

# MOABB

## Motivations and future

Sylvain Chevallier  
LISV - Université de Versailles, France  
[sylvain.chevallier@uvsq.fr](mailto:sylvain.chevallier@uvsq.fr)

Pedro Rodrigues & Marco Congedo  
Gipsa-Lab - Grenoble, France  
[pedro.rodrigues@gipsa-lab.fr](mailto:pedro.rodrigues@gipsa-lab.fr)  
[marco.congedo@gmail.com](mailto:marco.congedo@gmail.com)

16 September 2019



# What will this presentation be about?

- Less theoretical work
- Little bit more code explanation
- All you need to know for hands-on session

# Outline

MOABB: Why and What

MNE for neuro

Scikit-learn for ML

Pyriemann

# Why do we need MOABB?

Reproducible research in BCI has a long way to go...

- Unavailable code
- Exotic data format/language/toolboxes
- Preprocessed data (including errors)

⇒ **No comprehensive benchmark of BCI algorithms**

⇒ **Huge waste of time for everyone**

⇒ **MOABB aims to be the standard benchmark for any new paper**

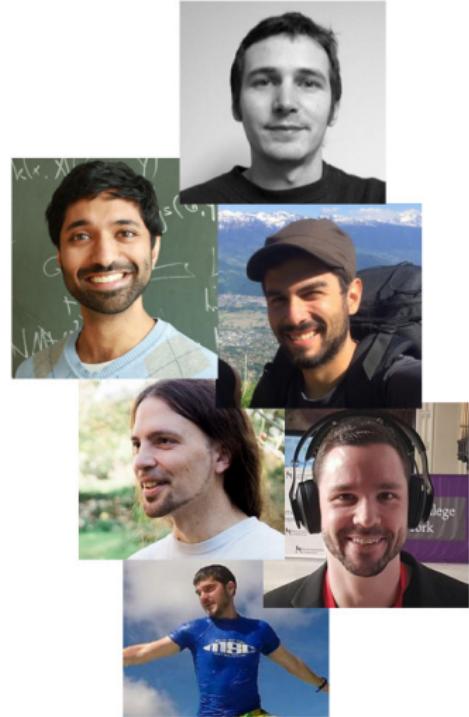
- Comprehensive benchmark of popular BCI algorithm
- Extensive list of freely available EEG datasets
- Ranking algorithms with fair evaluations

# Who is behind MOABB?

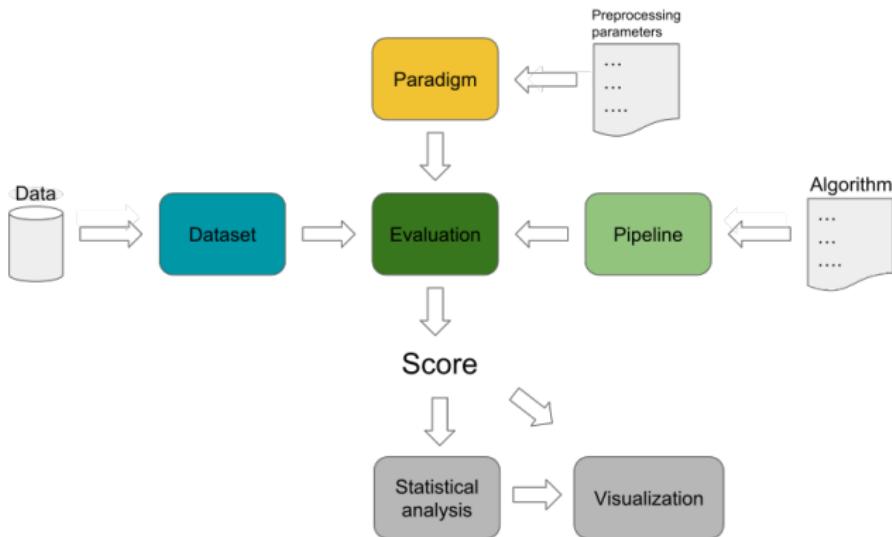
**Founder: Alexandre Barachant**

**Contributors:**

- Vinay Jayaram
- Pedro Rodrigues
- Sylvain Chevallier
- Justin D. Harris
- Yannick Roy & the NeuroTechX community
- ...
- You!



