## **Supplemental Figure 1**

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**Control Strains** 



## Group 1



Group 2







**Fig. S1. CCV biogenesis phenotypes displayed after low MOI infection.** HeLa cells were infected at an MOI of one and were analyzed five days post-infection. (A) The parental NMII strain (wild type) and a Dot/Icm-deficient *icmL*::Tn mutant were used as a positive and negative controls for CCV biogenesis, respectively. The previously described effector mutants in Group 1 displayed vacuoles that were equivalent in size to the vacuoles containing the NMII strain. The seven effector mutants described in this study were in Group 2, and the vacuoles displayed by these mutants were all reduced in size relative to the NMII-containing vacuoles, but larger than the vacuoles containing the *icmL*::Tn mutant. (B) Average vacuole area was determined for each mutant by averaging the size of all CCVs in the infected cells of eight random images acquired for each strain at 600X magnification. \*, P-value < 0.001 indicate a significant difference in vacuole size compared to the size of vacuoles created by both the control NMII strain and the control *icmL*::Tn mutant strain as determined by Student's t-test.

## **Supplemental Figure 2**



**Fig. S2.** Intracellular replication kinetics of *cbu2028*::Tn in CHO and Vero cells. Intracellular replication over a seven day time period was determined by fold-increase in genomic equivalents relative to the first day of infection (DAI=0). Intracellular replication for the parental NMII strain (Wild Type, circle), the *cbu2028*::Tn mutant (square), are shown for (A) CHO cells and (B) Vero cells.

Supplemental Table 1

Effector	Gene Name	Genomic Insertion	CCV phenotype relative to
Gene		Site(s) (NC_002971.3)	wild type
cbu0012		12286	no difference
cbu0021	cig2, cvpB	19192; 19280	multi-vacuolar
cbu0041	cirA, coxCC1	38223; 39905	no difference*
cbu0048		45802; 45910	no difference
cbu0049		46672; 46994	no difference
cbu0062	cem1	56843	no difference
cbu0069	ankP	63560; 64162	no difference*
cbu0072	ankA	65711; 66529	no difference
cbu0122		111925	no difference
cbu0128		117057; 116556	no difference
cbu0129	coxCC2	116697; 116719	no difference
cbu0144	ankB	132025; 131885	no difference
cbu0295		262533; 262532	no difference
cbu0344		312064; 312062	no difference
cbu0388		352392	no difference*
cbu0393		357149	no difference
cbu0414	coxH1	369171	small
cbu0425	cirB	377489	no difference*
cbu0447	ankF	394134	no difference
cbu0513		453570	small
cbu0534		479403; 479397	no difference
cbu0781	ankG	722990	no difference
cbu0885	cetCb4	837619	no difference
cbu0937	cirC, coxDFB1	887774; 887017	small
cbu0978	cem3	930641	small
cbu1043	gacA.4	984953; 985088	no difference
cbu1063	cem4	1008750	no difference
cbu1079		1023981; 1023913	no difference
cbu1110		1056726	no difference
cbu1198		1142539	no difference
cbu1217	coxU2	1163535; 1162531	no difference
cbu1251		1202529	no difference
cbu1292	ankK	1246725; 1246917	no difference
cbu1379	coxK2	1328215	no difference
cbu1387	cem6	1337685	small
cbu1406	coxDFB3	1358424	no difference
cbu1457	coxTPR1	1410662	no difference

cbu1461	coxCC8	1417098; 1417177	small
cbu1524	caeA	1475299	no difference
cbu1525		1475299	no difference
cbu1530		1481087	no difference
cbu1566		1514458	no difference
cbu1569	coxCC12	1516828; 1517306	no difference
cbu1576		1523162; 1524222	no difference
cbu1599	coxCC13	1545143	no difference
cbu1634	cem8	1575307	no difference
cbu1636	coxCC14, cig55	1576055; 157917	no difference
cbu1639		1579591, 1579074	no difference
cbu1665		1601500	no difference*
cbu1677		1611159	no difference
cbu1699		1632356; 1632105	no difference
cbu1724		1658482	no difference
cbu1751	cig57	1682157; 1682180	small
cbu1752		1683656; 1683656	small
cbu1754		1684955; 1685031	small
cbu1757	ankM, cig58	1687615; 1688124	no difference
cbu1769	coxH3	1700167	no difference
cbu1780		1708652; 1709402	no difference
cbu1789		1721435; 1721437	no difference
cbu1802		1735650	no difference
cbu2007	cem12	1913579; 1913478	no difference
cbu2013	cem13	1920462	no difference
cbu2028		1935712; 1935580	small
cbu2052	cirD	1958215; 1958615	no difference
cbu2056		1962315; 1962702	no difference
cbu2064		1968966	no difference
cbu2069		1973680	no difference
cbu2076		1978466; 1978391	no difference
cbu2078	coxFIC1	1980054; 1979704	no difference

\* Mutants with transposon insertions in these genes were reported previously (15, 17) to have severe intracellular replication defects that were not detectable in this screen with this mutant library.

## Supplemental Table 2

Primer Name	Primer Sequence	Primer features
Partially complementary adapter 1	AGTTCTCCAGGTCTTGCGTTGCTCTTCCGATC*T	*Phosphothiorate linkage
Partially complementary adapter 2	GATCGGAAGAGCTCGTATGCCGTCTTCTGCTTG	5' phosphorylation
Universal adapter	CAAGCAGAAGACGGCATACGAGCTCTTCCGATC*T	*Phosphothiorate linkage
Barcode primers	AATGATACGGCGACCACCGAGATCTACACTCTTTC CCTACACGACGCTCTTCCGATCTNNNNNCGGGG ACTTATCAGCCAACC	NNNNN = unique barcode
Barcode1	AAACAC	
Barcode2	TGAAGG	
Barcode3	AACATA	
Barcode4	CGCGTC	
Barcode5	GATACA	
Barcode6	GGTGTG	
Barcode7	TAAGAA	
Barcode8	AGCGAG	
Barcode9	CGGTTA	
Barcode10	AGCTTT	
Barcode11	TGGTCT	
Barcode12	TATCCC	
Barcode13	TGTCGT	
Barcode14	CCCCAC	
Barcode15	ATACGA	
Barcode16	CCCTTG	
Barcode17	ACCGGC	
Barcode18	TTACTG	
Barcode19	GGAACT	
Barcode20	GTTATT	
Barcode21	AAAAGT	
Barcode22	AAGGGA	
Barcode23	AAGTAT	
Barcode24	ACATCT	