

# Package ‘banter’

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**Type** Package

**Title** BioAcoustic EveNT ClassifiER

**Description** Create a hierarchical acoustic event species classifier out of multiple call type detectors as described in Rankin et al (2017) <doi:10.1111/mms.12381>.

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**Depends** magrittr

**Imports** dplyr, ggplot2, gridExtra, methods, parallel, plyr, randomForest, ranger, rfPermute, rlang, stats, tibble, tidyR

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<b>banter-package</b>	<i>BioAcoustic EveNT ClassifiER</i>
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## Description

**banter**

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<b>addBanterDetector</b>	<i>Add a BANTER Detector Model</i>
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## Description

Add a detector model to a BANTER classifier.

## Usage

```
addBanterDetector(x, data, name, ntree, sampsize = 1, importance = FALSE,
                  num.cores = NULL)

removeBanterDetector(x, name)
```

## Arguments

x	a <b>banter_model</b> object.
data	detector data.frame or named list of detector data.frames. If a data.frame, then name must be provided.
name	detector name.
ntree	number of trees.
sampsize	number or fraction of samples to use in each tree.
importance	retain importance scores in model? Defaults to FALSE and will be ignored if num.cores > 1.
num.cores	number of cores to use for Random Forest model.

## Value

a **banter\_model** object with the detector model added or removed.

## Author(s)

Eric Archer <eric.archer@noaa.gov>

## Examples

```
data(train.data)
# initialize BANTER model with event data
bant.mdl <- initBanterModel(train.data$events)
# add the 'bp' (burst pulse) detector model
bant.mdl <- addBanterDetector(
  x = bant.mdl,
  data = train.data$detectors$bp,
  name = "bp",
  ntree = 50, sampsize = 1, num.cores = 1
)
bant.mdl

# remove the 'bp' detector model
bant.mdl <- removeBanterDetector(bant.mdl, "bp")
bant.mdl
```

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getBanterModel	<i>Extract Random Forest Model</i>
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## Description

Extract BANTER event or detector Random Forest model.

## Usage

```
getBanterModel(x, model = "event")
```

## Arguments

- |       |  |
|-------|--|
| x     | a <a href="#">banter_model</a> object.   |
| model | name of model to extract. Default is "event" to summarize the event-level model. Can also be name of a detector. |

## Value

a [randomForest](#) model object.

## Author(s)

Eric Archer <eric.archer@noaa.gov>

## Examples

```

data(train.data)
# initialize BANTER model with event data
bant.mdl <- initBanterModel(train.data$events)
# add all detector models
bant.mdl <- addBanterDetector(
  bant.mdl, train.data$detectors,
  ntree = 50, sampsize = 1, num.cores = 1
)
# run BANTER event model
bant.mdl <- runBanterModel(bant.mdl, ntree = 1000, sampsize = 1)

# extract the event randomForest model
library(randomForest)
event.rf <- getBanterModel(bant.mdl)
event.rf

# extract the burst pulse (bp) detector model
bp.rf <- getBanterModel(bant.mdl, "bp")
bp.rf

```

`getBanterModelData`      *Extract Random Forest Model Data*

## Description

Extract BANTER event data used for the Random Forest model.

## Usage

```
getBanterModelData(x)
```

## Arguments

`x`                  a `banter_model` object.

## Value

a `randomForest` model object.

## Author(s)

Eric Archer <eric.archer@noaa.gov>

**Examples**

```
data(train.data)
# initialize BANTER model with event data
bant.mdl <- initBanterModel(train.data$events)
# add all detector models
bant.mdl <- addBanterDetector(
  bant.mdl, train.data$detectors,
  ntree = 50, sampsize = 1, num.cores = 1
)
# run BANTER event model
bant.mdl <- runBanterModel(bant.mdl, ntree = 1000, sampsize = 1)

event.df <- getBanterModelData(bant.mdl)
head(event.df)
```

---

getDetectorNames      *Detector Names*

---

**Description**

Return names of detectors loaded in BANTER model.

**Usage**

```
getDetectorNames(x)
```

**Arguments**

x                    a [banter\\_model](#) object.

**Value**

a vector of names.

**Author(s)**

Eric Archer <[eric.archer@noaa.gov](mailto:eric.archer@noaa.gov)>

**Examples**

```
data(train.data)
# initialize BANTER model with event data
bant.mdl <- initBanterModel(train.data$events)
# add all detector models
bant.mdl <- addBanterDetector(
  bant.mdl, train.data$detectors,
  ntree = 50, sampsize = 1, num.cores = 1
)
```

```
getDetectorNames(bant.mdl)
```

getSampSize	<i>Sample Size</i>
-------------	--------------------

## Description

Return sample size used for a BANTER model.

