

# Algebraic Statistics in A State of the Union



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- Historical Perspective
- The **algstat** Ecosystem
  - Installing the packages
  - **mpoly**
  - **latter**
  - **m2r**
  - **bertini**
  - **tropical**
  - **algstat**
- GitHub and Contributing
- Upcoming Projects

# Historical Perspective

Base R has essentially no support for symbolic computing

But it does have:

- a rich collection of base object types (S3, S4, R6, ...)

- a variety of ways to create new objects

- flexible mechanisms for implementing new methods

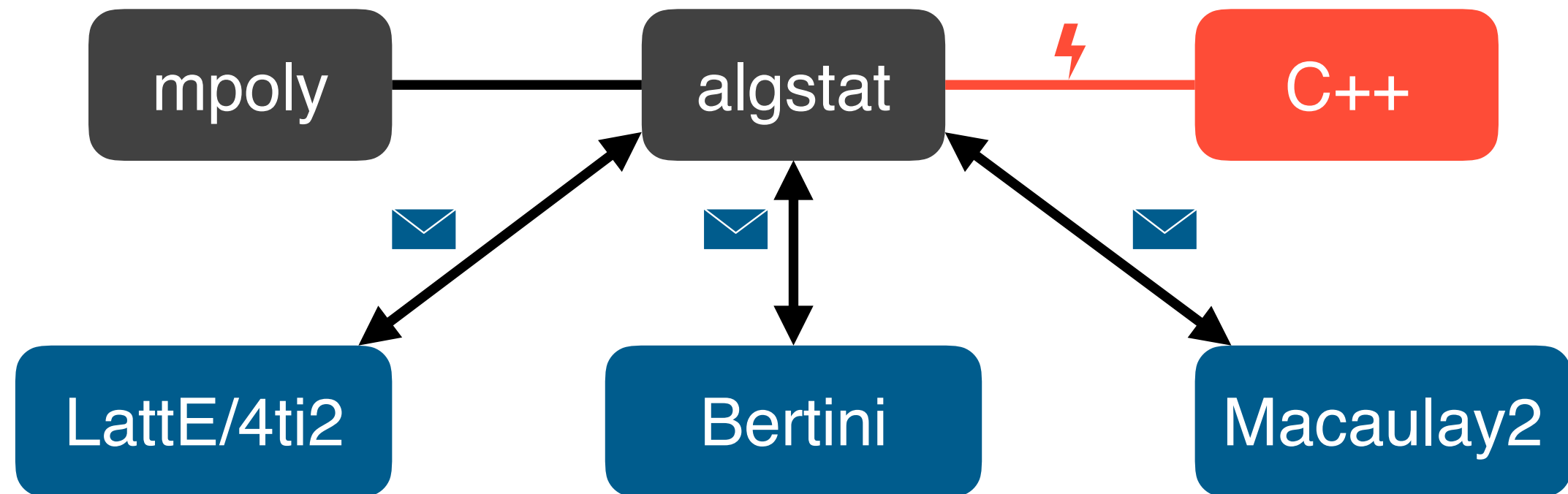
  - (including infix operators and overloading)

- access to the operating system (rw, sockets, ...)

- a simple way to incorporate C++ routines (Rcpp)

2011 : **mpoly** – data structures and methods for multivariate polynomials

2014 : **algstat** – algebraic statistical data analysis



Objects computed on in place



R writes / program executes / R reads

2011 : **mpoly** – data structures and methods for multivariate polynomials

2014 : **algstat** – algebraic statistical data analysis

Then, connections needed their own packages!

