











PSPTO0803	C			hypothetical protein	53.1	32								
PSPTO0819	C			conserved hypothetical protein	58.3	248	Rs	Xcc		At	Xf			
PSPTO0831	C	+	+	conserved hypothetical protein	56	642		Xcc	Xac		Xf			
PSPTO0837	C	+	+	conserved hypothetical protein	44.2	200			Xac					
PSPTO0840	C	+	+	hypothetical protein	50.2	138								
PSPTO0847	C	+	+	hypothetical protein	50	122								
PSPTO0848	C	+	+	conserved hypothetical protein	57.8	457		Xcc	Xac					
PSPTO0849	C	+	+	conserved hypothetical protein	59.5	312		Xcc	Xac					
PSPTO0850	C	+	+	conserved hypothetical protein	58.5	138			Xac					
PSPTO0851	C	+	+	conserved hypothetical protein	51.6	162						Yp	Ec	
PSPTO0854	C	+	+	hypothetical protein	51.1	73								
PSPTO0856	C	+	+	conserved hypothetical protein	52	272								
PSPTO0858	C	+	+	hypothetical protein	53.2	99								
PSPTO0859	C	+	+	conserved hypothetical protein	56.1	889			Xac					
PSPTO0860	C	+	+	conserved hypothetical protein	58.1	151								
PSPTO0861	C	+	+	conserved hypothetical protein	58.2	498			Xac					
PSPTO0862	C	+	+	conserved hypothetical protein	58.5	307			Xac					
PSPTO0863	C	+	+	conserved hypothetical protein	59.6	222								
PSPTO0866	C	+	+	hypothetical protein	59.8	121			Xac					
PSPTO0867	C	+	+	conserved hypothetical protein	59	122			Xac					
PSPTO0868	C	+	+	hypothetical protein	61.2	49								
PSPTO0870	C	+	+	hypothetical protein	49	66								
PSPTO0874	C	+	+	nikkomycin biosynthesis domain protein	49.2	417								
PSPTO0878	C	+	+	conserved domain protein	59.4	105								

PSPTO0880	C	+	+	conserved hypothetical protein	56.9	249			Xac					
PSPTO0881	C	+	+	conserved domain protein	57.9	293			Xac					
PSPTO0882	C	+	+	hypothetical protein	50.8	177								
PSPTO0891	C		+	hypothetical protein	57	76								
PSPTO0895	C	+	+	hypothetical protein	50.4	168								
PSPTO0899	C	+	+	conserved domain protein	55.3	155	Rs							
PSPTO0900	C	+	+	hypothetical protein	63.4	93								
PSPTO0904	C	+	+	mutT/nudix family protein	51.9	504								
PSPTO0907	C	+	+	hypothetical protein	53.1	403								
PSPTO0914	C	+	+	STAS domain protein	63.9	96		Xcc	Xac					
PSPTO0918	C	+	+	conserved hypothetical protein	60	35								
PSPTO0921	C	+	+	hypothetical protein	44.2	43								
PSPTO0930	C	+		conserved hypothetical protein	55.8	230								
PSPTO0931	C	+		hypothetical protein	51.9	135								
PSPTO0932	C	+		conserved hypothetical protein	56.2	181								
PSPTO0933	C	+		conserved hypothetical protein	60.1	332						Yp	Ec	
PSPTO0935	C	+		conserved hypothetical protein	62.7	377						Yp	Ec	
PSPTO0936	C	+		conserved hypothetical protein	59.5	252						Yp	Ec	
PSPTO0937	C	+		conserved hypothetical protein	62.5	364						Yp	Ec	
PSPTO0938	C	+		conserved hypothetical protein	60.1	316						Yp	Ec	
PSPTO0966	C			hypothetical protein	53.4	58								
PSPTO0974	C			hypothetical protein	54.3	35								
PSPTO0987	C			conserved domain protein	50.4	76								
PSPTO1003	C			acetyltransferase, GNAT family	58.2	145				At				St

