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## Research article

## Novel roles for two-component regulatory systems in cytotoxicity and virulence-related properties in *Pseudomonas aeruginosa*

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## **Supplementary**

**Table S1.** Effect of strain PA14 mutations in two-component regulatory system kinases and response regulators on cytotoxicity towards human bronchial epithelial cells. In general 3 or more experiments were performed assessing cytotoxicity (measured as LDH release after 8 hours of infection); results are expressed as the mean  $\pm$  standard error expressed as a percentage of cytotoxicity caused by WT.

PA14 locus ID	PAO1 homolog	Gene name	Description	Mean % of WT			
				± standard error			
Mutants with incr	Mutants with increased cytotoxicity						
PA14_07840	PA0601	agtR	Two-component response regulator; contains a CheY-like	$119.9 \pm 4.2$			
			receiver domain; amine uptake				
PA14_52260	PA0928	gacS	Sensor/response regulator hybrid; multi-host virulence through	$134.3 \pm 4.7$			
			regulation of small regulatory RNAs RsmZ and RsmY				

PA14 locus ID	PAO1 homolog	Gene name	Description	Mean % of WT
				± standard error
PA14_49170	PA1180	phoQ	Two-component sensor kinase PhoQ	$121.1 \pm 5.7^{\ddagger}$
PA14_45590	PA1458		Probable two-component sensor kinase; putative homolog of	$144.3 \pm 6.3$
			E. coli chemotaxis regulator CheA	
PA14_33780	PA2388	fpvR	Probable Fe <sup>2+</sup> -dicitrate sensor kinase	$121.6 \pm 7.9$
PA14_30650	PA2586	gacA	Response regulator; multi-host virulence through regulation of small regulatory RNAs RsmZ and RsmY	$144.7 \pm 15.8$
PA14_26810	PA2882		Probable two-component sensor kinase; has binding domain homologous to that found in sensors of C4-dicarboxylates	$139.3 \pm 15.7$
PA14_16470	PA3704	wspE	CheA-type sensor kinase; c-di-GMP regulation	$134.5 \pm 10.4$
PA14_62530	PA4725	cbrA	Two-component sensor kinase; required for carbon-nitrogen balance and control of catabolite repression	$158.8 \pm 9.1$
PA14_62540	PA4726	cbrB	Two-component response regulator; required for carbon- nitrogen balance and control of catabolite repression	$140.8 \pm 9.7$
PA14_67680	PA5125	ntrC	Two-component response regulator; regulates use of carbon and nitrogen	122.6 ±4.1
PA14_67670	PA5124	ntrB	Two-component sensor kinase; regulates use of carbon and nitrogen	117.1 ±5.8
PA14_70790	PA5364		Probable two-component response regulator	130.4 ±9.1
PA14_72740	PA5512	mifS	Two-component sensor kinase; regulates biofilm development	$119.6 \pm 13.2$
Mutants with decr	reased cytotoxicity			
PA14_05320	PA0408	pilG	Type IV pilus response regulator; required for pilus extension and twitching motility	$28.8 \pm 4.4$
PA14_05330	PA0409	pilH	Type IV pilus response regulator; required for pilus retraction	$25.8 \pm 1.5$
PA14_50220	PA1097	fleQ	Flagella major transcriptional regulator; cyclic-di-GMP responsive; potential FleSR modulator	77.6 ±12.7
PA14_50200	PA1098	fleS	Two-component sensor kinase; required for hook and basal body protein biosynthesis for flagellum assembly	$20.6 \pm 6.7^{\ddagger}$
PA14_50180	PA1099	fleR	Two-component response regulator; required for hook and basal body protein biosynthesis for flagellum assembly	$12.3 \pm 0.8^{\ddagger}$
PA14_16500	PA3702	wspR	Two-component response regulator with GGDEF domain; c-di-GMP regulation	$16.8 \pm 3.7$
PA14_12780	PA3948	rocA1	Two-component response regulator; cyclic-di-GMP regulation	$53.2 \pm 0.9$
PA14_60260	PA4547	pilR	Two-component response regulator; required for pilus expression and therefore for type IV pilus biosynthesis	$14.6 \pm 4.5$
PA14_68230	PA5165	dctB	Two-component sensor kinase; regulates a C4-dicarboxylate transport system with DctD	$14.9 \pm 3.5$
PA14_69470	PA5261	algR	Alginate biosynthesis regulatory protein	33.2 ±11.7
No major change	in cytotoxicity			
PA14_00430	PA0034		Probable two-component response regulator	$108.9 \pm 5.1$
PA14_01860	PA0150		Probable transmembrane sensor kinase	$101.4 \pm 3.0$
PA14_02250	PA0178	cheA	Probable two-component sensor kinase	$107.5 \pm 4.1$
PA14_02260	PA0179		Probable two-component response regulator	103.1 ±4.1