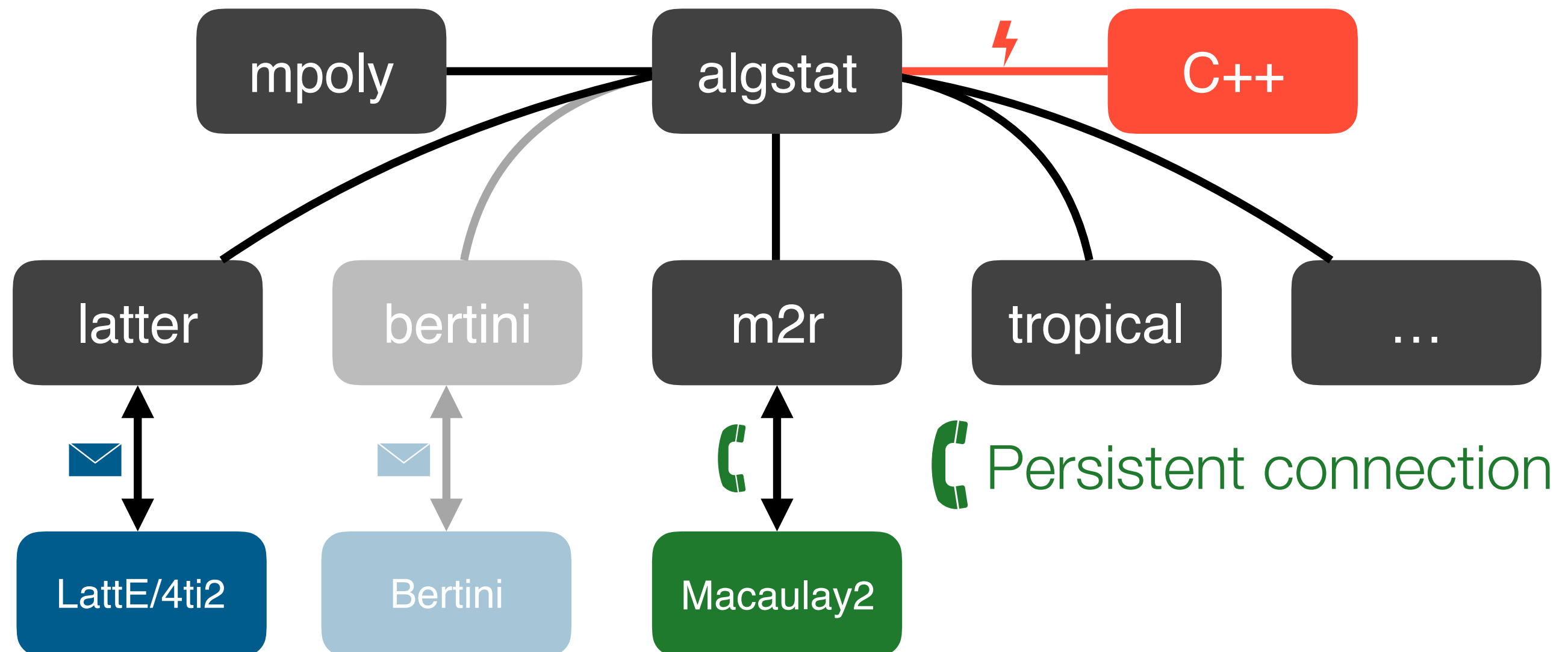
2015 : **latter** – LattE/4ti2

2016 : **m2r** – Macaulay2

2017 : **tropical** – Tropical geometry

2011 : **mpoly** – data structures and methods for multivariate polynomials

2014 : **algstat** – algebraic statistical data analysis



\* Many other interconnections not shown

# The algstat Ecosystem



To install the packages I'm using in this talk, copy/paste the following code into R

```
if(!requireNamespace("devtools"))  
  install.packages("devtools")  
devtools::install_github("dkahle/mpoly")  
devtools::install_github("dkahle/latter")  
devtools::install_github("dkahle/tropical")  
devtools::install_github("coneill-math/m2r")  
devtools::install_github("dkahle/algstat")
```

For latter, you'll need LattE/4ti2, which you can get [here](#)

For algstat, you'll want Bertini, which you can get [here](#)

`mpoly` contains data structures and methods for polynomials

Understanding `mpoly` requires understanding a bit about the R language

## homogeneous

must hold the same kinds of objects

## heterogeneous

can hold different kinds of objects

1d

atomic vector

int, num, char, logi, ...

list

2d

matrix

data frame

list of equal-length atomic vectors

nd

array

```
(x <- c(1, 2, 3)) # a vector  
# [1] 1 2 3
```

```
(y <- list(1:3, c("a", "b"), TRUE)) # a list  
# [[1]]  
# [1] 1 2 3  
#  
# [[2]]  
# [1] "a" "b"  
#  
# [[3]]  
# [1] TRUE  
#
```

R objects can also have a list of metadata attached to them called **attributes**

