Table 1: **Primers employed for qRT-PCR.** Three genes were used for evaluation, and one for normalization. T_M: Melting Temperature. GC%: Percent of G+C content

Type	**GenID**	**Amplicon length (pb)** [1]	**Direction**	**T_M**	**GC%**	**Sequence (5'->3')**
Evaluated	QR90_RS11755[1]	202	Forward	62.0	52.6	GCTGGACGGTGAGATTGTT
			Reverse	62.0	50.0	TTCCTCACGCACAATGTAGG
	QR90_RS09640[2]	209	Forward	62.1	50	TGATTCACGGCGAGAGATTG
			Reverse	62.2	55	CAGTTCGGGCAGTTCCTTAG
	QR90_RS11750[3]	223	Forward	62	52.4	GATCAGACCTTGGAGCAGTTC
			Reverse	62	55	CGTAGACCAGTTTGCGGTAG
Normalizer	QR90_RS09970[4]	210	Forward	62	40.9	AACGTTTATTGACGCGAAACAG
			Reverse	62	50	AGGATCAGCCAGTCGTAGAA

[1] GntR family transcriptional regulator.
[2] RNA helicase.
[3] Proline dehydrogenase.
[4] Succinate dehydrogenase.