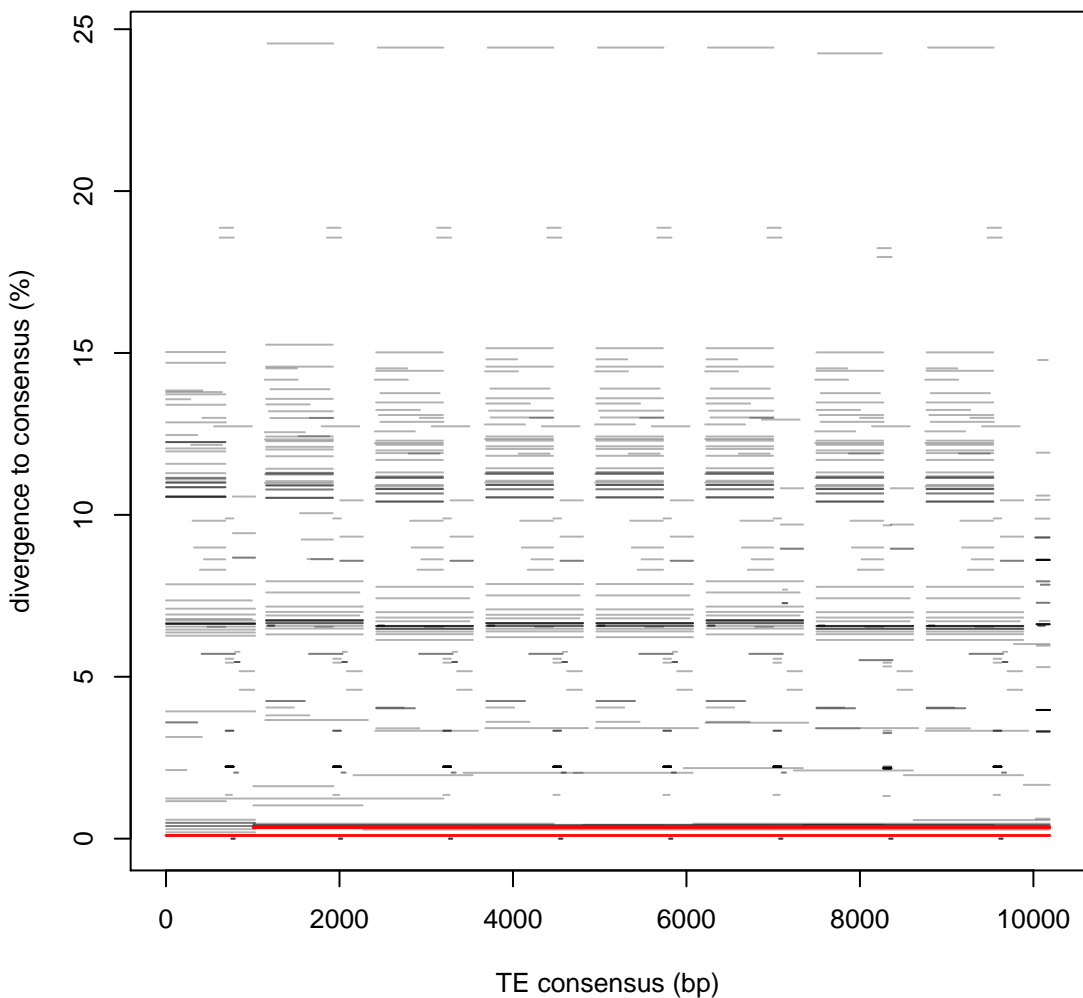
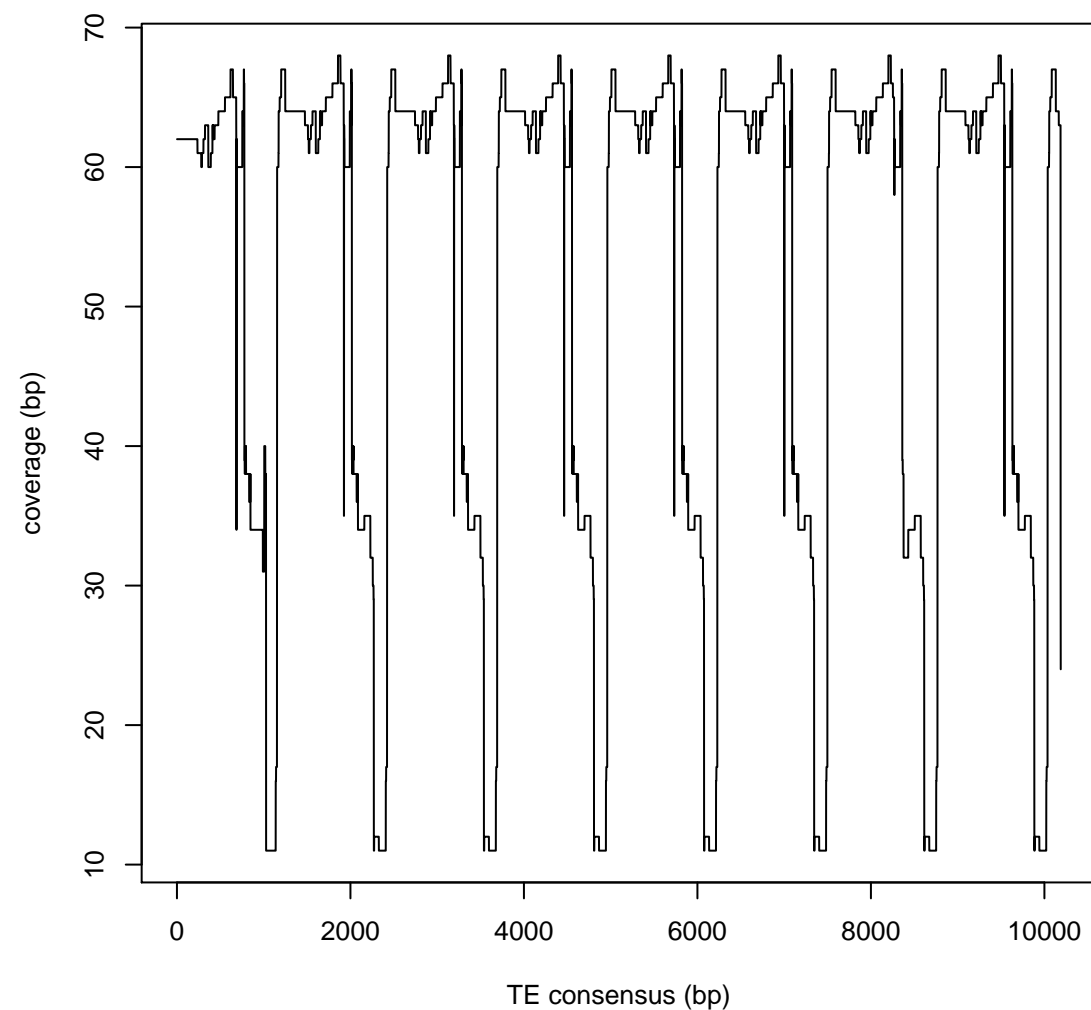


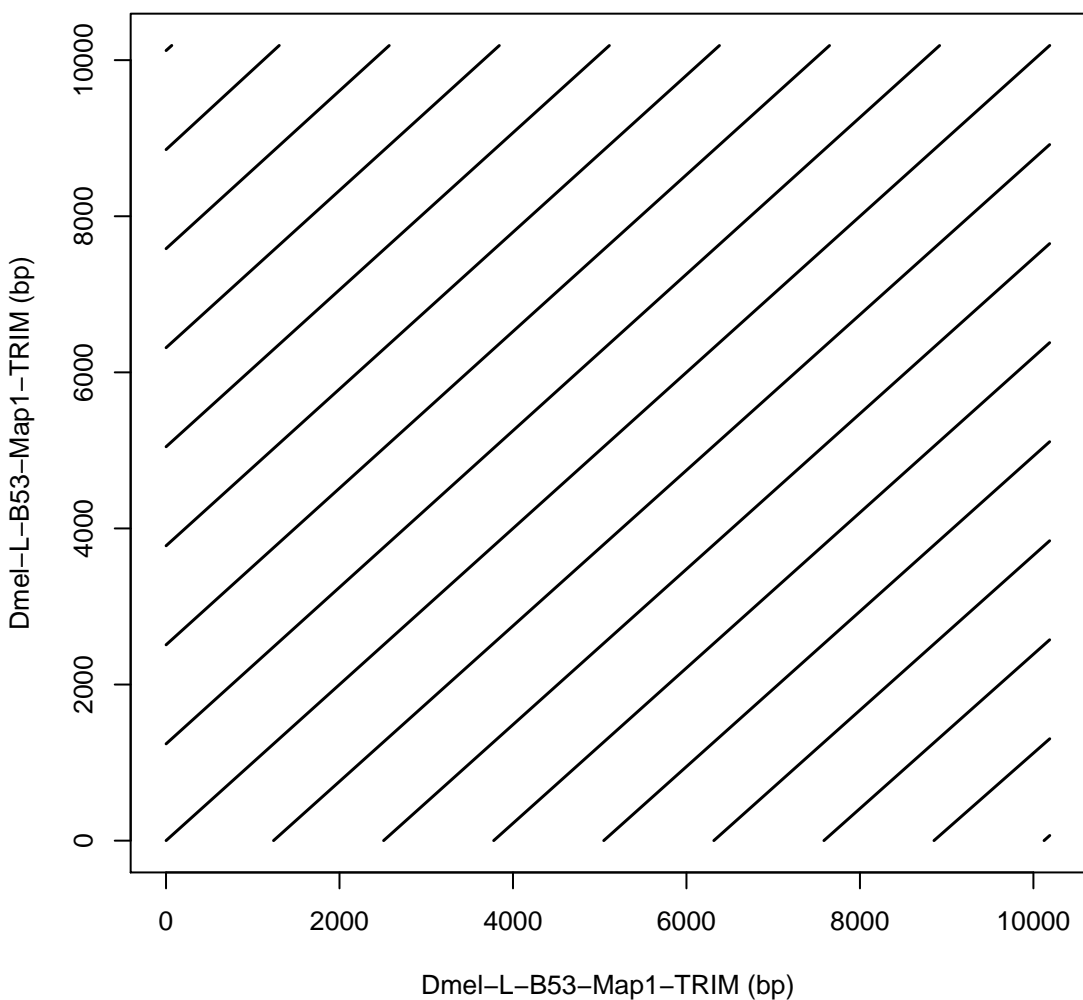
TE: Dmel-L-B53-Map1-TRIM
consensus size: 10188bp; fragments: 975; full length: 3 (>=9169.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

