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ATGTCAGGTCTGCGTCAGATCGA	TTCTACGGA	AAGGGGGG	CATCGGCAAGTCCACCACCTC	TCAAAATACGCTAGCCGC	CTGGTCGACCT	CGGA	CAGAAA	AATCCTGAT	CGTCGGGCTGC	120
ATGGCAGCTCTGCGTCAGATCGG	TTCTACGGT	AAGGGGGG	TATCGGCAAGTCCAC	GACCTC	CAAAATAC	ACT	CGCCGCGCT	TGTCGACCTGGG	CAAAAGATCCT	120
ATGGCTTCACTAAGACA	AATCGC	CTTCTACGG	GAAAGG	CGGA	AATCGGCAAGTCCACCAC	TTCG	CAGAA	CACGCTAGC	GCGCTGC	120
GACCCCAAAGC	CGACTCCAC	CGGCTG	ATCCTGAACGCAAAGGG	G	CAGGAT	ACCGTTCTGCATCT	C	GCCGC	CCAA	240
GATCCGAAAGCGGACTCCAC	GCGCTCATCCTGAACGCAAAGGCA	CAGGACACCGT	ACTGCATCTTGC	GGA	AC	CGAAGGTTG	GT	CGAAGACCT	CAGCTCGAGGACGTGCTCAA	240
GATCCGAAAGCGGACTCGACT	CGCCTTATTCTG	CACGC	CAAGGCT	CA	A	GACAC	GATTTTGAG	TCTTGCCGC	GAGCGCCGC	240
GGCTACAAAGACATCAAGTGC	TGGAGTCCGG	C	GGCCC	C	GAGCCGGGTGTCGGCTGCGCCGGCCGCGGCGT	CATCACCTCGATCAACTTCCT	C	GAG	GAGAACGGCGCCTAT	360
GGTTACAGAGGCATCAAGTGC	TGGAGTCCGGTGGCCCA	GAGCCGGG	CGTCGGCTGCGCCGA	CGCGCGT	TATCACCTCGATCAACTTCCT	G	GAAGAGAACGGCGC	T	TACAA	360
GGCTACCAAGGACATTCGCT	TGCGTT	GAGTCCGGTGGCCCT	GAGCC	A	GGTGTGCGCTGCGCCGGCCGCGG	TGTCATCACCTCGATCAATTTTCTT	G	AAGAGAACGG	AGCCTACGA	360
GACTATGTCTCTACGACGTGCT	G	GGCGAT	GTG	GTG	TGCGGCGGCTTTGCGATGCC	GATCCGCGAAAAACAAGGC	C	CAGGAAATCTACATCGTCATGTCCGG	C	480
GATTA	CGTCTCATACGACGTGCT	AGG	G	GACGT	AGTATGCGGCGGCTTTGCGATGCC	TAT	T	CGCGAAAAACAAGGC	T	480
GACTATGTTCTTACGATGTGCT	T	GGCGACGT	T	GT	TGCGGT	GGCTTTGCGATGCC	A	ATCCGCGAAAAACAAGGC	C	480
GCCGCCAACACATCGCCAAGGGTATCCTGAAATACGCCCATTCGGGCGGCGTGCGGCT	C	GGCGGGCTGATCTGCAA	T	GAGCGA	CAGACCGACCG	C	GAGCTCGACCTC	T	CCGAGGCGCTG	600
GCCGCCAACACATCGCGA	AAGGGTATCCTGAA	G	TACGCCAT	G	CGGCGGCGTGCGGCTGGG	G	GGG	TGATT	TGCAACGAGCG	600
GCCGC	AAACAA	TATTT	CCAAGGG	GATCCTGAAATACGC	GA	ACT	CA	AGT	GGG	600
GCCGCCAG	CTCAATTCCAAGCTCATCCACTTCGTGCCGCGCGACAA	CATCGT	C	CAGCACGC	C	GAA	CTGAGAAAGATGACGGT	A	ATCCAG	720
GCCGCC	CGCTCAATTCCAAGCTCATCCACTTCGTGCCGCGCGACAA	TATCGT	T	CAGCACGCAGAGCT	C	AGAAAAGATGAC	A	GTGATCCAATATGCGCCG	A	720
GCC	AAGA	AGCTTGGCA	CTCA	CTGATCTACTTCGTGCCGCGT	GACAAT	GT	GGT	G	CAGCAT	720
TAC	CGCGCGT	TGGC	C	GAG	AAGATCCATG	TC	AATTC	G	GGCCAGGGCACCATCCC	840
TATCGCGC	CTGGCT	GAA	AAGATCCATG	CA	AATTCGGCC	G	AGGCACC	GTCCCT	TACA	840
TATCGGAAA	CTAGCG	GCC	AAGGTT	CA	CAAT	AATGG	CGGC	A	AGGGCAT	840
CAAATGCTTGCCGAG	CTT	CAG	GCCAAGGAAGCG	AAAT	T	G	CCGT	C	GT	894
CAGATGCTTGCCGAA	CTCCAC	GCCAAGGAAGCG	AAAGG	T	AAT	AGCCCC	C	CAC	TGA	894
GAA	TCAAT	T	CAT	CGG	...	CA	AAA	CC	...	885