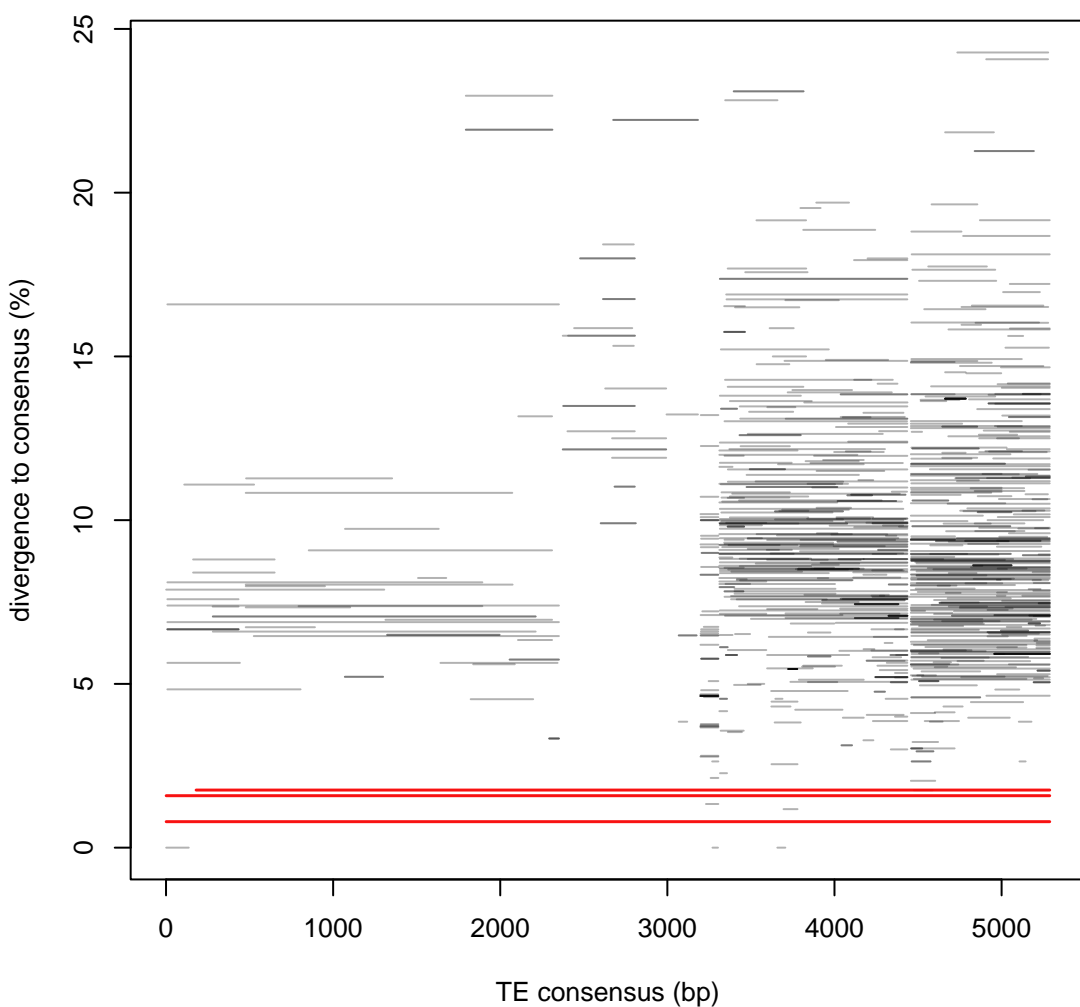
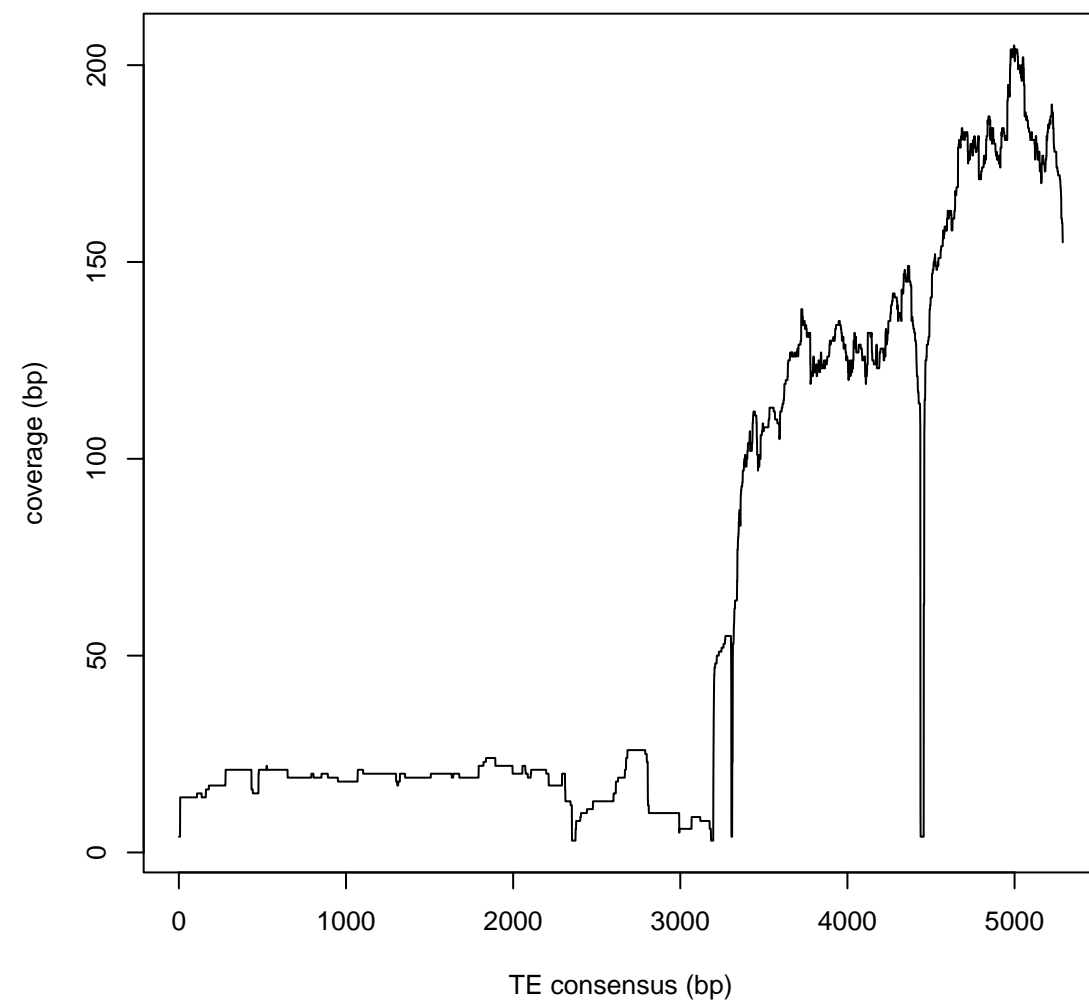


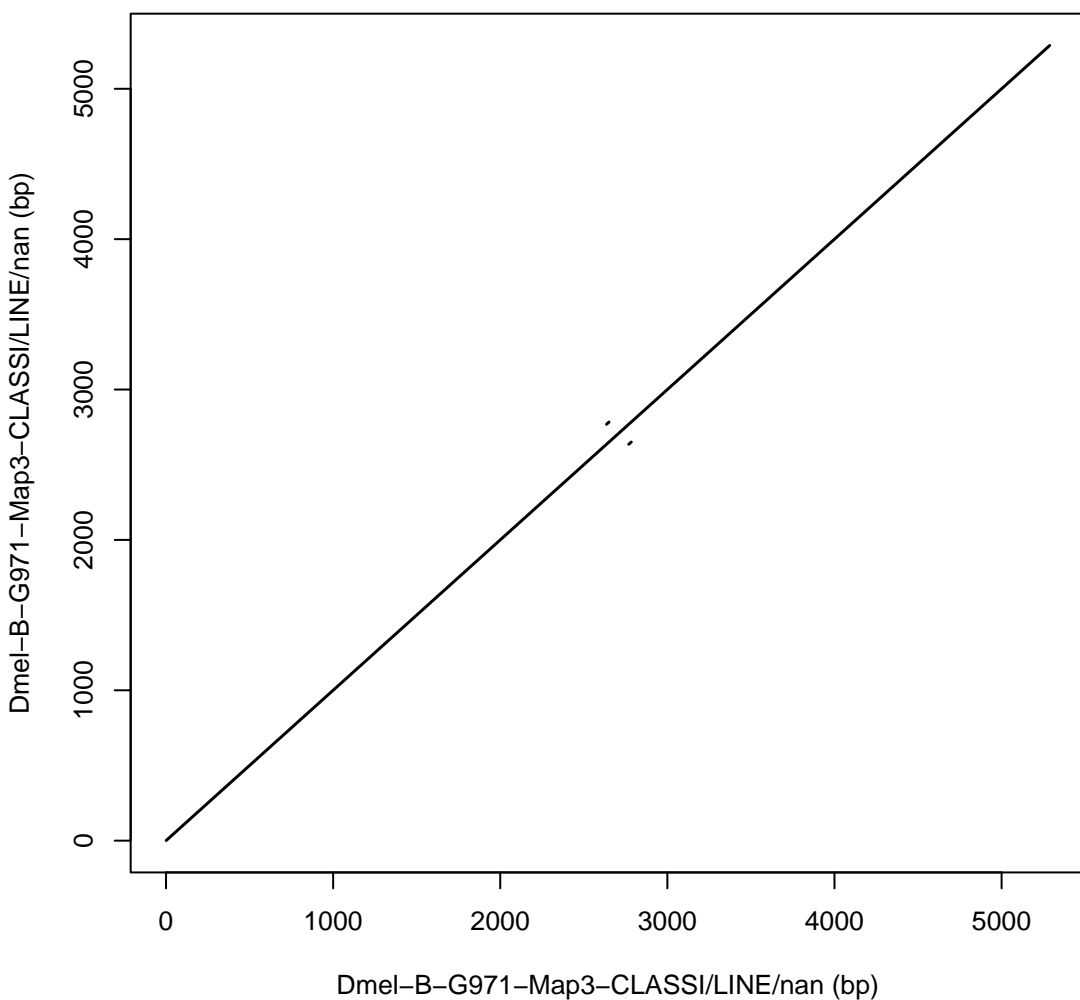
TE: Dmel-B-G971-Map3-CLASSI/LINE/nan
consensus size: 5288bp; fragments: 834; full length: 3 (>=4759.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

