RAMP DATA CHALLENGES WITH MODULARIZATION AND CODE SUBMISSION

BALÁZS KÉGL Université Paris-Saclay / CNRS



WHO AM !? Balázs Kégl

- Senior researcher CNRS
 - machine learning (20 years)
 interfacing with particle physics (10 years)
- Head of the Paris-Saclay Center for Data Science
 - interfacing with biology, economy, climatology, chemistry, etc. (4 years)





A multi-disciplinary initiative, building interfaces, matching people, helping them launching projects

345 affiliated researchers, 50 laboratories

Biology & bioinformatics

IBISC/UEvry LRI/UPSud Hepatinov CESP/UPSud-UVSO-Inserm IGM-I2BC/UPSud MIA/Agro MIAj-MIG/INRA LMAS/Centrale

Chemistry EA4041/UPSud

Earth sciences LATMOS/UVSO GEOPS/UPSud **IPSL/UVSQ** LSCE/UVSQ LMD/Polytechnique

Economy

LM/ENSAE RITM/UPSud LFA/ENSAE

Neuroscience

UNICOG/Inserm U1000/Inserm NeuroSpin/CEA

Particle physics astrophysics & cosmology

LPP/Polytechnique **DMPH/ONERA** CosmoStat/CEA IAS/UPSud AIM/CEA LAL/UPSud

Machine learning

LRI/UPSud LTCI/Telecom CMLA/Cachan LS/ENSAE LIX/Polytechnique MIA/Agro CMA/Polytechnique LSS/Supélec CVN/Centrale LMAS/Centrale DTIM/ONERA IBISC/UEvry LIST/CEA Visualization

INRIA LIMSI

Signal processing

LTCI/Telecom CMA/Polytechnique CVN/Centrale LSS/Supélec CMLA/Cachan LIMSI DTIM/ONERA

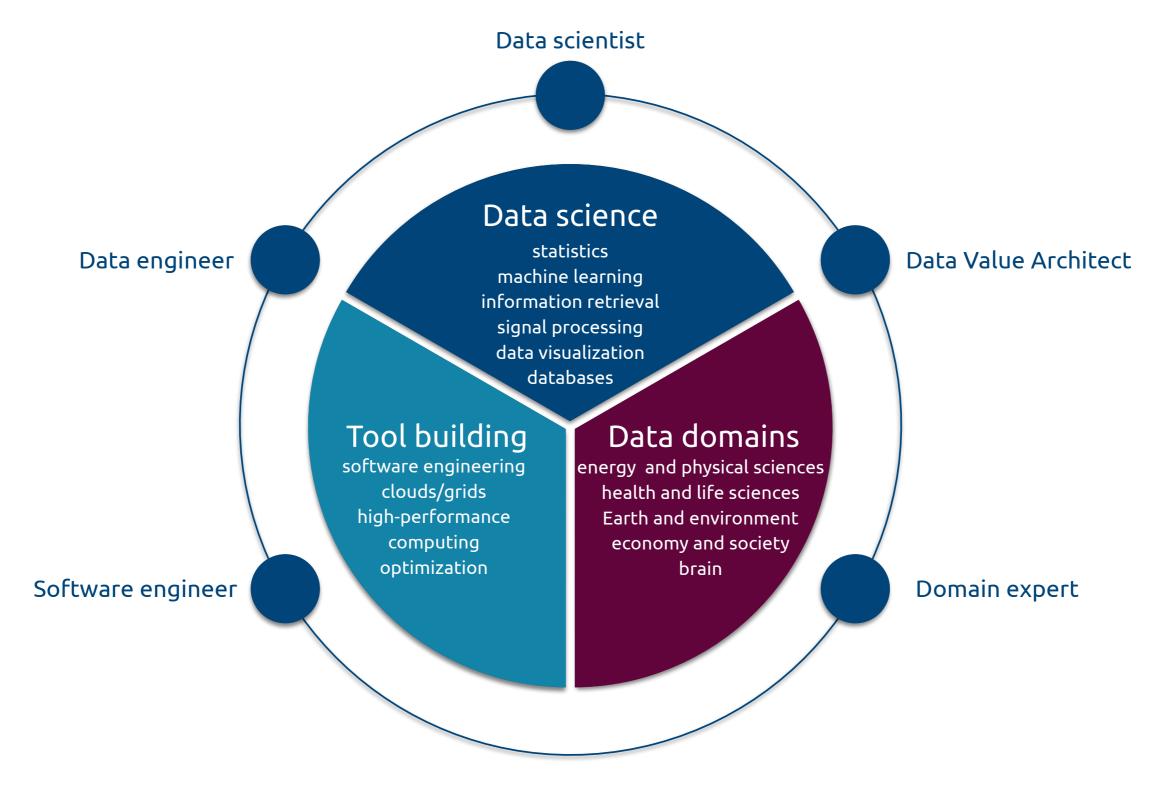
Statistics

LMO/UPSud LS/ENSAE LSS/Supélec CMA/Polytechnique LMAS/Centrale MIA/AgroParisTech



THE DATA SCIENCE ECOSYSTEM

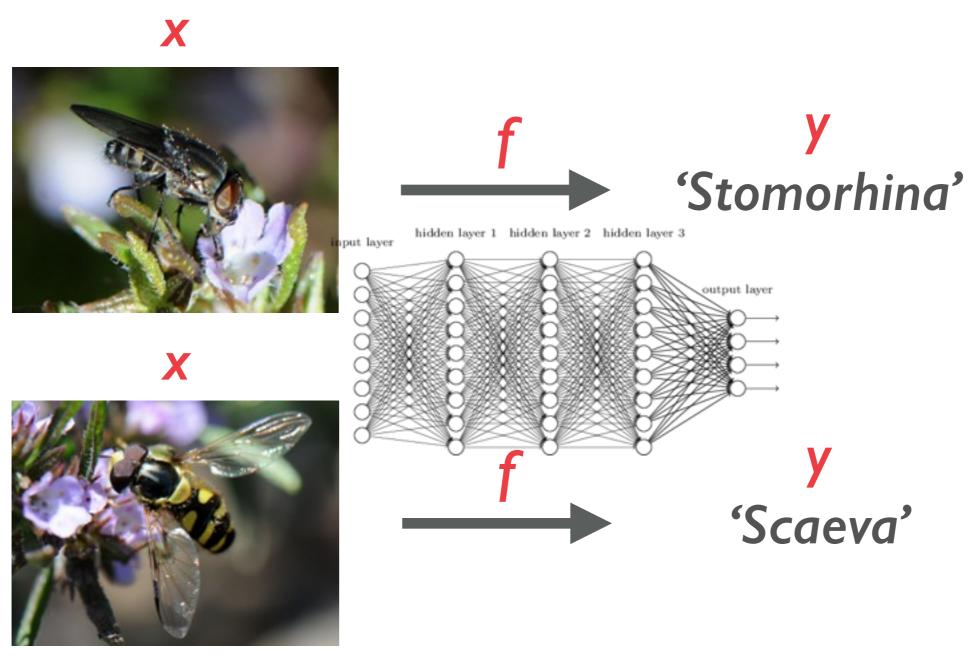
https://medium.com/@balazskegl





What does machine learning do

• Classification problem y = f(x)



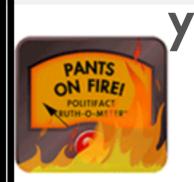
What does machine learning do



FREEDOMJUNKSHUN.COM

Says "FBI and Texas State Police are both confirming that Raymond Peter Littleberry, the man accused of shooting up a Texas church, was an avid atheist on the payroll of the Democrat National Committee."

PolitiFact National on Monday, November 6th, 2017



Not on DNC payroll



DAVID PERDUE

The diversity visa lottery is "plagued by fraud."

PolitiFact National on Monday, November 6th, 2017



Some fraud rings extract money from lottery winners



BARACK OBAMA

"Eight in 10 people this year can find plans for \$75 a month or less."

