

# Embrace R in Pharma -

## Building an Internal R Community and Establishing Fit-for-Purpose Pilots

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is becoming increasingly popular in pharma



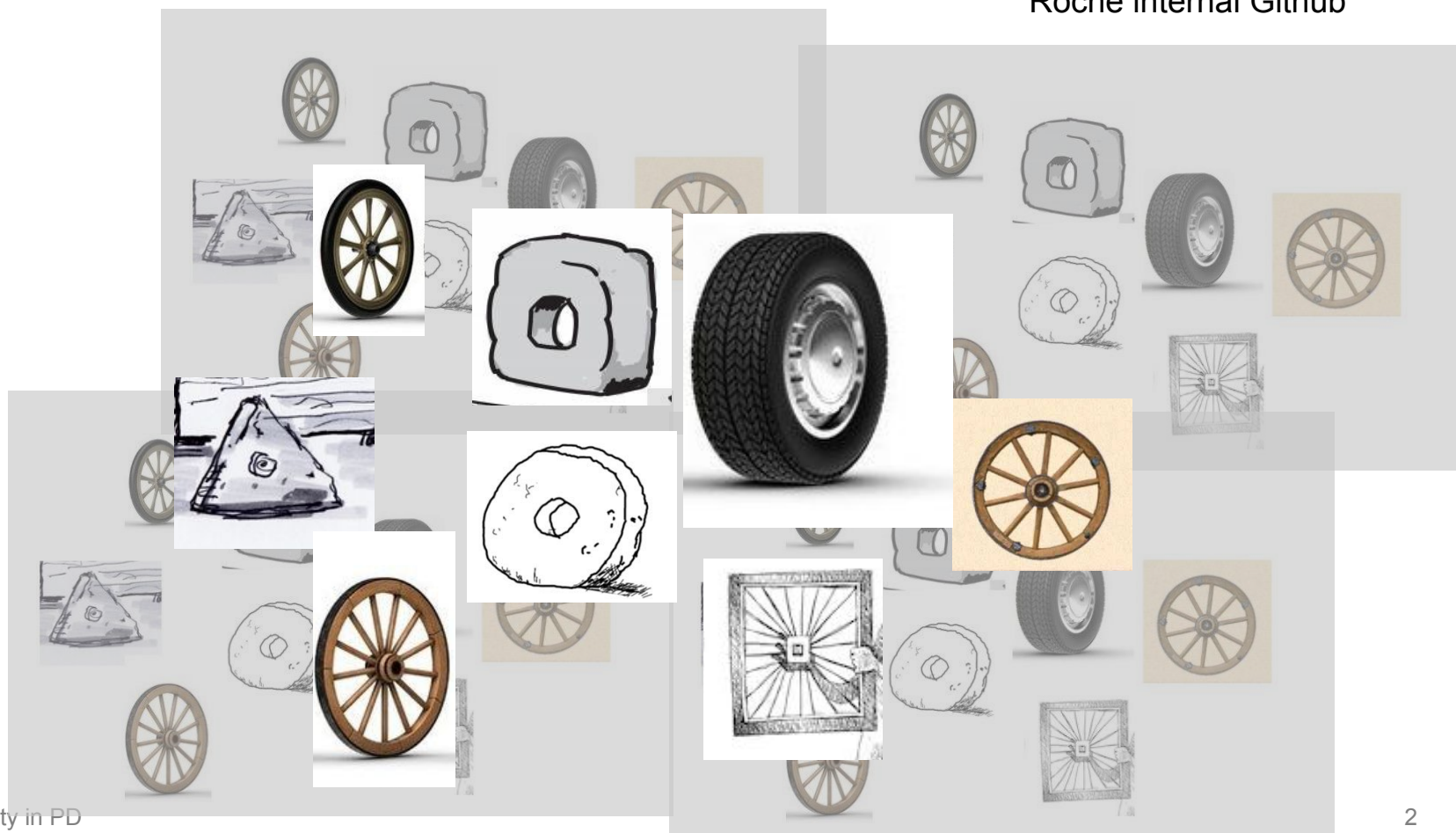
Enterprise

language:R

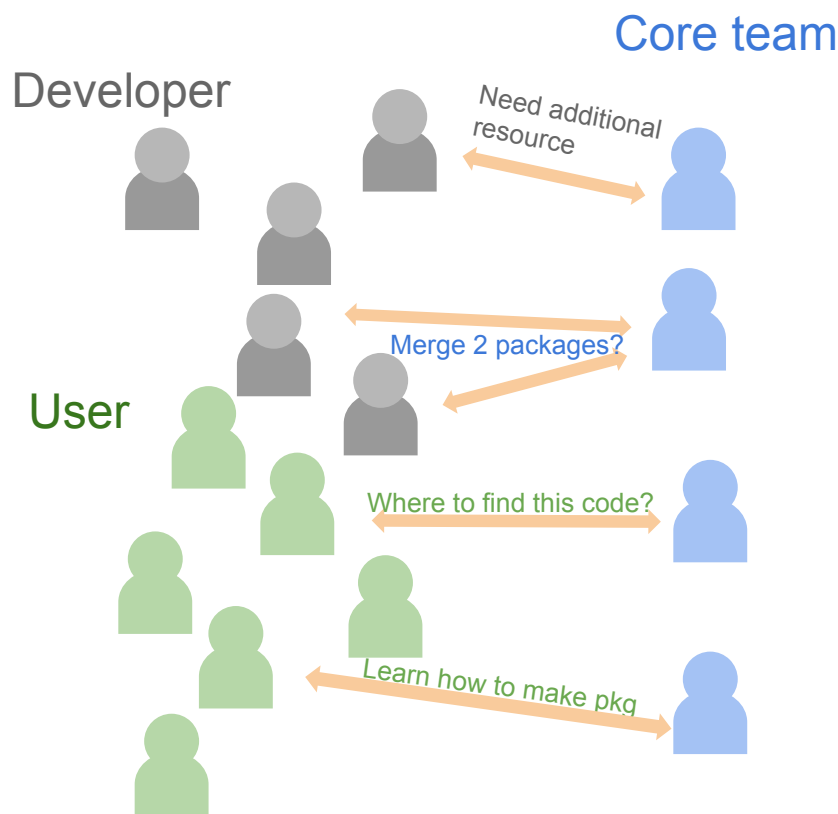
Users

1K

Roche internal Github



# Build an Internal R Community to Connect the Dots & Improve Reusability



Core team: volunteers  
representing different functions  
(statisticians, programmers, data  
scientists, data managers, etc.)

# What We do



## Connect R users/developers and advocate inner-source

R Community Website

Newsletters

Catalog existing tools

Electronic forum

## Further R expertise

Trainings

R user share and learn

Good practice guidelines

## Connect with external stakeholders

Roche/Genentech as the first Pharma company joining R Consortium

## Improve R adoption

Fit-for-purpose R pilots

# Example Pilot

## Project Sautoir: Semi-AUTOmated Interpretation and Report

On behalf of Sautoir Team: Heng Wang, Shengchun Kong, Paul Paczuski, Dominik Heinzmann, Antoine Francois

### The Need

- Biomarker analysis of >10 studies within one molecule (Post-Marketing Commitment)
  - Multiple outputs for each combination of:
    - >10 studies
    - 3 endpoints (OS, PFS, Response)
    - 2 modeling methods
    - >1 missing data imputation methods
  - Plus a comprehensive report

### The Solution

- “One-click” workflow with automated report writing