PSPTO1006	C			hypothetical protein	57.6	33											
PSPTO1016	C			hypothetical protein	53.2	534											
PSPTO1036	C	+		hypothetical protein	47.7	153											
PSPTO1038	C	+		conserved hypothetical protein	63.5	149								Yp			
PSPTO1044	C	+		hypothetical protein	55.9	59											
PSPTO1045	C	+		PIN domain protein	54.2	134				At	Xf						St
PSPTO1046	C	+		conserved hypothetical protein	52.2	76											St
PSPTO1055	C	+		hypothetical protein	53.7	72											
PSPTO1057	C	+		conserved hypothetical protein TIGR00305	56.4	136							Xf				
PSPTO1058	C	+		conserved hypothetical protein	53.2	79							Xf				
PSPTO1060	C	+		hypothetical protein	52.5	47											
PSPTO1062	C	+		hypothetical protein	54	87											
PSPTO1082	C			hypothetical protein	54.2	72											
PSPTO1088	C	+		conserved hypothetical protein	58.6	350	Rs		Xac						Ec	St	
PSPTO1090	C	+		hypothetical protein	38.1	325											
PSPTO1091	C	+		conserved hypothetical protein	39.1	138											
PSPTO1093	C	+		relaxase/mobilization nuclease domain protein	52.8	805											
PSPTO1094	C	+		conserved hypothetical protein	53.1	125											
PSPTO1113	C			hypothetical protein	54.1	194											
PSPTO1114	C			hypothetical protein	55.9	74		Xcc	Xac								
PSPTO1124	C			hypothetical protein	56.7	40											
PSPTO1125	C			hypothetical protein	58.7	71											
PSPTO1137	C			hypothetical protein	62	36											
PSPTO1142	C			hypothetical protein	60.5	86		Xcc	Xac								















PSPTO2615	C	+		GAF domain protein	54.3	165		Xcc	Xac	At				
PSPTO2619	C	+		hypothetical protein	60.6	218								
PSPTO2631	C			hypothetical protein	59.6	197								
PSPTO2647	C			hypothetical protein	57	246								
PSPTO2648	C			hypothetical protein	57	155								
PSPTO2669	C	+	+	conserved hypothetical protein	58.3	96	Rs			At				
PSPTO2675	C	+	+	conserved hypothetical protein	58.2	196								
PSPTO2679	C	+	+	hypothetical protein	60.8	40								
PSPTO2682	C	+	+	hypothetical protein	59.5	199			Xac	At				
PSPTO2695	C	+	+	conserved hypothetical protein	60.9	353		Xcc	Xac					
PSPTO2698	C	+	+	hypothetical protein	59.6	104								
PSPTO2700	C	+	+	hypothetical protein	56.9	184								
PSPTO2727	C			hypothetical protein	49.5	31								
PSPTO2768	C	+		hypothetical protein	50.6	58								
PSPTO2770	C	+		hypothetical protein	60	35								
PSPTO2772	C	+		hypothetical protein	55.1	141								
PSPTO2774	C	+		conserved hypothetical protein	57.5	131						Yp		
PSPTO2781	C	+		conserved hypothetical protein	62.8	411	Rs			At				
PSPTO2789	C			hypothetical protein	43.1	48								
PSPTO2796	C			conserved hypothetical protein	60.2	283				At				
PSPTO2797	C			hypothetical protein	63.1	56								
PSPTO2809	C	+		conserved hypothetical protein	64.6	486	Rs	Xcc	Xac	At			Ec	St
PSPTO2815	C	+		conserved hypothetical protein	61.3	173	Rs			At		Yp		
PSPTO2817	C	+		conserved domain protein	62.8	61		Xcc	Xac	At		Yp		

PSPTO2819	C	+		conserved hypothetical protein	58.9	262												
PSPTO2820	C	+		hypothetical protein	63.8	371												
PSPTO2821	C	+		hypothetical protein	65.6	302												
PSPTO2823	C	+		hypothetical protein	59.3	249												
PSPTO2824	C	+		auxin-responsive GH3-related protein	61.8	487												
PSPTO2826	C	+		conserved domain protein	58.9	99				At			Yp	Ec				
PSPTO2827	C	+		conserved domain protein	60.8	80				At			Yp		St			
PSPTO2835	C			conserved domain protein	58.7	50				At								
PSPTO2858	C	+	+	hypothetical protein	51.7	118												
PSPTO2859	C	+	+	conserved hypothetical protein	56.2	866							Yp					
PSPTO2860	C	+	+	helicase domain protein	57.5	459								Ec				
PSPTO2866	C	+	+	cupin family protein	57.1	133		Xcc	Xac	At			Yp					
PSPTO2867	C	+	+	TspO/MBR family protein	56.8	146		Xcc	Xac	At								
PSPTO2868	C	+	+	conserved hypothetical protein	59.4	392	Rs					Xf						
PSPTO2869	C	+	+	hypothetical protein	43.3	40												
PSPTO2871	C	+	+	conserved hypothetical protein	59.7	1044							Yp		St			
PSPTO2873	C	+	+	conserved hypothetical protein	61.2	345												
PSPTO2880	C	+	+	conserved hypothetical protein	61.6	423												
PSPTO2881	C	+	+	hypothetical protein	61.5	32												
PSPTO2888	C	+	+	conserved domain protein	45.1	48												
PSPTO2889	C	+	+	hypothetical protein	55	37												
PSPTO2892	C	+	+	hypothetical protein	52.1	32												
PSPTO2894	C	+	+	lectin repeat domain protein	50	801												
PSPTO2895	C	+	+	hypothetical protein	57.1	339				At								

