

Figure S1: Gating of flow cytometry data for single cell analyses. A-D: Forward scatter (FSC, x-axis) by side scatter (SSC, y-axis) plots identify *Salmonella*-specific events. A two-lobed SSC distribution characterizes *Salmonella* and other rod-shaped organisms (Hewitt, C.J. et al. and Rychlik I. et al.). Data were collected on a Becton Dickinson FACScan flow cytometer as indicated in the methods. A) FSCxSSC plot of total events for one sample of the GFP- WT strain (BC1662). 99.502% of events are within the *Salmonella*-specific gate. B) FSCxSSC plot of total events for one sample of the WT strain harboring a pBR322-based p*PtetA::gfp* constitutively expressed reporter (BC1967). 99.614% of events are within the *Salmonella*-specific gate. C) Overlay plot depicting GFP fluorescence of total events shown in (A) (GFP- sample, grey curve) and total events from (B) (constitutive GFP expression, black curve). 99.444% of total events from (B) are GFP+, indicating that they are *Salmonella* as opposed to particulate contaminants. D) FSCxSSC plots for representative WT 14028 p*PfliC::gfp* (BC2117) histograms from figure 2A. The *Salmonella*-specific gate was moved through the timecourse to accommodate the changing size, and therefore FSC, of the organisms. E) The *fliC*-OFF population was always defined as the range of fluorescence observed for the GFP- control (grey curve). For figures 2 and 4, the boundary between *fliC*-INT and *fliC*-HIGH was defined as the x-intercept of a line (orange) approximating the slope of the peak of highest expression for the 5 hour WT (BC2117) sample. For figure 5B, the *fliC*-HIGH gate was defined as the most intense peak of expression observed in those experiments (panel 16).

Hewitt, C.J., & Nebe-Von-Caron G. (2001) An Industrial Application of Multiparameter Flow Cytometry: Assessment of Cell Physiological State and its Application to the Study of Microbial Fermentations. *Cytometry* **44**: 179-187.

Rychlik, I., Cardova, L., Sevcik, M., & Barrow, P.A. (2000) Flow cytometry characterisation of *Salmonella typhimurium* mutants defective in proton translocating proteins and stationary-phase growth phenotype. *J Microbiol Methods* **42**: 255-263.

