Supplementary Information

Table S1. Bacterial strains and plasmids used in this study.

Strains or Plasmids	Genotype[a]	Source
E.coli		
K12	*endA1, hsdR17*(r_{K12}-, m_{K12+}), *supE44, thi-1, recA1, gyrA96, relA1, lacF'*[*proA*$^+$*B*$^+$, *lacI*^{q}*ZDM15*::tn10(tet^R)]	Clontech
DH5α	*endA1 hsdR17* (r_K^- m_K^-) *supE44 thi-1 recA1 gyrA relA1* Δ*(lacZYA-argF)U169 deoR [ϕ80dlac* Δ*(lacZ* Δ*M15)*]	Invitrogen
C. jejuni		
AS144	*C. jejuni* NCTC 11168	National Collection of Type Cultures
AS27	AS144 Δ*tonB1*::cam^R	(1)
AS211	AS144 ΔCj0178::cam^R	(2)
AS240	AS144 Δ*tonB2*::cam^R	(1)
AS241	AS144 Δ*tonB1*+*tonB2*::cam^{R}kan^R	(1)
AS242	AS144 Δ*tonB2*+*tonB3*::cam^{R}kan^R	(1)
AS256	AS144 Δ*p19*::cam^R	This study
AS260	AS144 ΔCj1658::cam^R	This study
AS265	AS144 Δ*ceuE*::cam^R	(2)
AS269	AS144 Δ*cfrA*::cam^R	(2)
AS287	AS144 Δ*spoT*::cam^R	(3)
AS335	AS144 Δ*chuA*::cam^R	(4)
AS450	AS144 Δ*p19*+ΔCj1658::cam^R	This study
AS498	AS144 Δ*acs*::cam^R	This study
AS735	AS144 Δ*truB*::cam^R	This study
AS736	AS144 ΔCj1340c::cam^R	This study
AS737	AS144 Δ*ald*::cam^R	This study
AS738	AS144 ΔCj1341c::cam^R	This study
AS739	AS144 ΔCj0020c::cam^R	This study
AS740	AS144 ΔCj0358::cam^R	This study
AS741	AS144 Δ*mdaB*::cam^R	This study
AS742	AS144 ΔCj1026c::cam^R	This study
AS743	AS144 Δ*ccoQ*::cam^R	This study
AS745	AS144 ΔCj1167::cam^R	This study
AS746	AS144 Δ*pstC*::cam^R	This study
AS747	AS144 ΔCj0073c::cam^R	This study
AS748	AS144 ΔCj0818::cam^R	This study
AS749	AS144 ΔCj1613c::cam^R	This study
AS750	AS144 ΔCj1663::cam^R	This study
AS751	AS144 Δ*flgR*::cam^R	This study
AS752	AS144 ΔCj1377c::cam^R	This study
AS753	AS144 Δ*exbD1:*:cam^R	This study
AS754	AS144 ΔCj0045c::cam^R	This study
AS755	AS144 ΔCj0202c::cam^R	This study
AS756	AS144 ΔCj0385c::cam^R	This study

AS757	AS144 Δ*flgH*::cam^R	This study
AS758	AS144 Δ*cfbpB*::cam^R	This study
AS759	AS144 ΔCj1335::cam^R	This study
AS760	AS144 ΔCj1036c::cam^R	This study
AS761	AS144 Δ*flhB*::cam^R	This study
AS762	AS144 Δ*cfbpA*::cam^R	This study
AS763	AS144 ΔCj0040::cam^R	This study
AS764	AS144 Δ*flgD*::cam^R	This study
AS765	AS144 Δ*flgK*::cam^R	This study
AS766	AS144 ΔCj0041::cam^R	This study
AS767	AS144 ΔCj0416::cam^R	This study
AS768	AS144 Δ*flgE*::cam^R	This study
AS769	AS144 ΔCj1342c::cam^R	This study
AS772	AS144 Δ*ceuB*::cam^R	This study
AS773	AS144 ΔCj0494::cam^R	This study
AS774	AS144 Δ*trpF*::cam^R	This study
AS775	AS144 ΔCj1241::cam^R	This study
AS776	AS144 ΔCj0295::cam^R	This study
AS777	AS144 ΔCj0524::cam^R	This study
AS778	AS144 Δ*chuB*::cam^R	This study
AS778	AS144 ΔCj0741::cam^R	This study
AS779	AS144 ΔCj1661::cam^R	This study
AS780	AS144 ΔCj0148c::cam^R	This study
AS781	AS144 ΔCj0554::cam^R	This study
AS782	AS144 ΔCj0561::cam^R	This study
AS783	AS144 ΔCj1406c::cam^R	This study
AS784	AS144 ΔCj0587::cam^R	This study
AS785	AS144 ΔCj0819::cam^R	This study
AS786	AS144 Δ*flgM*::cam^R	This study
AS787	AS144 ΔCj0672::cam^R	This study
AS789	AS144 ΔCj1209::cam^R	This study
AS790	AS144 ΔCj1356c::cam^R	This study
AS791	AS144 ΔCj0786::cam^R	This study
AS792	AS144 ΔCj0949c::cam^R	This study
AS793	AS144 ΔCj1211::cam^R	This study
AS794	AS144 ΔCj0814::cam^R	This study
AS795	AS144 ΔCj1242::cam^R	This study
AS796	AS144 ΔCj1623::cam^R	This study
AS797	AS144 Δ*flgG*::cam^R	This study
AS798	AS144 Δ*exbB2*::cam^R	This study
AS799	AS144 ΔCj0634::cam^R	This study
AS800	AS144 Δ*cfbpC*::cam^R	This study
AS801	AS144 Δ*hypC*::cam^R	This study
AS802	AS144 ΔCj0260c::cam^R	This study
AS803	AS144 Δ*flgI*::cam^R	This study
AS804	AS144 Δ*chuD*::cam^R	This study
AS805	AS144 Δ*exbB1*::cam^R	This study
AS806	AS144 ΔCj1207c::cam^R	This study
AS807	AS144 Δ*flgE2*::cam^R	This study
AS808	AS144 ΔCj0062c::cam^R	This study
AS809	AS144 Δ*chaN*::cam^R	This study

AS830	AS144 Δ*flaG*::cam^R	This study
AS831	AS144 Δ*flgG2*::cam^R	This study
AS832	AS144 ΔCj0309c::cam^R	This study
AS833	AS144 ΔCj1388::cam^R	This study
AS940	AS144 Δ*cmeA*::cam^R	This study
AS941	AS144 ΔCj1255::cam^R	This study
AS943	AS144 ΔCj0253::cam^R	This study
AS959	AS144 Δ*chuC*::cam^R	This study
AS961	AS786 + *flgM*::cam^{R}kan^R	This study
AS962	AS808 + Cj0062c::cam^{R}kan^R	This study
AS963	AS809 + *chaN*::cam^{R}kan^R	This study
AS964	AS747 + Cj0073c::cam^{R}kan^R	This study
AS965	AS794 + Cj0814::cam^{R}kan^R	This study
AS966	AS796 + Cj1623::cam^{R}kan^R	This study
AS967	AS742 + Cj1026c::cam^{R}kan^R	This study
AS968	AS774 + *trpF*::cam^{R}kan^R	This study
AS970	AS803 + *flgI*::cam^{R}kan^R	This study
AS971	AS802 + Cj0260c::cam^{R}kan^R	This study
AS972	AS760 + Cj1036c::cam^{R}kan^R	This study
AS973	AS757 + *flgH*::cam^{R}kan^R	This study
AS974	AS144 ΔCj0171::cam^R	This study
AS975	AS746 + *pstC*::cam^{R}kan^R	This study
AS976	AS805 + *exbB1*::cam^{R}kan^R	This study
AS977	AS766 + Cj0041::cam^{R}kan^R	This study
AS979	AS764 + *flgE*::cam^{R}kan^R	This study
AS980	AS765 + Δ*flgK*::cam^{R}kan^R	This study
AS981	AS752 + Cj1377c::cam^{R}kan^R	This study
AS982	AS751 + *flgR*::cam^{R}kan^R	This study
AS983	AS761 + *flhB*::cam^{R}kan^R	This study
AS984	AS740 + Cj0358::cam^{R}kan^R	This study
AS991	AS755 + Cj0202c::cam^{R}kan^R	This study
AS992	AS764 + *flgD*::cam^{R}kan^R	This study
AS993	AS798 + *exbB2*::cam^{R}kan^R	This study
AS1014	AS498 + *acs*::cam^{R}kan^R	This study
AS1030	AS144 ΔCj0977::cam^R	This study
AS1060	AS144 ΔCj1383c::cam^R	This study
AS1061	AS144 ΔCj0344::cam^R	This study
AS1062	AS144 Δ*flgL*::cam^R	This study
AS1063	AS144 ΔCj1159c::cam^R	This study
AS1064	AS144 Δ*folP*::cam^R	This study
AS1065	AS144 Δ*exbD2*::cam^R	This study
AS1066	AS144 ΔCj0947c::cam^R	This study
AS1067	AS144 Δ*pseB*::cam^R	This study
AS1068	AS144 ΔCj0900c::cam^R	This study
AS1071	AS144 Δ*rrc*::cam^R	This study
AS1123	AS1060 + Cj1383c::cam^{R}kan^R	This study
AS1124	AS1061 + Cj0344::cam^{R}kan^R	This study
AS1125	AS1066 + Cj0947c::cam^{R}kan^R	This study
AS1126	AS1071 + *rrc*::cam^{R}kan^R	This study
AS1127	AS1131 + *acnB*::cam^{R}kan^R	This study
AS1128	AS1062 + *flgL*::cam^{R}kan^R	This study

AS1129	AS1067 + *pseB*::cam^{R}kan^R	This study
AS1131	AS144 Δ*acnB*::cam^R	This study
AS1132	AS144 Δ*tonB3*::cam^R	(1)
AS1133	AS1132 + *tonB3*::cam^{R}kan^R	(1)
AS1330	AS144 Δ*motAB*::cam^R	This study
AS1334	AS144 Δ*ccoQ*::kan^R	This study
AS1337	AS144 Δ*ccoQ*Δ*flgH*::kan^{R}cam^R	This study
AS1338	AS144 Δ*ccoQ*Δ*flhB*::kan^{R}cam^R	This study
AS1339	AS144 Δ*ccoQ*Δ*flgD*::kan^{R}cam^R	This study
AS1340	AS144 Δ*ccoQ*Δ*motAB*::kan^{R}cam^R	This study
AS1341	AS1330 + *motAB*::kan^{R}cam^R	This study
Plasmids		
pRY111	Cam^R resistance gene	Yao (5)
pILL600	Kan^R resistance gene	Reid (6)
pRRK	Cloning vector used for complementation of mutants, kan^R	Reid (6)
pUC19	Cloning vector, amp^R	Biolabs

[a-] cam^R, chloamphenicol resistance gene, kan^R, kanamycin resistance gene, amp^R, ampicillin resistance gene.

