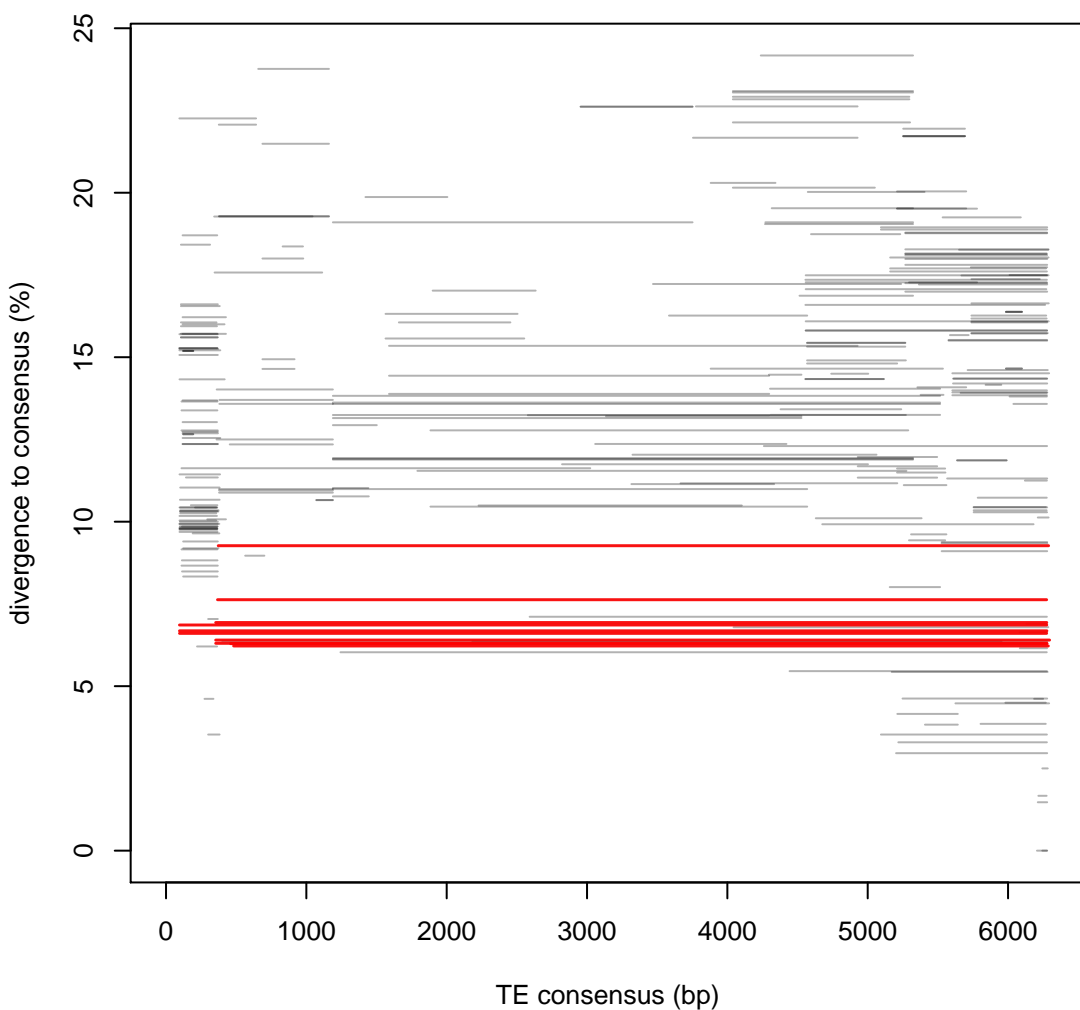
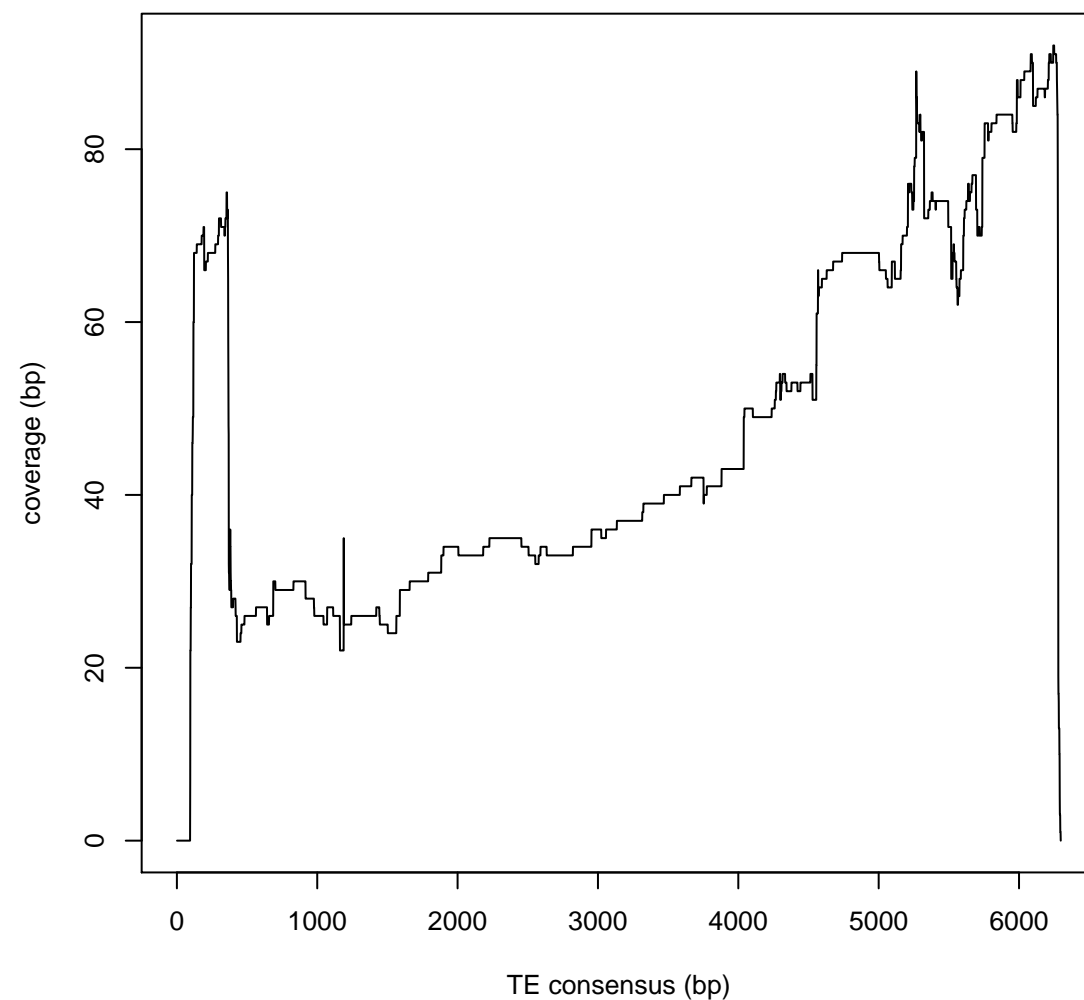


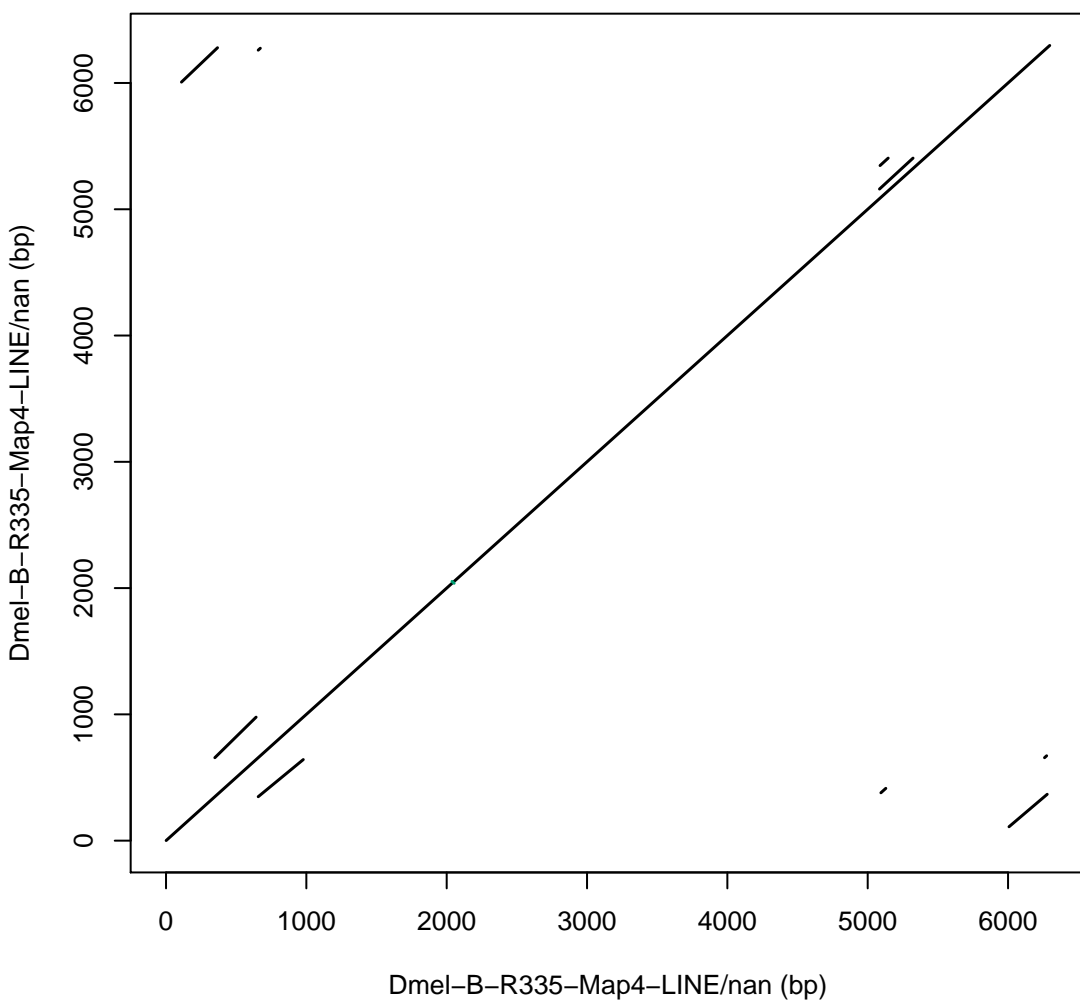
TE: Dmel-B-R335-Map4-LINE/nan
consensus size: 6297bp; fragments: 288; full length: 10 (>=5667.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