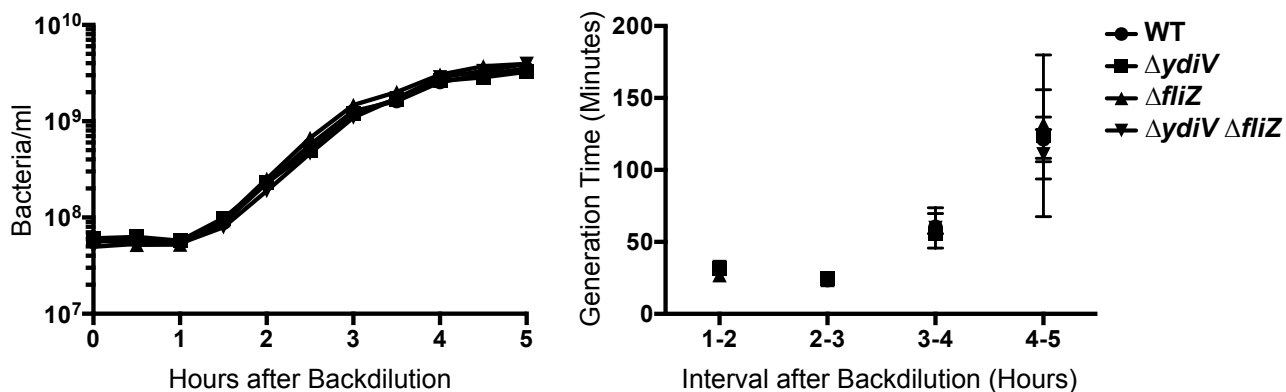


Figure S2: Growth curves for strains used in figures 2 and 4. (WT=BC2117, $\Delta ydiV$ =BC2118, $\Delta fliZ$ =BC2119 and $\Delta ydiV \Delta fliZ$ =BC2675). Strains were grown as for flow cytometry experiments, in 2ml LB cultures with 100 μ g/ml carbenicillin, shaking at 225 rpm. Samples for direct counting on a Beckman Coulter Multisizer 4 were taken every 30 minutes. Left panel: Representative set of growth curves. Strains are in exponential growth phase from 1.5-3 hours after backdilution and post-exponential phase for the remainder of the timecourse. Right panel: Generation times for the four strains between 1-2, 2-3, 3-4 and 4-5 hours after backdilution (not calculated for lag phase (0-1 hour) since no appreciable growth occurred, as seen in left panel). Mean and standard deviation for each strain from three independent experiments is plotted for each hour-long interval. There were no significant differences between the strains in generation time within intervals, as determined by 2-way ANOVA.

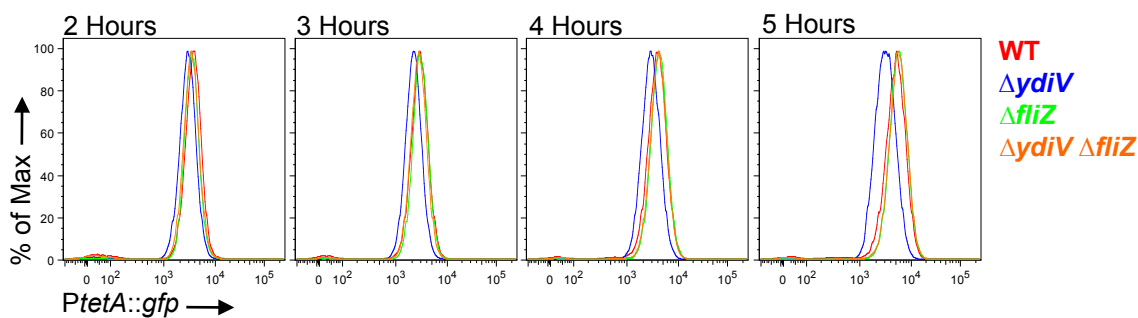


Figure S3: Monomodal expression of a constitutively expressed, plasmid-borne reporter fusion for WT and the three mutant strains across the timecourse. (WT: BC1967, $\Delta ydiV$: BC1966, $\Delta fliZ$: BC3703, $\Delta ydiV \Delta fliZ$: BC3704). Data were collected on a Becton Dickinson FACSCantoll flow cytometer using the following parameters: FSC-A 600, SSC-A 550, GFP 450, triggering on FSC-H 400 and SSC-H 200. Events shown fell within the *Salmonella*-specific gate, as shown in figure S1.

