

mRNAid

Input

Five Prime Flanking Sequence: AGGAATAAACTAGTATTCTTCTGGTCCCCACAGACTCAGAGA-GAACCCGCCACC

Three Prime Flanking Sequence:

ATCGTGGGCATTGTTGCTGGCCTGGCTGTCCTAGCAGTTGTGGTCATCGGAGCTGTGGTCGC-TACTGTGATGTGTAGGAGGAAGAGCTCAGGTGGAAAAGGAGGGAGCTACTCTCAGGCT-GCGTCCAGCGACAGTGCCCAAGGGCTCTGATGTGTCTCTCACAGCTTGATGAGCTGGAGC-CTCGGTGGCCTAGCTTCTTGCCCCCTTGGGCCTCCCCCAGCCCCCTCCTCCCCCTCCTGCAC-CCGTACCCCCGTGGTCTTTGAATAAAGTCTGAGTGGGCG-GCAA

mRNA Sequence:

AGAGAACCCGCCACCAUGGACGCAAUGAAGAGAGGC-CUGUGCUGCGUGCUGCUGCUGUGCGGCGCCGUGUUCGUGAGCCCCAUGCUGUGGCACGC-CAUGCCACCAGAGCUGAACACCGCCAGACUGAUGGCCGCGCAGGACCAGCCCCAUGCUG-GCCGCCGCCGCCGAUGGCAGACACUGAGCGCAGCCUGGACGCACAGGCCGUGGAACU-GACCGCCAGACUGAACAGCCUGGGAGAGGCCUGGACAGGCGGCGGCAGCGACAAGGC-CCUGGCGCGCCGCCACCAUGGUGGUGUGGUGCAGACCGCCAGCACCCAGGCCAAGA-CAAGAGCCAUGCAGGCCACCGCCCAGGCCGCCGCCUACACCCAGGCCAUGGCCACCAC-CCCCAGCCUGCCAGAGAUCCGCCCAACCAUACCCAGGCCGUGCUGACCGCCAC-CAACUUCUUCGGCAUCAACACCAUCCCCAUCGCCCUGACCGAGAUGGACUACUUAUCA-GAAUGUGGAACCAAGGCCGCCUGGCCAUGGAGGUGUACCAGGCCGAGACCGCCGUGAA-CACCCUGUUCGAGAAGCUGGAGCCCAUGGCCAGCAUCCUGGACCCCGGCGCCAGCCAGAG-CACCACCAACCCCAUCUUCGGCAUGCCCAGCCCCGGCAGCAGCACACCAGUG-GGCCAGCUGCCCCCAGCCGCCACCCAGACCCUGGGACAGCUGGGAGAGAUGAGCGGCC-CAUGCAGCAGCUGACCCAGCCACUGCAGCAGGUGACCAGCCUGUUCAGCCAGGUGGGCG-GCACCGGCGGCGGAACCCAGCCGACGAGGAAGCCGCCCAGAUGGGACUGCUGGGAA-CAAGCCCACUGAGCAACCACCCACUGGCCGGAGGCUCAGGACCAAGCGCCGGCGCCGGC-CUGCUGAGAGCCGAGUCACUGCCCGGCGCCGGAGGCAGCCUGACAAGAACCCCCCUGAU-GAGCCAGCUGAUCGAGAAACCCGUGGCCCCCAGCGUGAUGCCAGCCGCCGCCGCCG-GAAGCUCAGCAACCCGAGGAGCAGCACCCAGUGGGAGCCGGAGCCAUGGGACAGGGCG-CACAGAGCGGCGGAAGCACAAGACCAGGCCUGGUGGCCCCCCGCCCCACUGGCCCCAGGAGA-GAGAAGAGGACGACGAGGACGACUGGGACGAGGAGGACGACUGGGGCGGGCGGCG-GCAGCAUCGUGGGCAUUGUUGCUGGCCUGGCUGUCCUAGCAGUUGUGGUCAUCG-GAGCUGUGGUCGCUACUGUGAUGUGUAGGAGGAAGAGCUCAGGUGGAAAAGGAG-GGAGCUACUCUCAGGCUGCGUCCAGCGACAGUGCCCAGGGCUCUGAUGUGUCUCU-CACAGCUUGAUGAGCUGGAGCCUCGAGAGAACCCGCCACCAUGGACGCAAUGAAGAGAGGC-CUGUGCUGCGUGCUGCUGCUGUGCGGCGCCGUGUUCGUGAGCCCCAUGGCCAGAGGC-CUGCAGGGCGUGAUGCUGAGGAGCUUCGGCGCCAGGGACCACACCGCCACCGUGAUC-GAGACCAUCAGCAUCGCCCCCCACUUCGUGAGGGUGAGGAUGGUGAGCCCCACCCUGU-UCCAGGACGCCGAGGCCGAGCCGCCGCCUGGCUGAGAUUCUGGUUCCCCGACCCCAACG-GCAGCAACACCGAGAUUCCAGAGGGCCUACACCAUCAGCGAGGCCGACCCAGCCGCCGGCA-GAUUCGCCGUGGACGUGGUGCUGCACGACCCAGCCGGCCAGCCAGCAGCUGGGCCA-GAACCGUGAAGCCAGGCGCCACAUCGCCGUGAUGAGCCUGAUGGGCAGCAGCAGGUUC-GACGUGCCCGAGGAGCAGCCCGCCGGCUACCUGCUGAUCGGCGACAGCGCCAGCAUC-CCCGGCAUGAACGGCAUCAUCGAGACCGUGCCCAACGACGUGCCCAUCGAGAUGUACCUG-GAGCAGCACGACGACAACGACACCCUGAUCCCCCUGGCCAAGCACCCACAGACUGAGGGU-GAGGUGGGUGAUGAGGAGGGACGAGAAGAGCCUGGCCGAGGCCAUCGAGAACAGGGACUG-GAGCGACUGGUACGCCUGGGCCACCCCGAGGCCGCCGCCUGAAGUGCGUGAGAGUGAG-GCUGAGGGACGAGUUCGGCUUCCCCAAGAGCGAGAUCCACGCCCAGGCCUACUGGAACGC-CGGCAGAGCCAUGGGCACCCACAGAGCCACCGAGCCAGCCGCCACAGAACCAGAAGUG-GGCGCCGCCCCACAGCCAGAGUCAGCCGUGCCAGCCCCAGCCAGAGGCAGCUGGAGAGC-

