

Sample_name	Strains	single cells	dead single cells	Fraction of dead cells
C3_Data Source - 1.fcs	W3110	55167	220	0.00397205
C4_Data Source - 1.fcs	W3110	56628	182	0.00320366
C5_Data Source - 1.fcs	W3110	56192	199	0.00352893
C6_Data Source - 1.fcs	W3110	55989	161	0.0028673
C7_Data Source - 1.fcs	W3110	57565	392	0.00676363
E2_Data Source - 1.fcs	delta_bcp	66205	77	0.00116170
E3_Data Source - 1.fcs	delta_bcp	68879	93	0.00134837
E4_Data Source - 1.fcs	delta_bcp	69949	85	0.00121369
E5_Data Source - 1.fcs	delta_bcp	69820	55	0.0007871
E6_Data Source - 1.fcs	delta_bcp	68577	66	0.00096149
E7_Data Source - 1.fcs	delta_bcp	69171	73	0.00105424
G2_Data Source - 1.fcs	LN5	64104	526	0.00813863
G3_Data Source - 1.fcs	LN5	66231	618	0.00924471
G4_Data Source - 1.fcs	LN5	66798	649	0.0096223
G5_Data Source - 1.fcs	LN5	68158	502	0.00731138
G6_Data Source - 1.fcs	LN5	66369	560	0.00836707
G7 Data Source - 1.fcs	LN5	65786	613	0.00923206

S1 Fig. Selection of single cells in flow cytometry analysis. (A) one representative data of W3110. (B) LN3. (C) LN5. Number of fluorescence-signal positive dead cells in the boxed area was count and summarized in (D).