Table S2. Primers used in this study.

Primer Name	Primer Sequence (5'- 3')[a]
acs_AS	CGACTCTAGAGGATCCTCCACGCCCTGTGACAAATCC
acs_AS_inverse	GAACACCGCCGAGCAGCTCATTAGGTGCATCAGCA
acs_SE	CGGTACCCGGGGATCCAACGGCTGATGGTGCTTTTCG
acs_SE_inverse	GAACTAAAGGGCGCAGGCGAGTCATTGTTTGCTTT
ald_AS	CGACTCTAGAGGATCCCCATCAGCTCCGCCTATGC
ald_AS_inverse	GAACACCGCCGAGCATAAAGCAGGTGCCATCTTCC
ald_SE	CGGTACCCGGGGATCCATGGCAAGAGATGAAAGCACAC
ald_SE_inverse	GAACTAAAGGGCGCAAGTGTGCAAAGATGCGGATA
ccoQ_AS_cam	CGACTCTAGAGGATCCCATGACAAGCCGAACACTGAAC
ccoQ_AS_inverse_cam	GAACACCGCCGAGCACAAAAACATCACGAGTGCAAA
ccoQ_AS_inverse_kan	GTTCGCTGGGTTTATCCAAAAACATCACGAGTGCAAA
ccoQ_SE_cam	CGGTACCCGGGGATCCGGACCTGATCTTGCTCGTGTAG
ccoQ_SE_inverse_cam	GAACTAAAGGGCGCATGCAAACCTTGCCTTAAAAGA
ccoQ_SE_inverse_kan	GATGTCTAGAAAGCTTTGCAAACCTTGCCTTAAAAGA
ceuB_AS	CGACTCTAGAGGATCCATTAGCCGCACCAAAAGAAA
ceuB_AS_inverse	GAACACCGCCGAGCACTTCACCCATGCCAACTATG
ceuB_SE	CGGTACCCGGGGATCCATGCAACAACTCACGCAAAA
ceuB_SE_inverse	GAACTAAAGGGCGCATGATTGTAAGCGTTGGGATT
cfbpA_AS	CGACTCTAGAGGATCCAGCAGAACTCACACCCAAAATC
cfbpA_AS_inverse	GAACACCGCCGAGCATCGCAAACCATTCTTTATCCTT
cfbpA_SE	CGGTACCCGGGGATCCCTATGCGTTGGCTTTGCTTTTG
cfbpA_SE_inverse	GAACTAAAGGGCGCAAAGTGGGGAATTCTTTGGGTA
cfbpB_AS	CGACTCTAGAGGATCCATCCTCTTGCCATAGCTCTAGC
cfbpB_AS_inverse	GAACACCGCCGAGCACCAAATAAGCACTTGCACCA
cfbpB_SE	CGGTACCCGGGGATCCTGATTTTGGGTGTGAGTTCTGC
cfbpB_SE_inverse	GAACTAAAGGGCGCACAGCTTTGCCTTCCTTGTTT
cfbpC_AS	CGACTCTAGAGGATCCACAGTAGCCACACGACTTACCC
cfbpC_AS_inverse	GAACACCGCCGAGCACACTCTTTGCGCTTGTCCTC
cfbpC_SE	CGGTACCCGGGGATCCTGATGCAGCTTTGCCTTCCTTG
cfbpC_SE_inverse	GAACTAAAGGGCGCAAAAATCAAGGCATTACGGCTA
chuB_AS	CGACTCTAGAGGATCCTGCTTTTGCCTGAGCCATTTGG
chuB_AS_inverse	GAACACCGCCGAGCATCGCCATTATAATCCGTGGT
chuB_SE	CGGTACCCGGGGATCCGCTGGGCTTGAATACGCTTGG
chuB_SE_inverse	GAACTAAAGGGCGCATGGTTTCGTGTGGGAATTTT
chuC_AS	CGACTCTAGAGGATCCGGTGATTTGGCCTAAGGTTGTG
chuC_AS_inverse	GAACACCGCCGAGCATGAGGCACAAAGCCACAAAT
chuC_SE	CGGTACCCGGGGATCCGCGGACTTATAGGCTTTGTTGG
chuC_SE_inverse	GAACTAAAGGGCGCAAGCCGTTGTTGCTATCTTGC
chuD_AS	CGACTCTAGAGGATCCTGGTGCCTTGCTGTTTGGATAC
chuD_AS_inverse	GAACACCGCCGAGCATGAAAAAGTGCCTACGCTTG
chuD_SE	CGGTACCCGGGGATCCGATGAACCCACTTCAGCCCTTG
chuD_SE_inverse	GAACTAAAGGGCGCATTAAGCCCGCAAAGTCAAAT
Cj0012c_AS	CGACTCTAGAGGATCCATCGCTTTAGCTTTAGCTTCGC
Cj0012c_AS_comp	GGGGAAGCTTTCTAGCACCAAAGAGAAAAGCCGTTA
Cj0012c_AS_inverse	GAACACCGCCGAGCATTTCATTTTCAGCCGCTTCT
Cj0012c_SE	CGGTACCCGGGGATCCTCGCATCGCTTCTCATTGTAAC
Cj0012c_SE_comp	GATTTAGATGTCTAGAGCAAAGCTTAATTTTTGTGAAA
Cj0012c_SE_inverse	GAACTAAAGGGCGCATGCAAAAATCGCTGAAGATG

Cj0020c_AS	CGACTCTAGAGGATCCCGCTTGAACACTTGGATCGTAG
Cj0020c_AS_inverse	GAACACCGCCGAGCAGCCGTGTTGATCAAGCCTAT
Cj0020c_SE	CGGTACCCGGGGATCCAGAAGAACTTAGCGCAGGCATG
Cj0020c_SE_inverse	GAACTAAAGGGCGCAGCATCTCTTGCCATCAAGGT
Cj0040_AS	CGACTCTAGAGGATCCCGCTCTCATCAAAAGCTCCT
Cj0040_AS_inverse	GAACACCGCCGAGCAAAAAGCTCTAAAACTTGAAGCAAAA
Cj0040_SE	CGGTACCCGGGGATCCCGTTCTACTTCTCCGCCAAA
Cj0040_SE_inverse	GAACTAAAGGGCGCATGGCAAGAATTTCAAGTGAGC
Cj0041_AS	CGACTCTAGAGGATCCTGCGTTTTTGTTTGCTCTTG
Cj0041_AS_comp	GGGGAAGCTTTCTAGGGTAGCATTTGGATTGCTCA
Cj0041_AS_inverse	GAACACCGCCGAGCATCGTGTTTGCTTTCTTTGGA
Cj0041_SE	CGGTACCCGGGGATCCGAAGCGATGAGCAGACTTCA
Cj0041_SE_comp	GATTTAGATGTCTAGTGAAAGAAAAATTTCCTAATGTAATGG
Cj0041_SE_inverse	GAACTAAAGGGCGCAGAAGAAGATACCACAGATGCTAAAAA
Cj0044c_AS	CGACTCTAGAGGATCCCAACGGCAAATAGCGGTGATG
Cj0044c_AS_inverse	GAACACCGCCGAGCAAAATCCATAATTGGCGCTAAAA
Cj0044c_SE	CGGTACCCGGGGATCCTCCAACTCCAACCTAACCCATC
Cj0044c_SE_inverse	GAACTAAAGGGCGCAAGCCGAAAATATCGGAGAGG
Cj0045c_AS	CGACTCTAGAGGATCCAGATGGGTTAGGTTGGAGTTGG
Cj0045c_AS_inverse	GAACACCGCCGAGCATCAATCAAAGGAAAATCAATGC
Cj0045c_SE	CGGTACCCGGGGATCCAAACCCTTCCCCAGATAACTGC
Cj0045c_SE_inverse	GAACTAAAGGGCGCATTCTTTGCGTATCCATGCAG
Cj0062c_AS	CGACTCTAGAGGATCCTTTATCGTAACGGCGTGAAAGC
Cj0062c_AS_comp	GGGGAAGCTTTCTAGTTGAAGGCAAGCGTTCTTTT
Cj0062c_AS_inverse	GAACACCGCCGAGCATCCACAAACAGTGAAAAATGC
Cj0062c_SE	CGGTACCCGGGGATCCGCAGCAATCACTGATGCGTATG
Cj0062c_SE_comp	GATTTAGATGTCTAGGCCCTAGCTTCCAAGCTTTT
Cj0062c_SE_inverse	GAACTAAAGGGCGCATGAAGCCTTTGATATACTGATTTTT
Cj0073c_AS	CGACTCTAGAGGATCCCCTTGCTTGCAAGTTCACAA
Cj0073c_AS_comp	GGGGAAGCTTTCTAGTTGGAATTAGTTTATCTAAAAGCACCT
Cj0073c_AS_inverse	GAACACCGCCGAGCATTTTGCGACAATTTCATTGATT
Cj0073c_SE	CGGTACCCGGGGATCCGCCCTAGAAGCCTTTGCTTT
Cj0073c_SE_comp	GATTTAGATGTCTAGCGGTACAAAATTTCGCAAGC
Cj0073c_SE_inverse	GAACTAAAGGGCGCAGCCTTGCACCTAAACTTTGC
Cj0148c_AS	CGACTCTAGAGGATCCGCGTAGGAATAGAGCGAACACC
Cj0148c_AS_inverse	GAACACCGCCGAGCACAAGCCTTATCTTCCCCTAAAA
Cj0148c_SE	CGGTACCCGGGGATCCAGCAGGAGCAGGAGGAGAGG
Cj0148c_SE_inverse	GAACTAAAGGGCGCACCGAGCGTTTGGATAGAAAA
Cj0171_AS	CGACTCTAGAGGATCCTCGCTAAAGGAATTTGGCAAGG
Cj0171_AS_inverse	GAACACCGCCGAGCAGCAAGGATCTTTTTCCCAAA
Cj0171_SE	CGGTACCCGGGGATCCAATGCACGCCCTGCTTATTTAG
Cj0171_SE_inverse	GAACTAAAGGGCGCAGCTTTTAAAGAAGCTGTGCAAGA
Cj0177_AS	CGACTCTAGAGGATCCCACGCCCAAAACCACTCATACC
Cj0177_AS_comp	GGGGAAGCTTTCTAGTTTTTCATTCCTCCCCCTTA
Cj0177_AS_inverse	GAACACCGCCGAGCATCGCCAAGCAGTATCACATC
Cj0177_SE	CGGTACCCGGGGATCCAGCAAGCAAAAGCAAAGCCAAC
Cj0177_SE_comp	GATTTAGATGTCTAGAAGCAAAGCCAACGCATAGT
Cj0177_SE_inverse	GAACTAAAGGGCGCAACAGGCGTATGGCTGATGTA
Cj0202c_AS	CGACTCTAGAGGATCCTTTTCTGCCTAAATGTTCCAAA
Cj0202c_AS_comp	GGGGAAGCTTTCTAGTTCCTGCACGAAAGACAAGA
Cj0202c_AS_inverse	GAACACCGCCGAGCAGCTGCATTTTCAGCAACAAC

