



Fig. S1. Growth of *wt* and *dksA* strains on LB and minimal plates in the presence or absence of the vector control. A. The individual strains were streaked on LB or minimal plates; *ppGpp⁰* strain is included as a negative control. A Cannon PowerShot Pro1 camera was used. **B.** Images of individual colonies obtained by spreading cell culture dilutions on LB and minimal plates. A Leica MZ12 microscope with Jenoptik Progress C14 camera was used. Vector control (*pV*) is *pGB2*.

Table S1. Strains

CF1648	MG1655	(1)
CF6301	MG1655 but <i>relA251 rrnBP1':::lacZ ΔrelA25::kan</i>	(2)
CF7968	MG1655 but <i>rph+ ΔlacIZ</i>	(3)
CF9239	MG1655 but <i>ΔdksA::kan</i>	(3)
CF9240	MG1655 but <i>ΔdksA::tet</i>	(3)
CF10237	MG1655 but <i>ΔrelA256 ΔspoT212</i>	KK371 (4)
CF11657	MG1655 but <i>ΔgreA::cat</i>	(5)
CF11663	MG1655 but <i>ΔgreB::kan</i>	MG1655 x CLT255
CF11929	CF6301 <i>ΔdksA::kan</i>	(5)
CF11935	CF6301 <i>ΔspoT207 ΔlacIZ rrnBP1::lacZ</i>	(5)
CF11938	CF11935 <i>ΔdksA::tet</i>	(5)
CF12257	CF7968 but <i>rph+ ΔrelA256 ΔspoT212</i>	(6)
CF15615	CF10237 but <i>ΔrelA256 ΔspoT212 ΔlacIZ</i>	as CF12257 but for 10237
CF15617	MG1655 but <i>ΔlacIZ</i>	as CF6301 but for MG1655
CF12767	CF7968 (<i>λ PgreA::lacZ</i>)	Lysogen of CF7968, See Materials and Methods
CF12775	CF7968 (<i>λ PgreA::lacZ greA::cat</i>)	CF12767 x CF11657
CF12776	CF7968 (<i>λ PgreA::lacZ greB::kan</i>)	CF12767 x CF11663
CF12777	CF7968 (<i>λ PgreA::lacZ dksA::kan</i>)	CF12767 x CF9239
CF12773	CF7968 (<i>λ PgreB::lacZ</i>)	Lysogen of CF7968, See Materials and Methods
CF12790	CF7968 (<i>λ PgreB_B::lacZ greA::cat</i>)	CF12790 x CF11657
CF12791	CF7968 (<i>λ PgreB::lacZ greB::kan</i>)	CF12790 x CF11663
CF12792	CF7968 (<i>λ PgreB::lacZ dksA::kan</i>)	CF12790 x CF9239
CF12790	CF7968 (<i>λ PdksA::lacZ</i>)	Lysogen of CF7968, See Materials and Methods
CF12780	CF7968 (<i>λ PdksA::lacZ greA::cat</i>)	CF12790 x CF11657
CF12781	CF7968 (<i>λ PdksA::lacZ greB::kan</i>)	CF12790 x CF11663
CF12782	CF7968 (<i>λ PdksA::lacZ dksA::kan</i>)	CF12790 x CF9239
CF12602	CF12257 (<i>λ PgreA::lacZ</i>)	Lysogen of CF12257, See Materials and Methods
CF12719	CF12257 (<i>λ PgreA::lacZ greA::cat</i>)	CF12602 x CF11657
CF12720	CF12257 (<i>λ PgreA::lacZ greB::kan</i>)	CF12602 x CF11663
CF12721	CF12257 (<i>λ PgreA::lacZ dksA::kan</i>)	CF12602 x CF9239
CF12412	CF12257 (<i>λ PgreB::lacZ</i>)	Lysogen of CF12257, See Materials and Methods
CF12743	CF12257 (<i>λ PgreB::lacZ greA::cat</i>)	CF12412 x CF11657
CF12744	CF12257 (<i>λ PgreB::lacZ greB::kan</i>)	CF12412 x CF11663
CF12745	CF12257 (<i>λ PgreB::lacZ dksA::kan</i>)	CF12412 x CF9239
CF12604	CF12257 (<i>λ PdlsA::lacZ</i>)	Lysogen of CF12257, See Materials and Methods
CF12627	CF12257 (<i>λ PdksA::lacZ greA::cat</i>)	CF12604 x CF116574
CF12628	CF12257 (<i>λ PdksA::lacZ greB::kan</i>)	CF12604 x CF11663
CF12629	CF12257 (<i>λ PdksA::lacZ dksA::kan</i>)	CF12604 x CF9239
DV1618	CF15617 (<i>λ PgadA::lacZ</i>)	Lysogen of CF15617, See Materials and Methods
DV1640	CF15617 (<i>λ gadA::lacZ dksA::kan</i>)	DV1618 x CF9239
DV1622	CF15615 (<i>λ PgadA::lacZ</i>)	Lysogen of CF15615, See Materials and Methods
DV1641	CF15615 (<i>λ PgadA::lacZ dksA::kan</i>)	DV1622 x CF9239
DV1620	CF15617 (<i>λ PgadE::lacZ</i>)	Lysogen of CF15617, See Materials and Methods
DV1642	CF15617 (<i>λ PgadE::lacZ dksA::kan</i>)	DV1620 x CF9239
DV1624	CF15615 (<i>λ PgadE::lacZ</i>)	Lysogen of CF15615, See Materials and Methods
DV1643	CF15615 (<i>λ PgadE::lacZ dksA::kan</i>)	DV1624 x CF9239

