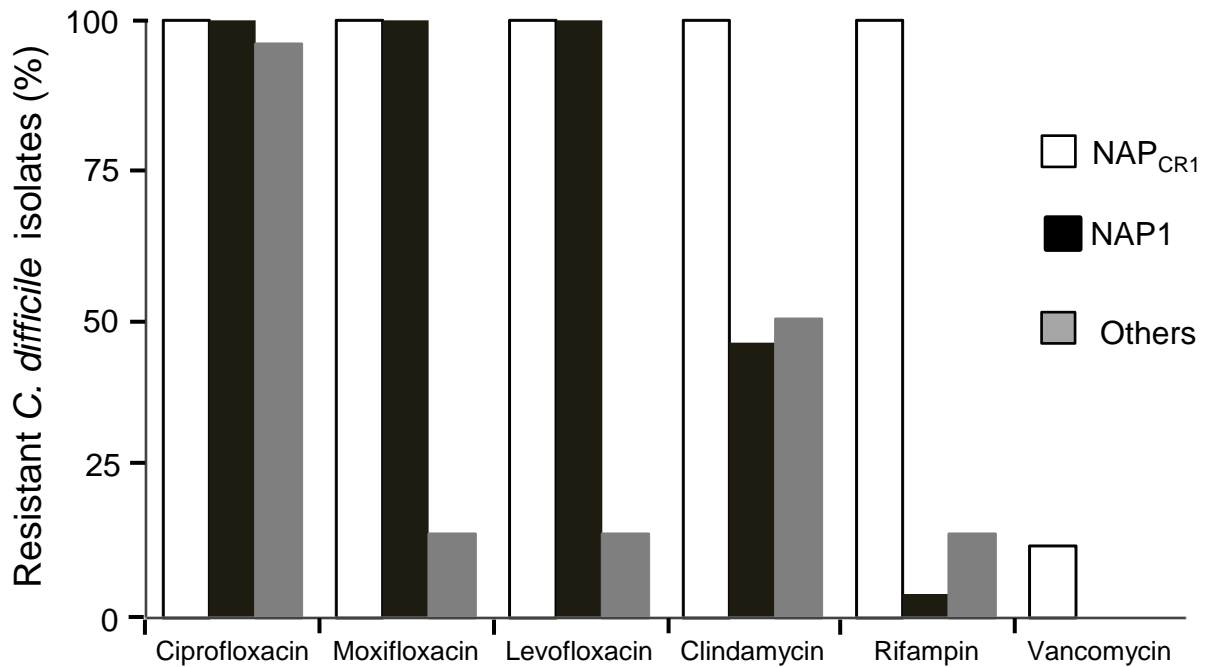


**FIGURE S1.** Effect of bacterial free supernatants on inflammatory cytokines and myeloperoxidase (MPO) activity in the ligated murine ileal model. Samples from ligated ileal groups prepared as indicated in figure 3 legend were processed for determination of MPO activity (A) or immunoenzymatic quantification of IL-6 (B), TNF-a (C) or IL-1b (D). \* $P < 0,05$ , compared to the groups without asterisk (One-way ANOVA with Bonferroni's correction).



**FIGURE S2** Antimicrobial resistance profile of the *C. difficile* genotypes identified during the outbreak. The 57 strains were divided in three groups: NAP<sub>CR1</sub> (n=18), NAP1 (n=26) and other genotypes (n=13). Antibiotic susceptibility test was determined by agar dilution. The percentage of resistant strains in each group is indicated.

**TABLE S1.** Comparison of genomic features of a representative NAP<sub>CR1</sub> strain and CD630. Counts indicate the number of genes assigned to each category of functional roles.

Subsystem Feature*	Counts	
	NAP <sub>CR1</sub>	630
Phages, Prophages, Transposable elements, Plasmids	55	34
DNA Metabolism	130	116
Cell Wall and Capsule	150	150
Virulence, Disease and Defense	100	100
Potassium metabolism	12	12
Miscellaneous	11	11
Cofactors, Vitamins, Prosthetic Groups, Pigments	252	249
Membrane Transport	54	68
Iron acquisition and metabolism	32	31
RNA Metabolism	105	102
Nucleosides and Nucleotides	150	146
Protein Metabolism	248	249
Cell Division and Cell Cycle	44	41
Motility and Chemotaxis	68	68
Regulation and Cell signaling	31	24
Fatty Acids, Lipids, and Isoprenoids	95	95
Nitrogen Metabolism	9	9
Dormancy and Sporulation	68	68
Respiration	80	80
Stress Response	64	68
Metabolism of Aromatic Compounds	11	11
Amino Acids and Derivatives	297	301
Sulfur Metabolism	24	24
Phosphorus Metabolism	45	43
Carbohydrates	422	427

\*Details for subsystem and functional role assignment for the genes of each strain using SEED and RAST

servers are listed in *Materials and Methods*.