Cj0202c_SE	CGGTACCCGGGGATCCAAAACCAAGCCGTTCAAGAAT
Cj0202c_SE_comp	GATTTAGATGTCTAGAAAACCAAGCCGTTCAAGAAT
Cj0202c_SE_inverse	GAACTAAAGGGCGCAAGCAAAAAGAATGGGTGGTG
Cj0253_AS	CGACTCTAGAGGATCCGCACTTTCAGCGGTTTCTTT
Cj0253_AS_inverse	GAACACCGCCGAGCACGCACTTGAATTCGAATGTTT
Cj0253_SE	CGGTACCCGGGGATCCCGGCTATAATGGGGGTTAAAA
Cj0253_SE_inverse	GAACTAAAGGGCGCAAAAAAGATCGTTTGGATATATTTGA
Cj0260c_AS	CGACTCTAGAGGATCCTTGTGCCGCAGGTGTATTTA
Cj0260c_AS_comp	GGGGAAGCTTTCTAGCGTGCCTTATATGGCTGGAG
Cj0260c_AS_inverse	GAACACCGCCGAGCACATCAAGAATCAAGCGTCCA
Cj0260c_SE	CGGTACCCGGGGATCCAAATTTATCTTGGTTGGGCTGA
Cj0260c_SE_comp	GATTTAGATGTCTAGTGGCATTTAAGTATGAACAAAATCA
Cj0260c_SE_inverse	GAACTAAAGGGCGCAGCATTGGATTTTTAGCTGGTTT
Cj0295_AS	CGACTCTAGAGGATCCACCATAGCCACGCAAGAAGAAG
Cj0295_AS_inverse	GAACACCGCCGAGCAGCAAATCAAGCAATTTCTGC
Cj0295_SE	CGGTACCCGGGGATCCTCAACAGGTGGAGCTAGAAAGC
Cj0295_SE_inverse	GAACTAAAGGGCGCATGAAATTTACACTTTCGCATCC
Cj0309c_AS	CGACTCTAGAGGATCCTTTGCTACCAAAATCGTAGGG
Cj0309c_AS_inverse	GAACACCGCCGAGCAGCAACGCTCATAGCGATATTT
Cj0309c_SE	CGGTACCCGGGGATCCGGCTTCATGTGAATGAAAAATTC
Cj0309c_SE_inverse	GAACTAAAGGGCGCATATGGACAGGAGCTGGAACC
Cj0344_AS	CGACTCTAGAGGATCCAATCCAGCAAAGTCGCAAAT
Cj0344_AS_comp	GGGGAAGCTTTCTAGAATCCAGCAAAGTCGCAAAT
Cj0344_AS_inverse	GAACACCGCCGAGCATTATATTTTATAATATTTTGAAACAT
Cj0344_SE	CGGTACCCGGGGATCCCATTCCGCTAAAGCTTGCTC
Cj0344_SE_comp	GATTTAGATGTCTAGACCAATGCCAAAAAGTCCAG
Cj0344_SE_inverse	GAACTAAAGGGCGCAATTTTCATAAAAGTCTTAATTTTTGA
Cj0358_AS	CGACTCTAGAGGATCCAATCGCAGGAGTTGGCATAGG
Cj0358_AS_comp	GGGGAAGCTTTCTAGCAGCCATTGCTAAACGCATA
Cj0358_AS_inverse	GAACACCGCCGAGCAAGCAGGAGTTGCCATTTCAG
Cj0358_SE	CGGTACCCGGGGATCCTGCTAATCGCATCCTTAGTTGC
Cj0358_SE_comp	GATTTAGATGTCTAGTTAGCCCAAAAGGGATTGAA
Cj0358_SE_inverse	GAACTAAAGGGCGCATGTTGCTGAAACTGCTCCAT
Cj0385c_AS	CGACTCTAGAGGATCCTGCTGCGGCTACAAGTAAATCG
Cj0385c_AS_inverse	GAACACCGCCGAGCAAGCCAAAAATATCCCCCTTC
Cj0385c_SE	CGGTACCCGGGGATCCTCACTTTCATCAAGCCCTCCAC
Cj0385c_SE_inverse	GAACTAAAGGGCGCAAAACAAGCAGGAGTGGTTGC
Cj0416_AS	CGACTCTAGAGGATCCGCTGATCCTAGCCCACAAGA
Cj0416_AS_inverse	GAACACCGCCGAGCATGTCTGCAGAGGATTATTTCCA
Cj0416_SE	CGGTACCCGGGGATCCTGGGCGCAGATAGAGAAACT
Cj0416_SE_inverse	GAACTAAAGGGCGCAGAGATGAAAGTATTGAAAATGCAAA
Cj0494_AS	CGACTCTAGAGGATCCTGCAAGGGTAAAAATTCCTGA
Cj0494_AS_inverse	GAACACCGCCGAGCAAATTTACTGGGCTTTGTGCTG
Cj0494_SE	CGGTACCCGGGGATCCTTGCTGCTTCTATGGGCTTT
Cj0494_SE_inverse	GAACTAAAGGGCGCAAAAGAACAGCTTCCAAAACG
Cj0524_AS	CGACTCTAGAGGATCCCGAAGCCGCTAAAACGATGC
Cj0524_AS_inverse	GAACACCGCCGAGCACGACTAAAACCTATAGCATGAGCA
Cj0524_SE	CGGTACCCGGGGATCCAGTGTTGGAGGAGTTGTAACGG
Cj0524_SE_inverse	GAACTAAAGGGCGCACACAACAGCTGAAGGAATTTCA
Cj0554_AS	CGACTCTAGAGGATCCCGCTCCTATCACTCACACTCAC
Cj0554_AS_inverse	GAACACCGCCGAGCATTCACCAAAAGCCAGTGAGA

Cj0554_SE	CGGTACCCGGGGATCCTTACTGCTGCTATTGCTTTGCC
Cj0554_SE_inverse	GAACTAAAGGGCGCATTCAAAACCCTGCACACAAA
Cj0561c_AS	CGACTCTAGAGGATCCAGCTTTTCCGTCTGCATCATTG
Cj0561c_AS_inverse	GAACACCGCCGAGCAAGCCGTAGCTGATTGGGTTA
Cj0561c_SE	CGGTACCCGGGGATCCAAGGAGTGGCAAGAATTTCTGC
Cj0561c_SE_inverse	GAACTAAAGGGCGCATTCAACCTTGGCAAACAGTG
Cj0587_AS	CGACTCTAGAGGATCCTGAACAAAACCGCCCGTACTAG
Cj0587_AS_inverse	GAACACCGCCGAGCAAAAAAGCCCATAAAATGCATAAA
Cj0587_SE	CGGTACCCGGGGATCCGGAGAAGAGGCAGGTTCTAAGC
Cj0587_SE_inverse	GAACTAAAGGGCGCATGCCTTAAAATCCGAACAAGA
Cj0634_AS	CGACTCTAGAGGATCCTGATCCACCCTTAGGAATGC
Cj0634_AS_inverse	GAACACCGCCGAGCATTGATCTAAACCATTGGCAAAA
Cj0634_SE	CGGTACCCGGGGATCCCGCAGTAAAACTCGCACAAA
Cj0634_SE_inverse	GAACTAAAGGGCGCATAAGGCTGTTGTGGTTGCAC
Cj0672_AS	CGACTCTAGAGGATCCTGCCTGTATTTTGCGAAAGC
Cj0672_AS_inverse	GAACACCGCCGAGCAGAAGCAAAGCAAAGATAATATCCA
Cj0672_SE	CGGTACCCGGGGATCCCGGCTGCTTGTTATGGGTAT
Cj0672_SE_inverse	GAACTAAAGGGCGCACGCGGATGATATTTTTACTAGTTTT
Cj0741_AS	CGACTCTAGAGGATCCTGTGGTTCATTTGCTTCAGGAG
Cj0741_AS_inverse	GAACACCGCCGAGCACAGCATCTGTTGCCTTGGTA
Cj0741_SE	CGGTACCCGGGGATCCGTGGGTGGCATAGGCAGAC
Cj0741_SE_inverse	GAACTAAAGGGCGCAGAAAAGCAGTTTGCGGATCT
Cj0786_AS	CGACTCTAGAGGATCCTGTCTTCTTTGCTCTTCCATGA
Cj0786_AS_inverse	GAACACCGCCGAGCATCAATTCCAAAAATAACACCAA
Cj0786_SE	CGGTACCCGGGGATCCATCGTTTCAGGCTTTGGTGA
Cj0786_SE_inverse	GAACTAAAGGGCGCAAAAAATCAAGATTTAAAAGTGCAAG
Cj0814_AS	CGACTCTAGAGGATCCATCACCTTCTATGCCACTCTGC
Cj0814_AS_comp	GGGGAAGCTTTCTAGACAGGTTGTCCACCTTGAGC
Cj0814_AS_inverse	GAACACCGCCGAGCATCTTGCTAGAGTTTGTGAATTTGC
Cj0814_SE	CGGTACCCGGGGATCCAAGGGCATTTAGGCATTTACGC
Cj0814_SE_comp	GATTTAGATGTCTAGCAACAAGCATGGGGATTGAT
Cj0814_SE_inverse	GAACTAAAGGGCGCAGAACGCAATGGATGGACTTT
Cj0818_AS	CGACTCTAGAGGATCCATGGTTGTAAGCTCAGTTTTAATGG
Cj0818_AS_inverse	GAACACCGCCGAGCAAACCCCAGCACCTACTACCC
Cj0818_SE	CGGTACCCGGGGATCCTTGGAAGTGATTCAAGGCAAG
Cj0818_SE_inverse	GAACTAAAGGGCGCAAATTGACATGATGGCTGCAA
Cj0819_AS	CGACTCTAGAGGATCCAGCTATGGGGACGCAAAGTA
Cj0819_AS_inverse	GAACACCGCCGAGCATTTTCTAGGGTCTTGATAGAATAAAAA
Cj0819_SE	CGGTACCCGGGGATCCCCATTTTAGCGCCTTTTCAA
Cj0819_SE_inverse	GAACTAAAGGGCGCAGTCTTTCTAGGGCGAACAGC
Cj0900c_AS	CGACTCTAGAGGATCCACAAGCAAACTCGCCCTCTA
Cj0900c_AS_inverse	GAACACCGCCGAGCACCAAAACAATCATTAAAAACACAAA
Cj0900c_SE	CGGTACCCGGGGATCCAAAATTACCGGTGTTCCTCGT
Cj0900c_SE_inverse	GAACTAAAGGGCGCATCAATCATCAAATGATGCAAAA
Cj0947c_AS	CGACTCTAGAGGATCCTTTTAAACCGCTTTGGGAAA
Cj0947c_AS_comp	GGGGAAGCTTTCTAGAAACGAACAAGAAAAAGAGCAAA
Cj0947c_AS_inverse	GAACACCGCCGAGCACCCCGCTTACATCTTTTTCA
Cj0947c_SE	CGGTACCCGGGGATCCGGAGCTGAGCTTGTTTGCTT
Cj0947c_SE_comp	GATTTAGATGTCTAGTGCAGAACAAGAAAGGCAAA
Cj0947c_SE_inverse	GAACTAAAGGGCGCATTTTGTTTTTGGCCCTCAAG
Cj0949c_AS	CGACTCTAGAGGATCCTCGCACTTGAAAGTACTGCAAC

