

# The `builtenvir` package for linear DLM's

(With intro to OOP)

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- OOP in R
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# Intro to OOP

## Object Oriented Programming

- So-called by analogy to objects in the real world
- For example, a physical door has a set of *attributes* like a handle or knob, a hinge-mechanism, the dimensions of the doorway, ...
- When you encounter a door, you always know just how to use it—it doesn't matter if it's the door to SPH, the door to your car, or the door on your oven, some similar action allows you to *open* it
- Programmers use this kind of abstraction to design efficient and easy to use software

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# Intro to OOP: Objects in R

Whether or not you've realized it, if you've used R, you've probably interacted with OOP design

For instance, take the “object” returned by `lm()`:

```
## simulate data:
set.seed(12345)
n <- 100
p <- 2
x <- rnorm(n)
b <- rnorm(p) # beta coefs
sigma.sq <- 1
y <- rnorm(n, cbind(1, x) %*% b, sqrt(sigma.sq))

fit <- lm(y ~ x)
```



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# Intro to OOP: Objects in R

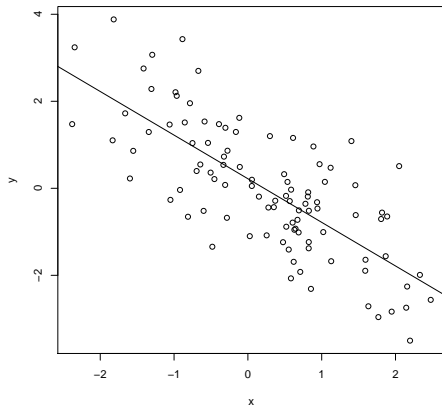
```
> fit
```

```
Call:
```

```
lm(formula = y ~ x)
```

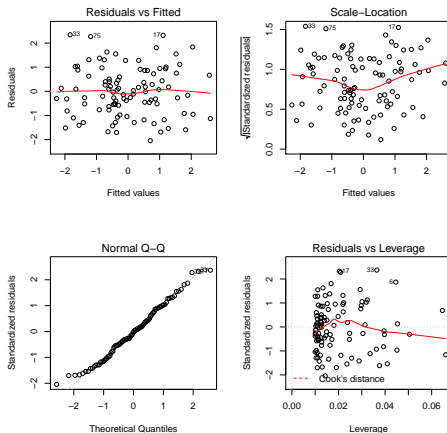
```
Coefficients:
```

(Intercept)	x
0.2201	-1.0024



# Intro to OOP: Objects in R

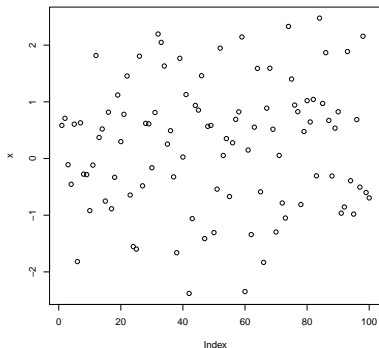
```
> plot(fit)
```



# Intro to OOP: Objects in R

And many more functions exist to interact with the objects `lm()` returns...  
How does R know exactly what to do with these commands? And why  
does the output of `plot(fit)` look different from  $\Downarrow$ ?

```
> plot(x)
```



# Intro to OOP: Objects in R

The answer is because the R developers chose to organize these kinds of functions as *methods* that act on different objects [e.g. `open(door)`]

```
> class (fit)
[1] "lm"
>
> class (x)
[1] "numeric"
```

