

PhoQ allele	Fold activation	PhoQ allele	Fold activation
R219C	2.27	R236S	15.63
P220H	3.16	L239F	7.09
I221N	1.29	L239P	5.60
R226W	3.02	P241L	1.84
E227V	2.74	P241T	1.46
E227K	15.10	E242K	1.37
V228I	3.64	T244K	9.68
R229L	4.97	R245C	2.35
E230A	13.29	R245L	9.63
E230G	19.33	E246D	2.02
E230K	12.48	L247Q	4.35
L231F	5.27	L247P	7.74
L231R	3.24	L250H	2.09
L231I	12.12	R252C	2.08
L231V	9.05	L254I	8.79
L231P	11.63	L254F	26.59
E232G	17.52	L254V	3.73
E232K	36.16	N255T	21.78
E232V	16.41	N255I	24.46
D233G	1.14	L257Q	0.84
D233Y	2.26	L258F	6.39
D233V	3.99	E261K	13.28
H234Q	5.69	Y265C	14.78
R236H	19.33	N266S	4.68

Table S1. Fold activation of the different PhoQ alleles when compared to wild-type grown under PhoQ repressive conditions. The mean activity of alkaline phosphatase was determined for each allele in at least three independent experiments and then compared to the mean wild-type activity. All strains were grown in N minimal media in the presence of 10 mM MgCl₂. Amino acid substitutions resulting in 10 fold increase in activity above the wild-type are shown over a grey background.