CF16124	CF15615 but $\Delta greA::kan$	CF15515 x $\Delta greA::kan$ JW3148 (8)
CF16126	CF15615 $\Delta greA$	CF15615 x CF16124 (cured)
CF16132	CF15615 but $\Delta greB::kan$	CF15515 x $\Delta greB::kan$ JW3679 (8)
CF16128	CF15615 $\Delta greB$	CF15615 x CF16132 (cured)
CF16138	CF15615 $\Delta dksA::kan$	CF15615 x $\Delta dksA::kan$ JW0141 (8)
CF16151	CF15615 $\Delta greA \Delta greB$	CF16126 x JW3679 (cured)
CF16152	CF15615 $\Delta greA \Delta dksA::kan$	This work
CF16180	CF15615 $\Delta greB \Delta dksA::kan$	CF16132 x CF16138
CF16186	CF15615 $\Delta greA \Delta greB \Delta dksA::tet$	CF16151 x CF16138
CF15971	CF15617 but $greA::cat$	CF15617 x CF11657
CF15977	CF15617 but $greB::kan$	CF15617 x CF11663
CF15989	CF15617 but $dksA::tet$	CF15617 x CF9240
CF16050	CF15617 but $greA::cat greB::kan$	CF15971 x CF11663
CF16035	CF11617 but $greA::cat dksA::tet$	CF15871 x CF9240
CF16041	CF15617 but $greB::kan dksA::tet$	CF15977 x CF9240
CF16101	CF15617 but $greA::cat greB::kan dksA::tet$	CF16050 x CF11657
CF11762	CF1693 but $rpoB114$ (S531F) $btuB::Tn10$	allele: (9)
CF11770	CF1693 but $rpoB$ (G534C) $btuB::Tn10$	allele: (10)
CF11754	CF1693 but $rpoB3443$ (L533P) $btuB::Tn10$ (<i>rif-2</i>)	allele: (9)
CF11774	CF1693 but $rpoB$ (L571P) $btuB::Tn10$ (<i>rif-1</i>)	allele: unpublished Murphy & Cashel
CF11758	CF1693 but $rpoB3449$ (A532 Δ) $btuB::Tn10$	allele: (8)
CF11766	CF1693 but $rpoB3370$ (T563P) $btuB::Tn10$	“ “
CF16377	CF15615 but $rpoB$ (S531F) $btuB::Tn10$	CF15615 x CF11762
CF16383	As CF16377 but $\Delta greA$	CF16126 x “
CF16339	As CF16377 but $\Delta greB$	CF16132 x “
CF16395	As CF16377 but $\Delta dksA::kan$	CF16138 x “
CF16401	As CF16377 but $\Delta greA \Delta greB$	CF16151 x “
CF16419	As CF16377 but $\Delta greA \Delta dksA::kan$	CF16152 x “
CF16409	As CF16377 but $\Delta greB \Delta dksA::kan$	CF16180 x “
CF16427	As CF16377 but $\Delta greA::cat \Delta greB \Delta dksA::kan$	CF16186 x “
CF16379	CF15615 but $rpoB$ (G534C) $btuB::Tn10$	CF15615 x CF11770
CF16385	As CF16379 but $\Delta greA$	CF16126 x “
CF16391	As CF16379 but $\Delta greB$	CF16132 x “
CF16397	As CF16379 but $\Delta dksA::kan$	CF16138 x “
CF16403	As CF16379 but $\Delta greA \Delta greB$	CF16151 x “
CF16421	As CF16379 but $\Delta greA \Delta dksA::kan$	CF16152 x “
CF16413	As CF16379 but $\Delta greB \Delta dksA::kan$	CF16180 x “
CF16431	As CF16379 but $\Delta greA::cat \Delta greB \Delta dksA::kan$	CF16186 x “
CF16375	CF15615 but $rpoB$ (L533P) $btuB::Tn10$	CF15615 x CF11754
CF16381	As CF16375 but $\Delta greA$	CF16126 x “
CF16387	As CF16375 but $\Delta greB$	CF16132 x “
CF16393	As CF16375 but $\Delta dksA::kan$	CF16138 x “
CF16399	As CF16375 but $\Delta greA \Delta greB$	CF16151 x “
CF16417	As CF16375 but $\Delta greA \Delta dksA::kan$	CF16152 x “
CF16405	As CF16375 but $\Delta greB \Delta dksA::kan$	CF16180 x “
CF16423	As CF16375 but $\Delta greA::cat \Delta greB \Delta dksA::kan$	CF16186 x “
CF16380	CF15615 but $rpoB$ (L571P) $btuB::Tn10$	CF15615 x CF11774
CF16386	As CF16380 but $\Delta greA$	CF16126 x “
CF16392	As CF16380 but $\Delta greB$	CF16132 x “
CF16398	As CF16380 but $\Delta dksA::kan$	CF16138 x “
CF16404	As CF16380 but $\Delta greA \Delta greB$	CF16151 x “