Cj0949c_AS	CGACTCTAGAGGATCCTCGCACTTGAAAGTACTGCAAC
Cj0949c_AS_inverse	GAACACCGCCGAGCACCAAAATCGCGTATCCAAGT
Cj0949c_AS_inverse	GAACACCGCCGAGCACCAAAATCGCGTATCCAAGT
Cj0949c_SE	CGGTACCCGGGGATCCATGCTTTGTCACCCAATGGATG
Cj0949c_SE	CGGTACCCGGGGATCCATGCTTTGTCACCCAATGGATG
Cj0949c_SE_inverse	GAACTAAAGGGCGCATGAGCATTATTTGCCTTTGC
Cj0949c_SE_inverse	GAACTAAAGGGCGCATGAGCATTATTTGCCTTTGC
Cj0977_AS	CGACTCTAGAGGATCCAAGGCTATGCTTGGGCTGATAC
Cj0977_AS_inverse	GAACACCGCCGAGCATCGTCAAAAAGTGCATGAGC
Cj0977_SE	CGGTACCCGGGGATCCGGCTATGAAGCACCCAAAAGAG
Cj0977_SE_inverse	GAACTAAAGGGCGCAGTTGTTAGCACCGATGAGCA
Cj1036c_AS	CGACTCTAGAGGATCCAAGCTTAAATCATAACGCTTCCA
Cj1036c_AS_comp	GGGGAAGCTTTCTAGGACGCCAACCTCGTTTTATT
Cj1036c_AS_inverse	GAACACCGCCGAGCACTTGCACCACTCTTGCTTGA
Cj1036c_SE	CGGTACCCGGGGATCCTCCAAGTCCTGGAAAAATAGGA
Cj1036c_SE_comp	GATTTAGATGTCTAGTGCAATAGCTGCTACGCTTAAA
Cj1036c_SE_inverse	GAACTAAAGGGCGCACAGGATGAATTTTCCGAATTT
Cj1159c_AS	CGACTCTAGAGGATCCAATGTGCAAAAAGCACTAACAA
Cj1159c_AS_inverse	GAACACCGCCGAGCAGCCTATATGATAAATCACGGGAGA
Cj1159c_SE	CGGTACCCGGGGATCCTGTGCTTTGCATTCCTATCG
Cj1159c_SE_inverse	GAACTAAAGGGCGCATTTATTGCCTTAACAGCTCTAAAAA
Cj1167_AS	CGACTCTAGAGGATCCACCAGGGATTCGTCAGTACATC
Cj1167_AS_inverse	GAACACCGCCGAGCAGGTGTCATTTGGATTTGTTGC
Cj1167_SE	CGGTACCCGGGGATCCACCTCAGCTATTCTTCCTACGC
Cj1167_SE_inverse	GAACTAAAGGGCGCAAGAACTCGAAAAAGCCGTGA
Cj1207c_AS	CGACTCTAGAGGATCCTAATTGCTCAATCGCACCTG
Cj1207c_AS_inverse	GAACACCGCCGAGCATTATTTTCACCATTAGCGACTTT
Cj1207c_SE	CGGTACCCGGGGATCCTTTGCAATGCCCTAAAATCTG
Cj1207c_SE_inverse	GAACTAAAGGGCGCAGCTTTAGGTGGGGTAAATGGA
Cj1209_AS	CGACTCTAGAGGATCCGCGAATTTCTCTTCCGGCATTG
Cj1209_AS_inverse	GAACACCGCCGAGCACTGCAAATTCACCTGCAAAA
Cj1209_SE	CGGTACCCGGGGATCCAGAACAAGCCAAAGCTAAAGCC
Cj1209_SE_inverse	GAACTAAAGGGCGCATACCGAGCAAGCTATGGACA
Cj1211_AS	CGACTCTAGAGGATCCCCTTTGGGGCTTGGGCTTAG
Cj1211_AS_inverse	GAACACCGCCGAGCATGGGTGATAATTCTAAGGCTCAA
Cj1211_SE	CGGTACCCGGGGATCCGCTAGTTTGGCATGGGCTATTG
Cj1211_SE_inverse	GAACTAAAGGGCGCATTTGCTATGGTTTTACCTGTGC
Cj1241_AS	CGACTCTAGAGGATCCAGCACCAAGATCAATGTCTTCG
Cj1241_AS_inverse	GAACACCGCCGAGCATCCTGCTATAAAGCGTGCAA
Cj1241_SE	CGGTACCCGGGGATCCACTCGCCAAACTTAAACCAAGC
Cj1241_SE_inverse	GAACTAAAGGGCGCATTGCCTTTTCAACCATACCTG
Cj1242_AS	CGACTCTAGAGGATCCTTCCTGCTTGACGCATCATCC
Cj1242_AS_inverse	GAACACCGCCGAGCATTGAAATTCATCTGCCGAAGT
Cj1242_SE	CGGTACCCGGGGATCCGCCTTTTCAACCATACCTGCAC
Cj1242_SE_inverse	GAACTAAAGGGCGCATCAAGCAAATTTACTAACGAAGACA
Cj1255_AS	CGACTCTAGAGGATCCTTTTTGAATTTGAGATCACTTTCTTG
Cj1255_AS_inverse	GAACACCGCCGAGCACCGTTACTCCTGCGATAAGC
Cj1255_SE	CGGTACCCGGGGATCCGAAACACTTTTTAATGCAAAATTAGA
Cj1255_SE_inverse	GAACTAAAGGGCGCACGGACTAGGTGGAAAAAGCA
Cj1335_AS	CGACTCTAGAGGATCCTGCCTTGAAGAGCATCTTTTG
Cj1335_AS_inverse	GAACACCGCCGAGCATTTTGGGTAGGGGTGAAGGT

Cj1335_SE	CGGTACCCGGGGATCCACCTACAACTGCACCGAAGG
Cj1335_SE_inverse	GAACTAAAGGGCGCACAAAAACATTGAGGCTTTAAACAA
Cj1340c_AS	CGACTCTAGAGGATCCGAGCCGCTTGAGTTGCCTTAG
Cj1340c_AS_inverse	GAACACCGCCGAGCATTGCTCCACCTTCAGTAGCA
Cj1340c_SE	CGGTACCCGGGGATCCATTGCTCTTGAGTGTGCTACGC
Cj1340c_SE_inverse	GAACTAAAGGGCGCAAGCCTTTGCTTATCGTGGAA
Cj1341c_AS	CGACTCTAGAGGATCCAGTCCTCGCTCATACATCACTG
Cj1341c_AS_inverse	GAACACCGCCGAGCATCATTAAGATTGAAGCGTTGG
Cj1341c_SE	CGGTACCCGGGGATCCGAGGATGTCTGGCATAAAGCAG
Cj1341c_SE_inverse	GAACTAAAGGGCGCAATGCAGGCAAAGGTGAAGTT
Cj1342c_AS	CGACTCTAGAGGATCCCTAACCAAGGCGCAATCTTAGC
Cj1342c_AS_inverse	GAACACCGCCGAGCAACCACAAACACCAAGGCTTC
Cj1342c_SE	CGGTACCCGGGGATCCAGCGGGGTTGAAAATGTTGATC
Cj1342c_SE_inverse	GAACTAAAGGGCGCATGTATTGGCAATCGTCCTCA
Cj1356c_AS	CGACTCTAGAGGATCCAGAACGTGCTCAAGGCATACTC
Cj1356c_AS_inverse	GAACACCGCCGAGCACCCGTTAAAGCAAAAAGCAA
Cj1356c_SE	CGGTACCCGGGGATCCGCAGCGGCAAATGGAATTAAAG
Cj1356c_SE_inverse	GAACTAAAGGGCGCATGTGCTTAGCGTAGGTTTTGG
Cj1377c_AS	CGACTCTAGAGGATCCACGCAACCATTCGCCACTTTC
Cj1377c_AS_comp	GGGGAAGCTTTCTAGCATCGGATTAGGTTCATTGGA
Cj1377c_AS_inverse	GAACACCGCCGAGCACAAGCCCCAAATCATAAACG
Cj1377c_SE	CGGTACCCGGGGATCCTGTGTATAACCACTCCGCTTGC
Cj1377c_SE_comp	GATTTAGATGTCTAGTGGAAAAGTTCTAAATTTGTTCATC
Cj1377c_SE_inverse	GAACTAAAGGGCGCATTGTTTGGGTTGTGGAGGAT
Cj1383c_AS	CGACTCTAGAGGATCCCACCACTTCCCCAAGTTGAA
Cj1383c_AS_comp	GGGGAAGCTTTCTAGGCCGCTCCTTCTGTATTTCC
Cj1383c_AS_inverse	GAACACCGCCGAGCATGAAAGTAAAAAGCGTATGTATTAAGC
Cj1383c_SE	CGGTACCCGGGGATCCGCCGATTCCTTGTGATACAGA
Cj1383c_SE_comp	GATTTAGATGTCTAGGAAGCTAAAAAGCGTAAAATTCCA
Cj1383c_SE_inverse	GAACTAAAGGGCGCATGTTTTGATGCGCAAATTTTA
Cj1388_AS	CGACTCTAGAGGATCCGTCATAAGTGTAAGTCCAAGCC
Cj1388_AS_inverse	GAACACCGCCGAGCATCTCCTGAAGCAGGGTTGAT
Cj1388_SE	CGGTACCCGGGGATCCGCAGTCAAGGGAGAATCTAAAC
Cj1388_SE_inverse	GAACTAAAGGGCGCACGGTGCTATCTTAGAAGAAAATGG
Cj1406c_AS	CGACTCTAGAGGATCCATTGACATGAGCGAGCAAGA
Cj1406c_AS_inverse	GAACACCGCCGAGCACACTTGCACCAAAAGCAAAA
Cj1406c_SE	CGGTACCCGGGGATCCAGAGCCTTGGAGCTTGTTTAT
Cj1406c_SE_inverse	GAACTAAAGGGCGCAAAAAATCGAAAAAGAACTTGATGC
mdaB_AS	CGACTCTAGAGGATCCCTCCTCCTTTGCCTGCCAAG
mdaB_AS_inverse	GAACACCGCCGAGCACACCAAGCTGGCATTTGATA
mdaB_SE	CGGTACCCGGGGATCCTGGGTGGAATTTGAGCGTATTC
mdaB_SE_inverse	GAACTAAAGGGCGCATGGTGTATTGGCATTTGCAT
Cj1613c_AS	CGACTCTAGAGGATCCTCACTCATCATCAAGCCCAAGC
Cj1613c_AS_inverse	GAACACCGCCGAGCACGTGTTTTCATCGGCTTTTT
Cj1613c_SE	CGGTACCCGGGGATCCTCCGCATCTTGTTCAAAACCTG
Cj1613c_SE_inverse	GAACTAAAGGGCGCATTGCTCGTATGCACCTTTTG
Cj1658_AS	ATGC**AGATCT**GAGGGATTAAAGTTTCATAGTAAGGA
Cj1658_AS_inverse	ATGC**GGATCC**ACCGCTTTTGGAGTTTCTTG
Cj1658_SE	ATGC**AGATCT**GCTCAGCTTTTGCTAGGGTAGA
Cj1658_SE_inverse	ATGC**GGATCC**GTTTGGCATTGGCTTCTAGG
Cj1661_AS	CGACTCTAGAGGATCCTGCCTAAAACCACCCCGCTTG