CCAGGCAGCAAGCAGACUGCUGGCACCCCUGAAGCUGCCCCUGGUGCUGAGCG-
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GAGCUGAGCAGGCUGCUGGUGAGCGGGCGCCGGAGCCCACAGACUGUUCACAGUGGGCU-
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GCUGCACGUGAUCGACGCCAGGUUCGCCAGGGCCCUGAGACUGAGGCUGCUGAGCAAGCU-
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GCCCUGGUGCUGUUCGGCCCCGUGCUGGUGUACCUGACCAUACCCAGCAGCCUGAC-
CAUCCAGAGCGGCCCCAGGAUUCGUGCAGGCCCCAGAGGUGGGCCGAGAAGAUGAACGGC-
GAGGCCCGGCAGCUACCUGGAGGGGCCAGCCCGUGAUCAGAGUGUUCGGCGC-
CGCCAGCAGCAGCUUCAGGAGGAGGCUGGACGAGUACAUCGGCUUCCUGGUGGCCUGGCA-
GAGGCCCCUGGCCGGCAAGAAGACCCUGAUGGACCUGGCCACCAGACCCGCCACCUUC-
CUGUGGCUGAUCGCCGCCACCGGCACCCUGCUGGUGGCCACCCACAGAAUGGACCCCGU-
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CUACGGCCUGGGCGGCCUGAGGACAGGCCUGCUGGCCGCCAGACACCUGCAGGUGA-
CACUGGACGAGACCGAGCUGGCCGUGAGAGAGCACCCAGGGAGCCACUGGACGGCGAG-
GCCCCCGCCACCGUGGUGUUCGACCACGUGACCUUCGGCUACAGACCCGGCGUGCCAGU-
GAUCCAGGACGUGAGCCUGACCCUGAGACCAGGCACCGUGACAGCCCUUGGUGGGC-
CCCAGCGGAAGCGGCAAGAGCACCCUGGCCACCCUGCUGGCCAGGUUCCACGACGUG-
GAGAGGGGGCGCCAUCAGAGUGGGCGGCCAGGACAUCAGGAGCCUGGCCGCCGAC-
GAGCUGUACACCAGAGUGGGCUUCGUGCUGCAGGAGGCCCAGCUGGUGCACGGAACCGC-
CGCCGAGAACAUCGCACUGGCCGUGCCAGACGCCCCAGCCGAGCAGGUGCAGGUG-
GCAGCCAGAGAGGCCCAGAUCCACGACAGGGUGCUGAGACUGCCCGACGGCUACGA-
CACAGUGCUGGGCGCCAACAGCGGCCUGAGCGGGCGGCGAGAGGCAGAGACUGAC-
CAUCGCCAGAGCCAUCCUGGGCGACACCCCCGUGCUGAUCUGGACGAGGCCACCGCCU-
UCGCCGACCCCGAGAGCGAGUACCUGGUGCAGCAGGCCCUGAACAGACU-
GACCAGGGACAGGACCGUGCUGGUGAUCGCCACAGGCUGCACACCAUACCCAGGGCC-
GACCAGAUUCGUGGUGCUGGACCACGGCAGGAUCGUGGAGAGAGGCACCCACGAG-
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GAGUGGCCGUGGCCGCCGCCAGGACGGCACCAAGAGGCGGCAUCGUGGGCAUUGU-
UGCUGGCCUGGCUGUCCUAGCAGUUGUGGUCAUCGGAGCUGUGGUCGCUACUGUGAU-
GUGUAGGAGGAAGAGCUCAGGUGGAAAAGGAGGGAGCUACUCUCAGGCUGCGUCCAGC-
GACAGUGCCCAGGGCUCUGAUGUGUCUCUCACAGCUUGAUGAGCUGGAGCCUCGAGA-
GAACCCGCCACCAUGGACGCAUUGAAGAGAGGCCUGUGCUGCGUGCUGCUGCUGUGCG-
GCGCCGUGUUCGUGAGCCCCAUGAGCUACAUGAUCGCCACCCAGCCGCCUGACCGC-
CGCCGCCACCGACAUCGACGGAUCGGCAGCGCAGUGAGCGUGGCCAACGCAGC-
CGCAGUGGCCGCAACAACAGGAGUGCUGGCCGCCGGAGGCGACGAGGUGCUGGCCGC-
CAUCGCCAGGCUGUUAACGCCAACGCCGAGGAGUACCACGCCCUAGAGCGCCAGGUGGC-
CGCCUUCAGACCCUGUUCGUGAGGACCCUGACCGGGCGGCUGCGGCGUGUUCAGAAGAA-
GAAGGGGCAGGCAGUGCGUGACCGCCGCCGAGCACAGAGCCGCCGGCGGCCGAGAA-
GACAGAGGAGAAGAAGAAGCGGCGACGGCCAGUGGAGACUGAGACAGCAGAGACACUUCG-
GCUGCGGGCGGCCAGCCCGAGUUCAGGCAGCACAGCGAGCACAGGAGAGGGCGGCGGCG-
GCAGCAUCGUGGGCAUUGUUGCUGGCCUGGCUGUCCUAGCAGUUGUGGUCAUCG-
GAGCUGUGGUCGCUACUGUGAUGUGUAGGAGGAAGAGCUCAGGUGGAAAAGGAG-
GGAGCUACUCUCAGGCUGCGUCCAGCGACAGUGCCCAGGGCUCUGAUGUGUCUCU-
CACAGCUUGAUGAGCUGGAGCCUCG

