# Lagged objects in package 'lagged'

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## 1 Class "Lagged"

```
"Lagged" is the base lagged class. It is virtual and defines a slot data from class "ANY".
```

Actual classes inherit from "Lagged" and restrict the data slot.

```
# setClass("X", slots = c(data = "structure"))
setClass("Lagged1d", contains = "Lagged", slots = c(data = "vector") )
setClass("Lagged2d", contains = "Lagged", slots = c(data = "matrix") )
setClass("Lagged3d", contains = "Lagged", slots = c(data = "array") )
# TODO: check validity for Lagged3d: 3 dimensional.
```

One special subclass of "Lagged" is "FlexibleLagged", which can represent objects from any subclass of "Lagged". This is achieved by setting the data slot to be "Lagged" along with methods for indexing, maxLag, and similar to ensure that the Lagged functionality is provided. (\*NOTE (2019-05-15): adding prototype since the default is puzzling and leaves the object invalid if slot data is not set during initialisation. In particular, new("FlexibleLagged"), was returning an invalid object without warning nor error.)

The initialisation method for "FlexibleLagged" objects does the obvious thing if argument data is "Lagged". Otherwise it converts argument data to a suitable "Lagged" object before assigning it to the data slot. If the function is not able to infer a suitable "Lagged" class it still passes data on to the next method which usually leads to an error.

The following convenience function is used to infer a suitable "Lagged" class for argument data:

```
.whichNativeLagged <- function(x){
   if(is(x, "Lagged"))
        "FlexibleLagged"
   else if(is.vector(x))
        "Lagged1d"
   else if(is.matrix(x))
        "Lagged2d"
   else if(is.array(x) && length(dim(x)) == 3)
        "Lagged3d"
   else
        NA</pre>
```

This is the initialisation method for "FlexibleLagged". Note that it gets rid of recursive "FlexibleLagged" data slots, i.e. the data slot of the returned object is "Lagged" but not "FlexibleLagged". **TODO:** This probably should be taken care of by a validation method.

!!! 2019-05-15 Note: Moving argument data after the "..." argument, so that it is matched only when explicitly supplied. Otherwise it makes it difficult to use unnamed arguments in initialisation of subclasses. The success may depend on the order of the arguments and the error message is not helpful. This change is potentially breaking change but it seems that data is always named in existing code.

```
setMethod("initialize", "FlexibleLagged",
          function(.Object, ..., data){
              if(missing(data)){
                  res <- callNextMethod(.Object, ...)</pre>
                  return(res)
              }
              while(is(data, "FlexibleLagged"))
                  data <- data@data
              if(!is(data, "Lagged")){
                  clname <- .whichNativeLagged(data)</pre>
                  if(!is.na(clname))
                       data <- new(clname, data = data)
                  ##else don't know what to do with data, pass it on
                          and let others complain if not appropriate.
              }
               .Object <- callNextMethod(.Object, data = data, ...)
```

```
.Object
}
```

In general, "FlexibleLagged" can be used as superclass of classes which wish to represent any possible subclasses of "Lagged". For slots, it is sufficient (and more efficient) to use "Lagged".

Since class "FlexibleLagged" is special, it has its own implementations of some core methods defined for "Lagged".

**TODO:** Decide what support to offer for the native S3 class "acf". Turn it into S4 using setOldClass? Or just adapt the various methods and constructors to convert it properly to Lagged? For now putting some code in Lagged() to accept "acf" objects.

#### 1.1 Core methods for lagged objects

The methods in this section are ok for objects inheriting from "Lagged". Where necessary, specialised methods are defined for "FlexibleLagged".

## 1.1.1 Subscripting with "["

Subscripting with i missing, returns the raw data.

```
setMethod("[", c(x = "Lagged", i = "missing"), function(x) x@data )
setMethod("[", c(x = "FlexibleLagged", i = "missing"), function(x) x@data[] )
```

When i is present, indexing depends on the type of the data slot and so is defined by subclasses. For indices larger than maxLag(x) the values are filled with NA's.

**TODO:** consider making the 1d method the default one?

For "[", the default for drop is FALSE. TODO: check that the existing methods follow this convention! Indexing "FlexibleLagged" simply transfers the operation to the data slot (it is "Lagged"). I created separate methods for drop since just omitting it from the method signature and setting drop = FALSE in the signature of the function seems to have no effect (and findMethods("FlexibleLagged") shows that the default for drop in the method is TRUE).

