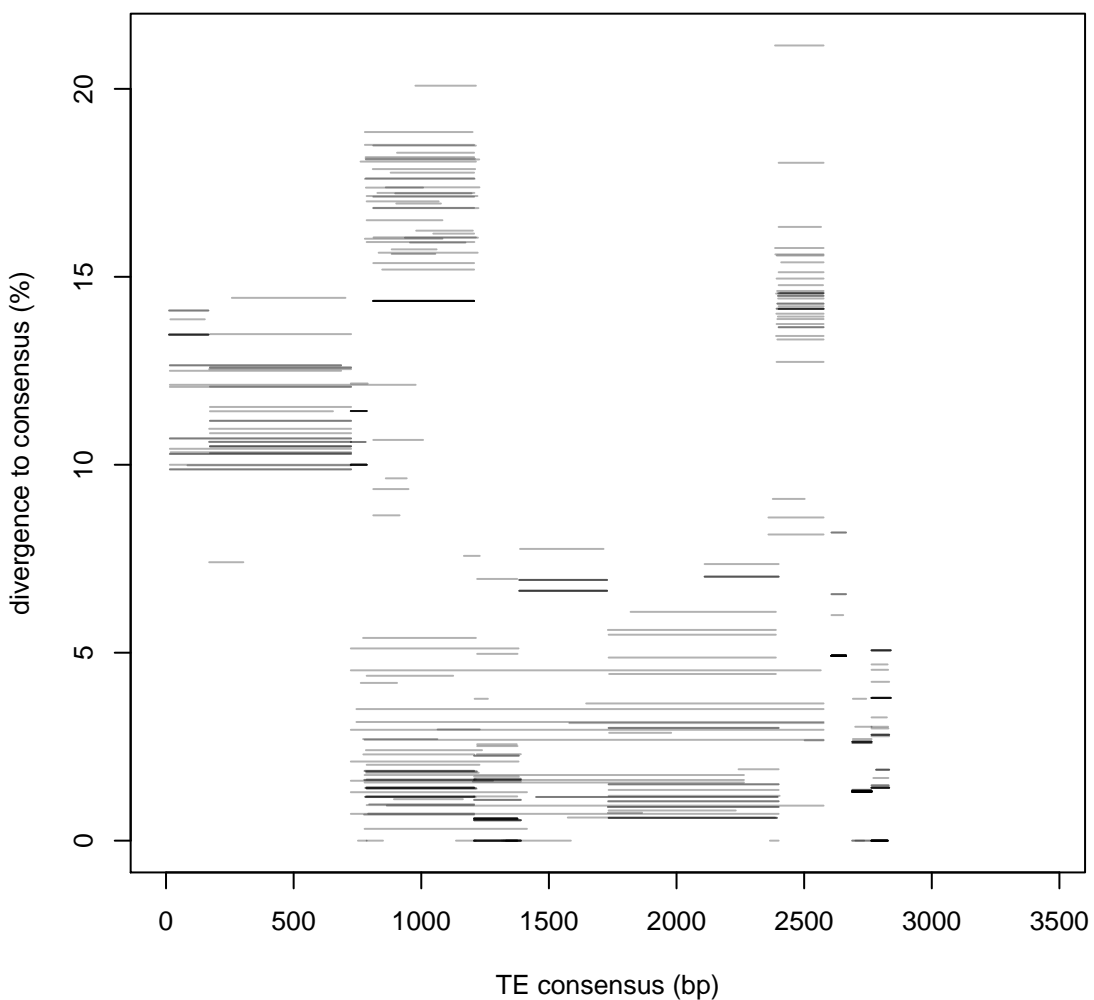
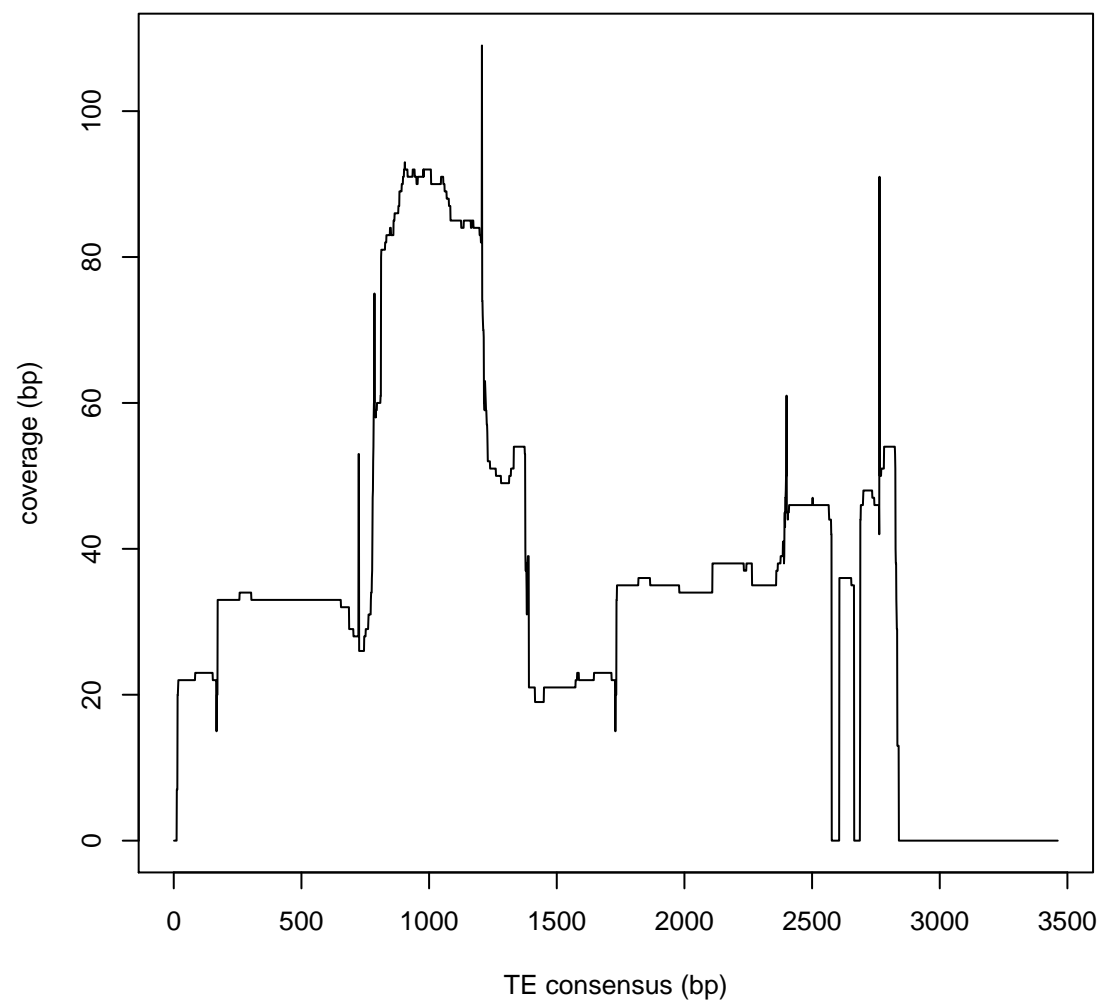


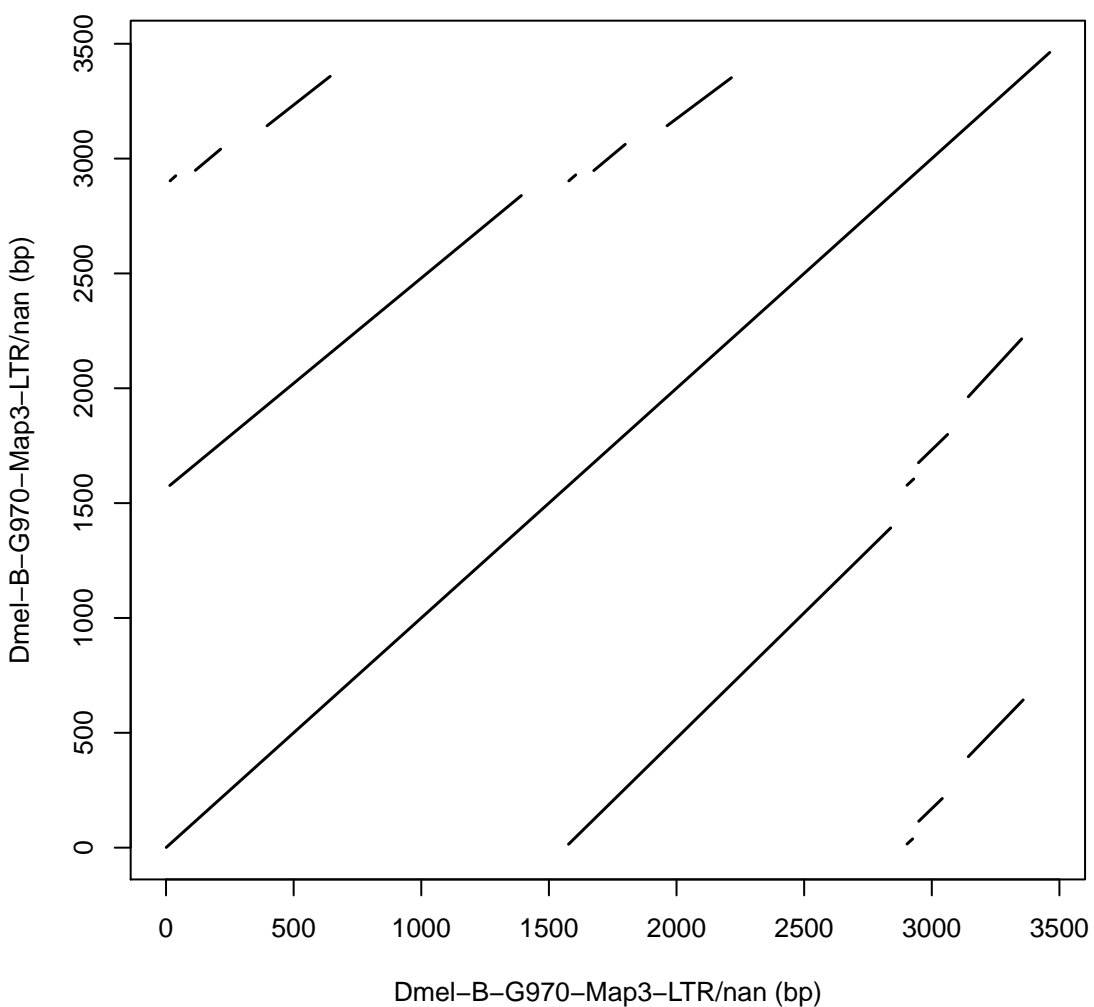
TE: Dmel-B-G970-Map3-LTR/nan  
consensus size: 3462bp; fragments: 420; full length: 0 ( $\geq 3115.8$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