Cj1661_AS_inverse	GAACACCGCCGAGCATAATTTTTCTTCGCCCATGC
Cj1661_SE	CGGTACCCGGGGATCCAACAACATCACCGCACTTGCTC
Cj1661_SE_inverse	GAACTAAAGGGCGCATGGCACAGCACTTTCTCAAA
Cj1663_AS	CGACTCTAGAGGATCCAACACAGCGTAACGATCATTGG
Cj1663_AS_inverse	GAACACCGCCGAGCATGCTCTTCATCCATTTGCTCT
Cj1663_SE	CGGTACCCGGGGATCCGCACTGCGTTTAGCCTTGGG
Cj1663_SE_inverse	GAACTAAAGGGCGCATCGCAGATCGAACCCTTATC
cmeA_AS	CGACTCTAGAGGATCCGGCTTGATCAGGATCTGTACCG
cmeA_AS_inverse	GAACACCGCCGAGCACATCAAAAGGAGCTTTTATTTCG
cmeA_SE	CGGTACCCGGGGATCCAATGCCGCCTCAACCTGTAAC
cmeA_SE_inverse	GAACTAAAGGGCGCATGGCTTTAAAGTGCCTCAAAT
exbB1_AS	CGACTCTAGAGGATCCCTCTGCTTGCTGCGAACTCTTG
exbB1_AS_comp	GGGGAAGCTTTCTAGGCTGCGAACTCTTGGGTAAT
exbB1_AS_inverse	GAACACCGCCGAGCATGGTATTTTGCAAACGCAAT
exbB1_SE	CGGTACCCGGGGATCCCAATTCTGCTCGCGGTCGTAC
exbB1_SE_comp	GATTTAGATGTCTAGGCTCGCGGTCGTACTTTTAT
exbB1_SE_inverse	GAACTAAAGGGCGCAGCAAATCACAGCAAGCAAAA
exbB2_AS	CGACTCTAGAGGATCCAAATTTTCATGTTCTTTGGCTTTT
exbB2_AS_comp	GGGGAAGCTTTCTAGCATGCGCAATAAAGGTTGAA
exbB2_AS_inverse	GAACACCGCCGAGCATCGCATCATCAAATTGCTCT
exbB2_SE	CGGTACCCGGGGATCCAATCACGCCCTCTTTGGTAA
exbB2_SE_comp	GATTTAGATGTCTAGGAAGAACAAAAGAGGTAAAATTGTGA
exbB2_SE_inverse	GAACTAAAGGGCGCATTGGCGGGCAATATAGATGT
exbD1_AS	CGACTCTAGAGGATCCTAGCGGAGCACTGAGCGATTC
exbD1_AS_inverse	GAACACCGCCGAGCAGCTGCGAACTCTTGGGTAAT
exbD1_SE	CGGTACCCGGGGATCCAAATGGCTTGAGTTTGCTTGCG
exbD1_SE_inverse	GAACTAAAGGGCGCAAAACCAAAGGCAATAAAGAAGAAA
exbD2_AS	CGACTCTAGAGGATCCTTGGTATTGGGTGTGGAGGT
exbD2_AS_inverse	GAACACCGCCGAGCACATGCGCAATAAAGGTTGAA
exbD2_SE	CGGTACCCGGGGATCCTGGCATTTATAGCTTTTTGGTG
exbD2_SE_inverse	GAACTAAAGGGCGCACACTTCCTTCAAGTGAAAATGC
flaG_AS	CGACTCTAGAGGATCCAGACTTGCTGGAGGATTATCGC
flaG_AS_inverse	GAACACCGCCGAGCATCGCCTTCTTGACCTTGACT
flaG_SE	CGGTACCCGGGGATCCTATATGCTTGTTTGGCGTGTGG
flaG_SE_inverse	GAACTAAAGGGCGCAACAGCAACGAGGTGTGAGTG
flgD_AS	CGACTCTAGAGGATCCCCGCTACCATAACCACCTTGAG
flgD_AS_comp	GGGGAAGCTTTCTAGAAAGCCCGTGGTATTGACAT
flgD_AS_inverse	GAACACCGCCGAGCAGGTAGCATTTGGATTGCTCA
flgD_SE	CGGTACCCGGGGATCCCAACTCCAATGCCAATGCAATG
flgD_SE_comp	GATTTAGATGTCTAGGAGCAAAATCAAAATCAAGGAAA
flgD_SE_inverse	GAACTAAAGGGCGCACTATGGAGTGGCCAGGAAGA
flgE_AS	CGACTCTAGAGGATCCATCACCGCTATTTGCCGTTGC
flgE_AS_comp	GGGGAAGCTTTCTAGGCGAAAGCGACATAGAAGAAA
flgE_AS_inverse	GAACACCGCCGAGCACCCATAAGCCCAGCAACTCT
flgE_SE	CGGTACCCGGGGATCCCAAGTGGTGGTTCAAGCGATGG
flgE_SE_comp	GATTTAGATGTCTAGTATGCAAAAATGGCTGGACA
flgE_SE_inverse	GAACTAAAGGGCGCAACCGCAAGAAGGCGATAATA
flgE2_AS	CGACTCTAGAGGATCCTGCAGGTTCAGGTACACGGATG
flgE2_AS_inverse	GAACACCGCCGAGCAGGATATGCATACCTGGATCG
flgE2_SE	CGGTACCCGGGGATCCTCTGGCGTAAGCGGACTACAAG
flgE2_SE_inverse	GAACTAAAGGGCGCAGCAGCTTATTGGGATGCTGT

flgG_AS	CGACTCTAGAGGATCCAACCATTCCTGTGGGTCCTATG
flgG_AS_inverse	GAACACCGCCGAGCAACCCATTACCTGCAATAGCC
flgG_SE	CGGTACCCGGGGATCCACTCCAATGCTATTCGCCAAGG
flgG_SE_inverse	GAACTAAAGGGCGCACAAGATGGGCTTGGAACAAT
flgG2_AS	CGACTCTAGAGGATCCCGCTGTTGGACGCACACC
flgG2_AS_inverse	GAACACCGCCGAGCAGTCATTGCCAAATCCAAAGG
flgG2_SE	CGGTACCCGGGGATCCAAGGTAAGCGGTGGTTACGATG
flgG2_SE_inverse	GAACTAAAGGGCGCACAATGCTATTCGCCAAGGTT
flgH_AS	CGACTCTAGAGGATCCTCATGTCCAACTTGCGGTAGG
flgH_AS_comp	GGGGAAGCTTTCTAGTGCTTTAGGCGAAGCTAAGG
flgH_AS_inverse	GAACACCGCCGAGCATTGCTTTGTTTTGGTGCAAG
flgH_SE	CGGTACCCGGGGATCCTCCTCCATTGCCTCTCTAAAGG
flgH_SE_comp	GATTTAGATGTCTAGTGCCATCATTTTCTCCTTGA
flgH_SE_inverse	GAACTAAAGGGCGCACAACGGAGAGAAGCAAATCA
flgI_AS	CGACTCTAGAGGATCCTTTCATCGCTGGCATCTTTTCC
flgI_AS_comp	GGGGAAGCTTTCTAGTGGCTGCACGATCAAGTAAA
flgI_AS_inverse	GAACACCGCCGAGCATACTGTGCGAGAGTCGATGG
flgI_SE	CGGTACCCGGGGATCCCAGCCAAACTTCCAGCCTTTG
flgI_SE_comp	GATTTAGATGTCTAGCAACCATAAAAACTCCCGAAA
flgI_SE_inverse	GAACTAAAGGGCGCACAGGAACAGTGATTGCTGGA
flgK_AS	CGACTCTAGAGGATCCAGGGCGGCTAATTCTTCATTTG
flgK_AS_comp	GGGGAAGCTTTCTAGGCAATTTGCAAAGGATCTGG
flgK_AS_inverse	GAACACCGCCGAGCATGCGTGTTCTGTTGGTAAGG
flgK_SE	CGGTACCCGGGGATCCAACAGGTGGAGTTCAAGTAGGC
flgK_SE_comp	GATTTAGATGTCTAGTGATGGAACAAATAATGCTTATGG
flgK_SE_inverse	GAACTAAAGGGCGCATATCATGCGCCAAATCAATG
flgL_AS	CGACTCTAGAGGATCCCCCATTCTCCCATGCCAGGTG
flgL_AS_comp	GGGGAAGCTTTCTAGAAGCGTGGAGCTGGTAAAAA
flgL_AS_inverse	GAACACCGCCGAGCAACCCCACTATTTCCAGTTGC
flgL_SE	CGGTACCCGGGGATCCTCCAAGCAGCACAGGATGAAGG
flgL_SE_comp	GATTTAGATGTCTAGTTTTTGGAACAGTTATTGCTTTTG
flgL_SE_inverse	GAACTAAAGGGCGCAGAGAGCAGACTCCGAAAGTGA
flgM_AS	CGACTCTAGAGGATCCGCCATCAAGAAAATCTTTGACA
flgM_AS_comp	GGGGAAGCTTTCTAGTTCCGCGATAAGTTTGTTTTT
flgM_AS_inverse	GAACACCGCCGAGCATTTTTGAGTATCGTTTGTTTTAGTTTC
flgM_SE	CGGTACCCGGGGATCCGCAGCCAAAGCAAGAAGTTT
flgM_SE_comp	GATTTAGATGTCTAGAAAAGATGCCAGCGATGAAA
flgM_SE_inverse	GAACTAAAGGGCGCAAGCGAGTAAAATCGCAGAGC
flgP_AS	CGACTCTAGAGGATCCATTCTACGGTAAAACGCAAGGG
flgP_AS_comp	GGGGAAGCTTTCTAGACATTTTCGCTTTGCGTCAT
flgP_AS_inverse	GAACACCGCCGAGCAAGCATCTGGAGCCAACATTT
flgP_SE	CGGTACCCGGGGATCCGCAAAGCAGAGGTGGTAAGGG
flgP_SE_comp	GATTTAGATGTCTAGAGGGCGTAATACGAGTGGTG
flgP_SE_inverse	GAACTAAAGGGCGCATGCGCAAGTAAATGGTTTGA
flgR_AS	CGACTCTAGAGGATCCGCACCAGTAGCACCCACAATG
flgR_AS_comp	GGGGAAGCTTTCTAGCACCAGTAGCACCCACAATG
flgR_AS_inverse	GAACACCGCCGAGCAGCCATCAATTCCTGGCATA
flgR_SE	CGGTACCCGGGGATCCATCCTTAGCAATGACGCAAAGC
flgR_SE_comp	GATTTAGATGTCTAGTTTACCCTTGCGTTTTACCG
flgR_SE_inverse	GAACTAAAGGGCGCAAAATGGCGAATTTCGTTCAG
flhB_AS	CGACTCTAGAGGATCCGCCTACCAAAGAGCGGAAAATG

