MACHINE LEARNING FOR SCIENTIFIC WORKFLOWS MANAGING THE DATA SCIENCE PROCESS

BALÁZS KÉGL

Head of Al research Huawei Paris

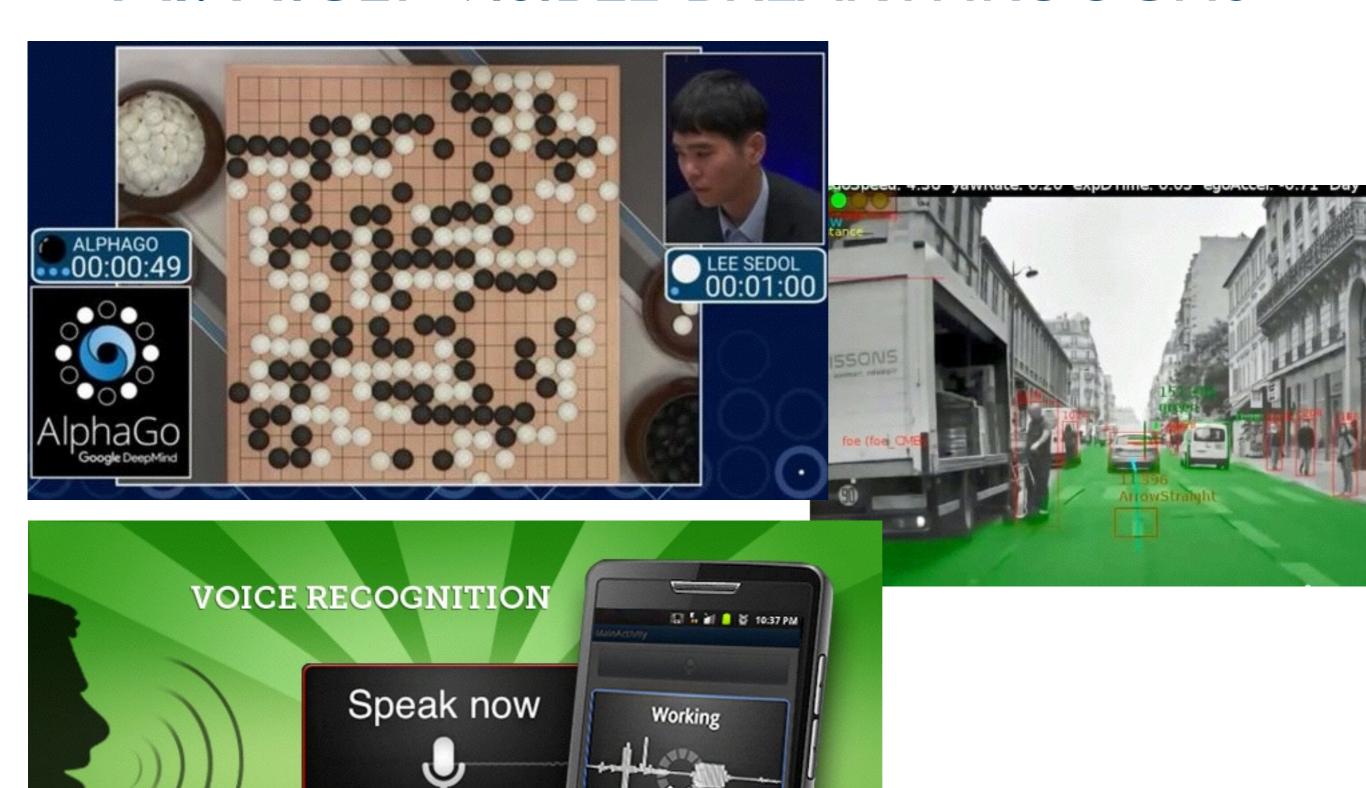
WHO AM I?

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)
 - industrial ML projects (4 years)
- Head of Al research, Huawei Paris
 - interfacing with telecom engineering applications (I year)



Al: HIGLY VISIBLE BREAKTHROUGHS



Cancel

Cancel

Why is the adoption of Al so slow?

THE HUMAN FACTOR

- Adopting Al will change the way we work
 - both the Al "consumer"
 - and the AI developer (engineers and data scientists)
- We have excellent tools to solve problems, but not very little know-how to manage the process
- Designing interfaces
 - formal APIs
 - human/human communication
 - human/Al communication
 - Al/Al communication

OUTLINE

- Machine learning for reusable scientific workflows
 - use cases
 - examples
- Managing the data science process: the RAMP framework
 - roles and tasks
 - building the workflow: who does what
 - what is a predictive workflow, what are the parametrizable components
 - how to make data scientists efficient



WHY IS DATA SCIENCE PROCESS MANAGEMENT RELEVANT IN RESEARCH?

- Typical applied research project
 - take an existing domain-scientific or industrial problem (e.g., galaxy deblending)
 - scan literature
 - install/develop experimental environment
 - collect data and establish benchmark
 - apply existing ML solution
 - optionally fine tune, explore a small number of alternatives
 - show that the ML solution is better than the classical "manual" solution on your own benchmark
 - publish



Takes years, typically

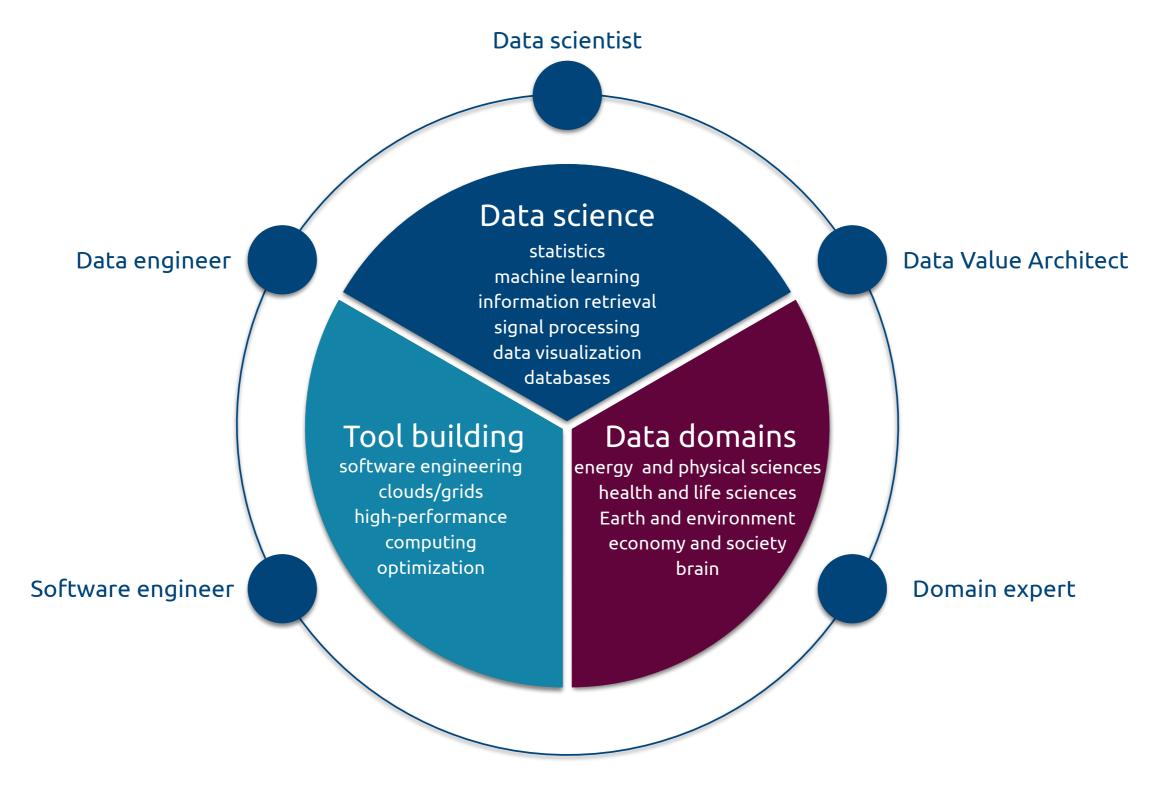
How to accelerate experimental projects?

How to explore a large number of ML solutions in a short time?

How to make solutions not only reproducible but also reusable?