**TABLE S2.** Genes in NAP<sub>CR</sub> that are not found in *C. difficile* 630 strain by genomic comparison

<b>Gene ID</b>	<b>Protein length</b>	<b>Contig</b>	<b>Function</b>
fig 1496.572.peg.8	43	1	hypothetical protein
fig 1496.572.peg.38	38		hypothetical protein
fig 1496.572.peg.46	49		hypothetical protein
fig 1496.572.peg.73	54		FIG00516771: hypothetical protein
fig 1496.572.peg.139	41		hypothetical protein
fig 1496.572.peg.231	40		hypothetical protein
fig 1496.572.peg.242	83		hypothetical protein
fig 1496.572.peg.248	51		hypothetical protein
fig 1496.572.peg.294	50		hypothetical protein
fig 1496.572.peg.374	52		FIG00513157: hypothetical protein
fig 1496.572.peg.400	39		hypothetical protein
fig 1496.572.peg.441	39		hypothetical protein
fig 1496.572.peg.460	53		FIG00517932: hypothetical protein
fig 1496.572.peg.490	39		hypothetical protein
fig 1496.572.peg.629	105	2	FIG00528816: hypothetical protein
fig 1496.572.peg.635	138		Recombinase
fig 1496.572.peg.637	290		FIG00630629: hypothetical protein
fig 1496.572.peg.639	303		putative aminoglycoside 6-adenylyltransferase
fig 1496.572.peg.641	265		Aminoglycoside phosphotransferase
fig 1496.572.peg.643	220		5-amino-6-(5-phosphoribosylamino)uracil reductase (EC 1.1.1.193)
fig 1496.572.peg.645	124		hypothetical protein
fig 1496.572.peg.646	150		hypothetical protein
fig 1496.572.peg.647	48		hypothetical protein
fig 1496.572.peg.648	89		Iron-dependent repressor IdeR/DtxR
fig 1496.572.peg.651	49		hypothetical protein

fig 1496.572.peg.652	99	hypothetical protein
fig 1496.572.peg.653	229	hypothetical protein
fig 1496.572.peg.656	238	hypothetical protein
fig 1496.572.peg.658	60	hypothetical protein
fig 1496.572.peg.659	200	hypothetical protein
fig 1496.572.peg.661	72	hypothetical protein
fig 1496.572.peg.663	60	hypothetical protein
fig 1496.572.peg.664	64	hypothetical protein
fig 1496.572.peg.670	41	hypothetical protein
fig 1496.572.peg.672	63	FIG00514266: hypothetical protein
fig 1496.572.peg.677	99	FIG00521858: hypothetical protein
fig 1496.572.peg.678	204	hypothetical protein
fig 1496.572.peg.679	162	ADP-heptose synthase (EC 2.7.-.-) / D-glycero-beta-D-manno-heptose 7-phosphate kinase
fig 1496.572.peg.687	53	hypothetical protein
fig 1496.572.peg.688	110	hypothetical protein
fig 1496.572.peg.689	200	Chromosome (plasmid) partitioning protein ParB
fig 1496.572.peg.695	99	hypothetical protein
fig 1496.572.peg.523	78	hypothetical protein
fig 1496.572.peg.525	109	hypothetical protein
fig 1496.572.peg.581	40	hypothetical protein
fig 1496.572.peg.592	45	hypothetical protein
fig 1496.572.peg.596	66	hypothetical protein
fig 1496.572.peg.597	57	hypothetical protein
fig 1496.572.peg.599	46	hypothetical protein
fig 1496.572.peg.600	56	Transcriptional regulator
fig 1496.572.peg.601	59	Phage protein
fig 1496.572.peg.606	40	hypothetical protein
fig 1496.572.peg.607	82	hypothetical protein
fig 1496.572.peg.618	53	hypothetical protein