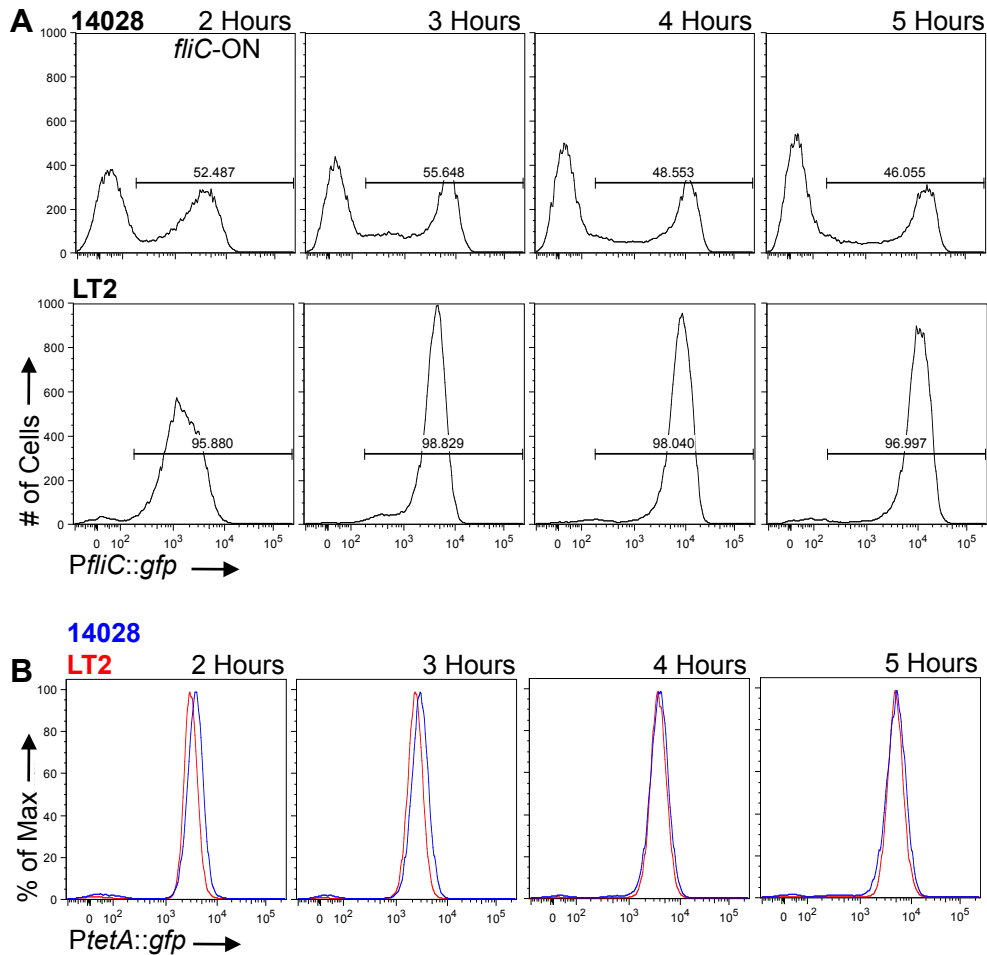


Figure S4: The census of *fliC* expression differs for *Salmonella* strains 14028 and LT2. All data were collected on a Becton Dickinson FACSCantoll flow cytometer using the following parameters: FSC-A 600, SSC-A 550, GFP 450, triggering on FSC-H 400 and SSC-H 200. Events shown fell within the *Salmonella*-specific gate, as shown in figure S1. A) *fliC* expression from 2 through 5 hours after backdilution for strains 14028 (BC2117, top panels) and LT2 (BC3616, bottom panels). 14028 maintains a *fliC*-OFF subpopulation throughout the timecourse, while most LT2 cells activate *fliC* expression by 2 hours. *fliC*-ON gate includes all fluorescence exceeding that of the GFP- control. B) The transcriptional pattern of a constitutively expressed promoter (p*PtetA::gfp*) fusion located on pBR322 is monomodal for both 14028 (BC1967) and LT2 (BC3705) throughout the timecourse.