THE DATA SCIENCE ECOSYSTEM

https://medium.com/@balazskegl





ML USE CASES IN SCIENCES

https://www.ramp.studio/problems

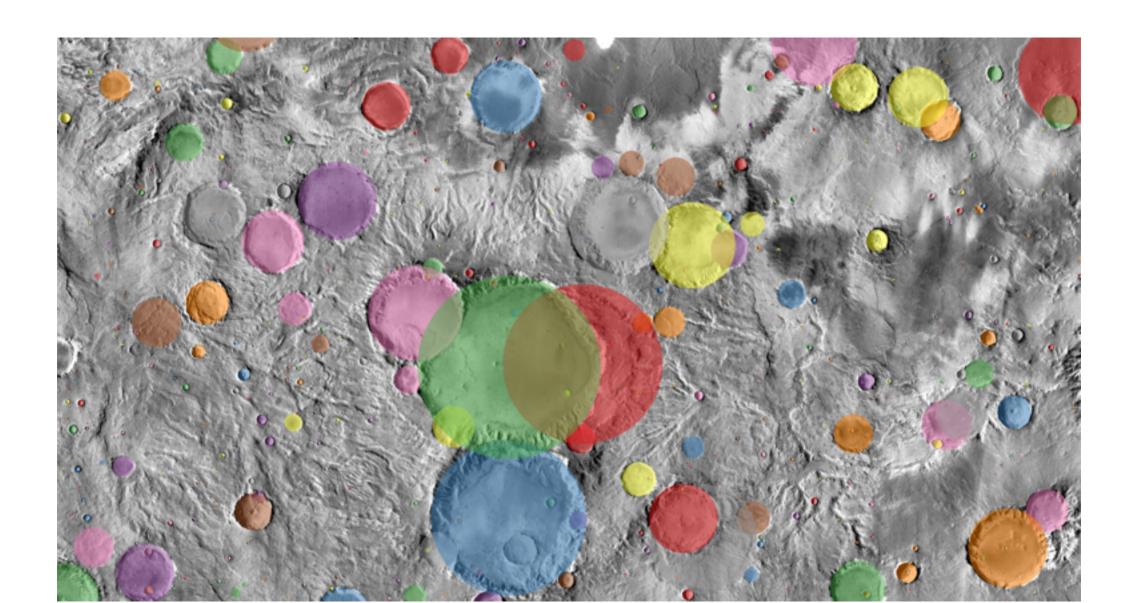
- Data collection: replace human or algorithmic collector or annotator
 - label insect photos, detect Mars craters, detect particle tracks
- Inference: to invert the generative model
 - "predict" a particle, detect an anomaly, infer a parameter y from observation x
- Generation, model reduction: to replace expensive simulations
 - "learn" a physics simulation or an agent based micro-economical model 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



Data collection

DETECTING MARS CRATERS

- collaboration with planetary geologists at Paris-Saclay
- new metrics and workflow
- great benchmark for detection in satellite imagery



DETECTING MARS CRATERS

Saclay MSc students competing and learning from each other while benchmarking state of the art deep learning detection algorithms

| rank 11 | move | † team ↓ | submission | ↓† ap ↓† | train time [s] | test time [s] |
|---------|------|-------------------|----------------------|----------|----------------|---------------|
| 1 | +1 | felix.larrouy | m-rcnn3 | 0.584 | 5947 | 2009 |
| 2 | -1 | nicolas.toussaint | thanks_felix | 0.577 | 6524 | 1964 |
| 3 | - | haquang.le | yolo_v3 | 0.567 | 35400 | 1793 |
| 4 | - | clement.hardy | mask2 | 0.565 | 12130 | 1512 |
| 5 | +1 | alann.cheral | hello_world_7 | 0.538 | 6039 | 2075 |
| 6 | +4 | guillaume.fradet | takeoff_COCO_augment | 0.519 | 11824 | 2098 |
| 7 | -2 | sidali.hamideche | eighth | 0.499 | 27407 | 285 |
| 8 | -1 | manon.cesaire | mrcnn_3 | 0.476 | 11734 | 1463 |
| 9 | - | hao.liu | NASA-V | 0.434 | 16140 | 200 |
| 10 | -2 | enzo.terreau | ssd_class_1 | 0.408 | 2551 | 125 |





Inference

Learning to discover: the Higgs boson machine learning challenge



Claire Adam-Bourdarios^a, Glen Cowan^b, Cécile Germain^c, Isabelle Guyon^d, Balázs Kégl^{a,c}, David Rousseau^a

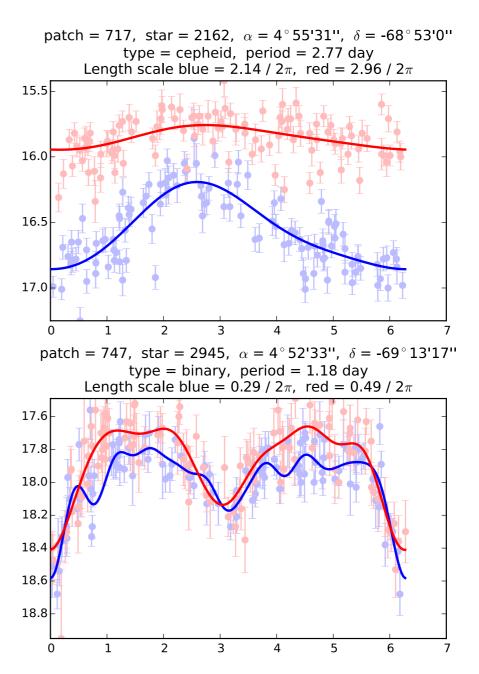
^a LAL, IN2P3/CNRS & University Paris-Sud, France ^b Physics Department, Royal Holloway, University of London, UK ^c TAO team, INRIA & LRI, CNRS & University Paris-Sud, France ^d ChaLearn

CLASSIFYING VARIABLE STARS

- collaboration with astrophysicists at Paris-Saclay
- variable-length functional data

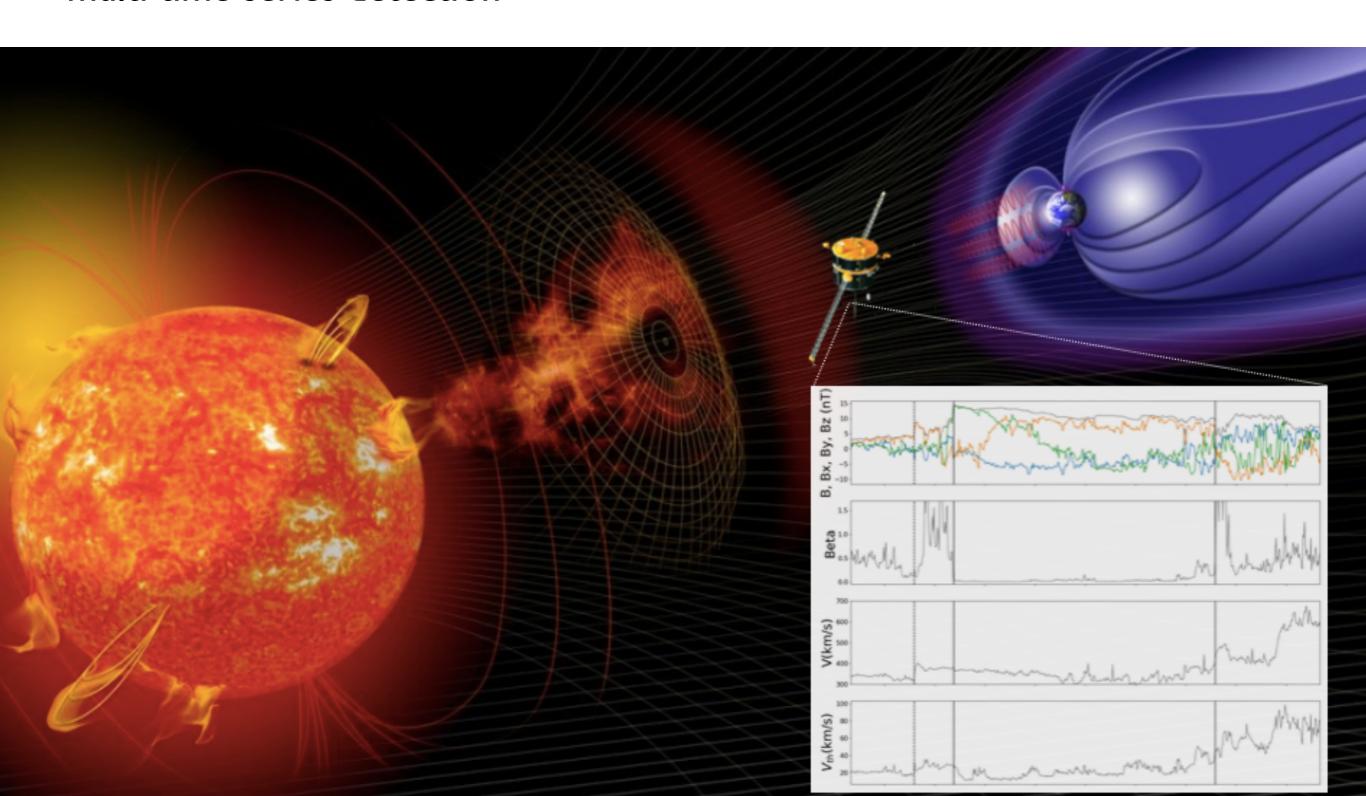


type = mira, period = 214.28 day Length scale blue = $2.48 / 2\pi$, red = $2.09 / 2\pi$

