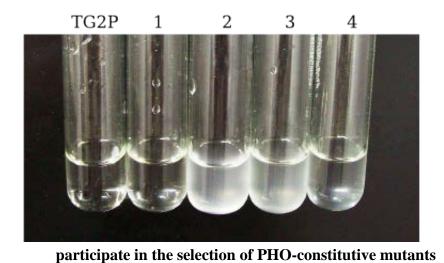
1. MG1655 pitA::Km 2. MG1655 pitA::Km pst81

4. MG1655 phoR129

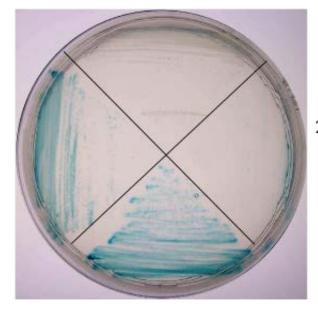
3. MG1655 *pitA*::Km *phoR*217



Ugp and PitA

Figure S1. The following derivatives of MG1655 were grown in TG2P: 1. *pitA*::Km (strain EG2); 2. *pitA*::Km pst81 (strain RI81); 3. *pitA*::Km *phoR*217 (strain RI65) and 4. *phoR*129 (strain TP01).

1. MG1655 pitA::Km



4. MG1655

phoR129

2. MG1655 pitA::Km pst81

3. MG1655 pitA::Km phoR217

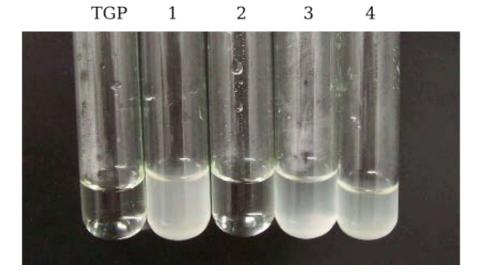


Figure S2. The following derivatives of MG1655 were grown in TGP: 1. *pitA*::Km (strain EG2); 2. *pitA*::Km *pst*81 (strain RI81); 3. *pitA*::Km *phoR*217 (strain RI65) and 4. *phoR*129 (strain TP01).

Figure S3.

CLUSTAL 2.1 sequence alignment between PhoR and EnvZ. The conserved residue Thr247 (EnvZ)/Thr217 (PhoR) is surrounded by a red square.