flhB_AS_comp	GGGGAAGCTTTCTAGAAGGTGGAGCAAATTTGAAAAA
flhB_AS_inverse	GAACACCGCCGAGCAAATAGCTGCCGCATCTTGAG
flhB_SE	CGGTACCCGGGGATCCCGTTCTTGCTTGATGCTGATGG
flhB_SE_comp	GATTTAGATGTCTAGCCTAAAGGCGTGGCTGAATA
flhB_SE_inverse	GAACTAAAGGGCGCAGTGGATTTTCTCGCTCTTCG
folP_AS	CGACTCTAGAGGATCCCTTCGCCATCGCCCCTAGTG
folP_AS_inverse	GAACACCGCCGAGCACTCACTCCCAGGTCTTGAGC
folP_SE	CGGTACCCGGGGATCCAGCGGGGTGCTTTTAGCTACTC
folP_SE_inverse	GAACTAAAGGGCGCATGCGTATTTTCAAAGCGAGA
hypC_AS	CGACTCTAGAGGATCCTCCAAAACCTCAAGTGGAGAA
hypC_AS_inverse	GAACACCGCCGAGCATCGCCTTGTTTTAAAGGTTCA
hypC_SE	CGGTACCCGGGGATCCAGCTGTCATTTGGACGCTTT
hypC_SE_inverse	GAACTAAAGGGCGCACATGTAGGCGTTGCTATGGA
motAmotB_AS	CGACTCTAGAGGATCCATAAGCTCAAGAACCGATTTTTGA
motAmotB_AS_comp	GGGGAAGCTTTCTAGGCTGTAGCTGAGGTGCTAGG
motAmotB_AS_inverse	GAACACCGCCGAGCA TGCTCTTCAAGTTGTTCGGT
motAmotB_SE	CGGTACCCGGGGATCCAGGAATGGTGCTTGCAGTTACT
motAmotB_SE_comp	GATTTAGATGTCTAGTGAAGTAAAAGACGATTTGTGTGA
motAmotB_SE_inverse	GAACTAAAGGGCGCA AGCGATAATCAGGTTGCTCTT
P19_AS	ATGC**AGATCT**AAACAAGCAAGGCTAAAGCAA
P19_AS_inverse	ATGC**GGATCC**GACGCCATGTTGATGAAGAA
P19_SE	ATGC**AGATCT**GGCTACTTTCAAACGCACAA
P19_SE_inverse	ATGC**GGATCC**ACCTCTTGGCTCCATTTCAA
pseB_AS	CGACTCTAGAGGATCCGCATTAGCTGTAGCGGCAAAGG
pseB_AS_comp	GGGGAAGCTTTCTAGTCGCTTTGATCGATGTTTTG
pseB_AS_inverse	GAACACCGCCGAGCATAACACCAAAGCGTGTTTGC
pseB_SE	CGGTACCCGGGGATCCTCAAGCCCAAACTGGCATAACC
pseB_SE_comp	GATTTAGATGTCTAGAGGCAAGTATCAAGCCCAAA
pseB_SE_inverse	GAACTAAAGGGCGCAGCCCATGCTCTAGCTCCTAA
pstC_AS	CGACTCTAGAGGATCCAGAAGCTGCCATCACCACTATC
pstC_AS_comp	GGGGAAGCTTTCTAGTGCTTAAAAGCAGGAAATCCA
pstC_AS_inverse	GAACACCGCCGAGCAGCTGCCCACTGACTTGAGA
pstC_SE	CGGTACCCGGGGATCCGTGTTACGCCAAGCGAAGAAAG
pstC_SE_comp	GATTTAGATGTCTAGTGATTTGGCAAAAAGTGGTG
pstC_SE_inverse	GAACTAAAGGGCGCAGCAGATGGCACGAGTAAACA
trpF_AS	CGACTCTAGAGGATCCGCAAGCCCGTGTTGACCTG
trpF_AS_comp	GGGGAAGCTTTCTAGCATTGCACTTTCAGGCAAAA
trpF_AS_inverse	GAACACCGCCGAGCATCCATTACCGCCCTTTAAAAT
trpF_SE	CGGTACCCGGGGATCCACAGCCAATAAAGCAAGTTCGC
trpF_SE_comp	GATTTAGATGTCTAGGTAGAATGCAAGGGGCAAAA
trpF_SE_inverse	GAACTAAAGGGCGCAATTAGCCGGAGGCATAGGTT
truB_AS	CGACTCTAGAGGATCCTGTCAACAAGCTCAAGCTCATC
truB_AS_inverse	GAACACCGCCGAGCATTGGCGAAAGGATCAAGAGT
truB_SE	CGGTACCCGGGGATCCCCGTCTTGCTTATGTTGCCATC
truB_SE_inverse	GAACTAAAGGGCGCAAGCGCATTAAAGAAGGCAAA
ak233-SE	GCAAGAGTTTTGCTTATGTTAGCAG
ak234-SE	GAAATGGGCAGAGTGTATTCTCCG
ak235-SE	GTGCGGATAATGTTGTTTCTG
AR56-AS	CATCCTCTTCGTCTTGGTAGC
Cat-AS	TGCGCCCTTTAGTTCCTAAAGGGT
Cat-SE	TGCTCGGCGGTGTTCCTTTCCAAG

Kan-AS	AAGCTTTCTAGACATCTAAATC
Kan-SE	GATAAACCCAGCGAACCATT

[a-] Restriction sites in bold.

Table S3. Genes selected for isogenic deletion mutant construction. Significantly differentially expressed genes from microarray analysis upon oxidant exposure or in a Δ*perR* background were targeted for deletion in *C. jejuni*. Genes successfully deleted are presented. Genes were attempted to be deleted at least 3 times.

	Microarray analysis[a] (condition assayed[b])				
Gene name	H_2O_2	CHP	MND	Δ*perR*	Mutant constructed?
cj0416	+[c]	+	+	+	Yes
mdaB	+	+	+	+	Yes
rpmF	+	+	+	+	No
cj1485c	+	+	+	+	No
cj0672	+	+	+	+	Yes
cj0148c	+	+	+	+	Yes
cj0202c	+	+	+	+	Yes
cj0344	+	+	+	+	Yes
maf4	+	+	+	+	Yes
cj0176c	+	+	+	+	No
cj0819	+	+	+	+	Yes
cj0253	+	+	+	+	Yes
folP	+	+	+	+	Yes
cj0877c	+	+		+	No
cj0295	+	+		+	Yes
cj0524	+	+		+	Yes
ald	+		+	+	Yes
cj1534c	+		+	+	No
cj0988c	+		+	+	No
cj1241	+			+	Yes
flaG	+			+	Yes
cj0878	+			+	No
cj1388	+			+	Yes
cj1383c	-[d]	-	-	+	Yes
chuC	-	-	-	+	Yes
chuD	-	-		+	Yes
aroC	-	-		+	No
chuB	-	-		+	Yes
flhB	-		-	+	Yes
cfbpB	-			+	Yes
chaN	-			+	Yes
exbB1	-			+	Yes

cj1386	-			+	Yes
cj1710c	-			+	No
cj0135		+	+	+	No
cj0260c		+	+	+	Yes
cj1667c		+	+	+	No
cj1159c		+		+	Yes
cj0554		+		+	Yes
cj1714		+		+	No
grpE		+		+	No
chuA		-		+	Yes
cj0786			+	+	Yes
rpmJ			+	+	No
cj1558			+	+	No
tonB2			-	+	Yes
fliK			-	+	Yes
exbD2			-	+	Yes
exbD1				+	Yes
cj1658				+	Yes
chuZ				+	Yes
cj1661				+	Yes
tonB3				+	Yes
exbB2				+	Yes
p19				+	Yes
cfbpA				+	Yes
ceuC				+	No
cj1384c				+	No
trxB				+	No
cfrA				+	Yes
cj0818				+	Yes
cj0040				+	Yes
cj0045c				+	Yes
flgD				+	Yes
flgK				+	Yes
cj0587				+	Yes
cj1664				+	No
pstC				+	Yes
cj1295				+	No
flgE				+	Yes
flgL				+	Yes
pstS				+	No
cj1663				+	Yes
cj0062c				+	Yes
cj0814				+	Yes

flgC				+	No
cj0428				+	No
flgP				+	Yes
pseB				+	Yes
flgI				+	Yes
asd				+	No
flgB				+	No
flgH				+	Yes
cj1294				+	No
flgG				+	Yes
flgG2				+	Yes
flgE2				+	Yes
flgR				+	Yes
cj0073c	+			-	Yes
spoT	-			-	Yes
cj1340c		-		-	Yes
maf7				-	Yes
maf6				-	Yes
fdxA				-	No
cj1345c				-	No
trpF	+	+	+		Yes
cj0309c	+	+	+		Yes
hypC	+	+			Yes
cj1677	+		+		No
cj1623	+		+		Yes
ccoQ	+		+		Yes
cft	+				No
cj0034c	+				No
cj1484c	+				Yes
cj1406c	+				Yes
cj1255	+				Yes
cj1036c	+				Yes
ktrB	-	-	-		No
tonB1	-				Yes
cfbpC	-				Yes
cj1356c		+	+		Yes
cj0011c		+	+		No
cj0900c		+	+		Yes
cj1211		+	-		No
cj0171		+			Yes
hrcA		+			No
dnaK		+			No
cj0561c		+			Yes

flgM	+		Yes
cj0367c	+		Yes
cj1207c	+		Yes
uvrC	-	-	No
rrc		+	Yes
cj1167		+	Yes
cj0741		+	Yes
acnB		+	Yes
truB		-	Yes
cj1343c[e]			No
dprA			Yes
cj0949c			Yes
acs			Yes
ceuB			Yes
murE			No
cj1377c			Yes
cj1375			No
cj0947c			Yes
cj0494			Yes
cj0977			Yes
cj0044c			Yes
cj1242			Yes
cj1209			Yes
cj0385c			Yes
cj0020c			Yes
cj0178			Yes
cj0358			Yes
motAB			Yes

[a] Genes considered significantly differentially expressed with fold changes of at least 1.5 fold, $p<10^{-4}$.