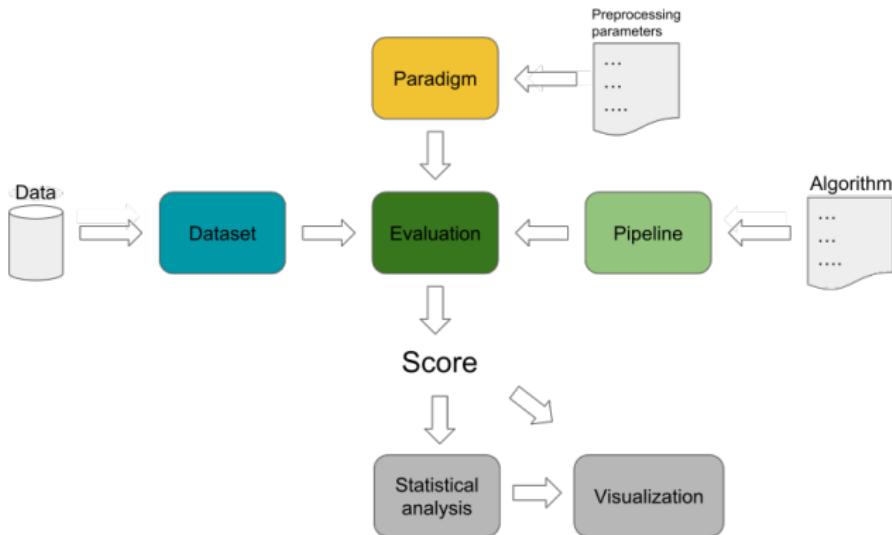
# MOABB architecture



## Dataset

- Stored locally, converted in MNE format
- Pick only subjects/sessions you need

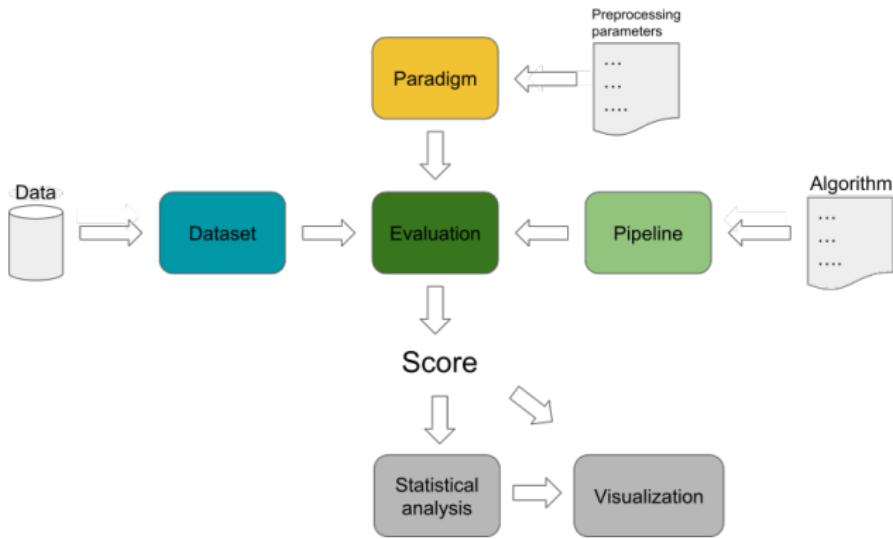
# MOABB architecture



## Paradigm

- Defines conversion to trial
- Preprocessing

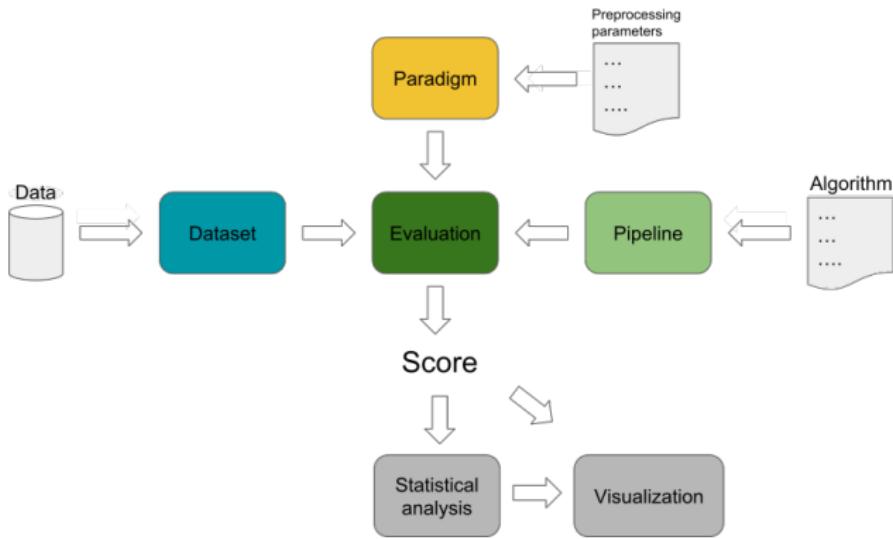
# MOABB architecture



## Evaluations

- Defines a scoring method (AUC, f-score, ...)
- within or across session, across-subject, ...

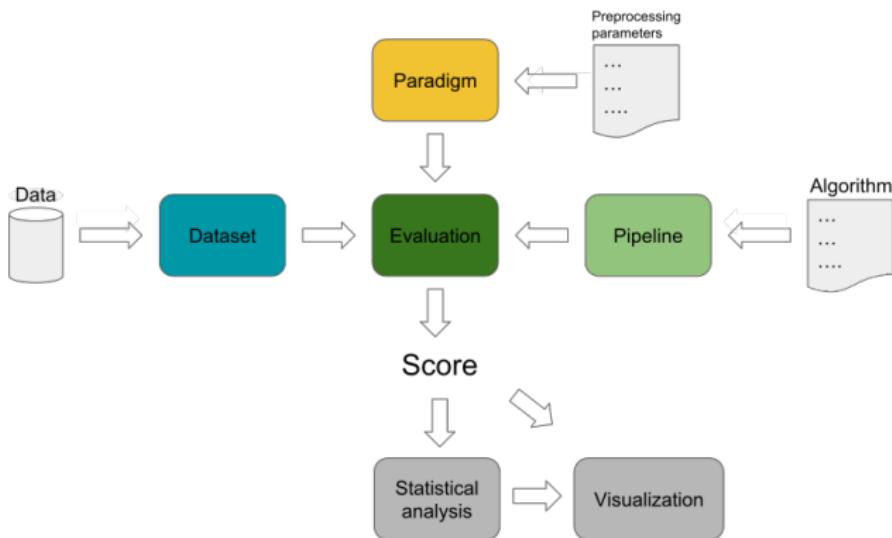
# MOABB architecture



## Pipelines

- All steps required for obtaining a prediction
- Scikit-learn style

# MOABB architecture



## Results

- Statistics & visualization
- Results are stored in a DataFrame

# Outline

MOABB: Why and What

MNE for neuro

Scikit-learn for ML

Pyriemann

# Reproducible research with neurophysiological data

**Freesurfer** Popular software for extracting features from MRI

<https://surfer.nmr.mgh.harvard.edu/>

⇒ Hardware and software differences can lead to different features/statistical results and scientific conclusions

**ICA** Popular matrix factorization problem

[https:](https://)