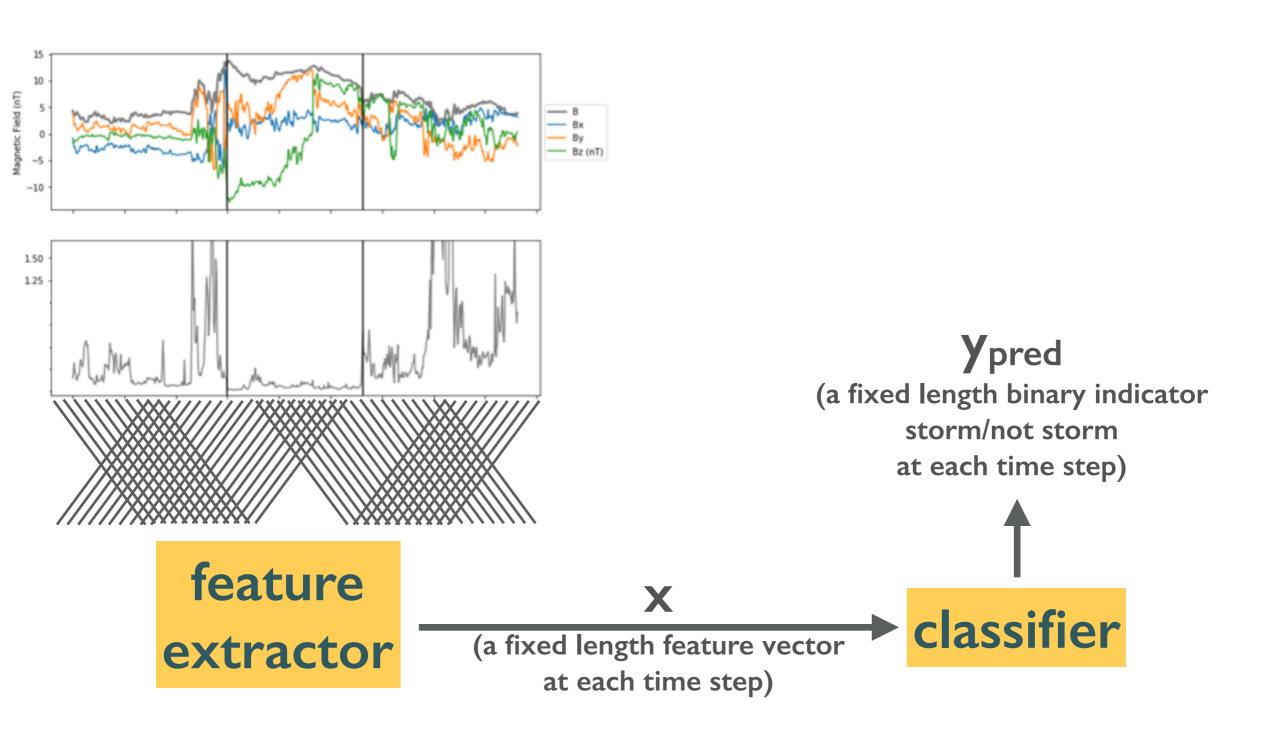


DETECTING SOLAR STORMS

- collaboration with plasma physicists at Paris-Saclay
- multi time series detection

