

**Supplementary Table 1.** Genes potentially involved in the mechanics of regulation by the Gac/Rsmnetwork present in the *Pseudomonas chlororaphis* O6

<b>Gene name</b>	<b>Locus PchlO6_</b>	<b>Potential function</b>
<i>gacS</i>	4573	Membrane sensor
<i>gacA</i> *	3799	Regulator
<i>rpeA</i> *	3453	Membrane sensor- negative regulator
<i>rpeB</i> *	3454	RpeB-P positive regulator
<i>rgsA</i> *	1906	Small regulatory RNA
<i>rsmX</i> *	4246	Small regulatory RNA, competes for RsmA/E
<i>rsmY</i> *	1201	Small regulatory RNA, competes for RsmA/E
<i>rsmZ</i> *	5703	Small regulatory RNA, competes for RsmA/E
<i>rsmA</i> *	4625	Translational repressor
<i>rsmE</i> *	2238	Translational repressor
<i>pip</i> *	5422	Regulator for <i>phzI</i> and <i>phzR</i>
<i>vfr</i> *	5645	Regulator
<i>phsR</i> *	1386	Regulator for Vfr-traits
<i>psrA</i>	2065	Regulator for <i>rpoS</i>
<i>ptrA</i>	1367	Master LysR transcriptional regulator
<i>Alternative sigma factors</i>		
<i>rpoS</i>	1200	Alternative sigma factor undergac control
<i>algU</i> *	1485	Alternative sigma factor undergac control