Supplementary Table 2. *E. coli* strains (n=108) isolated from the stool samples of indigenous individuals in Malaysia

Strain code	Strain profile[a]	REP-PCR profile	Strain cluster[b]	VTEC Detection				*mcr-1*	Antimicrobial Susceptibility Data[c]								
				eaeA	*vtx1*	*vtx2*	*vtx2f*		AMP	AMC	SAM	CRO	CXM	CIP	IPM	MEM	SXT
KG 31.2	1	REP-01	C1	No	No	No	No	No	R	I	R	S	S	S	S	S	R
KL 15.04	2	REP-02	C2	No	No	No	No	No	S	S	S	S	S	S	S	S	S
KL 10.1	3	REP-03	C3														
KL 10.12		REP-03		No	No	No	No	No	S	S	S	S	S	S	S	S	S
KG 29.2	4	REP-04	C4	No	No	No	No	No	S	S	S	S	S	S	S	S	S
KG 40.3	5	REP-05	C5	No	No	No	No	No	S	S	S	S	S	S	S	S	S
KL 16.06	6	REP-06	C6	No	No	No	No	No	R	S	S	S	S	S	S	S	R
KL 10.11	7	REP-07	C7	No	No	No	No	No	S	S	S	S	S	S	S	S	S
KL 10.09	8	REP-08	C8	No	No	No	No	No	S	S	S	S	S	S	S	S	S
KG 3.5	9	REP-09	C9	No	No	No	No	No	S	S	S	S	S	S	S	S	R
KG 10.3	10	REP-09		No	No	No	No	No	R	S	S	S	S	S	S	S	R
KG 10.4		REP-09															
KG 3.6	11	REP-10	C10	No	No	No	No	No	R	S	S	S	S	S	S	S	R
KG 27.5	12	REP-11		No	No	No	No	No	R	S	S	S	S	S	S	S	R
KG 27.6		REP-11															
KG 28.1	13	REP-11		No	No	No	No	No	R	S	S	S	S	S	S	S	R
KG 28.2		REP-11															
KG 25.3	14	REP-12		No	No	No	No	No	R	I	S	S	S	S	S	S	R
KG 25.6		REP-12															
KG 25.4	15	REP-13		No	No	No	No	No	R	S	S	S	S	I	S	S	R
KG 11.2	16	REP-13		No	No	No	No	No	R	I	R	S	S	S	S	S	R
KG 12.2	17	REP-13		No	No	No	No	No	S	S	S	S	S	S	S	S	S
KG 22.3	18	REP-14		No	No	No	No	No	R	S	R	S	S	S	S	S	R
KG 22.4		REP-14															
KG 12.6	19	REP-15	C11	No	No	No	No	No	R	I	R	S	S	S	S	S	R
KG 21.4	20	REP-16	C12	No	No	No	No	No	R	S	I	S	S	S	S	S	S
KG 8.6	21	REP-17	C13	No	No	No	No	No	R	I	S	S	S	S	S	S	R
KG 29.3	22	REP-18		No	No	No	No	No	S	S	S	S	S	S	S	S	S

KG 2.5	23	REP-19		No	No	No	No	No	S	S	S	S	S	S	S	S	S
KG 2.6		REP-19															
KG 4.5	24	REP-19	C14	No	No	No	No	No	S	S	S	S	S	S	S	S	S
KG 4.6		REP-19															
KG 15.4	25	REP-20		No	No	No	No	No	S	S	S	S	S	S	S	S	S
KG 33.1	26	REP-21	C15	No	No	No	No	No	R	S	S	S	S	S	S	S	R
KG 26.3	27	REP-22	C16	No	No	No	No	No	S	S	S	S	S	S	S	S	S
KG 26.4		REP-22															
KG 42.4	28	REP-23		No	No	No	No	No	R	S	I	S	S	S	S	S	R
KG 42.3		REP-23	C17														
KG 40.4	29	REP-23		No	No	No	No	No	R	S	I	S	S	S	S	S	R
KG 8.5	30	REP-24		No	No	No	No	No	S	S	S	S	S	S	S	S	S
KG 14.6	31	REP-25	C18	Yes	No	No	No	No	S	S	S	S	S	S	S	S	S
KG 17.3	32	REP-26	C19	No	No	No	No	No	S	S	S	S	S	S	S	S	S
SJ 3.06	33	REP-27	C20	No	No	No	No	No	S	S	S	S	S	S	S	S	S
KG 29.4	34	REP-28		No	No	No	No	No	S	S	S	S	S	S	S	S	S
KG 7.3	35	REP-29	C21	No	No	No	No	No	S	S	S	S	S	S	S	S	S
KG 7.4		REP-29															
KL 15.01		REP-30															
KL 15.02	36	REP-30	C22	No	No	No	No	No	S	S	S	S	S	S	S	S	S
KL 15.03		REP-30															
KG 5.5	37	REP-31	C23														
KG 5.6		REP-31		No	No	No	No	No	S	S	S	S	S	S	S	S	S
KG 12.5	38	REP-32	C24	No	No	No	No	No	S	S	S	S	S	S	S	S	S
KG 9.5	39	REP-33		Yes	No	No	No	No	S	S	S	S	S	S	S	S	S
KG 9.6		REP-33															
KG 32.4	40	REP-34	C25	No	No	No	No	No	S	S	S	S	S	S	S	S	S
KG 32.3		REP-34															
KG 6.3	41	REP-35		No	No	No	No	No	R	S	I	S	S	S	S	S	S
KG 6.4		REP-35															
KG 44.1	42	REP-36		No	No	No	No	No	R	S	S	S	S	S	S	S	S
KG 44.2		REP-36															
KG 43.5	43	REP-37	C26	No	No	No	No	No	S	S	S	S	S	S	S	S	S
KG 43.6		REP-37															
SJ 8.01	44	REP-38		No	No	No	No	No	R	S	S	S	S	S	S	S	R
SJ 8.03		REP-38															
KG 29.1	45	REP-39	C27	No	No	No	No	No	S	S	S	S	S	S	S	S	S

