

**Table S6: Leaderless mRNAs of *C. jejuni* NCTC 11168**

gene <sup>a</sup>	strand	TSS	Start	UTR length	RBS	Leaderless	Sequence -30 to startcodon <sup>b</sup>
<i>cj0153c</i>	-	157298	157298	0	no	yes	ttttaagtcaagattag <b>ggTAAAA</b> Tttttaat <u>ATG</u>
<i>cj0164c</i>	-	164091	164091	0	no	yes	tcatagcaaaaaatattt <b>TAAAG</b> Tagaaaa <u>ATG</u>
<i>cj0461c</i>	-	426422	426422	0	no	yes	taaaatataaattat <b>gcTATAG</b> Ttccgctt <u>ATG</u>
<i>cj0595c</i>	-	553151	553151	0	no	yes	ggtttaaagctaattaac <b>TAAAA</b> Ttaaaaa <u>ATG</u>
<i>cj0641</i>	+	602326	602326	0	no	yes	ttgctttgctaagattt <b>gaTATAA</b> Tttttt <u>ATG</u>
<i>cj0667</i>	+	622768	622768	0	no	yes	aaattgatttttttt <b>gcTAAAA</b> Ttgctttt <u>ATG</u>
<i>cj0846</i>	+	793305	793305	0	no	yes	tatttaattttactt <b>ggtTATAA</b> Tactttt <u>ATG</u>
<i>cj1031</i>	+	963479	963479	0	no	yes	aatttatataaagtatat <b>TAAAA</b> Tatcaaa <u>ATG</u>
<i>cj1152c</i>	-	1085255	1085255	0	no	yes	tttttaaaaaagctatat <b>TATAA</b> Tgttttt <u>ATG</u>
<i>cj1247c</i>	-	1176327	1176327	0	no	yes	aattaagaaaagaaat <b>gcTAAAA</b> Tcatctt <u>ATG</u>
<i>cj1257c</i>	-	1188360	1188360	0	no	yes	aaattataaaaaatata <b>TACA</b> Aatcc <u>ATG</u>
<i>cj1349c</i>	-	1282299	1282299	0	no	yes	aatcgccaaaaatattt <b>TATAA</b> Ttctttt <u>ATG</u>
<i>cj1565c</i>	-	1496400	1496400	0	no	yes	atgttataagagtttt <b>gcTATAA</b> Ttatttt <u>ATG</u>
<i>cj1620c</i>	-	1548761	1548760	1	no	yes	aattttaaaatttttt <b>gtTATA</b> CTctcatt <u>TATG</u>
<i>cj1711c</i>	-	1624461	1624459	2	no	yes	taattaagaatttttag <b>gaTAAAA</b> Ttatttc <u>TtATG</u>
<i>cj0430</i>	+	391708	391711	3	no	yes	atattttaagtttaattttg <b>TAAAA</b> Ttttc <u>AttATG</u>
<i>cj1449c</i>	-	1388187	1388184	3	no	yes	tttttaaaaaatattttag <b>TATAA</b> Tttttac <u>AatATG</u>
<i>cj1458c</i>	-	1396196	1396193	3	no	yes	aatttaaaaggaaatttta <b>TAAAA</b> Tcacgc <u>AttATG</u>
<i>cj0792</i>	+	742472	742477	5	no	yes	tccttataaaaaattt <b>gcTAAAA</b> Ttatacta <u>TaaaaATG</u>
<i>cj0748</i>	+	703019	703027	8	no	yes	ataatcagtaattttttta <b>TAGA</b> ATacctat <u>Gcgg</u> aataATG
<i>cj1441c</i>	-	1380906	1380897	9	maybe	no	ttttgaggatttttaatta <b>TAAAA</b> Taatgaa <u>AGG</u> taaaatATG
<i>cj0318</i>	+	288447	288457	10	yes	no	aaaaatagttaaaatt <b>gcTAA</b> ACTaaggg <u>AtGAGA</u> attaATG
<i>cj0945c</i>	-	886130	886119	11	yes	no	cttttagaaaaaata <b>gtTATAA</b> Ttaaagcc <u>AAGGA</u> aatataATG
<i>cj1638</i>	+	1562943	1562955	12	yes	no	attatagccaaatctttg <b>TATAA</b> Ttttaac <u>AAGG</u> tttgtaaaATG
<i>cj0829c</i>	-	776786	776773	13	yes	no	ttttttaatcattta <b>gaTAAAA</b> Taccaaa <u>AcaAAGG</u> ttaacaATG
<i>cj1350</i>	+	1282348	1282361	13	yes	no	ttttggcgaattttgt <b>ggTAGA</b> ATtacatt <u>AaaaAGGA</u> aatttGTG
<i>cj1618c</i>	-	1546343	1546330	13	yes	no	aaatttcataaatttt <b>ggTAAAA</b> Tgtttac <u>TctAGGA</u> aatataATG

a) Genes are sorted on length of the 5' UTR.

b) Capital letters indicate relevant motifs: promoters (bold, red typeface for the  $\sigma^{70}$  promoters with TAnaaT or gnTAnaaT motifs), TSS (green, underlined and bold typeface), putative ribosome binding site (blue, bold typeface) and translational startcodon.