# Intro to OOP: Objects in R

```
> methods(class = "lm")
```

[1] add1	alias	anova	case.names	coerce
[6] confint	cooks.distance	deviance	dfbeta	dfbetas
[11] drop1	dummy.coef	effects	extractAIC	family
[16] formula	hatvalues	influence	initialize	kappa
[21] labels	logLik	model.frame	model.matrix	nobs
[26] plot	predict	print	proj	qr
[31] residuals	rstandard	rstudent	show	simulate
[36] slotsFromS3	summary	variable.names	vcov	

see `'?methods'` for accessing help and source code

# Intro to OOP: Objects in R

Internally, when you type

```
> fit
> getAnywhere(print.lm)
A single object matching print.lm was found
It was found in the following places
  registered S3 method for print from namespace stats
  namespace:stats
with value

function (x, digits = max(3L, getOption("digits") - 3L), ...)
{
  cat("\nCall:\n", paste(deparse(x$call), sep = "\n", collapse = "\n"),
      "\n\n", sep = "")
  if (length(coef(x))) {
    cat("Coefficients:\n")
    print.default(format(coef(x), digits = digits), print.gap = 2L,
                  quote = FALSE)
  }
  else cat("No coefficients\n")
  cat("\n")
  invisible(x)
}
<bytecode: 0x7f8ccc3d5e00>
<environment: namespace:stats>
```

Call:

```
lm(formula = y ~ x)
```

Coefficients:

	x
(Intercept)	0.2201
	-1.0024

# Intro to OOP: Objects in R

- And the same with:
  - `plot(fit) → plot.lm`
  - `summary(fit) → summary.lm`
  - ...
- We can and should take advantage of this organization structure when we write R code
- Users should rightly expect `plot(fitted-model)` to behave similarly regardless of the specific program or type of model
- Not only can this approach produce user-friendly code, the structure can be more efficient in terms of what the *language* expects as well (more on this later)



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# Intro to OOP: Defining new R classes

- There are multiple ways of doing this in R, including: S3, S4, and Reference Classes
- Won't go into all these, but you can read more about them for example [here](#)
- Most of base R uses S3 classes and methods; newer packages often use some combination of the three

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# builtenvir: Class Structure

The primary purpose of `builtenvir` at the moment is to provide users a way to fit Distributed Lag Models

Similarly to `stats::lm`, where the output is an object of class `'lm'`, `builtenvir::dlm` outputs an object of class `'Dlm'`. The goal is to make the syntax as classically R as possible:

```
library (builtenvir)
data (simdata)

lag <- seq(0.1, 10, length.out = 100) # distances
X <- simdata[, -(1:3)] # measurements binned by lag distance

fit <- dlm(Y ~ Age * Gender + cr(lag, X), data = simdata)
```



# builtinvir: Class Structure

```
> fit
Call: dlm(formula = Y ~ Age * Gender + cr(lag, X), data = simdata)
(frequentist) distributed lag model fit via REML
Number of Observations: 1000
Log-Likelihood: -1261.647
Fixed Effects:
      (Intercept)      Age      Gender Age:Gender
[1,]      0.33080 -0.49092  1.15477      -0.009

      Random Effects Residuals
Std Error      0.016283      0.8299
>
> class(fit)
[1] "FreqDlm"
attr(,"package")
[1] "builtinvir"

> inherits(fit, "Dlm")
[1] TRUE
```

# builtinvir: Class Structure

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> inherits(fit, "Dlm")
[1] TRUE
```

# builtenvir: Class Structure