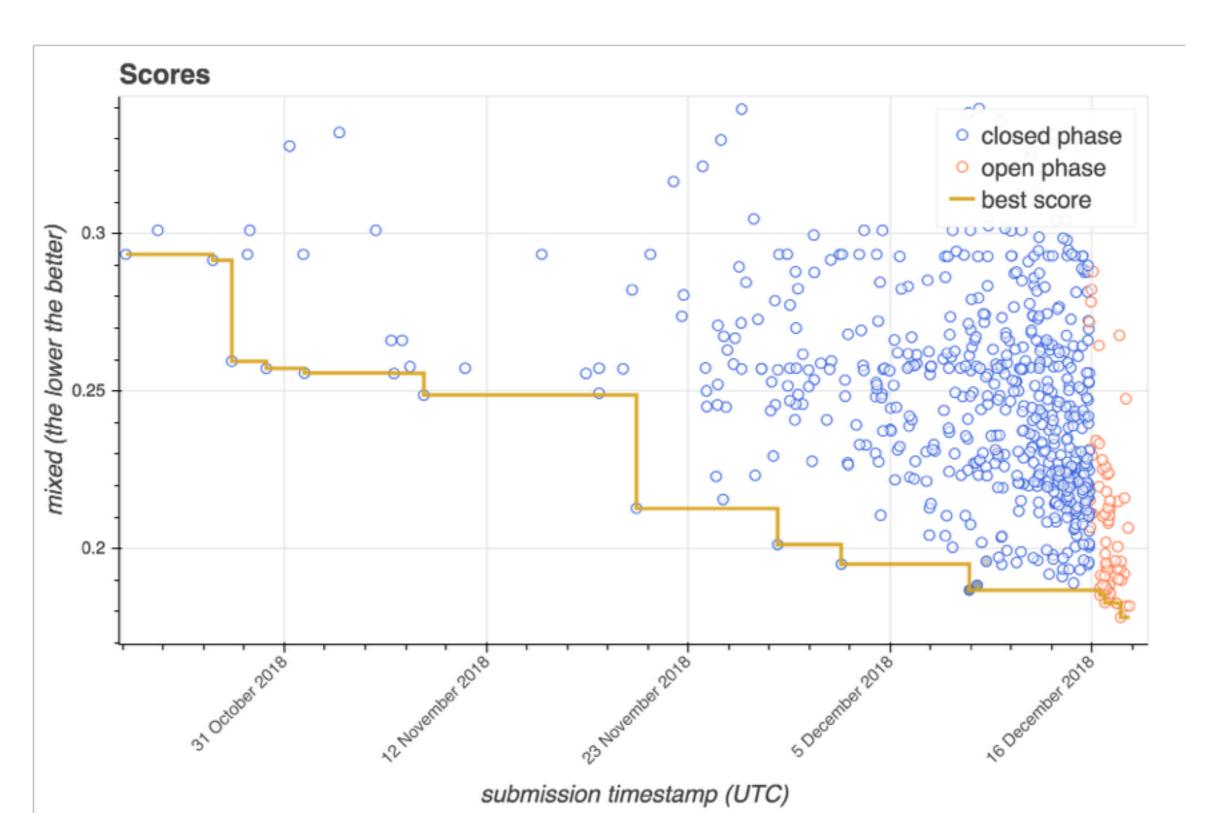


DETECTING SOLAR STORMS

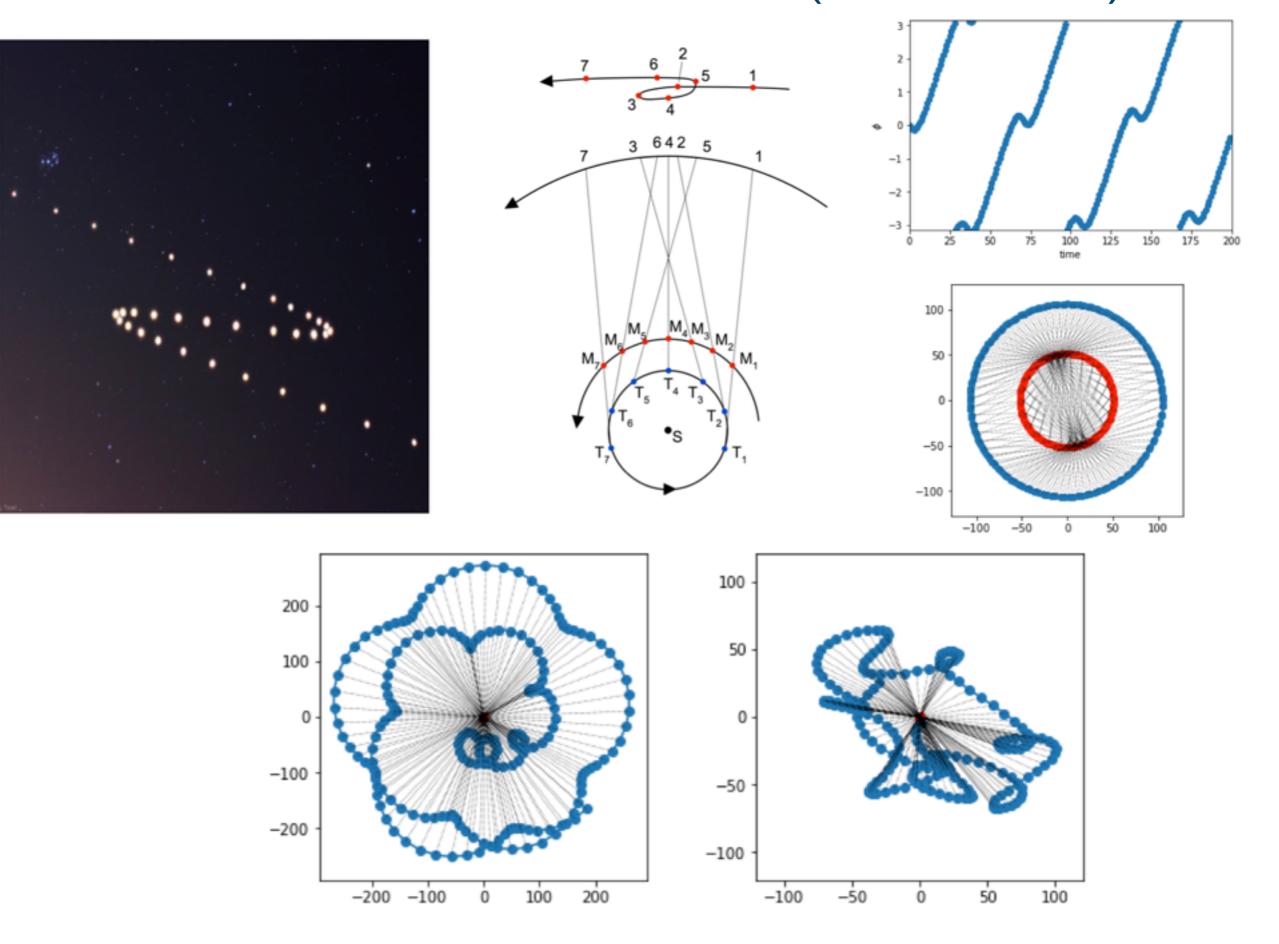


GRADUATE STUDENT DESCENT

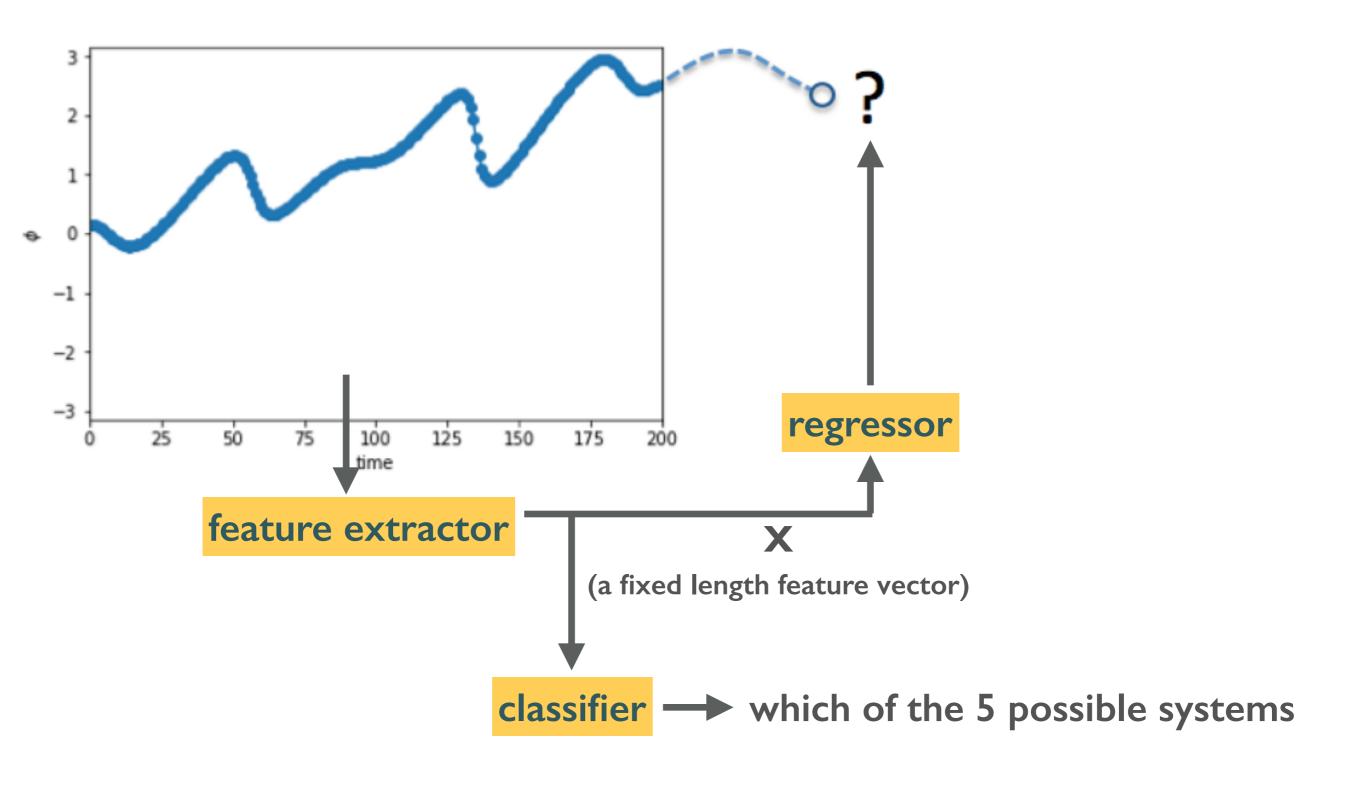
Saclay MSc students competing and learning from each other submitting 700 predictive models in three weeks



DISCOVERING PHYSICS (MECHANICS)



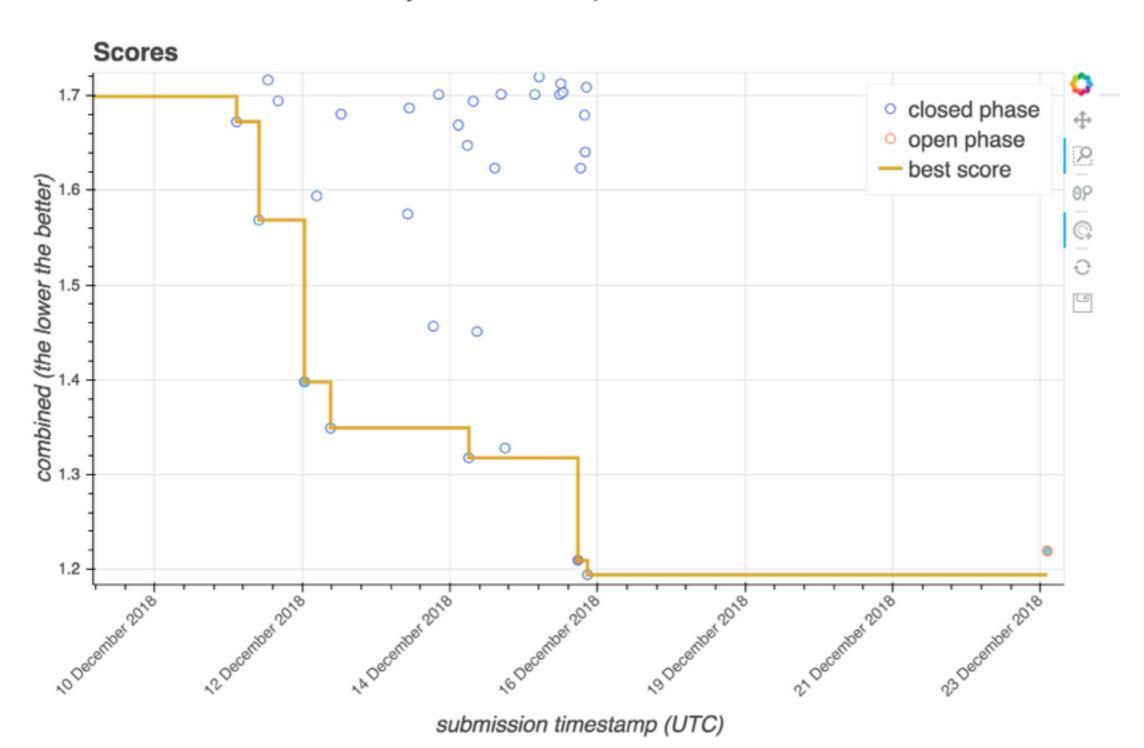
DISCOVERING PHYSICS (MECHANICS)