PSPTO2897	C	+	+	conserved hypothetical protein	58.9	380		Xcc	Xac						
PSPTO2899	C	+	+	conserved hypothetical protein	59.1	394		Xcc	Xac						
PSPTO2908	C	+		hypothetical protein	56.8	230									
PSPTO2918	C	+		conserved hypothetical protein	61.2	135	Rs	Xcc	Xac	At					
PSPTO2924	C	+		hypothetical protein	54.7	53									
PSPTO2929	C	+		hypothetical protein	53.5	320									
PSPTO2931	C	+		conserved hypothetical protein	59.3	394									
PSPTO2939	C	+		hypothetical protein	54.5	170									
PSPTO2942	C	+		hypothetical protein	57.8	34									
PSPTO2959	C	+		conserved hypothetical protein	64.2	551									
PSPTO2963	C	+		oxidoreductase, FAD-binding	65	379	Rs	Xcc	Xac	At					
PSPTO2965	C	+		conserved hypothetical protein	57.9	130	Rs								
PSPTO2977	C	+		conserved hypothetical protein	51.7	211									
PSPTO2979	C	+		conserved hypothetical protein	52.3	128									
PSPTO2982	C	+		conserved hypothetical protein	62.3	576		Xcc		At					
PSPTO2998	C			conserved hypothetical protein	58.7	142	Rs	Xcc	Xac	At	Xf				
PSPTO2999	C			hypothetical protein	54.4	250									
PSPTO3016	C			conserved hypothetical protein	48.4	117	Rs					Yp	Ec	St	
PSPTO3018	C			conserved hypothetical protein	58.6	307	Rs			At		Yp		St	
PSPTO3026	C			conserved hypothetical protein	55.1	81		Xcc							
PSPTO3055	C			hypothetical protein	54.4	49									
PSPTO3067	C		+	hypothetical protein	59.1	230									
PSPTO3078	C		+	conserved hypothetical protein	52.9	128									
PSPTO3092	C		+	conserved hypothetical protein	60.1	278		Xcc	Xac						

PSPTO3104	C			hypothetical protein	59	48											
PSPTO3117	C			hypothetical protein	57.2	60											
PSPTO3120	C			Cof-like hydrolase family protein	60.4	272							Yp	Ec	St		
PSPTO3151	C			conserved hypothetical protein	64.5	61				At							
PSPTO3166	C	+		conserved domain protein	58.5	320											
PSPTO3167	C	+		hypothetical protein	60.7	101											
PSPTO3168	C	+		hypothetical protein	39.1	46											
PSPTO3169	C	+		hypothetical protein	62.1	87											
PSPTO3177	C			conserved hypothetical protein	57.6	99											
PSPTO3187	C			hypothetical protein	53.9	86											
PSPTO3189	C			conserved hypothetical protein	62.1	317											
PSPTO3200	C	+		hypothetical protein	60.5	290											
PSPTO3201	C	+		hypothetical protein	60.6	259											
PSPTO3203	C	+		conserved hypothetical protein	51.7	154											
PSPTO3211	C			conserved hypothetical protein	40.5	171							Yp				
PSPTO3217	C	+		hypothetical protein	50.3	55											
PSPTO3218	C	+		hypothetical protein	52.6	398											
PSPTO3219	C	+		hypothetical protein	57.7	41											
PSPTO3222	C	+		hypothetical protein	53.1	71											
PSPTO3228	C	+		hypothetical protein	53	56											
PSPTO3236	C			hypothetical protein	51.9	131											
PSPTO3241	C			phytase domain protein	62.9	640											
PSPTO3246	C			hypothetical protein	59.3	232									Ec	St	
PSPTO3253	C			conserved hypothetical protein	55	240							Yp	Ec	St		

