

## Supplementary material

**Figure S1.** Alignment of the sequences of *Edwardsiella tarda* Ivy homologues. Dots denote gaps introduced for maximum matching. Numbers in brackets indicate overall sequence identities between CsEBI3 and the compared sequences. The consensus residues are in black; the residues that are  $\geq 75\%$  identical among the aligned sequences are in grey. The three residues that were subjected to mutagenesis analysis are boxed. The GenBank accession numbers of the aligned sequences are as follows: *Edwardsiella ictaluri*, ACR68365.1; *Serratia marcescens*, YP\_007345734.1; *Serratia proteamaculans*, YP\_001479800.1; *Aeromonas hydrophila*, YP\_857437.1; *Yersinia pestis*, WP\_002227858; *Yersinia pseudotuberculosis*, YP\_001873342.1; *Yersinia enterocolitica*, YP\_006007815.1; *Burkholderia pseudomallei*, YP\_002895840.1; *Escherichia coli*, BAL37520.1.

<i>Edwardsiella tarda</i>	.....MKTIVGALAALLLAYGAPGMADETIP.....	26
<i>Edwardsiella ictaluri</i> (88.7%)	.....MKTIVGALAALLLAYGAPGMADEAIP.....	26
<i>Serratia marcescens</i> (45.5%)	....MTGMHTLRYTLLGALLAVSAGSIAQQV.....	27
<i>Serratia proteamaculans</i> (41.6%)	....MTGQNALRGAIIFGALLSFSVGGFAQQI.....	27
<i>Aeromonas hydrophila</i> (40.8%)	.....MTRLSRALALSLLLVCAAFAFAAA.....	24
<i>Yersinia pestis</i> (37.8%)	.....MSISSGSFAQFA...AVVSSPGVTVP.E	24
<i>Yersinia pseudotuberculosis</i> (35.5%)	....MKGQKNLHCLLAGILLSISSGSFAQFA...AVVSSPGVTVP.E	39
<i>Yersinia enterocolitica</i> (34.8%)	....MKGHKLHCLVTATILSASSWVFAQFATNVALVSASSTPASSA	43
<i>Burkholderia pseudomallei</i> (30.4%)	MIRSFPPSSIRRAVLAATIAGIAGAAVADEFA.....	33
<i>Escherichia coli</i> (25.2%)	MGRISSGMMFEKAITTVAALVIATSAQAQDD.....	31

<i>Edwardsiella tarda</i>	.....PSLSEIQAQPSYQQSWKQALQGEKDLPSWVRSQGQTS	63
<i>Edwardsiella ictaluri</i>	.....PSLSEIQAQPSYQQSWKQALQGEKDLPSWVRSQGQTS	63
<i>Serratia marcescens</i>	.....VTADLVQQPGYQSSWQCMVKGQCARLPGWARKGSGTS	64
<i>Serratia proteamaculans</i>	.....VTTSDVIQQPGYQSSWQCMVKGQCARLPGWARKGSGTS	64
<i>Aeromonas hydrophila</i>	.....TYTSDLLLSADFKAFWAKATQGIKNIKPKWVRSNGTS	61
<i>Yersinia pestis</i>	PVAST.....PTMAELLQQPVYKARWQKMKVKGQKNLEAWARKGAGTS	66
<i>Yersinia pseudotuberculosis</i>	PVAST.....PTMAELLQQPVYKARWQKMKVKGQKNLEAWARKGAGTS	81
<i>Yersinia enterocolitica</i>	PVQPTLAKPLPTADLLQQPVYKNSWQKMKVKSQKNLEAWARKGAGTS	90
<i>Burkholderia pseudomallei</i>	.....PTFSEAIQAQSAHRAEWFKRMISSERRVPGWLKSDNRVS	70
<i>Escherichia coli</i>	.....LTISSIAKGETTKAAFNQMVQG.HKLEAWVMKG.GTY	66

