Package 'qgshiny'

October 13, 2022

Interest A shiny Application for Active Learning Instruction in Introductory Quantitative Genetics	
Version 0.1.0	
Description A 'shiny' application for teaching introductory quantitative genetics and plant breeding through interactive simulations. The application relies on established plant breeding and quantitative genetic theory found in Falconer and Mackay (1996, ISBN:0582243025) and Bernardo (2010, ISBN:978-0972072427).	
License MIT + file LICENSE	
Encoding UTF-8	
LazyData true	
RoxygenNote 7.1.0	
Depends R (>= 3.5.0)	
Imports stats, shiny, rmarkdown	
NeedsCompilation no	
Author Jeffrey Neyhart [aut, cre] (https://orcid.org/0000-0002-1991-5310)	
Maintainer Jeffrey Neyhart <neyhartje@gmail.com></neyhartje@gmail.com>	
Repository CRAN	
Date/Publication 2020-07-02 11:20:02 UTC	
R topics documented:	
run_qgshiny	2
Index	3

run_qgshiny

run_qgshiny

Run the application

Description

Runs the 'shiny' application interactively.

Usage

```
run_qgshiny(display.mode = c("normal", "showcase"))
```

Arguments

display.mode

The mode in which to display the application. If "showcase", application code is displayed along with the output. If "normal" (default), the code is hidden.

Examples

```
if (interactive()) {
  run_qgshiny()
}
```

Index

run_qgshiny, 2