PolitiFact National on Monday, November 6th, 2017



Under 4 percent of Americans

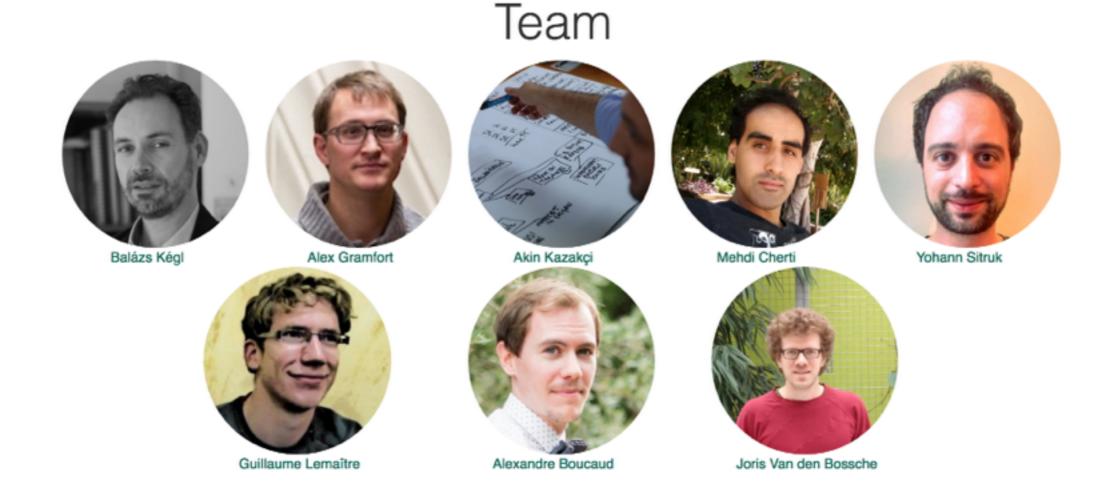
ML IN NATURAL SCIENCES

- Inference: to invert the generative model
 - "predict" a particle, detect an anomaly, infer a parameter y, from observation x
- Generation: to replace expensive simulations
 - "learn" a GEANT4 simulation with a neural net
- Hypothesis generation: to replace theoreticians:)
 - learn, represent structural knowledge and generate novelty in model space, e.g., molecule generation in drug discovery

RAMP is a tool for

- I. Collaborative prototyping
- 2. Teaching aid
- 3. Data science process management

Funded by Université Paris-Saclay and CNRS



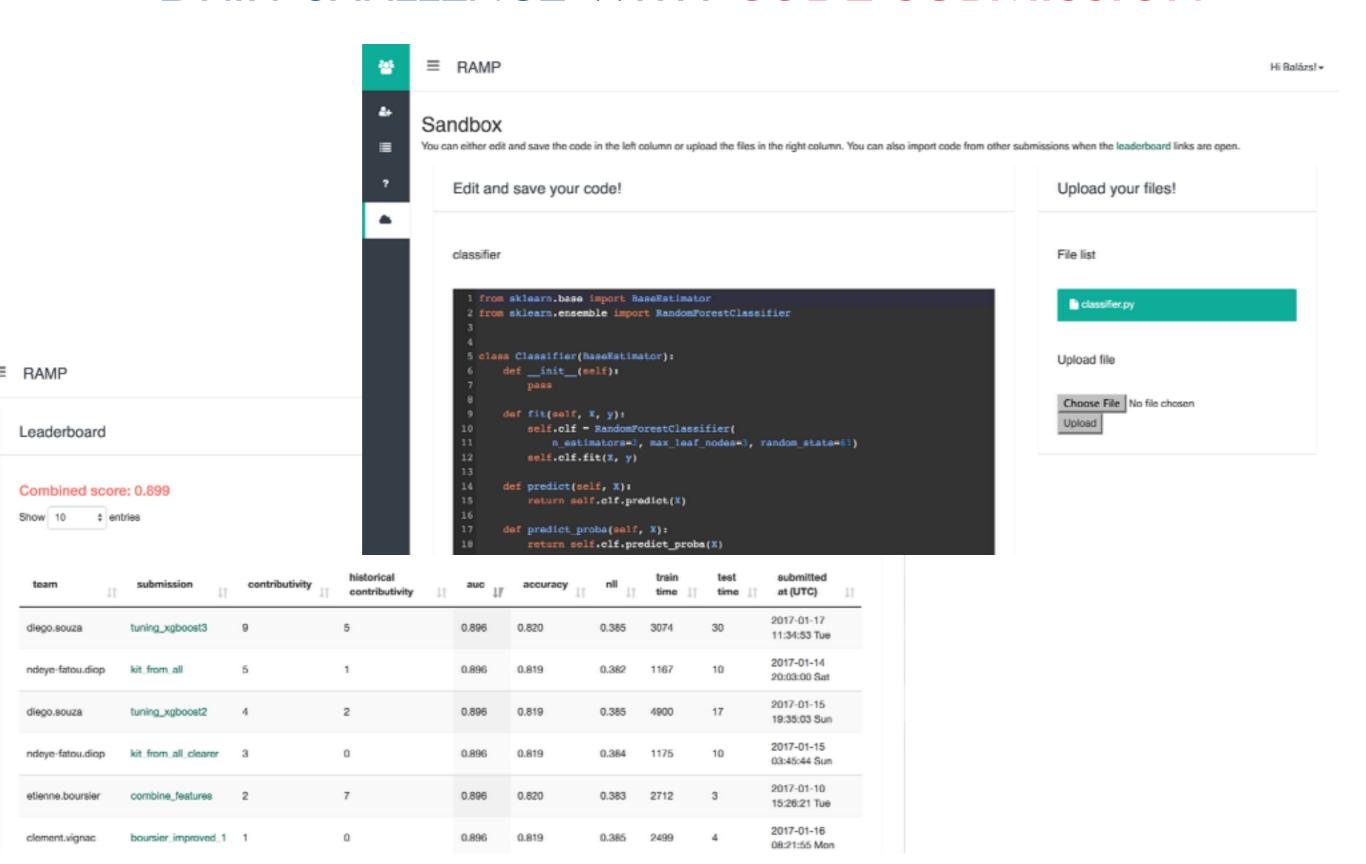
Alumni



Djalel Benbouzid

RAMP.STUDIO

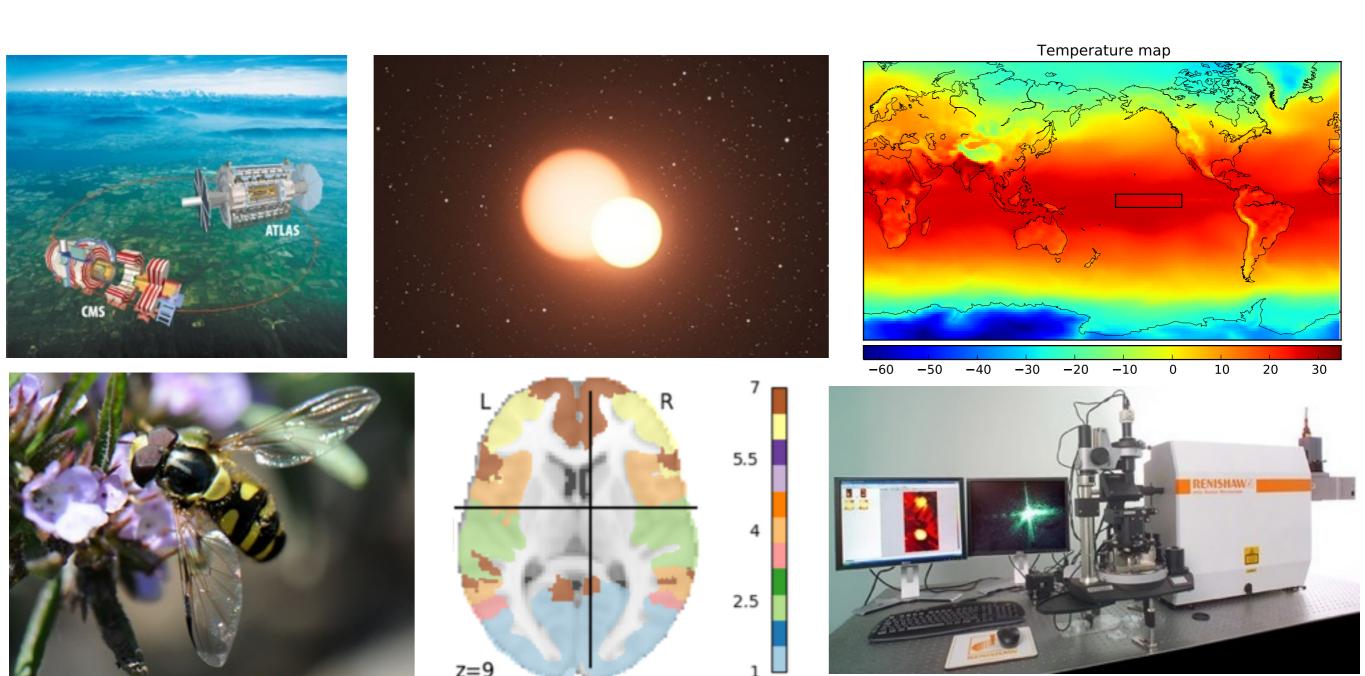
DATA CHALLENGE WITH CODE SUBMISSION



Code submission

- 1. lets us deliver a working prototype
- 2. lets the participants collaborate
- 3. makes the backend challenging to run (cloud management)

