

**Table S2. Genes of the VreI (PA0675) regulon upregulated in wild-type PAO1 exposed to epithelial cells versus grown in TSB (adapted from reference [21])**

Gene*	Description*	Fold change†
PA0674/ <i>pigC</i>	hypothetical protein (PigCDE CSS receptor)	99.8
PA0675/ <i>pigD</i>	sigma-70 factor, ECF subfamily	9.0
PA0676/ <i>pigE</i>	transmembrane sensor	9.5
PA0679/ <i>hxcP</i>	hypothetical protein	8.3
PA0680/ <i>hxcV</i>	HxcV putative pseudopilin (Type II secretion system)	24.5
PA0683/ <i>hxcY</i>	probable type II secretion system protein	3.9
PA0688	low molecular weight alkaline phosphatase	3.8
PA0691	hypothetical protein (similar to transposase)	11.9
PA0692	outer membrane transporter of TPS pathway	8.9
PA0693/ <i>exbB2</i>	transport protein ExbB2	6.9
PA0694/ <i>exbD2</i>	transport protein ExbD	5.3
PA0696	hypothetical protein	60.0
PA0697	hypothetical protein	8.5
PA0698	hypothetical protein	13.1
PA0699	probable peptidyl-prolyl cis-trans isomerase, PpiC-type	6.3
PA2384	Probable Fur, Fe <sup>2+</sup> /Zn <sup>2+</sup> uptake regulation protein	27.7

\*PA gene number, gene name and description are according to *Pseudomonas* Genome Project website (<http://www.pseudomonas.com>).

†Fold change in gene expression of wildtype PAO1 exposed to epithelial cells versus grown in TSB.