fig 1496.572.peg.620	42	hypothetical protein
fig 1496.572.peg.624	80	hypothetical protein
fig 1496.572.peg.627	38	hypothetical protein
fig 1496.572.peg.764	48	3 hypothetical protein
fig 1496.572.peg.798	48	hypothetical protein
fig 1496.572.peg.822	65	hypothetical protein
fig 1496.572.peg.922	48	4 FIG00520514: hypothetical protein
fig 1496.572.peg.959	43	hypothetical protein
fig 1496.572.peg.1021	38	5 hypothetical protein
fig 1496.572.peg.1059	52	FIG00527335: hypothetical protein
fig 1496.572.peg.1063	43	hypothetical protein
fig 1496.572.peg.1190	38	6 hypothetical protein
fig 1496.572.peg.1242	40	hypothetical protein
fig 1496.572.peg.1266	40	7 hypothetical protein
fig 1496.572.peg.1270	237	hypothetical protein
fig 1496.572.peg.1271	387	hypothetical protein
fig 1496.572.peg.1275	53	hypothetical protein
fig 1496.572.peg.1276	96	hypothetical protein
fig 1496.572.peg.1277	120	hypothetical protein
fig 1496.572.peg.1282	146	hypothetical protein
fig 1496.572.peg.1283	150	hypothetical protein
fig 1496.572.peg.1284	268	Phage protein
fig 1496.572.peg.1285	273	Phage protein
fig 1496.572.peg.1286	268	hypothetical protein

fig 1496.572.peg.1287	165	hypothetical protein
fig 1496.572.peg.1288	331	hypothetical protein
fig 1496.572.peg.1289	212	hypothetical protein
fig 1496.572.peg.1290	198	hypothetical protein
fig 1496.572.peg.1291	324	hypothetical protein
fig 1496.572.peg.1292	163	hypothetical protein
fig 1496.572.peg.1294	76	hypothetical protein
fig 1496.572.peg.1295	495	hypothetical protein
fig 1496.572.peg.1296	591	unknown
fig 1496.572.peg.1297	287	unknown
fig 1496.572.peg.1298	398	Phage protein
fig 1496.572.peg.1299	63	hypothetical protein
fig 1496.572.peg.1300	206	hypothetical protein
fig 1496.572.peg.1301	185	hypothetical protein
fig 1496.572.peg.1302	211	hypothetical protein
fig 1496.572.peg.1303	102	hypothetical protein
fig 1496.572.peg.1304	401	hypothetical protein
fig 1496.572.peg.1305	65	hypothetical protein
fig 1496.572.peg.1306	80	hypothetical protein
fig 1496.572.peg.1307	209	hypothetical protein
fig 1496.572.peg.1308	315	hypothetical protein
fig 1496.572.peg.1309	46	hypothetical protein
fig 1496.572.peg.1313	63	hypothetical protein
fig 1496.572.peg.1314	86	hypothetical protein
fig 1496.572.peg.1315	47	hypothetical protein
fig 1496.572.peg.1316	115	hypothetical protein
fig 1496.572.peg.1317	79	hypothetical protein
fig 1496.572.peg.1319	113	hypothetical protein
fig 1496.572.peg.1320	195	hypothetical protein

fig 1496.572.peg.1321	162	hypothetical protein
fig 1496.572.peg.1322	39	hypothetical protein
fig 1496.572.peg.1323	361	hypothetical protein
fig 1496.572.peg.1324	169	hypothetical protein
fig 1496.572.peg.1325	101	hypothetical protein
fig 1496.572.peg.1326	111	hypothetical protein
fig 1496.572.peg.1327	178	hypothetical protein
fig 1496.572.peg.1328	175	hypothetical protein
fig 1496.572.peg.1331	58	hypothetical protein
fig 1496.572.peg.1332	123	Death on curing protein, Doc toxin
fig 1496.572.peg.1334	298	hypothetical protein
fig 1496.572.peg.1335	113	hypothetical protein
fig 1496.572.peg.1336	47	hypothetical protein
fig 1496.572.peg.1337	78	hypothetical protein
fig 1496.572.peg.1338	138	hypothetical protein
fig 1496.572.peg.1339	65	hypothetical protein
fig 1496.572.peg.1340	39	hypothetical protein
fig 1496.572.peg.1341	181	RNA:NAD 2'-phosphotransferase
fig 1496.572.peg.1342	109	hypothetical protein
fig 1496.572.peg.1344	49	hypothetical protein
fig 1496.572.peg.1345	62	hypothetical protein
fig 1496.572.peg.1346	428	DNA ligase (ATP) (EC 6.5.1.1)
fig 1496.572.peg.1347	67	hypothetical protein
fig 1496.572.peg.1348	196	hypothetical protein
fig 1496.572.peg.1349	63	hypothetical protein
fig 1496.572.peg.1350	337	hypothetical protein
fig 1496.572.peg.1351	132	hypothetical protein
fig 1496.572.peg.1352	179	hypothetical protein
fig 1496.572.peg.1353	123	hypothetical protein



