logo
Proband TGGCAAATTACCCTGCTAGCACGAGGCCTTGTTCTCAGGGGCCTGCTTCTCAAATGAAGGCGGATCTGCTTGAGGTTGGCATGGGGCCCAATGCCCACCGGGATCACAATGACCTTCTTC 120
RefSeq   0

logo
Proband TTCTTCAGGCCCTGGACGTAGCGGACAAAGTTCCGGGACATCCGTTGGGGCTCCTGGCTGGCCATCAGGAGCAGGGCGATGCGGAAGGCTTCAGGGCGG..TCGATCTTGCTGAAGATTT 238
RefSeq   ....GTGGCAGC..TCACAGCTATTG..TGGTGGGAAGGGAGGGGTGGTTGGTGGATGTCACAGCTTGGGCTTTATCTCCCCAGCAGTG 82

logo
Proband GGAACAGTGTGTATTTCAAGGCCTCGCTGGTGGAGGCCACCTGGCTGCCCGCATACTTCACCTGGCTGGCAATGCGCCGCAGCTCTGACGGTCGCTTCCGGTCCTTGAGCCGATGT..A 356
RefSeq   GGGACTCCACA.GCCCCTGGCGTACATAACAGCAAGACAGTCCGGAGCT...GTAGCAGACCTGATTGAGCCTTTGCAGCAGCTGAGAGCATGGCCTAGGTGGGCGGCACCATTGTCCA 198

logo
Proband GGCGTGGGAGCCGTCGTGGTACTCCACCACGGCCACGCGGACCCACTTCTGGGAGATGCGCAGCCGCTCCATCATGTCCACCACAAAGGCCTTCAGCACTTCAAACTCAGCCTCGGACAG 476
RefSeq   GCAGCTGAGTTTCCCAGGGACCTTGGAGATAGCCGCA.GCCCTCATTTGCAGGGGAAG.. 255

logo
Proband CCTGGAGGAGCCATCCAGCAGGAAGACCAGGTCCAGTAGCCTGCTGCAGTAGAAATCGTGCAACGGCGGTTCCGAGATGTCCTCCACATACAGAGTGGTGGGGCTCACCGGGGCATCTGT 596
RefSeq   255

logo
Proband GGGAGGCACCACCAGGCCTCCCGGCTCCTGGCAGGCTTCACAGGTGAGGTTGACAACATCACAGTGGCTGCAGAAAGAGCGAAGAAATTAAAATGGTTCAGAAGAACCTGTGGACACT 716
RefSeq   255

logo
Proband TCTGAGCCCTACAGTGTACGACGACTTCCATATTCCCACAGAATCTCCTCTGTTCCACCTGAACTTGAGATCCCATGGACCATTCCACATCCAAGTGAGATGTCACTTGTTTAACATCTG 836
RefSeq   255

logo
Proband TCCCCAAATAGCATGCCCCCCCCACCTTCAAAACACACACATAGCCGACTTCCTTTGATTCTAGAAACCAAAGCTTTAGCCCATCCTAGGATATGAAAAAATATACTCGTTGTTCTAGGC 956
RefSeq   255

```
logo
        CATCCACACAACACCTCAAAGCTA
Proband CATCCACACAACACCTCAAAGCTA   980
RefSeq  .......................   255


        ⊠  non conserved
        ☒  ≥ 50% conserved
```