Package 'greencrab.toolkit'

September 12, 2023

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Title Run 'Stan' Models to Interpret Green Crab Monitoring Assessments
Version 0.2
Maintainer Abigail G. Keller <agkeller@berkeley.edu></agkeller@berkeley.edu>
Description These Bayesian models written in the 'Stan' probabilistic language can be used to interpret green crab trapping and environmental DNA monitoring data, either independently or jointly. Detailed model information is found in Keller (2022) <doi:10.1002 eap.2561="">.</doi:10.1002>
License GPL-3
Encoding UTF-8
RoxygenNote 7.2.3
Biarch true
Depends R (>= 3.4.0)
Imports Rcpp (>= 0.12.0), RcppParallel (>= 5.0.1), rstan (>= 2.26.23), rstantools (>= 2.2.0), methods
LinkingTo BH (>= 1.66.0), Rcpp (>= 0.12.0), RcppEigen (>= 0.3.3.3.0), RcppParallel (>= 5.0.1), rstan (>= 2.26.23), StanHeaders (>= 2.26.22)
SystemRequirements GNU make
NeedsCompilation yes
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Repository CRAN
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R topics documented:
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greencrab.toolkit-package

The 'greencrab.toolkit' package.

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References

Stan Development Team (2023). RStan: the R interface to Stan. R package version 2.21.8. https://mc-stan.org

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