fig 1496.572.peg.1354	95	hypothetical protein
fig 1496.572.peg.1355	115	hypothetical protein
fig 1496.572.peg.1356	41	hypothetical protein
fig 1496.572.peg.1357	156	hypothetical protein
fig 1496.572.peg.1358	127	hypothetical protein
fig 1496.572.peg.1359	149	hypothetical protein
fig 1496.572.peg.1360	55	hypothetical protein
fig 1496.572.peg.1361	117	hypothetical protein
fig 1496.572.peg.1362	156	hypothetical protein
fig 1496.572.peg.1363	120	hypothetical protein
fig 1496.572.peg.1365	60	hypothetical protein
fig 1496.572.peg.1366	51	hypothetical protein
fig 1496.572.peg.1367	121	hypothetical protein
fig 1496.572.peg.1368	77	hypothetical protein
fig 1496.572.peg.1369	59	hypothetical protein
fig 1496.572.peg.1370	70	hypothetical protein
fig 1496.572.peg.1371	61	hypothetical protein
fig 1496.572.peg.1372	117	hypothetical protein
fig 1496.572.peg.1373	106	hypothetical protein
fig 1496.572.peg.1374	95	hypothetical protein
fig 1496.572.peg.1375	56	hypothetical protein
fig 1496.572.peg.1376	45	hypothetical protein
fig 1496.572.peg.1377	109	hypothetical protein
fig 1496.572.peg.1378	201	COG0030: Dimethyladenosine transferase (rRNA methylation)
fig 1496.572.peg.1379	70	hypothetical protein
fig 1496.572.peg.1380	84	hypothetical protein
fig 1496.572.peg.1382	374	Phage protein
fig 1496.572.peg.1383	144	hypothetical protein
fig 1496.572.peg.1384	52	hypothetical protein

fig 1496.572.peg.1386	222	hypothetical protein
fig 1496.572.peg.1387	57	hypothetical protein
fig 1496.572.peg.1388	282	hypothetical protein
fig 1496.572.peg.1389	192	Phage-associated recombinase
fig 1496.572.peg.1390	186	hypothetical protein
fig 1496.572.peg.1391	531	hypothetical protein
fig 1496.572.peg.1392	38	hypothetical protein
fig 1496.572.peg.1393	259	Phage toprim domain containing protein, YorJ B.subtilis homolog
fig 1496.572.peg.1396	101	hypothetical protein
fig 1496.572.peg.1397	380	hypothetical protein
fig 1496.572.peg.1399	125	hypothetical protein
fig 1496.572.peg.1400	44	hypothetical protein
fig 1496.572.peg.1401	82	hypothetical protein
fig 1496.572.peg.1402	178	hypothetical protein
fig 1496.572.peg.1403	269	hypothetical protein
fig 1496.572.peg.1404	51	hypothetical protein
fig 1496.572.peg.1405	65	hypothetical protein
fig 1496.572.peg.1406	209	serine/threonine protein phosphatase
fig 1496.572.peg.1407	85	hypothetical protein
fig 1496.572.peg.1408	95	hypothetical protein
fig 1496.572.peg.1409	270	Adenine-specific methyltransferase (EC 2.1.1.72)
fig 1496.572.peg.1410	121	hypothetical protein
fig 1496.572.peg.1411	191	Guanylate kinase (EC 2.7.4.8)
fig 1496.572.peg.1412	165	Crossover junction endodeoxyribonuclease RuvC (EC 3.1.22.4)
fig 1496.572.peg.1414	40	hypothetical protein
fig 1496.572.peg.1415	437	Il-IS_2, transposase # fragment
fig 1496.572.peg.1416	281	hypothetical protein
fig 1496.572.peg.1418	615	hypothetical protein
fig 1496.572.peg.1419	44	hypothetical protein

fig 1496.572.peg.1420	74	hypothetical protein
fig 1496.572.peg.1425	104	hypothetical protein
fig 1496.572.peg.1436	52	hypothetical protein
fig 1496.572.peg.1437	643	hypothetical protein
fig 1496.572.peg.1439	52	hypothetical protein
fig 1496.572.peg.1444	52	8 FIG00519437: hypothetical protein
fig 1496.572.peg.1447	40	hypothetical protein
fig 1496.572.peg.1465	50	FIG00520773: hypothetical protein
fig 1496.572.peg.1497	40	hypothetical protein
fig 1496.572.peg.1658	43	10 hypothetical protein
fig 1496.572.peg.1682	39	hypothetical protein
fig 1496.572.peg.1690	45	hypothetical protein
fig 1496.572.peg.1692	45	hypothetical protein
fig 1496.572.peg.1754	93	11 hypothetical protein
fig 1496.572.peg.1776	40	hypothetical protein
fig 1496.572.peg.1786	43	hypothetical protein
fig 1496.572.peg.1787	40	hypothetical protein
fig 1496.572.peg.1795	40	hypothetical protein
fig 1496.572.peg.1799	44	hypothetical protein
fig 1496.572.peg.1800	154	FIG00513238: hypothetical protein
fig 1496.572.peg.1825	45	12 hypothetical protein
fig 1496.572.peg.1856	48	hypothetical protein
fig 1496.572.peg.1868	43	hypothetical protein
fig 1496.572.peg.1903	49	hypothetical protein
fig 1496.572.peg.1918	41	hypothetical protein

