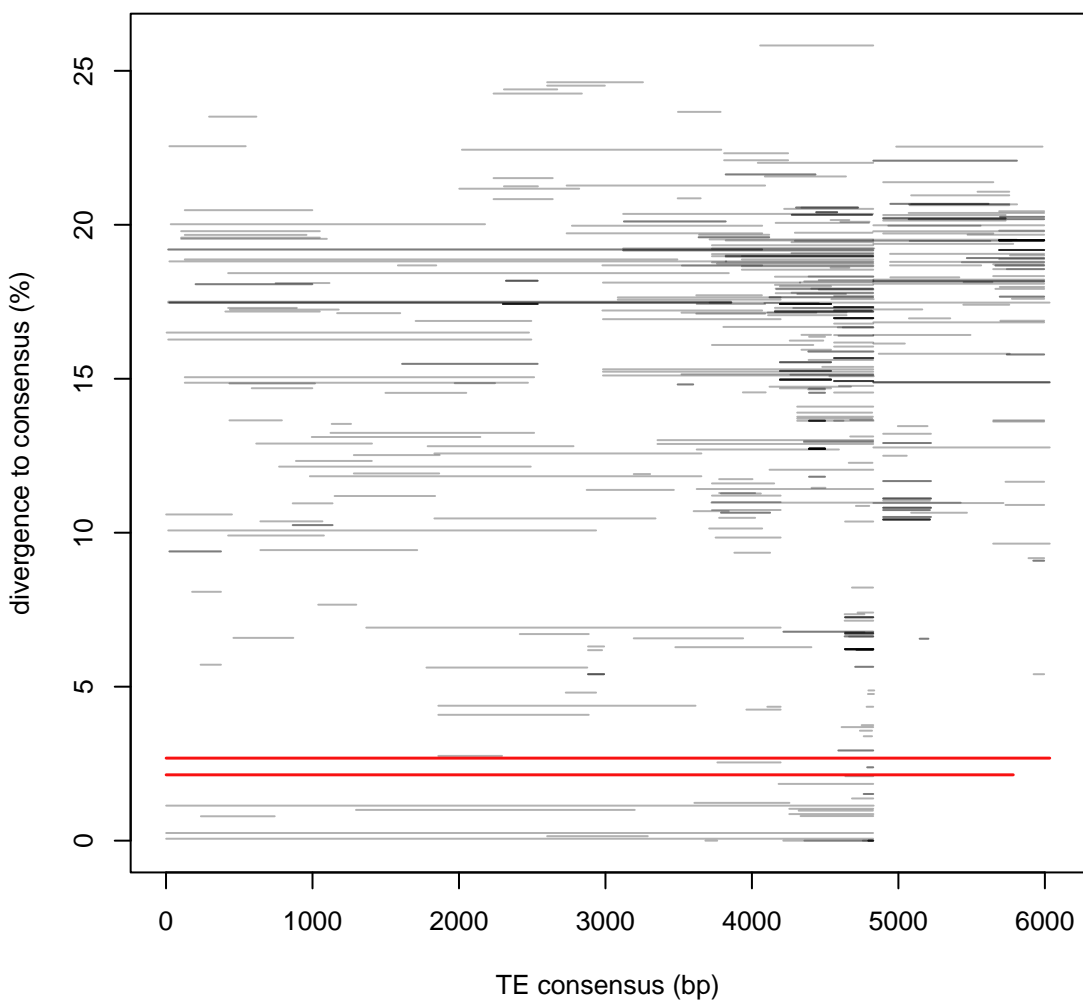
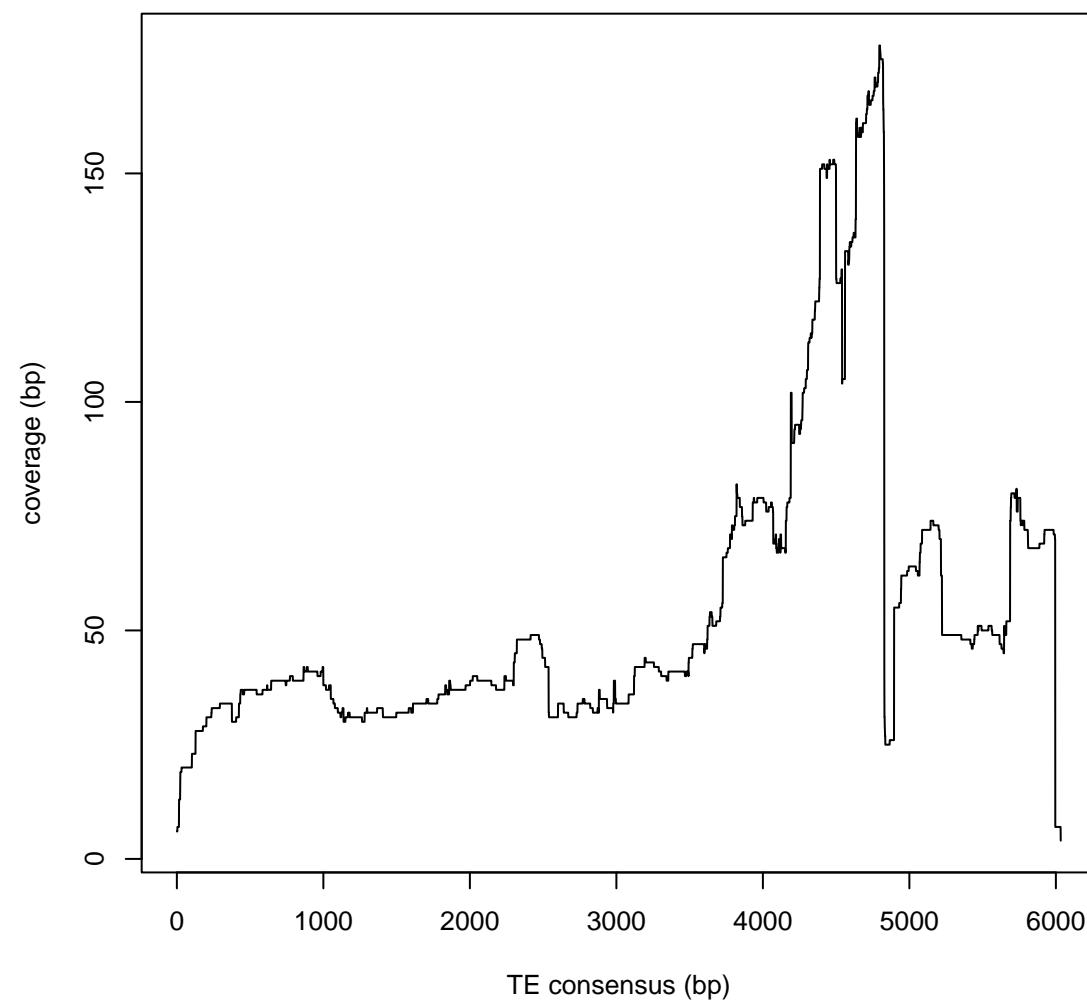


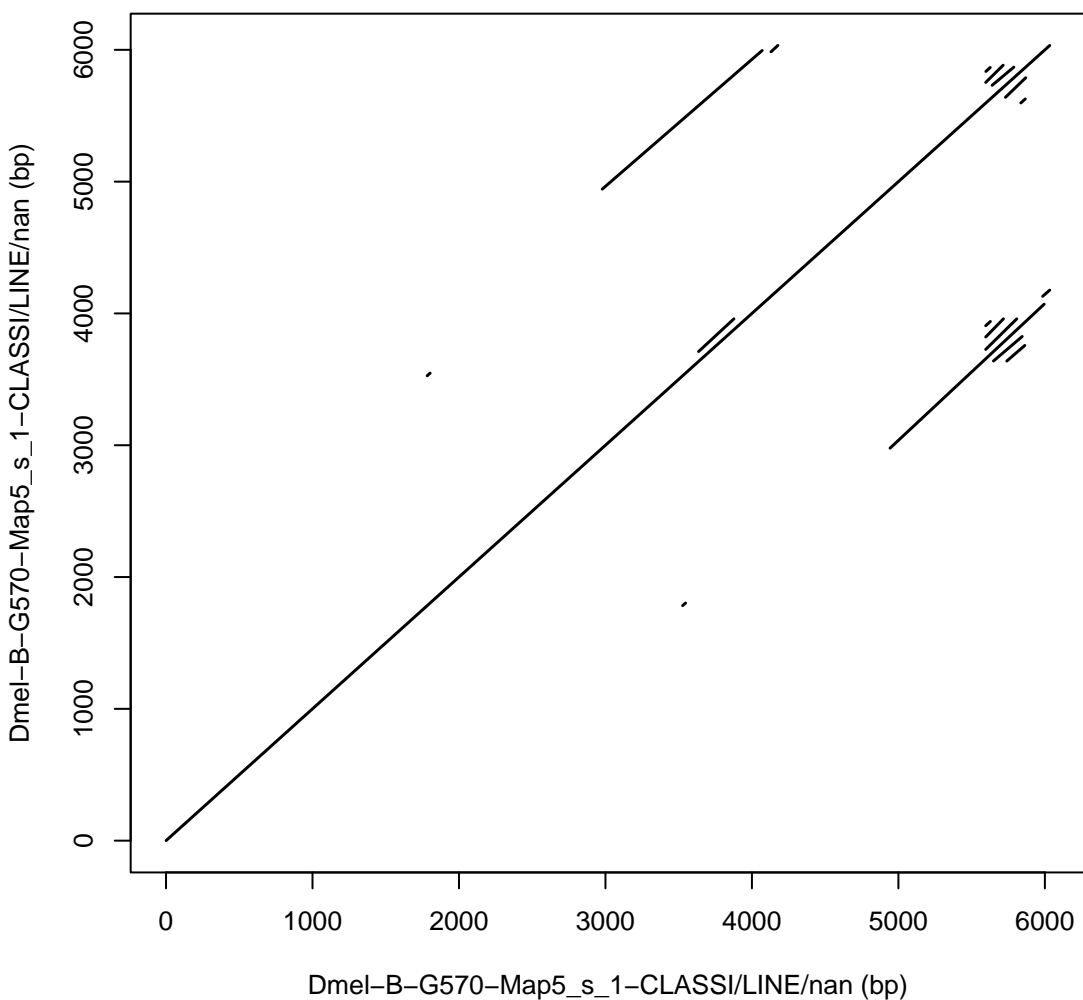
TE: Dmel-B-G570-Map5_s_1-CLASSI/LINE/nan
consensus size: 6033bp; fragments: 522; full length: 2 (>=5429.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

