## **Supplementary**

S2: Complete list of all identified Open reading frames (ORFs) for each of the 10 annotated plasmids. ND: not determined

Contig (circular	#	# of	Plasmid/Phage/Transposons related proteins	Potential accessory
element)	identified	hypothetic		proteins
	ORFs	al		
pWWcol	32	29	ND	Collagen like protein,
				Primase, VirE
RNODE_1158	26	16	Recombinase, Type IV secretory system	Thermonuclease
			Conjugative DNA transfer, TraD/Mobilization	
			protein, MobA/MobL, Putative toxin VapC6,	
			Recombinase x2, ParA, and RepB. Tn3 family	
			transposase	
pWWtox	25	10	Site-specific integrase, Toxin-antitoxin x2, AbiEii,	PEP-CTERM domain, CbiA,
			Nicking enzyme, TraD, MobA/MobL, and RepB.	DNA (cytosine-5-
				)methyltransferase,
				endonuclease, EcoRII
Temperate	29	5	Recombinase, YmfE, Lit, Integrase, Excisionase-	mcrA, Isocitrate
phage e14			like proteins, LexA repressor, YmfL, Terminase,	dehydrogenase, SAM-
			YmfR, Baseplate J-like protein, YmfQ, Tfp x2 and	dependent
			Tfap x2	methyltransferase
pWWvir	15	6	KfrA_N, ParG, ParA, VirD1, Mob, Rep, and Site-	NotI and DNA-cytosine
			specific recombinase	methyltransferase
pWWmer	13	7	Protelomerase. YafQ toxin. RelB antitoxin, Helicase	merR.
			domaine, and DNA primase	
RNODE_1311	8	3	DNaB replicative DNA helicase, Toxin-antitoxin,	ND
			Recombinase, ParA, and DNA primase.	
RNODE_1264	6	1	KfrA_N, MobC, RepB, RepA, and RepC.	ND
RNODE_961	7	5	Replication initiator protein	Integrase
RNODE_1224	3	1	IS481 family transposase	ND