

1                           **Supplemental Information**

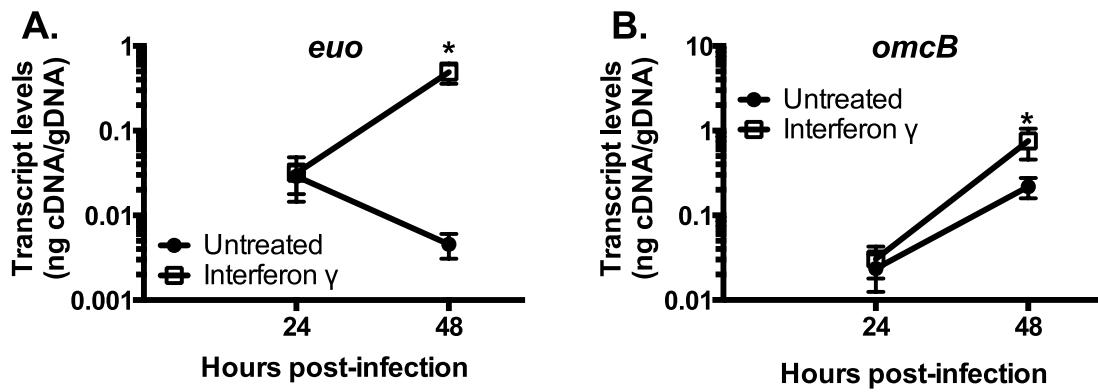
2     **Supplemental Figure S1.** Data from Figure 1C&D were plotted as average values with standard  
3     deviation similar to data representations elsewhere in the text. See legend of Figure 1 for more  
4     information.

5     **Supplemental Figure S2.** Comparison of transcript levels of selected genes between untreated and  
6     IFN $\gamma$ -treated Cpn at 24hpi. Samples were collected and analyzed as described in the legend of Figure  
7     1.

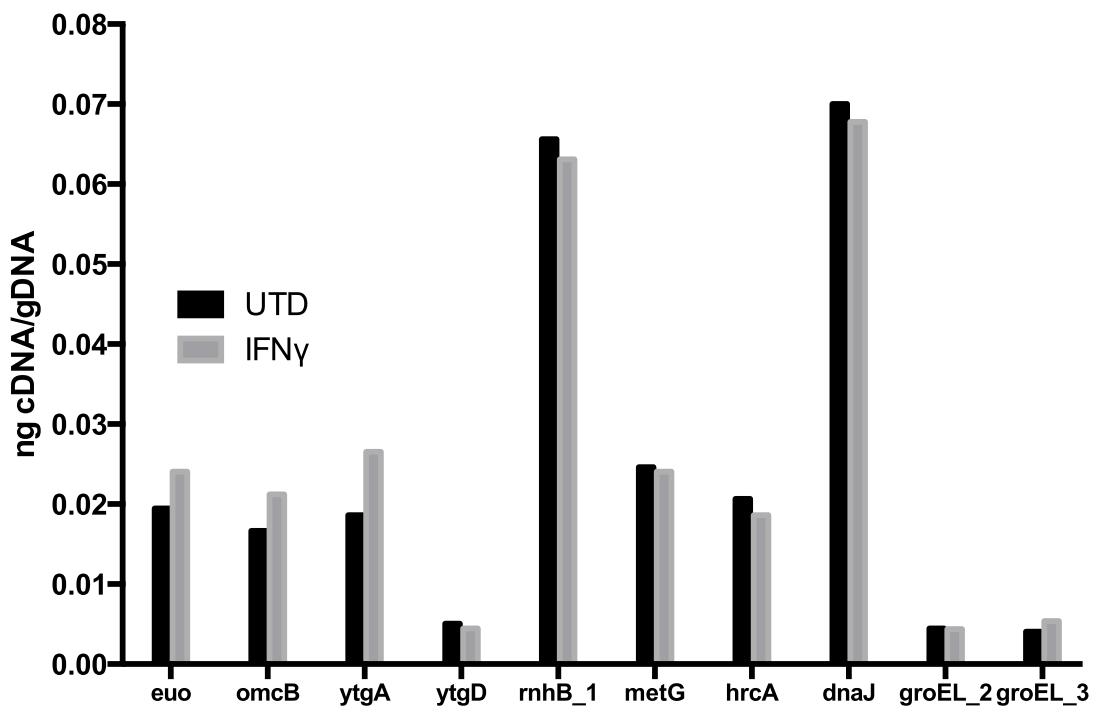
8     **Supplemental Table 1.** A list of primer sets used in the study for qPCR.

9     **Supplemental Table 2.** Codon content analysis of Cpn microarray data.

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Supplementary Figure 1



**Supplementary Figure 2**

Supplemental Table 1. List of Primers Used in Study.