PSPTO3261	C			hypothetical protein	58.3	84											
PSPTO3270	C	+		conserved hypothetical protein	55.7	94											
PSPTO3271	C	+		hypothetical protein	60.3	110											
PSPTO3276	C	+		hypothetical protein	44.4	39											
PSPTO3289	C	+		hypothetical protein	54.6	72											
PSPTO3292	C	+		hypothetical protein	53.1	404	Rs										
PSPTO3293	C	+		hypothetical protein	57.2	416											
PSPTO3303	C			hypothetical protein	55.4	62											
PSPTO3305	C			hypothetical protein	54.2	280											
PSPTO3324	C			hypothetical protein	50.5	35											
PSPTO3337	C			hypothetical protein	53.2	195											
PSPTO3384	C	+		hypothetical protein	52.9	68											
PSPTO3386	C	+		hypothetical protein	50.4	784				At							
PSPTO3387	C	+		conserved hypothetical protein	41.6	109											Ec
PSPTO3402	C			hypothetical protein	62.7	92											
PSPTO3414	C	+		hypothetical protein	61.1	48											
PSPTO3415	C	+		conserved domain protein	57.6	128											
PSPTO3420	C	+		hypothetical protein	56.8	199	Rs										
PSPTO3422	C	+		hypothetical protein	55	191											
PSPTO3424	C	+		hypothetical protein	54.6	124											
PSPTO3426	C	+		hypothetical protein	53.3	140											
PSPTO3428	C	+		conserved domain protein	55.7	152	Rs										
PSPTO3429	C	+		DNA-binding protein, putative	53.8	80	Rs		Xac		Xf	Yp	Ec				
PSPTO3468	C			conserved hypothetical protein	65.2	467		Xcc	Xac	At							



PSPTO3623	C			hypothetical protein	60.3	121											
PSPTO3640	C			conserved hypothetical protein	63.1	378	Rs			At							
PSPTO3650	C	+		hypothetical protein	42.3	426											
PSPTO3654	C	+		acetyltransferase, GNAT family	59.3	162											
PSPTO3655	C	+		conserved hypothetical protein	60.7	388	Rs			At							
PSPTO3673	C			hypothetical protein	53.3	40											
PSPTO3682	C	+		conserved hypothetical protein	63.4	375	Rs	Xcc	Xac								
PSPTO3683	C	+		hypothetical protein	54.6	61											
PSPTO3684	C	+		hypothetical protein	54.6	133											
PSPTO3690	C			conserved hypothetical protein	48.7	286			Xac								
PSPTO3693	C			hypothetical protein	60.4	95											
PSPTO3697	C			hypothetical protein	61.4	38											
PSPTO3710	C			acetyltransferase, GNAT family	57.4	184											
PSPTO3728	C			hypothetical protein	56.1	38											
PSPTO3732	C			hypothetical protein	52.3	79											
PSPTO3736	C			conserved hypothetical protein	57	90	Rs			At							
PSPTO3754	C			hypothetical protein	50.7	297											
PSPTO3767	C			hypothetical protein	52.7	86											
PSPTO3775	C	+		hypothetical protein	59.9	94											
PSPTO3776	C	+		hypothetical protein	47.5	148											
PSPTO3780	C	+		glutathionylspermidine synthase family protein	58.2	385		Xcc	Xac	At		Yp	Ec	St			
PSPTO3781	C	+		conserved hypothetical protein	61.4	240		Xcc	Xac				Ec				
PSPTO3782	C	+		conserved hypothetical protein	59.3	135	Rs	Xcc	Xac	At			Ec	St			
PSPTO3783	C	+		conserved hypothetical protein	59.5	218		Xcc	Xac				Ec	St			

























The inferred proteomes of eight pathogenic bacteria were searched with the 811 genes unique to DC3000 by using a BLASTP cutoff criterion of  $E < 10^{-5}$  which will identify putative homologs as well as paralogous gene family members. Rs = *Ralstonia solanacearum*, Xac = *Xanthomonas axonopodis* pv. *citri*, Xcc = *Xanthomonas campestris* pv. *campestris*, At = *Agrobacterium tumefaciens* (U. Wash seq.), Xf = *Xylella fastidiosa*, Yp = *Yersinia pestis* CO92, St = *Salmonella typhimurium* LT2, and Ec = *Escherichia coli* O157:H7 EDL933.

<sup>a</sup> Individual unique genes that cluster with one another at a density of at least three genes within a 10-kb region.

<sup>b</sup> Genes or clusters (as defined in <sup>a</sup>) that are present within 20 kb of TTSS components and effectors.

<sup>c</sup> The annotation of some of the genes in the table is based on multiple alignments to a nonredundant amino acid database as well as Hidden Markov Models (HMMs), which identify domains in protein families (Pfam and TIGRFAMs). Additional information on the individual loci and protein families is available in The Institute for Genomic Research Comprehensive Microbial Resource (<http://www.tigr.org/tigr-scripts/CMR2/CMRHomePage.spl>).