<https://github.com/mne-tools/mne-python/issues/4922>

⇒ Different results with different machines

**eigs/eigsh** Popular solver for eigenvalues decomposition

<https://github.com/scikit-learn/scikit-learn/issues/5545>

⇒ Even on the same machine numerical solvers can lead to different outcome

# MNE

<https://github.com/mne-tools/mne-python>

## History

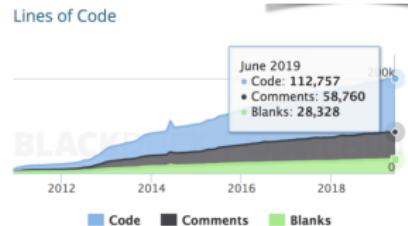
- based on C code developed for 18 years by Matti Hämäläinen
- Python started in 2010 at MGH, Boston

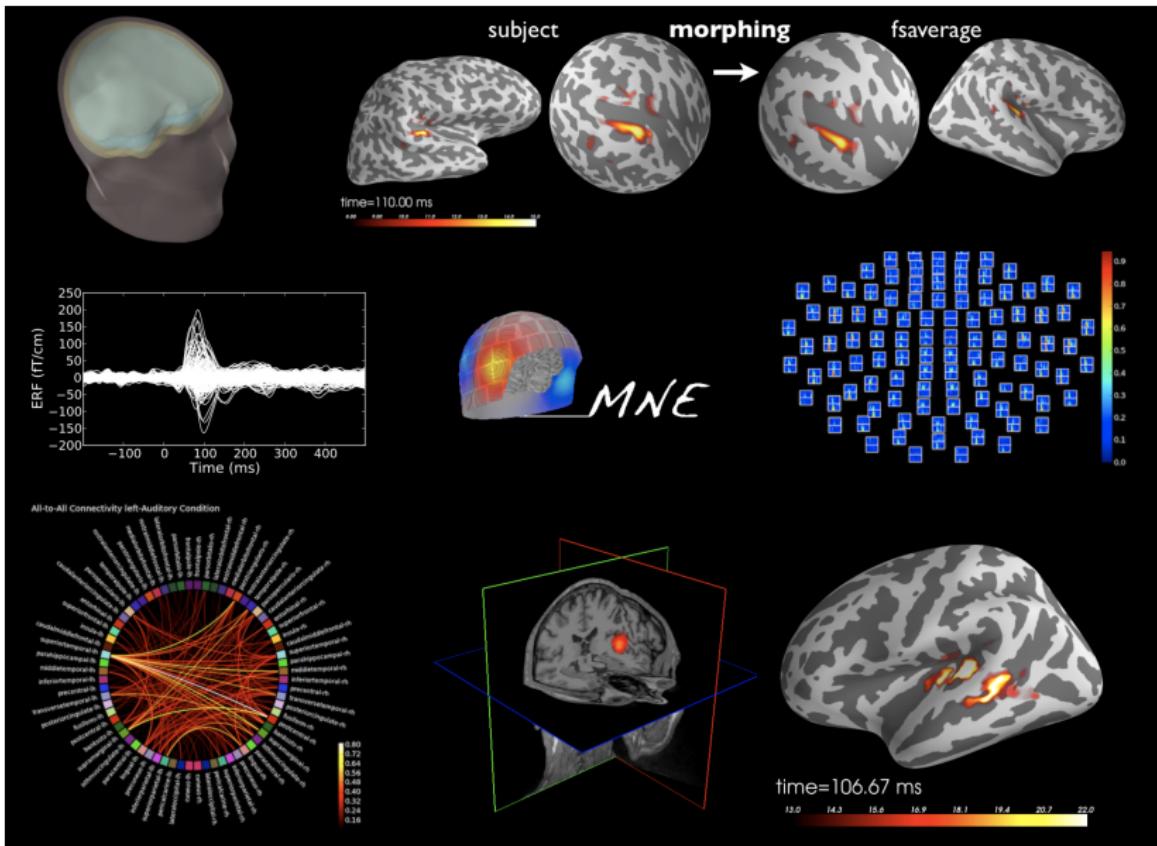
## In a nutshell

- 236 contributors, 100k LOC
- mature codebase, large dev team
- 29 years of efforts (COCOMO)

