**Sequences of sponge-derived *Chlorella-like* algae**

18S

ATTCTTGGATTTATGAAAGACGAACTACTGCGAAAGCATTTGCCAAGGATGTTTTCATTAATCAAGAACGAAAGTTGGGGGCTCGAAGACGATTAGATACCGTCCTAGTCTCAACCATAAACGATGCCGACTAGGGATCGGCGGATGTTTCTTCGATGACTCCGCCGGCACCTTATGAGAAATCAAAGTTTTTGGGTTCCGGGGGGAGTATGGTCGCAAGGCTGAAACTTAAAGGAATTGACGGAAGGGCACCACCAGGCGTGGAGCCTGCGGCTTAATTTGACTCAACACGGGAAAACTTACCAGGTCCAGACATAGTGAGGATTGACAGATTGAGAGCTCTTTCTTGATTCTATGGGTGGTGGTGCATGGCCGTTCTTAGTTGGTGGGTTGCCTTGTCAGGTTGATTCCGGTAACGAACGAGACCTCAGCCTGCTAAATAGTCACGGTTGGCTCGCCAGCCGGCGGACTTCTTAGAGGGACTATTGGCGACTAGCCAATGGAAGCATGAGGCAATAACAGGTCTGTGATGCCCTTAGATGTTCTGGGCCGCACGCGCGCTACACTGATGCATTCAACGAGCCTAGCCTTGGCCGAGAGGCCCGGGTAATCTTTGAAACTGCATCGTGATGGGGATAGATTATTGCAATTATTAATCTTCAAC

5’ region of SSU ITS

TTAGGGTTCGATTCCGGAGAGGGAGCCTGAGAAACGGCTACCACATCCAAGGAAGGCAGCAGGCGCGCAAATTACCCAATCCTGACACAGGGAGGTAGTGACAATAAATAACAATACTGGGCCTTTTCAGGTCTGGTAATTGGAATGAGTACAATCTAAACCCCTTAACGAGGATCAATTGGAGGGCAAGTCTGGTGCCAGCAGCCGCGGTAATTCCAGCTCCAATAGCGTATATTTAAGTTGCTGCAGTTAAAAAGCTCGTAGTTGGATTTCGGGTGGGGCCTGCCGGTCCGCCGTTTCGGTGTGCACTGGCAGGGCCCACCTTGTTGCCGGGGACGGGCTCCTGGGCTTCACTGTCCGGGACTCGGAGTCGGCGCTGTTACTTTGAGTAAATTAGAGTGTTCAAAGCAGGCCTACGCTCTGAATACATTAGCATGGAATAACACGATAGGACTCTGGCCTATCCTGTTGGTCTGTAGGACCGGAGTAATGATTAAGAGGGACAGTCGGGGGCATTCGTATTTCATTGTCAGAGGTGAA

3’ region of SSU ITS

AGGTGAACCTGCGGAAGGATCATTGAATCGATCGAATCCACTCTGTGAACCAAACGTCCCCCCTTGGGTGCGGGCTTCGGTCTGCCCCAAGGCGTCGGTTCCCTGGCTGGGGTCTTCGGACCGCAGTTAGGTCCGGCGGGCGCGCCCTCTGGCGTGTCGGCCCTCGTGGCTGCCGCCAGTTGGGTTCGCTGGAAATTGTATCCAACTCAACCCACCCCAAACCACAACTTATACTGAAGCAATCGGTGAGTGCACTCTGGTGCCTCGCTCTAACCAAAGACAACTCTCAACAACGGATATCTTGGCTCCCGTATCGATGAAGAACGCAGCGAAATGCGATACGTAGTGTGAATTGCAGAATTCCGTGAACCATCGAATCTTTGAACGCAAATTGCGCCCAAGGCTTCGGCCGAGGGCATGTCTGCCTCAGCGTCGGCTTACCCCCTCGCTCCCCCTCTCCTTTGGAGTGGGTGAACGGATCTGGTTTTCCCGGCTACGTGCTTCTGCACGCCCGGGTTGACTGAAGTGTAGAGGCTTGAGCATGGACCCCGTTTGTAGGGCAATGGCTTGGTAGGTAGCTTAGCTACACCGCCTGCCGTGGTCCGAGGGGACTTTGCTGGCGGCCCAGCAGGAATTCGGGTGTTGGGTTTCCCACCCCGAAAGCTTCAAACCTTCGACCTGAGCTCAGGCAAGACTACCCGCTGAACTTAAGCATATCAATAAGCGGAGGAAAAGAAACTAACTAGGATGCCCTTAGTAACGGCGAGCGAACCGGGCAAAGCCCAACTTGAAAATCTCCAGCCTCCGGCTGGCGAATTGTAGTCTAGAGAAGTGCTCTCTGCCTCAGCCTGTTCCCAAGTCCCCTGGAAAGGGGCGTCAGAGAGGGTGAGAACCCCGTTGGGATCGGATCCTGAGGCTCC