#### 1.1.2 Subscript-replacement with "[<-"

Similarly to "[", subscript-replacement "[<-" replaces the contents of the data. The method for "Lagged" does not check the validity of argument value but the assignment will raise an error if it is not appropriate. Subclasses that wish to provide finer control over this can define suitable methods (e.g. to coerce value appropriately).

Assignment to "FlexibleLagged", when i is missing, attempts to coerce value to a suitable lagged class before assigning it (using .whichNativeLagged(), as the initialisation function does, but raising an error if unsuccessful). Further methods can be defined using "value = xxx" in the signature to accommodate additional types or overwrite the default method here.

```
setReplaceMethod("[", c(x = "FlexibleLagged", i = "missing"),
                  function(x, i, value){
                      if(is(value, "FlexibleLagged"))
                          x@data <- value@data
                      else if(is(value, "Lagged"))
                          x@data <- value
                      else{
                          clname <- .whichNativeLagged(value)</pre>
                          if(is.na(clname))
                              stop("Don't know what Lagged class to use for this value")
                          else
                              x@data <- new(clname, data = value) # as(value, clname)</pre>
                      }
                      х
                 })
When i is present, no attempt is made to coerce it:
setReplaceMethod("[", c(x = "FlexibleLagged", i = "numeric"),
                 function(x, i, value){
                      x@data[i] <- value # not i+1, since x@data is a "Lagged" object here.
                      х
          })
## Ne, tezi zasega ne gi pravya, pravya vischko bez "value = xxx" - tova pozvolyava da se
## definirat metodi ako tryabva.
##
## setReplaceMethod("[", c(x = "FlexibleLagged", i = "missing", value = "vector"),
##
             function(x, i, value){
                 x@data <- as(value, "Lagged1d")
##
##
##
             })
##
##
   setReplaceMethod("[", c(x = "FlexibleLagged", i = "missing", value = "matrix"),
             function(x, i, value){
##
                 x@data <- as(value, "Lagged2d")
##
##
##
             })
```

### 1.1.3 Methods for "[[" and "[]<-"

Indexing with "[[" returns the value for the specified lag. This is the recommended way to extract the value at a single index.

This defines a default method. For efficiency specific classes can define versions that avoid calling the generic "[[". If multi-seasons are supported the check for length equal to one should be adapted accordingly.

```
else
     stop("length of argument 'i' must be equal to one")
}
)
```

Note the use of drop = TRUE.

**TODO:** The use of drop = TRUE maybe needs some further thought. Maybe something that drops only the index corresponding to the lag is preferable and such behaviour should be documented!

The replace method works similarly:

#### 1.1.4 Arithmetic and other operations (Ops group)

Operations in the Ops group involving lagged objects are defined "naturally" on their data. However, they are more restrictive than base R's conventions for atomic objects and do not follow the recycling rules.

The binary "Ops" methods return values from one of the core lagged classes, even if the objects are from classes inheriting from "Lagged". The reason is that, for example, the difference between autocovariance functions is not necessarily autocovariance, but it is still a lagged object. It would be very confusing if the result was not guaranteed to be "Lagged". Also, if a policy of preserving the actual class were to be adopted, what would be the rule for the class of the result from binary operations between lagged objects from different classes (it seems not possible to have a simple one).

Of course, methods defined for subclasses of lagged objects may preserve the actual classes when appropriate but should not introduce confusion on indexing.

In the default methods below, the result of these operations is a strict lagged object, i.e. an object from the core lagged classes (**TODO**: explain). The exact type of lagged object is determined by the data. The net effect is that the value of the Ops operation is also a lagged object, a core one, with indexing starting from zero but additional structure is lost.

**TODO:** Should operations between "Lagged" and base R objects be permitted at all? For users of "Lagged" the returned "Lagged" value is natural and expected. But what about users who are not aware that there are "Lagged" objects among the arguments? What to do when the "ordinary" argument is of length one - should this be an exception? But then the user may not know that the length is one, leading to surprises. Also, there is a conceptual difference here between the additive and multiplicative operations. (All this should be documented in a vignette. It seems sufficient that the recycling rule is banned. Need to finalise operation with singletons.)

Operations between two lagged objects give a lagged object. If their maxLag() properties are different, the shorter data slot is extended with NA's before applying the binary operation.

#### 1. "Ops" involving "Lagged"

The unary operators preserve the class of the object:

**TODO:** the current mechanism to decide the lagged class of the return value is not very satisfactory, see whichLagged() which encapsulates it. Also, forbid mixing 1d with 2d, etc.?

