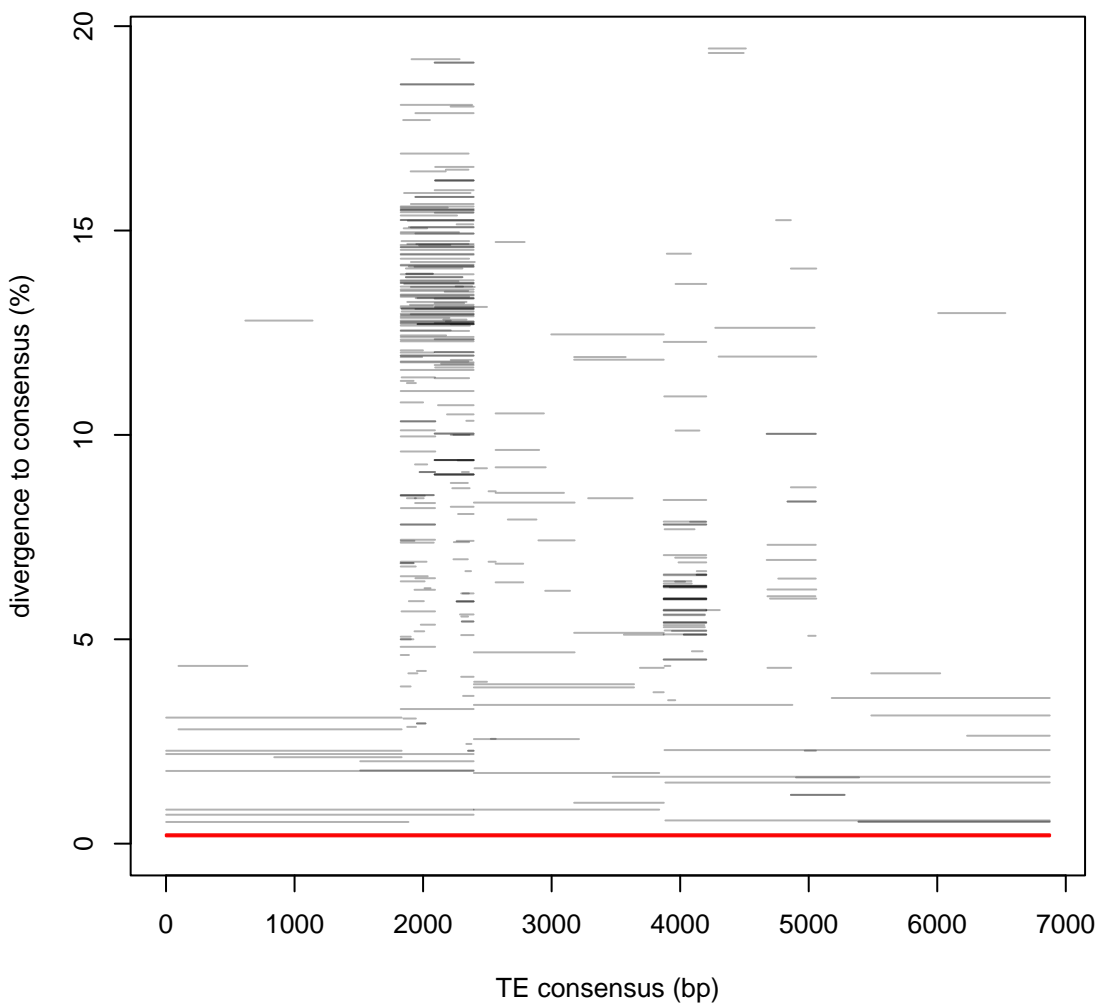
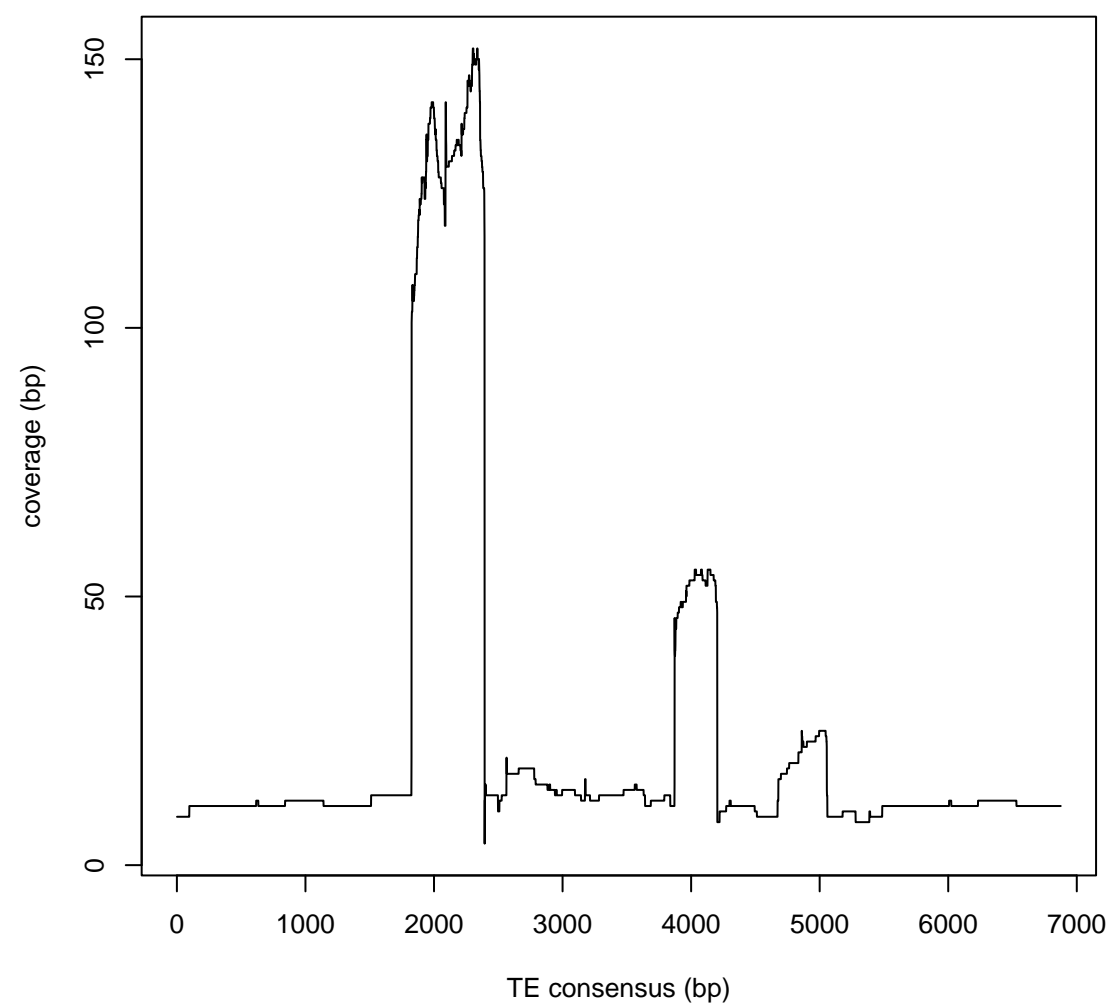


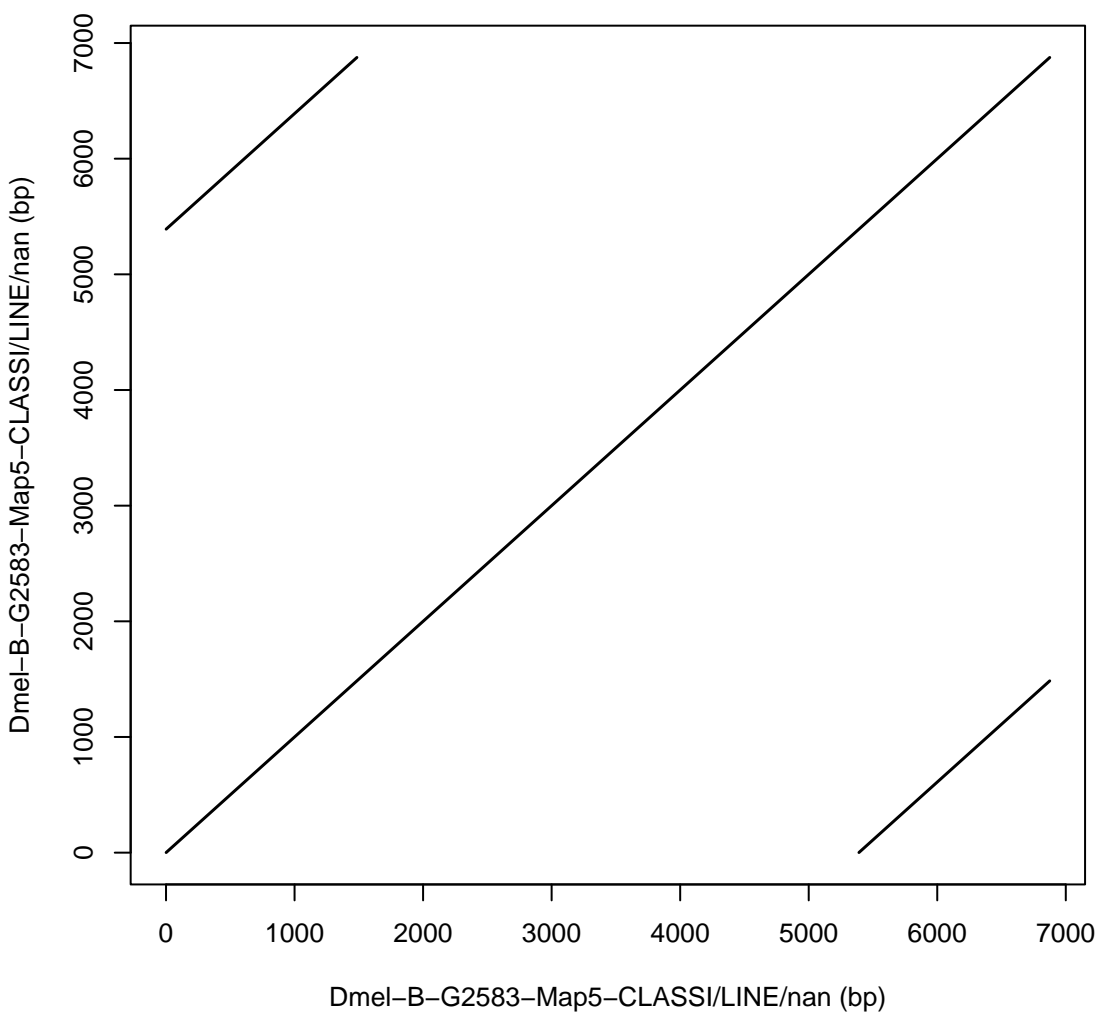
TE: Dmel-B-G2583-Map5-CLASSI/LINE/nan
consensus size: 6875bp; fragments: 357; full length: 2 (>=6187.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

