1 AGTTCTAAAGTCCCCACGCACCCGGACTCAGAATCTCCTCAGACGCCGAGATGCGG 1 AGTTTTAAAGTTTTTACGTATTTATTCGGATTTAGAATTTTTTTAGACGTCGAGATGCGG 61 GTCACGGCGCCCGAACCCTCCTGCTGCTGCTGGGGGGGCAGTGGCCCTGACCGAGACC 61 GTTACGGCGTTTCGAATTTTTTTTTTTTTTTTGTGGGGGGTAGTGGTTTTGATCGAGATT ||||:++|:|:::|:|::|:||||||||||||:||:|:++::|||||::++|::++|:++++|| 121 TGGGTCGGTTTTTATTTTATGAGGTATTTTTATATCGTTATGTTTCGGTTCGGTCGCGGG 181 GAGCCCCGCTTCATCACCGTGGGCTACGTGGACGACACGCTGTTCGTGAGGTTCGACAGC |||:::++:||:||:||:++||||:||++||||++:||||++:||||++|||||++|:||+ 181 GAGTTTCGTTTATTATCGTGGGTTACGTGGACGATACGTTGTTCGTGAGGTTCGATAGC 241 GACGCCACGAGTCCGAGGAAGGAGCCGCGGGCGCCATGGATAGAGCAGGAGGGGCCGGAG 241 GACGTTACGAGTTCGAGGAAGGAGTCGCGGGCGTTATGGATAGAGTAGGAGGGGTCGGAG 301 TATTGGGATCGGGAGATATAGATTTTTAAGATTAATATATAGATTTATCGAGAGAATTTG 361 CGCACCGCGCTCCGCTACTACAACCAGAGCGAGGCCGGGTCTCACATCATCCAGAGGATG 361 CGTATCGCGTTTCGTTATTATAATTAGAGCGAGGTCGGGTTTTATATTATTTAGAGGATG 421 TACGGCTGCGACGTGGGGCCGGACGGCGCCTCCTCCGCGGGTATGACCAGGACGCCTAC 421 TACGGTTGCGACGTGGGGTCGGACGGGCGTTTTTTTCGCGGGTATGATTAGGACGTTTAC 481 GACGGCAAGGATTACATCGCCCTGAACGAGGACCTGAGCTCCTGGACCGCGGCGGACACC +|++|:|||||:++++|++||:||| 481 GACGGTAAGGATTATATCGTTTTGAACGAGGATTTGAGTTTTTGGATCGCGGCGGATATC 541 GCGGCTCAGATCACCCAGCGCAAGTGGGAGGCGGCCCGTGTGGCGGAGCAGAGCCC 541 GCGGTTTAGATTATTTAGCGTAAGTGGGAGGCGGTTCGTGTGGCGGAGTAGGATAGAGTT 601 TACCTGGAGGGCCTGTGCGTGGAGTCGCTCCGCAGATACCTGGAGAACGGGAAGGAGACG 601 TATTTGGAGGGTTTGTGCGTGGAGTCGTTTCGTAGATATTTGGAGAACGGGAAGGAGACG 721 GTCACCCTGAGGTGCTGGGCCCTGGGCTTCTACCCTGCGGAGATCACACTGACCTGGCAG

721 GTTATTTTGAGGTGTTGGGTTTTGGGTTTTATTTTGCGGAGATTATATTGATTTGGTAG

- 841 AGAACCTTCCAGAAGTGGGCAGCTGTGGTGGTGCCTTCTGGAGAAGAGCAGAGATACACA

- 961 TAGTTTATCGTTGTGGGTATTGTTGGTTTGGTTGTTTTAGTAGTTGTGGTT

- 1081 TACTCTCAGGCTGCGTGCAGCGACAGTGCCCAGGGCTCTGATGTGTCTCTCACAGCTTGA
- 1081 TATTTTTAGGTTGCGTGTAGCGATAGTGTTTTAGGGTTTTGATGTGTTTTTATAGTTTGA
- 1141 AAAGCCTGAGACAGCTGTCTTGTGAGGGACTGAGATGCAGGATTTCTTCACGCCTCCCCT
- 1201 TTGTGACTTCAAGAGCCTCTGGCATCTCTTTCTGCAAAGGCACCTGAATGTGTCTGCGTC
- 1201 TTGTGATTTTAAGAGTTTTTGGTATTTTTTTTTTTTTAAAGGTATTTGAATGTGTTTGCGTT
- 1261 CCTGTTAGCATAATGTGAGGAGGTGGAGAGACAGCCCACCCCGTGTCCACCGTGACCCC
- ::|||||:||:||++|||::::
 1261 TTTGTTAGTATAATGTGAGGAGGTGGAGAGATAGTTTATTTTCGTGTTTATCGTGATTTT
- 1321 TGTTCCCATGCTGACTTGTGTTTCCTCCCCAGTCATCTTTCCTGTTCCAGAGAGGTGGGG

- 1441 TACTGAAAATAAGAATCTGAATATAAATTTGTTTTCTCAAATATTTGCTATGAGAGGTTG
- 1501 ATGGATTAATTAAATAAGTCAATTCCTGGAATTTGAGAGAGCAAATAAAGACCTGAGAAC

1501 ATGGATTAATTAAATAAGTTAATTTTTGGAATTTGAGAGAGTAAATAAAGATTTGAGAAT

1561 CTTCCAGAA

:||::|||

1561 TTTTTAGAA