

Supplementary Information

Common and distinct structural features of *Salmonella* injectisome and flagellar basal body

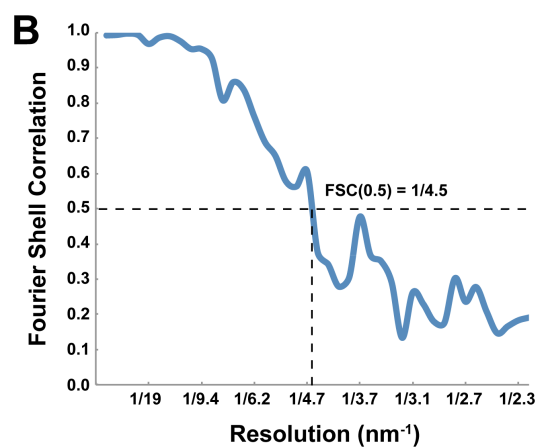
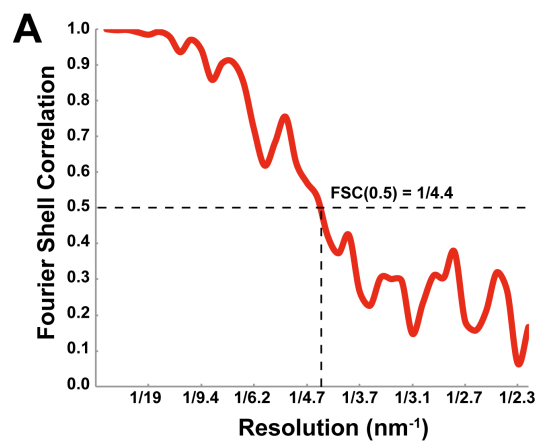
Akihiro Kawamoto^{1,2}, Yusuke V. Morimoto^{1,2}, Tomoko Miyata¹, Tohru Minamino¹, Kelly T. Hughes³, Takayuki Kato¹ & Keiichi Namba^{1,2*}

¹Graduate School of Frontier Biosciences, Osaka University, 1-3 Yamadaoka, Suita, Osaka 565-0871, Japan.

²Riken Quantitative Biology Center, 1-3 Yamadaoka, Suita, Osaka 565-0871, Japan.

³Department of Biology, University of Utah, Salt Lake City, UT, USA.

*Correspondence and requests for materials should be addressed to
K.N. (keiichi@fbs.osaka-u.ac.jp)



Supplementary Fig. 1. The resolution of 3D image reconstructions after subtomogram average. The resolution was estimated by the Fourier shell correlation at 0.5, and it was 4.4 nm for NC (A) and 4.5 nm for HBB (B).

Supplementary Table 1. Collection of T3SS homologs of the flagellum and injectisome from *Salmonella*, *Shigella*, *Yersinia* and enterohemorrhagic *E. coli* (2).

function	flagellar T3SS	nonflagellar T3SS			function		
	<i>Salmonella</i> protein (amino acid)	<i>Salmonella</i> protein (amino acid)	<i>Shigella</i> protein (amino acid)	<i>Yersinia</i> protein (amino acid)		<i>E. coli</i> * protein (amino acid)	
hook		SipD (343)	IpaD (332)	LcrV (324)	EspA (192)	needle tip	
		FlgK (553)	PrgI (80)	MxiH (83)	YscF (87)	EscF (73)	needle
			InvG (562)	MxiD (566)	YscC (607)	EscC (512)	outer membrane ring
			PrgJ (101)	MxiI (97)	YscI (115)	EscI (142)	inner rod
			InvH (147)	MxiM (142)	YscW (131)		assist in secretin insertion into OM
MS ring protein	FliF (560)	PrgK (252)	MxiJ (241)	YscJ (244)	EscJ (190)	inner membrane ring	
C ring protein	FliG (331)	PrgH (392)	MxiG (371)	YscD (418)	EscD (406)	inner membrane ring	
export apparatus	FliO (125)	-	-	-	-		
export apparatus	FliP (245)	SpaP (224)	Spa24 (216)	YscR (217)	EscR (217)	export apparatus	
export apparatus	FliQ (245)	SpaQ (86)	Spa9 (86)	YscS (88)	EscS (89)	export apparatus	
export apparatus	FliR (264)	SpaR (263)	Spa29 (256)	YscT (261)	EscT (258)	export apparatus	
export apparatus	FlhA (692)	InvA (685)	MxiA (686)	YscV (704)	EscV (675)	export apparatus	
export apparatus	FlhB (383)	SpaS (356)	Spa40 (342)	YscU (354)	EscU (345)	export apparatus	
C ring protein	FliM (334)	SpaO (303)	Spa33 (293)	YscQ (307)	EscQ (305)	sorting platform	
export apparatus	FliH (235)	OrgA (199)	MxiK (172)	YscK (209)	Orf4 (109)	sorting platform	
		OrgB (226)	MxiN (226)	YscL (223)	EscL (204)	sorting platform	
ATPase	FliI (456)	InvC (431)	Spa47 (430)	YscN (439)	EscN (446)	ATPase	
export apparatus	FliJ (147)	InvI (147)	Spa13 (112)	YscO (154)		?	
		InvE (372)	MxiC (355)	YopN (293)	SepL (351)	regulator translocation	
hook length control	FliK (405)	InvJ (336)	Spa32 (292)	YscP (453)	Orf16 (138)	regulator	
		OrgC (150)	MxiL (135)			?	

**E. coli* is EHEC O157

Supplementary Table 2. Image processing statistics of subtomogram averaging.

Strain/plasmid	Relevant properties	sample	Tilt-series showing NC or HBB	Extracted particles	Contributed to averaging structure	Pixel size (nm)	defocus (μm)	Resolution (FSC 0.5) (nm)
SJW1103/pYVM031	Minicell wild-type	Needle complex	52	142	75	1.1	-4 ~ -7	4.4
		Hook basal body	90	105	48			4.5

Supplementary Movie 1. *Salmonella* mini-cells actively swimming in the M9 medium nearly at the same speed as the cells of normal size.