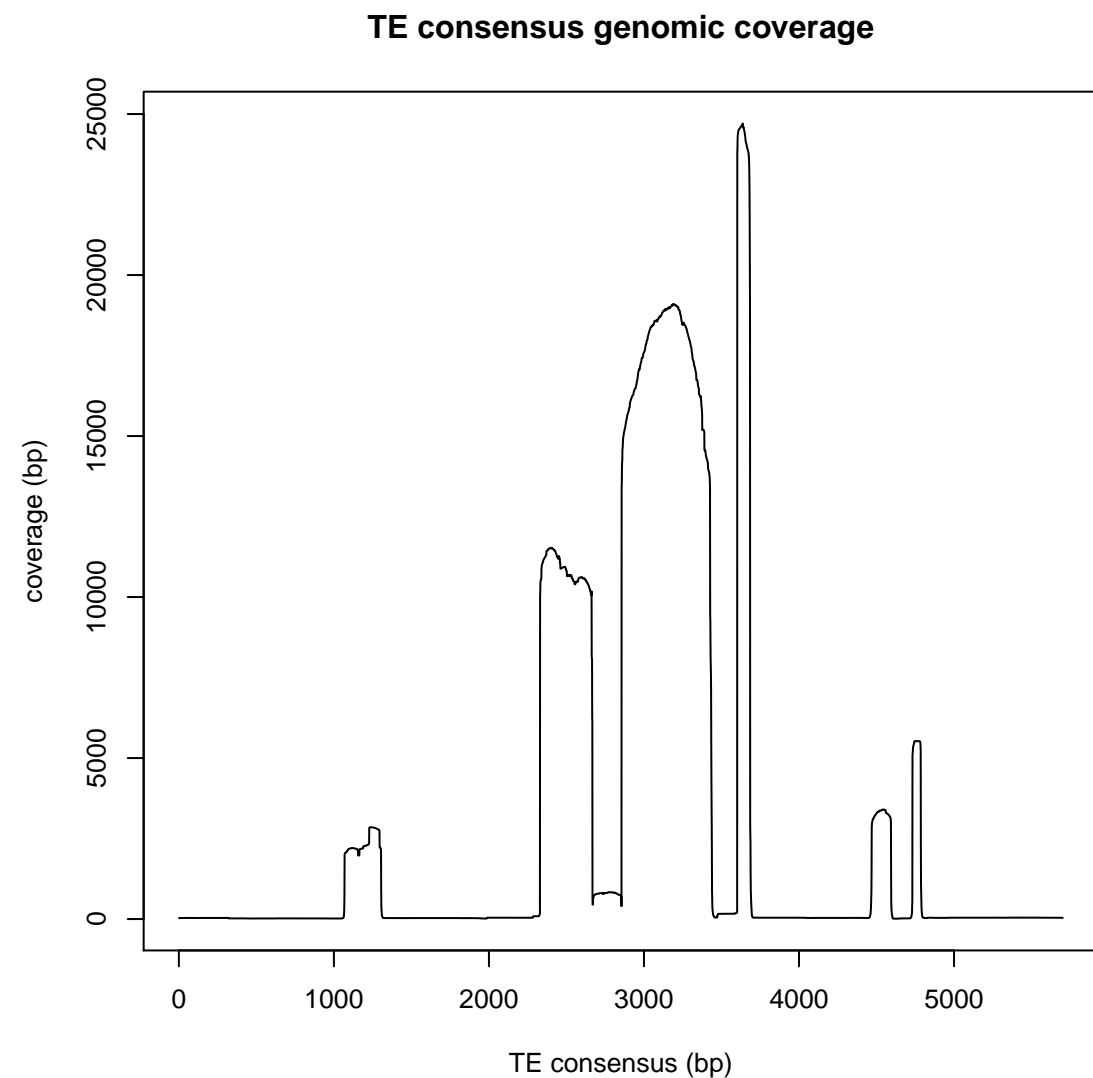
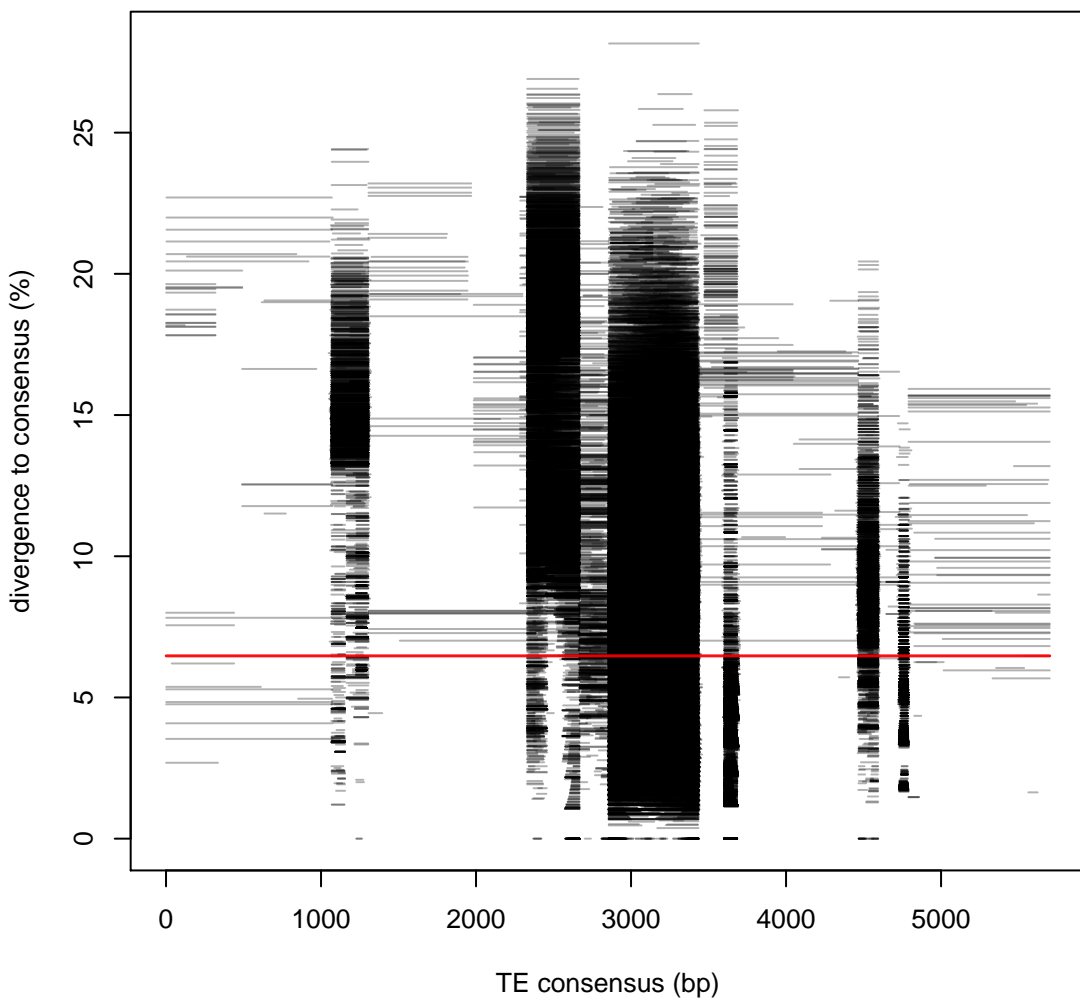
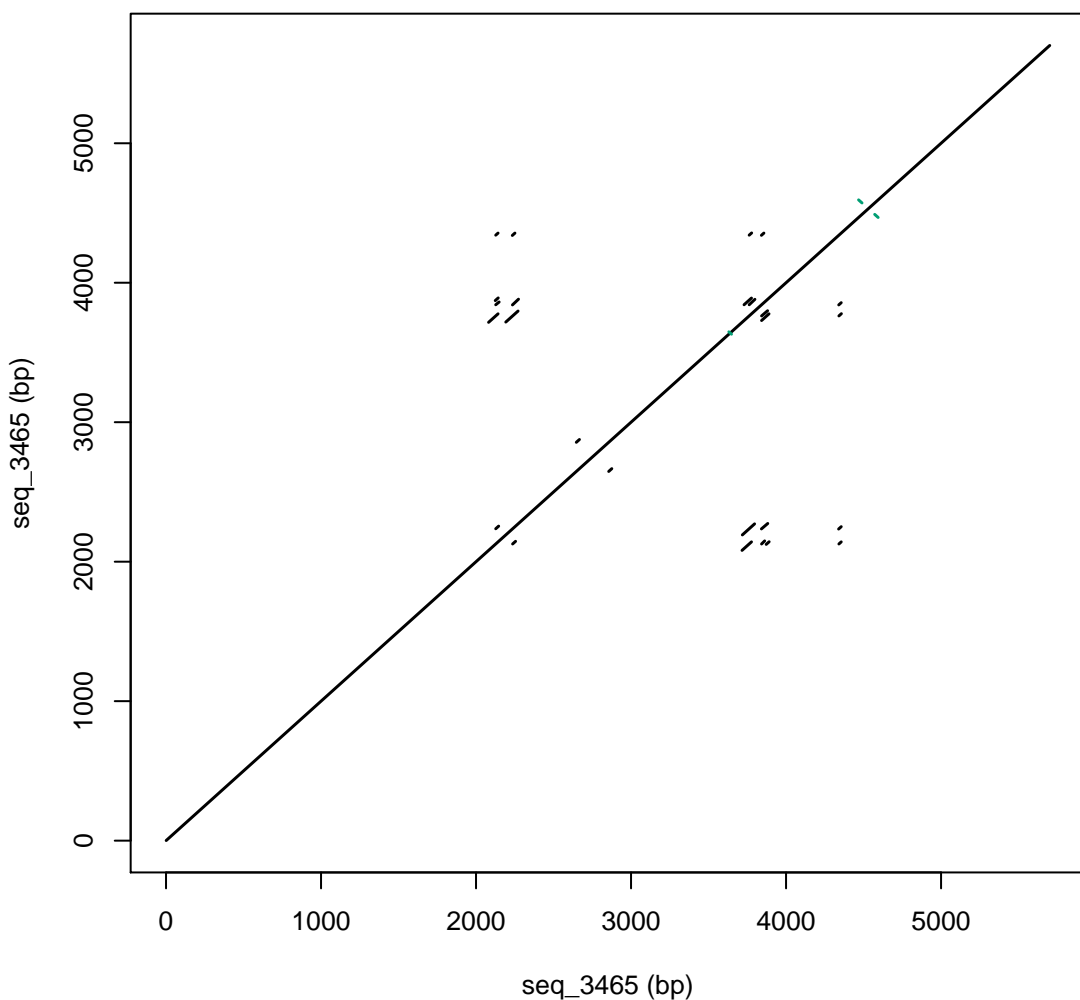


TE: seq\_3465  
consensus size: 5701bp; fragments: 73931; full length: 1 (>=5130.9bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