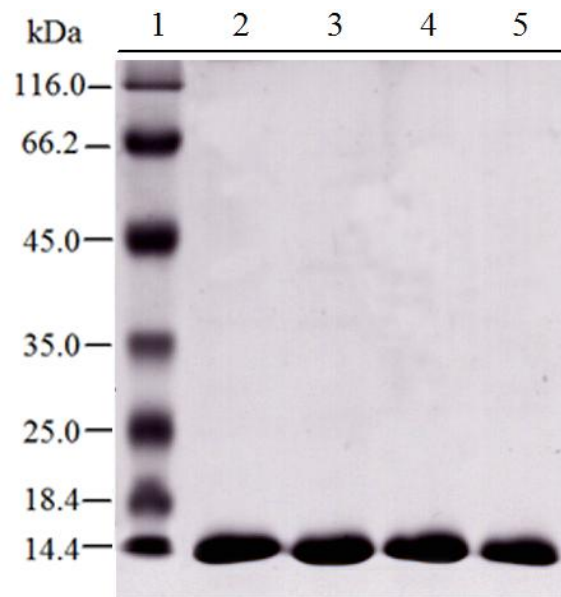
W

<i>Edwardsiella tarda</i>	AEPQTLTWQKAYQVGMVQCPHNCGNRQLFIAFSADRR...QAWALR	107
<i>Edwardsiella ictaluri</i>	TEPQTLTWQDKAYQVGMICQCPHNCGNRQVFIAFSGDRR...QAWALR	107
<i>Serratia marcescens</i>	TEAETVSWQGGQYQVGNICKPHDCGNFLIVAFKADKS...QAWGVR	108
<i>Serratia proteamaculans</i>	TEAQSLSWKGNLVLGNICKPHDCANNFMMVAFKEDKS...QAWAVR	108
<i>Aeromonas hydrophila</i>	SP...LEAVAGTFYQSGSVCKPHDCASHYMQLLEDGKAK...RVWVGL	103
<i>Yersinia pestis</i>	GPYENITWGAQQYKVGSIKPHDCSNNFMWVAFSNDKK...HVVGLR	110
<i>Yersinia pseudotuberculosis</i>	GPYENITWGAQQYKVGSIKPHDCSNNFMWVAFSNDKK...HVVGLR	125
<i>Yersinia enterocolitica</i>	APYEVINWEGQQYKIGRIKPHDCSNNFMWVAFSRDEKQVWQAWGMR	137
<i>Burkholderia pseudomallei</i>	SEYRREQVEGASYLVGWCKPHDCANQFYGVIDEEDSH...RMWGML	114
<i>Escherichia coli</i>	TEAQTVTLLGDETYQVMSACKPHDCGSQRIAMWSEKSN...QMTGLF	110

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<i>Edwardsiella tarda</i>	VTLPDSLAAMLQFAKYARFQWLGNEFASLKAALMQQVTSQED...WK	151
<i>Edwardsiella ictaluri</i>	VTLPDSLAAMLHPSKHVHFQWLGNEFASLKTLLMKQVTSQEG...WK	151
<i>Serratia marcescens</i>	VEVEDRPEAVAHPPKYYAKYQWLGKFDNSNMQALRQQFENSED...WQ	152
<i>Serratia proteamaculans</i>	VEVANKPEAIDHPKYYAKYQWLGKFNDEMKAALKQQFENNED...WK	152
<i>Aeromonas hydrophila</i>	IDLPDSLAARETPSKFARYTWLGQPDKGMCAALMKQVEADEN...WQ	147
<i>Yersinia pestis</i>	VTVADKPEALDNPSKFATYQWLGRENENIQAMTAQLEKDEN...WR	154
<i>Yersinia pseudotuberculosis</i>	VTVADKPEALDNPSKFATYQWLGRENENIQAMTAQLEKDEN...WR	169
<i>Yersinia enterocolitica</i>	VSVDDKPEALDAPSKFATYQWLGRESESVQALKKQLEQDEN...WH	181
<i>Burkholderia pseudomallei</i>	VTLPQTPGAYCAPSKYASFRWFGKPEDERMKTYLQDQLKQDEN...WK	158
<i>Escherichia coli</i>	STIDEKTSQEKLTWLNVDALSIDGKTVLFAATGSGLENHEDGFNEK	157

**Figure S2.** SDS-PAGE analysis of rIvy and its mutants. Purified rIvy, rIvyW55A, rIvyC82S, and rIvyH85D (lanes 2 to 5 respectively) were analyzed by SDS-PAGE and viewed after staining with Coomassie blue. Lane 1, protein markers.



**Table S1.** Data of one representative experimental repeat of Figure 6.

		Blood (CFU/ml)			
		TX01	TXivy	TXivy+rIvy	TXivy+rIvyC82S
<b>Fish 1</b>	plate 1	7670	680	5880	1220
	plate 2	6430	480	7120	1890
	plate 3	5630	920	7600	900
	Mean	6576.667	693.3333	6866.666667	1336.666667
<b>Fish 2</b>	plate 1	2770	390	6340	2990
	plate 2	3760	110	3210	4210
	plate 3	4170	540	6740	400
	Mean	3566.667	346.6667	5430	2533.333333
<b>Fish 3</b>	plate 1	3830	890	3210	1320
	plate 2	5110	1470	2070	890
	plate 3	3430	1350	1190	1150
	Mean	4123.333	1236.667	2156.666667	1120
<b>Fish 4</b>	plate 1	780	750	11200	230
	plate 2	1220	620	8810	100
	plate 3	1790	1050	10600	250
	Mean	1263.333	806.6667	10203.33333	193.3333333
<b>Fish 5</b>	plate 1	12300	670	6570	540
	plate 2	10800	350	8200	280
	plate 3	9810	690	8740	150
	Mean	10970	570	7836.666667	323.3333333
<b>Mean of five fish</b>		5300	730.6667	6498.666667	1101.333333

