

Table S1: Results of qRT-PCR. Differential gene expression was determined using REST software (1). *adk* and *gatB* (2) were used as reference genes (REF). With this method, the results of the RNA sequencing experiments can be confirmed or refuted on a “yes-or-no” level (UP or DOWN or - not regulated). The differential expression values do not necessarily correlate, as there are fundamental differences between the two methods, regarding e.g. primer design for qRT-PCR or different analysis software (3). TRG = target. C.I. = confidence interval. P(H1) = significance. REST sets maximum primer efficiency = 1.

Gene	Type	Efficiency	Expression	Std. Error	95% C.I.	P(H1)	Result
<i>adk</i>	REF	0.928	1.035				
<i>gatB</i>	REF	0.943	0.966				
bcf_04565	TRG	0.941	26.854	20.779 - 39.002	18.149 - 45.283	0.011	UP
bcf_04560	TRG	0.989	6.729	5.189 - 8.544	4.698 - 11.681	0.01	UP
bcf_27370	TRG	0.868	32.601	28.173 - 38.185	24.969 - 39.559	0.02	UP
bcf_09850	TRG	0.974	9.407	7.697 - 11.016	7.175 - 12.229	0.006	UP
bcf_16240	TRG	0.985	3.619	3.057 - 4.100	2.864 - 4.287	0.006	UP
bcf_08380	TRG	0.969	5.091	4.208 - 5.977	4.116 - 6.089	0.001	UP
bcf_13835	TRG	1	3.605	2.534 - 5.347	1.949 - 6.600	0	UP
bcf_05660	TRG	0.949	2.942	2.172 - 4.706	1.956 - 5.338	0.016	UP
bcf_17385	TRG	0.998	3.313	2.928 - 3.662	2.741 - 4.193	0.006	UP
bcf_17575	TRG	1	1.24	0.921 - 1.669	0.861 - 1.834	0.272	-
bcf_01910	TRG	1	5.766	4.376 - 7.379	4.153 - 9.144	0.024	UP
bcf_24405	TRG	0.91	3.653	2.415 - 4.596	2.180 - 5.146	0.006	UP
bcf_24230	TRG	0.982	1.872	1.457 - 2.384	1.201 - 2.885	0.011	UP
bcf_17530	TRG	0.76	1.399	1.198 - 1.623	1.063 - 1.752	0.015	UP
bcf_18185	TRG	0.891	0.907	0.523 - 1.398	0.404 - 2.098	0.804	-
bcf_17340	TRG	1	2.317	2.040 - 2.638	1.934 - 2.774	0.005	UP
bcf_15285	TRG	0.883	0.807	0.603 - 1.055	0.523 - 1.212	0.235	-
bcf_00530	TRG	0.925	3.021	2.429 - 3.748	2.214 - 4.081	0.017	UP
bcf_09265	TRG	1	1.29	1.026 - 1.547	1.003 - 1.578	0.037	UP
bcf_08015	TRG	0.809	0.68	0.539 - 0.946	0.514 - 0.994	0.051	-
bcf_03465	TRG	1	0.025	0.016 - 0.039	0.013 - 0.048	0.013	DOWN
bcf_04545	TRG	0.987	0.345	0.292 - 0.410	0.269 - 0.429	0.029	DOWN
bcf_16685	TRG	0.988	0.359	0.292 - 0.452	0.273 - 0.466	0.017	DOWN
bcf_15280	TRG	0.939	0.368	0.236 - 0.573	0.214 - 0.633	0.01	DOWN
bcf_13360	TRG	1	0.089	0.034 - 0.228	0.028 - 0.280	0.021	DOWN
bcf_18650	TRG	0.985	0.256	0.223 - 0.323	0.205 - 0.354	0.017	DOWN

1. Pfaffl MW, Horgan GW, Dempfle L. 2002. Relative expression software tool (REST) for group-wise comparison and statistical analysis of relative expression results in real-time PCR. *Nucleic Acids Res* 30:e36.
2. Reiter L, Kolsto AB, Piehler AP. 2011. Reference genes for quantitative, reverse-transcription PCR in *Bacillus cereus* group strains throughout the bacterial life cycle. *J Microbiol Methods* 86:210-7.
3. Dallas PB, Gottardo NG, Firth MJ, Beesley AH, Hoffmann K, Terry PA, Freitas JR, Boag JM, Cummings AJ, Kees UR. 2005. Gene expression levels assessed by oligonucleotide microarray analysis and quantitative real-time RT-PCR -- how well do they correlate? *BMC Genomics* 6:59.