Supplemental Data

Crystal Structure of a Functional Dimer of the PhoQ Sensor Domain J. Cheung, C. A. Bingman, M. Reyngold, W. A. Hendrickson and Carey D. Waldburger

Constructions and Cloning for Mutant Variants

Variants of PhoQ in which amino acids at residues 50, 54, or 179 were substituted were made using pLPQ3 (HindIII-NcoI) and pLPQ3 λ as follows. pLPQ3 (HindIII-NcoI) [1] is a pBR322-derived plasmid in which the phoP-phoQ operon is driven by the lacUV5 promoter and HindIII and NcoI restriction enzyme sites have been introduced at codons 42-44 and 189-191 of phoQ, respectively. The mutations that create the HindIII site are silent and the mutations that create the NcoI site result in a serine in place of the wild type tyrosine at codon 189. pLPQ3 λ (1) is a derivative of pLPQ3 (HindIII-NcoI) in which a ~1.2 kb HindIII-NcoI stuffer fragment corresponding to bp 23901-25157 of λ has been cloned into the HindIII-NcoI backbone of pLPQ3 (HindIII-NcoI). The following primers were used for the mutagenesis:

P1: 5'-GATCGGTTATAGCGTAAGCTTCGATAAAACTACGTTTCGGC-3'

P2: 5'-CCAGCTCCAGAACATGTAGGAACTTTTTAGCTCCACCGG-3'

P3: 5'-GATCGGTTATAGCGTAAGCTTCGATAAAACTACGTTTCGGCTGTTACGTGACGAGAGC-3'

P4: 5'-GATCGGTTATAGCGTAAGCTTCGATAAAACTACGTTTGACCTGTTACGTGGC-3'

P5: 5'-CCAGCTCCAGAACATGTAGGAACTTTTTAGCTCCACCGGAATGGTGCGCACCACC-3'

Restriction enzyme recognition sites are italicized in each primer (*Hind*III in P1, P3, and P4; *Afl*III in P2 and P4). Mutagenized codons are underlined. PCR products were generated using pLPQ3 (*Hind*III-*Nco*I) as a template and the following primer pairs for each listed mutation:

G54D	P3/P2
R50D	P4/P2
D179R	P1/P5
R50D/D179R	P4/P5

The resulting DNA fragments were then digested with HindIII and AfIIII and cloned into the HindIII-NcoI backbone of pLPQ3 λ . Note that this procedure results in restoration of the wild type tyrosine codon in the phoQ gene at position 189. All mutations were confirmed by double stranded DNA sequencing of the resulting plasmid. The mutations were then moved into a lower copy number vector by sub-cloning the EcoRI-SaII $P_{1ac}UV5$ -phoP-phoQ fragment (2) into the EcoRI-SaII backbone of pGB2 (3).

References

- 1. Regelmann, A. G, Lesley, J. A., Mott, C., Stokes, L., and Waldburger, C. D. (2002) *J. Bacteriol.* **184**, 5468-5478.
- 2. Waldburger, C. D., and Sauer, R. T. (1996) J. Biol. Chem. 271, 26630-26636.
- 3. Churchward, G., Belin, D., and Nagamine, Y. (1984) Gene 31, 165-171.