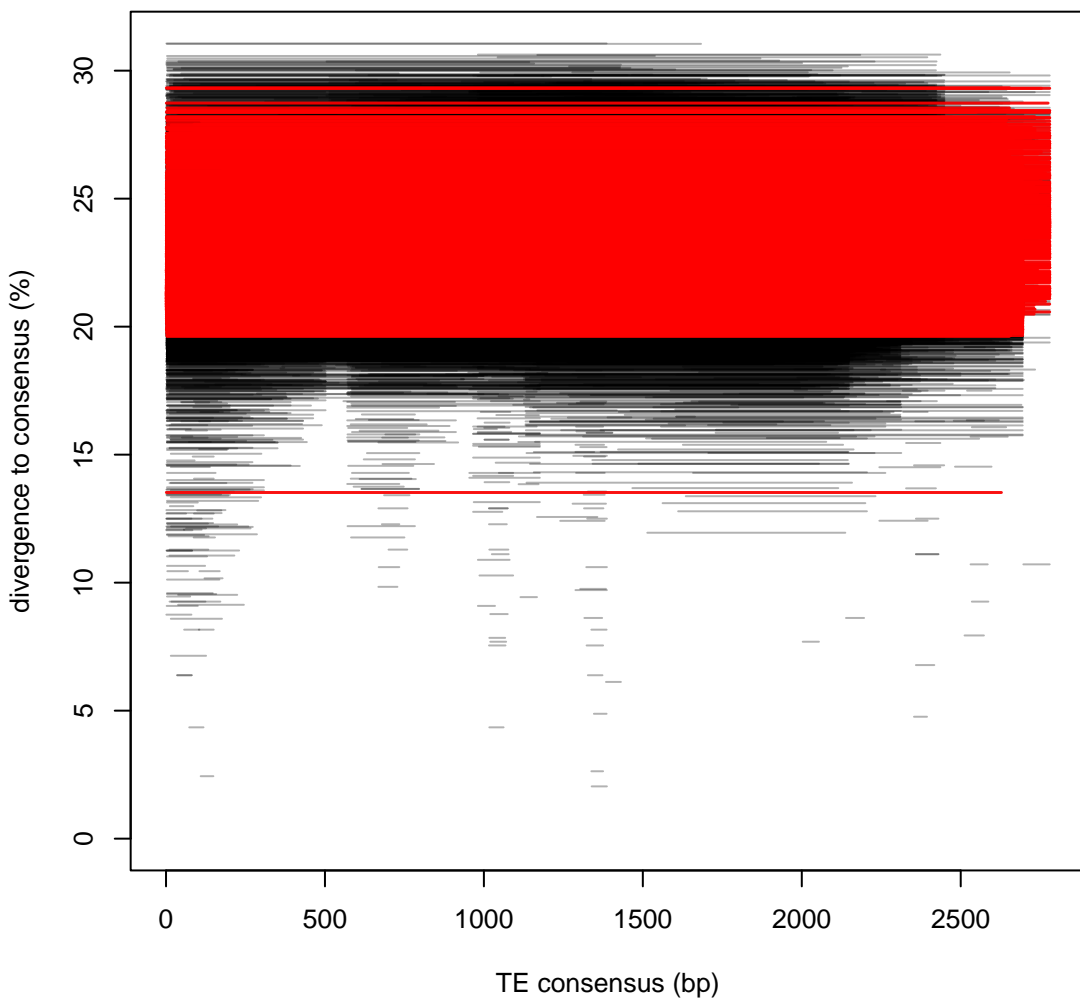
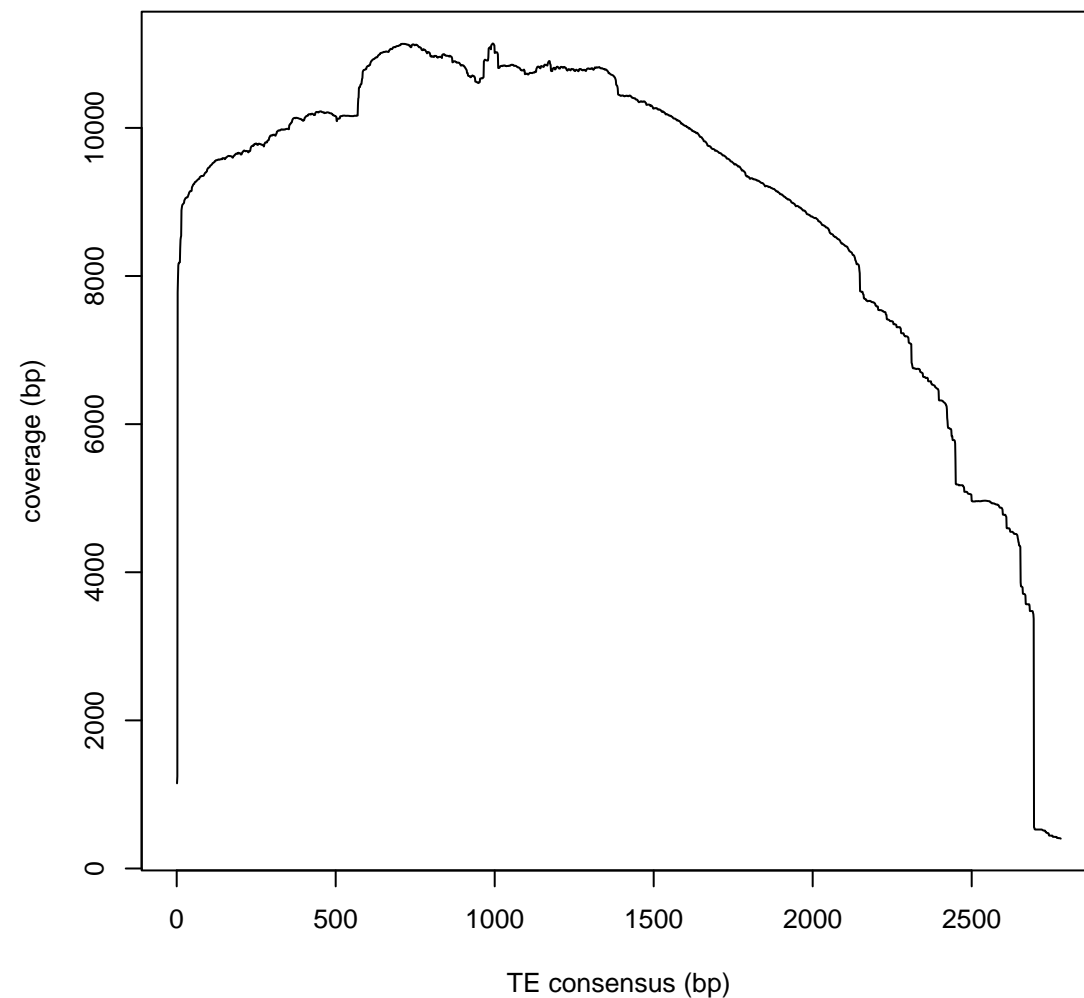


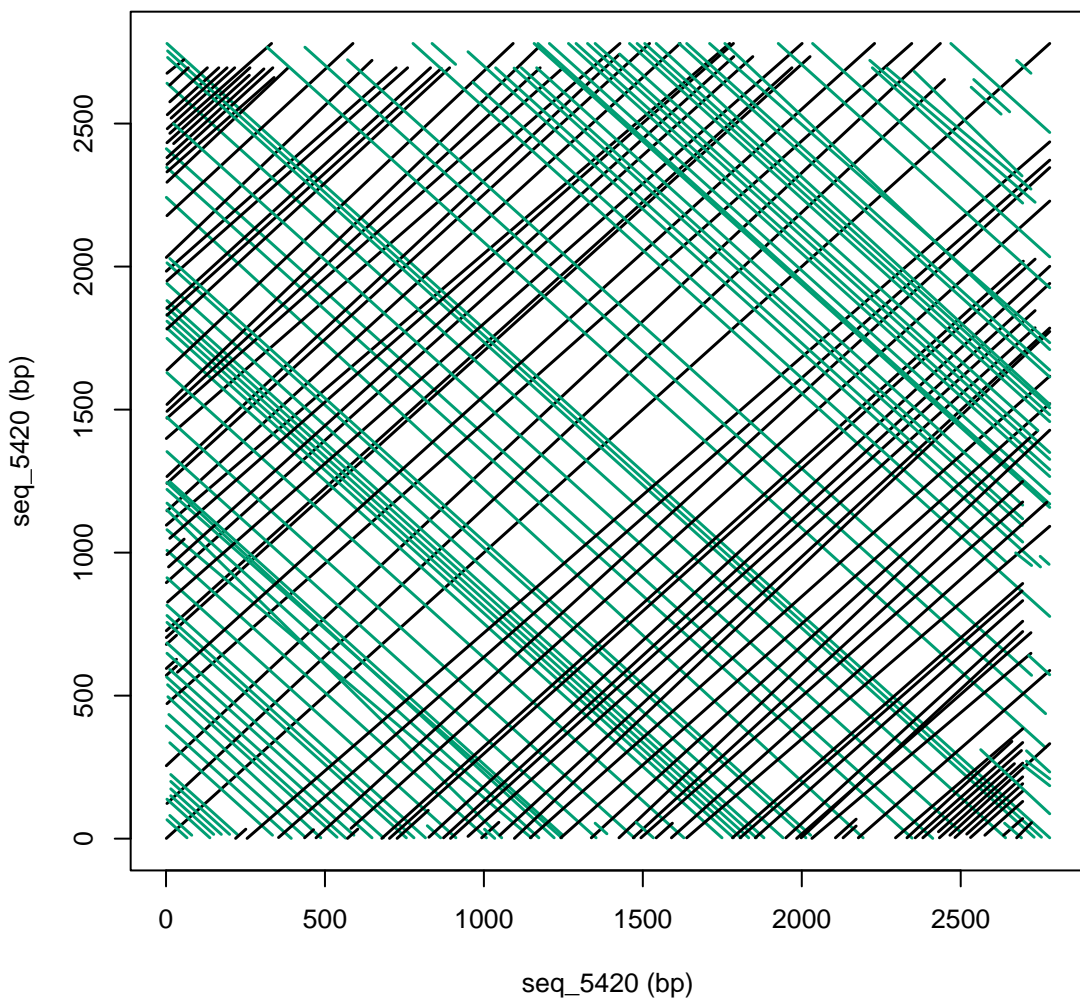
TE: seq\_5420  
consensus size: 2780bp; fragments: 14089; full length: 3580 (>=2502bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