CF16415	As CF16380 but $\Delta greA \Delta dksA::kan$	CF16152x “
CF16422	As CF16380 but $\Delta greB \Delta dksA::kan$	CF16180 x “
CF16433	As CF16380 but $\Delta greA::cat \Delta greB \Delta dksA::kan$	CF16186 x “
CF16376	CF15615 but <i>rpoB</i> (A532 Δ) <i>btuB::Tn10</i>	CF15615 x CF11758
CF16382	As CF16376 but $\Delta greA$	CF16126 x “
CF16388	As CF16376 but $\Delta greB$	CF16132 x “
CF16394	As CF16376 but $\Delta dksA::kan$	CF16138 x “
CF16400	As CF16376 but $\Delta greA \Delta greB$	CF16151 x “
CF16418	As CF16376 but $\Delta greA \Delta dksA::kan$	CF16152 x “
CF16407	As CF16376 but $\Delta greB \Delta dksA::kan$	CF16180 x “
CF16425	As CF16376 but $\Delta greA::cat \Delta greB \Delta dksA::kan$	CF16186 x “
CF16378	CF15615 but <i>rpoB</i> (T563P) <i>btuB::Tn10</i>	CF15615 x CF11766
CF16384	As CF163768 but $\Delta greA$	CF16126 x “
CF16390	As CF16378 but $\Delta greB$	CF16132 x “
CF16396	As CF16378 but $\Delta dksA::kan$	CF16138 x “
CF16402	As CF16378 but $\Delta greA \Delta greB$	CF16151 x “
CF16411	As CF16378 but $\Delta greA \Delta dksA::kan$	CF16152 x “
CF16420	As CF16378 but $\Delta greB \Delta dksA::kan$	CF16180 x “
CF16429	As CF16378 but $\Delta greA::cat \Delta greB \Delta dksA::kan$	CF16186 x “

references.

1. Xiao H, Kalman M, Ikehara K, Zemel S, Glaser G et al. (1991) Residual guanosine 3',5'-bispyrophosphate synthetic activity of *relA* null mutants can be eliminated by *spoT* null mutations. *J Biol Chem.* 266:5980-5990.
2. Vinella D, Cashel M, D'Ari R (2000) Selected Amplification of the cell division genes *ftsQ-ftsA-ftsZ* in *Escherichia coli*. *Genetics.* 156:1483-1492.
3. Brown L, Gentry D, Elliott T, Cashel M (2002) DksA affects ppGpp induction of RpoS at a translational level. *J Bacteriol* 184: 4455-4465.
4. Jishage M, Kvint K, Shingler V, Nyström T. (2002) Regulation of sigma factor competition by the alarmone ppGpp. *Genes Dev.* 16: 1260-1270.
5. Potrykus K, Vinella D, Murphy H, Szalewska-Palasz A, D'Ari R et al. (2006) Antagonistic regulation of *Escherichia coli* ribosomal RNA *rrnB* P1 promoter activity by GreA and DksA. *J Biol Chem.* 281:15238-15248.
6. Harinarayanan R, Murphy H, Cashel M (2008) Synthetic growth phenotypes of *Escherichia coli* lacking ppGpp and transketolase A (*tktA*) are due to ppGpp-mediated transcriptional regulation of *tktB*. *Mol. Microbiol.* 69: 882-899.
7. Baba T, Ara T, Hasegawa M, Takai Y, Okumura Y et al. (2006) Construction of *Escherichia coli* K-12 in-frame, single-gene knockout mutants: the Keio collection. *Mol Syst Biol* 2: 2006 0008 10.1038/msb4100050.
8. Jin DJ, Gross CA (1988) Mapping and sequencing of mutations in the *Escherichia coli rpoB* gene that lead to rifampicin resistance. *J Mol Biol.* 202: 45-58.
9. Murphy H, Cashel M (2003) Isolation of RNA polymerase suppressors of a (p)ppGpp deficiency. *Methods Enzymol* 371: 596-601.

Table S2. Primers used for *lacZ* fusion construction and *greA* mutagenesis.

PRIMERS	SEQUENCE (5' TO 3')
TRANSCRIPTIONAL FUSIONS	
GREAUP (-250)*	GGCAGAATTCATAATCTCGCGCTAACCAACCTGGAATCGA
GREADO (-13)	CACAGGATCCTGAATATTCCTGATAGGGCAAGTCTTCACCCACC
GREBUP (239)	GGCAGAATTCGTTCTCTTGCATTGTTTGTACTCCCAAAGG
GREBDO (-18)	CAGAGGATCCTGTTTGATAATGCGCACATTGGGTATAACG
DKSAUP (-256)	GGCAGAATTCGGTAGAAATTCTGGCTTACAAAGCGGAAA
DKSADO (-16)	CACAGGATCCTTAACACGCACTATCGATCCCCATGTTTCG
GADAUP (-313)	AAAGAATTCAGGTAGGCAAAGAGCTGCC
GADADO (-1)	AAAGGATCCTTCGAACTCCTTAAATTTATTG
GAD EUP (-806)	AAAGAATTCAAGCAGCAATAATTACCCCG
GAD EDO (-1)	AAAGGATCCAACCTTGCTCCTTAGCCGTTATC
GRE A MUTAGENESIS	
HMPR277	
FORWARD D41A + E44Y	GCGCGTGAGCATGGCGCCCTGAAATATAACGCCGAATACCAC
HMPR278	
REVERSE D41A + E44A	GTGGTATTCGGCGTTATATTTTCAGGGCGCCATGCTCACGCGC