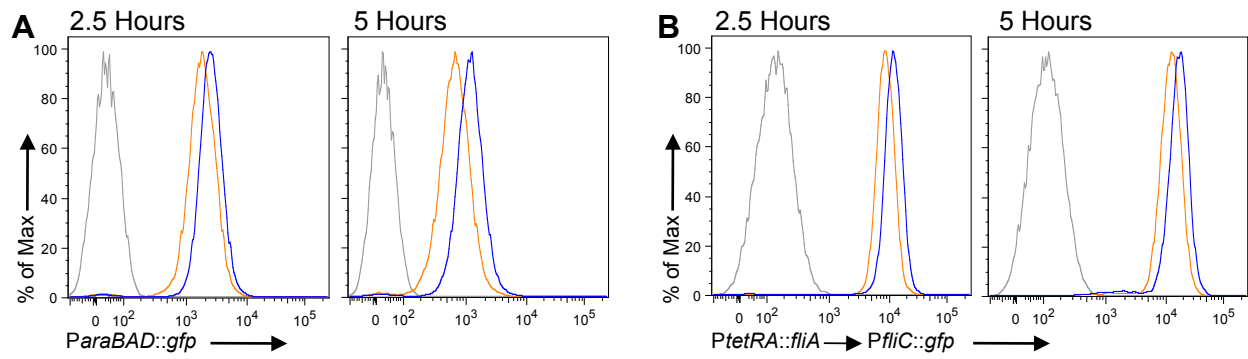


Figure S5: Unimodal expression from inducible promoters used in this study. Data were collected on a Becton Dickinson FACSCantoll flow cytometer using the following parameters: FSC-A 600, SSC-A 550, GFP 450, triggering on FSC-H 400 and SSC-H 200. Events shown fell within the *Salmonella*-specific gate, as shown in figure S1. For A and B, the left panel depicts GFP expression 2.5 hours after backdilution, and the right panel depicts GFP expression 5 hours after backdilution. A) Expression of GFP from *ParaBAD* in the pJN105 vector (BC2414). Grey trace: no L-arabinose (L-ara), orange trace: 0.01% L-ara, blue trace: 0.1% L-ara. L-ara was added 2.5 hours before sampling for both timepoints, as for the experiments reported in figure 5B. B) *tetRA*-induced FliA drives unimodal *PfiC::gfp* expression (BC2804) Grey trace: 0 μ g ml⁻¹ chlortetracycline (ct), orange trace: 1.6 μ g ml⁻¹ ct, blue trace: 3.1 μ g ml⁻¹ ct).

Table S1: Strains and plasmids used in this study, with construction intermediates

