Package 'boggy'

July 22, 2025
Title Real-Time PCR Data Sets by Boggy et al. (2010)
Version 0.0.1
Description Real-time quantitative polymerase chain reaction (qPCR) data sets by Boggy et al. (2008) <doi:10.1371 journal.pone.0012355="">. This package provides a dilution series for one PCR target: a random sequence that minimizes secondary structure and off-target primer binding. The data set is a six-point, ten-fold dilution series. For each concentration there are two replicates. Each amplification curve is 40 cycles long. Original raw data file: https://journals.plos.org/plosone/article/file?type=supplementary&id=10.1371/journal.pone.0012355.s004>.</doi:10.1371>
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Encoding UTF-8
RoxygenNote 7.3.1
Imports tibble
Depends R (>= 2.10)
LazyData true
<pre>URL https://rmagno.eu/boggy/, https://github.com/ramiromagno/boggy</pre>
<pre>BugReports https://github.com/ramiromagno/boggy/issues</pre>
NeedsCompilation no
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qPCR data set by Boggy et al. (2010)

s1

Description

The qPCR data set named s1 comprises a six-point, ten-fold dilution series, replicated in duplicates targeting an amplicon designed with a random sequence that minimizes secondary structure and off-target primer binding. Please read the *Quantitative PCR data* section, qPCR assays subsection of Boggy et al. (2010) for more details.

Format

```
A tibble with 480 rows and 10 variables:
```

plate Plate identifier. For this data set it is always missing (NA).

well Well identifier.

dye The type of dye used. In this data set the values are always "SYT0 13", meaning 2 mM of the dsDNA dye SYTO-13 (Invitrogen, Carlsbad, CA).

target Target identifier: the amplicon used, a random sequence dubbed "S1" by the original authors.

sample_type Sample type (all curves are standards, i.e. "std").

replicate Replicate identifier: either 1 or 2.

copies Standard copy number.

dilution Dilution factor. Higher number means greater dilution.

cycle PCR cycle.

fluor Raw fluorescence values.

Source

doi:10.1371/journal.pone.0012355

Examples

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