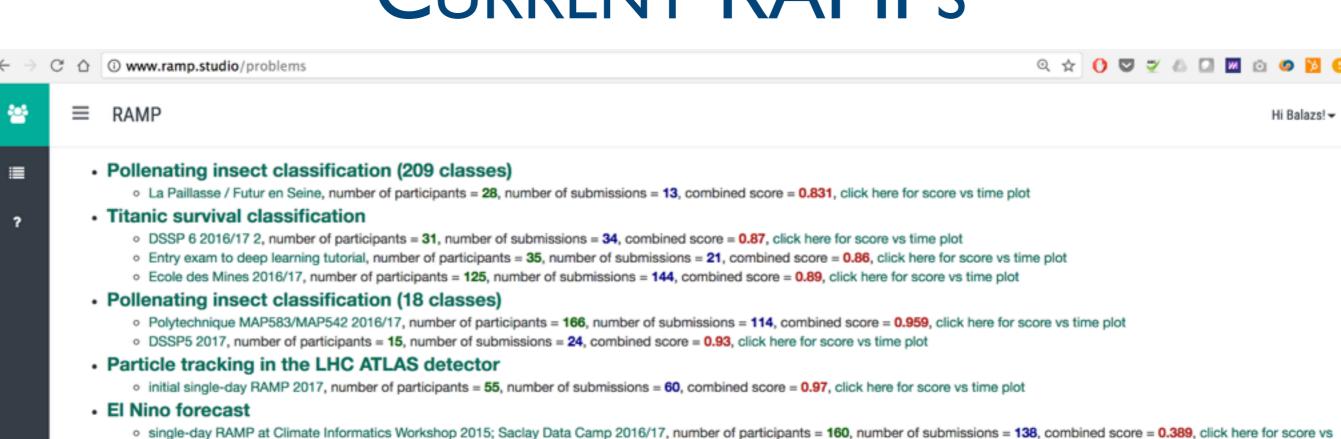
RAPID ANALYTICS AND MODEL PROTOTYPING



functional data, time series, data augmentation, deep learning, learning on simulations, nonstandard and multi-objective losses



CURRENT RAMPS



· Arctic sea ice forecast

time plot

- single-day RAMP at Climate Informatics Workshop 2016, number of participants = 46, number of submissions = 83, combined score = 0.31, click here for score vs time plot
- Polytechnique MAP542 2016/17, number of participants = 20, number of submissions = 52, combined score = 0.268, click here for score vs time plot
- Polytechnique MAP583 2016/17, number of participants = 123, number of submissions = 252, combined score = 0.259, click here for score vs time plot

· Number of air passengers prediction

- DSSP4/5 2016, number of participants = 95, number of submissions = 242, combined score = 0.236, click here for score vs time plot
- DSSP6 2017, number of participants = 23, number of submissions = 59, combined score = 0.268, click here for score vs time plot

· Drug classification and concentration estimation from Raman spectra

- Polytechnique MAP583 2016/17, number of participants = 125, number of submissions = 258, combined score = 0.048, click here for score vs time plot
- o initial single-day RAMP 2016; Saclay Data Camp 2016/17, number of participants = 242, number of submissions = 554, combined score = 0.027, click here for score vs time plot
- Ecole des Mines 2016/17, number of participants = 124, number of submissions = 560, combined score = 0.023, click here for score vs time plot

Detecting anomalies in the LHC ATLAS detector

- Polytechnique MAP542 2016/17, number of participants = 29, number of submissions = 47, combined score = 0.865, click here for score vs time plot
- Polytechnique MAP583 2016/17, number of participants = 133, number of submissions = 275, combined score = 0.899, click here for score vs time plot
- initial single-day RAMP 2016, number of participants = 49, number of submissions = 19, combined score = 0.677, click here for score vs time plot

Epidemium cancer mortality rate prediction (2nd RAMP)

- initial single-day RAMP 2016, number of participants = 39, number of submissions = 46, combined score = 21.79, click here for score vs time plot
- Polytechnique MAP583 2016/17, number of participants = 128, number of submissions = 192, combined score = 18.59, click here for score vs time plot
- Polytechnique MAP542 2016/17, number of participants = 22, number of submissions = 57, combined score = 19.31, click here for score vs time plot

DATA SCIENCE THEMES

C' 1

① www.ramp.studio/data_science_themes



RAMP

Data science themes

classification

- Iris classification
- Detecting anomalies in the LHC ATLAS detector
- Drug classification and concentration estimation from Raman spectra
- Titanic survival classification
- Pollenating insect classification (18 classes)
- Pollenating insect classification (209 classes)

convolutional networks

- Pollenating insect classification (18 classes)
- Pollenating insect classification (209 classes)

external data

Number of air passengers prediction

feature engineering

- El Nino forecast
- Arctic sea ice forecast
- Drug classification and concentration estimation from Raman spectra
- Detecting anomalies in the LHC ATLAS detector

forests

- Iris classification
- Detecting anomalies in the LHC ATLAS detector
- Titanic survival classification
- · Boston housing price regression
- El Nino forecast
- Arctic sea ice forecast
- Number of air passengers prediction
- Epidemium cancer mortality rate prediction (2nd RAMP)

functional data

Drug classification and concentration estimation from Raman spectra

image data

- Pollenating insect classification (18 classes)
- o Pollenating insect classification (209 classes)
- El Nino forecast

missing data

RAMP

- Epidemium cancer mortality rate prediction (2nd RAMP)
- Titanic survival classification

neural networks (deep learning)

- · Drug classification and concentration estimation from Raman spectra
- Pollenating insect classification (18 classes)
- Pollenating insect classification (209 classes)

regression

- Boston housing price regression
- El Nino forecast
- Arctic sea ice forecast
- Number of air passengers prediction
- Drug classification and concentration estimation from Raman spectra
- Epidemium cancer mortality rate prediction (2nd RAMP)

· small data

- Drug classification and concentration estimation from Raman spectra
- Epidemium cancer mortality rate prediction (2nd RAMP)
- o Detecting anomalies in the LHC ATLAS detector
- El Nino forecast
- Arctic sea ice forecast
- Number of air passengers prediction
- · Particle tracking in the LHC ATLAS detector

supervised clustering (unsupervised classification)

· Particle tracking in the LHC ATLAS detector

tabular data

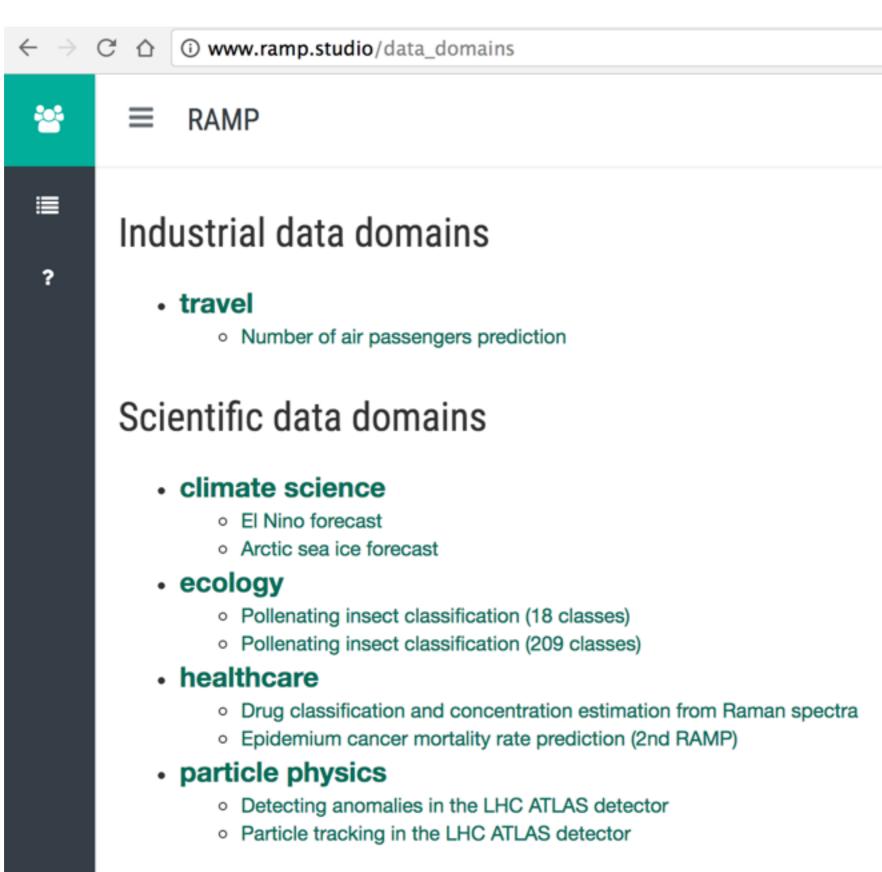
- Iris classification
- · Detecting anomalies in the LHC ATLAS detector
- Titanic survival classification
- · Boston housing price regression
- · Number of air passengers prediction
- · Epidemium cancer mortality rate prediction (2nd RAMP)

· time series forecasting

- El Nino forecast
- Arctic sea ice forecast

.....