Strains		Source
BC156	<i>Salmonella</i> Typhimurium 14028	ATCC
BC1662	<i>Salmonella</i> Typhimurium 14028 <i>fijBA</i> ::FRT	Stewart et al., 2011
BC1643	<i>Salmonella</i> Typhimurium 14028 <i>fijBA</i> ::FRT <i>ydiV</i> ::FRT	Stewart et al., 2011
BC1722	<i>Salmonella</i> Typhimurium 14028 <i>fijBA</i> ::FRT <i>fliZ</i> ::FRT	This Study
BC2047	<i>Salmonella</i> Typhimurium 14028 <i>fijBA</i> ::FRT <i>fliZ</i> ::FRT <i>ydiV</i> ::FRT	This Study
BC2117	<i>Salmonella</i> Typhimurium 14028 <i>fijBA</i> ::FRT <i>pPflIC</i> :: <i>gfp</i> , <i>carbR</i>	This Study
BC2118	<i>Salmonella</i> Typhimurium 14028 <i>fijBA</i> ::FRT <i>ydiV</i> ::FRT <i>pPflIC</i> :: <i>gfp</i> , <i>carbR</i>	This Study
BC2119	<i>Salmonella</i> Typhimurium 14028 <i>fijBA</i> ::FRT <i>fliZ</i> ::FRT <i>pPflIC</i> :: <i>gfp</i> , <i>carbR</i>	This Study
BC2675	<i>Salmonella</i> Typhimurium 14028 <i>fijBA</i> ::FRT <i>fliZ</i> ::FRT <i>ydiV</i> ::FRT <i>pPflIC</i> :: <i>gfp</i> , <i>carbR</i>	This Study
BC2307	<i>Salmonella</i> Typhimurium 14028 <i>fijBA</i> ::FRT <i>ydiV</i> ::FKF::pKG136, <i>kanR</i>	This Study
BC3276	<i>Salmonella</i> Typhimurium 14028 <i>fijBA</i> ::FRT <i>fliZ</i> ::FRT <i>ydiV</i> ::FKF::pKG136 pJN105, <i>kanR</i> <i>gentR</i>	This Study
BC3277	<i>Salmonella</i> Typhimurium 14028 <i>fijBA</i> ::FRT <i>fliZ</i> ::FRT <i>ydiV</i> ::FKF::pKG136 <i>pfliz</i> , <i>kanR</i> <i>gentR</i>	This Study
BC2264	<i>Salmonella</i> Typhimurium 14028 <i>fijBA</i> ::FRT <i>FliC</i> ::3XFLAG, <i>kanR</i>	Stewart et al., 2011
BC2265	<i>Salmonella</i> Typhimurium 14028 <i>fijBA</i> ::FRT <i>ydiV</i> ::FRT <i>FliC</i> ::3XFLAG, <i>kanR</i>	Stewart et al., 2011
BC2266	<i>Salmonella</i> Typhimurium 14028 <i>fijBA</i> ::FRT <i>fliZ</i> ::FRT <i>FliC</i> ::3XFLAG, <i>kanR</i>	This Study
BC2638	<i>Salmonella</i> Typhimurium 14028 <i>fijBA</i> ::FRT <i>fliZ</i> ::FRT <i>ydiV</i> ::FRT <i>FliC</i> ::3XFLAG, <i>kanR</i>	This Study
BC2298	<i>Salmonella</i> Typhimurium LT2 <i>ydiV</i> ::FKF::pKG136, <i>kanR</i> (TH13742)	Wozniak et al., 2009
BC1697	<i>Salmonella</i> Typhimurium LT2 <i>ydiV</i> 240::Tn10dTc(del-25) (<i>ydiV</i> T-POP, TH8757), <i>tetR</i>	Wozniak et al., 2009
BC3314	<i>Salmonella</i> Typhimurium 14028 <i>fijBA</i> ::FRT <i>fliZ</i> ::FRT <i>ydiV</i> 240::Tn10dTc(del-25) <i>pfliz</i> <i>pPflIC</i> :: <i>gfp</i> , <i>tetR</i> <i>gentR</i> <i>carbR</i>	This Study
BC2414	<i>Salmonella</i> Typhimurium 14028 <i>fijBA</i> ::FRT <i>pgfp</i> , <i>gentR</i>	This Study
BC1091	<i>Salmonella</i> Typhimurium 14028 pKAS32::PflIC:: <i>gfp</i> , <i>carbR</i>	Cummings et al., 2006
BC2804	<i>Salmonella</i> Typhimurium 14028 <i>fijBA</i> ::FRT <i>flgM</i> ::FRT pKAS32::PflIC:: <i>gfp</i> <i>tetRA</i> :: <i>fliA</i> tet-driven <i>fliA</i> , <i>carbR</i> <i>tetR</i>	This Study
BC1967	<i>Salmonella</i> Typhimurium 14028 <i>fijBA</i> ::FRT <i>pTetA</i> :: <i>gfp</i> <i>carbR</i>	This Study
BC1966	<i>Salmonella</i> Typhimurium 14028 <i>fijBA</i> ::FRT <i>ydiV</i> ::FRT <i>pTetA</i> :: <i>gfp</i> <i>carbR</i>	This Study
BC3703	<i>Salmonella</i> Typhimurium 14028 <i>fijBA</i> ::FRT <i>fliZ</i> ::FRT <i>pTetA</i> :: <i>gfp</i> <i>carbR</i>	This Study
BC3704	<i>Salmonella</i> Typhimurium 14028 <i>fijBA</i> ::FRT <i>fliZ</i> ::FRT <i>ydiV</i> ::FRT <i>pTetA</i> :: <i>gfp</i> <i>carbR</i>	This Study
BC3616	<i>Salmonella</i> Typhimurium LT2 <i>fijBA</i> ::FRT <i>pPflIC</i> :: <i>gfp</i> , <i>carbR</i>	This Study
BC3705	<i>Salmonella</i> Typhimurium LT2 <i>fijBA</i> ::FRT <i>pTetA</i> :: <i>gfp</i> <i>carbR</i>	This Study
Plasmids		
<i>pPflIC</i> :: <i>gfp</i>	pSRB1, PflIC:: <i>gfp</i> construct in pBR322 vector, <i>carbR</i>	Cummings et al., 2006
pJN105	Vector, <i>gentR</i>	Newman et al., 1999
<i>pfliz</i>	<i>Salmonella</i> Typhimurium 14028 <i>fliZ</i> and native RBS cloned into pJN105, arabinose-induced expression, <i>gentR</i>	This Study
<i>pgfp</i>	<i>gfp</i> and RBS from pDW5 (Cummings et al., 2006) cloned into pJN105, arabinose-induced expression, <i>gentR</i>	This Study
<i>pTetA</i> :: <i>gfp</i>	pDW5, <i>carbR</i>	Cummings et al., 2006

