

```

1
GFP_AEQVI... ATGAGCAAAGGCGAAGAAGCTGTTTACCGGCGTGGTGCCGATTCTGGTGGAAGCTGGATGGCGATGTGAACGGCCATAAATTTA
GFP_RNA      AUGAGCAAAGGCGAAGAACUGUUUACCGGCGUGGUACCGAUUCUGGUGGAACUGGAUGGCGAUGUGAACGGCCAUAUUUUUA
.....

83
GFP_AEQVI... GCGTGAGCGGCGAAGGCGAAGGCGATGCGACCTATGGCAAAGCTGACCCGTGAAATTTATTTGCACCACCGGCAAAGCTGCCGGT
GFP_RNA      GCGUGAGCGGCGAAGGCGAAGGCGAUGCGACCUAUGGCAAACUGACCCUGAAAUUUAUUUUGCACCACCGGCAAACUGCCGGU
.....

165
GFP_AEQVI... GCCGTGGCCGACCCCTGGTGACCACCTTTAGCTATGGCGTGCAGTGCTTTAGCCGCTATCCGGATCATATGAAACAGCATGAT
GFP_RNA      GCCGUGGCCGACCCUGGUGACCACCUUUAGCUAUGGCGUGCAGUGCUUUAGCCGCUAUCCGGAUCAUAUGAAACAGCAUGAU
.....

247
GFP_AEQVI... TTTTTTAAAAGCGCGATGCCGGAAGGCTATGTGCAGGAACGCACCATTTTTTTTTTTAAAGATGATGGCAACTATAAAAACCCGCG
GFP_RNA      UUUUUUAAAAGCGCGAUGCCGGAAGGCUAUGUGCAGGAACGCACCAUUUUUUUAAAGAUGAUGGCAACUAUAAAACCCGCG
.....

329
GFP_AEQVI... CGGAAGTGAAATTTGAAGGCGATAACCTGGTGAACCGCATTGAACTGAAAAGGCATTGATTTTTAAAGAAGATGGCAACATTCT
GFP_RNA      CGGAAGUGAAAUUUGAAGGCGAUACCCUGGUGAACCGCAUUGAACUGAAAAGGCAUUGAUUUUAAAGAAGAUUGGCAACAUUCU
.....

411
GFP_AEQVI... GGGCCATAAACTGGAATATAACTATAACAGCCATAACGTGTATATTATGGCGGATAAACAGAAAAACGGCATTAAAGTGAAC
GFP_RNA      GGGCCAUAAACUGGA---AUAACUAUAACAGCCAUAACGUGUAUAUUUUGGCGGAUAAACAGAAAAACGGCAUUAAGUGAAC
.....

493
GFP_AEQVI... TTTAAAATTCGCCATAACATTGAAGATGGCAGCGTGCAGCTGGCGGATCATTATCAGCAGAACACCCCGATTGGCGATGGCC
GFP_RNA      UUUAAAAUUCGCCAUAAACAUUGAAGAUUGGCGAGCGUGCAGCUGGCGGAUCAUUAUCAGCAGAACACCCCGAUUGGCGAUGGCC
.....

575
GFP_AEQVI... CGGTGCTGCTGCCGATAACCATTTATCTGAGCACCCAGAGCGCGCTGAGCAAAGATCCGAACGAAAAACGCGATCATATGGT
GFP_RNA      CGGUGCUGCUGCCGGAUAACCAUUAUCUGAGCACCCAGAGCGCGCUGAGCAAAGAUCCGAACG---AAAACGCGAUCUAUUGGU
.....

657
GFP_AEQVI... GCTGCTGGAATTTGTGACCGCGGCGGGCATTACCCATGGCATGGATGAACTGTATAAA
GFP_RNA      GCUGCUGGAAUUUGUGACCCGCGGCGGGCAUUAACCAUGGCAUGGAUGAACUGUAUAAA
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714

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