* numbers in parenthesis indicate the extent of the fusion relative to translational start site

Table S3. Effects of GreA and GreB on ppGpp⁰ and ppGpp⁰ *dksA* strains.
(d- = *dksA*-; pv= vector control (pGB2); pA = pGreA; pB= pGreB)

A. inhibited in ppGpp⁰*dksA*-, unaffected by GreA or GreB

pv/ d- pv	pv/ d- pv	pv/ d- pv
90.40 <i>dksA</i>	3.08 <i>sdhD</i>	2.68 <i>yncE</i>
4.78 <i>yadB</i>	3.07 <i>ilvN</i>	2.68 <i>pheL</i>
3.72 <i>glnH</i>	3.07 <i>ydjN</i>	2.65 <i>aroF</i>
3.51 <i>exbB</i>	3.03 <i>fepA</i>	2.64 <i>artP</i>
3.46 <i>artQ</i>	2.98 <i>sdhA</i>	2.56 <i>proW</i>
3.46 <i>gltS</i>	2.90 <i>mglB</i>	2.54 <i>b0725</i>
3.33 IG_2238369_2238647-r	2.82 <i>artM</i>	2.53 <i>sdhC</i>
3.32 IG_761963_762236-r	2.70 <i>guaB</i>	2.52 IG_3530078_3530455-f
3.25 IG_1298469_1299205-f	2.70 <i>manX</i>	

B. activated in ppGpp⁰*dksA*-, unaffected by GreA or GreB

pv/ d- pv	pv/ d- pv	pv/ d- pv
0.06 <i>fimC</i>	0.32 <i>yciT</i>	0.37 <i>purH</i>
0.07 <i>fimA</i>	0.33 <i>beeE'</i>	0.38 <i>purD</i>
0.12 <i>fimG</i>	0.33 <i>metA</i>	0.38 <i>yfbT</i>
0.15 <i>ydfO</i>	0.33 <i>purA</i>	0.38 <i>sprT</i>
0.16 <i>fimD</i>	0.33 <i>yihI</i>	0.38 <i>sibD</i>
0.19 <i>fimF</i>	0.33 <i>aceB</i>	0.38 <i>yi81</i>
0.19 <i>aroA</i>	0.34 <i>psiA</i>	0.38 <i>jayE'</i>
0.22 <i>fimH</i>	0.34 <i>ybjP</i>	0.38 <i>ldhA</i>
0.25 <i>pheP</i>	0.34 <i>ydil</i>	0.38 IG_2468483_2468780-r
0.27 <i>pgaD</i>	0.34 <i>ycaK</i>	0.38 <i>msrA</i>
0.27 <i>ycgN</i>	0.34 <i>arnB</i>	0.38 <i>mdtI</i>
0.27 <i>rstB</i>	0.34 <i>yhhJ</i>	0.39 <i>treB</i>
0.27 <i>yceJ</i>	0.35 <i>ycaD</i>	0.39 <i>basR</i>
0.27 IG_804988_805220-r	0.35 <i>ybhC</i>	0.39 <i>yifN'</i>
0.27 <i>yfjV'</i>	0.35 <i>ybcW</i>	0.39 <i>alkB</i>
0.28 <i>ybhE</i>	0.35 IG_2267588_2267998-f	0.39 <i>yedF</i>
0.28 IG_2494586_2495076-f	0.36 IG_1671526_1671936-r	0.39 <i>mdtJ</i>
0.28 <i>mokB</i>	0.36 <i>dcuA</i>	0.39 <i>kup</i>
0.29 <i>yegD</i>	0.36 <i>yciK</i>	0.39 <i>Int</i>
0.29 IS183_b4452	0.36 <i>yicH</i>	0.39 <i>insZ'</i>
0.29 <i>ilvH</i>	0.36 <i>serA</i>	0.39 <i>cysQ</i>
0.30 <i>yeeA</i>	0.36 <i>ryjA</i>	0.40 <i>yifE</i>
0.30 <i>kdpB</i>	0.36 <i>ygjR</i>	0.40 <i>ycel</i>
0.30 <i>yjbE</i>	0.37 <i>adhE</i>	0.40 <i>yfaE</i>
0.31 <i>leuB</i>	0.37 <i>xthA</i>	0.40 <i>trmL</i>
0.31 <i>rhtB</i>	0.37 IG_668152_668518-r	0.40 <i>cbrC</i>
0.32 <i>gadW</i>	0.37 IG_986206_986807-f	
0.32 <i>yfiE</i>	0.37 <i>yzgL</i>	
0.32 IG_953690_954094-r	0.37 <i>cydA</i>	