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PA14 locus ID	PAO1 homolog	Gene name	Description	Mean % of WT ± standard error
PA14_06060	PA0463	creB	Two-component response regulator; catabolism and motility	94.2 ±3.3
PA14_06070	PA0464	creC	Two-component sensor kinase; catabolism and motility	$109.9 \pm 5.4$
PA14_07820	PA0600	agtS	Two component ensor kinase; amine uptake	$95.9 \pm 7.0$
PA14_54510	PA0756		Probable two-component response regulator	$103.3 \pm 1.3$
PA14_54500	PA0757		Probable two-component sensor kinase	109.4 ±4.9
PA14_52240	PA0930	pirS	Two-component sensor kinase; iron acquisition	$109.9 \pm 3.5$
PA14_49440	PA1157		Probable two-component response regulator	$107.5 \pm 3.8$
PA14_49420	PA1158		Probable two-component sensor kinase	$113.8 \pm 5.7$
PA14_49180	PA1179	phoP	Two-component response regulator; regulates cytotoxicity and resistance to cationic peptides	$104.4 \pm 1.8$
PA14_49160	PA1243		Probable sensor kinase /regulator hybrid	$94.0 \pm 8.5$
PA14_47390	PA1301		Probable transmembrane sensor kinase	$111.2 \pm 8.2$
PA14_46980	PA1336	aauS	Two-component sensor kinase; amino acid uptake	$110.1 \pm 0.3$
PA14_46850	PA1347		Probable transcriptional regulator	$106.6 \pm 1.3$
PA14_46370	PA1396		Probably two-component sensor kinase	$98.5 \pm 3.5$
PA14_46060	PA1422	gbuR	Transcriptional regulator	$95.5 \pm 4.5$
PA14_45880	PA1437		Probable two-component response regulator	$106.7 \pm 3.1$
PA14_45870	PA1438		Probable two-component sensor kinase	$111.6 \pm 4.5$
PA14_43350	PA1636	kpdD	Two-component sensor kinase; potassium transport regulation	$109.8 \pm 4.5$
PA14_42220	PA1727	mucR	Two component sensor kinase; alginate regulation	$113.0 \pm 4.5$
PA14_41490	PA1785	nasT	Response regulator; nitrate/nitrite assimilation	$103.2 \pm 2.6$
PA14_41270	PA1798	parS	Two-component sensor kinase; resistance to cationic peptides	$93.9 \pm 8.5$
PA14_41260	PA1799	parR	Probable two-component response regulator; resistance to cationic peptides	107.7 ±4.3
PA14_40570	PA1851		Probable two-component response regulator	$104.4 \pm 1.9$
PA14_38970	PA1976	ercS'	Probable two-component sensor kinase; ethanol oxidation	$112.7 \pm 5.1$
PA14_38900	PA1980	eraR	Probable two-component response regulator; ethanol oxidation	$98.7 \pm 5.0$
PA14_37980	PA2051		Probable transmembrane sensor kinase	$103.9 \pm 5.7$
PA14_36420	PA2177		Probable histidine sensor kinase	$100.6 \pm 4.6$
PA14_32580	PA2479		Probable two-component response regulator	$107.3 \pm 4.7$
PA14_31960	PA2523	czcR	Two-component response regulator; metal and imipenem resistance	$107.4 \pm 4.1$
PA14_31950	PA2524	czcS	Two-component sensor kinase; metal and imipenem resistance	$107.5 \pm 1.0$
PA14_30830	PA2572		Probable two-component response regulator	111.9 ±5.2
PA14_30700	PA2583		Probable sensor kinase/response regulator hybrid	$100.3 \pm 6.5$
PA14_29740	PA2656		Probable two-component sensor kinase	101.6 ±6.5
PA14_29360	PA2687	pfeS	Two-component sensor kinase; iron acquisition	$88.0 \pm 4.8$
PA14_27940	PA2798		Probable two-component response regulator	$108.6 \pm 3.4$
PA14_27810	PA2809	copR	Two-component response regulator; copper resistance	$106.4 \pm 5.3$
PA14_27800	PA2810	copS	Two-component sensor kinase; copper resistance	$93.7 \pm 2.3$