KG 21.5	46	REP-40	C28														
KG 21.6		REP-40		Yes	No	No	No	No	R	S	S	S	S	S	S	S	R
KG 33.5	47	REP-41	C29	No	No	No	No	No	R	S	S	S	S	S	S	S	S
KG 33.2		REP-41															
KG 20.7	48	REP-42	C30														
KG 20.8		REP-42		No	No	No	No	No	S	S	S	S	S	R	S	S	S
KG 9.3	49	REP-43	C31	No	No	No	No	No	S	S	S	S	S	S	S	S	S
KG 9.4		REP-43															
KG 18.3	50	REP-44	C32														
KG 18.4		REP-44		No	No	No	No	No	S	S	S	S	S	S	S	S	R
KG 36.8		REP-45		No	No	No	No	No	S	S	S	S	S	S	S	S	S
KG 36.2	51	REP-45	C33														
KG 36.1		REP-45															
KL 15.05	52	REP-46	C34	No	No	No	No	No	S	S	S	S	S	S	S	S	S
KG 11.1	53	REP-47	C35	No	No	No	No	No	S	S	S	S	S	S	S	S	S
KL 16.07	54	REP-48	C36	No	No	No	No	No	S	S	S	S	S	S	S	S	S
KG 31.1	55	REP-49		No	No	No	No	No	S	S	S	S	S	S	S	S	S
KL 9.07	56	REP-50	C37	No	No	No	No	No	S	S	S	S	S	S	S	S	S
SJ 13.05	57	REP-51	C38	No	No	No	No	No	S	S	S	S	S	S	S	S	S
KG 20.5	58	REP-52	C39	No	No	No	No	No	S	S	S	S	S	S	S	S	S
KG 33.6	59	REP-53	C40	No	No	No	No	No	S	S	S	S	S	S	S	S	S
KG 17.4	60	REP-54	C41	No	No	No	No	No	S	S	S	S	S	S	S	S	S
KG 20.6	61	REP-55	C42	No	No	No	No	No	R	S	S	S	S	S	S	S	S
KG 21.3	62	REP-55		No	No	No	No	No	R	S	S	S	S	S	S	S	R
KG 1.5	63	REP-56	C43	No	No	No	No	No	S	S	S	S	S	S	S	S	S
KG 1.6		REP-56															
KG 36.3	64	REP-57	C44	No	No	No	No	No	S	S	S	S	S	S	S	S	S
SJ 3.07		REP-58		No	No	No	No	No	S	S	S	S	S	S	S	S	S
SJ 3.08	65	REP-58	C45														
SJ 3.10		REP-58															
SJ 3.09		REP-58															
SJ 8.02	66	REP-59	C46	No	No	No	No	No	R	S	S	S	S	S	S	S	R
KL 9.05		REP-60															
KL 9.06	67	REP-60	C47	No	No	No	No	No	S	S	S	S	S	S	S	S	S
KL 9.08		REP-60															
KL 9.09		REP-60															
SJ 9.05	68	REP-61	C48	No	No	No	No	No	S	S	S	S	S	S	S	S	S

KG 14.5	69	REP-62	C49	Yes	No	No	No	No	S	S	S	S	S	S	S	S	S
SJ 9.07	70	REP-63	C50	No	No	No	No	No	S	S	S	S	S	S	S	S	S
SJ 13.07	71	REP-64	C51	No	No	No	No	No	R	S	I	S	S	S	S	S	S
KG 30.8	72	REP-65	C52	No	No	No	No	No	S	S	S	S	S	S	S	S	S
KG 30.7		REP-65															
KG 29.9	73	REP-66	C53	No	No	No	No	No	S	S	S	S	S	S	S	S	S

Footnote:

[a] Strain profile indicates *E. coli* strain that was isolated from an individual with a unique REP-PCR profile

[b] *E. coli* strains were grouped into clusters based on REP-PCR profiles at similarity cut-off F=0.85

[c] Susceptibility phenotype is indicated as S: susceptible; I: intermediate; R: resistant