C. activated in ppGpp⁰ *dksA*-, but GreA activates even more

pv/ d- pv	d-pv/d-pA	
0.12	0.06	hdeB
0.15	0.07	ybaS
0.29	0.07	hdeD
0.16	0.07	hdeA
0.40	0.17	rpsV
0.06	0.18	slp
0.31	0.22	rmf
0.26	0.40	aceK

D. Inhibited in ppGpp⁰ *dksA*-, but GreA inhibits even more

pv/ d- pv	d-pv/d-pA	
2.72	6.77	mgtA

E. GreA and/or GreB inhibit in ppGpp⁰ *dksA*-

d-pv/d-pA	d-pv/d-pB		d-pv/d-pA	d-pv/d-pB	
8.21	4.16	emrD	2.90	-	ycjF
7.34	4.46	deaD	2.89	-	IG_296321_296604-f
6.35	3.44	IG_1306670_1307039-r	2.89	-	IG_262171_262551-f
5.56	-	IG_1308294_1308592-r	2.87	-	IG_3636772_3637014-f
5.38	-	rarD	2.86	-	menA
5.06	-	ecnA	2.86	-	yeiE
4.33	2.84	hslV	2.85	-	yhdT
4.24	3.19	rdlA	2.83	-	IG_1903284_1903657-f
4.07	-	purP	2.83	-	dnaJ
4.02	2.82	hslU	2.82	-	nplI
4.01	-	ygeQ'	2.80	-	tdh
4.00	-	recX	2.79	-	IG_643191_643419-r
3.99	-	yhbE	2.77	-	ytfL
3.97	3.09	hspQ	2.76	-	yihA
3.79	-	puuC	2.76	-	rlmB
3.75	-	yicG	2.75	-	dnaT
3.70	-	IG_3326342_3326603-r	2.74	-	rhIE
3.67	-	IG_1269147_1269460-f	2.74	-	ycgF
3.67	-	IG_1269684_1269971-r	2.72	-	rpIS
3.63	-	IG_1927732_1928057-r	2.72	-	IG_3402255_3403072-f
3.60	-	IG_1360493_1360766-f	2.71	2.62	ucpA
3.59	-	yihG	2.71	-	erpA
3.50	2.83	yfcZ	2.71	-	selU
3.47	-	IG_3645282_3646157-f	2.67	-	umuC
3.39	-	IG_259325_259611-f	2.67	-	pitA

3.36	-	IG_831460_831690-f	2.66	-	htpX
3.33	-	tpr	2.65	-	yebE
3.32	-	bioA	2.65	-	cgtA
3.24	-	sibC	2.63	3.10	hokD
3.22	-	cyaR	2.63	-	yhjJ
3.18	-	IG_2925695_2926250-f	2.62	-	sbcD
3.16	-	IG_2660152_2660602-r	2.62	-	slmZ
3.16	-	yiiX	2.62	-	ade
3.15	-	rnk	2.61	2.58	glmZ
3.15	-	IG_2438141_2438404-r	2.61	2.98	yhhQ
3.14	-	IG_2689361_2689675-r	2.59	-	emrE
3.14	-	hslR	2.58	2.25	ryfA
3.07	-	ndK	2.58	-	ansB
3.07	-	gcd	2.57	-	ydiY
3.06	-	uvrD	2.56	-	ychE
3.05	2.75	recN	2.55	2.62	IG_856779_857018-f
3.04	-	sdaC	2.54	-	lldP
3.04	-	IG_631223_631611-r	2.54	-	prfC
3.03	2.52	IG_2815526_2815805-r	2.53	-	yahM
3.02	2.83	zntA	2.51	-	hscA
3.02	2.54	yfiP	2.51	-	mnmG
3.02	2.72	gfcA	2.51	-	lon
3.01	-	IG_1120179_1120464-r	-	3.16	uspG
3.00	-	IG_3679571_3679790-r	-	2.75	glmY
3.00	-	sdaB	-	2.84	degP
2.99	2.57	nusB	-	2.73	bssS
2.97	3.54	mutL	-	2.93	4.5SRNA
2.92	2.76	atpC	-	2.68	IG_1019277_1019632-r
2.91	-	IG_1994856_1995084-f	-	2.66	nhaA
2.91	-	yhgN	-	2.60	clpB
2.90	-	htpG	-	2.57	fabA
2.90	-	IG_3575186_3575415-r	-	2.51	recA

F. GreA and/or GreB activate in ppGpp⁰ *dksA*-

d-pv/d-pA	d-pv/d-pB		d-pv/d-pA	d-pv/d-pB	
0.01	0.23	gadB	0.27	-	cfa
0.01	0.35	gadA	0.27	0.28	gsiA
0.02	-	gadC	0.27	-	sufC
0.02	-	fliA	0.28	-	aroG
0.03	0.31	gadE	0.28	-	blc
0.04	0.17	flgC	0.28	-	IG_302830_303076-f
0.04	0.14	flgE	0.28	-	fic
0.04	-	IG_3656524_3656861-f	0.28	0.37	dld
0.04	0.16	flgB	0.28	-	sufB
0.05	0.19	flgD	0.28	0.28	pqqL
0.05	0.17	flgF	0.29	0.36	hisH