```
> summary(fit)
(frequentist) distributed lag model fit via REML
Call: dlm(formula = Y ~ Age * Gender + cr(lag, X), data = simdata)
```

Number of observations: 1000

Standardized Residuals:

	Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
	-3.25001	-0.70212	-0.02999	0.00000	0.68596	2.87362

Random Effects Residuals

Var		
	0.00026513	0.6888

Fixed Effects:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	0.3307981	0.3600354	0.9188	0.358427
Age	-0.4909218	0.0048446	-101.3336	< 2.2e-16 ***
Gender	1.1547739	0.4331537	2.6660	0.007801 **
Age:Gender	-0.0089778	0.0066326	-1.3536	0.176175

---

Signif. codes: 0 \*\*\* 0.001 \*\* 0.01 \* 0.05 . 0.1 1

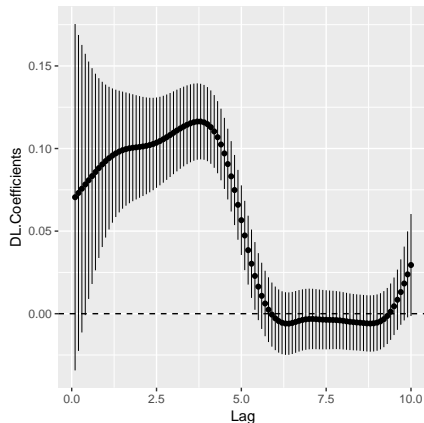
Correlation:

	(Intercept)	Age	Gender
Age	-0.88016		
Gender	-0.65179	0.72642	
Age:Gender	0.64604	-0.73193	-0.9926

# builtinvir: Class Structure

Plot doesn't work exactly the same way (no diagnostics), but that's by design:

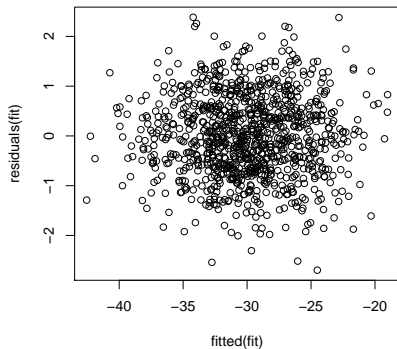
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> plot(fit)
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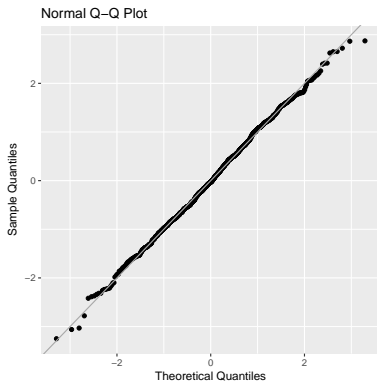
# builtenvir: Class Structure

Can still get residuals and QQ plots easily from similar methods:

```
> plot(fitted(fit), residuals(fit))
```



```
> qqnorm(fit)
```



# builtinvir: Lazy Programming

One of the first rules of good programming is to do no more work than necessary. This goes hand in hand with using a system of classes to be more consistent with what the *language* expects

Quick example: just by defining an R-consistent `logLik` method, we can get the functions `AIC` and `BIC` for free

```
logLik.Dlm <- function(object, ...) {  
  ll <- object@logLik  
  attr(ll, "nall") <- attr(ll, "nobs") <- object@N  
  attr(ll, "df") <- object@K$fixed + 2  
  ## fixed effects + 2 variance parameters  
  class(ll) <- "logLik"  
  return (ll)  
}  
  
> logLik(fit)  
'log Lik.' -1261.647 (df=8)  
>  
> AIC(fit)  
[1] 2539.294  
> BIC(fit)  
[1] 2578.556
```

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# builtenvir: Extending the Package

Ultimately, most of the methods defined for `stats::lm` and `nlme::lme` should probably be implemented for `builtenvir::Dlm` + a few extras

```
> methods(class = "Dlm")  
[1] coef      coerce<-  deviance  fitted    fixef      logLik     plot  
[8] predict   qqnorm    ranef     residuals scaleMat   se.fixef   se.ranef  
[15] show      sigma     theta     vcov      vcovTheta  
see '?methods' for accessing help and source code
```

# builtenvir: Extending the Package

- We'll also want to have `gdlm`'s for logistic models, etc
- This should involve creation of `Gdlm` and `SummaryGdlm` classes and possibly sub-classes
- I've kept with the strategy of using `S4` classes and a combination of `S3` and `S4` methods defined on those classes

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I'd very much appreciate your help  
with beta-testing!

Thanks!