		Kidney (CFU/g)			
		TX01	TXivy	TXivy+rIvy	TXivy+rIvyC82S
<b>Fish 1</b>	plate 1	24439.66	7826.087	21864.95177	9142.857143
	plate 2	21077.59	5296.443	23183.27974	13500
	plate 3	34482.76	7391.304	25401.92926	8142.857143
	Mean	26666.67	6837.945	23483.38692	10261.90476
<b>Fish 2</b>	plate 1	15411.26	12403.43	31448.76325	4885.496183
	plate 2	20216.45	5751.073	15229.68198	8435.114504

<b>Fish 2</b>	plate 3	20519.48	8626.609	18409.89399	3129.770992
	Mean	18715.73	8927.039	21696.11307	5483.46056
<b>Fish 3</b>	plate 1	20156.25	7877.358	24782.6087	2977.777778
	plate 2	21927.08	5801.887	26576.08696	2488.888889
	plate 3	10885.42	3396.226	54891.30435	5822.222222
	Mean	17656.25	5691.824	35416.66667	3762.962963
<b>Fish 4</b>	plate 1	27966.8	7959.184	15357.14286	6326.530612
	plate 2	34066.39	10255.1	17103.1746	10728.86297
	plate 3	35020.75	2755.102	17936.50794	9154.51895
	Mean	32351.31	6989.796	16798.9418	8736.637512
<b>Fish 5</b>	plate 1	19334.81	11551.72	48015.87302	14662.37942
	plate 2	15232.82	6250	35317.46032	19967.84566
	plate 3	30177.38	13965.52	38809.52381	20257.23473
	Mean	21581.67	10589.08	40714.28571	18295.81994
<b>Mean of five fish</b>		23394.33	7807.137	27621.87884	9308.157147

### Liver (CFU/g)

		<b>TX01</b>	<b>TXivy</b>	<b>TXivy+rIvy</b>	<b>TXivy+rIvyC82S</b>
<b>Fish 1</b>	plate 1	5966.736	611.8547	8069.105691	2300.319489
	plate 2	6569.647	898.6616	9491.869919	2843.450479
	plate 3	9771.31	1051.625	8048.780488	750.798722
	Mean	7435.897	854.0472	8536.585366	1964.85623
<b>Fish 2</b>	plate 1	4005.525	804.1958	5525.81262	4107.883817
	plate 2	4917.127	472.028	3193.116635	2116.182573
	plate 3	4033.149	734.2657	4990.439771	3132.780083
	Mean	4318.6	670.1632	4569.789675	3118.948824
<b>Fish 3</b>	plate 1	5536.398	1482.301	4620.253165	2675.276753
	plate 2	4904.215	929.2035	2215.189873	1642.066421
	plate 3	5536.398	1725.664	7468.35443	2011.070111
	Mean	5325.67	1379.056	4767.932489	2109.471095
<b>Fish 4</b>	plate 1	7973.485	1423.895	3121.495327	1070.28754
	plate 2	4640.152	916.5303	1495.327103	1421.72524
	plate 3	8522.727	1423.895	2485.981308	1102.236422
	Mean	7045.455	1254.774	2367.601246	1198.083067

	plate 1	7987.013	5214.844	12248.62888	499.2199688
	plate 2	10584.42	6093.75	16819.0128	234.0093604
	plate 3	9069.264	1718.75	12248.62888	265.2106084
<b>Fish 5</b>	Mean	9213.564	4342.448	13772.09019	332.8133125
<b>Mean of five fish</b>		6667.837	1700.098	6802.799793	1744.834506

		Spleen (CFU/g)			
		TX01	TXivy	TXivy+rIvy	TXivy+rIvyC82S
<b>Fish 1</b>	plate 1	69146.34	19565.22	95357.14286	51780.82192
	plate 2	52560.98	44347.83	111071.4286	13972.60274
	plate 3	125365.9	9565.217	166785.7143	30547.94521
	Mean	82357.72	24492.75	124404.7619	32100.45662
<b>Fish 2</b>	plate 1	103939.4	47142.86	202058.8235	42678.57143
	plate 2	56666.67	15952.38	270588.2353	30535.71429
	plate 3	43030.3	30714.29	164411.7647	43214.28571
	Mean	67878.79	31269.84	212352.9412	38809.52381
<b>Fish 3</b>	plate 1	211250	50952.38	119423.0769	45211.26761
	plate 2	145312.5	42380.95	87692.30769	56478.87324
	plate 3	180312.5	26984.13	148076.9231	22112.67606
	Mean	178958.3	40105.82	118397.4359	41267.60563
<b>Fish 4</b>	plate 1	89807.69	26046.51	193442.623	25769.23077
	plate 2	100192.3	38837.21	255737.7049	36346.15385
	plate 3	99423.08	23720.93	87704.91803	27500
	Mean	96474.36	29534.88	178961.7486	29871.79487
<b>Fish 5</b>	plate 1	249253.7	31964.29	76666.66667	71111.11111
	plate 2	344776.1	46964.29	106666.6667	49444.44444
	plate 3	262238.8	12857.14	89375	49166.66667
	Mean	285422.9	30595.24	90902.77778	56574.07407
<b>Mean of five fish</b>		142218.4	31199.71	145003.9331	39724.691