DATA DOMAINS





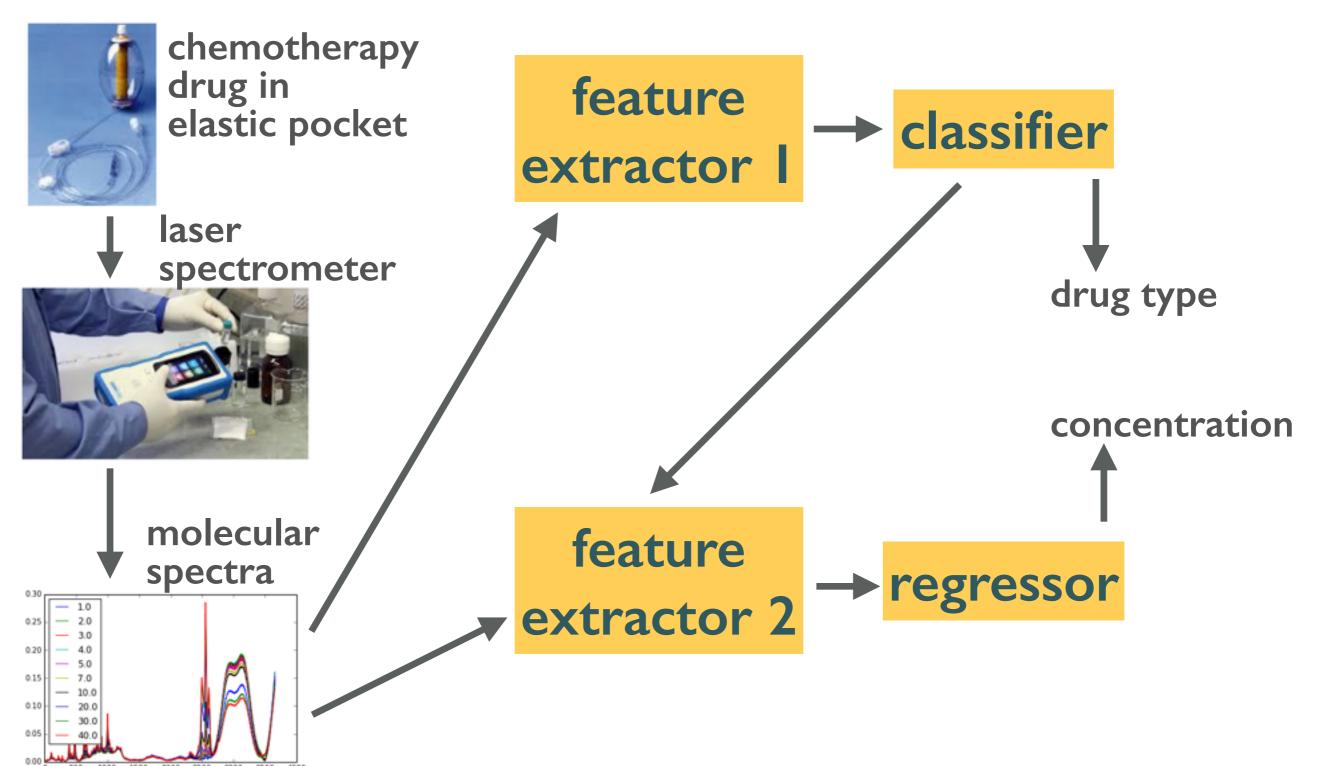
RAMP.STUDIO DATA CHALLENGE WITH CODE SUBMISSION

20+ challenges
40+ events
1200+ users
7000+ predictive models

RAMP.STUDIO DATA CHALLENGE WITH CODE SUBMISSION

12 hackathons6 remote data challenges11 course data camps

CLASSIFYING AND REGRESSING ON MOLECULAR SPECTRA





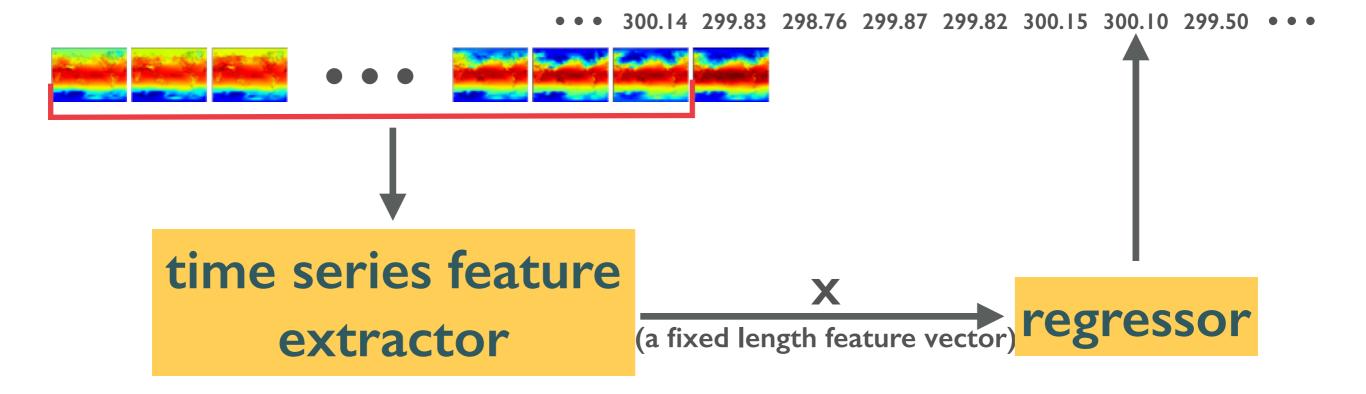
Classifying and quantifying monoclonal antibody preparations for cancer therapy using machine learning

Laetitia Le ^{ab}, Camille Marini ^{ce}, Alexandre Gramfort ^{cfg}, David Nguyen ^a, Mehdi Cherti ^{ch}, Sana Tfaili ^b, Ali Tfayli ^b, Arlette Baillet-Guffroy ^b, Eric Caudron ^{ab}, Balázs Kégl ^{ch}

^a European Georges Pompidou Hospital (AP-HP), Pharmacy department, Paris, France department, Paris, France Lip(Sys) Chimi Analytique Pharmaceutique, Univ. Paris-Sud, Universit Paris Saclay, F92290 Chatenay-Malabry, France (EA4041 Groupe de Chimie Analytique de Paris Sud)
^c Center of Data Science, Université Paris-Saclay de Université Paris-Sud e CMAP, Ecole Polytechnique, Palaiseau, France f INRIA, Parietal team, Saclay, France gent EtAL, CNRS, France

