TE: seq\_6392 consensus size: 2302bp; fragments: 5810; full length: 1 (>=2071.8bp) TE consensus genomic coverage divergence to consensus (%) coverage (bp) TE consensus (bp) TE consensus (bp) TE consensus self dotplot (blastn) TE consensus structure and protein hits Sola2-1\_BM\_\_tp TIR/Sola-2 seq\_6392 (bp) seq\_6392 (bp) seq\_6392 (bp)