GRADUATE STUDENT DESCENT

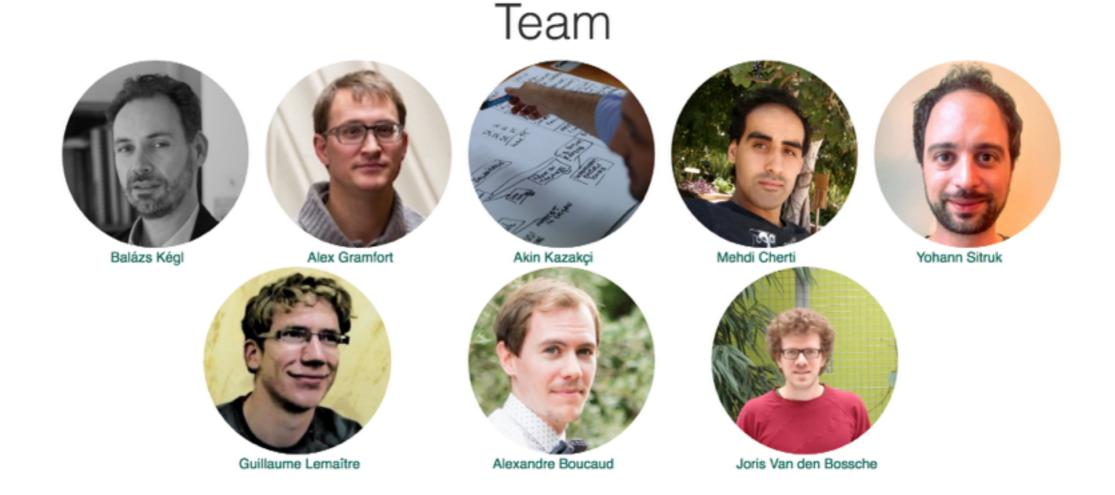
Saclay MSc students competing and learning from each other submitting 40 predictive models in three weeks

Mechanics classification, Saclay M2 Data Camp 2018/19



We built and optimized ~20 predictive workflows for three years

Funded by Université Paris-Saclay and CNRS



Alumni



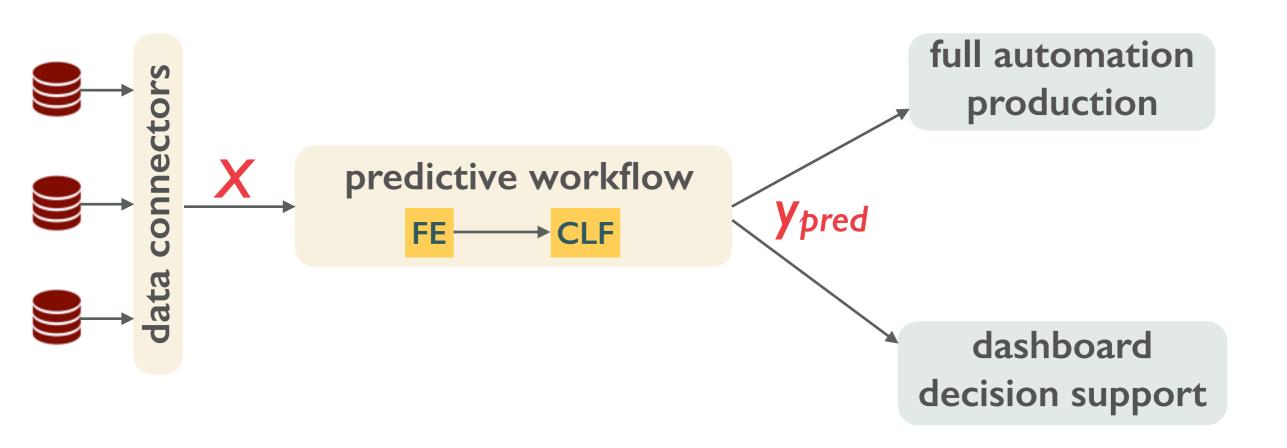
Djalel Benbouzid

What have we learned?

BUILDING PREDICTIVE WORKFLOWS WHAT HAVE WE LEARNED?

Building the workflow: what are the tasks and who does what

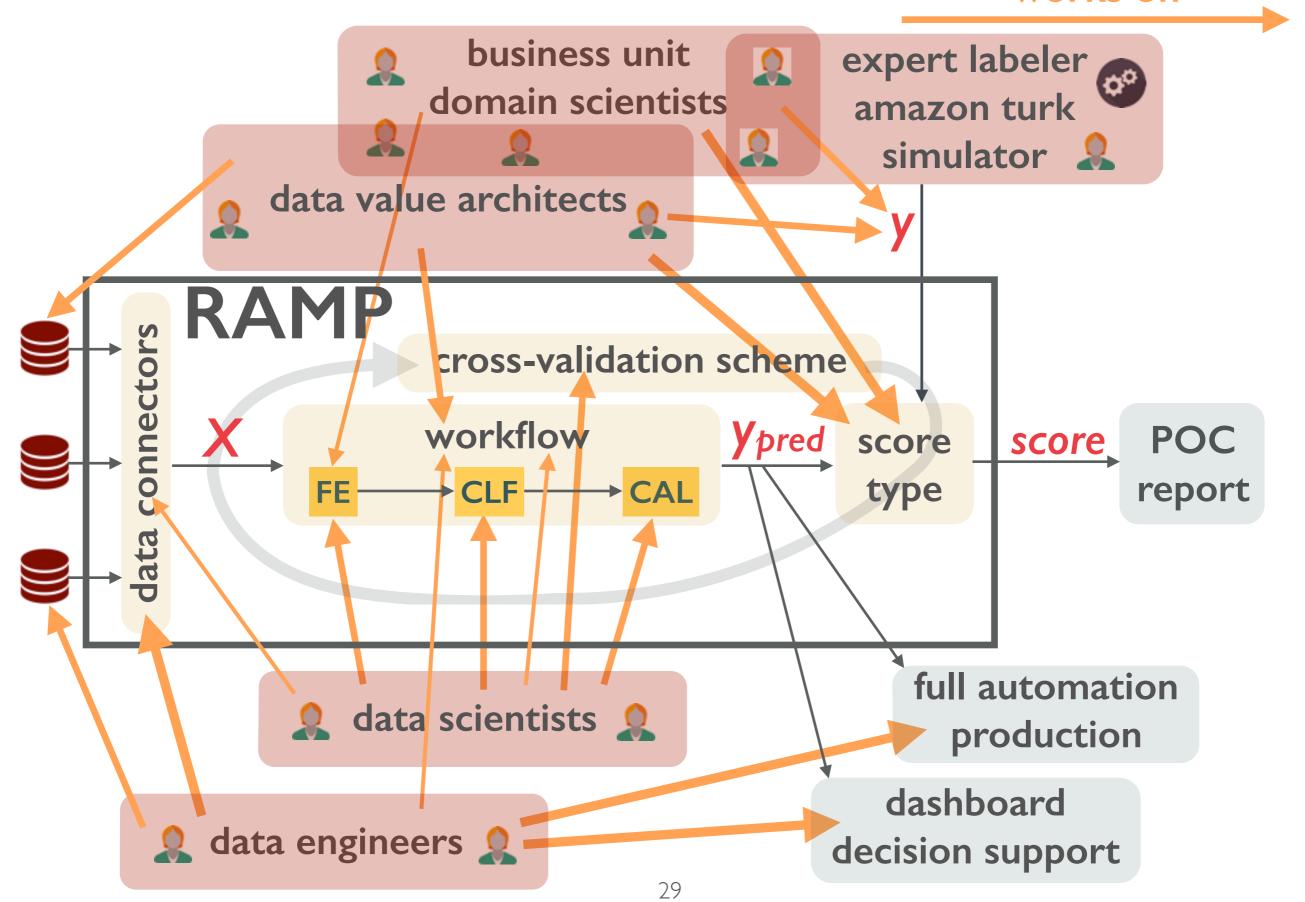
THE PREDICTIVE WORKFLOW



THE IDEAL SEQUENCE

data flow

"works on"



BUILDING PREDICTIVE WORKFLOWS WHAT HAVE WE LEARNED?