Codon Usage: h_sapiens

Codon Usage freq.threshold: 0.1

Uridine depletion:

avoid Motifs: HindIII

Parameters

Min GC Content: 0.3

Window Size for local GC Content: 100

Max GC Content: 0.7

Entropy Window size: 30

Optimized Outputs

Output 1

RNA Sequence:

AGGAAUAAACUAGUAUUCUUCUGGUCCCCACAGACUCAGAGAGAACCCGCCACCA-
GAGAACCAGCAACAAUGGACGCAUGAAAAGAGGC-
CUGUGCUGCGUGCUGCUGCUGUGCGGCGCCGUGUUUGUUUCCCCGAUGCUCUG-
GCAUGCGAUGCCCGCCUGAACUCAAUACAGCUCGAUUGAUGGCCGGUGCAGGGGCCUG-
CACCGAUGCUUGCUGCAGCGGCGGGUUGGCAAACACUCAGUGCUGCGUUGGAUGCG-
CAAGCAGUCGAACUCACUGCACGGCUUAAUUCCCUCGGAGAAGCGUGGACUGGUGGAG-
GAUCCGAUAAAGCGCUCGCGUGCGGCUACACCUAUGGUAGUAUGGUUGCAAACUGCAU-
CAACGCAAGCGAAAACUCGGGCUAUGCAAGCAACUGCUCAAGCGGCAGCAUAUA-
CACAAGCGAUGGCUACGACACCUUCUCUUCGGAUAGCUGCAAUUAUUA-
CACAAGCGGUUCUCACUGCGACGAACUUCUUCGGCAUCAACACCAUCCCCAUCGCCCCU-
GACCGAGAUGGACUACUUAUCAGAAUGUGGAACCAGGCCGCCUGGCCAUGGAG-
GUGUACCAGGCCGAGACCGCCGUGAACACCCUGUUCGAGAAACUUGAACCUAUGGCAU-
CAAUACUUGAUCCUGGUGCUUCACAAUCCACAACUAAUCCUAUAUUUGGAUUGCCUUC-
CCCUGGUUCUUCACGCCGGUUGGACAACUUCGCGCCGCGCAGCUACUCAACUCUCGGU-
CAAUUGGGGGAAUUGUCCGGUCCGAUGCAACAACUCACGCAACCGCUUCAACAGGU-
UACUAGUUUGUUUAGUCAGGUAGGGGGGAACAGGGGGGGGGAAUCCUGCAGAUCAA-
GAAGCUGCGCAAUUGGGGUUGCUUGGGACAUCCCCUCUCUCAAAUUAUCCUCU-
UGCAGGAGGUUCAGGGCCUAGUGCUGGAGCUGGUUCUUCUCCGAGCUGAAUCUCUUC-
CUGGGGGCGGGGGGUUCACUCACGCGGACACCAUUGAUGUCACAGUUGAUUGAAAAAC-
CUGUCGCACCAUCUGUUAUGCCAGCAGCAGCAGCAGGAUCUUCAGCGACGGGAG-
GUGCUGCACCUUGUUGGUGCUGGUGCAAUGGGACAGGGUGCACAGAGUGGUGGUU-
CAACUCGGCCUGGACUUGUUGCGCCAGCGCCUCUUGCUCAAGAACGGGAAGAAGAU-
GAUGAAGAUGAUUGGGAUGAGGAAGAUGAUUGGGGGGGGGCGGGUCAAUUGUCG-
GGAUAGUUGCUGGCCUGGCUGUCCUGGCAGUUGUGGUCAUCGGAGCUGUG-
GUCGCUACUGUGAUGUGUAGGAGGAAGAGCUCAGGUGGAAAAGGAGGGAGCUACU-
