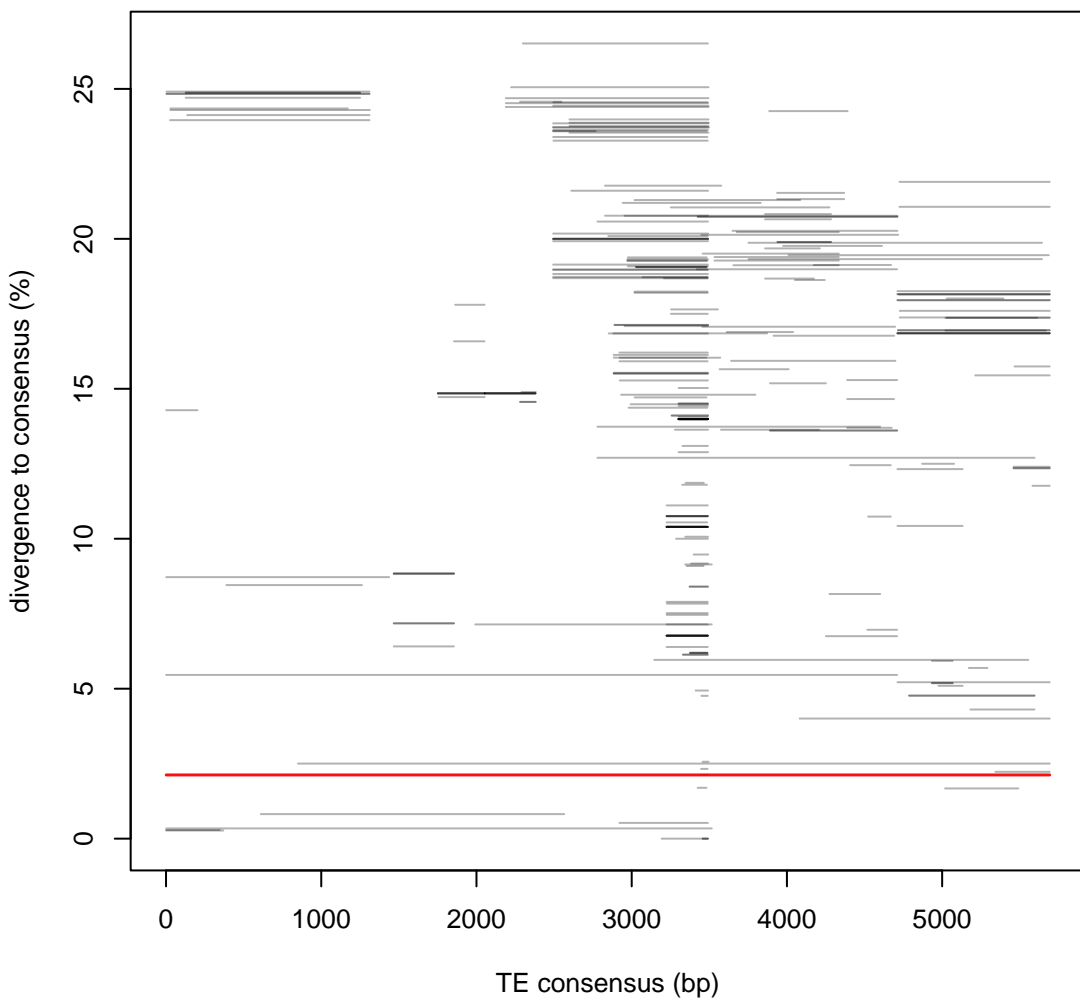
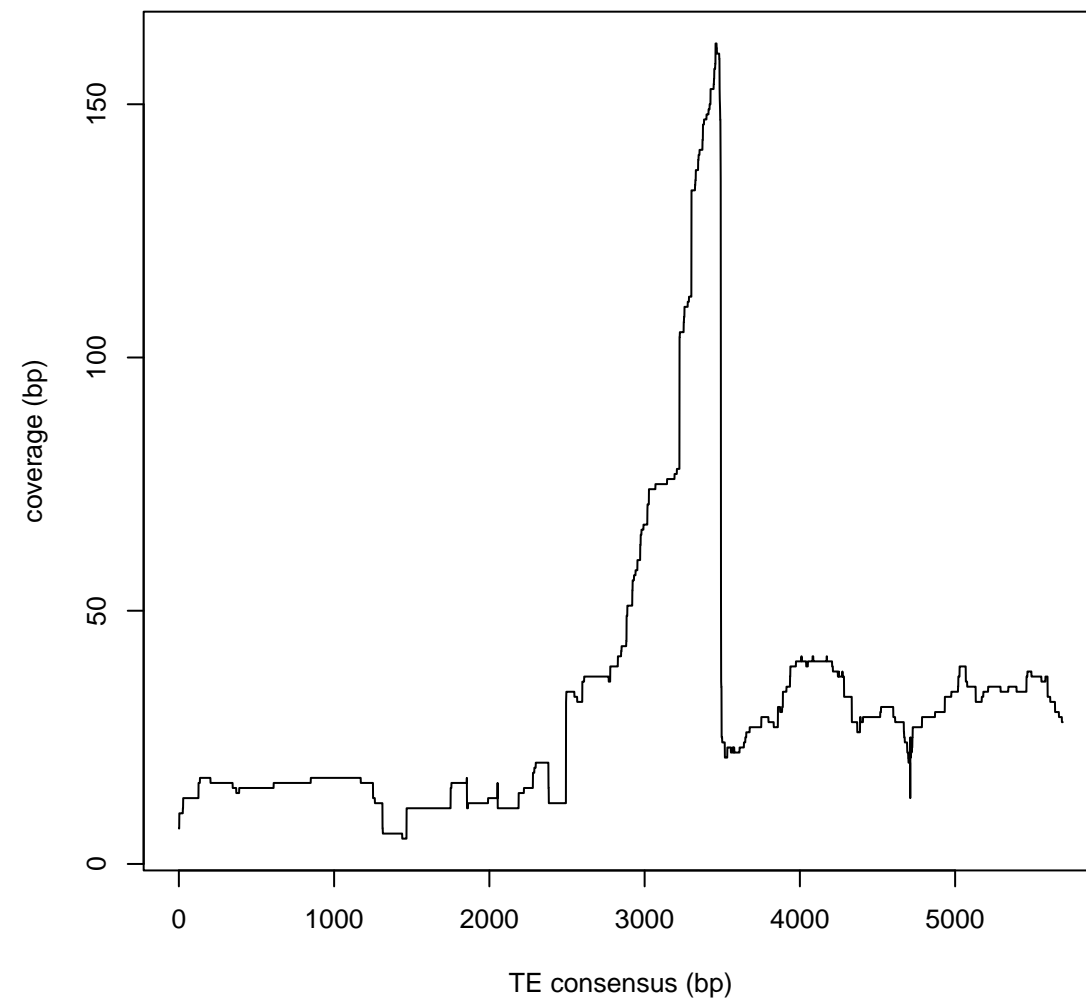


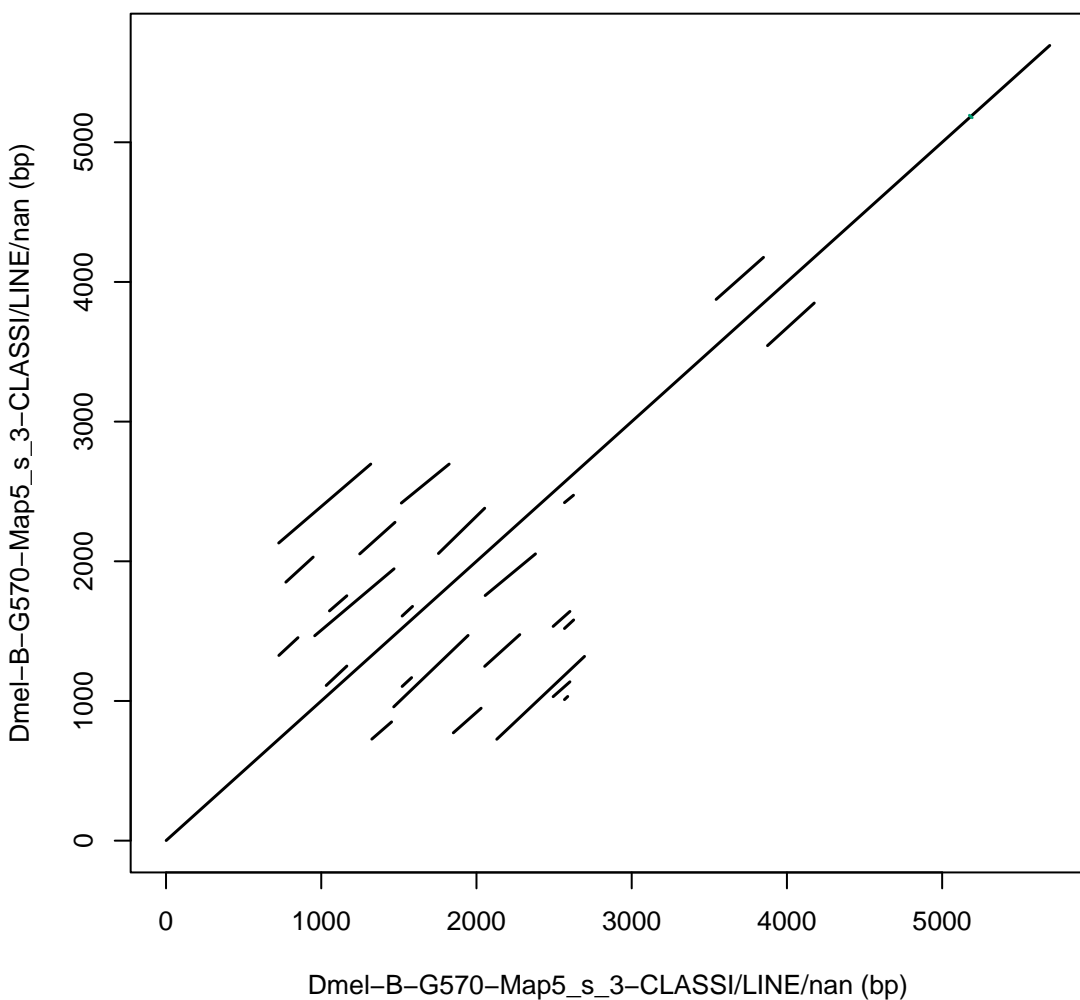
TE: Dmel-B-G570-Map5\_s\_3-CLASSI/LINE/nan  
consensus size: 5693bp; fragments: 278; full length: 1 (>=5123.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