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PA14 locus ID	PAO1 homolog	Gene name	Description	Mean % of WT
				± standard error
PA14_27550	PA2824	sagS	Two-component sensor/regulator hybrid; motile/sessile switch, activates Gac/Hpt/Rsm system	$88.9 \pm 8.6$
PA14_24720	PA3044	rocS2	Two-component sensor kinase; c-di-GMP regulation	$96.9 \pm 4.5$
PA14_24710	PA3045	rocA2	Two-component response regulator; c-di-GMP regulation	$103.2 \pm 4.0$
PA14_24350	PA3077	cprR	Probable two-component system regulatory protein; cationic peptide resistance	112.8 ±9.5
PA14_24340	PA3078	cprS	Probable two-component sensor kinase; cationic peptide resistance	$98.4 \pm 6.2$
PA14_22960	PA3191	gtrS	Probable two-component sensor kinase; virulence	$107.3 \pm 4.4$
PA14_22940	PA3192	gltR	Two-component response regulator; virulence	$98.6 \pm 2.6$
PA14_22730	PA3206		Probable two-component sensor kinase	$102.5 \pm 6.3$
PA14_21700	PA3271		Probable two-component sensor kinase	$105.9 \pm 11.3$
PA14_20820	PA3343		Probable two-component response regulator	$108.8 \pm 7.1$
PA14_20780	PA3346		Probable two-component response regulator	$105.7 \pm 3.9$
PA14_20000	PA3409	hasS	Probable Fe <sup>2+</sup> -dicitrate sensor kinase; diguanylate cyclase	$102.5 \pm 5.7$
PA14_17670	PA3604	erdR	Two-component response regulator; ethanol oxidation	99.4 ±10.1
PA14_16350	PA3714		Probable two-component response regulator	110.9 ±5.3
PA14_13740	PA3878	narX	Two-component sensor kinase; nitrate respiration, biofilm, motility	$100.5 \pm 10.9$
PA14_12820	PA3900	fec <b>R</b>	Probable transmembrane sensor protein	$110.9 \pm 5.6$
PA14_12820	PA3946	rocS1	Two-component sensor kinase; c-di-GMP regulation	$103.4 \pm 7.4$
PA14_12810	PA3947	rocR	Antagonist of rocA1 response regulator; c-di-GMP regulation with EAL domain	$93.5 \pm 1.9$
PA14_11680	PA4032		Probable two-component response regulator	$106.0 \pm 2.6$
- PA14_11630	PA4036		Probable two-component sensor kinase	110.9 ±3.8
– PA14_10770	PA4112		Probable sensor/regulator hybrid	97.4 ±9.4
PA14_09690	PA4196	bfiR	Probable two-component response regulator; biofilm formation	90.8 ±9.9
PA14_09680	PA4197	bfiS	Two-component sensor kinase; biofilm formation	97.5 ±2.9
PA14_55780	PA4293	pprA	Two-component sensor kinase; outer membrane permeability and resistance	$108.8 \pm 4.1$
PA14_55810	PA4296	pprB	Probable two-component response regulator; outer membrane permeability and resistance	$100.5 \pm 6.8$
PA14_56950	PA4381	colR	Probable DNA-binding response regulator; polymyxin resistance	$103.8 \pm 2.5$
PA14_57140	PA4396		Probable two-component response regulator	$103.0 \pm 10.0$
PA14_57170	PA4398		Probable two-component response regulator	$105.5 \pm 12.7$
PA14_58300	PA4493	roxR	Two-component response regulator; regulates production of cyanide insensitive oxidase	$107.0 \pm 3.8$
PA14_58320	PA4494	roxS	Two-component sensor kinase; regulates production of cyanide insensitive oxidase	$108.1 \pm 6.4$
PA14_60250	PA4546	pilS	Two-component sensor kinase; type IV pili production	113.5 ±5.3

