

logo										
Father	..	AAGCAA	GTAGC	CTCCTCCTTTTAGTT	AAATGAGACTTCCTTGTGGGAAAGGGGAGCGTTTTGGTTCTTAATGAGAGCTTTCTTTTGCAGGGAAAAATCCTGGATGAGCTTTTGCAGA	116				
Mother	.	TAGCCAAT	GTTGG	CTCCTCCTTTTAGTT	AAATGAGACTTCCTTGTGGGAAAGGGGAGCGTTTTGGTTCTTAATGAGAGCTTTCTTTTGCAGGGAAAAATCCTGGATGAGCTTTTGCAGA	118				
Sister	..	AACCACT	GTTGC	CTCCTCCTTTTAGTT	AAATGAGACTTCCTTGTGGGAAAGGGGAGCGTTTTGGTTCTTAATGAGAGCTTTCTTTTGCAGGGAAAAATCCTGGATGAGCTTTTGCAGA	118				
Brother	TTTCTATAG	GAGGG	CTCCTCCTTTTAGTT	AAATGAGACTTCCTTGTGGGAAAGGGGAGCGTTTTGGTTCTTAATGAGAGCTTTCTTTTGCAGGGAAAAATCCTGGATGAGCTTTTGCAGA	119					
Proband	...	CCCGCAT	TCTG	CTCATCTATTTTCGCC	AAATGAGACTTCCTTGTGGGAAAGGGGAGCGTTTTGGTTCTTAATGAGAGCTTTCTTTTGCAGGGAAAAATCCTGGATGAGCTTTTGCAGA	117				
RefSeqGGAAAAATCCTGGATGAGCTTTTGCAGA	27				

logo											
Father	CCTGCGTTGACCCTGAAGACTGCCCGGTGTGTGAGGTGGCTGGCCGGCGTTTTGCCTCAAGAAAGAAAGTCACCTTGAATCCCAGTGACCCTGAGCACTGCCAGATTTGGTAAAAACAGAT										236
Mother	CCTGCGTTGACCCTGAAGACTGCCCGGTGTGTGAGGTGGCTGGCCGGCGTTTTGCCTCAAGAAAGAAAGTCACCTTGAATCCCAGTGACCCTGAGCACTGCCAGATTTGGTAAAAACAGAT										238
Sister	CCTGCGTTGACCCTGAAGACTGCCCGGTGTGTGAGGTGGCTGGCCGGCGTTTTGCCTCAAGAAAGAAAGTCACCTTGAATCCCAGTGACCCTGAGCACTGCCAGATTTGGTAAAAACAGAT										238
Brother	CCTGCGTTGACCCTGAAGACTGCCCGGTGTGTGAGGTGGCTGGCCGGCGTTTTGCCTCAAGAAAGAAAGTCACCTTGAATCCCAGTGACCCTGAGCACTGCCAGATTTGGTAAAAACAGAT										239
Proband	CCTGCGTTGACCCTGAAGACTGCCCGGTGTGTGAGGTGGCTGGCCGGCGTTTTGCCTCAAGAAAGAAAGTCACCTTGAATCCCAGTGACCCTGAGCACTGCCAGATTTGGTAAAAACAGAT										237
RefSeq	CCTGCGTTGACCCTGAAGACTGTCCAAGTGTGTGAGGTGGCTGGCCGGCGTTTTGCCTCAGGAAAGAAAGTCACCTTGAATCCCAGTGACCCTGAGCACTGCCAGATTTG.....										136

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Father	TCCTGGGTTGTTTGAAGTGACGAATCTTATTGCTTTCCAGGGGGCACATAAC.GTTAGGTGTTTAGTTATCCTAATTGGTGTTTTTGCGGTTTCATGG.GTTC AAGGCTAGGAGGAGGA	353								
Mother	TCCTGGGTTGTTTGAAGTGACGAATCTTATTGCTTTCTCCAGGGGGCAGCATAGTCCAGTGCTTCCTTTTACTATTATCTTGTTTTTGCGGGCCATG.TGGACAAATA.....	345								
Sister	TCCTGGGTTGTTTGAAGTGACGAATCTTATTGCTTTCCAGGGGGCACAAACGAGTCC...GCTTCCTTTTAGTAGTAACCTGCTGTGGCTCTTGATGATTGTCTAGATCCCTGAGCGAG	355								
Brother	TCCTGGGTTGTTTGAAGTGACGAATCTTATTGCTTTCCAGGGGGCAGAAA....CAGGGGAATGGCGAACGTTTCCTGGCGCAATGGAGGTAGAGGGCATTCTGAGGAAGAGGAGGAGGA	353								
Proband	TCCTGGGTTGTTTGAAGTGACGAATCTTATTGCTTTTCCAGGGGGCACAAAATGTTA....CCTCCAATTGCCAGTTGTCAAGACTACACAGCGCTA.TAGTGGTGCAAAACCGATAAAGA	352								
RefSeq	136								

logo		
Father	GTG.....	356
Mother	345
Sister	CTGGGGGG.....	363
Brother	GCGGGACAGAGACTCAGAGGAGGTCCGAGGTGAGCAGCAAAGCCTTGCCATTGCCAAGACCTATTCTGAGCTTCCTTGCTGGCCCCATTTTATTTGAAGCTATGGAAACAGAGAGAGGA	473
Proband	TGAGAATCCTCCCCTCTATTGACTCTCTCCTGGAAGGAGAGGGGTGACGG.....	402
RefSeq	136

non conserved

≥ 50% conserved