Figure 2: Subpopulation Percentages			
Experiment 1			
Sample	Percent in fliC-HIGH Gate	Percent in fliC-INT Gate	Percent in fliC-OFF Gate
BC2117 2 Hr	4.91	51.2	43.9
BC2117 3 Hr	22.9	35.1	42.1
BC2117 4 Hr	31.1	18.8	50.1
BC2117 5 Hr	31.7	14.3	54
BC2119 2 Hr	5.25	45.7	49
BC2119 3 Hr	20.4	31.6	48
BC2119 4 Hr	21.1	28.4	50.5
BC2119 5 Hr	16.8	26.8	56.3
Experiment 2			
Sample	Percent in fliC-HIGH Gate	Percent in fliC-INT Gate	Percent in fliC-OFF Gate
BC2117 2 Hr	7.65	50.4	41.9
BC2117 3 Hr	25.6	33.5	40.8
BC2117 4 Hr	37	20.3	42.7
BC2117 5 Hr	38.8	16.2	45
BC2119 2 Hr	6.26	44.4	49.3
BC2119 3 Hr	26.6	28.4	45.1
BC2119 4 Hr	27.6	26.3	46.1
BC2119 5 Hr	21.5	27.8	50.8
Experiment 3 (Representative Histograms)			
Sample	Percent in fliC-HIGH Gate	Percent in fliC-INT Gate	Percent in fliC-OFF Gate
BC2117 2 Hr	7.23	42.7	50.1
BC2117 3 Hr	24.2	30.1	45.7
BC2117 4 Hr	35.3	17.4	47.2
BC2117 5 Hr	34.1	15	51
BC2119 2 Hr	6.29	45.3	48.5
BC2119 3 Hr	26.3	30.3	43.4
BC2119 4 Hr	26.2	26.6	47.2
BC2119 5 Hr	23	29.8	47.2
Averages			
Sample	Percent in fliC-HIGH Gate	Percent in fliC-INT Gate	Percent in fliC-OFF Gate
BC2117 2 Hr	6.6	48.1	45.3
BC2117 3 Hr	24.2	32.9	42.9
BC2117 4 Hr	34.5	18.8	46.7
BC2117 5 Hr	34.9	15.2	50.0
BC2119 2 Hr	5.9	45.1	48.9
BC2119 3 Hr	24.4	30.1	45.5
BC2119 4 Hr	25.0	27.1	47.9
BC2119 5 Hr	20.4	28.1	51.4

