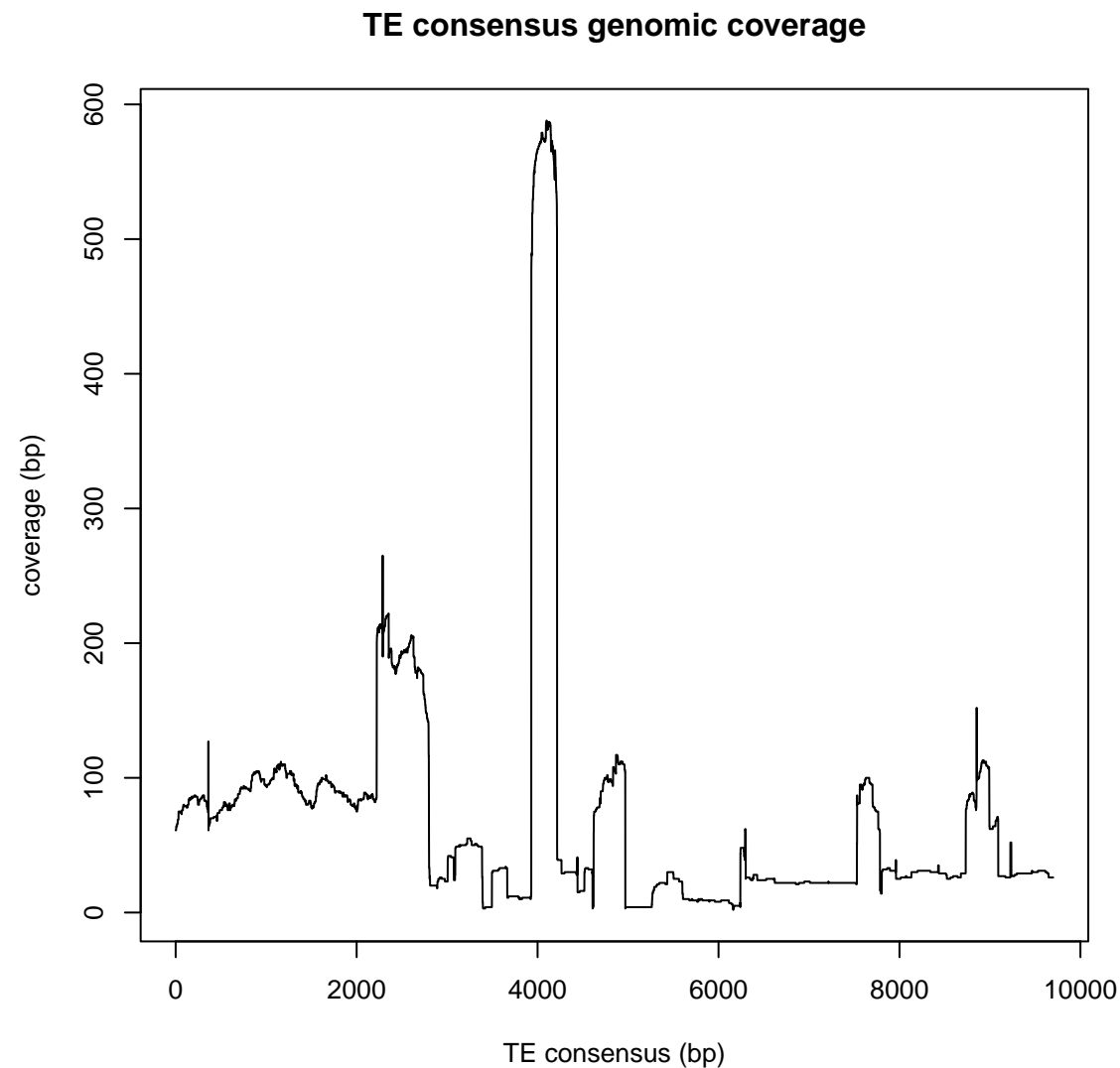
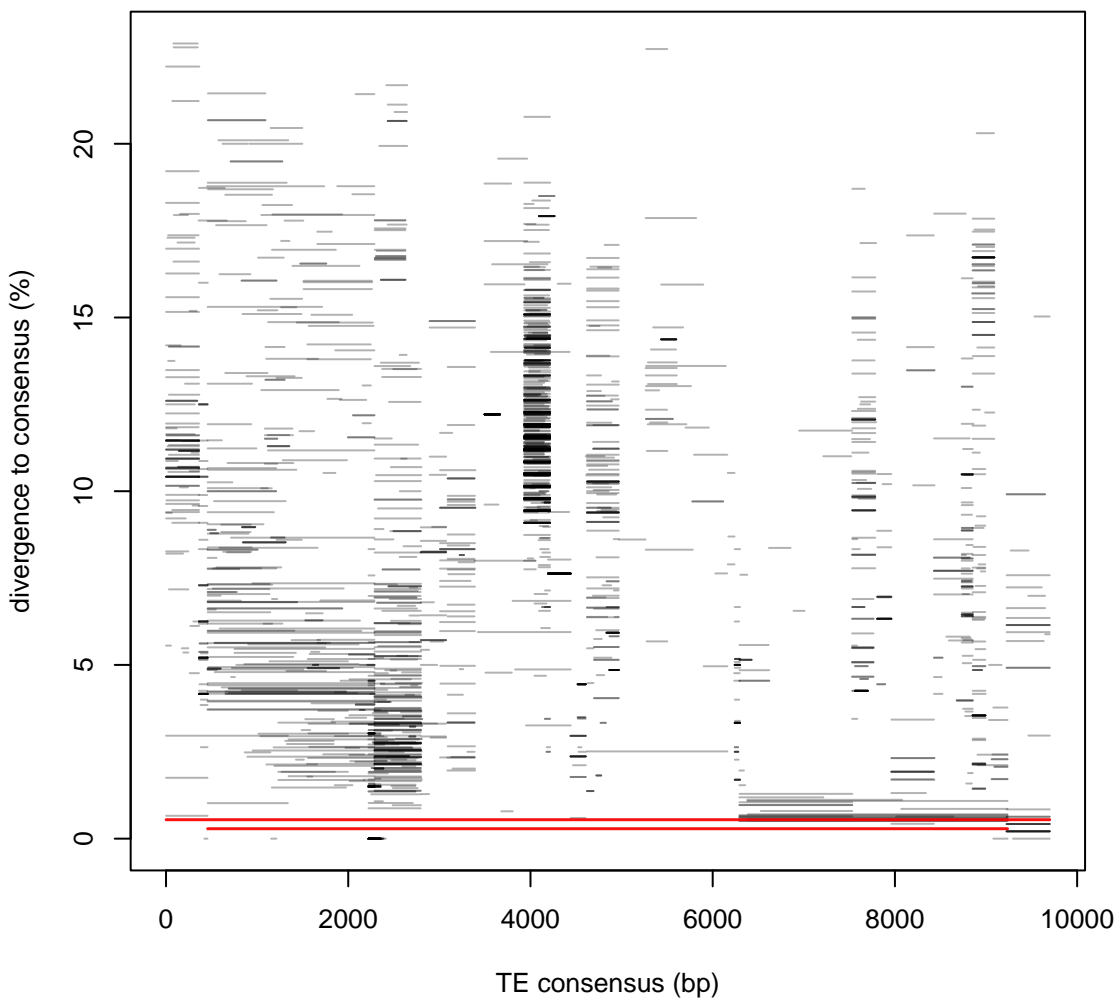
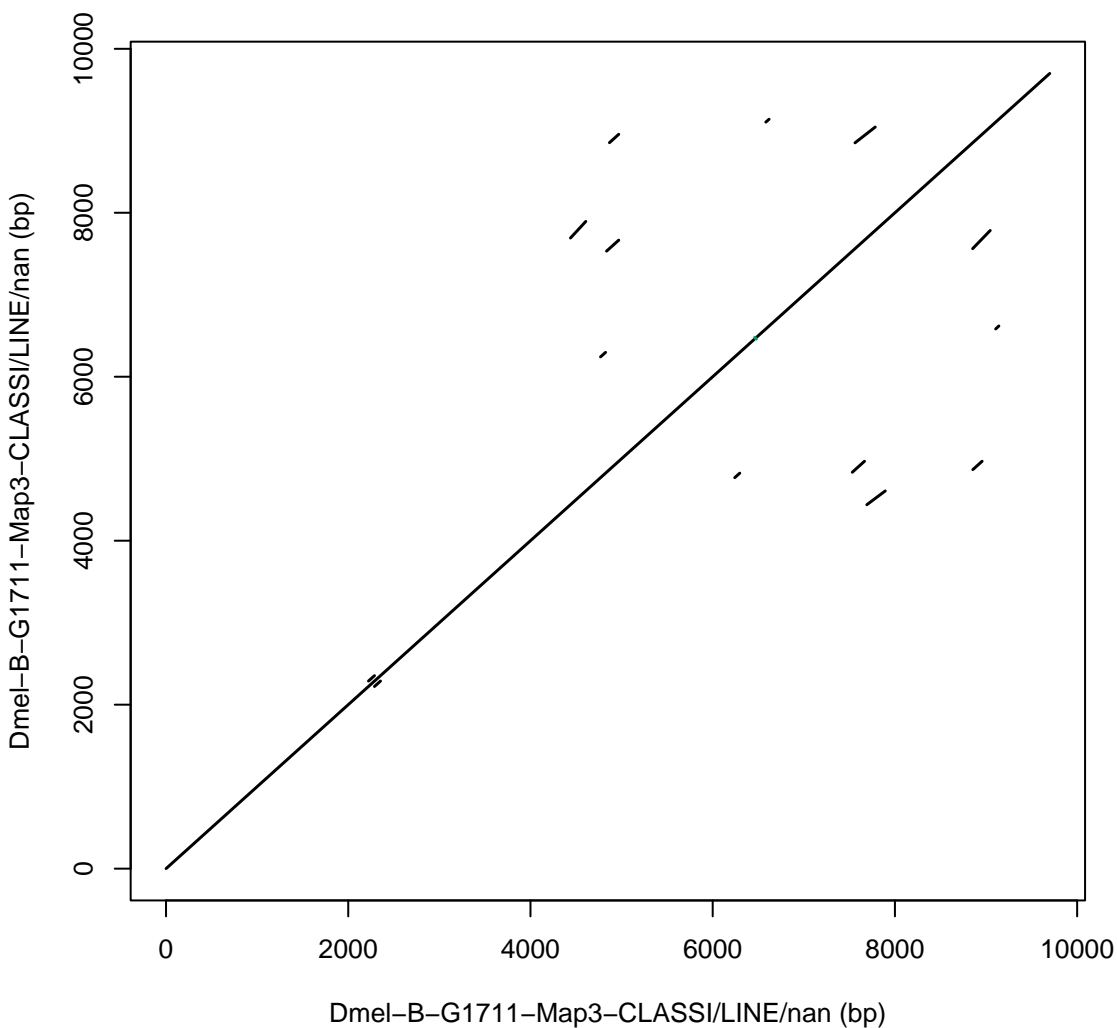


TE: Dmel-B-G1711-Map3-CLASSI/LINE/nan
consensus size: 9699bp; fragments: 2223; full length: 2 (>=8729.1bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