CUCAGGCUGCGUCCAGCGACAGUGCCCAGGGCUCUGAUGUGUCUCUCACAGCUUGAU-
GAGCUGGAGCCAGUAGAGAACCCGCCACCAUGGACGCAAUGAAGAGAGGGUUGUGU-
UGUGUCUUGUUGCUUUGUGGAGCGGUUUUUGUUUCUCCUAUGGCGCGGGGU-
CUCCAGGGAGUAAUGCUUCGGUCAUUUUGGGGCCCCGCGAUCAUACGGCGACGGUUAUC-
GAGACCAUCAGCAUAGCUCCUCAUUUUGUCCGGGUACGAAUGGUCUCCCCUACACUUUU-
UCAGGAUGCGGAAGCUGAACCUGCAGCGUGGCUUCGGUUUUGGUUUCGGAUC-
CAAACGGCAGCAAUACUGAAUUUCAGCGCGCAUAUACGAUUUCAGAAGCAGAUC-
CUGCUGCAGGACGCUUUGCGGUCGACGUCGUUCUCCAUGAUCCUGCUG-
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GAGCCUGAUGGGCAGCAGCAGGUUCGACGUGCCCGAGGAGCAGCCCGCCG-
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CAUCGGCUUUUUGGUAGCCUGGCAGCGACCCCUUGCCGGGAAGAAGACUCUGAUGGAU-
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GAGUACCCUGGCCACAUUGCUCGCCCCGCUUCCACGAUGUGGAGCGGGGGGCCAUUCG-
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GGGUUCGUGCUUCAGGAGGCCCAGCUGGUGCACGGAACCGCAGCCGAGAAUAUUG-
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CCGCCUGUUCAAUGCAAACGCCGAGGAGUACCACGCCCUUGUCCGCCAGGUGGCCGC-
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GAAGAGGCCGCCAGUGCGUGACAGCCGCCGAACACAGAGCCGCCGGUGCCGGCA-
GACGCCAGCGCCGCAGAAAGAUUCUGGCGACGGCCAGUGGAGACUGAGACAGCAGAGA-
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GCGGCGGCAGUAUCGUGGGCAUUGUUGCUGGCCUGGCUGUCCUCGCAGUUGUGGU-
CAUCGGAGCUGUGGUCGCUACUGUGAUGUGUAGGAGGAAGAGCUCAGGCGGAAAAG-
GAGGGAGCUACUCUCAGGCUGCGUCCAGCGACAGUGCCCAGGGCUCUGAUGUGU-
CUCUCACAGCUUGAUAAAGCUGGAGCCUCUAUCGUGGGCAUUGUUGCUGGCCUG-
GCUGUCCUAGCAGUUGUGGUCAUCGGAGCUGUGGUCGCUACUGUGAUGUGUAGGAG-
GAAGAGCUCAGGUGGAAAAGGAGGGAGCUACUCUCAGGCUGCGUCCAGCGACAGUGC-
CCAGGGCUCUGAUGUGUCUCUCACAGCUUGAUGAGCUGGAGCCUCGGUGGCCUAGCU-
UCUUGCCCCUUGGGCCUCCCCCAGCCCCUCCUCCCUUCCUGCACCCGUACCCC-
CGUGGUCUUUGAAUAAAGUCUGAGUGGGCG-
GCAA