What is a predictive workflow?

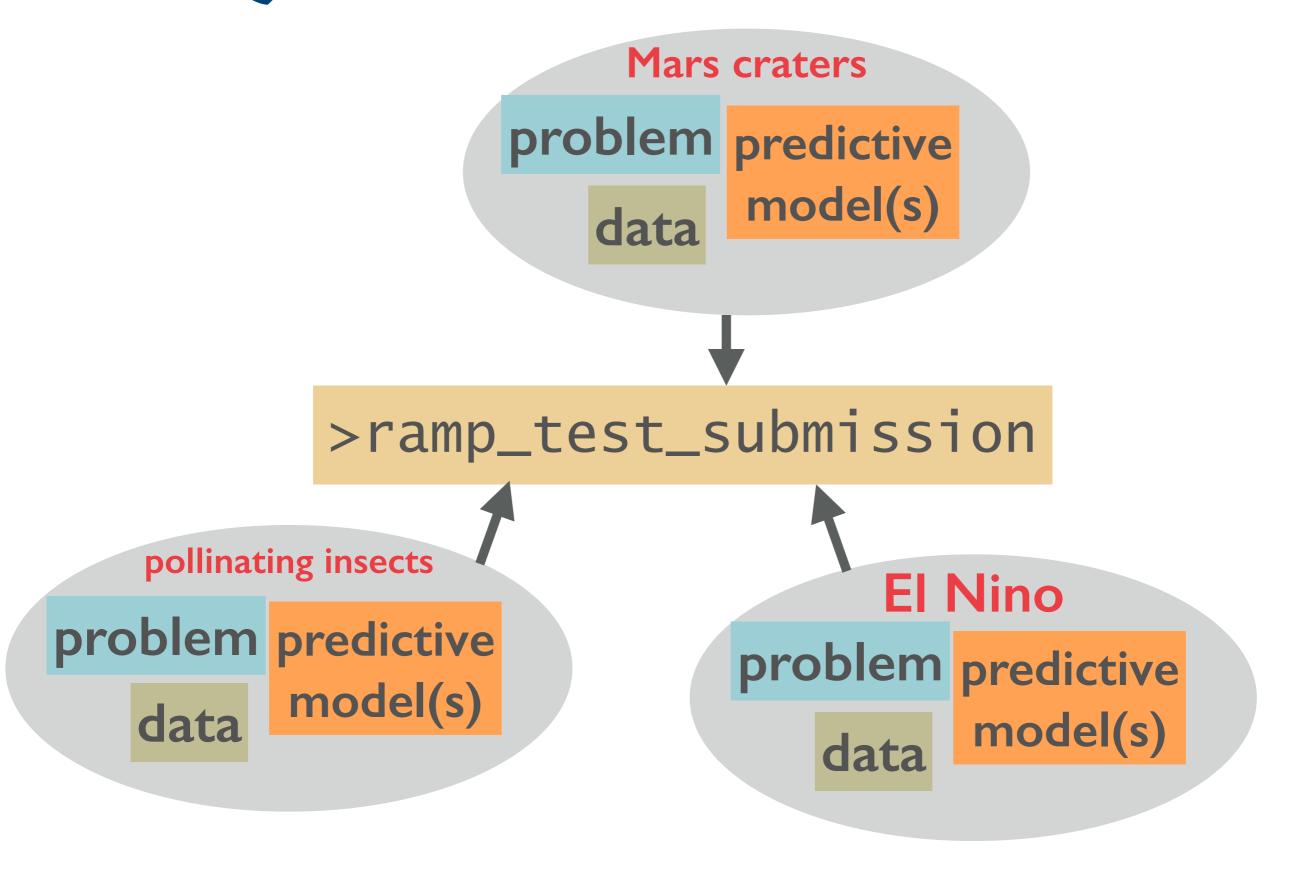
What are the parametrizable components?

What can be put into a unique training/scoring script?

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
      X df = data.drop([ target column name], axis=1)
41
                                                               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 UNIQUE SCRIPT TO RUN THE BUNDLES



A UNIQUE SCRIPT TO RUN THE BUNDLES

```
1 read training and test data
2 read submission
3 create train and valid folds
  on training data
4 for all train and valid folds:
        train submission on train
6        score submission on train,
        valid, and test
7 summarize scores
```

```
silver6:autism kegl$ ramp_test_submission
Testing Autism Spectrum Disorder classification
Reading train and test files from ./data ...
Reading cv ...
Training ./submissions/starting_kit ...
Couldn't re-order the score matrix...
              0.696 0.765
        train 0.767 0.847
       valid 0.611 0.647
CV fold 1
Couldn't re-order the score matrix..
              0.478 0.659
        train 0.766 0.842
        valid 0.628 0.662
CV fold 2
Couldn't re-order the score matrix..
                     0.720
        train 0.786 0.854
       valid 0.615 0.645
CV fold 3
Couldn't re-order the score matrix...
              0.565 0.758
        train 0.769 0.849
        valid 0.619 0.645
Mean CV scores
Couldn't re-order the score matrix..
              0.587 \pm 0.0784
                               0.725 \pm 0.042
        train 0.772 ± 0.0081 0.848 ± 0.0042
        valid 0.618 ± 0.0065
                               0.65 \pm 0.0072
Bagged scores
Couldn't re-order the score matrix..
        score
```

valid 0.647

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
 - ramp_test_submission: unique command-line test script
- examples: https://github.com/ramp-kits
 - a zoo of problems, experiments, workflows
 - (at least) one initial solution



BUILDING PREDICTIVE WORKFLOWS WHAT HAVE WE LEARNED?

How to make (novice) data scientists efficient

HOW TO MAKE DATA SCIENTISTS EFFICIENT

Principles

- incite them to work on the problem
- give them a working (but unoptimized) model to start with
- make incremental contributions easy
- gamification
- help them to collaborate and to learn from each other
- "hide" heavy engineering and computational obstacles

THE JUPYTER NOTEBOOK

(http://www.datascience-paris-saclay.fr)

RAMP on Mars craters detection

Alexandre Boucaud (CDS), Joris van den Bossche (CDS), Balazs Kegl (CDS), Frédéric Schmidt (GEOPS), Anthony Lagain (GEOPS)

- 1. Introduction
- Preprocessing
- Workflow
- Evaluatio
- Local testing/exploration
- Submission

Introduction

Impact craters in planetary science are used to date planetary surfaces, to characterize surface processes and to study the upper crust of terrestrial bodies in our Solar System (Melosh, 1989). Thanks to the Martian crater morphology, a wide amount of information could be deduced on the geological history of Mars, as for example the evolution of the surface erosion rate, the presence of liquid water in the past, the volcanic episodes or the volatiles layer in the subsurface (Carr & Head, 2010). These studies are widely facilitated by the availability of reference crater databases.

Surveying impact craters is therefore an important task which traditionally has been achieved by means of visual inspection of images. The enormous number of craters smaller than one kilometer in diameter, present on high resolution images, makes visual counting of such craters impractical. In order to overcome this problem, several algorithms have been developed to automatically detect impact structures on planetary images (Bandeira et al., 2007; Martins et al., 2009). Nevertheless, these method allow to detect only 70-80 % of craters (Urbach & Stepinski, 2009).

