

Supporting Information

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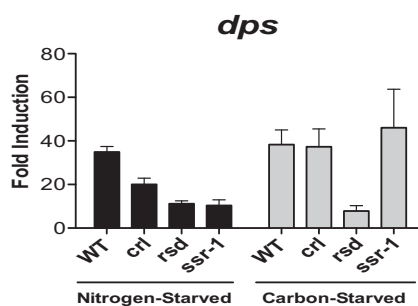


Fig. S1. *dps* expression under carbon-starvation and nitrogen-starvation conditions. Induction of the *dps* promoter in wild-type, *cri*, *rsd*, and *ssr* strains under nitrogen-starvation and carbon-starvation conditions. Error bars represent the SEM.

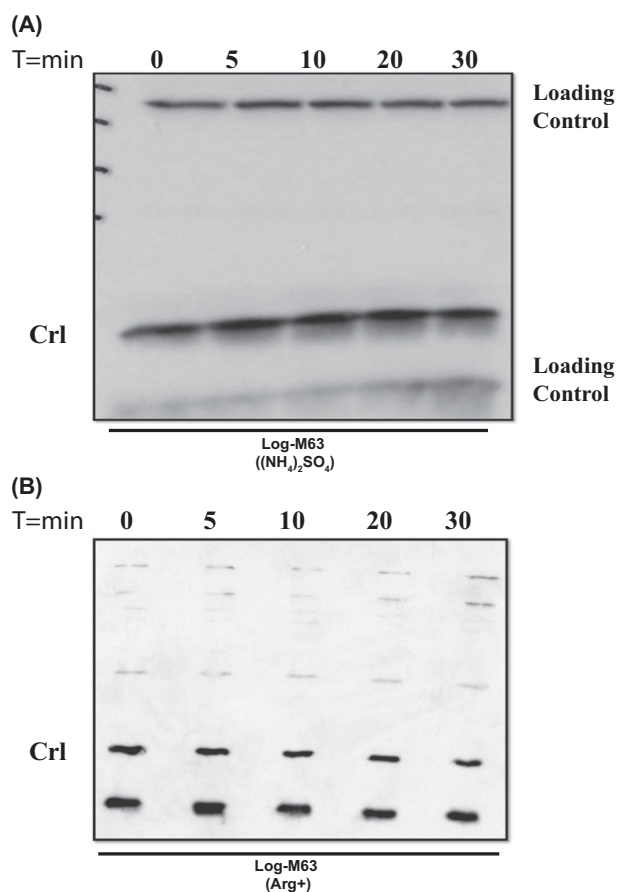


Fig. S2. Crl is a stable protein. Western blot of the Crl protein from MC4100 grown in M63 media supplemented with glucose (A) and from strain AZ-360 under nitrogen limiting conditions M63 (Arg+) (B). Cells were grown to midlog phase, and growth was stopped by adding chloramphenicol (250 μ g/mL). Cells were collected at the time points indicated and resuspended in sample buffer, boiled, and loaded on an SDS gel.

Table S1. List of strains used in this study

Strain/plasmid	Relevant genotype or phenotype	Source
MC4100	F ⁻ <i>araD139</i> Δ(<i>argF-lac</i>)U169 <i>rpsL150</i>	(1)
MG1655	F ⁻ λ ⁻ <i>ilvG</i> ⁻ <i>rfb-50</i> <i>rph-1</i>	(2)
HME6mutS	W3110Δ(<i>argF-lac</i>)U169 { λcl1857 Δ <i>crl-bioA</i> } <i>galK</i> _{TYR145UAG} <i>mutS::amp</i>	(3)
AZ-9	MC4100 <i>crl::kan</i>	This study
AZ-35	MC4100 Δ <i>crl</i>	This study
AZ246	MG1655 <i>crl::kan</i>	This study
AZ-309	MC4100 <i>rpoS::kan</i>	This study
AZ-310	MC4100 Δ <i>crl</i> , <i>rpoS::kan</i>	This study
AZ-14	MC4100 <i>ntrC::tn5</i>	(4)
VC-67	MC4100 <i>rsd::kan</i>	This study
VC-167	MC4100 <i>rpoN::kan</i>	This study
MJM-106	MC4100 <i>ssr-1</i>	(5)
MJM-372	MC4100 <i>crl</i> <i>c285 c375 c415</i> (<i>crl-serine</i>)	This study
AZ-400/pZS*11	AZ-35 Amp ^R	This study
AZ-360/pZS*11 <i>crl-1000</i>	AZ-35 <i>crl</i> ⁺ Amp ^R	This study
AZ-362/ pZS*11 <i>crl-1000 gc-12AT</i>	AZ-35 <i>crl</i> ⁺ Amp ^R	This study
pCP20	FLP ⁺ λcl857 ⁺ λp _R Rep ^{ts} Amp ^R Cam ^R	(6)

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- Costantino N, Court DL (2003) Enhanced levels of lambda Red-mediated recombinants in mismatch repair mutants. *Proc Natl Acad Sci USA* 100(26):15748–15753.
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- Cherepanov PP, Wackernagel W (1995) Gene disruption in *Escherichia coli*: TcR and KmR cassettes with the option of Flp-catalyzed excision of the antibiotic-resistance determinant. *Gene* 158(1):9–14.

Table S2. List of primers used in this study

Primer name	Sequence (5'→3')
5'ompA-RT	TACGCGATCACTCCTGAAATC
3'ompA-RT	GTAGGAAACACCCAGGCTCA
5'crl beg-RT	CACCCGAAGAGCAGATTGAT
3'crl beg-RT	GTTTCACGTTGACGCATACAG
5'dps-RT	ACCTGAAAGAAGACTGGCTGACC
3'dps-RT	AGGATATCTGCGGTGTCGTC
3'Crl-RT	GTTAACTTCACCGGCTCGTC
MJM335ss	CCGGTGCCGGTTTCACGTTGACGCTTACAGCCAGACTATC GAAAAAGAATCGATTATCTTTGCTCTTACCTTCACGAATATACGGGCC
MJM67F	GAGCATGTCCATATGACGTTACCGAGTGGACACC
MJM68R	GTGACGTCAGTCGACTCAGCCGTTAACTTCACCCGG
pZS-KpnI-Crl-c-terminal	GGGGTACCTCAGCCGTTAACTTCACCCGGC
pZS-XbaI-crl-1000upstream	CCTCTAGAGCCACCCAGGCGATGGTGTGGC
Crl-200-F	TCAGCCCGAAGAGGACTCACGC
Crl(-)gc-AT	TTGCTATCTCCTGTTGTGATATAACTGTTTTACCAAATTGGC
Crl(+)-gc-AT	GCCAAATTTGGTAAAACAGTTATATACACAACAGGAGATAGCAA

Underlined sequences are either substitutions or restriction sites.