DNA Sequence:

AGGAATAAACTAGTATTCTTCTGGTCCCCACAGACTCAGAGAGAACCCGCCACCA-
GAGAACCAGCAACAATGGACGCAATGAAAAGAGGCCTGTGCTGCGTGCTGCTGTGCG-
GCGCCGTGTTTGTTCCTCCCGATGCTCTGGCATGCGATGCCGCCTGAACTCAATACAGCTC-
GATTGATGGCCGGTGCAGGGCCTGCACCGATGCTTGCTGCAGCGGCGGGTTGGCAA-
CACTCAGTGCTGCGTTGGATGCGCAAGCAGTCAACTCACTGCACGGCTTAATTCCTCG-

GAGAAGCGTGGACTGGTGGAGGATCCGATAAAGCGCTCGCTGCGGCTACACCTATGGTAG-
TATGGTTGCAAACCTGCATCAACGCAAGCGAAAACCTCGGGCTATGCAAGCAACTGCT-
CAAGCGGCAGCATATACACAAGCGATGGCTACGACACCTTCTCTTCCGGAAATAGCTG-
CAAATCATATTACACAAGCGGTTCTCACTGCGACGAACTTCTTCGGCATCAACACCATCCC-
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GAGGTGTACCAGGCCGAGACCGCCGTGAACACCCTGTTTCGAGAACTTGAACCTATG-
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CCAGGGCTCTGATGTGTCTCTCACAGCTTGATGAGCTGGAGCCAGTAGAGAACCCGCCAC-
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CAAGCACCCAGACTGAGGGTGAGGTGGGTGATGAGGAGGGACGAGAAGAGCCTGGCC-
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CGCCCTGAAGTGCGTGAGAGTGAGGCTGAGGGACGAGTTCGGCTTCCCCAAGAGT-
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CCAGCTCCAGCTCGAGGTAGTTGGCGAGCTCAGGCAGCTTCCCGGTTGCTTGCTCCACT-
CAAACCTTCCACTCGTACTCAGTGGTGTTCTCGCTGCATTGGTTACCCTTGACAGTTG-
GCTCCATTTGTATTGTTGGTCAATTGTCCCGGCTCCTCGTTTCCGGAGCAGGAG-
CACATCGCCTCTTTACAGTCGGATTTGCTGCAGTAGGGCTTCTCGGGACAGGTGCTCTC-
CTCGCTGCAGCATTGACTTTGTGGCTCCATGTTATAGATGCTCGGTTTGCTCGAGCTTTGC-
GATTGAGGCTGCTGAGCAAGCTGAGCAGGCTGCCCCCTGGGCTGGTTCACCAGCAG-
GGGCAGCGGCAGCATAAAAAAGCTCGTAACAGATGATACTCGCACTCCATTATCTCGTAA-
CACATGCCGTCCCAGATGCTGTGCTGCTGTAGTCGCTCCAGTAGGTGTTCTTGTCTATTT-
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GGCCGAGAAGATGAACGGCGAGGCCGGCAGCTACCTGGAGGGCCAGCCCGTGATCA-
GAGTGTTTCGGCGCCGCCTCCAGCAGCTTCAGGAGGAGGCTGGACGAGTACATCGGCT-
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GAATGGACCCCGTGAACCTGCTGCCCTTCATGTTCTCGGGACTACTTTTGGAGCCC-
GACTCCTGGGAATAGCCTATGGGCTTGGGGGACTGCGCACTGGGCTGTTGGCCGCCCG-
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CCGGAGTCCCAGTGATCCAGGACGTGAGCCTGACCCTGCGCCCAGGGACCGTGACCG-
