

© Quant-iT™ RiboGreen™ RNA Quantification V.3

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1 Works for me

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ABSTRACT

The following protocol is intended for the quantification of RNA using Quant-iT™ RiboGreen™ RNA Assay Kit (ThermoFisher). This protocol is a simplified and condensed version of the <u>full protocol</u> from the manufacturer. The procedure described here is for 96 reactions. If samples are run in duplicates, then this should allow quantifying 40 samples.

EXTERNAL LINK

https://sfamjournals.onlinelibrary.wiley.com/doi/abs/10.1111/1462-2920.12140

THIS PROTOCOL ACCOMPANIES THE FOLLOWING PUBLICATION

Angel, R., and Conrad, R. (2013). Elucidating the microbial resuscitation cascade in biological soil crusts following a simulated rain event. Environ Microbiol 15, 2799–2815. doi:10.1111/1462-2920.12140.

ATTACHMENTS

mp11490.pdf

MATERIALS **NAME**

Quant-iT™ RiboGreen™ RNA Assay Kit	R11490	Invitrogen - Thermo Fisher
STEPS MATERIALS		
NAME	CATALOG #	VENDOR
Quant-iT™ RiboGreen® RNA Assay Kit	R11490	Thermo Scientific
EQUIPMENT		

CATALOG #

VENDOR

NAME	CATALOG #	VENDOR
Infinite M Nano	TEC006436I	
Multipette® E3	4987000010	
96-Well microtiter plate	165305	

SAFETY WARNINGS

Quant-iT™ RiboGreen® RNA reagent is classified as Not Hazardous. Nevertheless, the user should always consult the MSDS accompanying any of the reagents and apparatus described in this protocol.

BEFORE STARTING

- This protocol is optimised for measuring an entire 96-well plate. It assumes that 10 wells will be used for measuring the standards and the blank samples (in duplicates) and 86 wells will be used for measuring unknown RNA samples (typically in duplicates).
- The protocol can be easily adjusted for a lower number of samples by reducing the volume of the working solutions of the reagents. Note though that enough TE should be retained for diluting the standard stock solution (490 or 680 μl), for potentially diluting the unknown samples, if their concentration is too high, and for

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accounting for pipetting errors. To fill the plate, 19.2 ml of TE is needed. So if only 40 wells are to be used for measuring unknown samples prepare about $\,50/96 imes 22 pprox 11.5\,$ ml of TE buffer.

- 3. The dynamic range of the assay is between 1 ng ml $^{-1}$ to 50 ng ml $^{-1}$ in the "low-range" version of the assay and 20 ng ml $^{-1}$ to 1 μ g ml $^{-1}$ in the "high-range" version of the assay. This translates into RNA sample concentrations of 0.2-10 ng μ l $^{-1}$ and 4-200 ng μ l $^{-1}$ in the low-range and high-range assays, respectively. Samples with higher RNA concentration need to be diluted (e.g. in RNase-free water or TE buffer).
- 4. Note that some compounds that often contaminate RNA are known to interfere with the measurement and produce a lower observed measurement. Please refer to the full protocol for a list of these compounds and their effect on the measurement.
- 5. Quant-iT™ RiboGreen® reagent also binds to DNA. Samples containing DNA should be pre-treated with an RNase-free DNase before using this protocol to ensure that the fluorescent signal is solely due to binding with RNA.

Prepare the reaction

Take out all reagents from the fridge and bring them to room temperature.
Take out the RNA samples from the freezer. RNA samples should be slowly thawed on ice.

20m

- Quant-iT™ RiboGreen® RNA reagent is dissolved in dimethylsulfoxide (DMSO), which freezes below 19 °C. The reagent must be completely thawed before using it by bringing it to room temperature. After the reagent thawed, it is advisable to briefly vortex the tube to make sure it is adequately mixed and to spin it down in a centrifuge or a mini centrifuge.
- Quant-iT™ RiboGreen® RNA reagent is light sensitive and should be protected from light at all times.
- Quant-iT™ RiboGreen® RNA Assay Kit
 by Thermo Scientific
 Catalog #: R11490
- 2 Prepare 22 ml 1X TE buffer by pipetting 1.1 ml of 20X TE buffer into 20.9 ml of nuclease-free water into a sterile and nuclease-free 50 ml tube.

Mix by inverting the tube several times.

- ■1.1 ml 20X TE buffer
- ■20.9 ml nuclease-free water
- 3 For high-range quantification:

2m

Dilute the *E.coli* rRNA-standard stock solution (100 μ g ml⁻¹) to a final concentration of 2 ng μ l⁻¹ by mixing 10 μ l of rRNA-standard stock solution with 490 μ l 1X TE buffer.

