

TABLE S2. RNA sequencing results for *V. cholerae* grown in the presence of indole.

Number	Locus number	Gene ID	Log2 fold change (0.5mM indole/0mM indole)	P-value	FDR P-value	Gene
1	VC0018	2614875	-1.25881	1.28E-10	2.33E-08	ibpA
2	VC0143	2615264	-0.69368	6.18E-05	1.65E-03
3	VC0164	2612953	-0.7571	1.27E-05	4.38E-04	vexB
4	VC0165	2612954	-0.77811	7.71E-06	2.91E-04	vexA
5	VC0166	2614082	-1.448	0	0	vexR
6	VC0188	2614711	-0.57801	6.00E-04	0.011427	prlC
7	VC0280	2615003	0.72149	1.26E-03	0.021222	cadB
8	VC0486	2615280	0.577671	8.70E-09	7.30E-07
9	VC0513	2615805	-0.75862	2.20E-05	6.98E-04
10	VC0515	2615807	-0.66973	4.28E-08	2.94E-06
11	VC0574	2615251	0.595842	3.31E-09	3.30E-07	petB
12	VC0589	2615377	-0.69519	5.54E-04	0.01088
13	VC0590	2615378	-0.70909	4.65E-04	9.33E-03
14	VC0784	2615327	0.582814	6.35E-08	4.05E-06
15	VC0818	2614485	-0.92453	4.77E-07	2.52E-05
16	VC0819	2614486	-1.37999	1.72E-13	7.96E-11	aldA-1
17	VC0820	2614487	-1.20419	4.86E-10	6.19E-08	tagA
18	VC0821	2614488	-1.21528	4.56E-10	6.19E-08
19	VC0823	2614490	-0.80036	3.53E-08	2.51E-06	mop
20	VC0824	2614491	-0.98214	4.54E-08	2.94E-06	tagD
21	VC0825	2614492	-0.74122	1.50E-06	6.99E-05	tcpI
22	VC0828	2614495	-0.65985	1.77E-06	8.09E-05	tcpA
23	VC0829	2614496	-0.74848	1.26E-06	6.06E-05	tcpB
24	VC0830	2614497	-0.73982	2.34E-05	7.27E-04	tcpQ
25	VC0831	2614498	-0.84438	5.50E-09	5.02E-07	tcpC
26	VC0832	2614499	-1.06118	4.55E-09	4.43E-07	tcpR
27	VC0833	2614500	-1.1206	1.14E-12	3.81E-10	tcpD
28	VC0834	2614501	-1.04103	1.31E-08	1.01E-06	tcpS
29	VC0835	2614502	-1.01403	7.59E-10	9.04E-08	tcpT
30	VC0836	2614503	-1.03839	1.41E-07	8.41E-06	tcpE
31	VC0837	2614504	-1.13419	2.03E-11	4.41E-09	tcpF
32	VC0838	2614505	-0.66928	5.44E-06	2.19E-04	toxT
33	VC0842	2614509	-0.57967	1.70E-04	3.96E-03
34	VC0844	2614511	-0.71133	5.22E-05	1.47E-03	acfA
35	VC0845	2614512	-0.6745	2.74E-05	8.36E-04	acfD
36	VC0846	2614513	-0.58867	2.25E-03	0.032029
37	VC0856	2614523	-0.77836	9.40E-06	3.42E-04	dnaJ
38	VC0880	2614547	0.822142	3.45E-07	1.91E-05
39	VC0884	2614113	-0.65217	1.58E-05	5.20E-04
40	VC0885	2614114	-0.7041	6.63E-05	1.75E-03
41	VC0886	2614115	-0.66364	5.28E-05	1.48E-03
42	VC0916	2614136	2.315326	2.71E-04	5.76E-03	vpsU
43	VC0917	2614137	2.058211	2.21E-05	6.98E-04	vpsA
44	VC0918	2614138	1.527912	2.65E-04	5.75E-03	vpsB
45	VC0930	2614150	1.313227	5.81E-04	0.011238	rbmC
46	VC0932	2614152	2.432418	1.53E-05	5.08E-04	rbmE