<u>Gene</u>	<u>Forward Sequence</u>	<u>Reverse Sequence</u>	<u>Notes</u>
<i>Cpn0561/euo</i>	GGA ATA CCT GTG CAG AAG GTC TAC T	CCC AAG CGG CTC CCT TAC	Early gene
<i>Cpn0557/omcB</i>	CTG GAG ACT TGG TTC TTC ATG ATG	CCA CCA GGA GCT TCG AGT ACT G	Late gene
<i>Cpn0850/mreB</i>	GAT TGC TGA GGG CAT GTT AAA G	CCA GTA ATC CCA GAA GGA ACA G	0W
<i>Cpn0534/dksA</i>	GGG AAC GCT CAA GAG GTA AA	CTA GGC TAA TAG TCC GAT CAA AGG	0W
<i>Cpn0316/nusA</i>	CAG TCG TAG TCA CGC AGA AT	CCA GCT TCA CGA GCT ATC TT	0W
<i>Cpn0610/rho</i>	CTG ATA GGT GAA GGC GTG TTA G	CGA ATC TGA GCT GGA GAA ACA	0W
<i>Cpn1069/rodZ</i>	GCT CAT CAC GAG AGT CTC AAT C	CCC TGA GCA TAA ACT GGA GAA A	Upstream of all 5W residues
<i>Cpn0265/ubiA</i>	TGC AAG GAC TAC GGG AAT TG	ACA AGA TTC GCA GGA AGG AC	Upstream of all 5W residues
<i>Cpn0076/nusG</i>	ATC TGG CCT GGA TAC CTC TTA	GCC TCC AAG AAA CTC GAC TAT AC	Downstream of 2W/4W
<i>Cpn0544/obg</i>	GGT CAG TCA GGA GCT ACA AAT AA	TTC ACC AGT CTC AGC ATC AC	Downstream of 1W/2W
<i>Cpn0134/groEL_1</i>	TCC AGC TAC TTC TCC ACA AAT C	GCG TCC AGA TTC TGC TAC TT	0W
<i>Cpn0777/groEL_2</i>	TGG ATG TTG TGT GGG AAG AAG	GGG TGT TCT GTT CAG AGA TGA G	Downstream of 2W/2W
<i>Cpn0898/groEL_3</i>	GCG CGA CTA CTG GAC TTA TTT	TGA GCT TGT GTG TGG AGA TTC	Upstream of single W residue
<i>Cpn0520/clpP_1</i>	TAA CCT CAC CCG TCA CTA CT	CTT CCA GGA GCT GCA CAT AA	Downstream of 2W/3W
<i>Cpn0847/clpP_2</i>	CCA TGG GAG CCC TCT TAT TAT C	GTT CCG ATA ATG CCT CCA GAA	0W
<i>Cpn0511/rsbV_1</i>	CTC ACC TCG ATT ACA TGT CCA G	TCC CAG AAT GTT GTC CTA CTT G	0W
<i>Cpn0909/rsbV_2</i>	CTC AGT GCC TAG TGT TCA AGA G	GAA TCC CTG CGC TAC TGA TAT AC	0W
<i>Cpn0503/dnaK</i>	AAA CGT CAA GCA GTG ACA AAT C	AGA TCC GGA GGT GAC TGT ATA A	0W
<i>Cpn0501/hrcA</i>	GAA TCC CTA CAG ACC TAG CAT TAC	GCT AGG AAG CTG ACT GAT CTT ATC	0W
<i>Cpn0032/dnaJ</i>	TCT GGT CAA GGA GCT GTA AAC	CAT GTA GAG GCC ATG GAG AAA	0W
<i>Cpn0349/ytgA</i>	GGA TGC AAA TTC ACG TCC TTG T	CAG CGG TAG CAA GCC TAT T	6W
<i>Cpn0346/ytgD</i>	GCG AAT GCT GTC TCT CAT ACT	CAA GAG TCA AGG TAC CCA AAG A	1W
<i>Cpn0119/rnhB_1</i>	CGT AGT TGC TAG TGC CTG TAT	GAA CTT GGG CTC GTT GTT TAG	0W
<i>Cpn0122/metG 5'</i>	CCA ACT TAA AGC GTC TGG ATT G	GGA CAC GTC CCT TCT ACA TAA C	Upstream of all 8W residues
<i>Cpn0122/metG 3'</i>	CGT AAG GCT ACG AGT GTG ATT AT	TCA CGA GTC CCT TCT TTC AAT AG	Downstream of 5W/8W
<i>Cpn0741/greA 5'</i>	TCA GCA GAA GGT GTT GAT AGA G	CCG TCG TGA AAG GAA GTG AT	Downstream of 3W/11W
<i>Cpn0741/greA 3'</i>	GCT GTG CGT CAG ATG ATA GAA	CGG GCA TTT GGT GGA TAG TAA	Downstream of 8W/11W