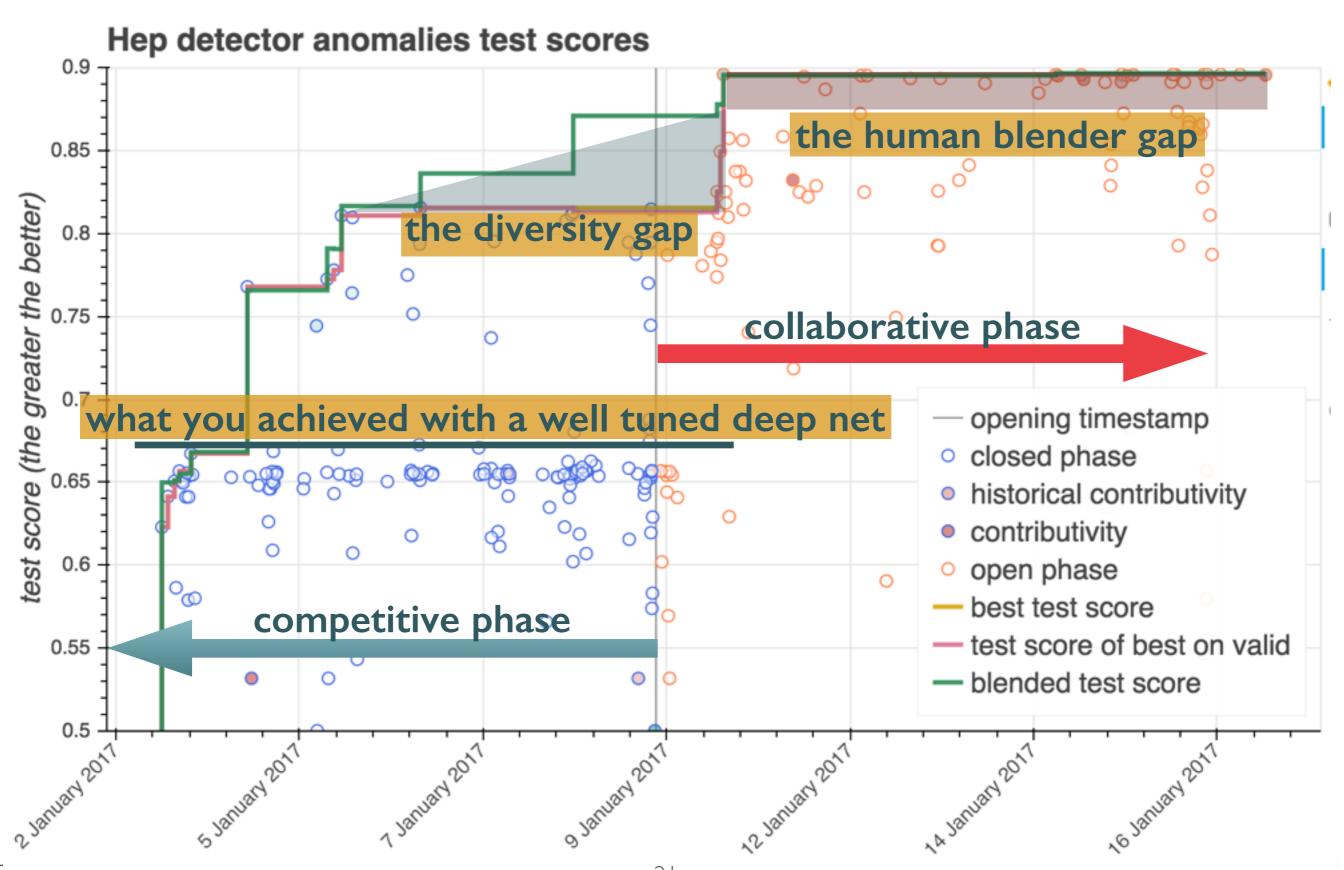


FORECASTING EL NINO SIX MONTHS AHEAD

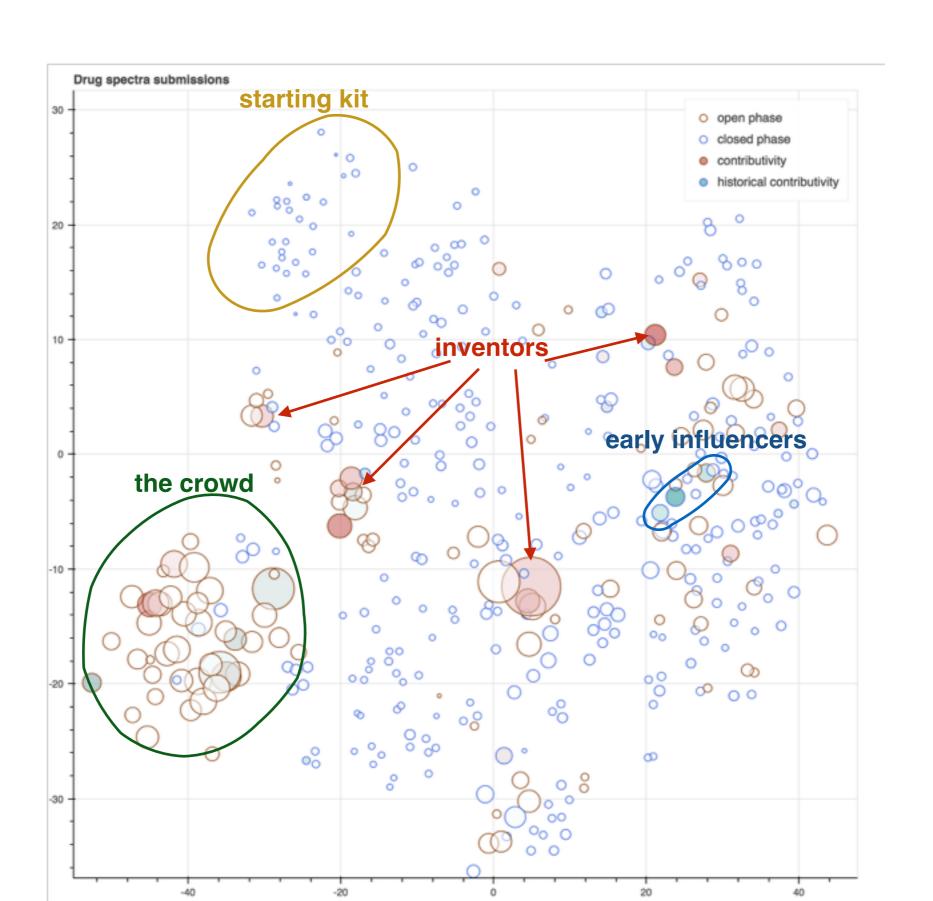




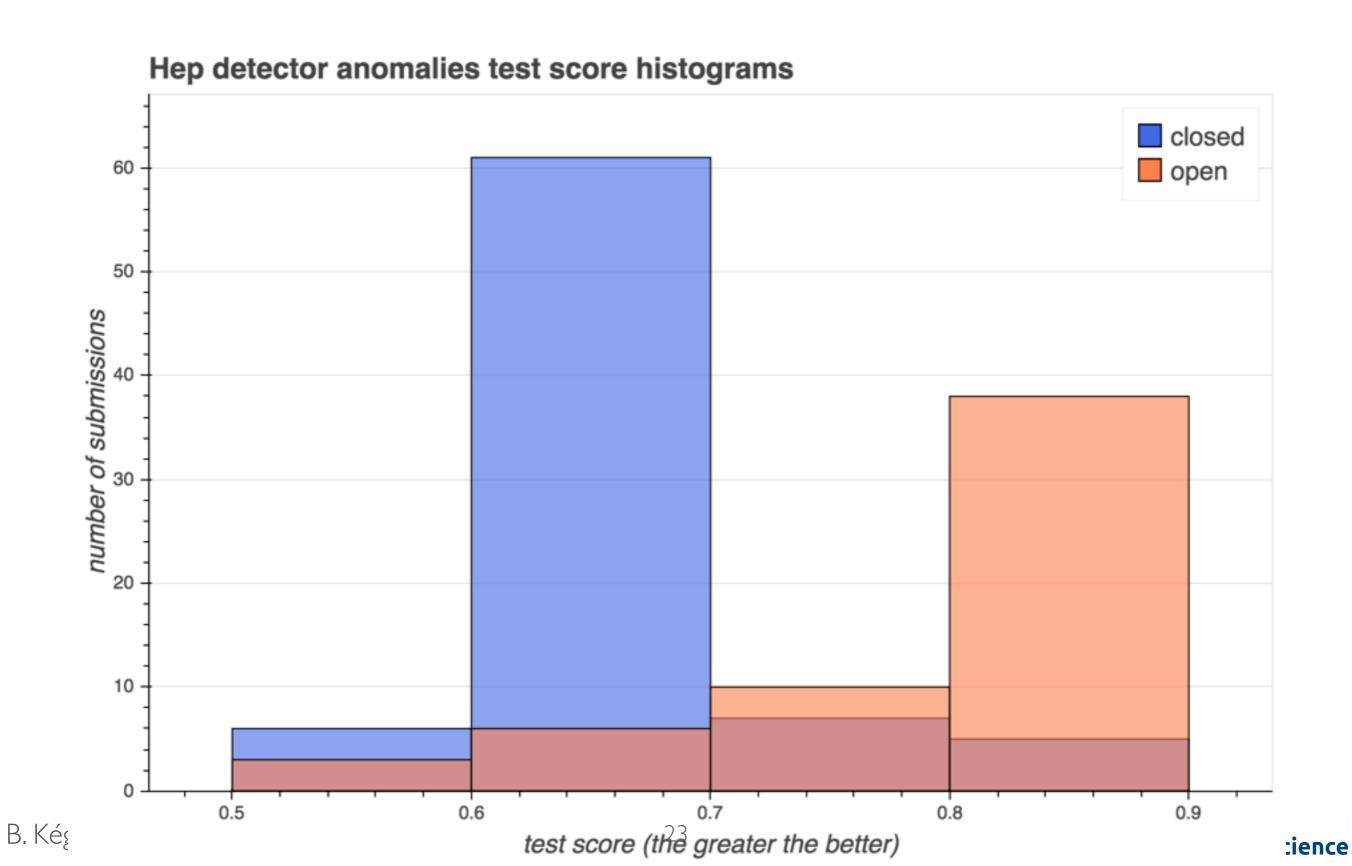
THE POWER OF THE (COLLABORATING) CROWD



THE DYNAMICS OF COLLABORATION

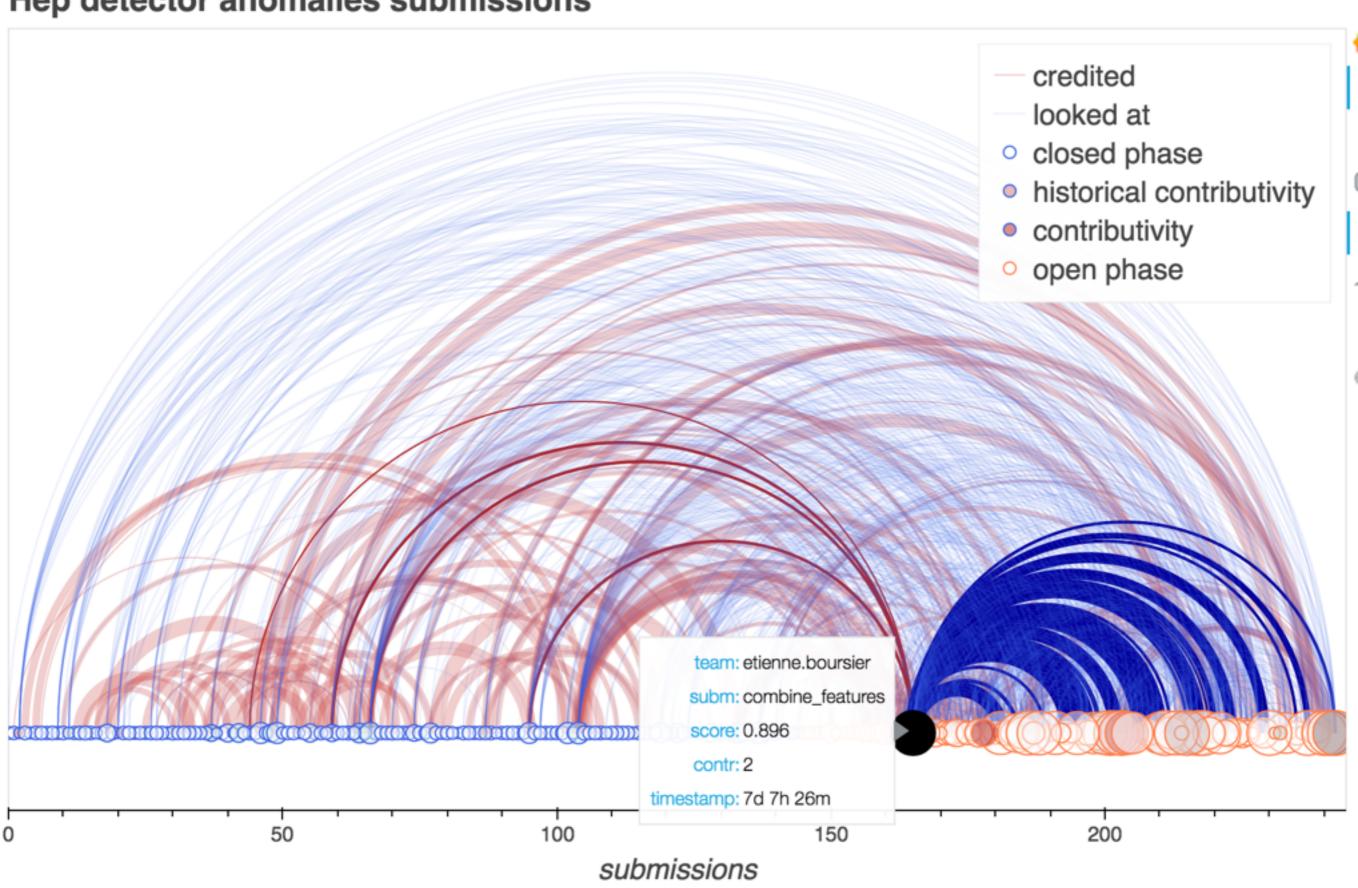


OPEN PHASE LETS PARTICIPANTS CATCH UP THE GOAL OF TEACHING