fig 1496.572.peg.1933	38	hypothetical protein
fig 1496.572.peg.1935	54	FIG00520743: hypothetical protein
fig 1496.572.peg.1953	46	hypothetical protein
fig 1496.572.peg.1966	43	hypothetical protein
fig 1496.572.peg.2062	65	hypothetical protein
fig 1496.572.peg.2067	134	hypothetical protein
fig 1496.572.peg.2083	42	hypothetical protein
fig 1496.572.peg.2098	48	hypothetical protein
fig 1496.572.peg.2100	57	hypothetical protein
fig 1496.572.peg.2112	51	hypothetical protein
fig 1496.572.peg.2213	50	13 hypothetical protein
fig 1496.572.peg.2252	39	14 hypothetical protein
fig 1496.572.peg.2259	188	FIG00517580: hypothetical protein
fig 1496.572.peg.2285	53	15 Transcriptional regulator
fig 1496.572.peg.2286	57	Phage protein
fig 1496.572.peg.2287	48	hypothetical protein
fig 1496.572.peg.2289	39	hypothetical protein
fig 1496.572.peg.2290	145	Phage protein
fig 1496.572.peg.2291	85	hypothetical protein
fig 1496.572.peg.2292	251	Phage antirepressor protein
fig 1496.572.peg.2295	46	hypothetical protein
fig 1496.572.peg.2296	74	Phage protein
fig 1496.572.peg.2297	69	Phage protein
fig 1496.572.peg.2393	40	16 hypothetical protein

fig 1496.572.peg.2452	51	18 thioredoxin family protein
fig 1496.572.peg.2459	38	hypothetical protein
fig 1496.572.peg.2472	39	hypothetical protein
fig 1496.572.peg.2481	105	hypothetical protein
fig 1496.572.peg.2483	95	hypothetical protein
fig 1496.572.peg.2499	52	19 FIG00522519: hypothetical protein
fig 1496.572.peg.2530	50	hypothetical protein
fig 1496.572.peg.2534	38	20 hypothetical protein
fig 1496.572.peg.2561	134	Phage-related protein
fig 1496.572.peg.2562	46	hypothetical protein
fig 1496.572.peg.2564	70	hypothetical protein
fig 1496.572.peg.2565	58	Phage protein
fig 1496.572.peg.2566	57	Phage protein
fig 1496.572.peg.2567	263	Phage antirepressor protein
fig 1496.572.peg.2568	65	Phage protein
fig 1496.572.peg.2569	66	hypothetical protein
fig 1496.572.peg.2570	67	hypothetical protein
fig 1496.572.peg.2571	66	hypothetical protein
fig 1496.572.peg.2577	70	21 hypothetical protein
fig 1496.572.peg.2578	150	hypothetical protein
fig 1496.572.peg.2579	212	Lmo2276 protein
fig 1496.572.peg.2580	105	hypothetical protein
fig 1496.572.peg.2581	51	hypothetical protein
fig 1496.572.peg.2582	47	hypothetical protein
fig 1496.572.peg.2584	111	hypothetical protein
fig 1496.572.peg.2588	113	hypothetical protein

fig 1496.572.peg.2589	125	Transcriptional regulator, Cro/CI family
fig 1496.572.peg.2590	61	hypothetical protein
fig 1496.572.peg.2591	142	hypothetical protein
fig 1496.572.peg.2592	54	hypothetical protein
fig 1496.572.peg.2593	115	hypothetical protein
fig 1496.572.peg.2595	49	hypothetical protein
fig 1496.572.peg.2596	90	hypothetical protein
fig 1496.572.peg.2598	154	hypothetical protein
fig 1496.572.peg.2603	130	FIG00516442: hypothetical protein
fig 1496.572.peg.2604	550	Phage terminase, large subunit
fig 1496.572.peg.2605	385	Portal protein, phage associated
fig 1496.572.peg.2608	96	Phage protein
fig 1496.572.peg.2609	113	hypothetical protein
fig 1496.572.peg.2610	138	hypothetical protein
fig 1496.572.peg.2611	139	conserved protein
fig 1496.572.peg.2614	128	FIG00521161: hypothetical protein
fig 1496.572.peg.2615	56	hypothetical protein
fig 1496.572.peg.2625	508	22 Phage protein
fig 1496.572.peg.2626	540	Exonuclease SbcC
fig 1496.572.peg.2627	51	hypothetical protein
fig 1496.572.peg.2629	119	hypothetical protein
fig 1496.572.peg.2630	343	Phage major capsid protein
fig 1496.572.peg.2631	69	hypothetical protein
fig 1496.572.peg.2632	106	hypothetical protein
fig 1496.572.peg.2633	128	hypothetical protein
fig 1496.572.peg.2634	137	hypothetical protein
fig 1496.572.peg.2635	275	FIG00513450: hypothetical protein
fig 1496.572.peg.2636	68	hypothetical protein

