Table S2. Nucleotide sequence, annealing temperature and amplification efficiency for primer pairs used during quantitative reverse transcription polymerase chain reaction (RT-qPCR).

Gene symbol	Accession nr (NCBI)[1]	Primer sequence (5′-3′)	Amplicon size (base pairs)	Annealing temp (°C)	RT-qPCR efficiency (%)
18S[2]	AH001810.2	F: GTGACGGGTGACGGAGAATT R: GACACTAATGCGCCCGGTAT	151	60	109
Pst β-TUB[3]	EF570842.1	F: CTCGGACGAAACCTTCTGTATC R: CGTAGGTAGGTGTAGCCAATTT	83	60	98.5

[1]NCBI = National Centre for Biotechnology Information [48].
[2]*18S = 18S ribosomal RNA* [49].
[3]*Pst β-TUB = Puccinia striiformis* f. sp. *tritici β-tubulin*.

References

48. National Centre for Biotechnology Information. Available online: http://www.ncbi.nlm.nih.gov/ (accessed on 7 December 2018).
49. Jarošová, J.; Kundu, J.K. Validation of reference genes as internal control for studying viral infections in cereals by quantitative real-time RT-PCR. BMC *Plant Biol.* **2010**, *10*, 146, doi:10.1186/1471-2229-10-146.