0.06	0.27	ycaC	0.29	-	sufA
0.07	0.31	fbaB	0.29	0.34	IG_779613_779776-r
0.07	0.28	katE	0.29	-	yjdJ
0.08	-	yhiM	0.29	-	IG_1739147_1739436-f
0.08	-	IG_3655198_3655995-f	0.30	-	ygaU
0.08	-	ybaT	0.30	-	ydiJ
0.08	-	otsB	0.30	-	yjjU
0.08	0.33	yegP	0.30	-	ivy
0.08	0.33	amyA	0.30	-	ybeM'
0.08	-	fliE	0.30	-	yehY
0.08	-	yhiD	0.31	-	IG_1860454_1860794-f
0.09	-	fliL	0.31	0.36	ompF
0.09	-	mdtE	0.31	-	gapC'
0.10	-	yiaG	0.31	0.30	gsiD
0.10	-	fliK	0.31	-	fhuF
0.10	-	yfcG	0.31	-	IG_338968_339388-r
0.10	-	flgH	0.31	-	asd
0.11	-	ygaM	0.31	-	yqjD
0.11	-	IG_1639579_1639878-r	0.31	-	yncB
0.11	-	msyB	0.31	-	IG_1066932_1067140-r
0.11	-	fliZ	0.31	-	ecnB
0.12	-	otsA	0.32	-	IG_836660_836887-f
0.12	-	flgL	0.32	-	fiil
0.12	0.32	ybgA	0.32	0.39	nadE
0.12	-	yceK	0.32	-	efeO
0.12	-	yeaG	0.32	-	ydcH
0.12	-	ycgB	0.32	-	IG_1488390_1488889-r
0.13	0.38	flgG	0.32	-	IG_2792037_2792273-f
0.13	-	poxB	0.32	-	yjjV
0.13	0.38	talA	0.32	-	flxA
0.13	-	mdtF	0.33	-	hisB
0.13	-	appC	0.33	-	chaC
0.14	-	elaB	0.33	-	IG_3119296_3119649-r
0.14	-	aidB	0.33	-	yphA
0.14	-	dctR	0.33	0.35	dppA
0.15	-	ydaM	0.33	-	IG_2769637_2769860-r
0.15	-	IG_3663441_3663809-r	0.33	0.38	potG
0.15	-	ygdl	0.33	-	IG_921814_922135-r
0.16	-	appB	0.34	-	uspB
0.16	-	yhcO	0.34	-	ybeM'
0.16	-	flgA	0.34	-	yniA
0.16	-	yahK	0.34	-	ygiW
0.16	0.38	ybhP	0.34	-	tsr
0.16	-	bfr	0.34	-	yjgH
0.16	-	fliG	0.34	-	yjdl
0.17	-	tam	0.34	-	IG_3182483_3182795-r
0.17	-	gapC'	0.34	-	efeU'
0.17	-	fliC	0.35	-	ypjF
0.17	-	flgJ	0.35	-	efeB
0.17	0.24	rfaL	0.35	-	cueR

0.17	-	IG_1864497_1864931-f	0.35	-	IG_1239173_1239557-r
0.18	0.22	yfjW	0.35	-	bfd
0.18	-	osmF	0.35	-	fliS
0.18	0.18	gtrS	0.35	-	mug
0.18	-	yciF	0.35	-	evgS
0.19	-	yodD	0.36	-	mtfA
0.19	0.40	chaB	0.36	-	IG_3662249_3662493-r
0.19	-	osmY	0.36	-	potF
0.19	-	fliM	0.36	-	ynhG
0.19	-	wrbA	0.36	-	IG_3663441_3663809-f
0.20	-	IG_3571136_3571407-f	0.36	-	ygaC
0.20	-	yehE	0.37	-	yneE
0.20	-	osmE	0.37	-	yjgF
0.20	-	yncA	0.37	-	holE
0.21	-	ybil	0.37	-	tyrB
0.21	-	cheA	0.37	-	flgN
0.21	0.24	gatC	0.37	-	tatE
0.21	-	gdhA	0.37	-	pptA
0.21	-	yccJ	0.37	-	ybiX
0.21	-	yfcF	0.37	-	ydcZ
0.22	-	tar	0.38	-	sufD
0.22	-	yghU	0.38	-	ghrA
0.22	-	gadX	0.38	-	ilvL
0.22	-	yhfG	0.38	-	ygiV
0.22	-	fliO	0.38	-	rssB
0.22	-	cspB	0.38	-	glcG
0.22	-	yddM	0.38	0.18	yliF
0.22	-	yebV	0.38	-	fhuC
0.22	0.38	yeiG	0.38	-	dcp
0.23	0.28	hisC	0.38	-	IG_2378439_2378741-r
0.23	-	ykgC	0.38	-	phrB
0.23	-	psiF	0.38	-	yhjY
0.23	-	tktB	0.39	-	hisQ
0.23	-	IG_1515027_1515337-r	0.39	-	shiA
0.23	-	flgM	0.39	-	sufS
0.23	0.31	hisG	0.39	-	IG_3154533_3154753-f
0.23	0.23	gsiC	0.39	-	grxB
0.23	0.26	gsiB	0.39	0.36	zinT
0.24	0.30	ybiC	0.39	-	IG_1050399_1050683-f
0.24	-	cheW	0.39	-	yagU
0.24	-	yghA	0.39	-	ybgl
0.24	-	ybaY	0.39	-	aroD
0.24	-	yeaQ	0.40	-	IG_3571136_3571407-r
0.25	0.29	gatB	0.40	-	fruL
0.25	-	ydiZ	0.40	-	inaA
0.25	-	IG_1565165_1565527-r	-	0.22	IG_2411153_2411489-f
0.25	-	ybeL	-	0.28	tauA
0.25	-	yegS	-	0.29	yihL
0.25	-	dkgA	-	0.29	IG_3358252_3358810-f
0.25	0.24	iaaA	-	0.29	ackA