23S

AGCCGGCGACTTAGAAAACGTGGCAAGGTTAAGGAAATGTATCCGGAGCCGAAGCGAAAGCAAGTCTGAATAGGGCGCGTAAGTCATTTTTTCTAGACCCGAACCCGGGTGATCTAACCATGACCAGGATGAAGCTTGGGTGACACCAAGTGAAGGTCCGAACCGACCGATGTTGAAAAATCGGCGGATGAGTTGTGGTTAGCGGTGAAATACCAGTCGAACTCGGAGCTAGCTGGTTCTCCCCGAAATGCGTTGAGGCGCAGCGGTTCATAAGGCTGTCTAGGGGTAAAGCACTGTTTCGGTGCGGGCTGCGAAAGCGGTACCAAATCGTGGCAAACTCTGAATACTAGATATGCTATTTATGGGCCAGTGAGACGGTGGGGGATAAGCTTCATCGTCGAGAGGGAAACAGCCCAGATCACTAGCTAAGGCCCCAAAATGATCGTTAAGTGACAAAGGAGGTGAGAATGCTGAAACAACCAGGAGGTTTGCTTAGAAGCAGCCACCCTTTAAAGAGTGCGTA

16S

TGCAGTCGTACGAATGAATTTTGGCTTGCCAAAATTTAATGAGTGGCGGACGGGTGAGTAACACGTAAGAACCTACCTTTTGGAGAGGGACAACCATTGGAAACGATGGCTAATACCTCGTAATGCTGATAAGTTAAATGATGAACAATCGCCAAAAGATGGGCTTGCGGCTGATTAGCTAGTTGGTGGGGTAAAGGCTTACCAAGGCGATGATCAGTAGCTGGTCTGAGAGGATGATCAGCCACACTGGGACTGAGACACGGCCCAACTCCTACGGGAGGCAGCAGTGAGGAATTTTCCGCAATGGGCGAAAGCCTGACGGAGCAATGCCGCGTGAAGGATGAATGCCTATGGGTTGTAAACTTCTTTTCTCAGAGAAGAATTCTTGACGGTATCTGAGGAATAAGCATCGGCTAACTCTGTGCCGCAGCCGCGGTAAGACAGAGGATGCAAGCGTTATCCGGAATGATTGGGCGTAAAGCGTCTGTAGTGGCTTAAAAAGKCTTAAGTCTGTCAAAGATCAGGCKTAACCCTGGGCCGGCAGGAGAAACTCTTATGCTAGAGTTTGGTAGGGGCAGAGGGAATTCCCGGTGGAGCGGTGAAATGCGTAGAGATCGGGAGGAACACCAAAGGCGAAAGCACTCTGCTGGGCCACAACTGACACTGAGAGACGAAAGCGAGGGGAGCAAAAGGGATTAGATACCCCTGTAGTCCTCGCCGTAAACGATGGATACTAGATGTTGGATAGGTTAAAGCATTCAGTATCGTAGCTAACGCGTGAAGTATCCCGCCTGGGGAGTATGCTCGCAAGAGGAAACTCAAAGGAATTACGGGGGCCCGCACAAGCGGTGGAGCATGTGGTTTAATTCGATGCAACGCGAAGAACCTTACCAGGACTTGACATGCCACTTTTTCCCTGAAAGGGGAAGTTCCCGAGTGGACACAGGTGGTGCATGGCTGTCGTCAGCTCGTGTCTGAGATGTTGGGTTAAGTCCCGCAACGAGCGCAACCCTTGTTTTGAATTGCCATTATTGGGAAATTCAAAAGACTGCCGGTGACAAGCCGGAGGAAGGTGAGGATGACGTCAAGTCAGCATGCCCCTTACGTCCTGGGCGACACACGTGCTACAATGGCCGGGACAAAGAGATGCAAACCCGCGAGGGCTAGCCAACCTCAAAAACCCGGTTTYARTTTGGATTGCAGGTTGCAAATCGCATGCMTGAARTTGGAATCGCTAGTAATCGCAGGTCAGCCATACTGCGGTGAATACGTTCCCGGGCCTTGTACACACCGCCCGTCACACCATGGGAGCTGGTATGCCCAAGTCGTTACCCCAACC