When only one of the objects is "Lagged", the operations are defined if the following cases:

- (a) the length of the other object is equal to the length of the data part of the "Lagged" object,
- (b) the other object is of length one,

})

})

(c) the other object is a singleton with the same dimensions as a single element of the "Lagged" object.

old todo: document behaviour if length(object@data) == 0 (minor issue)?

2017-05-20 TODO: Change length(e1[[0]]) == length(e2)) below to dim(e1[[0]]) == dim(e2)) but needs more care (note though that the scalar case is covered by length(e2) == 1.

Notice that "vector" in the signatures is the S4 class "vector" (TODO: check!), see showClass("vector") for its subclasses.

```
setMethod("Ops", c(e1 = "vector", e2 = "Lagged"),
             function(e1, e2){
                 wrk <- if(length(e1) == 1 || length(e1) == length(e2@data)</pre>
                                 # 2017-05-20 was:
                                      || length(e1) > 0 && (length(e2@data) %% length(e1)) == 0
                            || length(e1) > 0 && length(e2[[0]]) == length(e1))
                             callGeneric(e1, e2@data)
                        else
                             stop("Incompatible length of operands in a binary operation")
                 new(whichLagged(e2), data = wrk)
             })
2. "Ops" involving "FlexibleLagged"
  Operations involving "FlexibleLagged" objects use those defined for "Lagged" by operating on the
```

data slot (which is "Lagged").

```
setMethod("Ops", c(e1 = "FlexibleLagged", e2 = "Lagged"),
          function(e1, e2){
              callGeneric(e1@data, e2)
          })
setMethod("Ops", c(e1 = "Lagged", e2 = "FlexibleLagged"),
          function(e1, e2){
              callGeneric(e1, e2@data)
          })
setMethod("Ops", c(e1 = "FlexibleLagged", e2 = "FlexibleLagged"),
          function(e1, e2){
              callGeneric(e1@data, e2@data)
          })
setMethod("Ops", c(e1 = "FlexibleLagged", e2 = "vector"),
          function(e1, e2){
              callGeneric(e1@data, e2)
          })
setMethod("Ops", c(e1 = "vector", e2 = "FlexibleLagged"),
          function(e1, e2){
              callGeneric(e1, e2@data)
          })
```

TODO: methods for "matrix", "array", these probably should be for specific "Lagged" subclasses, like "Lagged2d".

#### 1.1.5 "Math" and "Math2" group methods

"Math" and "Math2" methods return the object with its data part transformed by the corresponding function.

```
TODO: Does this work for FlexibleLagged?
```

```
setMethod("Math", c(x = "Lagged"),
          function(x){
```

```
x@data <- callGeneric(x@data)
x
})
setMethod("Math2", c(x = "Lagged"),
    function(x, digits){
        x@data <- callGeneric(x@data, digits)
        x
})</pre>
```

#### 1.1.6 "Summary" group methods

The "Summary" methods operate on the data part of the "Lagged" object.

### 1.2 S3 methods for as.vector() and related functions for "Lagged"

```
## TODO: check if the S3 methods understand S4 inheritance (I think they do)
as.vector.Lagged <- function(x, mode) as.vector(x@data) # todo: use mode?
as.double.Lagged <- function(x, ...) as.double(x@data) # note: this is for as.numeric()
as.matrix.Lagged <- function(x, ...) as.matrix(x@data)
as.array.Lagged <- function(x, ...) as.array(x@data)</pre>
```

Converting from "Lagged" to base atomic or structure objects applies the requested operation to the data slot. Define first the generic S3 methods:

Somewhat more efficient methods for these:

```
as.vector.Lagged1d <- function(x, mode) x@data
as.matrix.Lagged2d <- function(x, ...) x@data
as.array.Lagged3d <- function(x, ...) x@data
```

## 1.3 setAs() methods for "Lagged"

These methods call the corresponding S3 methods defined above:

```
setAs("Lagged", "vector", function(from) as.vector(from))
setAs("Lagged", "matrix", function(from) as.matrix(from))
setAs("Lagged", "array", function(from) as.array(from))
```

#### 1.4 Generic function maxLag()

The default method for maxLag() handles objects inheriting from S3 class "acf". In all other cases it raises an error. Notice that in "acf" the lag is in the first dimension.

```
maxLag <- function(object, ...){
   if(inherits(object, "acf"))
      dim(acf$acf)[1] - 1
   else
      stop("No applicable method to compute maxLag")
}
setGeneric("maxLag")</pre>
```

The convention for "Lagged" objects is that the last dimension carries the lag. So, the methods for basic objects compute the maximal lag as the last dimension minus one.