0.26	-	caiF	-	0.30	yecS
0.26	0.30	hisF	-	0.34	appY
0.26	-	cheY	-	0.34	yjhC
0.26	0.37	hisD	-	0.36	gltF
0.26	-	yodC	-	0.37	IG_1796967_1797249-r
0.26	-	entA	-	0.37	murQ
0.27	-	marR	-	0.38	pta
0.27	-	nhoA	-	0.38	sppA
0.27	-	yqjC	-	0.39	yecC
0.27	-	ycaP	-	0.39	IG_3043922_3044187-r
0.27	-	entE	-	0.40	potH
0.27	-	entH	-	0.40	IG_2097636_2097883-r
0.27	-	IG_1388683_1388956-r			

G. inhibited in ppGpp⁰ *dkSA*⁻, but GreA and/or GreB prevents that

pv/ d- pv	d-pv/d-pA	d-pv/d-pB	
19.73	0.07	0.04	cysH
15.01	0.09	0.05	cysI
14.67	0.21	0.14	yeeE
12.72	0.06	0.04	cysD
12.18	0.22	0.18	yeeD
11.79	0.10	0.07	cysJ
10.76	0.08	0.05	cysC
10.54	0.08	0.06	cysN
9.95	0.20	0.39	nac
9.88	0.13	0.20	argF
9.00	0.15	0.24	argA
8.98	0.22	-	glnK
8.94	0.13	0.10	cysW
8.91	0.10	0.19	argI
8.74	0.13	0.28	ppsA
8.63	0.11	0.09	cysA
8.60	0.14	0.12	cysP
8.31	0.20	0.28	argH
8.03	0.18	0.22	argC
7.72	0.34	0.11	sbp
7.56	0.08	0.16	nadA
7.53	0.16	0.11	cysM
6.77	-	0.26	nmpC'
6.77	0.23	0.29	argB
6.56	0.23	0.21	uraA
6.48	0.29	0.25	asnA
5.94	0.19	0.14	fliY
5.50	0.13	0.21	nadB
5.46	0.36	-	amtF
5.40	0.22	0.19	cysU
5.38	0.15	0.14	IG_127588_127911-r

5.19	0.14	0.20	gltB
5.09	0.34	0.30	argG
5.07	0.15	0.13	cbl
5.00	0.25	0.19	IG_2539274_2539698-r
4.77	0.45	0.47	argE
4.75	0.10	0.10	yciW
4.20	0.17	0.13	ndh
4.14	0.15	0.22	pyrB
3.97	0.14	0.17	gatA
3.96	0.14	0.21	pyrI
3.80	0.20	0.17	ymgG
3.79	0.11	0.26	pnuC
3.70	0.15	0.18	codB
3.64	0.19	0.23	gatZ
3.62	-	0.30	tyrP
3.61	-	0.38	pheA
3.58	0.17	0.19	IG_356679_357014-r
3.49	0.36	-	rbsB
3.47	0.27	0.31	gatY
3.39	0.20	-	entB
3.38	0.20	0.27	gatD
3.32	0.25	0.31	artJ
3.25	0.26	0.22	aceE
3.25	0.33	-	fhuA
3.22	0.32	-	glnG
3.21	0.21	-	fiu
3.17	0.34	-	pyrD
3.16	0.22	0.36	gltD
3.12	0.39	-	glnP
3.11	0.37	0.35	cysK
3.05	0.28	0.25	oppC
3.01	0.31	0.26	pdhR
2.99	0.18	0.20	codA
2.85	0.26	0.21	ygbE
2.84	0.27	0.33	carB
2.76	0.39	-	yagl
2.75	0.27	0.19	lysA
2.69	0.15	0.13	yddB
2.64	0.17	0.24	hisA
2.63	0.20	-	entC
2.61	0.19	0.16	nlpA
2.58	0.30	0.36	carA
2.57	0.29	0.19	yfiD
2.55	-	0.41	yqgB
2.53	0.34	0.30	ymgD