fig 1496.572.peg.2638	138		Phage-like element PBSX protein xkdM
fig 1496.572.peg.2640	51		hypothetical protein
fig 1496.572.peg.2641	348		Phage tail length tape-measure protein
fig 1496.572.peg.2642	95		hypothetical protein
fig 1496.572.peg.2643	1049		Phage tail length tape-measure protein
fig 1496.572.peg.2646	321		Phage-like element PBSX protein xkdQ
fig 1496.572.peg.2647	122		hypothetical protein
fig 1496.572.peg.2654	57		hypothetical protein
fig 1496.572.peg.2655	189		FIG00513626: hypothetical protein
fig 1496.572.peg.2656	97		FIG00515668: hypothetical protein
fig 1496.572.peg.2733	40	23	hypothetical protein
fig 1496.572.peg.2743	38		hypothetical protein
fig 1496.572.peg.2838	56		FIG00514757: hypothetical protein
fig 1496.572.peg.2872	43		hypothetical protein
fig 1496.572.peg.2881	42		hypothetical protein
fig 1496.572.peg.2886	77		hypothetical protein
fig 1496.572.peg.2894	166		hypothetical protein
fig 1496.572.peg.2916	50		hypothetical protein
fig 1496.572.peg.2923	258	24	conserved domain protein
fig 1496.572.peg.2924	40		hypothetical protein
fig 1496.572.peg.2933	340		Phage major capsid protein #Fam0021
fig 1496.572.peg.2934	127		hypothetical protein
fig 1496.572.peg.2936	77		hypothetical protein
fig 1496.572.peg.2937	91		hypothetical protein
fig 1496.572.peg.2938	163		Phage protein
fig 1496.572.peg.2946	169		hypothetical protein
fig 1496.572.peg.2949	40		Phage protein

fig 1496.572.peg.2951	118	hypothetical protein
fig 1496.572.peg.2955	232	Phage protein
fig 1496.572.peg.2982	63	26 Phage protein
fig 1496.572.peg.2984	308	Phage protein
fig 1496.572.peg.2985	195	Phage protein
fig 1496.572.peg.2986	151	Phage protein
fig 1496.572.peg.2987	799	Phage terminase, large subunit
fig 1496.572.peg.2990	210	Phage protein
fig 1496.572.peg.2991	65	Phage protein
fig 1496.572.peg.2992	77	hypothetical protein
fig 1496.572.peg.2995	74	hypothetical protein
fig 1496.572.peg.2996	117	Phage protein
fig 1496.572.peg.2997	117	Phage protein
fig 1496.572.peg.3006	50	hypothetical protein
fig 1496.572.peg.3007	64	hypothetical protein
fig 1496.572.peg.3009	91	Phage protein
fig 1496.572.peg.3010	81	Phage protein
fig 1496.572.peg.3049	40	27 hypothetical protein
fig 1496.572.peg.3064	174	29 hypothetical protein
fig 1496.572.peg.3091	264	31 FIG00513202: hypothetical protein
fig 1496.572.peg.3092	40	hypothetical protein
fig 1496.572.peg.3094	86	Phage protein
fig 1496.572.peg.3097	245	32 hypothetical protein
fig 1496.572.peg.3098	155	hypothetical protein



fig 1496.572.peg.3099	164	hypothetical protein
fig 1496.572.peg.3101	76	FIG00632701: hypothetical protein
fig 1496.572.peg.3106	39	33 hypothetical protein
fig 1496.572.peg.3110	47	34 hypothetical protein
fig 1496.572.peg.3181	47	hypothetical protein
fig 1496.572.peg.3185	52	FIG00522352: hypothetical protein
fig 1496.572.peg.3188	46	hypothetical protein
fig 1496.572.peg.3203	422	Plasmid recombination, MobE mobilization protein
fig 1496.572.peg.3204	45	hypothetical protein
fig 1496.572.peg.3206	208	Chloramphenicol acetyltransferase (EC 2.3.1.28)
fig 1496.572.peg.3207	69	TnpV
fig 1496.572.peg.3242	46	hypothetical protein
fig 1496.572.peg.3257	188	hypothetical protein
fig 1496.572.peg.3272	105	FIG00528816: hypothetical protein
fig 1496.572.peg.3310	40	hypothetical protein
fig 1496.572.peg.3314	51	hypothetical protein
fig 1496.572.peg.3329	39	41 hypothetical protein
fig 1496.572.peg.3332	40	hypothetical protein
fig 1496.572.peg.3361	49	hypothetical protein
fig 1496.572.peg.3432	56	Dihydrofolate reductase (EC 1.5.1.3)
fig 1496.572.peg.3447	42	FIG00516082: hypothetical protein
fig 1496.572.peg.3469	49	hypothetical protein
fig 1496.572.peg.3506	110	42 Phage protein
fig 1496.572.peg.3512	63	44 Phage protein

