

Locus tag ATCC9714PCS11_“x”	ORF	Predicted function	pCP13 homologue (aa identity)	pCLL homologue (aa identity)	<i>C. difficile</i> P49 homologue (aa identity)	pCP13 vs. pCLL (aa identity)	pCP13 vs. CdP49 (aa identity)	pCLL vs. CdP49 (aa identity)
00231	<i>cnaB</i>	Surface protein, putative adhesion factor	<i>pcp57/cnaB</i> (43.0%)	-	-	-	-	-
00241	<i>241</i>	Unknown	-	-	-	-	-	-
00251	<i>251</i>	Unknown	-	-	-	-	-	-
00261	<i>topA</i>	Type I Topoisomerase	<i>pcp47/topA</i> (50.9%)	-	QUA_0964 (58.8%)	-	-	-
00271	<i>271</i>	Unknown	-	-	-	-	-	-
00281	<i>281</i>	Transmembrane protein	-	-	-	-	-	-
00291	<i>cstA</i>	Membrane protein	<i>pcp53</i> (40.8%)	<i>cll4</i> (35.5%)	QUA_0952 (35.5%)	68.0%	56.4%	55.1%
00301	<i>cstD4</i>	VirD4-like ATPase (Coupling Protein)	<i>pcp51</i> (49.3%)	<i>cll5</i> (51.9%)	QUA_0953 (55.5%)	55.4%	49.2%	51.8%
00311	<i>cstB6</i>	VirB6-like membrane channel protein	<i>pcp50</i> (35.5%)	<i>cll6</i> (28.7%)	QUA_0954 (32.2%)	37.4%	29.1%	28.5%
00321	<i>cstE</i>	Unknown	<i>pcp49</i> (38.0%)	<i>cll7</i> (34.8%)	QUA_0955 (52.7%)	52.2%	46.7%	40.5%
00331	<i>cstF</i>	Unknown	<i>pcp48</i> (41.5%)	<i>cll8</i> (38.6%)	QUA_0956 (45.0%)	53.1%	41.6%	45.9%
00341	<i>cstB4</i>	VirB4-like ATPase	<i>pcp46</i> (51.9%)	<i>cll9</i> (52.8%)	QUA_0957 (53.3%)	63.9%	48.9%	50.9%
00351	<i>cstH</i>	Phage protein	-	-	-	-	-	-
00361	<i>cstI</i>	Unknown	-	-	QUA_0958 (27.2%)	-	-	-
00371	<i>cstB1</i>	Muramidase	<i>pcp44</i> (57.7%)	<i>cll11</i> (56.9%)	QUA_0959 (42.6%)	59.3%	45.7%	42.3%
00381	<i>cstK</i>	Single stranded DNA binding protein	-	<i>cll19</i> (30.4%)	QUA_0963 (36.0%)	-	-	27.0%
00391	<i>cstL</i>	Unknown	-	-	QUA_0965 (31.8%)	-	-	-
00401	<i>cstM</i>	CopG family protein	<i>pcp41</i> (28.6%)	<i>cll13</i> (40.3%)	QUA_0966 (27.3%)	48.3%	38.1%	39.8%

00411	<i>cstN</i>	CopG family protein	<i>pcp41</i> (40.2%)	<i>cll13</i> (41.9%)	QUA_0966 (39.3%)	48.3%	38.1%	39.8%
00421	<i>cstO</i>	Unknown	-	-	-	-	-	-
00431	<i>cstD2</i>	Relaxase	<i>pcp39</i> (43.5%)	<i>cll15</i> (44.9%)	QUA_0968 (50.9%)	53.1%	45.5%	47.1%
00441	<i>cstQ</i>	Unknown	-	-	QUA_0969 (32.3%)	-	-	-
00451	<i>cstR</i>	Unknown	<i>pcp36</i> (43.5%)	-	QUA_0971 (44.0%)	-	40.6%	-