Two basic types of attributes are **names** and **class**

**Names** are used for easy, non-index based referencing

**Classes** are R's most basic object oriented framework

**Classes** are also used for method dispatch

(Methods are not contained in objects in R)

```
attr(x, "names") <- c("x", "y", "z")
```

```
x
```

```
# x y z
```

```
# 1 2 3
```

```
attr(x, "class") <- "foo"
```

```
x
```

```
# x y z
```

```
# 1 2 3
```

```
# attr("class")
```

```
# [1] "foo"
```

```
print(x)
```

```
# x y z
```

```
# 1 2 3
```

```
# attr("class")
```

```
# [1] "foo"
```

Typing x at the command line implicitly calls `print(x)`

```
print(x)
```

```
# x y z
```

```
# 1 2 3
```

```
# attr("class")
```

```
# [1] "foo"
```

```
print.foo <- function(.) {  
  cat( paste0(names(.), "^", .) )  
}
```

```
x
```

```
# x^1 y^2 z^3
```

```
str(x)
```

```
# Class 'foo' Named num [1:3] 1 2 3
```

```
# ..- attr(*, "names")= chr [1:3] "x" "y" "z"
```

mpoly's main basic data structure is the mpoly,  
a list of numeric vectors representing terms

```
library(mpoly)
mp("x^2 + 2 x y - 1")
# x^2 + 2 x y - 1

str( mp("x^2 + 2 x y - 1") )
# List of 3
# $ : Named num [1:2] 2 1
# ..- attr(*, "names")= chr [1:2] "x" "coef"
# $ : Named num [1:3] 1 1 2
# ..- attr(*, "names")= chr [1:3] "x" "y" "coef"
# $ : Named num -1
# ..- attr(*, "names")= chr "coef"
# - attr(*, "class")= chr "mpoly"
```



`mpoly` provides all basic arithmetic operations as well as well as related tools

```
p <- mp("x + y"); q <- mp("x - y")
```

```
p + q
```

```
# 2 x
```

```
p * q
```

```
# x^2 - y^2
```

```
p^2
```

```
# x^2 + 2 x y + y^2
```

```
f <- as.function(p, vector = FALSE)
```

```
# f(x, y)
```

```
f(1, 2)
```

```
# [1] 3
```

Vectors of mpolys are also defined with similar methods

```
(ps <- mp(c("x + y", "x - y^2")))
```

```
# x + y
# x - y^2
```

```
g <- as.function(ps, vector = FALSE)
```

```
g(1, 2)
```

```
# [1] 3 -3
```

These kinds of lists are used by most other packages in the **algstat** ecosystem

`latter` implements back-end connections to LattE/4ti2

It is used mostly for computing Markov and related bases

```
library(latter)
```

```
# LattE found in /Applications/latte/dest/bin
```

```
# 4ti2 found in /Applications/latte/dest/bin
```

```
(A <- genmodel(c(2, 2), 1:2)) # 2x2 independence model
```

```
#      [,1] [,2] [,3] [,4]
```

```
# [1,]    1    0    1    0
```

```
# [2,]    0    1    0    1
```

```
# [3,]    1    1    0    0
```

```
# [4,]    0    0    1    1
```

```
markov(A)
```

```
#      [,1]
```

```
# [1,]    1
```

```
# [2,]   -1
```

```
# [3,]   -1
```

```
# [4,]    1
```

```
graver(A)
```

```
#      [,1]
```

```
# [1,]    1
```

```
# [2,]   -1
```

```
# [3,]   -1
```

```
# [4,]    1
```

LattE functions are also available

```
spec <- c("x + y <= 10", "x >= 1", "y >= 1")
```

```
count(spec)
```

```
# [1] 45
```

```
count(spec, dilation = 10)
```

```
# [1] 3321
```

```
latte_max(
```

```
  "-2 x + 3 y",
```

```
  c("x + y <= 10", "x >= 0", "y >= 0")
```

```
)
```

```
# $par
```

```
#   x   y
```

```
#   0 10
```

```
#
```

```
# $value
```

```
# [1] 30
```

m2r connections, data structures, and methods for Macaulay2

Since Macaulay2 is a full computer algebra system, a persistent connection is needed

So... stay tuned for Chris O'Neill's talk next!