## Usage

```
getSampSize(x, model = "event")
```

## Arguments

x	a <a href="#">banter_model</a> object.
model	name of model to extract. Default is "event" to summarize the event-level model. Can also be name of a detector.

## Value

a vector of sample sizes.

## Author(s)

Eric Archer <eric.archer@noaa.gov>

## Examples

```
data(train.data)
# initialize BANTER model with event data
bant.mdl <- initBanterModel(train.data$events)
# add all detector models
bant.mdl <- addBanterDetector(
  bant.mdl, train.data$detectors,
  ntree = 50, sampsize = 2, num.cores = 1
)
# run BANTER event model
bant.mdl <- runBanterModel(bant.mdl, ntree = 1000, sampsize = 1)

# sample size for the event model
getSampSize(bant.mdl)

# sample size for the burst pulse (bp) detector model
getSampSize(bant.mdl, "bp")
```

---

initBanterModel	<i>Initialize BANTER model</i>
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---

## Description

Initialize a BANTER model with event data.

## Usage

```
initBanterModel(x)
```

## Arguments

- x a data.frame of events. Every row is a unique event. Must have columns named event.id and species. All other columns will be used as predictor variables for the BANTER event classifier model.

## Value

a `banter_model` object without any detector models.

## Note

Values in the column species are passed through the `make.names` function on creation to ensure they don't include invalid characters.

## Author(s)

Eric Archer <eric.archer@noaa.gov>

## Examples

```
data(train.data)
# initialize BANTER model with event data
bant.mdl <- initBanterModel(train.data$events)
bant.mdl
```

<code>modelPctCorrect</code>	<i>Model Percent Correct</i>
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### Description

Extract percent correctly classified by species for detector and event models.

### Usage

```
modelPctCorrect(x)
```

### Arguments

`x` a `banter_model` object.

### Value

a `data.frame` with the percent correctly classified for each model in `x`.

### Author(s)

Eric Archer <eric.archer@noaa.gov>

### Examples

```
data(train.data)
# initialize BANTER model with event data
bant.mdl <- initBanterModel(train.data$events)
# add all detector models
bant.mdl <- addBanterDetector(
  bant.mdl, train.data$detectors,
  ntree = 50, sampsize = 1, num.cores = 1
)
# run BANTER event model
bant.mdl <- runBanterModel(bant.mdl, ntree = 1000, sampsize = 1)
modelPctCorrect(bant.mdl)
```

<code>numCalls</code>	<i>Number and Proportion of Calls</i>
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### Description

Number and proportion of calls in BANTER detector models

**Usage**

```
numCalls(x, by = c("species", "event"))

propCalls(x, by = c("species", "event"))
```

**Arguments**

x                a `banter_model` object.  
by              return summary by "species" or "event".

**Author(s)**

Eric Archer <eric.archer@noaa.gov>

**Examples**

```
data(train.data)
# initialize BANTER model with event data
bant.mdl <- initBanterModel(train.data$events)
# add all detector models
bant.mdl <- addBanterDetector(
  bant.mdl, train.data$detectors,
  ntree = 50, sampsize = 1, num.cores = 1
)
# run BANTER event model
bant.mdl <- runBanterModel(bant.mdl, ntree = 1000, sampsize = 1)

# number of calls by species and event
numCalls(bant.mdl, "species")
numCalls(bant.mdl, "event")

# proportion of calls by species and event
propCalls(bant.mdl, "species")
propCalls(bant.mdl, "event")
```

---

numEvents

*Number of Events*

---

**Description**

Number of events in BANTER model by species.

**Usage**

```
numEvents(x, model = "event")
```

**Arguments**

- x a `banter_model` object.  
 model name of model to extract. Default is "event" to summarize the event-level model. Can also be name of a detector.

**Value**

a data.frame giving the number of events available for each species.

**Author(s)**

Eric Archer <eric.archer@noaa.gov>

**Examples**

```
data(train.data)
# initialize BANTER model with event data
bant.mdl <- initBanterModel(train.data$events)
# add all detector models
bant.mdl <- addBanterDetector(
  bant.mdl, train.data$detectors,
  ntree = 50, sampsize = 1, num.cores = 1
)
# run BANTER event model
bant.mdl <- runBanterModel(bant.mdl, ntree = 1000, sampsize = 1)

# number of events in event model
numEvents(bant.mdl)

# number of events in burst pulse (bp) detector model
numEvents(bant.mdl, "bp")
```

`plotDetectorTrace`      *Plot BANTER Detector Traces*

**Description**

Plot traces of OOB error rates for detector Random Forest models

**Usage**

```
plotDetectorTrace(x, detector = NULL)
```

**Arguments**

- x a `banter_model` object.  
 detector names of models to plot. If set to NULL, traces for all models will be shown.

