TE divergence for Zoophthora_radicans_ATCC_208865.v1.divsum 2 · DNA DNA.PiggyBac LINE.Deceiver DNA.Sola.2 LINE.L1 DNA.Academ.H DNA.CMC.EnSpm DNA.TcMar.Ant1 LINE.L1.Tx1 DNA.Crypton.F DNA.TcMar.Fot1 LINE.Penelope DNA.Ginger DNA.TcMar.ISRm11 LINE.Tad1 DNA.Ginger.2 DNA.TcMar.Pogo LTR DNA.IS3EU DNA.TcMar.Sagan LTR.Copia DNA.Kolobok.H DNA.TcMar.Tc1 LTR.Gypsy DNA.MULE.MuDR DNA.TcMar.Tigger LTR.Unknown RC.Helitron DNA.MULE.MuDR. DNA.hAT.Ac DNA.Maverick DNA.hAT.Charlie Satellite DNA.Merlin DNA.hAT.hAT19 Simple_repeat DNA.P DNA.hAT.hATw Unknown rRNA DNA.P.Fungi LINE.CR1 DNA.PIF.Harbinger LINE.CRE

40

Percent of the genome

0 -

0

20

Kimura substitution level