- ■10 µl rRNA-standard stock solution
- ■490 µl 1X TE buffer

For low-range quantification:

Prepare a 20-fold dilution of the 2 ng μ l⁻¹ rRNA-standard work solution by mixing 10 μ l of the 2 ng μ l⁻¹ rRNA-standard work solution with 190 μ l 1X TE buffer to yield 0.1 ng μ l⁻¹ rRNA-standard work solution.

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■10 µl diluted rRNA-standard solution

■190 µl 1X TE buffer

4 If needed, prepare a dilution of each sample in 1X TE so that the reading will be within the dynamic range.



It is advisable to run samples in duplicates for a more accurate quantification

5 Prepare the RiboGreen® work solution:

2m

For the **high-range assay** pipette 50 μ l Quant-iT^M RiboGreen® RNA Reagent and 9950 μ l of 1X TE into a sterile and nucleic-acids free 50 ml tube.

For the **low-range** assay pipette 5 μ l Quant-iTTM RiboGreen® RNA Reagent and 9995 μ l of 1X TE into a sterile and nucleic-acids free 50 ml tube.

Mix by inverting and keep the solution away from light.

6 Prepare one of the following standard mixtures in the first two columns of a black, sterile, 96-well plate:

10m

Assay version	Diluted RNA std. (µI)	1X TE (μl)	Final
			RNA
			amount
			(ng)
High-range (4-200 ng µl-1)	100	0	200
Use 2 ng μl-1 standard	50	50	100
	10	90	20
	2	98	4
	0	100	0
Low-range (200 pg µl - 10 ng µl-1)	100	0	10
Use 0.1 ng μl-1 standard	50	50	5
	10	90	1
	2	98	0.2
	0	100	0



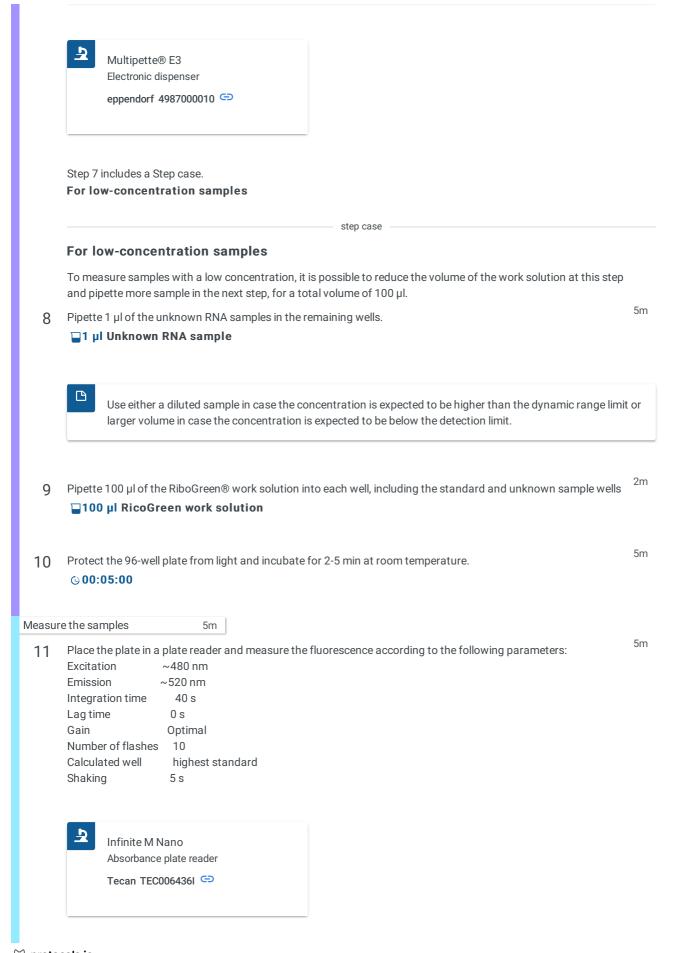
7 Pipette 99 μl of 1X TE buffer in the remaining wells.

2m

■99 µl 1X TE buffer



Tip: use a mechanical or electronic dispenser during this step and step no. 9 to speed up the work.





It is also possible to set the gain to a fixed value (e.g. 100). If the fluorescence values of the standard drop over time this could indicate damage to the reagents or the RNA standard.

Plot the measured fluorescent values of the standard samples against their known concentrations and fit a linear curve using linear regression. Make sure that the coefficient of determination (R^2) is close to 1 (typically >0.98). Calculate the RNA concentrations in the unknown samples using the slope and intercept parameters of the linear equation. Output values you obtained are in ng μ l⁻¹, assuming 1 μ l of each sample was used.



Do not forget to account for any dilutions when calculating the concentration of the RNA in the unknown samples