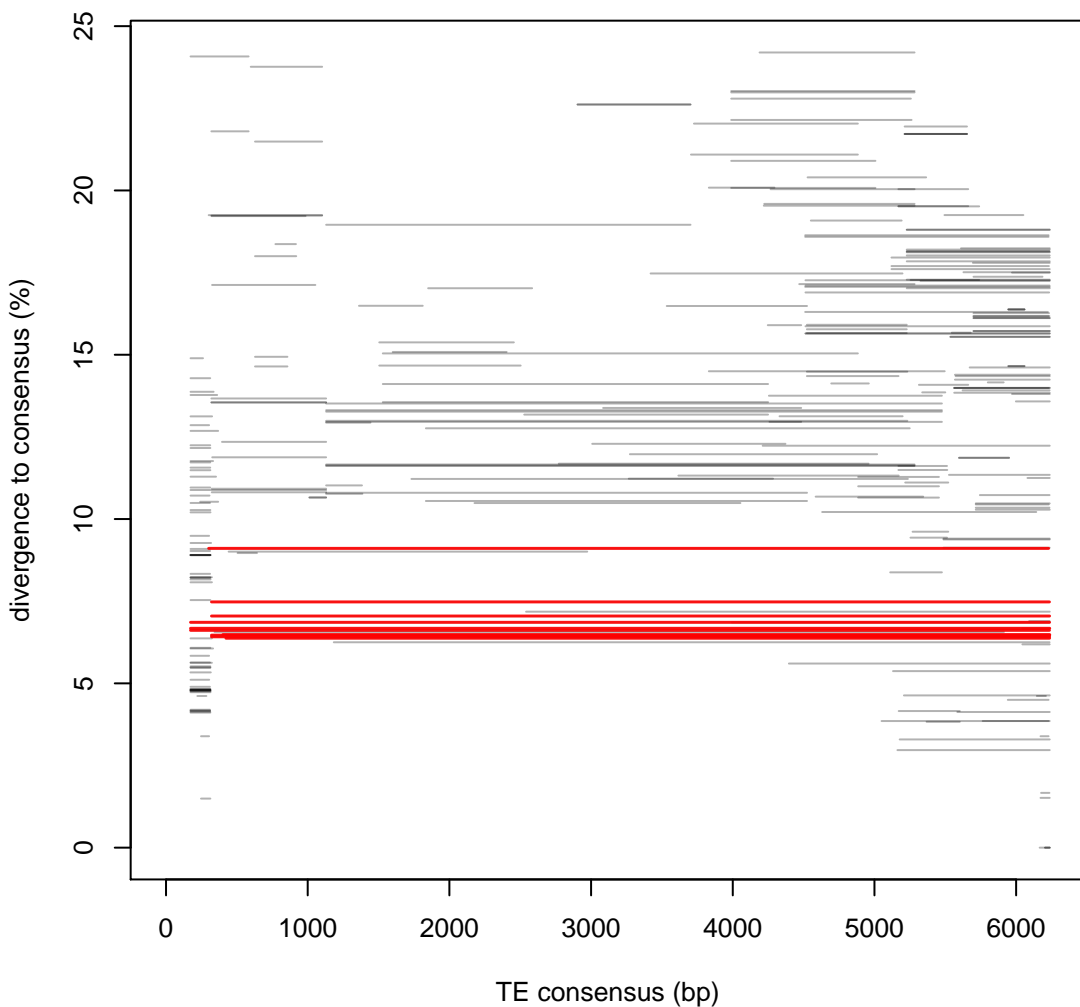
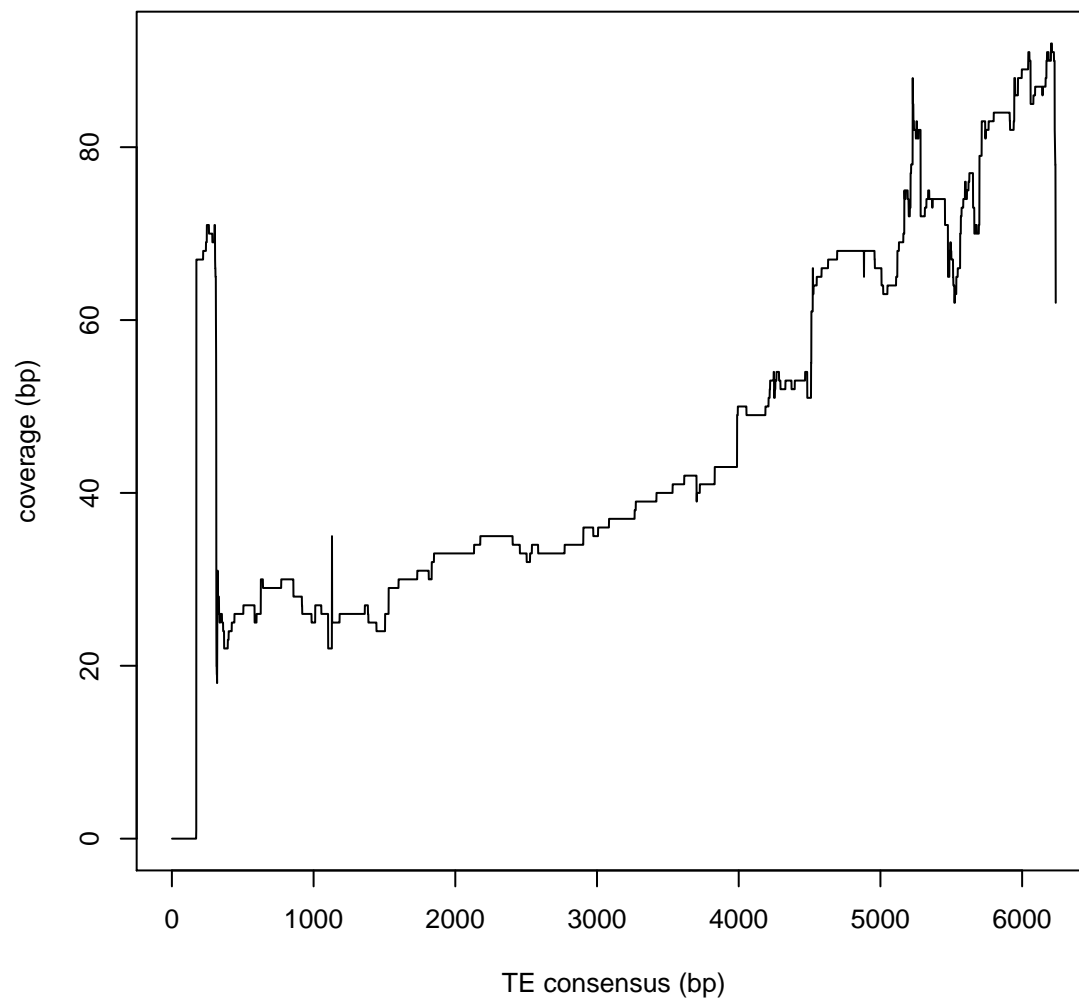


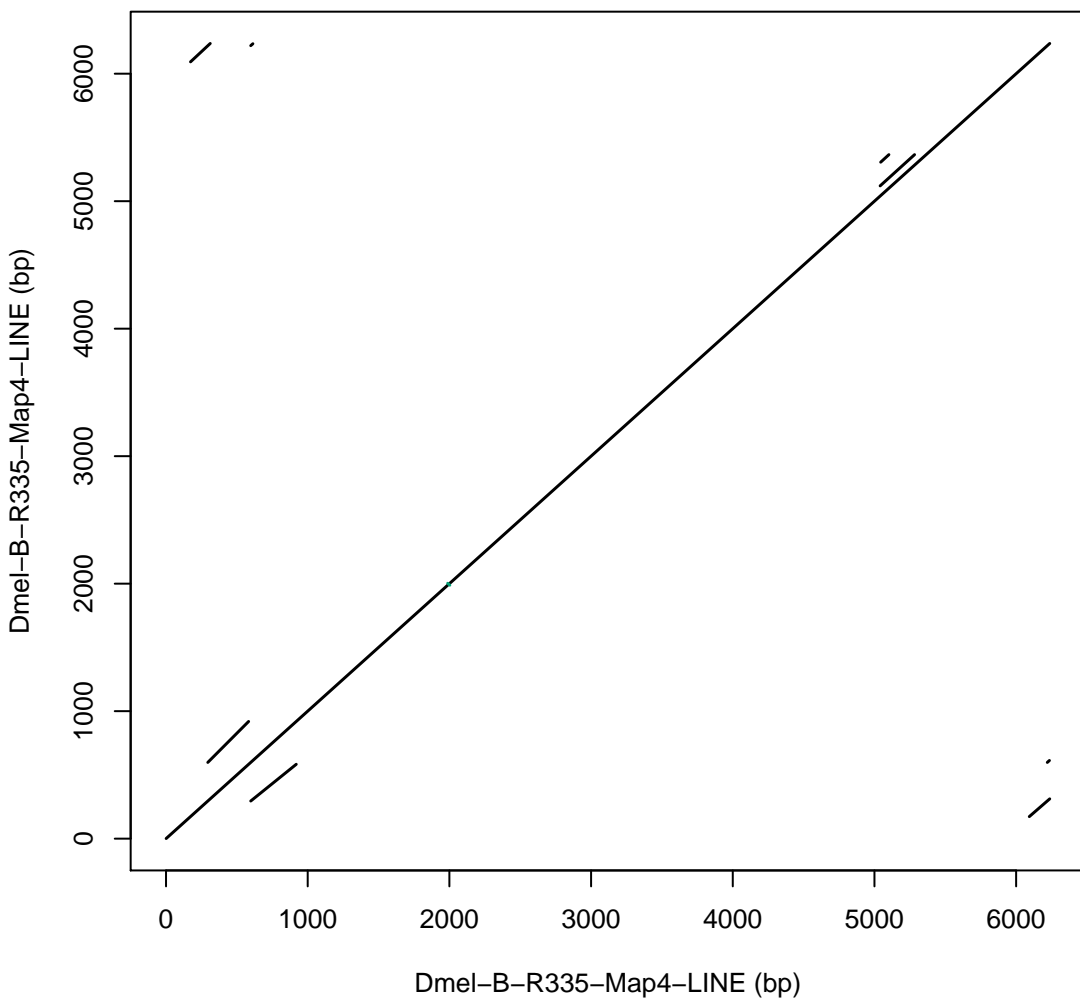
TE: Dmel-B-R335-Map4-LINE
consensus size: 6237bp; fragments: 281; full length: 10 (>=5613.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