The prediction task

This challenge proposes to design the best algorithm to detect crater position and size starting from the most complete Martian crater database containing 384 584 verified impact structures larger than one kilometer of diameter (Lagain et al. 2017). We propose to give to the users a subset of this large dataset in order to test and calibrate their algorithm.

labase containing 384 584 verified impact structures larger than one kilometer of diar le to give to the users a subset of this large dataset in order to test and calibrate their

- Concise description of the
 - scientific / business goal
 - the data
 - what are the steps, what do I have to do
 - initial solution
- Make the data scientist operational in a couple of hours

THE INITIAL SOLUTION

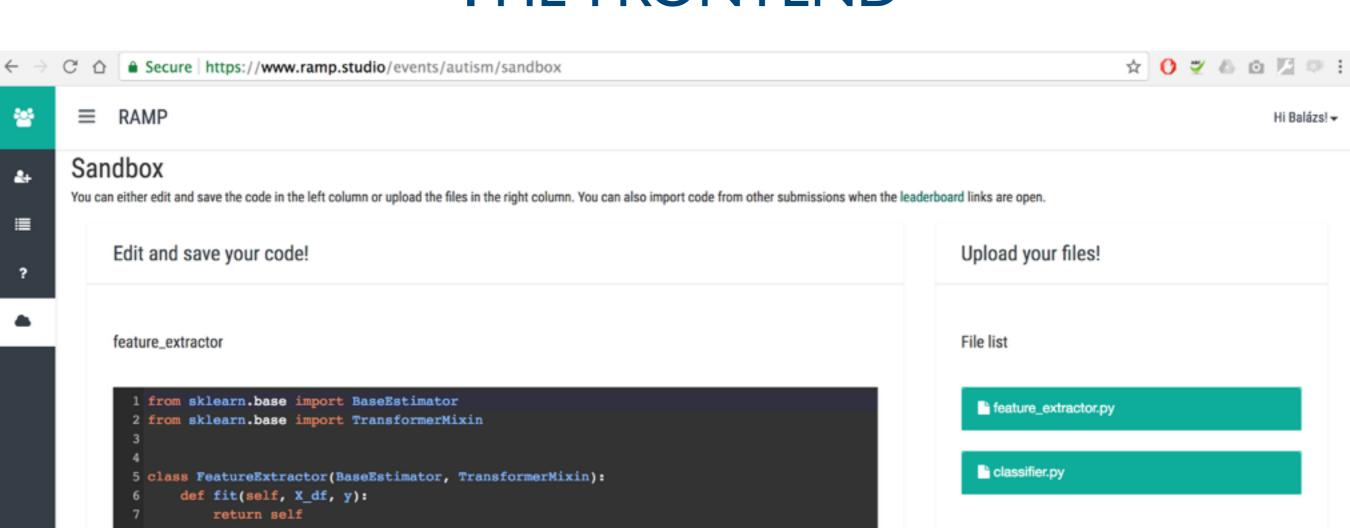
feature_extractor.py

```
1 from sklearn.base import BaseEstimator
 2 from sklearn.base import TransformerMixin
 3
  class FeatureExtractor(BaseEstimator, TransformerMixin):
       def fit(self, X df, y):
           return self
 9
       def transform(self, X df):
           # get only the anatomical information
10
11
           X = X df[[
12
               col for col in X df.columns
               if col.startswith('anatomy')]]
13
           return X.drop(columns='anatomy select')
14
15
```

classifier.py

```
1 from sklearn.base import BaseEstimator
 2 from sklearn.preprocessing import StandardScaler
 3 from sklearn.linear model import LogisticRegression
  from sklearn.pipeline import make pipeline
 5
  class Classifier(BaseEstimator):
       def init (self):
 9
           self.clf = make pipeline(
10
               StandardScaler(), LogisticRegression(C=1.))
11
12
       def fit(self, X, y):
1.3
           self.clf.fit(X, y)
           return self
14
15
16
       def predict(self, X, y):
17
           return self.clf.predict(X)
18
19
       def predict proba(self, X):
           return self.clf.predict proba(X)
20
21
```

THE FRONTEND



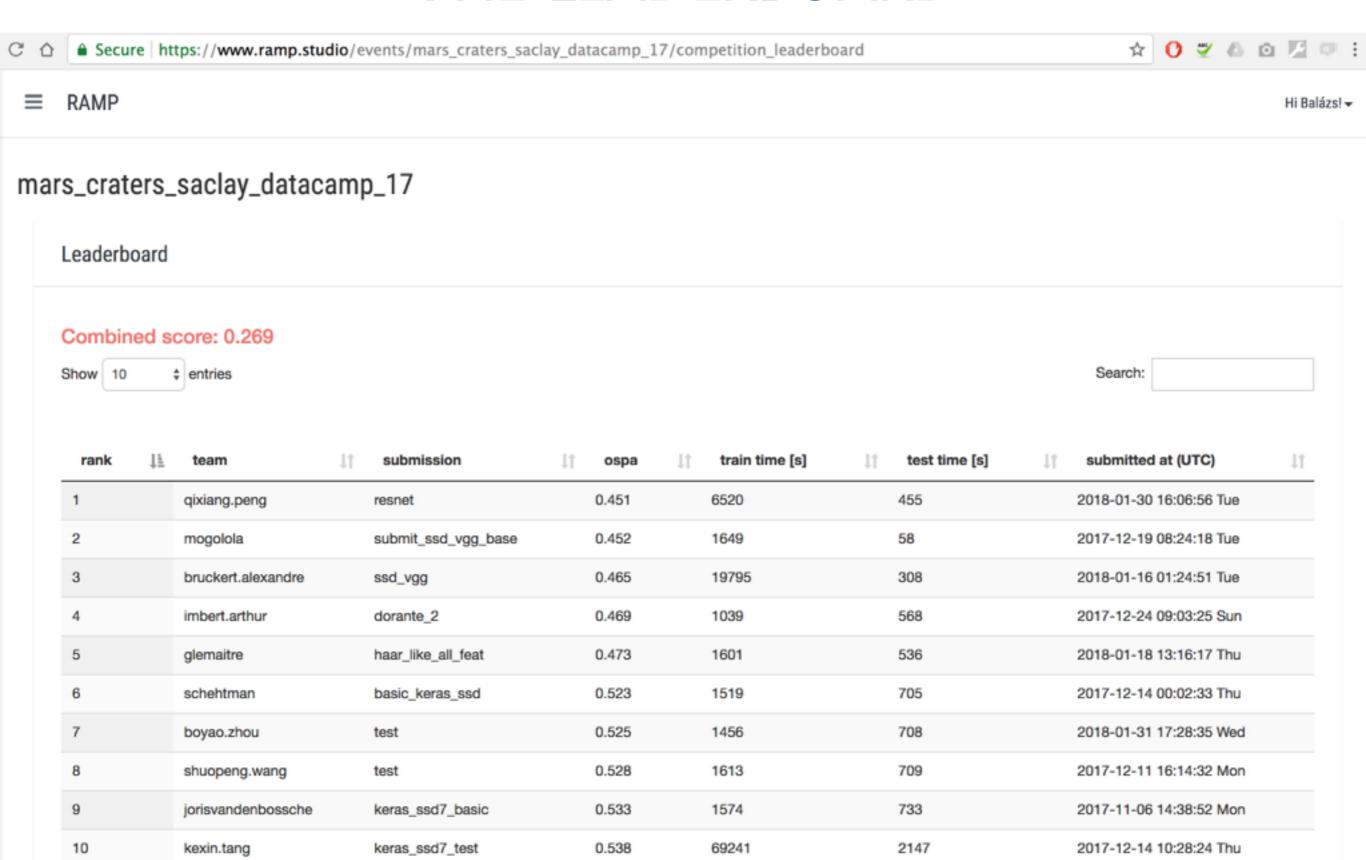
9 def transform(self, X_df): 10 # get only the anatomical information 11 X = X_df[[col for col in X_df.columns if col.startswith('anatomy')]] 12 return X.drop(columns='anatomy_select') 13

Upload file Choose File No file chosen Upload

classifier

```
1 from sklearn.base import BaseEstimator
2 from sklearn.preprocessing import StandardScaler
3 from sklearn.linear_model import LogisticRegression
4 from sklearn.pipeline import make_pipeline
```

