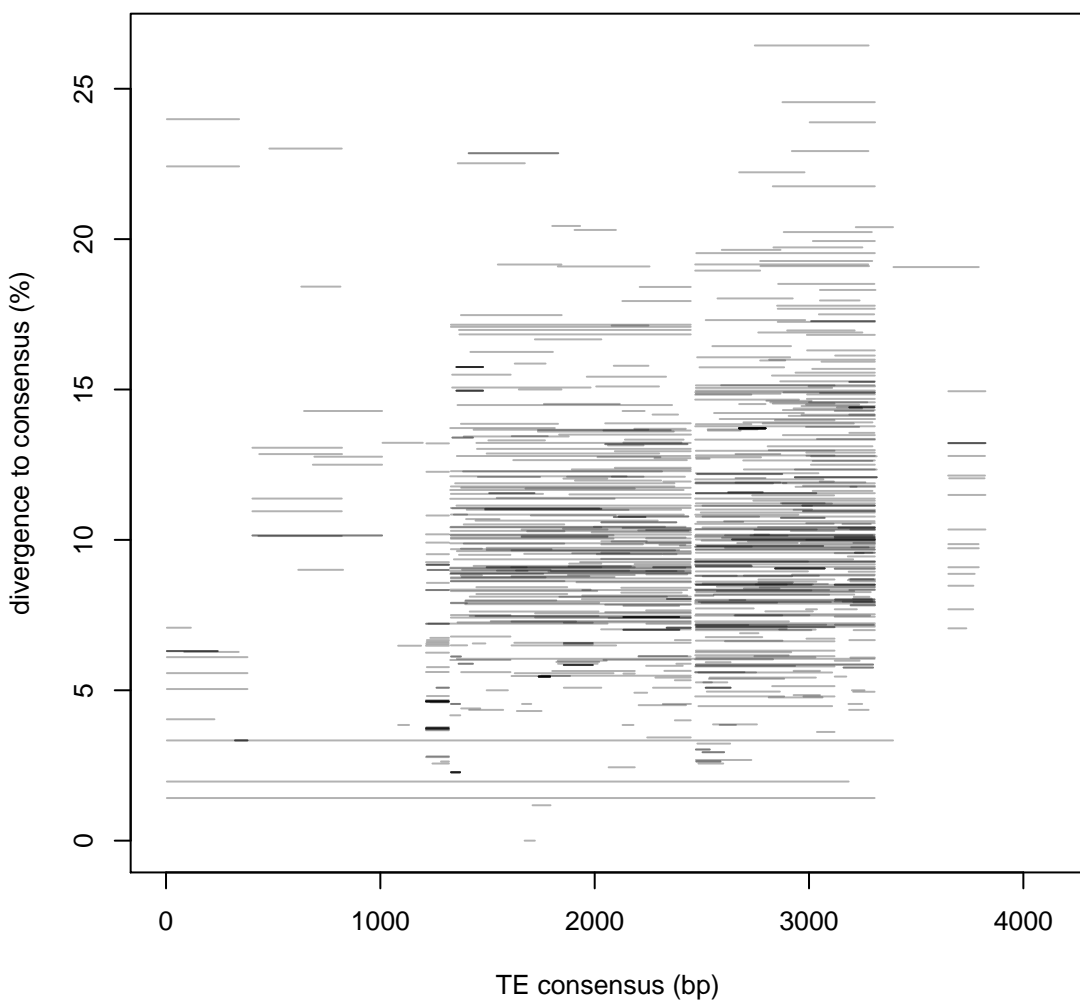
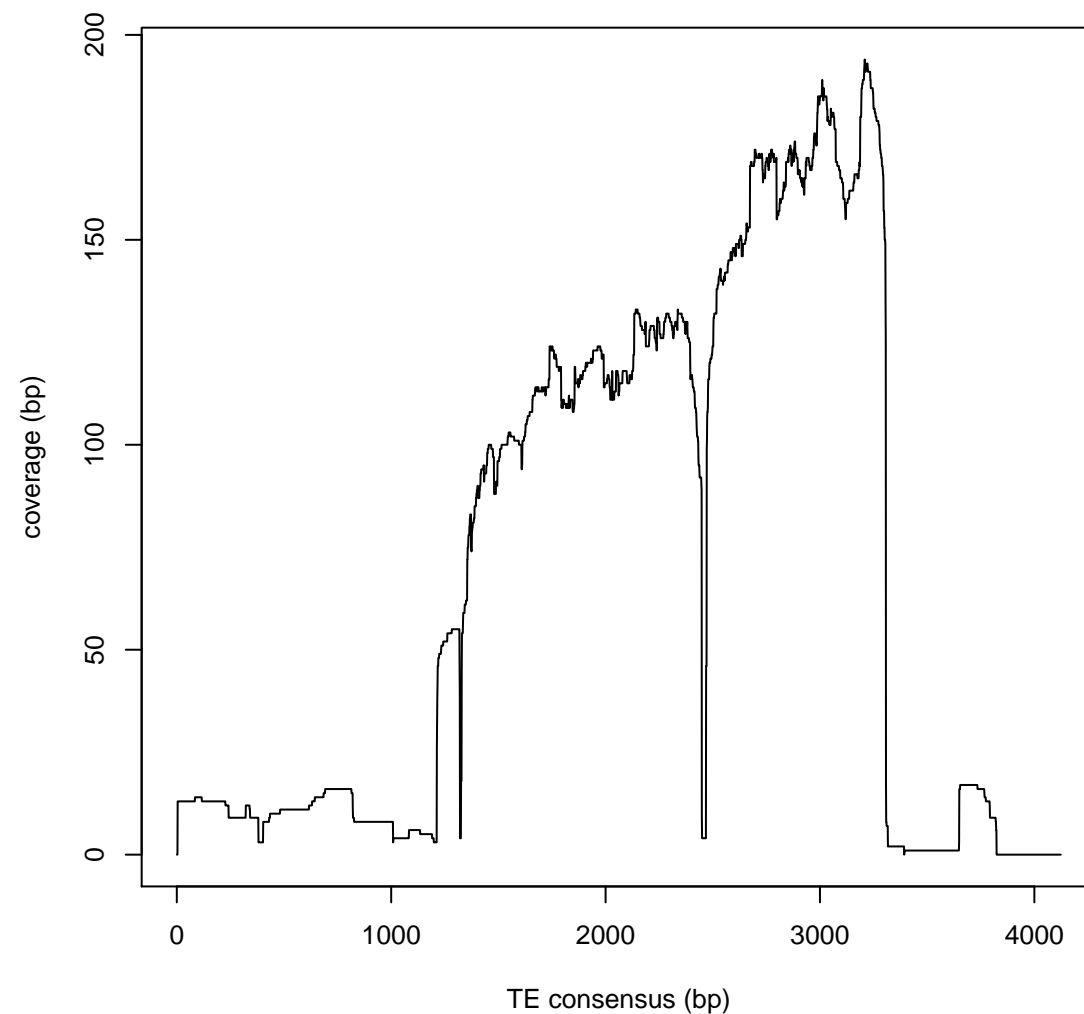


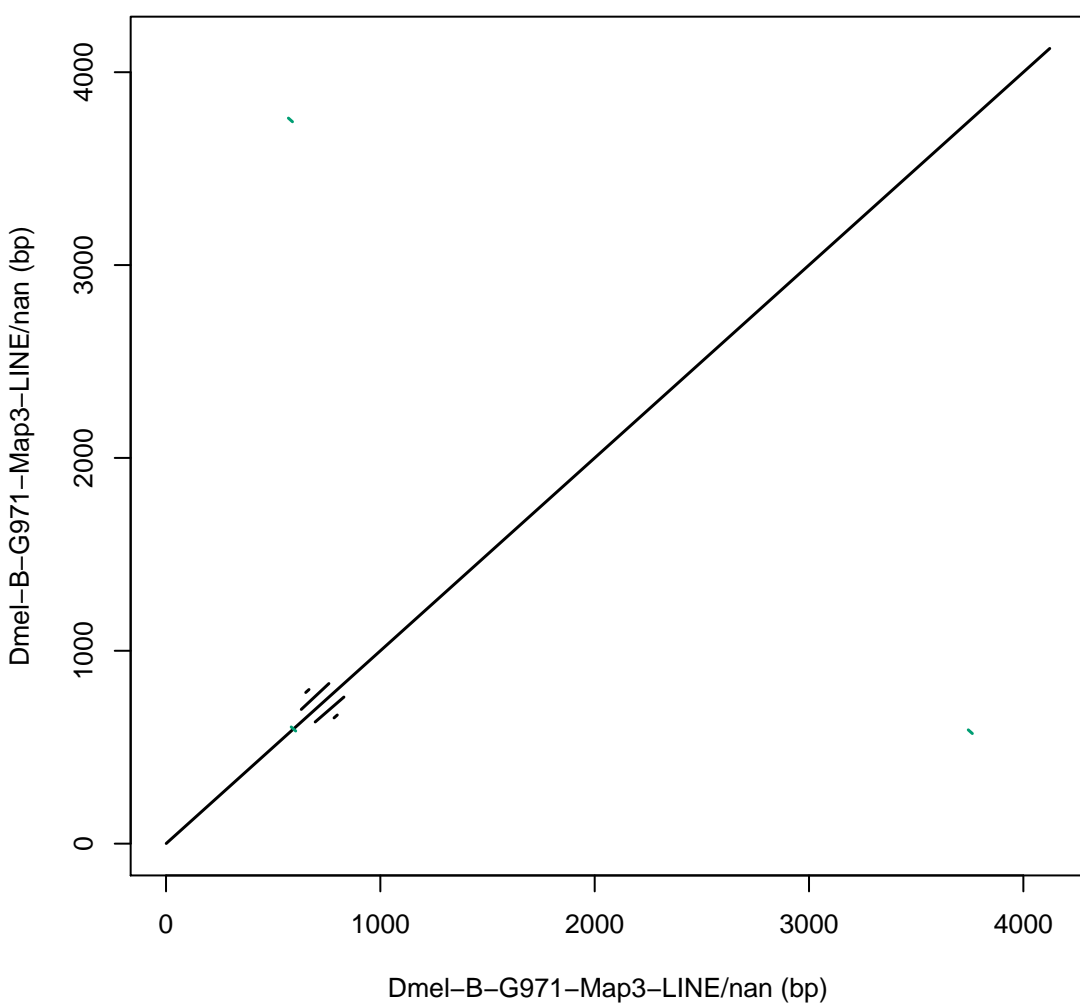
TE: Dmel-B-G971-Map3-LINE/nan  
consensus size: 4123bp; fragments: 749; full length: 0 (>=3710.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

