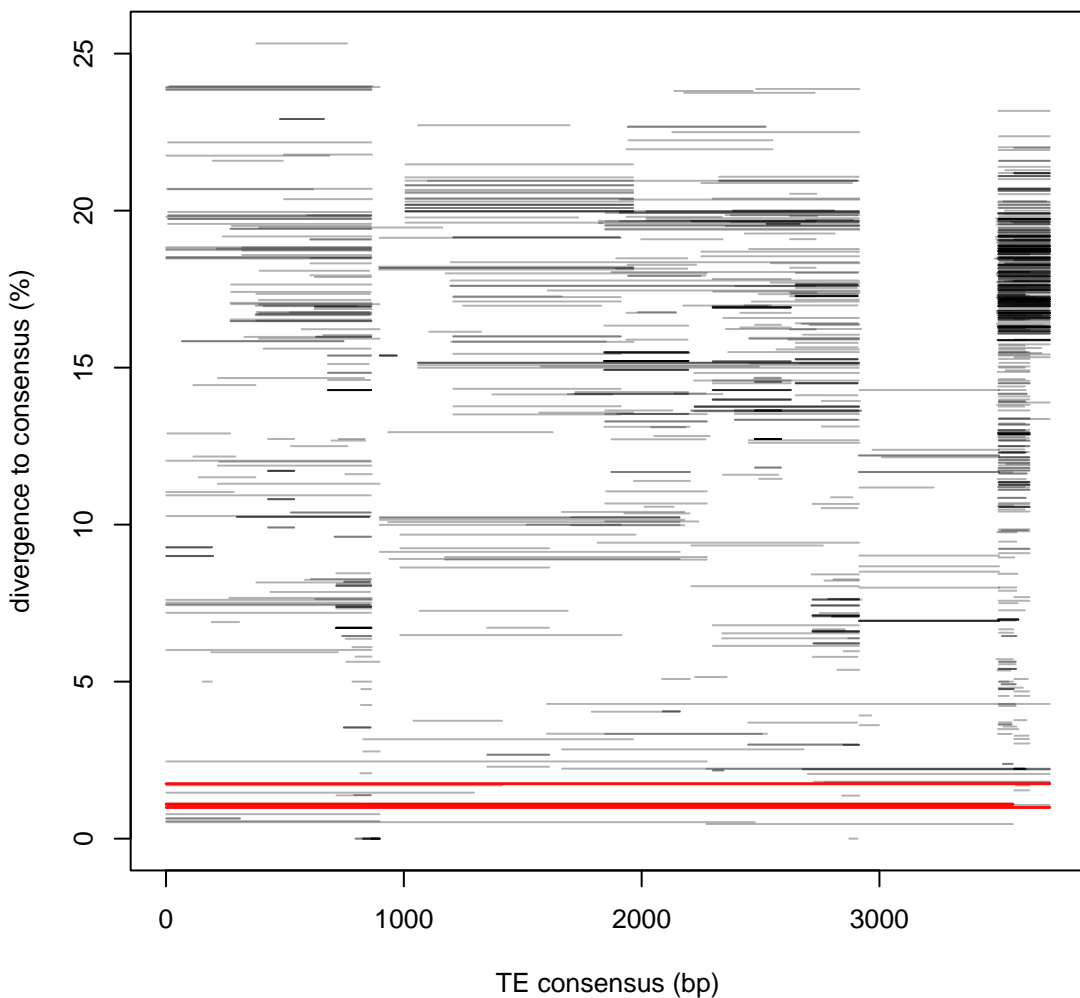
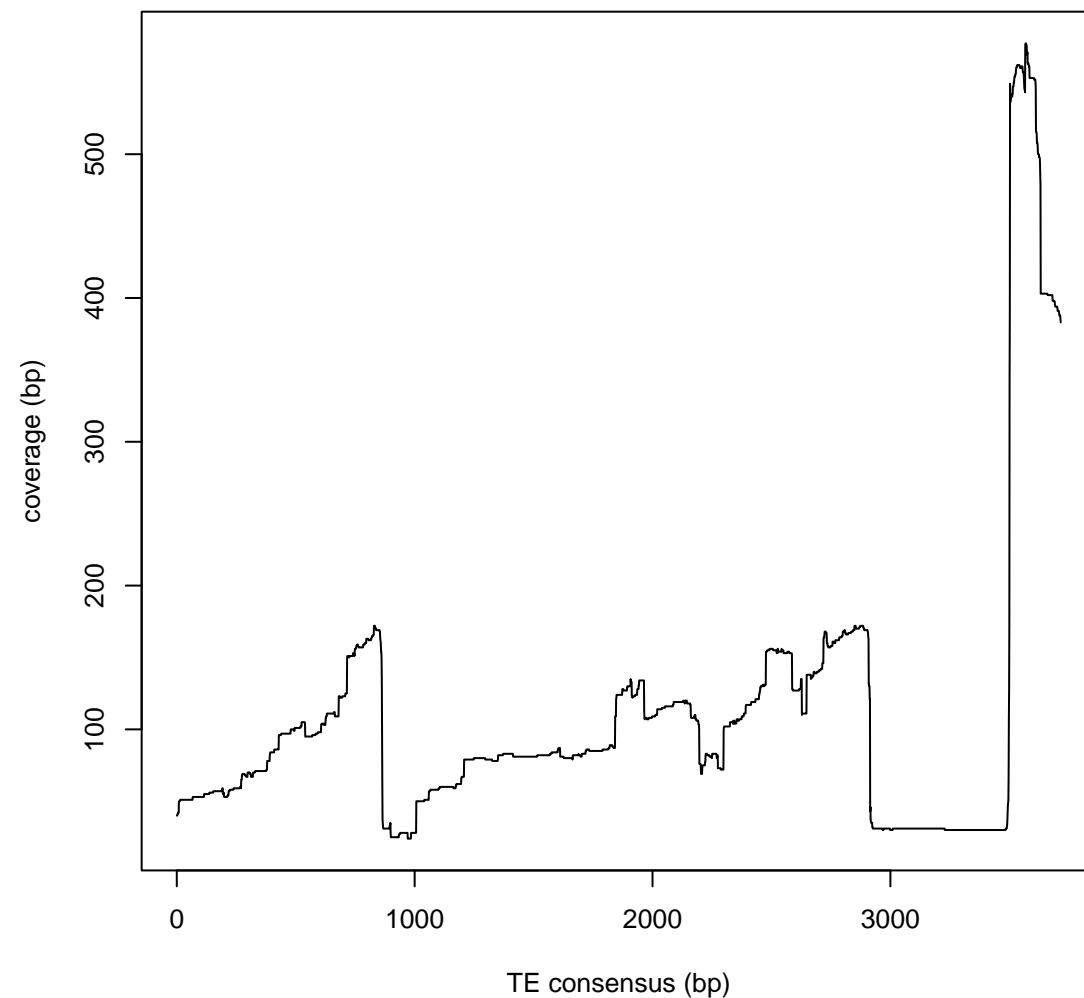


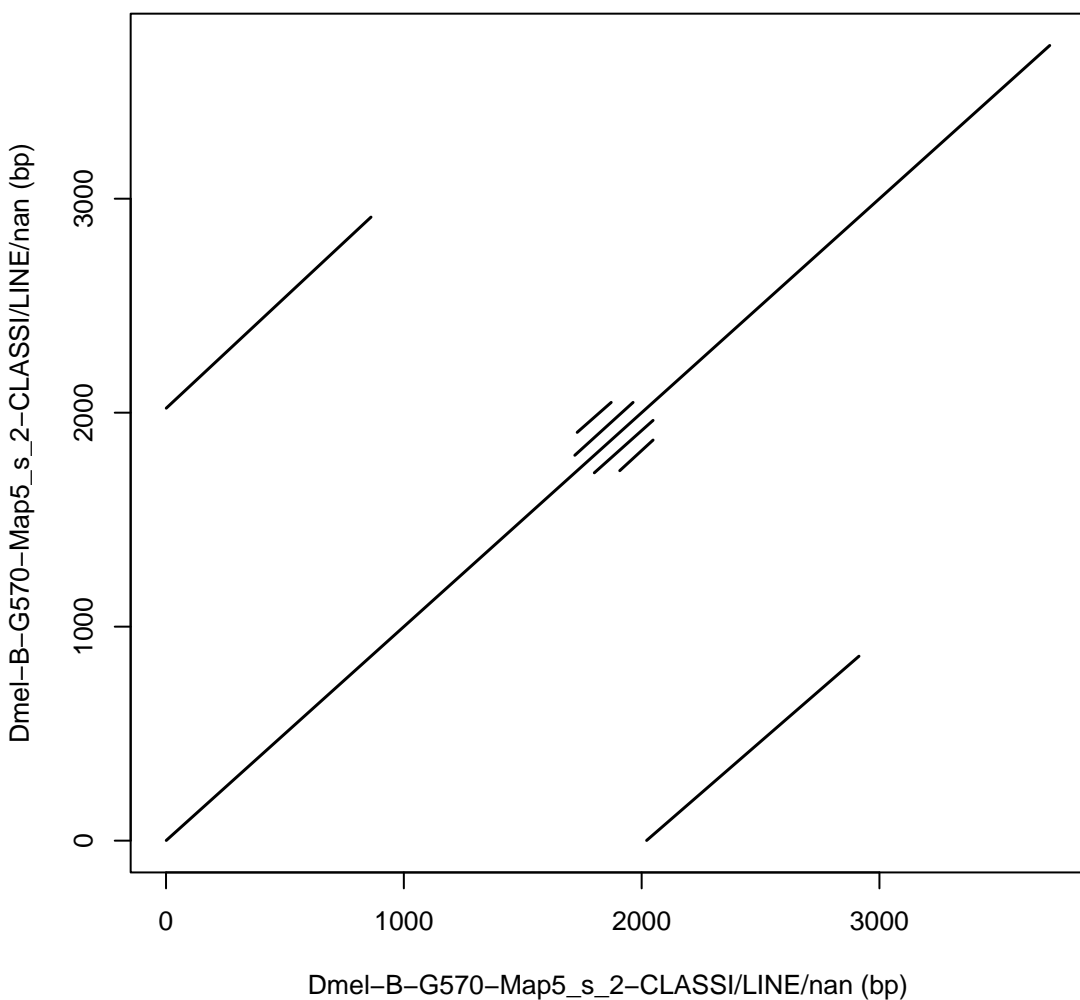
TE: Dmel-B-G570-Map5\_s\_2-CLASSI/LINE/nan  
consensus size: 3716bp; fragments: 1215; full length: 4 (>=3344.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