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CGATGAGCTGTATACCCGGGTGGGGTTCGTGCTTCAGGAGGCCCAGCTGGTGCACGGAAC-
CGCAGCCGAGAATATTGCACTGGCCGTGCCAGACGCCCCAGCCGAGCAGGTGCAGGTG-
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CTTCGCCGACCCCGAGTCTGAGTACCTGGTGCAGCAGGCCCTGAACAGACTGACCCG-
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CCAGGGCTCTGATGTGTCTCTCACAGCTTAGTAAGCTGGAGCCAGCAGAGAACCCGCCAC-
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CCTGTTCTGTGCGCACCCCTGACCGGCGGCTGCGGCGTGTTTCGCCGCAGAAGAGGC-
CGCCAGTGCGTGACAGCCGCCGAACACAGAGCCGCCGGTGCCGGCAGACGCCAGCGC-
CGCAGAAGATCTGGCGACGGCCAGTGGAGACTGAGACAGCAGAGACACTTCGGCTGCG-
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CGTGGTCTTTGAATAAAGTCTGAGTGGGCG-
GCAA

Parameters

A Ratio:0.20654817586529467	AT Ratio:0.4112254443405051
T/U Ratio:0.20467726847521048	GC Ratio:0.5887745556594949
G Ratio:0.3073900841908326	MFE:-2230.5
C Ratio:0.2813844714686623	Score:0.6444563473332711
5 MFE:-12.100000381469727	CAI:0.7504599714596822

Output 2

RNA Sequence:
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GCUCCGAUGCUUGCAGCUGCUGCUGGAUGGCAAACUUUGUCCGCGGCUUUGGACG-
CACAAGCGGUAGAACUUACUGCGCGACUCAAUAGUCUCGGAGAAGCGUGGACUGGAG-
GGGGAUCAGAUAAAGCAUUGGCUGCGGCGACGCCGAUGGUCGUUUGGCUU-
CAAACAGCAUCAACACAAGCGAAAACUCGAGCUAUGCAAGCAACAGCGCAAGCAGCG-
GCGUAUACACAAGCGAUGGCGACUACGCCUUCACUCCUGAAAUAGCAGCGAAU-
CAUAUAACGCAAGCUGUACUCACAGCAACGAUUUCUUCGGCAUCAACACCAUCCC-
CAUCGCCCUGACCGAGAUGGACUACUUCAUCAAGAAUGUGGAACCAGGCCGCCUGGC-
CAUGGAGGUGUACCAGGCCGAGACCGCCGUGAACACCCUGUUCGAGAAAUUGGAAC-
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GUAUGCCUAGUCCGGGGAGUUCUACACCGGUAGGGCAAUUGCCGCCUGCAGCAACU-
CAAACUUUGGGUCAACUUGGAGAAAUGUCCGGUCCGAUGCAACAAUUGACUCAAC-
CGCUCCAGCAAGUCACUAGUCUUUUUCCCAAGUCGGAGGUACAGGUGGAGGGAAUC-
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CAUCCUUUGGCUGGGGGGAUCUGGACCUUCCGCGGGGGCCGGGCUCCUCCGAGCA-
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UAUAGAAAAACCAGUCGCAACCUUCCGUUAUGCCAGCAGCUGCGGGCAGGGAGU-
UCAGCAACUGGAGGAGCAGCACCAGUUGGGGGCGGGAGCAAUG-
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GCAA

