

# ReadBiotic

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ReadBiotic

*Read biotic XML files*

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## Description

This function reads multiple biotic file to a list with a list of tables for each file.

## Usage

```
ReadBiotic(FileNames)
```

## Arguments

FileNames      The paths of the biotic files.

## Details

This function is awesome and does excellent stuff.

## Value

An object of StoX data type BioticData: A list of a list of data.tables of the different levels of the input biotic files.

## See Also

[readXmlFile](#).

## Examples

```
x <- 1
```