**Author(s)**

Eric Archer <eric.archer@noaa.gov>

**See Also**

[plotRFtrace](#)

**Examples**

```
data(train.data)
# initialize BANTER model with event data
bant.mdl <- initBanterModel(train.data$events)
# add all detector models
bant.mdl <- addBanterDetector(
  bant.mdl, train.data$detectors,
  ntree = 50, sampsize = 1, num.cores = 1
)
plotDetectorTrace(bant.mdl)
```

---

predict

*Predict BANTER events*

---

**Description**

Predict species of events for novel data from a BANTER model.

**Usage**

```
## S3 method for class 'banter_model'
predict(object, new.data, ...)
```

**Arguments**

- |          |  |
|----------|--|
| object   | a <a href="#">banter_model</a> object.   |
| new.data | a list of event and detector data that has the same predictors as in the banter_model. It must contain elements called events and detectors. The events element must be a data.frame that has a column called event.id and the same predictor columns as the event data used to initialize the banter model (see <a href="#">initBanterModel</a> ). The detectors element must be a named list with the same detectors used to build the model (see <a href="#">addBanterDetector</a> ). |
| ...      | unused.  |

**Value**

A list with the following elements:

**events** the data frame used in the event model for predictions.

**predict.df** data.frame of predicted species and assignment probabilities for each event.

**Author(s)**

Eric Archer <eric.archer@noaa.gov>

**Examples**

```
data(train.data)
# initialize BANTER model with event data
bant.mdl <- initBanterModel(train.data$events)
# add all detector models
bant.mdl <- addBanterDetector(
  bant.mdl, train.data$detectors,
  ntree = 50, sampsize = 2, num.cores = 1
)
# run BANTER event model
bant.mdl <- runBanterModel(bant.mdl, ntree = 1000, sampsize = 1)

# predict test data
data(test.data)
test.pred <- predict(bant.mdl, test.data)
test.pred
```

**runBanterModel**

*Run BANTER Model*

**Description**

Build full event classifier model

**Usage**

```
runBanterModel(x, ntree, sampsize = 1)
```

**Arguments**

<b>x</b>	a <b>banter_model</b> object.
<b>ntree</b>	number of trees.
<b>sampsize</b>	number or fraction of samples to use in each tree.

**Value**

a **banter\_model** object with the complete BANTER model.

**Author(s)**

Eric Archer <eric.archer@noaa.gov>

**Examples**

```
data(train.data)
# initialize BANTER model with event data
bant.mdl <- initBanterModel(train.data$events)
# add all detector models
bant.mdl <- addBanterDetector(
  bant.mdl, train.data$detectors,
  ntree = 50, sampsize = 1, num.cores = 1
)
# run BANTER event model
bant.mdl <- runBanterModel(bant.mdl, ntree = 1000, sampsize = 1)
summary(bant.mdl)
```

summary

*BANTER Classifier Model Summary***Description**

Display summaries for event and detector models

**Usage**

```
## S3 method for class 'banter_model'
summary(object, model = "event", n = 0.1,
        bins = 20, ...)
```

**Arguments**

- object** a `banter_model` object.
- model** name of model to summarize. Default is "event" to summarize the event-level model. Can also be name of a detector.
- n** number of final iterations to summarize OOB error rate for. If between 0 and 1 is taken as a proportion of chain.
- bins** number of bins in inbag histogram.
- ...** ignored.

**Value**

In the plot that is created, the upper panel shows the trace of the Random Forest model OOB rate across sequential trees in the forest. The lower plot shows a frequency histogram of the number of times each sample was inbag (used as training data in a tree in the forest). The vertical red lines indicate the expected inbag rate for samples of each species.

**Author(s)**

Eric Archer <eric.archer@noaa.gov>

**Examples**

```
data(train.data)
# initialize BANTER model with event data
bant.mdl <- initBanterModel(train.data$events)
# add all detector models
bant.mdl <- addBanterDetector(
  bant.mdl, train.data$detectors,
  ntree = 50, sampsize = 1, num.cores = 1
)
# run BANTER event model
bant.mdl <- runBanterModel(bant.mdl, ntree = 1000, sampsize = 1)
summary(bant.mdl)
```

**test.data***Testing events and detectors***Description**

A list of events and call data from detectors for testing BANTER model

**Usage**

```
data(test.data)
```

**Format**

list

**train.data***Training events and detectors***Description**

A list of events and call data from detectors for training BANTER model

**Usage**

```
data(train.data)
```

**Format**

list

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