

Table S1. Fluctuation analysis of the number of colicin resistant *E.coli* (EC^{CR}) obtained on selective plates containing colicins E3+E8 *

Culture no.	Number of EC ^{CR} colonies per plate					
	Small culture	Bulk culture	Small culture	Bulk culture	Small culture	Bulk culture
1	0	82	188	68	8	48
2	26	80	11	53	3	49
3	5	87	0	37	128	45
4	87	80	34	45	5	44
5	9	93	13	59	82	45
6	1	97	5	48	5	44
7	32	68	21	35	7	51
8	4	82	3	53	17	50
9	86	69	15	48	4	54
10	4	71	0	65	4	35
Mean	25.4	80.9	29.0	51.1	26.3	46.5
Variance	1150.3	94.8	3231.1	118.1	1853.8	27.4

* Data represented here is obtained from three independent experiments

Table S2. Fluctuation analysis of the number of colicin resistant *E.coli* (EC^{CR}) obtained on selective plates containing colicins E2+E3 *

Culture no.	Number of EC ^{CR} colonies per plate					
	Small culture	Bulk culture	Small culture	Bulk culture	Small culture	Bulk culture
1	97	39	17	60	152	53
2	6	42	8	99	71	34
3	21	36	80	75	102	35
4	6	25	2	56	7	44
5	32	33	111	66	12	54
6	62	21	26	61	16	47
7	19	20	13	59	4	46
8	83	39	6	40	63	55
9	120	29	6	74	173	45
10	7	34	117	76	166	49
Mean	45.3	31.8	38.6	66.6	76.6	46.2
Variance	1774.2	60.2	2087.2	246.3	4643.6	52.6

* Data represented here is obtained from three independent experiments

Table S3. Isolation of *S. sonnei* as a sole pathogen and in polymicrobial infections from diarrheal patients admitted to the Infectious Diseases Hospital, Kolkata, over the past five years

Etiology	Frequency	Percentage
<i>S. sonnei</i>	48	57.8
<i>S. sonnei</i> + <i>Aeromonas</i> sp.	1	1.2
<i>S. sonnei</i> + <i>Campylobacter jejuni</i>	6	7.2
<i>S. sonnei</i> + <i>Campylobacter jejuni</i> + EAEC	1	1.2
<i>S. sonnei</i> +Adenovirus	2	2.4
<i>S. sonnei</i> +Astrovirus+ <i>Entamoeba histolytica</i>	1	1.2
<i>S. sonnei</i> + <i>Cryptosporidium</i> sp.	1	1.2
<i>S. sonnei</i> +EAEC	2	2.4
<i>S. sonnei</i> +EPEC	1	1.2
<i>S. sonnei</i> + <i>Giardia lamblia</i>	3	3.6
<i>S. sonnei</i> + <i>Giardia lamblia</i> + <i>Cryptosporidium</i> sp.	1	1.2
<i>S. sonnei</i> +Rotavirus	2	2.4
<i>S. sonnei</i> +Rotavirus+Adenovirus	3	3.6
<i>S. sonnei</i> +Sapovirus	1	1.2
<i>S. sonnei</i> +Sapovirus+Astrovirus	1	1.2
<i>S. sonnei</i> + <i>Vibrio cholera</i> Non O1 Non O139+ <i>Aeromonas</i> sp.	1	1.2
<i>S. sonnei</i> + <i>Vibrio cholera</i> Non O1 Non O139+ <i>Campylobacter jejuni</i>	1	1.2
<i>S. sonnei</i> + <i>Vibrio cholera</i> Non O1 Non O139	1	1.2
<i>S. sonnei</i> + <i>Vibrio cholera</i> O1	2	2.4
<i>S. sonnei</i> + <i>Vibrio cholera</i> O1+ <i>Entamoeba histolytica</i>	1	1.2
<i>S. sonnei</i> + <i>Vibrio cholera</i> O1+Rotavirus	1	1.2
<i>S. sonnei</i> + <i>Vibrio cholera</i> O1+Rotavirus+Adenovirus+ <i>Giradia lamblia</i>	1	1.2
<i>S. sonnei</i> + <i>Vibrio cholera</i> O1+Sapovirus	1	1.2
Total	83	100.0