

Table S2. *C. jejuni* genes with primary and secondary promoters.

Gene	Promoter sequences from upstream -26 to ATG. Sigma factor recognition sequences are indicated ^a
Cj0054c	acctattaaactta TATAAT aaaga <u>A</u> cttat<N10>tag CTAGAAT ggaat <u>T</u> tattttaactaAAGGttgaaaaATG σ_{70} σ_{70}
Cj0102	tttagaaaatatt gaTAAAAT taaaa <u>A</u> att<N40>ttttat ggTAAAAT agtcct <u>T</u> ca<N10>ccaAGGAgaattATG σ_{70} σ_{70}
Cj0205	GGCA cgctttt TGCTT cagggcttat <u>A</u> gtg<N212>ttct gaTATAAT ttttc <u>T</u> tttgc<N10>caAGGAttaaaATG σ_{54} σ_{70}
Cj0342c	tatccaagttcttt TAAATAT tattat <u>G</u> c TAAAAT cttaaa <u>T</u> ttatttaatgttTAGGAaagactATG σ_{70} σ_{70}
Cj0476	aagaacaattat ggGATAAT gtagc <u>A</u> ctttt<N150>gttt TAAAAT aaacat <u>T</u> ga<N95>aaAGGAgaaaaATG σ_{70} σ_{70}
Cj0547	GGCA cagtttt TGCTT attatTTTT <u>T</u> tatta<N45>ttgt CGATAT aagctttt <u>A</u> ac<N20>aAGGAttttaaaATG σ_{54} σ_{28}
Cj0897c	tattggttggttt gaTACAAT tatat <u>T</u> ttta<N35>aa TAGAAT tagaac <u>T</u> tga<N10>ttAAGGtaagaaaaATG σ_{70} σ_{70}
Cj1001	ataaacataataac CGATTT tatatag <u>A</u> aagtt<N10>tag aTAAAAT ccaaa <u>G</u> ctt<N10>caAGGAtaatcaATG σ_{28} σ_{70}
Cj1103	aaagcttaaatgt gtTAAAAT acatt <u>A</u> accta<N135>aatacag TAAAAT atcttct <u>A</u> aatAAGGttgaaaaATG σ_{70} σ_{70}
Cj1156	ttaaaaaaatattttt TATAAT tata <u>A</u> aca<N25>at TAAAAT caca <u>T</u> ttttac<N20>ttacGAGGacttATG σ_{70} σ_{70}
Cj1316c	atatataaaatca TAAAAT aaaattt <u>A</u> attataa CGATAT tatatattt <u>A</u> agactaattaaAGGAaatttATG σ_{70} σ_{28}
Cj1361c	aataagattttag gcTAAAAT tt gcTAGAAT ttacatt <u>A</u> aactttaatgattATGGtttaaaaATG σ_{70} σ_{70}

a) Promoter sequences indicated are: σ^{28} , CGATwT at -10 position; σ^{54} , GGcA-N7-TGCTT at -24/-12 positions; σ^{70} , gnTAnaAT at -10 position. All relevant sequences are shown in capital letters; bold typeface indicates promoter sequences, bold and underlined typeface indicate TSS, capital letters alone indicate ribosome binding sites and startcodons. <Nxx> indicates length of sequences not shown.