

RT-qPCR Checklist Experimental Design

Check off items once complete or ready



	<i>Description of the step (builds your materials & methods section)</i>	<i>Check ✓</i>
Defined experimental and control group		<input type="checkbox"/>
Number within each group		<input type="checkbox"/>

RT-qPCR Checklist Sample

Check off items once complete or ready

	<i>Description of the step (builds your materials & methods section)</i>	<i>Check ✓</i>
Sample descriptions provided		<input type="checkbox"/>
Sample macro or micro dissection		<input type="checkbox"/>
Processing procedure		<input type="checkbox"/>
Check if frozen. If so, how was it frozen and how quickly?		<input type="checkbox"/>
Check if fixed. If so, with what and how quickly?		<input type="checkbox"/>

RT-qPCR Checklist

Nucleic Acid Extraction

Check off items once complete or ready



	<i>Description of the step (builds your materials & methods section)</i>	<i>Check ✓</i>
Procedure and/or instrumentation ready & described		<input type="checkbox"/>
Name of kit and details of any modifications		<input type="checkbox"/>
Contamination assessment (DNA or RNA)		<input type="checkbox"/>
Nucleic acid quantification completed		<input type="checkbox"/>
Instrument and method described		<input type="checkbox"/>
RIN/RQI or Co of 3' and 5' transcripts		<input type="checkbox"/>
Inhibition testing (Co dilutions, spike or other) completed		<input type="checkbox"/>

RT-qPCR Checklist

Reverse Transcription

Check off items once complete or ready



	<i>Description of the step (builds your materials & methods section)</i>	<i>Check ✓</i>
Complete reaction conditions		<input type="checkbox"/>
Amount of RNA and reaction volume defined		<input type="checkbox"/>
Priming oligonucleotide (if using GPS) and concentration		<input type="checkbox"/>
Reverse transcriptase and concentration		<input type="checkbox"/>
Temperature and time		<input type="checkbox"/>

RT-qPCR Checklist

qPCR Target Information

Check off items once complete or ready



	<i>Description of the step (builds your materials & methods section)</i>	<i>Check ✓</i>
If multiplex, efficiency and LOD of each assay		<input type="checkbox"/>
Sequence accession number		<input type="checkbox"/>
Amplicon length		<input type="checkbox"/>
In silico specificity screen (BLAST, etc)		<input type="checkbox"/>
Sequence alignment		<input type="checkbox"/>
Location of each primer by exon or intron (if applicable)		<input type="checkbox"/>
Targeted splice variants noted		<input type="checkbox"/>

RT-qPCR Checklist

Oligonucleotides

Check off items once complete or ready



	<i>Description of the step (builds your materials & methods section)</i>	<i>Check ✓</i>
Primer sequences		<input type="checkbox"/>
Location and identity of any modifications		<input type="checkbox"/>

RT-qPCR Checklist

qPCR Protocol

Check off items once complete or ready



	<i>Description of the step (builds your materials & methods section)</i>	<i>Check ✓</i>
Complete reaction conditions		<input type="checkbox"/>
Reaction volume and amount of cDNA/DNA		<input type="checkbox"/>
Primer, (probe), Mg ²⁺ and dNTP concentrations		<input type="checkbox"/>
Polymerase identity and concentration		<input type="checkbox"/>
Buffer/kit identity and manufacturer		<input type="checkbox"/>
Additives (SYBR Green I, DMSO, etc.)		<input type="checkbox"/>
Complete thermocycling parameters		<input type="checkbox"/>
Manufacturer of qPCR instrument		<input type="checkbox"/>

RT-qPCR Checklist

qPCR Validation

Check off items once complete or ready



	<i>Description of the step (builds your materials & methods section)</i>	<i>Check ✓</i>
	Specificity (gel, sequence, melt, or digest)	<input type="checkbox"/>
	For SYBR Green I, Co of the NTC	<input type="checkbox"/>
	Standard curves with slope and y-intercept	<input type="checkbox"/>
	PCR efficiency calculated from slope	<input type="checkbox"/>
	R2 of standard curve	<input type="checkbox"/>
	Linear dynamic range	<input type="checkbox"/>
	Cq variation at lower limit	<input type="checkbox"/>
	Evidence for limit of detection	<input type="checkbox"/>
	If multiplex, efficiency and LOD of each assay	<input type="checkbox"/>

RT-qPCR Checklist

Data Analysis



Check off items once complete or ready

	<i>Description of the step (builds your materials & methods section)</i>	<i>Check ✓</i>
qPCR analysis program (source, version)		<input type="checkbox"/>
Cq method determination		<input type="checkbox"/>
Outlier identification and disposition		<input type="checkbox"/>
Results of NTCs (No template controls)		<input type="checkbox"/>
Justification of number and choice of reference genes		<input type="checkbox"/>
Number and stage (RT or qPCR) of technical replicates		<input type="checkbox"/>
Repeatability (intra-assay variation)		<input type="checkbox"/>
Statistical methods for result significance		<input type="checkbox"/>
Software (source, version)		<input type="checkbox"/>

References

Bustin, S. A., Benes, V., Garson, J. A., Hellems, J., Huggett, J., Kubista, M., Mueller, R., Nolan, T., Pfaffl, M. W., Shipley, G. L., Vandesompele, J., & Wittwer, C. T. (2009). The MIQE guidelines: Minimum information for publication of quantitative real-time PCR experiments. *Clinical Chemistry*, 55(4), 611–622. <https://doi.org/10.1373/clinchem.2008.112797>

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Oswald, N. (2016). 10 Tips for Consistent Real-Time PCR. *BiteSizeBio*.

S.A. Deepak, K.R. Kottapalli, R. Rakwal, G. Oros, K.S. Rangappa, H. Iwahashi, Y. Masuo, & G.K. Agrawal. (2007). Real-Time PCR: Revolutionizing Detection and Expression Analysis of Genes. *Current Genomics*, 8(4), 234–251. <https://doi.org/10.2174/138920207781386960>

Sundquist, T., & Saldanha, G. (2012). 10 Tips to Improve Your qPCR and RT-qPCR Results (pp. 15–17). <http://www.genengnews.com/gen-articles/10-tips-to-improve-your-qpcr-and-rt-qpcr-results/4594>