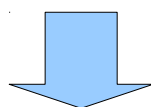


Q₁: GCUAAUAUCGCUGUGGAAACACCUGGAACCAUCCCGAACCCAGCAGU



A₁:

U39694.1/9296-9392GCUA.AUAUCGCUGUGGAAACACCUGGAACCAUCCCGAACCCAGC.AGU
M35167.1/2-111	UCCGGUGACUUUACGCGUGAGGAAACACUCGUUCCCAUCCGGAACACGAC.AGU
AJ131602.1/3-115	CCUGAUGACCAUAGCGACGUGGUACCACUCCUUCCCAUCCCGAACAGGAC.AGU
X52302.1/2-117	GCUGGUGGCUAUGGCGGAAGGGCCACACCCGAUCCCAUCCCGAACUCGGU.CGU
Y00159.1/2-117	CUUGGCGACUAUAGCGAUUUGGAACCACCUGAUACCAUCUCGAACUCAGA.AGU
M58387.1/5-112ACGGCCACAGUCAGCUGAAAACUGGGCAUCCCGUCC.GCUCUGCCAUACA
X67494.1/1-118	AUCCUCGGCCAUAGAAUGACGAAAACGACGCGUCCCGUCC.GAUCUGCGA.AUC
M36316.1/2-120	UCUUACGGCCAUUCACACCAGAAAGCACCAAUCCCGUCC.GAUCUUUGA.AGU
X02706.1/2-120	UGGAUCGUUCAAACCUUCAAGGCCCCUCCCAUCCCAUCA.GCACUGGGA.AGA
X05535.1/1-118	AGGAACGGCCAUACCACGUCGAUCGCACCACAUCCCGUCC.GCUCUGUGA.AGU
#=GC SS_cons	S: <<<<<<<<...<<.<<<<<...<<..<<<<<.....>>..>>>>...>>



S₁: <<<<...<<.<<<<<...<<..<<<<<.....>>..>>>>..>>