COMMUNICATION AND REUSE

Hep detector anomalies submissions



CLASSIFY POLLENATING INSECTS

https://www.ramp.studio/events/pollenating_insects_3_JNI_2017

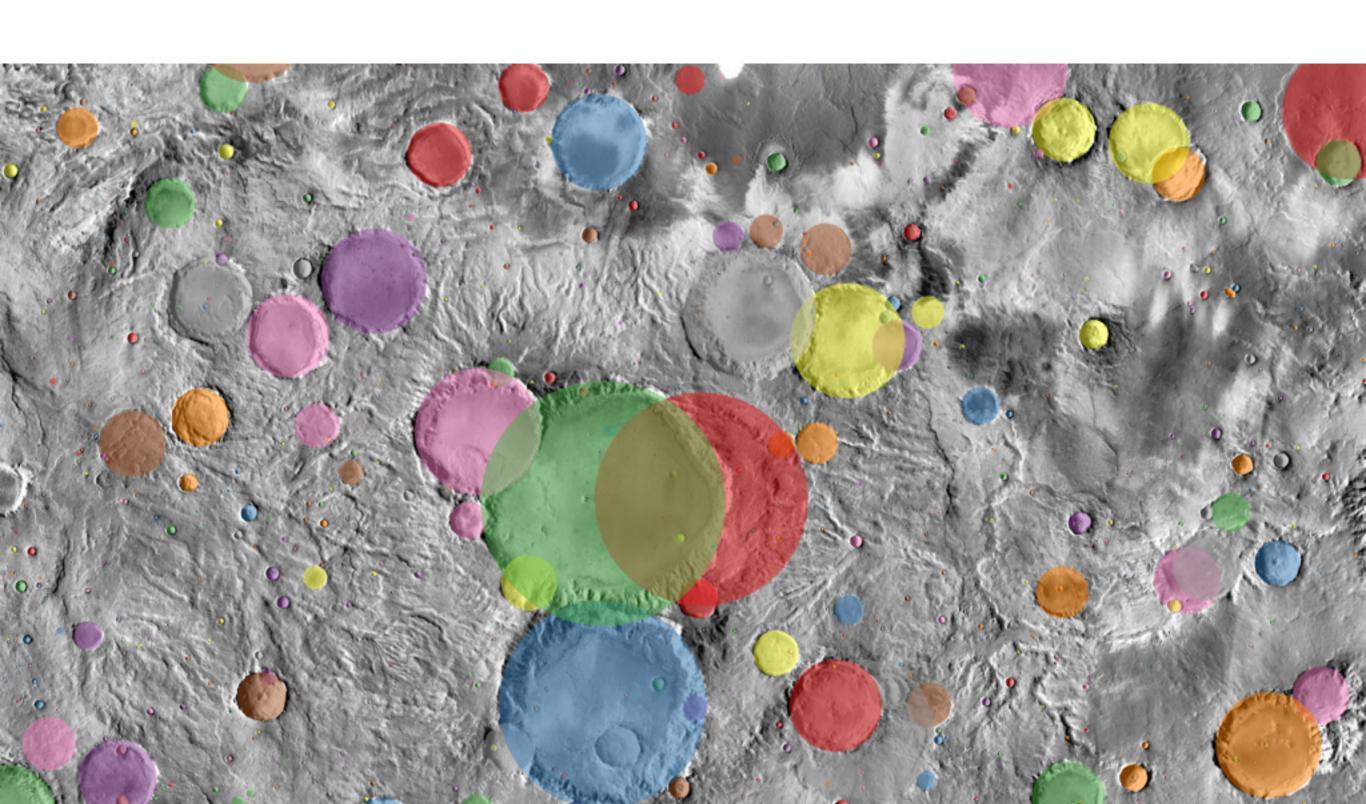
4.5K€ for the competitive phase 3K€ for the collaborative phase 50 GPU hours per participant