⇒ BSD licensed (commercial use ok)

⇒ Mac / Linux / Windows





# Outline

MOABB: Why and What

MNE for neuro

Scikit-learn for ML

Pyriemann



# Scikit-learn vision

<http://scikit-learn.org>

- **Machine learning for all**  
⇒ No specific application domain  
⇒ No requirements in machine learning
- **High-quality Pythonic software library**  
⇒ Interfaces designed for users
- **Community-driven development**  
⇒ BSD licensed, very diverse contributors

Easy as py:

```
from sklearn import svm
classifier = svm.SVC()
classifier.fit(X_train, Y_train)
Y_test = classifier.predict(X_test)
```



# Specifying a model

A central concept: **the estimator**

- Instanciated without data
- But specifying the parameters

Example:

```
from sklearn.neighbors import KNearestNeighbors  
  
estimator = KNearestNeighbors( n_neighbors=2)
```

# Training a model

## Training from data

```
estimator.fit(X_train , Y_train)
```

with:

- X a numpy array with shape  $n\_samples \times n\_features$
- y a numpy 1D array, of ints or float, with shape  $n\_samples$

# Using a model

Prediction classification, regression

```
Y_test = estimator.predict(X_test)
```

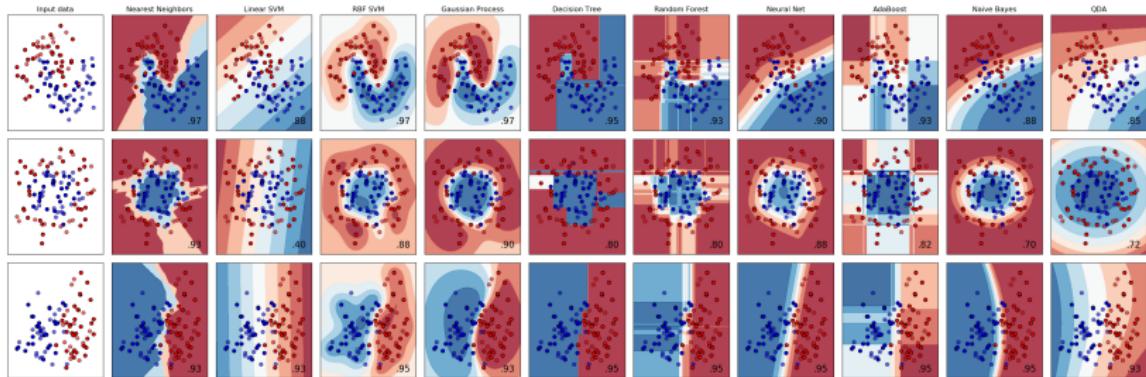
Transforming dimension reduction, filter

```
X_new = estimator.transform(X_test)
```

