TE: rnd-1\_family-200-LINE/I-Jockey TE consensus genomic coverage consensus size: 3778bp; fragments: 141; full length: 3 (>=3400.2bp) divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits UN-4755191\_Ath\_pol LINE/L1 rnd-1\_family-200-LINE/I-Jockey (bp) rnd-1\_family-200-LINE/I-Jockey (bp) rnd-1\_family-200-LINE/I-Jockey (bp)