47	VC0934	2614154	2.96624	8.34E-07	4.12E-05	vpsL
48	VC0935	2614155	2.108297	1.87E-05	6.06E-04	vpsM

49	VC0936	2614156	1.666023	8.72E-04	0.015634	vpsN
50	VC0937	2614157	1.010451	1.81E-03	0.027471	vpsO
51	VC1046	2614316	0.601258	7.84E-05	2.05E-03	fadI
52	VC1093	2614363	-0.67837	1.73E-06	7.97E-05	oppC
53	VC1172	2614605	0.61079	2.68E-04	5.75E-03	trpD
54	VC1173	2614606	0.857778	2.70E-04	5.76E-03	trpG
55	VC1174	2614607	1.109521	2.20E-10	3.69E-08	trpE
56	VC1183	2614616	0.616977	7.28E-06	2.77E-04
57	VC1217	2614654	-0.61884	7.25E-04	0.013258
58	VC1316	2614770	-1.14891	0	0
59	VC1317	2614771	-1.13309	1.22E-14	6.44E-12
60	VC1318	2614772	-0.87578	5.34E-09	5.02E-07	ompV
61	VC1319	2614773	-1.06683	9.62E-12	2.22E-09	carS
62	VC1320	2614774	-1.00602	4.13E-10	5.87E-08	carR
63	VC1325	2614779	0.608037	6.32E-06	2.51E-04	mglB
64	VC1409	2614041	-1.25889	0	0	vceC
65	VC1410	2614042	-1.12355	9.99E-16	9.23E-13	vceA
66	VC1411	2614043	-0.97588	1.13E-14	6.44E-12	vceB
67	VC1442	2614074	0.67803	2.16E-11	4.44E-09	ccoN
68	VC1448	2614080	-0.60189	1.03E-05	3.64E-04	rtxB
69	VC1449	2614081	-0.77061	4.49E-08	2.94E-06
70	VC1450	2613956	-0.65722	2.50E-06	1.11E-04	rtxC
71	VC1456	2613962	-0.75861	9.44E-06	3.42E-04	ctxB
72	VC1457	2613963	-0.59964	1.24E-05	4.33E-04	ctxA
73	VC1492	2613998	0.761978	6.66E-10	8.20E-08
74	VC1510	2614016	1.116413	3.56E-10	5.25E-08
75	VC1511	2614017	1.026774	1.89E-07	1.11E-05	fdnI
76	VC1512	2614018	1.058322	2.30E-09	2.36E-07	fdhB
77	VC1513	2613892	1.0086	7.30E-08	4.57E-06
78	VC1514	2613893	0.973247	8.34E-10	9.62E-08
79	VC1515	2613894	0.718859	7.50E-07	3.79E-05
80	VC1516	2613895	0.633804	2.72E-10	4.36E-08
81	VC1517	2613896	0.708429	3.28E-12	1.01E-09
82	VC1518	2613897	0.784926	1.07E-08	8.55E-07
83	VC1523	2613902	0.739097	2.18E-13	8.04E-11
84	VC1524	2613903	0.718371	4.21E-12	1.15E-09
85	VC1576	2613955	-0.90315	2.16E-09	2.28E-07
86	VC1577	2613831	-0.88935	4.42E-08	2.94E-06	almG
87	VC1578	2613832	-1.05571	1.97E-13	8.04E-11	almF
88	VC1579	2613833	-0.94349	4.47E-08	2.94E-06	almE
89	VC1584	2613838	-0.69782	3.13E-04	6.56E-03	ankB
90	VC1741	2613746	0.640273	1.25E-04	3.09E-03
91	VC1807	2613687	-1.05129	1.77E-05	5.78E-04
92	VC1825	2613579	-0.88546	3.42E-05	1.03E-03	marA
93	VC1888	2613517	1.019879	1.94E-03	0.028956	bap1
94	VC2078	2613334	-0.59922	2.31E-04	5.08E-03	feoA
95	VC2089	2613345	0.809977	1.67E-15	1.23E-12	sdhA
96	VC2090	2613346	0.816658	1.21E-07	7.30E-06	sdhD
