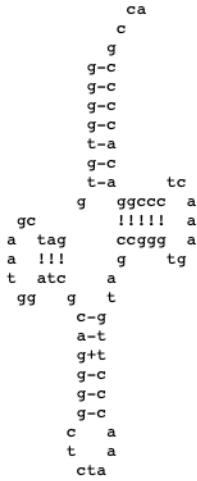
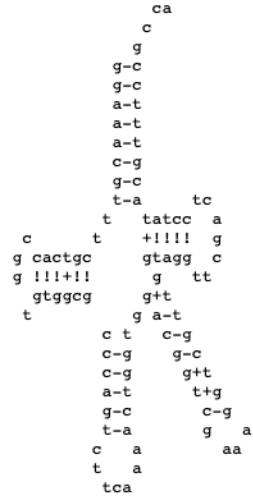


A**Bilophila sp. 4_1_30**

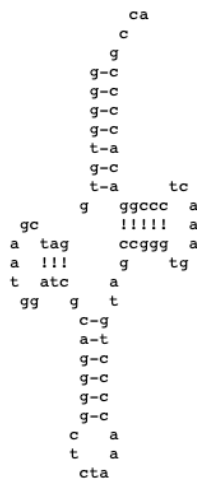
tRNA-Pyl(cta)
72 bases, %GC = 62.5
Sequence [437,508]

Primary sequence for tRNA-Pyl(cta)
1 . 10 . 20 . 30 . 40 . 50
ggggtgtggatcgaatggatcgagggtctaaaccctgtagccgggt
gaaactcccggacaccgccca



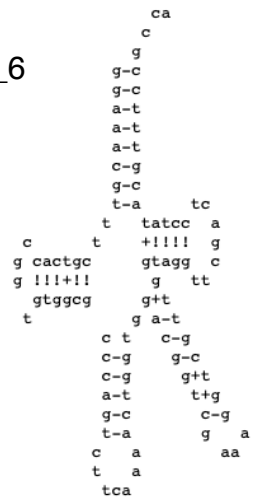
tRNA-SeC(tca)
94 bases, %GC = 55.3
Sequence c[862704,862797]

Primary sequence for tRNA-SeC(tca)
1 . 10 . 20 . 30 . 40 . 50
ggaacgtttcgtcacgggtgtggcggccagtcctcaaaactggtggacg
gtcgaaaggtcgttggtaggttcgactcctatacgtttccgccca

B**Bilophila wadsworthia 3_1_6**

tRNA-Pyl(cta)
72 bases, %GC = 63.9
Sequence [735263,735334]

Primary sequence for tRNA-Pyl(cta)
1 . 10 . 20 . 30 . 40 . 50
ggggtgtggatcgaatggatcgagggtctaaaccctgtagccgggt
gaaactcccggacaccgccca



tRNA-SeC(tca)
94 bases, %GC = 55.3
Sequence c[70717,70810]

Primary sequence for tRNA-SeC(tca)
1 . 10 . 20 . 30 . 40 . 50
ggaacgtttcgtcacgggtgtggcggccagtcctcaaaactggtggacg
gtcgaaaggtcgttggtaggttcgactcctatacgtttccgccca