

1 10 20 30 40 50 60 70 80 90 100 110 120 130
ATGCTCGGGCCCGCCACATTCCTGCCGCTGACCGGGCGCTCTTGCTGCTTATACCTGGAGGGCGAGCGCAGTTCCTCCCGCCAGTGCCTGACCCCGAGGCTCTCCGAGTGGCCAGTGTGCCTGGCCCTCTTCC

140 150 160 170 180 190 200 210 220 230 240 250 260 270
CTGCGCTGACACCGGACCCAGCAGACAGATGCGGGGCTTCCGTGGGCCGGGGCGTTGTGCGCCTCTGCAGGTGGATGCCGACCCACACGGGCCCCAGTATCCACACGATGGAGTGGATGACCGGGAGCAATGGCC

280 290 300 310 320 330 340 350 360 370 380 390 400
GACCCGCTTCTTCAACAACCTCTTGCCTGTGTGCAGAAAACCTCTCTGGCTACGACTGTGGGTCTCTGCAAAACCAGGCTGGGTCTGGAGTCAACTGCAACCAGAGGGTACTGGCAGTCAGGCGGAACATCTTGGATCTG

410 420 430 440 450 460 470 480 490 500 510 520 530 540
ACTGCACAGGAGAGGGCGTCGCTTTATTGCTGCCCTGGACTTGCCAAAAGAACCACACATCTCACTACGTCATCGCTCGGGAGATATGCAGAGATCATGGGACCCGATGGCAACAGCACCCAGTTTGAGAATG

550 560 570 580 590 600 610 620 630 640 650 660 670 680
TGTCGATTTACAACCTCTTTGTATGGACCCATTACTACTCCATTGGCAAGACTTTTCTTGTTGCCGTTCTGGGAGAGTTTTGGTGGCATAGACTTCTCCACGAGGGCCCTGCATTTGTTACCTGGCACAGGTACCA

690 700 710 720 730 740 750 760 770 780 790 800 810
CCTGCTACAACCTGAAAGGGATATGCAGGAGATGCTTCAAGATCCCACCTTTGCTCTTCCATATTGGAATTTTGTATTGGTGGCAATGAATGCGACATCTGCACTGACGACTTCATGGGTGCTCGGAGCAACTTC

820 830 840 850 860 870 880 890 900 910 920 930 940 950
GATTCATTCTTTTGAAGTTCCAACCTGTTTTTTCCCAATGGCGCGTTCTTTGCGAAAGCTTGGAAAGATTATGACACACTAGGAAACAATATGTAAACAGCACAGAGGGTGGTCCAAATCGAAGGAACCCGGAGGCA

960 970 980 990 1,000 1,010 1,020 1,030 1,040 1,050 1,060 1,070 1,080
ATGTAGCCAGGCCGATGGTTTCAGCGCCTCCCTGAACCACAGGATGTTGCCCTTTGCCCTTGGAAAGTTGGTTTATTGACACGCCACCTTCTATTCCAACCTCGTCTGAAAGCTTTGAAATACAGTTGAAGGATATAG

1,090 1,100 1,110 1,120 1,130 1,140 1,150 1,160 1,170 1,180 1,190 1,200 1,210 1,220
TGAGCCTTCTGGAAAATATGATCCTTCAGTTTGAAGCCTCCACAATTTGGCGCACCTGTTCTGAAATGGAACTGGAGGACAACTCACGTGTCCCTAATGACCCTATATTTGTTCTTCTGCACACGTTTACTGAT

1,230 1,240 1,250 1,260 1,270 1,280 1,290 1,300 1,310 1,320 1,330 1,340 1,350 1,360
GCAGTATTTGACGAATGGCTGAGGAGACATAATGCTGATATTTTCGCTTTACCCACTGGAGAATGCCCTATTGGACACAACAGACAGTACAACATGGTGCCTTCTGGCCCCAGTTTCCAACAATGAGATGTTTG

1,370 1,380 1,390 1,400 1,410 1,420 1,430 1,440 1,450 1,460 1,470 1,480 1,490
TTACTGCACCGGAGAGCCTCGGATATTCGATGAAGTCCAGTGGCCTAGTCGTGCTTTGAATTTACGGAGATCATAACCATCGCTGTGGTGGCAGCTTTGTGCTGGTTGCTGTGATCTTCGCTGCTGCTTCTTG

1,500 1,510 1,520 1,530 1,540 1,550 1,560 1,570 1,580 1,590 1,600 1,605
CGCGGTGCATCGGAGCAGGAAAGATGACGTTTCAACCAACCTCTTCTTGGCGAGCAGTATCCTCGGTACTCGGAAGAATATGAAAGGGATGCAAGTCAAGTCTGCTGTTTGA

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Deleted Nucleotide Copper Sequence