Chloroplast CS1\_4

TCGGCTGATTTGTTGGTGGGGTAAAGGCTTACCAAGGCGATGATCAGTAGCTGGTCTGAGAGGATGATCAGCCACACTGGGACTGAGACACGGCCCAGACTCCTACGGGAGGCAGCAGTGAGGAATTTTCCGCAATGGGCGAAAGCCTGACGGAGCAATGCCGCGTGAAGGATGAAGGCCTATGGGTTGTAAACTTCTTTTCTCAGAGAAGAATTCTTGACGGTATCTGAGGAATAAGCATCGGCTAACTCTGTGCCAGCAGCCGCGGTAAGACAGAGGATGCAAGCGTTATCCGGAATGATTGGGCGTAAAGCGTCTGTAGGTGGCTTAAAAAGTCTTCTGTCAAAGATCAGGGCTTAACCCTGGGCCGGCAGGAGAAACTCTTAGGCTAGAGTTTGGTAGGGGCAGAGGGAATTCCCGGTGGAGCGGTGAAATGCGTAGAGATCGGGAGGAACACCAAAGGCGAAAGCACTCTGCTGGGCCACAACTGACACTGAGAGACGAAAGCGAGGGGAGCAAAAGGGATTAGATACCCCTGTAGTCCTCGCCGTAAACGATGGATACTAGATGTTGGATAGGTTAAAGCATTCAGTATCGTAGCTAACGCGTGAAGTATCCCGCCTGGGGAGTATGCTCGCAAGAGTGAAACTCAAAGGAATTGACGGGGGCCCGCACAAGCGGTGGAGCATGTGGTTTAATTCGATGCAACGCGAAGAACCTTACCAGGACTTGACATGCCACTTTTTCCCTGAAAGGGGAAGTTCCCGAGTGGACACAGGTGGTGCATGGCTGTCGTCAGCTCGTGTCTTGAGATGTTGGGTTAAGTCCCGCAACGAGCGCAACCCTTGTTTTGAATTGCCATTATTGGGAAATTCAAAAGACTGCCGGTGATAAGCCCGGAGGAAA

Nuclear NS1\_4

CTTGTCTCAAAGATTAAGCCATGCATGTCTAAGTATAAACTGCTTTATACTGTGAAACTGCGAATGGCTCATTAAATCAGTTATAGTTTATTTGATGGTACCTACTACTCGGATACCCGTAGTAAATCTAGAGCTAATACGTGCGTAAATCCCGACTTCTGGAAGGGACGTATTTATTAGATAAAAGGCCGACCGGGCTCTGCCCGACTCGCGGTGAATCATGATAACTTCACGAATCGCATGGCCTCGTGCCGGCGATGTTTCATTCAAATTTCTGCCCTATCAACTTTCGATGGTAGGATAGAGGCCTACCATGGTGGTAACGGGTGACGGAGGATTAGGGTTCGATTCCGGAGAGGGAGCCTGAGAAACGGCTACCACATCCAAGGAAGGCAGCAGGCGCGCAAATTACCCAATCCTGACACAGGGAGGTAGTGACAATAAATAACAATACTGGGCCTTTTCAGGTCTGGTAATTGGAATGAGTACAATCTAAACCCCTTAACGAGGATCAATTGGAGGGCAAGTCTGGTG