fig 1496.572.peg.3534	45	hypothetical protein
fig 1496.572.peg.3622	45	hypothetical protein
fig 1496.572.peg.3677	38	hypothetical protein
fig 1496.572.peg.3689	391	Mobile element protein
fig 1496.572.peg.3691	480	6'-aminoglycoside N-acetyltransferase (EC 2.3.1.-) (AAC(6')) / 2"-aminoglycoside phosphotransferase (EC
fig 1496.572.peg.3711	51	45 hypothetical protein
fig 1496.572.peg.3773	51	hypothetical protein
fig 1496.572.peg.3855	74	FIG00513646: hypothetical protein
fig 1496.572.peg.4017	174	46 FIG01032097: hypothetical protein
fig 1496.572.peg.4027	47	hypothetical protein
fig 1496.572.peg.4054	51	hypothetical protein
fig 1496.572.peg.4063	64	FIG00517121: hypothetical protein
fig 1496.572.peg.4097	280	47 Diadenylate cyclase spyDAC; Bacterial checkpoint controller DisA with nucleotide-binding domain
fig 1496.572.peg.4129	42	hypothetical protein
fig 1496.572.peg.4142	86	4-hydroxyphenylacetate decarboxylase, small subunit (EC 4.1.1.83)
fig 1496.572.peg.4189	42	hypothetical protein

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1 **TABLE S3.** Genes in *C. difficile* 630 strain that are not found in NAP<sub>CR1</sub> by genomic comparison

<b>Gene ID</b>	<b>Protein Length</b>	<b>Function</b>
fig 1496.1.peg.15	98	
fig 1496.1.peg.41	80	
fig 1496.1.peg.72	54	
fig 1496.1.peg.90	59	
fig 1496.1.peg.128	68	
fig 1496.1.peg.131	75	
fig 1496.1.peg.145	71	
fig 1496.1.peg.153	69	
fig 1496.1.peg.158	34	
fig 1496.1.peg.161	77	
fig 1496.1.peg.298	59	
fig 1496.1.peg.342	76	
fig 1496.1.peg.357	62	
fig 1496.1.peg.378	69	
fig 1496.1.peg.400	81	
fig 1496.1.peg.403	46	
fig 1496.1.peg.416	66	
fig 1496.1.peg.452	77	
fig 1496.1.peg.498	70	
fig 1496.1.peg.514	190	
fig 1496.1.peg.515	187	hypothetical protein
fig 1496.1.peg.516	61	
fig 1496.1.peg.517	64	
fig 1496.1.peg.526	55	
fig 1496.1.peg.529	77	

fig 1496.1.peg.534	76	
fig 1496.1.peg.540	252	
fig 1496.1.peg.541	81	
fig 1496.1.peg.547	110	hypothetical protein
fig 1496.1.peg.581	67	
fig 1496.1.peg.585	76	
fig 1496.1.peg.593	79	
fig 1496.1.peg.594	64	
fig 1496.1.peg.663	73	
fig 1496.1.peg.668	59	
fig 1496.1.peg.711	77	
fig 1496.1.peg.736	59	
fig 1496.1.peg.816	70	
fig 1496.1.peg.834	78	
fig 1496.1.peg.836	104	
fig 1496.1.peg.862	59	
fig 1496.1.peg.865	52	
fig 1496.1.peg.880	59	
fig 1496.1.peg.932	61	
fig 1496.1.peg.960	43	
fig 1496.1.peg.965	99	
fig 1496.1.peg.980	95	hypothetical protein
fig 1496.1.peg.1010	68	
fig 1496.1.peg.1014	125	
fig 1496.1.peg.1017	120	
fig 1496.1.peg.1025	64	
fig 1496.1.peg.1046	58	
fig 1496.1.peg.1092	73	
fig 1496.1.peg.1144	61	