[b] microarray conditions: H_2O_2, hydrogen peroxide; CHP, cumene hydroperoxide; MND, menadione sodium bisulphite.

[c] '+' denotes gene is significantly upregulated.

[d] '-' denotes gene is significantly downregulated.

[e] Genes included in the mutant construction which were slightly below the cutoff threshold or of functional interest.

Table S4. Sensitivity of wild-type *C. jejuni*, isogenic deletion mutants and corresponding complemented strains to H_2O_2, cumene hydroperoxide or menadione bisulphite. The diameter of the zone of inhibition is represented as the mean clear zone ± standard error for each strain (in mm) after exposure to 10 µl of 3% H_2O_2, 3% cumene hydroperoxide, or 90 mM menadione bisulphite. Each experiment was repeated in quadruplicate. Values were considered significant (*) at $P < 0.001$ using Bayesian statistical analysis.

	Oxidant[a]		
Strain	H_2O_2	CHP	MND
C. jejuni NCTC11168	19.08 ± 0.19	24.50 ± 0.22	31.63 ± 0.36
Detoxification			
ΔCj0358	16.44 ± 0.72*	23.50 ± 0.53	29.63 ± 0.80
ΔCj0358+Cj0358	17.40 ± 0.40	22.50 ± 0.52	31.93 ± 1.13
Δ*rrc*	21.08 ± 1.08	23.92 ± 1.08	34.50 ± 1.83
Δ*rrc*+*rrc*	18.55 ± 0.38	20.94 ± 0.67	32.27 ± 1.60
Cation Transport/Binding Proteins			
Δ*chaN*	17.37 ± 0.30	21.47 ± 0.17*	29.83 ± 0.61
Δ*chaN*+*chaN*	17.26 ± 0.75	22.82 ± 0.66	34.08 ± 1.97
Δ*exbB1*	16.90 ± 0.31*	23.13 ± 0.64	28.30 ± 0.58
Δ*exbB1*+*exbB1*	17.33 ± 0.60	22.63 ± 0.37	32.90 ± 1.16
Δ*cfbpB*	19.30 ± 0.25	25.33 ± 0.83	29.98 ± 1.52
Δ*exbB2*	18.20 ± 0.31	24.37 ± 0.87	28.20 ± 0.50
Δ*exbB2*+*exbB2*	18.17 ± 0.17	23.53 ± 0.44	30.90 ± 0.90
Δ*tonB3*	18.16 ± 0.76	22.33 ± 0.58	28.83 ± 0.76
Δ*tonB1*	20.53 ± 0.17	24.37 ± 0.07	31.07 ± 0.32
Δ*tonB1*+Δ*tonB2*	21.10 ± 0.44	23.77 ± 0.67	33.27 ± 1.46
Δ*tonB2*	21.10 ± 0.46*	26.47 ± 0.67	33.07 ± 1.39
Δ*tonB2*+Δ*tonB3*	19.83 ± 0.49	24.20 ± 0.56	32.10 ± 1.33
Δ*ceuB*	19.15 ± 0.17	23.95 ± 0.45	31.60 ± 1.51
Δ*ceuE*	18.77 ± 0.29	22.90 ± 0.15	31.83 ± 0.61
Δ*cfrA*	19.40 ± 0.49	23.57 ± 0.35	32.37 ± 0.30
Δ*cfbpC*	19.08 ± 0.27	25.18 ± 0.64	30.50 ± 1.53
Δ*cfbpA*	19.13 ± 0.23	24.33 ± 0.57	30.27 ± 0.54
Δ*chuA*	17.70 ± 0.67	23.27 ± 0.43	31.47 ± 1.22
Δ*chuB*	20.13 ± 0.54	23.58 ± 0.48	35.08 ± 1.20

Δ*chuC*	19.33 ± 1.42	24.83 ± 0.85	34.29 ± 2.01
Δ*chuD*	18.88 ± 0.28	22.68 ± 0.50	34.30 ± 1.50
ΔCj0045c	19.15 ± 0.09	24.25 ± 0.88	30.38 ± 1.43
ΔCj0178	20.00 ± 0.25	25.17 ± 0.69	33.40 ± 0.83
Δ*exbD1*	18.23 ± 0.29	24.07 ± 0.79	28.57 ± 0.83
Δ*exbD2*	19.61 ± 0.97	22.33 ± 0.59	31.50 ± 0.66
Δ*chuZ*	20.30 ± 0.38	23.50 ± 0.17	34.90 ± 0.20
ΔCj1658	19.10 ± 0.15	22.90 ± 0.20	32.17 ± 0.44
ΔCj1661	19.10 ± 0.17	23.64 ± 0.76	33.00 ± 1.15
ΔCj1663	17.83 ± 0.52	23.67 ± 0.74	28.67 ± 0.38
Δ*p19*	19.67 ± 0.20	23.87 ± 0.47	32.67 ± 0.88
Δ*p19* +ΔCj1658	20.43 ± 0.54	25.38 ± 0.95	33.93 ± 0.07
Energy Metabolism			
Δ*ald*	17.70 ± 0.95	22.57 ± 0.23	32.07 ± 1.55
ΔCj0073c	17.43 ± 0.23	22.33 ± 0.68	29.67 ± 1.67
ΔCj0073c+Cj0073c	17.95 ± 0.65	23.43 ± 0.77	34.33 ± 1.92
ΔCj1377c	20.70 ± 1.07	25.30 ± 0.56	35.57 ± 2.50
ΔCj1377c+Cj1377c	17.93 ± 0.32	23.17 ± 0.42	32.83 ± 0.24
Δ*hypC*	19.40 ± 0.53	26.03 ± 0.92	33.03 ± 2.40
Δ*acnB*	21.22 ± 0.45*	23.28 ± 0.24	36.28 ± 0.72
Δ*acnB+acnB*	20.27 ± 0.25	21.33 ± 0.44	33.44 ± 1.49
Δ*ccoQ*	17.83 ± 0.33	24.10 ± 0.91	28.97 ± 0.73
ΔCj1207c	18.27 ± 0.27	24.73 ± 0.79	29.43 ± 0.47
Surface Structures			
Δ*flaG*	19.17 ± 0.53	26.08 ± 1.06	35.50 ± 1.53
Δ*flgD*	20.47 ± 0.47	26.27 ± 0.36	36.50 ± 0.31*
Δ*flgD+flgD*	18.40 ± 0.31	23.60 ± 0.80	31.70 ± 0.58
Δ*flgE*	20.17 ± 1.29	24.93 ± 1.12	35.85 ± 1.35
Δ*flgE+flgE*	17.93 ± 0.37	22.53 ± 0.17	35.20 ± 1.05
Δ*flgE2*	19.64 ± 0.72	24.57 ± 0.62	33.75 ± 1.04
Δ*flgG*	20.28 ± 0.71	25.80 ± 0.40	34.50 ± 1.07
Δ*flgG2*	19.83 ± 1.07	25.63 ± 1.01	32.96 ± 1.31
Δ*flgH*	21.00 ± 0.42*	26.55 ± 0.55	37.00 ± 1.30*
Δ*flgH+flgH*	19.07 ± 0.37	23.37 ± 0.23	32.33 ± 1.19
Δ*flgI*	21.35 ± 0.43*	26.00 ± 0.30	35.38 ± 1.89
Δ*flgI+flgI*	18.95 ± 0.34	23.95 ± 0.21	31.70 ± 1.13
Δ*flgK*	21.53 ± 0.13*	26.13 ± 0.46	34.90 ± 2.04
Δ*flgK+flgK*	19.67 ± 0.17	24.73 ± 0.15	31.10 ± 0.49
Δ*flgL*	21.00 ± 0.23*	25.08 ± 0.32	34.67 ± 1.73
Δ*flgL+flgL*	18.33 ± 1.17	21.25 ± 0.12	30.67 ± 0.10
Δ*flgM*	20.27 ± 0.50	24.33 ± 0.33	36.43 ± 1.40

Δ*flgM+flgM*	19.27 ± 0.48	24.20 ± 0.67	32.93 ± 2.02
Δ*flgP*	21.63 ± 1.24*	26.25 ± 0.86	35.25 ± 2.21
Δ*flgP+flgP*	19.94 ± 0.59	25.74 ± 0.62	33.08 ± 1.26
Δ*flgR*	20.47 ± 0.75	25.63 ± 0.34	36.78 ± 1.35*
Δ*flgR+flgR*	17.90 ± 0.61	22.17 ± 0.35	32.83 ± 0.87
Δ*flhB*	19.80 ± 0.44	23.18 ± 0.32	36.53 ± 1.86*
Δ*flhB+flhB*	18.18 ± 0.76	20.86 ± 0.82	32.15 ± 0.99
Δ*fliK*	20.43 ± 0.11	26.20 ± 0.45	35.95 ± 1.86
Δ*fliK+fliK*	17.83 ± 0.17	22.73 ± 0.62	32.67 ± 0.52
Δ*maf4*	18.97 ± 0.33	22.63 ± 0.35	29.77 ± 0.91
Δ*maf6*	18.40 ± 0.15	22.87 ± 0.17	30.93 ± 0.41
Δ*maf7*	19.50 ± 0.50	23.03 ± 0.15	29.90 ± 1.17
Δ*pseB*	20.50 ± 0.32	25.92 ± 0.53	35.95 ± 1.40*
Δ*pseB+pseB*	18.95 ± 1.13	23.67 ± 1.20	26.94 ± 2.96
Drug Efflux			
ΔCj0309c	19.71 ± 0.95	25.05 ± 0.51	34.53 ± 1.41
Δ*cmeA*	20.79 ± 0.85	25.60 ± 1.26	34.58 ± 1.15
Membranes, Lipoproteins and Porins			
ΔCj0385c	19.07 ± 0.32	23.30 ± 0.21	30.67 ± 0.67
ΔCj0587	18.93 ± 0.13	23.73 ± 1.02	30.03 ± 0.15
ΔCj0818	19.40 ± 0.21	23.63 ± 0.81	29.73 ± 0.72
ΔCj1211	20.27 ± 0.26	22.93 ± 0.30	34.27 ± 1.43
ΔCj1356c	20.35 ± 0.96	23.55 ± 0.63	33.03 ± 1.44
ΔCj1484c	20.60 ± 0.50	25.67 ± 0.57	32.17 ± 1.66
Hypothetical Unknown Proteins			
ΔCj0040	18.98 ± 0.79	25.93 ± 0.64	34.55 ± 1.63
ΔCj0044c	19.17 ± 0.44	22.80 ± 0.38	32.53 ± 0.77
ΔCj0148c	18.90 ± 0.30	22.50 ± 0.40	33.57 ± 0.72
ΔCj0171	19.92 ± 1.28	25.17 ± 1.16	34.00 ± 1.75
ΔCj0202c	18.77 ± 0.37	22.27 ± 0.32	31.53 ± 0.62
ΔCj0202c+Cj0202c	17.77 ± 0.39	23.60 ± 0.55	31.10 ± 1.46
ΔCj0253	18.88 + 1.31	25.00 ± 0.85	36.04 ± 1.76
ΔCj0260c	17.42 ± 0.19	21.67 ± 0.44*	30.87 ± 0.75
ΔCj0260c+Cj0260c	17.50 ± 0.35	22.83 ± 0.57	30.85 ± 0.35
ΔCj0344	20.92 ± 0.14*	23.92 ± 0.45	33.33 ± 1.01
ΔCj0344+Cj0344	18.12 ± 0.73	21.25 ± 1.24	30.36 ± 1.33
ΔCj0416	18.30 ± 0.60	23.47 ± 1.30	32.00 ± 0.93
ΔCj0524	20.53 ± 0.48	24.03 ± 0.72	32.18 ± 0.76
ΔCj0554	18.57 ± 0.23	23.07 ± 0.75	34.07 ± 0.41
ΔCj0741	17.80 ± 0.27	22.40 ± 0.62	28.90 ± 0.59