DETECTING MARS CRATERS



FAKE NEWS

PREDICT THE TRUTHFULNESS OF NEWS



EDITIONS V

TRUTH-O-METER™ ∨

PEOPLE V

PROMISES V

PANTS ON FIRE

ABOUT US



Hillary Clinton, Russia, and uranium: What you need to know

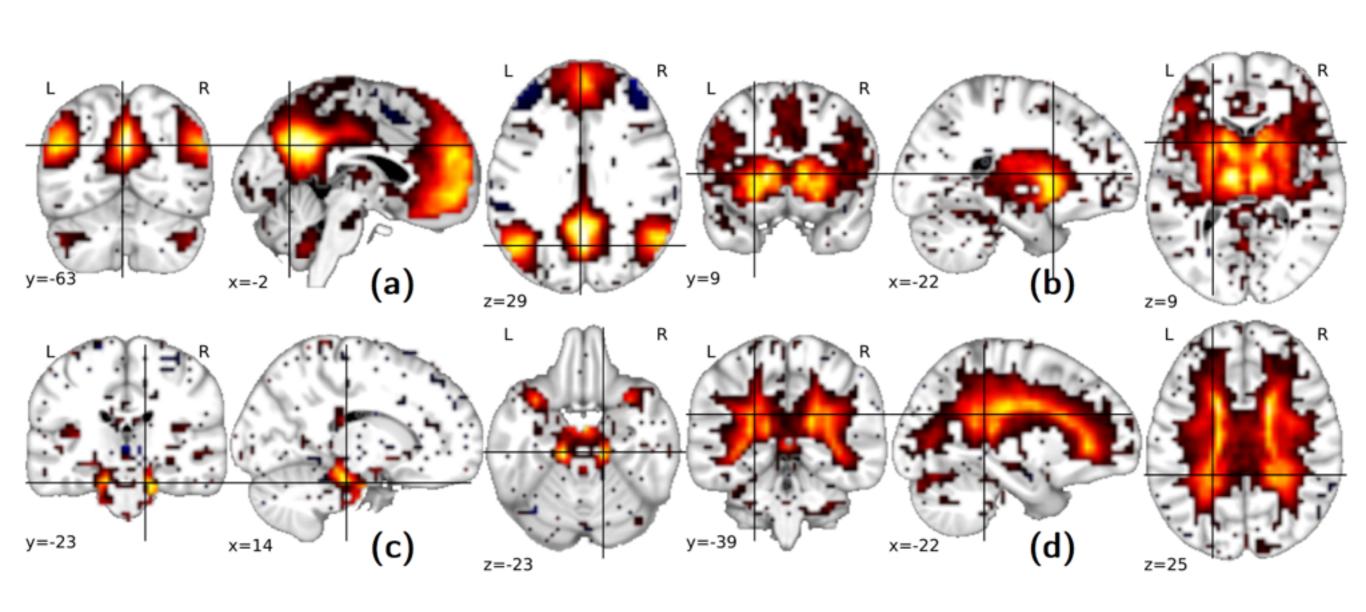


Can Americans expect a \$4,000 "raise" from Trump tax plan?

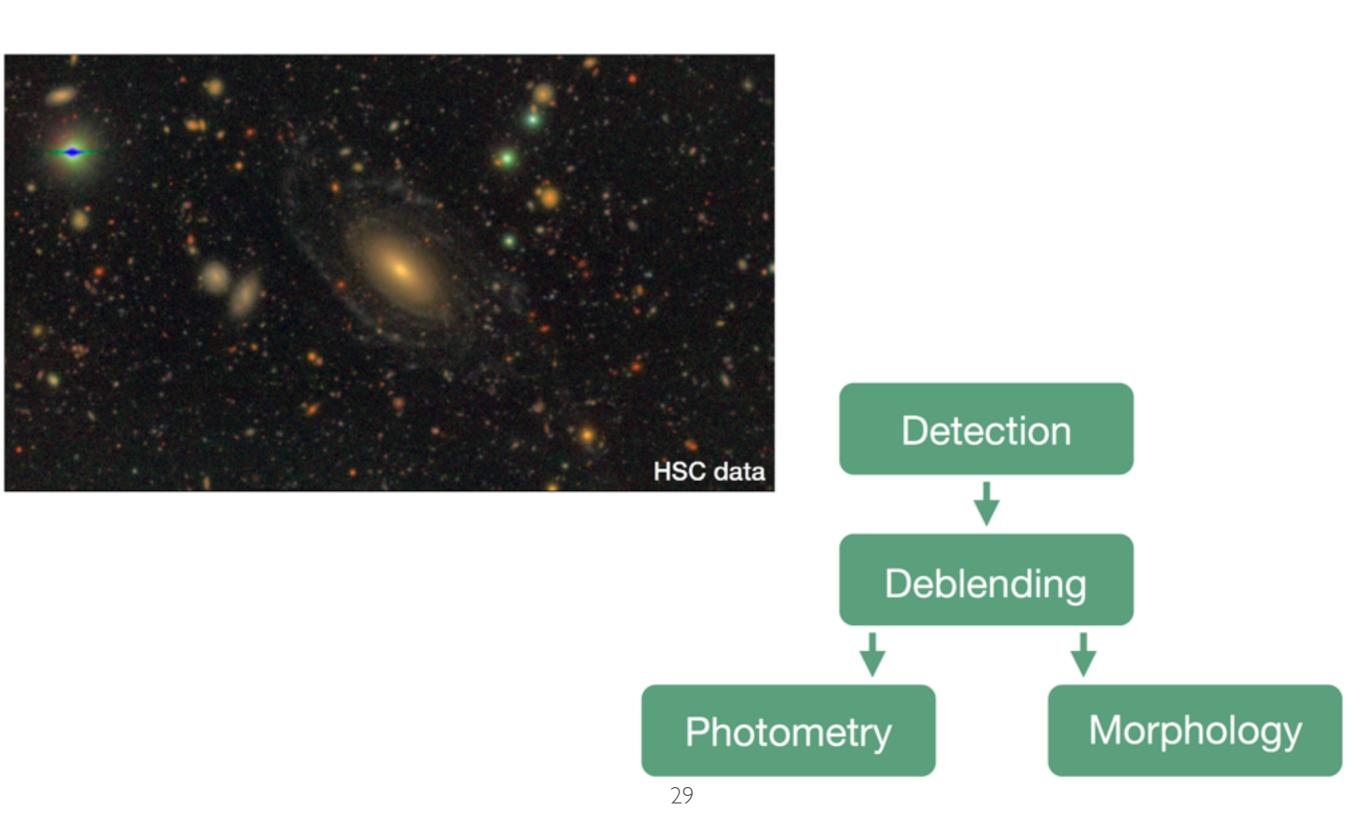


The big picture: Niger and what we know about what happened

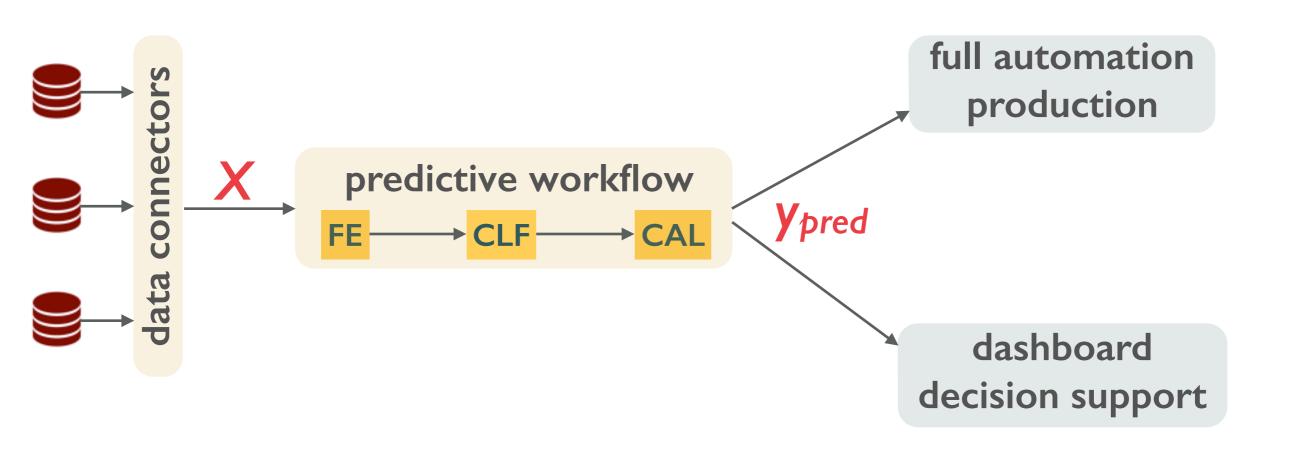
UPCOMING CHALLENGE PREDICT AUTISM FROM BRAIN SCANS