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A14 locus ID	PAO1 homolog	Gene name	Description	Mean % of WT ± standard error
PA14_63150	PA4776	pmrA	Two-component response regulator; cationic peptide resistance	104.3 ±2.3
PA14_63160	PA4777	pmrB	Two-component sensor kinase; cationic peptide resistance	$92.0 \pm 10.8$
PA14_64050	PA4843	gcbA	Probable two-component response regulator; diguanylate cyclase	$108.8 \pm 3.0$
PA14_64880	PA4983		Probable two-component response regulator	$108.5 \pm 5.4$
PA14_68250	PA5166	dctD	Two-component response regulator; C4-dicarboxylate metabolism	97.1 ±3.6
PA14_68680	PA5199	amgS	Two-component sensor kinase; membrane stress, aminoglycoside resistance	91.5 ±3.3
PA14_70750	PA5360	phoB	Two component response regulator; phosphate regulation and quorum sensing	$105.9 \pm 3.5$
PA14_70760	PA5361	phoR	Two-component sensor kinase; phosphate regulation and quorum sensing	$108.8 \pm 3.3$
PA14_72380	PA5483	algB	Two-component response regulator; alginate production	96.1 ±3.2
PA14_72390	PA5484	kinB	Two-component sensor kinase; alginate production	93.6 ±5.6
PA14_59770		rcsB	Two component response regulator; cup fimbria and biofilm formation	$104.0 \pm 5.2$
PA14_59790		pvrR	Two component response regulator; EAL domain, c-di-GMP regulation, cup fimbria and biofilm	103.6 ±4.7
PA14_59800		pvrS	Two component sensor kinase; c-di-GMP regulation, cup fimbria and biofilm	110.0 ±4.2

†Mutants are listed in order of the PAO1 homolog. Mutants not in the screen included PA1637 *kpdE*—response regulator with KpdD, potassium transport; PA1786 *nasS*—sensor with *nasT*, nitrate/nitrite assimilation; PA2686 *pfeR*—response regulator with *pfeS*, iron acquisition; PA0929 *pirR*—response regulator with *pirS*, iron acquisition; PA3345 *htpB*—response regulator with *sagS*, activates Gac/Htp/Rsm system; PA3879 *narL*—response regulator with *narX*, nitrate respiration, biofilm, motility; PA3974 *ladS*—sensor, virulence, biofilm formation, cytotoxicity; PA4380 *colS*—sensor with *colR*, polymyxin resistance; PA4101 *bfmR*—response regulator with *bfmS*, biofilm development; PA4102 *bfmS*—sensor with *bfmR*, biofilm development; PA4856 *retS*—sensor, virulence, biofilm formation, cytotoxicity; PA4959 *fimX*—sensor (orphan?); contains GGDEF and EAL domains; cytotoxicity; PA5200 *amgR*—response regulator with *amgS*, membrane stress and aminoglycoside resistance; PA5262 *fimS*—sensor with *algR*, virulence, alginate biosynthesis, motility, biofilm; PA5511 *mifR*—response regulator with *mifS*, biofilm development; PA14\_59780 *rcsC*—sensor with *rcsB*, biofilm formation. †The phenotype for these mutants was different between PA14 and PAO1 as described in the text.



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