

Supplemental Table S4. *Vibrio cholerae* T6SS structural genes and *C. jejuni* homologs

<i>V. cholerae</i> T6SS Gene	T6SS Gene	Function/Location	<i>C. jejuni</i> str. M129			% Present in <i>Campylobacter</i> Genomes ¹
			Protein Identity	Protein Coverage	Locus Tag	
TssA	VasJ/ImpA	Baseplate complex recruiter, Polymerization of TssD with sheath	20.2%	74.41%	CJM129_4895	22.33%
TssB	VipA/ImpB	Contractile sheath	50.6%	91.67%	CJM129_4900	24.59%
TssC	VipB/ImpC	Contractile sheath	60.3%	96.33%	CJM129_4905	24.71%
TssD	Hcp	Secreted tube, effector	63.3%	67.67%	CJM129_4875	24.84%
TssE	HsiF	Baseplate complex	24.2%	62.76%	CJM129_4910	24.38%
TssF	VasA/ImpG	Baseplate complex	30.4%	96.77%	CJM129_4915	24.25%
TssG	VasB/ImpH	Baseplate complex	26.2%	92.31%	CJM129_4920	23.77%
TssH	ClpV/VasG	AAA+ATPase, sheath recycling	40.9%	69.74%	CJM129_2355*	99.74%*
TssI	VgrG	Spike, effector	24.0%	48.12%	CJM129_5040	26.78%
TssJ	VasD/Lip/SciN	Membrane complex	0.0%	0.0%	N/A	0.00%
TssK	VasE/ImpJ	Baseplate	24.4%	98.42%	CJM129_4885	23.74%
TssL	IcmH/DotU/VasF	Membrane complex	29.7%	31.86%	CJM129_4880	23.80%
TssM	IcmF/VasK	Membrane complex	22.7%	40.39%	CJM129_4870	21.59%
TagD	PAAR	Tip of spike	0.0%	0.0%	N/A	0.06%

*Note: Hit is another ATPase (ClpB protein) not present in the CJIE3, but located elsewhere on the chromosome

¹Percentage of matches based on screening PubMLST database