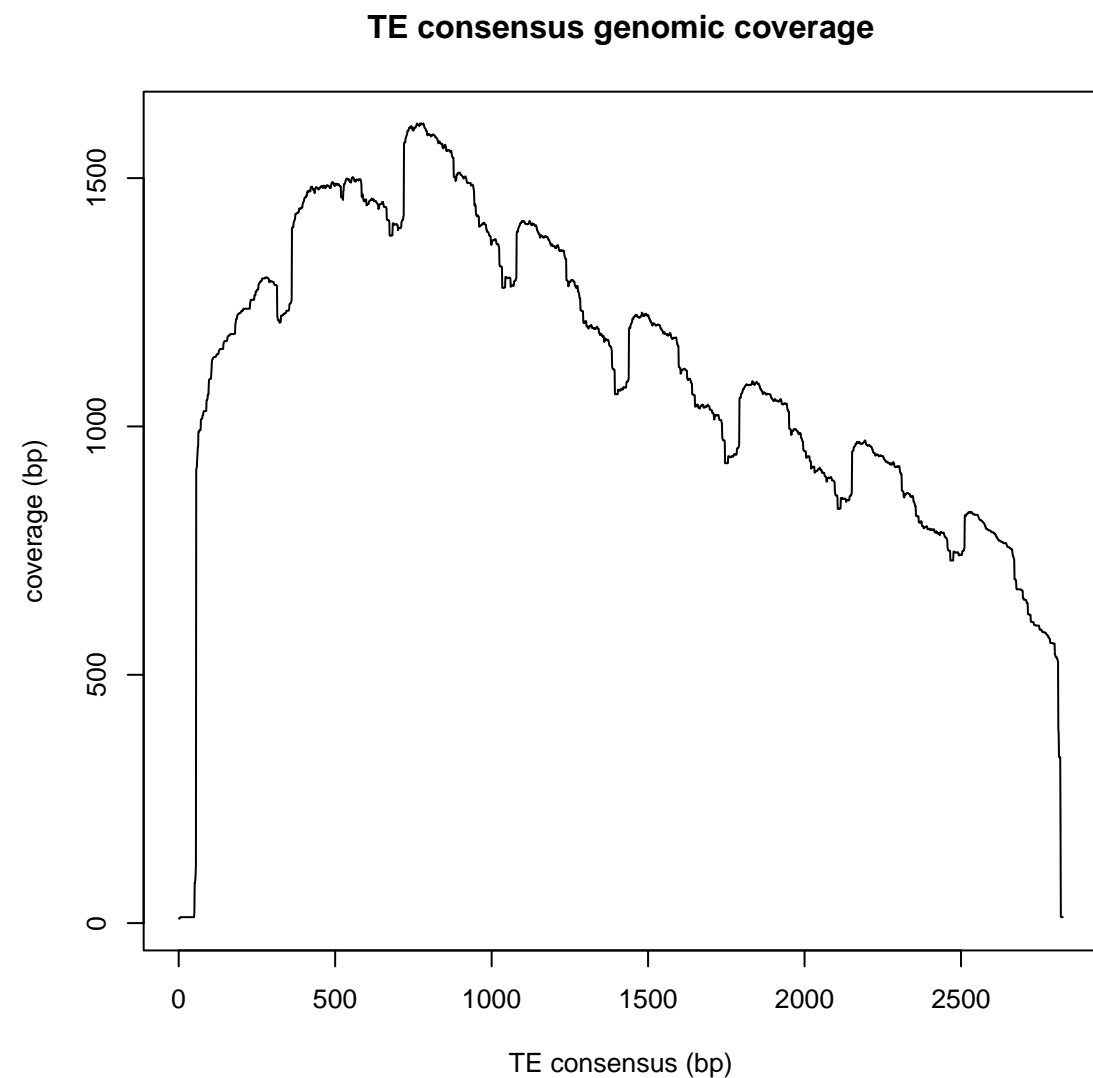
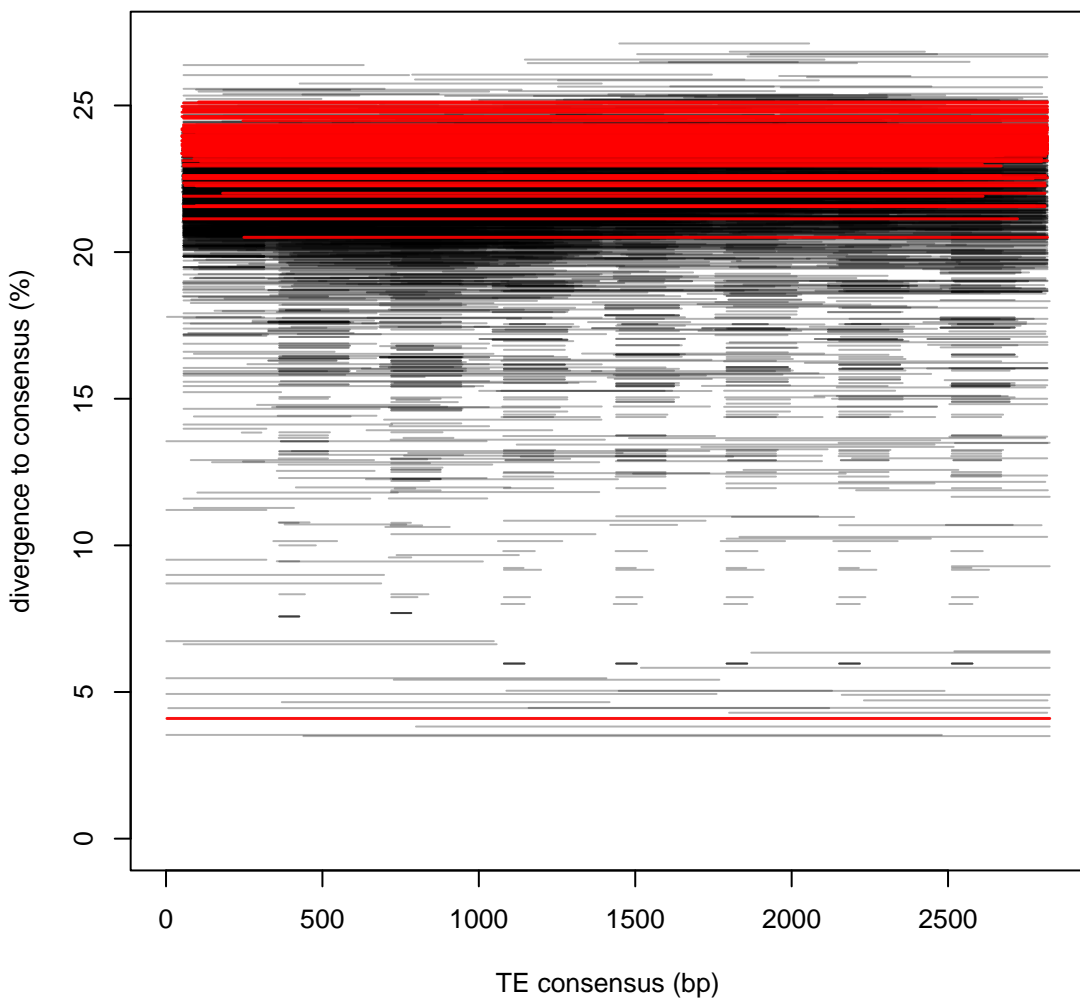
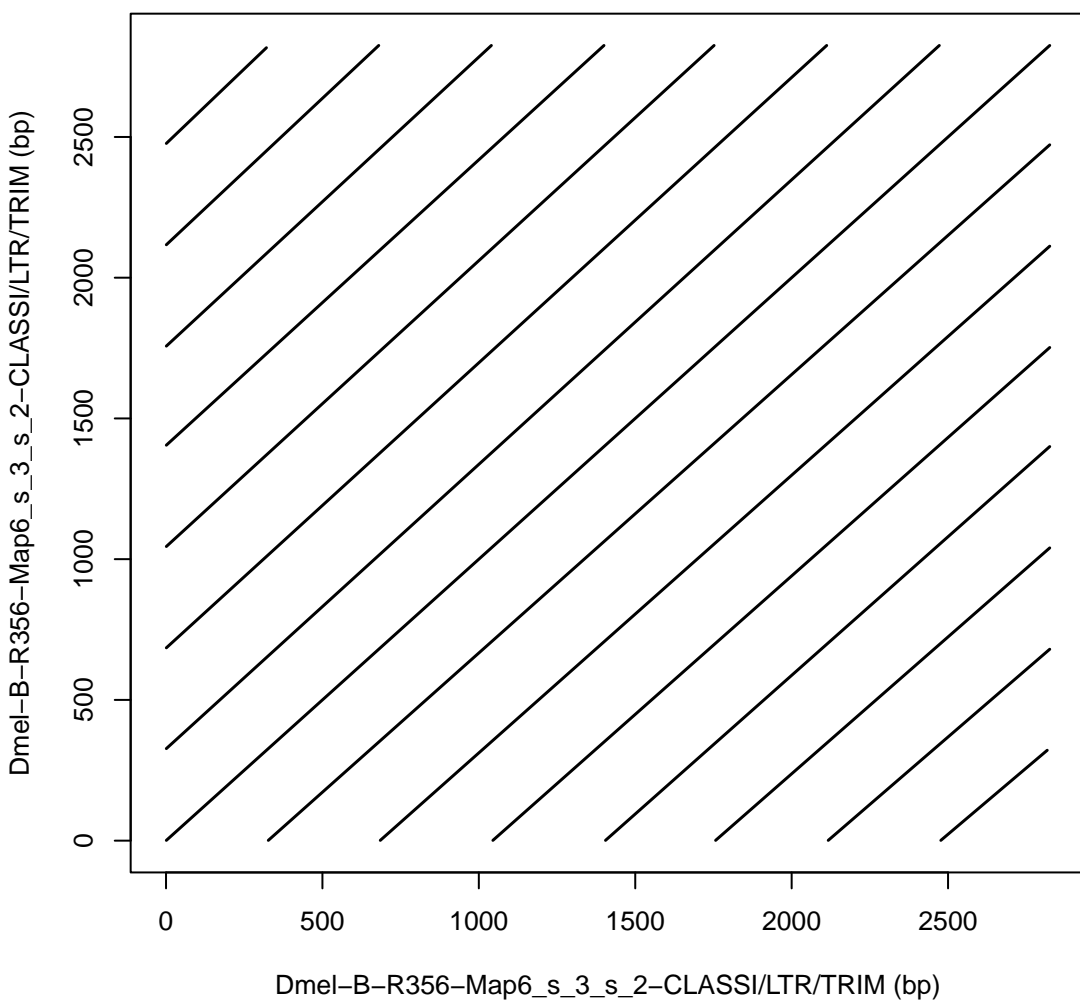


TE: Dmel-B-R356-Map6\_s\_3\_s\_2-CLASSI/LTR/TRIM  
consensus size: 2825bp; fragments: 3296; full length: 103 (>=2542.5bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