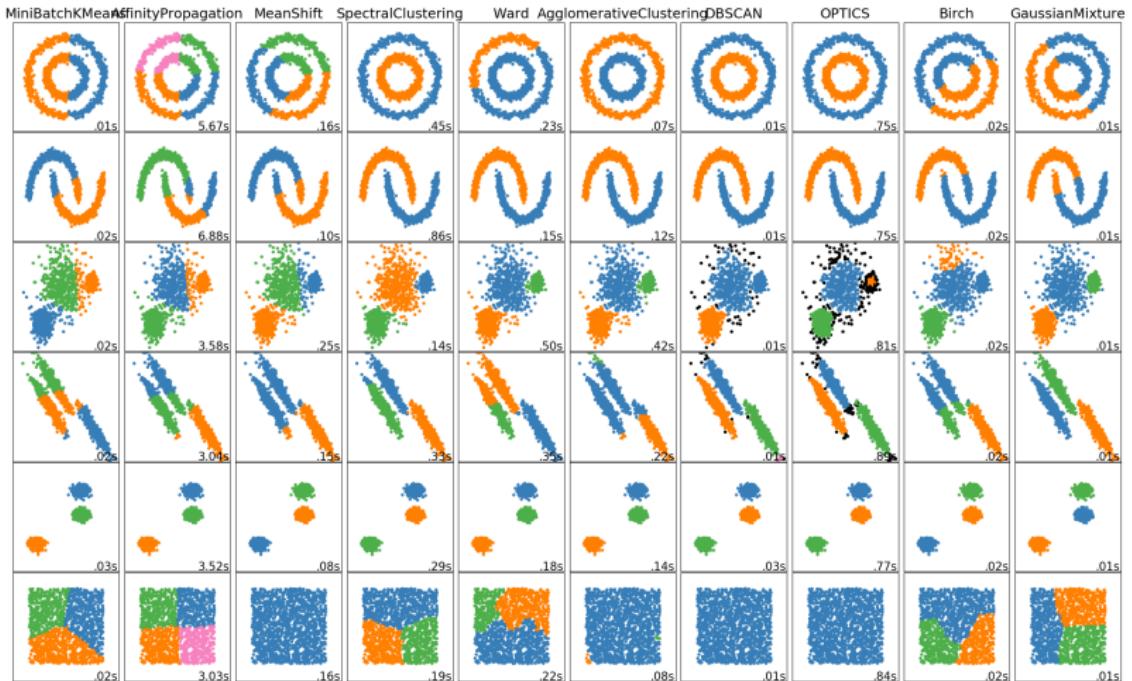
Test score, density estimation

```
test_score = estimator.score(X_test)
```

# Classification models

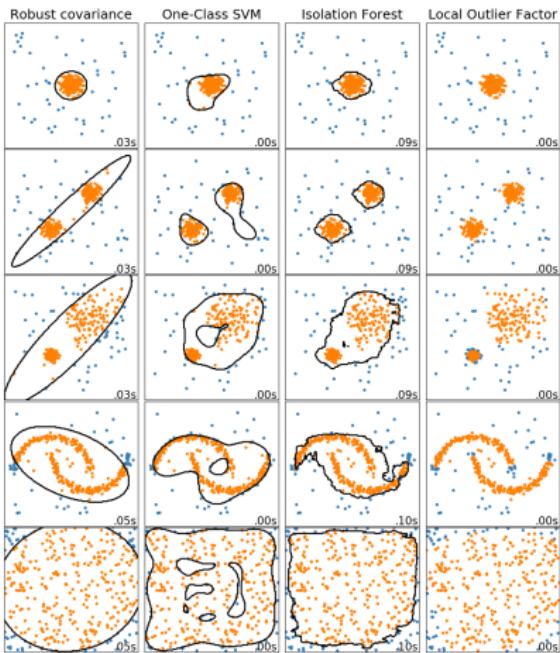


# Clustering models





# Anomaly detection



# Outline

MOABB: Why and What

MNE for neuro

Scikit-learn for ML

Pyriemann

# Pyriemann

<https://github.com/alexandrebarachant/pyRiemann>

- Fully integrated in scikit
- Could use MNE output

All you need in one toolbox

- Covariance estimation
- Classification
  - MDM
  - Tangent Space
  - K-Nearest Neighbor

- Clustering
- Tangent Space projection
- Spatial filtering
  - CSP
  - xDawn

# One example script

```
from scipy.io import loadmat
from pyriemann.estimation import Covariances
from pyriemann.classification import MDM
from sklearn.model_selection import train_test_split
from sklearn.metrics import accuracy_score

data = loadmat('data.mat')
X = data['X']
y = data['y'].squeeze()

covs = Covariances(estimator='scm').fit_transform(X)
covs_train, covs_test, y_train, y_test = \
    train_test_split(covs, y, train_size=0.8)

clf = MDM(metric='riemann')

clf.fit(covs_train, y_train)
y_pred = clf.predict(covs_test)
print(accuracy_score(y_test, y_pred))
```



Thank you !