Note again that acf() puts the lag in the first index.

The maxLag() method for "Lagged" objects simply calls maxLag() on the data slot. Classes inheriting from "Lagged" may define specific methods if the (in)efficiency of this method is a concern.

```
setMethod("maxLag", c(object = "Lagged"), function(object) maxLag(object@data) )
```

#### 1.5 Length of "Lagged" objects - S3 method for length()

The length of "Lagged" objects is defined to be maxLag(x)+1, not the length of the data in the "Lagged" object. In most cases of direct use maxLag(x) is more appropriate.

This defines an S3 method for function length() for "Lagged" objects.

```
length.Lagged <- function(x) maxLag(x) + 1</pre>
```

**TODO:** Check if other base R functions need S3 methods for "Lagged" objects.

## 2 Methods for "["

**TODO:** Don't give option to change argument drop and raise error if it is present?

**TODO:** Currently silently ignores argument j. Throw error if it is present? Another alternative is to define the current methods with j = "missing" but then explicit methods for "ANY" will be needed to avoid some other method matching quietly and doing something even less relevant.

TODO: For "Lagged2d", add argument type to accommodate indexing like "slMatrix". Need some thought to streamline that old stuff. However, it may be better not to put this on "Lagged2d" but introduce subclass, say "LaggedSL" of ="Lagged2d" and define modified methods for that, streamlining in the process.

## 2.1 whichLagged()

For now whichLagged() is not exported. It could be exported to allow core "Lagged" classes defined in other packages to add functionality. But if it is to be exported, it would need streamlining. Currently it is a hack.

Making it generic is lazy but avoids writing obscure code but see note above. The default returns "FlexibleLagged".

```
.matLagged <- matrix("FlexibleLagged", 4, 4)
diag(.matLagged) <- c("FlexibleLagged", "Lagged1d", "Lagged2d", "Lagged3d")
rownames(.matLagged) <- c("FlexibleLagged", "Lagged1d", "Lagged2d", "Lagged3d")
colnames(.matLagged) <- c("FlexibleLagged", "Lagged1d", "Lagged2d", "Lagged3d")

whichLagged <- function(x, y){
    .matLagged[whichLagged(x), whichLagged(y)]
}
setGeneric("whichLagged")

## TODO: define methods for "numeric", "matrix", etc?
setMethod("whichLagged", c(x = "ANY" , y = "missing"), function(x) "FlexibleLagged")
setMethod("whichLagged", c(x = "Lagged1d", y = "missing"), function(x) "Lagged1d")
setMethod("whichLagged", c(x = "Lagged2d", y = "missing"), function(x) "Lagged2d")
setMethod("whichLagged", c(x = "Lagged3d", y = "missing"), function(x) "Lagged3d")</pre>
```

## 2.2 Methods for "[<-"

Missing index is equivalent to replacing all data:

The above method just calls "[<-" again, so it applies to any lagged objects.