H. activated in ppGpp⁰ *dksA*-, but GreA and/or GreB prevents that

pv/ d- pv	d-pv/d-pA	d-pv/d-pB	
0.07	-	3.23	leuL
0.09	6.83	10.04	IG_77300_77620-f
0.10	25.05	19.12	ibpB
0.10	10.94	7.28	IG_2241673_2241929-r
0.16	4.36	-	IG_1294422_1294668-r
0.18	-	3.76	hisL
0.19	3.63	3.42	ycdS
0.19	8.10	2.94	IG_1293368_1293648-r
0.20	2.76	-	ycdR
0.20	2.64	3.37	leuC
0.20	-	2.88	IG_1695077_1695296-r
0.22	2.95	2.47	yfaZ
0.22	3.60	-	ilvC
0.22	3.13	-	mmuP
0.23	8.63	5.54	ibpA
0.25	4.29	2.69	ldrD
0.25	-	3.00	leuD
0.25	3.15	2.87	yaaX
0.26	2.77	-	rhuA
0.27	3.58	3.29	IG_2702084_2702354-r
0.27	3.27	2.86	yaaH
0.27	4.52	4.13	bamE
0.28	2.86	3.11	ybdL
0.28	2.33	2.02	mmuM
0.28	2.96	-	IG_2428784_2429041-r
0.28	3.03	-	IG_3697829_3698191-f
0.29	3.44	4.71	IG_3483456_3483756-f
0.29	3.04	2.84	rdlB
0.29	3.38	-	IG_3119296_3119649-f
0.29	2.58	-	rapA
0.29	-	2.52	leuA
0.29	3.47	-	IG_3407589_3407916-f
0.30	-	2.65	osmC
0.30	2.79	-	IG_578860_579102-f
0.31	2.94	-	IG_2547427_2547665-r
0.31	6.83	7.36	IG_3717398_3717677-f
0.31	3.99	4.38	hslO
0.32	3.26	3.32	rdlC
0.32	4.06	3.42	rfaE
0.32	2.53	-	yehH
0.34	3.51	-	fadL
0.34	3.24	-	ymfQ
0.34	-	3.10	marB
0.34	5.71	3.34	ybeD
0.34	-	2.82	slyX
0.36	3.04	-	IG_3267244_3267856-f
0.36	-	2.78	uspA
0.36	4.13	3.76	yjcB

0.37	2.94	2.61	metL
0.37	2.82	-	IG_1223131_1223501-r
0.38	4.47	3.00	yhiQ
0.38	-	3.62	IG_2302414_2303127-f
0.38	2.51	-	ydiV
0.40	4.30	-	yebN
0.40	0.60	2.07	aceA
0.40	3.68	3.29	IG_376536_376758-r
0.40	2.79	-	prmA

I. GreB inhibits in ppGpp⁰ *dksA*+

pv/pB

5.00	amtB
4.82	glnK
3.95	nac
3.59	ddpX
3.16	ydcZ
3.13	ilvG
3.09	astC
2.53	IG_1308294_1308592-f
2.51	IG_925667_925950-f

J. GreB activates in ppGpp⁰ *dksA*+

pv/pB

0.39	yddB
0.37	yjiX
0.34	aroA
0.26	yjiY

K. GreA inhibits in ppGpp⁰ *dksA*⁺

pv/pA		pv/pA		pv/pA	
7.63	pdhR	3.26	cysM	2.80	cysJ
5.70	yfiD	3.25	IG_1360493_1360766-f	2.76	IG_3717398_3717677-f
5.11	sbp	3.11	IG_1689385_1689609-f	2.74	ycfJ
4.88	IG_127588_127911-r	3.09	trpE	2.73	yeeE
4.42	gspC	3.08	trpB	2.72	IG_3326342_3326603-r
4.39	yjaY	3.00	IG_262171_262551-f	2.71	argH
4.29	aceE	2.94	IG_2539274_2539698-r	2.69	IG_3198607_3198847-f
4.10	aldH	2.93	ecnA	2.67	cysH
3.91	ndh	2.92	speA	2.65	tpr
3.81	yqgB	2.92	argI	2.65	ygiA
3.75	argA	2.89	cysD	2.64	lysC
3.71	fliY	2.88	IG_406395_406651-f	2.64	cysI
3.59	asnA	2.87	trpC	2.63	ygeQ"
3.56	iraD	2.87	cysP	2.62	aroM
3.43	purP	2.86	ibpB	2.60	IG_2310770_2311103-r
3.42	fepC	2.83	speB	2.59	argC
3.40	IG_2302414_2303127-r	2.83	ilvG_2	2.56	sibC
3.31	rnk	2.83	ilvG_1	2.56	IG_3662249_3662493-r
3.30	cysU	2.82	fepD	2.55	IG_1150628_1150837-r
3.28	IG_192581_192871-r	2.81	mgtA	2.53	trpD

L. GreA activates in ppGpp⁰ *dksA*⁺

pv/pA		pv/pA		pv/pA	
0.12	ynfM	0.33	aceB	0.38	aldA
0.17	flgE	0.33	yegD	0.38	dadA
0.18	flgC	0.34	metA	0.39	fruA
0.22	poxA	0.34	fimA	0.39	rpsV
0.22	mmuP	0.34	fruB	0.39	nfrA
0.23	ycaC	0.34	pheP	0.39	gcvP
0.30	aceK	0.35	mdtJ	0.39	dkgA
0.30	yehH	0.35	gntT	0.39	talA
0.30	nfrB	0.35	yjeA	0.39	ybdL
0.31	aceA	0.35	mmuM	0.39	blc
0.31	ilvC	0.36	dadX	0.39	yiaG
0.32	katE	0.36	otsB	0.40	aroD
0.32	ydhO	0.36	fimC	0.40	rstB
0.32	ybgS	0.37	fbaB	0.40	serA
0.33	otsA	0.37	yjiX	0.40	ycaK