Figure 2: Mean Fluorescent Intensities			
Experiment 1			
Sample	fliC-HIGH,Mean,FL1-H	fliC-INT,Mean,FL1-H	fliC-OFF,Mean,FL1-H
BC2117 2 Hr	807	209	6.64
BC2117 3 Hr	908	257	5.05
BC2117 4 Hr	1351	208	4.78
BC2117 5 Hr	1641	191	4.66
BC2119 2 Hr	826	196	6.54
BC2119 3 Hr	926	273	4.77
BC2119 4 Hr	1253	245	4.44
BC2119 5 Hr	1438	223	4.4
Experiment 2			
Sample	fliC-HIGH,Mean,FL1-H	fliC-INT,Mean,FL1-H	fliC-OFF,Mean,FL1-H
BC2117 2 Hr	877	226	5.87
BC2117 3 Hr	936	250	5.14
BC2117 4 Hr	1367	207	4.81
BC2117 5 Hr	1528	215	4.38
BC2119 2 Hr	877	197	6.46
BC2119 3 Hr	996	272	4.93
BC2119 4 Hr	1300	253	4.55
BC2119 5 Hr	1422	242	4.35
Experiment 3 (Representative Histograms)			
Sample	fliC-HIGH,Mean,FL1-H	fliC-INT,Mean,FL1-H	fliC-OFF,Mean,FL1-H
BC2117 2 Hr	826	232	6.3
BC2117 3 Hr	971	239	4.86
BC2117 4 Hr	1421	196	4.52
BC2117 5 Hr	1596	193	4.16
BC2119 2 Hr	832	200	6.4
BC2119 3 Hr	973	270	4.49
BC2119 4 Hr	1253	255	4.13
BC2119 5 Hr	1386	241	3.88
Averages			
Sample	fliC-HIGH,Mean,FL1-H	fliC-INT,Mean,FL1-H	fliC-OFF,Mean,FL1-H
BC2117 2 Hr	836.7	222.3	6.3
BC2117 3 Hr	938.3	248.7	5.0
BC2117 4 Hr	1379.7	203.7	4.7
BC2117 5 Hr	1588.3	199.7	4.4
BC2119 2 Hr	845.0	197.7	6.5
BC2119 3 Hr	965.0	271.7	4.7
BC2119 4 Hr	1268.7	251.0	4.4
BC2119 5 Hr	1415.3	235.3	4.2

Figure 4: Mean Fluorescent Intensities			
Experiment 3			
Sample	fliC-HIGH,Mean,FL1-H	fliC-INT,Mean,FL1-H	fliC-OFF,Mean,FL1-H
BC2117 2 Hr	813	238	6.27
BC2117 3 Hr	967	255	4.89
BC2117 4 Hr	1381	204	4.48
BC2117 5 Hr	1575	200	4.12
BC2118 2 Hr	1129	244	7.77
BC2118 3 Hr	1236	207	7.42
BC2118 4 Hr	1671	170	7.28
BC2118 5 Hr	1783	145	6.73
BC2119 2 Hr	826	209	6.26
BC2119 3 Hr	1051	276	4.49
BC2119 4 Hr	1255	260	4.08
BC2119 5 Hr	1384	244	3.83
BC2675 2 Hr	1262	245	7.87
BC2675 3 Hr	1466	203	7.48
BC2675 4 Hr	2013	159	7.53
BC2675 5 Hr	2401	136	7.31
Averages			
Sample	fliC-HIGH,Mean,FL1-H	fliC-INT,Mean,FL1-H	fliC-OFF,Mean,FL1-H
BC2117 2 Hr	811.3	234.7	6.0
BC2117 3 Hr	953.3	272.3	4.8
BC2117 4 Hr	1319.3	212.7	4.4
BC2117 5 Hr	1507.7	207.0	4.1
BC2118 2 Hr	1157.7	246.7	7.4
BC2118 3 Hr	1239.0	202.0	7.4
BC2118 4 Hr	1658.0	166.3	7.2
BC2118 5 Hr	1846.7	144.7	6.7
BC2119 2 Hr	839.3	212.0	6.2
BC2119 3 Hr	1057.3	274.3	4.6
BC2119 4 Hr	1284.7	252.3	4.1
BC2119 5 Hr	1412.7	244.7	3.9
BC2675 2 Hr	1296.0	239.7	7.2
BC2675 3 Hr	1464.0	197.7	6.8
BC2675 4 Hr	2095.0	152.0	7.0
BC2675 5 Hr	2395.3	131.7	6.7