ΔCj0786	18.27 ± 0.15	23.23 ± 0.65	31.30 ± 0.56
ΔCj0814	17.27 ± 0.54	23.90 ± 0.35	29.33 ± 1.07
ΔCj0814+Cj0814	17.64 ± 0.39	23.80 ± 0.34	33.38 ± 1.62
ΔCj0819	19.03 ± 0.10	23.50 ± 0.87	31.33 ± 0.46
ΔCj0900c	19.17 ± 0.40	22.08 ± 0.45	30.92 ± 0.99
ΔCj0977	19.77 ± 0.35	22.50 ± 0.28	34.00 ± 1.00
ΔCj1159c	19.71 ± 0.90	22.25 ± 0.40*	30.42 ± 0.72
ΔCj1209	20.60 ± 0.40	24.17 ± 0.19	36.00 ± 1.08
ΔCj1242	19.77 ± 0.33	22.73 ± 0.03	32.63 ± 0.67
ΔCj1383c	20.30 ± 0.40	23.30 ± 0.45	30.73 ± 0.66
ΔCj1383c+Cj1383c	18.44 ± 0.51	21.11 ± 0.19	31.61 ± 1.42
Δ*mdaB*	18.40 ± 1.02	26.43 ± 0.43	33.00 ± 1.62
Miscellaneous			
ΔCj0062c	21.73 ± 0.21*	26.00 ± 0.39	35.67 ± 0.33
ΔCj0062c+Cj0062c	19.44 ± 0.39	25.20 ± 0.50	33.63 ± 1.80
ΔCj0295	19.10 ± 0.31	23.80 ± 0.31	34.10 ± 0.76
ΔCj0947c	23.94 ± 1.20*	26.44 ± 0.80	43.00 ± 3.76*
ΔCj0947c+Cj0947c	20.66 ± 1.50	25.55 ± 0.50	32.33 ± 0.19
ΔCj1036c	19.17 ± 0.20	22.27 ± 0.26	30.53 ± 0.50
ΔCj1036c+Cj1036c	18.73 ± 0.48	23.83 ± 0.48	33.83 ± 1.91
ΔCj1388	21.17 ± 0.41*	25.38 ± 0.61	35.33 ± 2.20
ΔCj1623	18.33 ± 0.50	21.95 ± 0.19*	34.78 ± 2.50
ΔCj1623+Cj1623	17.44 ± 0.49	22.70 ± 0.39	32.63 ± 1.44
Δ*folP*	19.06 ± 0.36	22.11 ± 0.39	33.39 ± 0.25
Δ*pstC*	20.75 ± 0.31*	27.25 ± 1.00*	32.33 ± 1.24
Δ*pstC+pstC*	20.60 ± 0.25*	25.30 ± 0.41	36.84 ± 1.79
Δ*spoT*	20.07 ± 0.41	25.00 ± 0.17	33.83 ± 1.04
Δ*trpF*	20.04 ± 0.68	25.67 ± 0.96	34.94 ± 2.24
Δ*trpF+trpF*	18.77 ± 0.45	23.85 ± 0.34	32.72 ± 1.07
Δ*truB*	19.43 ± 0.49	25.37 ± 0.36	34.00 ± 1.31
Δ*acs*	19.12 ± 0.24	23.89 ± 0.27	33.74 ± 1.12
ΔCj0494	20.47 ± 0.15	24.83 ± 0.71	31.47 ± 1.36
ΔCj0561c	18.63 ± 0.23	22.65 ± 0.60	30.08 ± 0.43
ΔCj0672	18.60 ± 0.29	22.93 ± 0.63	30.33 ± 1.02
ΔCj0949c	20.57 ± 0.41	23.57 ± 0.19	33.90 ± 0.90
ΔCj1167	18.40 ± 0.70	23.10 ± 0.40	33.07 ± 1.18
ΔCj1241	19.30 ± 0.38	24.65 ± 0.65	30.08 ± 0.82
ΔCj1255	19.33 ± 0.84	25.35 ± 1.30	33.38 ± 1.81
ΔCj1340c	18.80 ± 0.67	22.77 ± 0.32	29.43 ± 0.81
ΔCj1406c	19.60 ± 0.20	22.43 ± 0.30	35.33 ± 1.70
Δ*dprA*	18.70 ± 0.49	24.77 ± 0.82	28.83 ± 0.49

[a-] H_2O_2, hydrogen peroxide; CHP, cumene hydroperoxide; MND, menadione

Table S5. Sensitivity of wild-type *C. jejuni*, isogenic single and double deletion mutants to H_2O_2, cumene hydroperoxide or menadione bisulphite in the presence of 20 mM sodium fumarate. The diameter of the zone of inhibition is represented as the mean clear zone ± standard error for each strain (in mm) after exposure to 10 μl of 3% H_2O_2, 3% cumene hydroperoxide, or 90 mM menadione bisulphite. Each experiment was repeated in at least quadruplicate. Values were considered significant (*) at $P < 0.05$ using one way ANOVA.

	Fumarate		Oxidant[a]	
Strain	(+/-)	H_2O_2	CHP	MND
C. jejuni NCTC11168	-	18.34 ± 0.32	23.33 ± 0.20	26.59 ± 0.41
C. jejuni NCTC11168	+	19.00 ± 0.26	23.25 ± 0.24	25.99 ± 0.61
Detoxification				
Δ*sodB*	-	22.24 ± 0.29*	nd	34.00 ± 0.70*
Δ*sodB*	+	22.41 ± 0.11*	nd	33.38 ± 0.55*
Surface Structures				
Δ*flgD*	-	20.54 ± 0.40*	24.99 ± 0.82*	33.27 ± 1.56*
Δ*flgD*	+	nd	nd	28.83 ± 1.26
Δ*flgD*Δ*ccoQ*	-	19.30 ± 0.35	23.43 ± 0.35	26.23 ± 0.63
Δ*flgH*	-	20.50 ± 0.25*	25.57 ± 0.19*	29.81 ± 0.41*
Δ*flgH*	+	20.30 ± 0.22*	25.23 ± 0.24*	26.43 ± 0.77
Δ*flgH*Δ*ccoQ*	-	21.30 ± 0.52*	25.00 ± 0.34*	24.00 ± 0.52
Δ*flgP*	-	21.50 ± 0.97*	25.21 ± 0.23*	28.88 ± 0.66
Δ*flgP*	+	21.08 ± 1.25*	25.50 ± 0.83*	26.89 ± 0.31
Δ*flhB*	-	20.63 ± 0.23*	23.30 ± 0.14	28.94 ± 0.45*
Δ*flhB*	+	20.47 ± 0.22*	23.33 ± 0.17	26.21 ± 0.66
Δ*flhB*Δ*ccoQ*	-	20.60 ± 0.27*	22.33 ± 0.20	23.36 ± 1.05*
Δ*maf7*	-	21.91 ± 0.75*	24.16 ± 0.83	28.46 ± 0.53
Δ*maf7*	+	21.25 ± 0.92*	23.92 ± 0.58	27.79 ± 0.56
Δ*motAB*	-	20.80 ± 0.12*	24.76 ± 0.09*	30.02 ± 0.92*
Δ*motAB*	+	20.54 ± 0.18*	24.50 ± 0.10	24.17 ± 0.67
Δ*motAB*Δ*ccoQ*	-	19.92 ± 0.17	24.58 ± 0.28	24.04 ± 0.22
Δ*motAB+motAB*	-	17.87 ± 0.27	22.55 ± 0.21	29.12 ± 0.30*
Energy Metabolism				

Δ*ccoQ::kan*	-	19.22 ± 0.34	23.44 ± 0.38	25.38 ± 0.59

[a-] H_2O_2, hydrogen peroxide; CHP, cumene hydroperoxide; MND, menadione

Nd, not determined

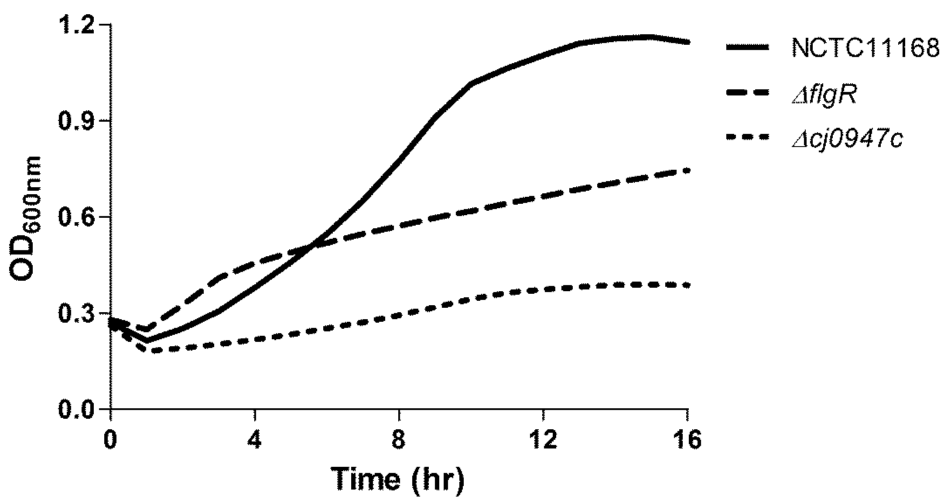

Figure S1. Growth of *C. jejuni* NCTC11168, Δ*flgR*, and Δ*cj0947c* in MH media over 16 hours. OD_{600nm} readings were taken every 15 min using a 96 well plate reader with continual shaking at 37 °C.

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