UPCOMING CHALLENGE(S) ASTROIMAGING



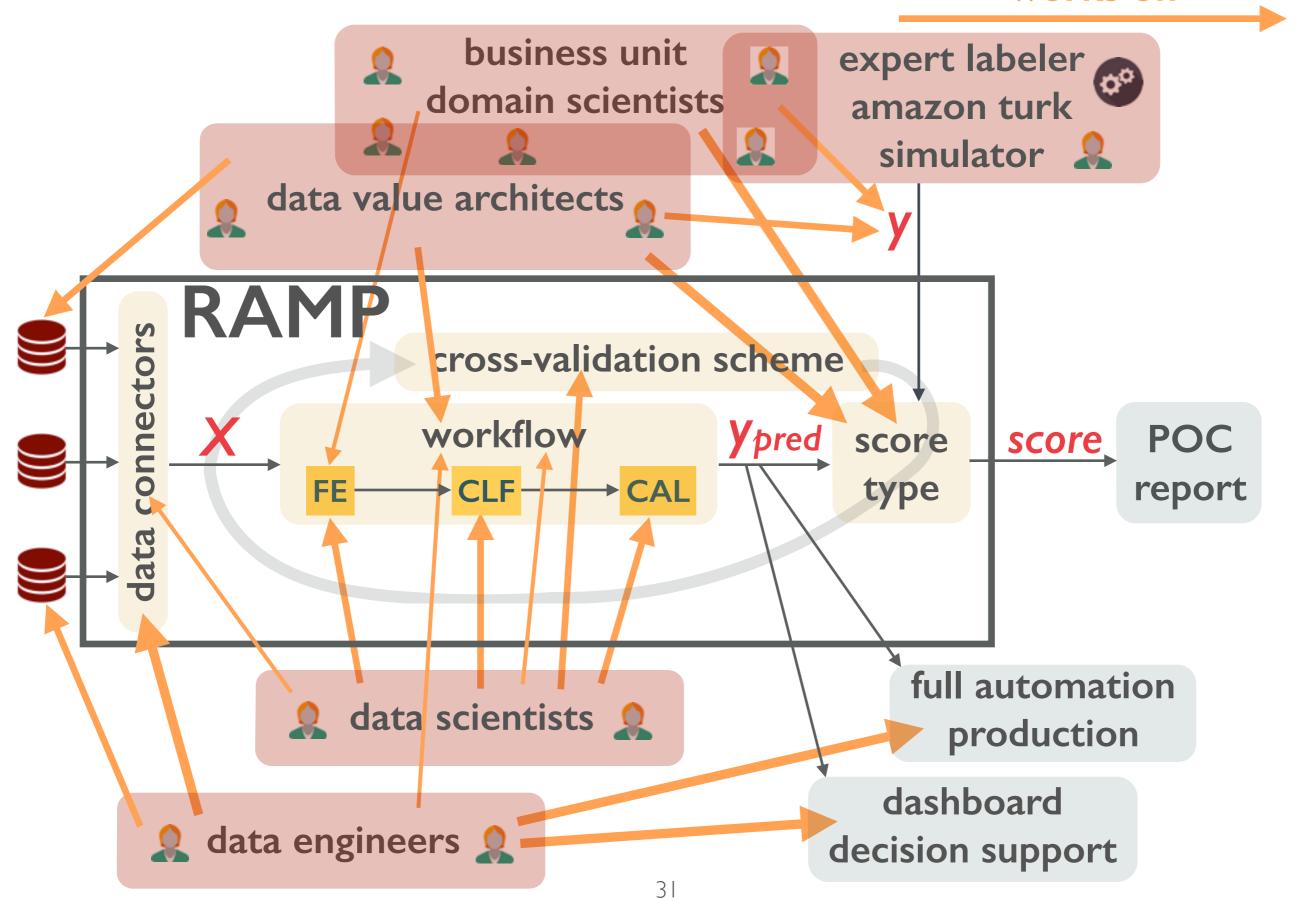
THE DATA FLOW



THE IDEAL SEQUENCE

data flow

"works on"



Before solving the problem, set it up (even put it into production)

Setting up the RAMP is was long and hard.

Separate workflow building and workflow optimization

RAMP

DATA-CHALLENGES-WITH

MODULARIZATION-AND-CODE-SUBMISSION

FRUGAL DATA SCIENCE
PROCESS MANAGEMENT

BALÁZS KÉGL

Université Paris-Saclay / CNRS



RAMP-WORKFLOW & RAMP-KITS

- toolkit: https://github.com/paris-saclay-cds/ramp-workflow
 - for designing workflows
 - set of ready-made metrics, workflows, CV schemes, data readers
 - unique command-line test script
- examples: https://github.com/ramp-kits
 - a zoo of problems, experiments, workflows
 - (at least) one initial solution

A SINGLE SCRIPT TO DEFINE THE BUNDLE

```
1 import os
28
                                                                2 import numpy as np
29 def get cv(X, y):
                                                                3 import pandas as pd
      unique replicates = np.unique(X['replicate'])
30
                                                                4 import rampwf as rw
31
      r = np.arange(len(X))
      for replicate in unique replicates:
32
                                                                6 problem title =
          train_is = r[(X['replicate'] != replicate).values]
33
                                                                      'Cell population identification from single-cell mass cytometry data'
34
          test_is = r[(X['replicate'] == replicate).values]
                                                                8 target column name = 'cell type'
35
          yield train is, test
                                                                  prediction label names = [
36
                                                                      'B-cell Frac A-C (pro-B cells)', 'Basophils', 'CD4 T cells', 'CD8 T cells',
37
                                                                      'CLP', 'CMP', 'Classical Monocytes', 'Eosinophils', 'GMP', 'HSC',
                                                               11
38 def _read_data(path, f_name):
                                                                      'IgD- IgMpos B cells', 'IgDpos IgMpos B cells', 'IgM- IgD- B-cells',
      data = pd.read_csv(os.path.jo.n(path, 'data', f_name))
39
                                                               13
                                                                      'Intermediate Monocytes', 'MEP', 'MPP', 'Macrophages', 'NK cells',
      y_array = data[ target column name]
40
                                                                      'NKT cells', 'Non-Classical Monocytes', 'Plasma Cells', 'gd T cells',
                                                               14
41
      X df = data.drop([ target column name], axis=1)
                                                               15
                                                                      'mDCs', 'pDCs']
      return X df, y array
                                                               16 # A type (class) which will be used to create wrapper objects for y pred
43
                                                               17 Predictions = rw.prediction_types.make_multiclass(
                                                                      label names= prediction label names)
45 def get_trair lata(path='.'):
                                                               19 # An object implementing the workflow
       f name = 'train.csv.gz'
                                                               20 workflow = rw.workflows.FeatureExtractorClassifier()
      return read data(path, f name)
47
48
                                                               22 score_types [
                                                                      rw.scor types.BalancedAccuracy(name='bac', precision=3),
                                                                      rw.score types.Accuracy(name='acc', precision=3),
50 def get test data (path='.'):
                                                               24
                                                                      rw.score types.NegativeLogLikelihood ...me='nll', precision=3),
       f name = 'test.csv.qz'
                                                               25
       return read data(path, f name)
                                                               26 1
                                                        cross-validation scheme
                                                       workflow
                                                                                       y pred
                                                                                                     score
                                                                                                                     score
                                                                                                      type
```

A SINGLE EXECUTABLE TO TEST THE SUBMISSIONS

- Keep your different submissions in a simple file structure
- Communicate them on git
- Execute them also from the notebook

```
silver6:mouse_cytometry kegl$ ramp_test_submission
Testing Cell population identification from single-cell mass cytometry data
Reading train and test files from ./data ...
Reading cv ...
Training ./submissions/starting_kit ...
CV fold 0
        train bac = 0.042
        valid bac = 0.042
        test bac = 0.042
        train acc = 0.427
        valid acc = 0.416
        test acc = 0.396
        train nll = 1.715
        valid nll = 1.71
        test nll = 1.779
CV fold 1
        train bac = 0.042
        valid bac = 0.046
        test bac = 0.042
        train acc = 0.415
        valid acc = 0.453
        test acc = 0.396
        train nll = 1.729
        valid nll = 1.657
        test nll = 1.775
CV fold 2
        train bac = 0.042
        valid bac = 0.042
        test bac = 0.042
        train acc = 0.408
        valid acc = 0.471
        test acc = 0.394
        train nll = 1.738
        valid nll = 1.61
        test nll = 1.772
CV fold 3
        train bac = 0.042
        valid bac = 0.043
        test bac = 0.042
        train acc = 0.448
        valid acc = 0.357
        test acc = 0.396
        train nll = 1.655
        valid nll = 1.915
        test nll = 1.789
train bac = 0.042 \pm 0.0001
train acc = 0.425 ± 0.0152
train nll = 1.709 \pm 0.0325
valid bac = 0.043 \pm 0.0016
valid acc = 0.424 \pm 0.0437
valid nll = 1.723 \pm 0.1165
test bac = 0.042 \pm 0.0001
test acc = 0.395 \pm 0.0006
test nll = 1.779 \pm 0.0062
```

You can

- 1. Participate in upcoming RAMPs
- 2. Use RAMP in teaching or training
- 3. Use the toolkit for your own workflows
- 4. Submit it to us if you want to run a data challenge

frontend:

www.ramp.studio

toolkit:

github.com/paris-saclay-cds/ramp-workflow

examples:

github.com/ramp-kits

slack:

ramp-studio.slack.com

blogs:

medium.com/@balazskegl

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balazs.kegl@gmail.com