The methods which work on the data, need to know their layout, so we need several methods.

```
## Include value = "array" in the signature? Will still need to check the dimensions
setReplaceMethod("[", c(x = "Lagged3d", i = "numeric"),
          function(x, i, value){
                       # was: x@data[i+1, , ] <- value</pre>
               x@data[,,i+1] \leftarrow value
          })
    show() methods
3
   .printVecOrArray <- function(x){</pre>
       if(is.vector(x)){
##
##
            if(is.null(names(x)) || length(names(x)) == 0)
##
                names(x) <- paste0("Lag_", 0:(length(x) - 1))</pre>
##
           print(x)
       }else if(is.matrix(x)){
##
           ## TODO:
##
           print(x)
##
##
       }else if(is.array(x)){
##
           ## TODO:
##
           print(x)
##
       }else
##
           print(x)
## }
setMethod("show", "Lagged1d",
          function(object){
               .reportClassName(object, "Lagged1d")
               cat("Slot *data*:", "\n")
               ## 2017-05-24 was:
               ##
                      x <- object@data
                      if(is.null(names(x)) || length(names(x)) == 0)
               ##
                          names(x) <- paste0("Lag_", 0:(length(x) - 1))</pre>
               x <- dataWithLagNames(object)</pre>
               print(x)
               ## cat("\n")
          }
          )
setMethod("show", "Lagged2d",
          function(object){
               .reportClassName(object, "Lagged2d")
               cat("Slot *data*:", "\n")
               x <- dataWithLagNames(object)</pre>
               print(x)
               ## cat("\n")
          }
          )
setMethod("show", "Lagged3d",
          function(object){
```

```
.reportClassName(object, "Lagged3d")
              cat("Slot *data*:", "\n")
              ## x <- object@data
              ## if(is.null(dimnames(x)) || length(dimnames(x)) == 0){
                      d <- dim(x)
              ##
                      dimnames(x) <- list(rep("", d[1]), rep("", d[2]),</pre>
              ##
                                           paste0("Lag_", 0:(d[3] - 1)) )
              ## }
              x <- dataWithLagNames(object)</pre>
              print(x)
              ## cat("\n")
          }
          )
## Commenting out since causes trouble by precluding default methods from printing.
##
  setMethod("show", "Lagged",
##
##
             function(object){
                  ## .reportClassName(object, "Lagged") # this is silly: never writes!
##
                 ## callNextMethod()
##
                  wrk <- object@data
##
                  cat("Slot *data*:", "\n")
##
                  .printVecOrArray(wrk)
##
                  cat("\n")
##
                  ## callNextMethod() # in case the object inherits from other classes
##
##
                                       # unfortunately, it prints slot data again.
             }
##
##
setMethod("show", "FlexibleLagged",
          function(object){
               .reportClassName(object, "FlexibleLagged")
              cat("Slot *data*:", "\n")
              show(object@data)
          }
          )
```

## 4 Further constructors for lagged objects

Function new() can be used to create objects from the lagged classes. In this section we define some functions to make this more convenient.

First, a function to convert objects from S3 class "acf" (created by acf()) to "Lagged":

```
acf2Lagged <- function(x){
   acv <- x$acf
   d <- dim(acv)
   if(d[2] == 1 && d[3] == 1){
      data <- as.vector(acv)
      if(x$type == "partial") # lag-0 is missing, insert it
            data <- c(1, data)
      new("Lagged1d", data = data)
}else{
      ## transpose to make the 3rd index corresponding to lag.</pre>
```

```
##
              (taken from acfbase2s1() in package pcts, see the comments there)
        ##
        ## TODO: test!
        ## Note: in pcts:::acfbase2s1() the analogous command is aperm(acv, c(3,2,1))
                  i.e. R[k] is transposed => check if that is correct!
        ##
        data \leftarrow aperm(acv, c(2, 3, 1))
        if(x$type == "partial"){ # lag-0 is missing, insert it
             datanew <- array(NA_real_, dim(data) + c(0,0,1) )</pre>
            datanew[ , , -1] \leftarrow data
            data <- datanew
        }
        new("Lagged3d", data = data)
    }
}
```

Function "Lagged" looks at the supplied data argument and chooses an appropriate class inheriting from "Lagged". **TODO:** Make Lagged() generic?

```
Lagged <- function(data, ...){
   if(is.vector(data)){
      new("Lagged1d", data = data, ...)
}else if(is.matrix(data)){
      new("Lagged2d", data = data, ...)
}else if(is.array(data)){
      new("Lagged3d", data = data, ...)
}else if(is(data, "Lagged")){
      new("FlexibleLagged", data = data, ...)
}else if(inherits(data, "acf")){  # for S3 class "acf"
      acf2Lagged(data)
}else
      stop("I don't know how to create a Lagged object from the given data")
}</pre>
```

TODO: Tests!

#### 5 New

The functions in this section are temporarily here during development and should be move to more appropriate places eventually.

\*\*

provideDimnames is new in R-3.0.0

dataWithLagNames(object) is a convenience function which works like object[] but also ensures that the lag dimension has names. It is exported for use in other packages. Occasionally users may wish to use it too.

```
dataWithLagNames <- function(object, prefix = "Lag_"){
    x <- object[]
    if(length(x) == 0)
        return(x)

if(is.array(x)){
        d <- dim(x)
        nd <- length(d)</pre>
```

```
xwithnams <- provideDimnames(x, base = list(""), unique = FALSE)
    dimnames(xwithnams)[[nd]] <- pasteO(prefix, 0:(d[nd] - 1))
    xwithnams
}else{
    if(is.null(names(x)) || length(names(x)) == 0)
        names(x) <- pasteO(prefix, 0:(length(x) - 1))
    x
}</pre>
```