DNA Sequence:

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CCAGGGGCTCTGATGTGTCTCTCACAGCTTGATGAGCTGGAGCCTCGGTGGCC-
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GCAA

Parameters

A Ratio:0.20654817586529467	AT Ratio:0.41159962581852194
T/U Ratio:0.20505144995322733	GC Ratio:0.5884003741814781
G Ratio:0.3079513564078578	MFE:-2224.5
C Ratio:0.2804490177736202	Score:0.6438388548291909
5 MFE:-12.100000381469727	CAI:0.7495788466671152

Output 3

RNA Sequence:
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GCAA

DNA Sequence:

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GCAA

Parameters

A Ratio:0.20654817586529467	AT Ratio:0.4112254443405051
T/U Ratio:0.20467726847521048	GC Ratio:0.5887745556594949
G Ratio:0.307577174929841	MFE:-2239.60009765625
C Ratio:0.2811973807296539	Score:0.6438159619548145

Output 4

RNA Sequence:

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GCAA

DNA Sequence:

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GGCG-
GCAA

Parameters

A Ratio:0.20654817586529467	AT Ratio:0.41216089803554723
T/U Ratio:0.20561272217025256	GC Ratio:0.5878391019644528
G Ratio:0.3081384471468662	MFE:-2193.199951171875
C Ratio:0.27970065481758655	Score:0.6437747645483248
5 MFE:-12.100000381469727	CAI:0.7494376136187404

Output 5

RNA Sequence:
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DNA Sequence:

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GCAA

Parameters

A Ratio:0.20673526660430308	AT Ratio:0.41216089803554734
T/U Ratio:0.20542563143124415	GC Ratio:0.5878391019644527
G Ratio:0.3072029934518241	MFE:-2201.300048828125
C Ratio:0.2806361085126286	Score:0.6433475459403518
5 MFE:-12.899999618530273	CAI:0.7496233653960439

Output 6

RNA Sequence:
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DNA Sequence:

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GCAA

Parameters

A Ratio:0.20636108512628626	AT Ratio:0.41216089803554723
T/U Ratio:0.205799812909261	GC Ratio:0.5878391019644528
G Ratio:0.30907390084190833	MFE:-2229.699951171875
C Ratio:0.27876520112254444	Score:0.6431697115470376
5 MFE:-12.100000381469727	CAI:0.749233309243293

Output 7

RNA Sequence:

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GCAA

Parameters

A Ratio:0.20636108512628626	AT Ratio:0.4117867165575304
T/U Ratio:0.20542563143124415	GC Ratio:0.5882132834424696
G Ratio:0.30851262862488305	MFE:-2230
C Ratio:0.27970065481758655	Score:0.6431042533429341
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Output 8

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CACAGTGCTGGGGGCCAATTCGGGGCTTTCCGGTGGGGAGCGGCAGCGCCTGAC-
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GGCG-
GCAA

Parameters

A Ratio:0.20673526660430308	AT Ratio:0.41216089803554734
T/U Ratio:0.20542563143124415	GC Ratio:0.5878391019644527
G Ratio:0.30851262862488305	MFE:-2240.300048828125
C Ratio:0.27932647333956967	Score:0.6430130207903054
5 MFE:-12.100000381469727	CAI:0.7496233653960439

Output 9

RNA Sequence:
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CAAACAUUGGGGCAACUCGGAGAAUUGUCUGGUCCAUGCAACAAUUGACGCAACCU-
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GCAA

DNA Sequence:

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GCAA

Parameters

A Ratio:0.20710944808231993	AT Ratio:0.41253507951356405
T/U Ratio:0.20542563143124415	GC Ratio:0.587464920486436
G Ratio:0.307577174929841	MFE:-2220.5
C Ratio:0.27988774555659496	Score:0.6426744525579702
5 MFE:-12.899999618530273	CAI:0.7500853943037809

Output 10

RNA Sequence:

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DNA Sequence:

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Parameters

A Ratio:0.20710944808231993	AT Ratio:0.41253507951356405
T/U Ratio:0.20542563143124415	GC Ratio:0.587464920486436
G Ratio:0.3073900841908326	MFE:-2218.5
C Ratio:0.28007483629560337	Score:0.6426731628000976
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