fig 1496.1.peg.1159	71
fig 1496.1.peg.1199	99
fig 1496.1.peg.1261	63
fig 1496.1.peg.1271	66
fig 1496.1.peg.1324	70
fig 1496.1.peg.1345	61
fig 1496.1.peg.1366	26
fig 1496.1.peg.1375	63
fig 1496.1.peg.1386	67
fig 1496.1.peg.1422	58
fig 1496.1.peg.1481	92
fig 1496.1.peg.1508	84
fig 1496.1.peg.3944	26 Peptide chain release factor 2 unshifted fragment
fig 1496.1.peg.1545	61
fig 1496.1.peg.1559	63
fig 1496.1.peg.1590	64
fig 1496.1.peg.1655	59
fig 1496.1.peg.1694	88
fig 1496.1.peg.1700	62
fig 1496.1.peg.1707	64
fig 1496.1.peg.1713	99
fig 1496.1.peg.1714	68
fig 1496.1.peg.1715	93
fig 1496.1.peg.1716	64
fig 1496.1.peg.1727	77
fig 1496.1.peg.1788	46
fig 1496.1.peg.1795	73
fig 1496.1.peg.1836	68
fig 1496.1.peg.1839	79

fig 1496.1.peg.1891	74	
fig 1496.1.peg.1896	66	
fig 1496.1.peg.1910	85	
fig 1496.1.peg.1957	63	
fig 1496.1.peg.1960	70	
fig 1496.1.peg.1969	130	
fig 1496.1.peg.1980	72	
fig 1496.1.peg.1982	76	
fig 1496.1.peg.2002	63	
fig 1496.1.peg.2006	61	
fig 1496.1.peg.2027	86	threonine dehydratase biosynthetic
fig 1496.1.peg.2040	93	
fig 1496.1.peg.2054	77	
fig 1496.1.peg.2060	117	
fig 1496.1.peg.2061	63	
fig 1496.1.peg.2067	78	
fig 1496.1.peg.2094	73	
fig 1496.1.peg.2111	67	
fig 1496.1.peg.2183	30	
fig 1496.1.peg.2205	58	
fig 1496.1.peg.2274	78	
fig 1496.1.peg.2300	84	
fig 1496.1.peg.2349	78	
fig 1496.1.peg.2455	69	
fig 1496.1.peg.2490	87	
fig 1496.1.peg.2513	84	
fig 1496.1.peg.2546	96	
fig 1496.1.peg.2547	55	
fig 1496.1.peg.2551	98	Transposase

fig 1496.1.peg.2554	78
fig 1496.1.peg.2577	95
fig 1496.1.peg.2633	63
fig 1496.1.peg.2634	75
fig 1496.1.peg.2638	37
fig 1496.1.peg.2737	59
fig 1496.1.peg.2738	70
fig 1496.1.peg.2774	44 MSL leader peptide 2
fig 1496.1.peg.2776	46 Omega protein
fig 1496.1.peg.2778	44 MSL leader peptide 2
fig 1496.1.peg.2779	62 hypothetical protein
fig 1496.1.peg.2804	64
fig 1496.1.peg.2849	47
fig 1496.1.peg.2859	93
fig 1496.1.peg.2873	93
fig 1496.1.peg.2899	68
fig 1496.1.peg.2971	355 3-dehydroquinate synthase (EC 4.2.3.4)
fig 1496.1.peg.3011	67
fig 1496.1.peg.3023	59
fig 1496.1.peg.3193	94
fig 1496.1.peg.3204	65
fig 1496.1.peg.3228	33
fig 1496.1.peg.3234	182
fig 1496.1.peg.3236	34
fig 1496.1.peg.3240	269 Methyltransferase
fig 1496.1.peg.3241	80
fig 1496.1.peg.3242	65
fig 1496.1.peg.3244	110 hypothetical protein
fig 1496.1.peg.3246	242 sulfate adenylate transferase, subunit 2 (cysD)

fig 1496.1.peg.3375	70	
fig 1496.1.peg.3399	61	
fig 1496.1.peg.3467	69	
fig 1496.1.peg.3513	62	
fig 1496.1.peg.3537	36	
fig 1496.1.peg.3538	34	
fig 1496.1.peg.3539	28	
fig 1496.1.peg.3654	35	
fig 1496.1.peg.3675	66	
fig 1496.1.peg.3727	97	
fig 1496.1.peg.3776	77	hypothetical protein
fig 1496.1.peg.3787	68	
fig 1496.1.peg.3886	80	
fig 1496.1.peg.3906	70	
fig 1496.1.peg.3922	190	
fig 1496.1.peg.3923	187	hypothetical protein
fig 1496.1.peg.3924	61	
fig 1496.1.peg.3925	64	
fig 1496.1.peg.3934	55	
fig 1496.1.peg.3936	108	hypothetical protein

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