

TABLE S1. Nucleotide sequences of the *novel candidate (nc)* genes

<b>nc1</b>	originates from metagenome (MG-RAST: 4441580)
1	ATGGGTGAGGTAATGGACATACAAAATAGGCGAAGGCTATACCAACTTAGATCTGCGCAAT
61	AAGGACTTCAGCGGCCAAGATTTAACCGACTGCACCTTTAGCCAATGCCAATTTGCGGGC
121	AGTGTCTTCCGCGGCGCCAACCTGACTAACTGTCAGTTTGTATCGATGCCAATTCACAAC
181	AGTGATCCAGACGAGCCCGCAAGCTTTGAATGGGCAAACCTTGCGCGAAGCGGTGTTTAGC
241	CATTGCGACCTAAACATGGTGGCCTTCAATCATTGCCTGGGTACGACCTGAGCTTTAAA
301	GAATGTCAAATGCAGGGCGCAGATTTATCTCTAGGAGATTTCAAATGCCTGTAGGCAAT
361	CTGGATTTGGCCGCACTGACCATTACCGACTGTAATTTTTCTACGGCAATTTATCCAAT
421	ACCTTTTTGGTTGGCTGCAAGCTTTCCAACAATCGACTAACTGAATGCTTGTAGATTTT
481	GCAGATCTAAGTGATGCGGATTTAAGCGACAGCGAAATGTTAATGTTTCAGCACGCAAC
541	TGCAAACCTCAAAGGCGCAGACTTGCGCGGTGCAAGCTTCAACAACCTTAGATTTGAAAAGC
601	TTAGACCTAGAGGGTGTGCGCCTGTACTACAACCAGTTACCTGCGTTAGTAGATTGA
<b>nc2</b>	originates from <i>Vibrio furnissii</i> genome (CP002378.1)
1	ATGAAATTAACCATCACACTTACCAACACACCGATTTCTCAAATCGGGATTGTGCGAAC
61	TTAGTTTTCAACGATTGTTGCTTCTATCAATGCAACTTTAGCCGTGCCGACATGCGTGAT
121	TCCAAATTCATCAATTGCAGCTTTATTGAGAAAAGCGATGTGGAGGGATGTAATTTTGCG
181	TATGCCGATCTCAGAGATGCGTCTTTCAACACTGCCGATTGGCGATGGCTCATTTCAGT
241	GGGGCGGACTGTTTTGGAATTGAACTGAGAGATTGTGATTTAAAAGGTGCCAATTTTTCT
301	CGCGCCCGCTTTGTTAATCAGGTCTCGAACAAAATGTTTTTTTTGTTCCGCGTACATCACG
361	GGATGTAATTTATCCTATGCCAACTTGGAGCGGCAGTGCCTCGAAAAATGTGATCTGTTT
421	GAAAATCGCTGGATTGGCGCAACTTGCACGGGGTTTCTTTTAAAGAGTCTGATTTAAGC
481	CGGGGTTCATTTTAGAGGACTGCTGGGAACAAGCGCGCTTCCAAGGTTGTGACTTAACC
541	CATATTGAACTGACCGGGTTGGATCCCCGCAGAGTGGATTTAACGGGGGTAAAAATTTGC
601	GATTGGCAGCAAGAACAACCTGCTAGAGAAGCTGGGTGTGATTGTGGTTCCGGACTAA
<b>nc3</b>	originates from metagenome (SRA: SRX032366)
1	ATGAGTCTGATGTTGAAGGGTGAAAAGATTGACCGTAACCGCTTTACCGGCGAGAAGATT
61	GAAAACGGCAGCTTTATGCTGTGTGATTTTTCGGGCGCCGATCTGACCGGCACCGAGTTT
121	ATCGGCTGCCAGTTTTACGATCGTGACAGCCGACAGGGCGGCAATTTACGCCGTGCGATA
181	CTCAAAGACGCCAGCTTCAGAAGCTGCGATTTGTGATGGCCGACTTCCGCTACGCCAGC
241	GCCCTGGGGCTGGAGATCCGCGAATGCCGCGCGCAGGGCGCCGATTTTCGCGGCGCCAGT
301	TTTATGAACATGATCACCAGCCGCATCGGGTTTTGCAGCGCCTATATCACAAAAGCAAC
361	CTGAGCTACGCCAACTTCGCCAAGGTGGTGTGCTGGAGAAATGCGAGCTGTGGGAAAACCGC
421	TGGCAAGGCACACAGGGACTCGGCGCCAGCTTCAGCGGCTCCGATCTGTCCGGCGGCGAG
481	TTCTCCGATTTCGACTGGCGCGCCGCGACGTACCCAGTGCGATTTGAGCAATGCGGAA
541	CTGGGCGAATTGGATTTACGCACCACCGATCTGCAGGGCGTCAAATGGACAGCCACCAG
601	GCCGCGCAGCTGCTGGAGCGGCTGGGGATCGCGATCGTCGGCTGA
<b>nc4</b>	originates from metagenome (SRA: SRX032366)
1	ATGAGTCTGATGTTGAAGGGTGAAAAGATTGACCGTAACCGCTTTACCGGCGAGAAGATT
61	GAAAACGGCAGCTTTATGCTGTGTGATTTTTCGGGCGCCGATCTGACCGGCACCGAGTTT
121	ATCGGCTGCCAGTTTTACGATCGTGACAGCCGACAGGGCGGCAATTTACGCCGTGCGATA
181	CTCAAAGACGCCAGCTTCAGAAGCTGCGATTTGTGATGGCCGACTTCCGCTACGCCAGC
241	GCCCTGGGGCTGGAGATCCGCGAATGCCGCGCGCAGGGCGCCGATTTTCGCGGCGCCAGT
301	TTTATGAACATGATCACCAGCCGCATCGGGTTTTGCAGCGCCTATATCACAAAAGCAAC
361	CTGAGCTACGCCAACTTCGCCAAGGTGGTGTGCTGGAGAAATGCGAGCTGTGGGAAAACCGC
421	TGGCACGGCGCACAGGTGCTCGGCGCCAGCTTCAGCGGCTCCGATCTGTCCGGCGGCGAG
481	TTCTCCGATTTCGACTGGCGCGCCGCGACGTACCCAGTGCGATTTGAGCAATGCGGAA
541	CTGGGCGAATTGGATTTACGCACCACCGATCTGCAGGGCGTCAAATGGACAGCCACCAG
601	GCCGCGCAGCTGCTGGAGCGGCTGGGGATCGCGATCGTCGGCTGA