Bertini was one of the original algstat connections, but it hasn't been implemented in its "Version 2" form yet

```
library(algstat)
polySolve(
  c("y == x^2", "y == 2 - x^2"),
  varOrder = c("x", "y")
)
# 2 solutions (x,y) found.
# (2 real, 0 complex; 2 nonsingular, 0 singular.)
#      (-1,1) (R)
#      ( 1,1) (R)
```

A new direction of **algstat** applications is phylogenetics

**tropical** is a (very!) new package intended to supply the necessary tropical geometry computations

```
library(tropical)  
# Using min-plus algebra.
```

$$x \oplus y = \min(x, y)$$

$$x \otimes y = x + y$$

Max-plus is available with `set_plus_max()`



```
# basic tropical arithmetic
```

```
1 %+% 5
```

```
# [1] 1
```

```
1 %.% 5
```

```
# [1] 6
```

```
5 %^% 3
```

```
# [1] 15
```

```
# vectorized for R-users
```

```
1:3 %+% 3:1
```

```
# [1] 1 2 1
```

```
1:3 %.% 3:1
```

```
# [1] 4 4 4
```

```
# tropical mat. mult
```

```
1:3 %..% 4:6
```

```
# [1] 5
```

```
(m1 <- matrix(1:6, 2, 3))
```

```
#      [,1] [,2] [,3]
```

```
# [1,]    1    3    5
```

```
# [2,]    2    4    6
```

```
(m2 <- matrix(6:1, 3, 2))
```

```
#      [,1] [,2]
```

```
# [1,]    6    3
```

```
# [2,]    5    2
```

```
# [3,]    4    1
```

```
m1 %..% m2
```

```
#      [,1] [,2]
```

```
# [1,]    7    4
```

```
# [2,]    8    5
```

```
data(politics)
```

```
politics
```

```
#           Party
# Personality Democrat Republican
#  Introvert           3           7
#  Extrovert           6           4
```

```
(A <- hmat(c(2, 2), list(1, 2))) # alternative to genmodel
```

```
#      11 12 21 22
# 1+   1  1  0  0
# 2+   0  0  1  1
# +1   1  0  1  0
# +2   0  1  0  1
```

```
countTables(politics, A)
```

```
# [1] 10
```

```
loglinear(~ Personality + Party, data = politics)
# Computing Markov moves (4ti2)... done.
# Running chain (C++)... done.
# Call:
# loglinear(model = ~Personality + Party, data = politics)
#
# Fitting method:
# Iterative proportional fitting (with stats::loglin)
#
# MCMC details:
# N = 10000 samples (after thinning), burn in = 1000, thinning = 10
#
#           Distance      Stat      SE p.value      SE mid.p.value
#           P(samp)                0.3677 0.0048      0.2201
#   Pearson X^2 1.8182 0.0146 0.3677 0.0048      0.2201
# Likelihood G^2 1.848 0.0155 0.3677 0.0048      0.2201
# Freeman-Tukey 1.8749 0.0167 0.3677 0.0048      0.2201
# Cressie-Read 1.8247 0.0148 0.3677 0.0048      0.2201
```

# GitHub and Contributing

To submit a feature request or report a bug:

- Go to <https://github.com/> and create a free account
- Go to <https://github.com/dkahle/algstat>
- Click *Issues*
- Click *New Issue*

To join the fray, submit a pull request (PR)!

- Go to <https://github.com/> and create a free account
- Go to <https://github.com/dkahle/algstat>
- Click *Fork* to make your own copy of the repository
- In RStudio...
  - File > New Project... > Version Control > Git
  - Enter the URL of the repo, <https://github.com/dkahle/algstat.git>
  - Make changes to the code and commit them, see [tutorial here](#)
  - Push changes to GitHub
- On GitHub, click *Submit a Pull Request*

# Upcoming Projects

tropical with Grant Innerst, Rudy Yoshida, Leon Zhang  
and Xu Zhang

m2r with Chris O'Neill and Jeff Sommars

algstat with Luis Garcia, Rudy Yoshida, ...

bertini with Grant Innerst

...?



# Thank you!!

[www.kahle.io](http://www.kahle.io)

<https://github.com/dkahle/2017-SIAM-Talk>