# Project Sautoir: “One-click” Workflow Implementation

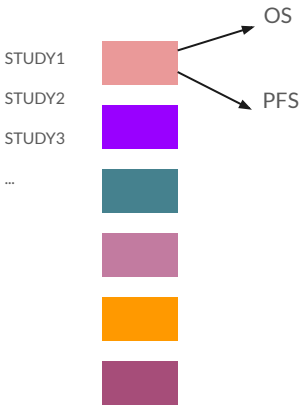
## metadata

name	study	endpoint	...
STUDY1	GO12345	OS, PFSINV	...
STUDY2	GO12346	OS, PFSINV	...

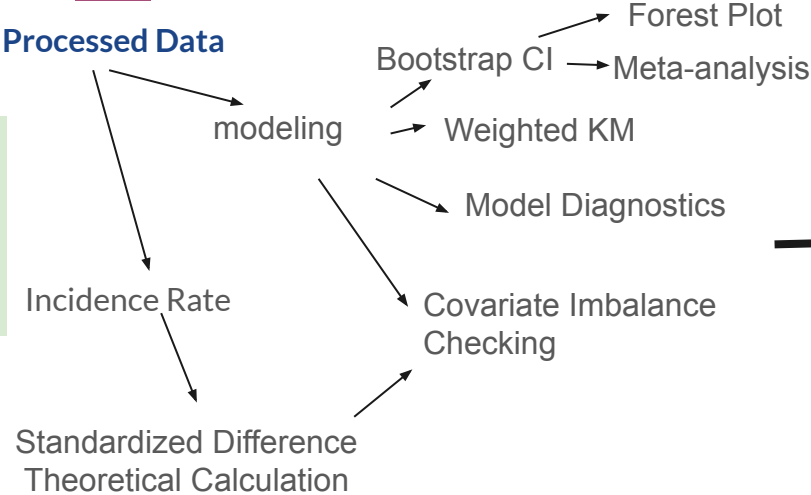
## Data



Sautoir::  
auto\_process()



## Processed Data



- Developed 3 R packages
- PropensitySub: analysis modules
  - gofaiWrites: auto-writing modules
  - Sautoir: Workflow package

Sautoir::auto\_analysis()  
Backend: PropensitySub, tern

## Auto Writing

OS and PFS landmark analysis template  
02 March, 2019

1 POPLAR

1.1 POPLAR OS

1.1.1 option1

1.1.1.1 POPLAR OS option1 : Sample Size

	Control	ADA-	ADA+
unadjusted	120	85	50
adjusted	122	81	48

1.1.1.2 POPLAR OS option1 : Variables in model  
EGORR + SEX + TBRK + METSTES + BALBUN + BLDH + BASLD + RACE1 + AGE55 + ENLR + BWL + HST +  
TC1C1 + LNER + PPROTY + BOP

1.1.1.3 POPLAR OS option1 : Summary Statistics and CI

Sautoir::report.Rmd  
Backend: gofaiWrites

- Meta data driven pipeline
- Automated writing
- Github workflow/project board
- Regression test to ensure backward compatibility (to “ground truth”)
- Make file & CI (ongoing)

# Project Sautoir: Current Stage

- Submitted multiple reports to FDA
  - Auto-generated languages incorporated
  - Submitted R packages for esub
- R package esub challenges
  - All program files are required to be in .txt, file names need to be in lower case
  - fun.R -> fun.r.txt; DESCRIPTION -> description.txt



STRONGER TOGETHER

*Doing now what patients need next*



*Doing now what patients need next*