97	VC2091	2613347	0.748507	3.26E-08	2.36E-06	sdhC
98	VC2221	2613261	0.870114	1.56E-04	3.69E-03
99	VC2283	2613205	1.0376	4.67E-12	1.15E-09

100	VC2305	2613101	0.858466	6.95E-06	2.70E-04	ompK
101	VC2352	2613148	0.817865	5.46E-07	2.80E-05	NupC-f
102	VC2392	2613061	-0.59057	4.11E-06	1.71E-04	mutT
103	VC2485	2615142	0.587067	6.70E-09	5.75E-07	leuO
104	VC2544	2615561	0.607502	1.98E-09	2.15E-07	fbp
105	VC2674	2615502	-0.65356	1.75E-04	3.98E-03	hslU
106	VCA0025	2612526	0.871835	1.32E-10	2.33E-08
107	VCA0029	2612470	-0.69412	4.90E-04	9.78E-03	oscR
108	VCA0052	2612167	0.665344	5.01E-06	2.03E-04
109	VCA0053	2612153	0.715214	5.56E-05	1.51E-03	deoD-2
110	VCA0136	2612747	0.692371	6.86E-06	2.70E-04	glpQ
111	VCA0137	2612732	0.769744	3.25E-10	5.00E-08	glpT
112	VCA0161	2612777	0.775051	1.00E-05	3.57E-04	tnaA
113	VCA0244	2612209	0.650181	2.95E-03	0.039936	araD
114	VCA0246	2612421	0.585303	6.85E-04	0.012648	ulaA
115	VCA0276	2612035	0.704095	4.51E-12	1.15E-09
116	VCA0278	2612058	0.608111	1.41E-05	4.74E-04	glyA-2
117	VCA0280	2612010	0.709558	3.40E-06	1.48E-04
118	VCA0363	2612729	-0.76565	7.66E-08	4.71E-06
119	VCA0447	2611936	-0.69493	1.22E-04	3.04E-03
120	VCA0451	2611922	-0.60697	1.12E-03	0.01924
121	VCA0453	2611920	-0.64392	2.20E-03	0.031596
122	VCA0495	2612922	-0.63036	4.79E-10	6.19E-08
123	VCA0556	2612312	0.629457	1.13E-09	1.26E-07
124	VCA0657	2612772	0.645192	1.90E-04	4.26E-03	glpD
125	VCA0658	2612769	0.601229	1.06E-04	2.67E-03
126	VCA0732	2611948	-0.83917	2.32E-05	7.26E-04
127	VCA0744	2611914	0.887196	1.32E-08	1.01E-06	glpK
128	VCA0745	2611909	0.938236	1.34E-08	1.01E-06	glpF
129	VCA0747	2611906	0.80673	6.50E-04	0.012065	glpA
130	VCA0758	2611835	0.578872	2.30E-04	5.08E-03	artQ
131	VCA0759	2611826	0.575315	1.14E-03	0.01949	artI
132	VCA0811	2611865	0.799123	8.91E-09	7.31E-07	gbpA
133	VCA0819	2612375	-0.96205	1.58E-03	0.024753	groES-2
134	VCA0843	2612256	0.915307	6.96E-11	1.35E-08	gapA-2
135	VCA0849	2612210	0.724404	7.34E-04	0.013345
136	VCA0864	2612924	0.615492	1.86E-03	0.028078
137	VCA0867	2612873	0.918618	5.81E-09	5.11E-07	ompW
138	VCA0983	2612816	0.669776	1.89E-04	4.25E-03
139	VCA0984	2612817	0.914808	2.70E-03	0.037254	lldD
140	VCA0998	2612248	-0.72791	3.04E-08	2.24E-06
141	VCA1042	2612135	-0.58683	5.57E-09	5.02E-07
142	VCA1043	2612128	-0.62412	4.44E-05	1.28E-03	tagE-2
143	VCA1060	2612048	-0.64697	8.46E-06	3.12E-04	ribB