THE LEADERBOARD



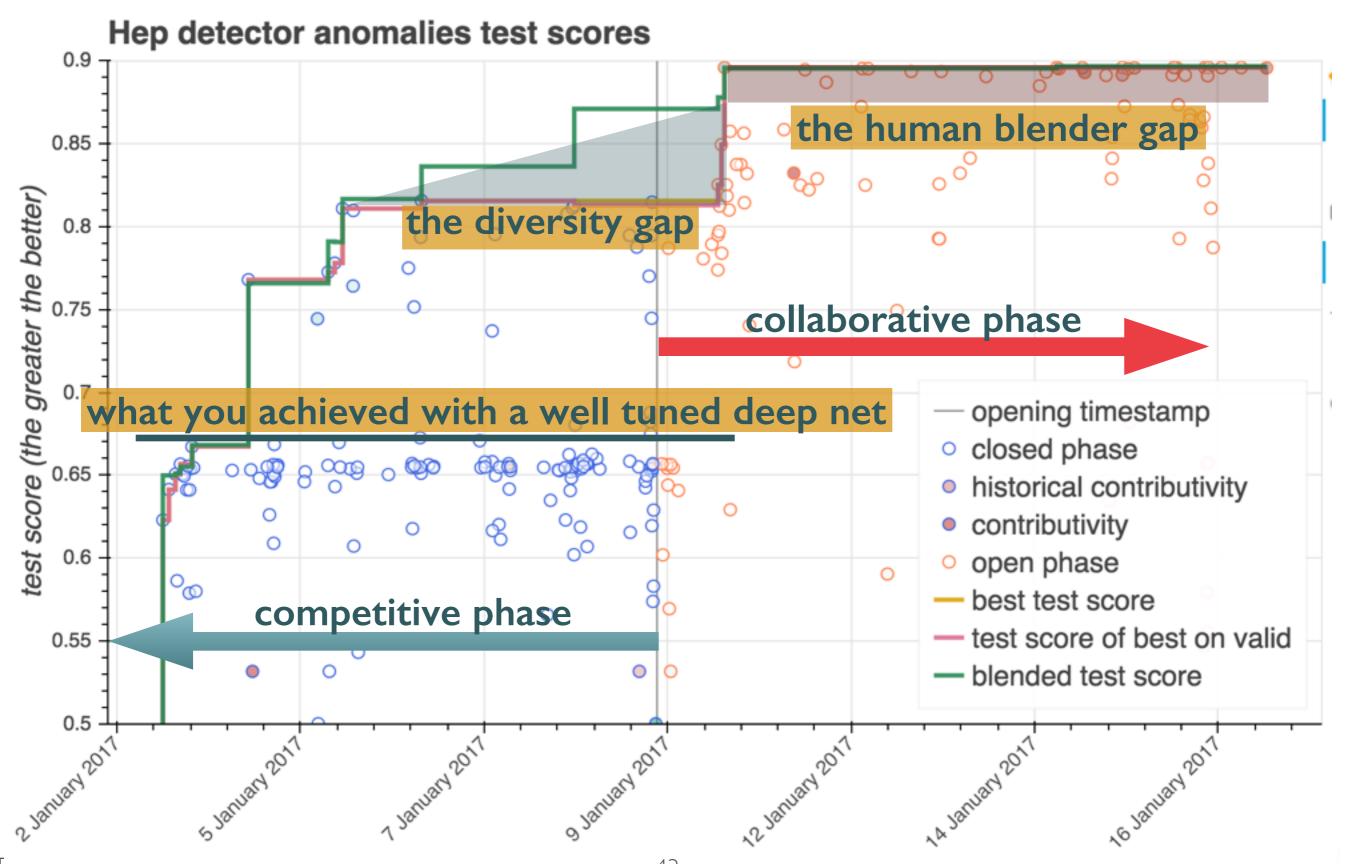
THE BACKEND ON AMAZON WEB SERVICES

| Laun | ch Instance | Connect | Actions * | | | | | | | | |
|------|--|---------|---------------------|-----------------|---------------------|---------------|--|--|--|--|--|
| Q | Q Filter by tags and attributes or search by keyword | | | | | | | | | | |
| | Name | - | Instance ID | Instance Type 🔻 | Availability Zone - | Instance Stat | | | | | |
| | 8475_submission_id | 0 | i-0f9411820a116930e | m5.xlarge | us-west-2a | terminated | | | | | |
| | 8474_test2 | | i-0a8afc1be7d824cb1 | m5.xlarge | us-west-2a | terminated | | | | | |
| | 8473_test | | i-0322ebb27a7e4e5 | m5.xlarge | us-west-2a | terminated | | | | | |
| | 8473_test | | i-09f50b72b6a3a0155 | m5.xlarge | us-west-2a | terminated | | | | | |
| | 8472_TunedClassif+ | svc | i-04dca6a8fd8738b5e | t2.small | us-west-2c | terminated | | | | | |
| | 8472_TunedClassif+ | svc | i-0f0529a6ab890f17c | t2.small | us-west-2b | terminated | | | | | |
| | 8469_test2 | | i-049ba0f2c8075fdd6 | m5.xlarge | us-west-2a | terminated | | | | | |
| | 8467_TunedClassif+ | svm | i-08018a0ee2b90b2ec | t2.small | us-west-2b | running | | | | | |

Why code submission

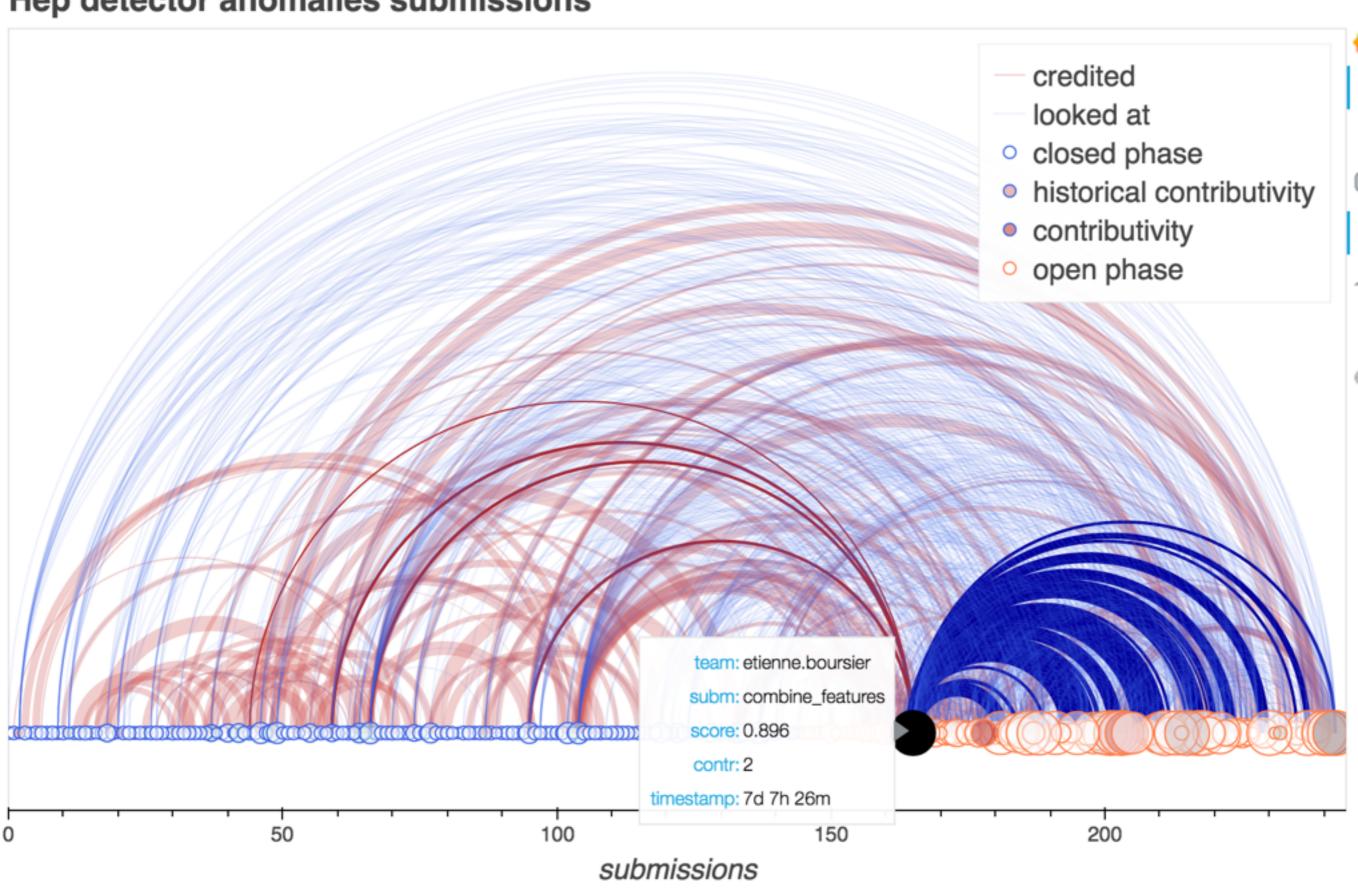
- I. lets us deliver a working prototype
- 2. lets the participants collaborate

THE POWER OF THE (COLLABORATING) CROWD OPTIMIZING GRADUATE STUDENT DESCENT



COMMUNICATION AND REUSE

Hep detector anomalies submissions



You can

- I. Use RAMP in teaching or training
- 2. Use the toolkit for your own workflows

LINKS

frontend:

www.ramp.studio

toolkit:

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

server:

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

examples:

github.com/ramp-kits

slack:

ramp-studio.slack.com

LINKS

- · medium.com/@balazskegl
 - The data science ecosystem (industrial edition)
 - Teaching the data science process
 - How to build a data science pipeline
- RAMP paper
 - https://openreview.net/forum?id=Syg4NHz4eQ

