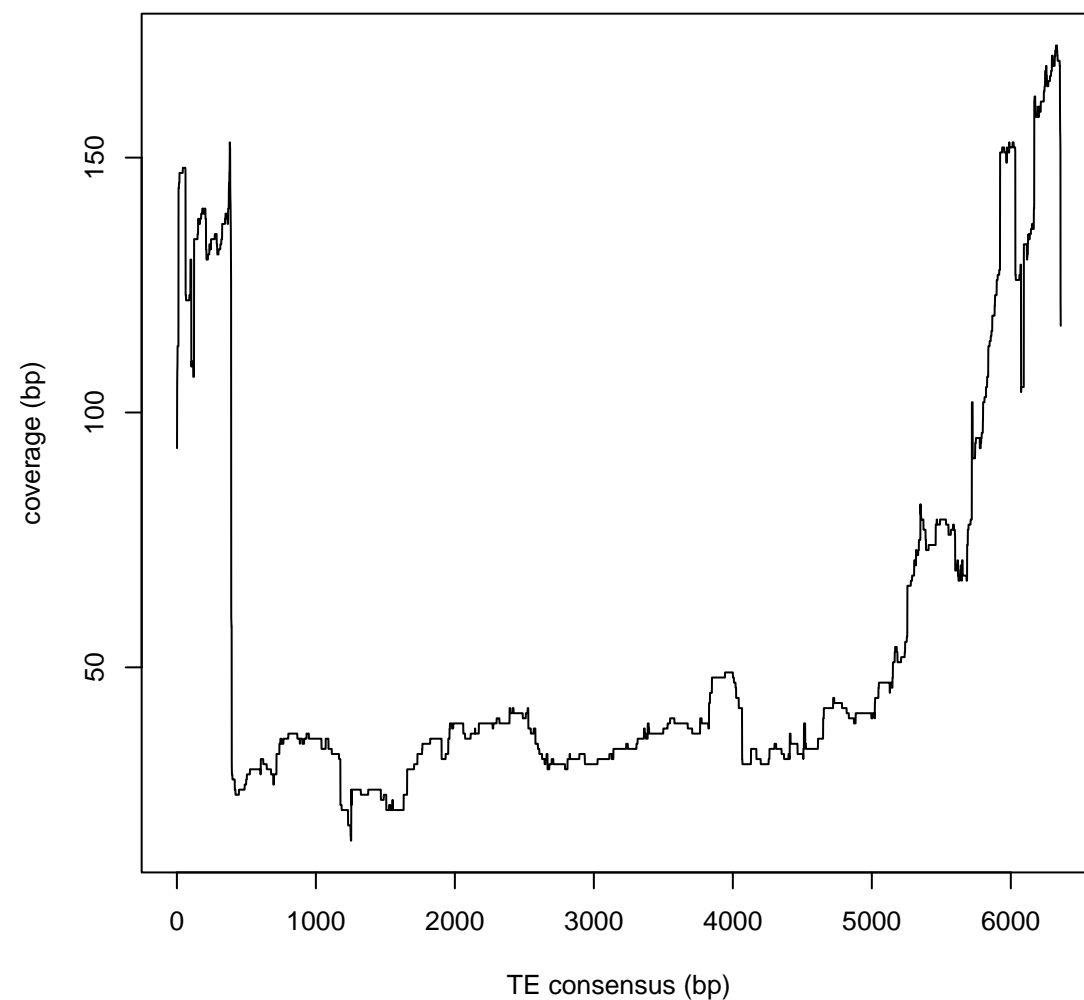


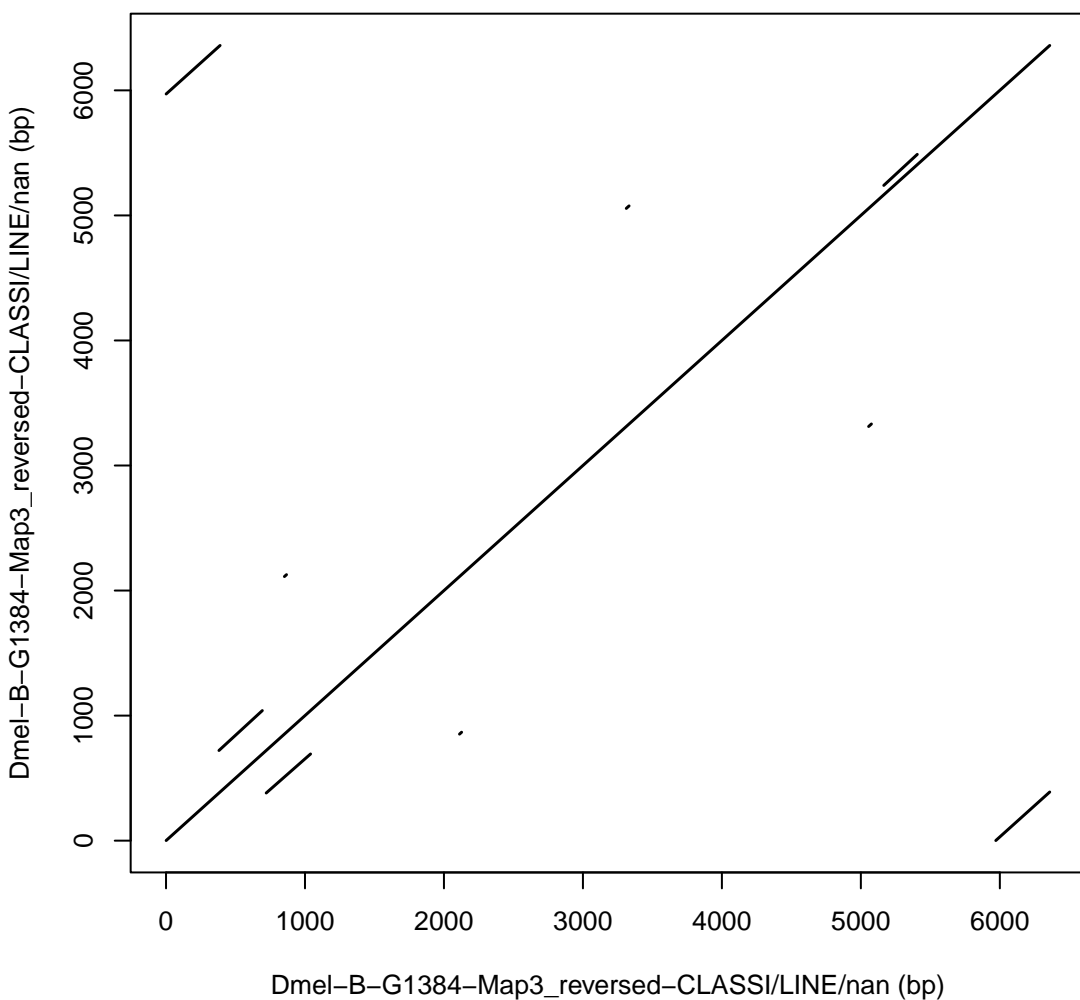
TE: Dmel-B-G1384-Map3_reversed-CLASSI/LINE/nan
consensus size: 6359bp; fragments: 641; full length: 4 (>=5723.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

