

# Modeling Brain Waveforms with Convolutional Dictionary Learning and Point Processes

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Cédric Allain, Lindsey Power, Tim Bardouille



MIND



**Goal:** Study the brain mechanisms while it is functioning.

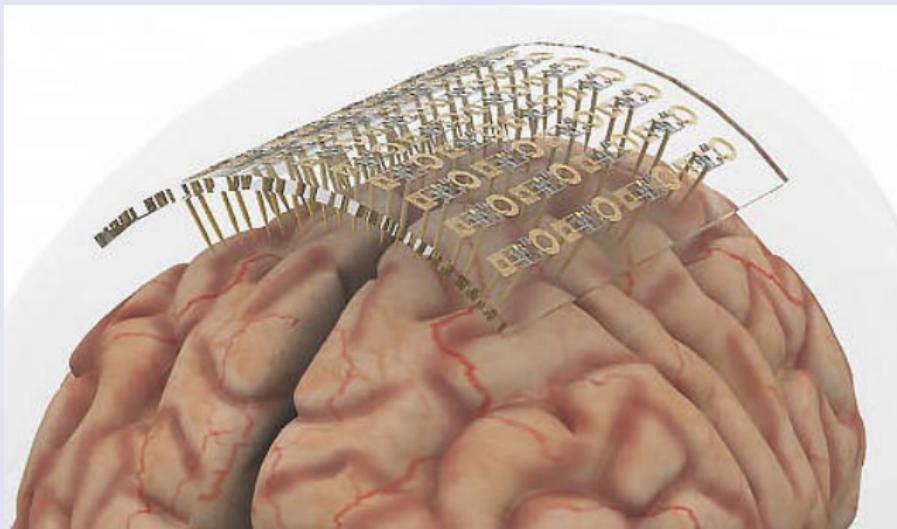
**Outputs:**

- ▶ **Functional Atlases:** Link areas of the brain to specific cognitive functions.
- ▶ **Functional Connectivity:** Highlight the information flow in the brain.
- ▶ **Healthcare:** Develop bio-markers for neurological disorders.

## Context: functional Neuroimaging

How to record living brains electrical activity: **Electrophysiology**

Direct measurement: intracranial EEG.



High Localization

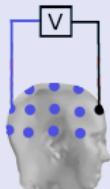
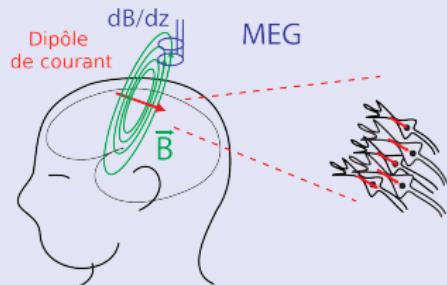
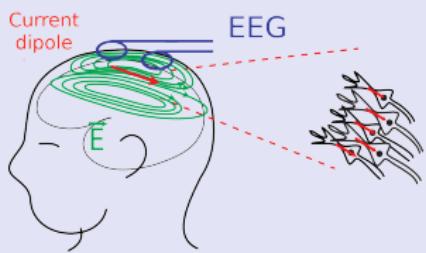
Low Resolution

Invasive

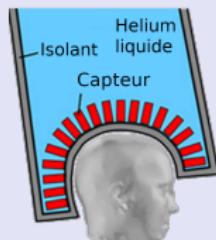
# Context: functional Neuroimaging

How to record living brains electrical activity: **Electrophysiology**

Remote measurement: M/EEG.



No Localization

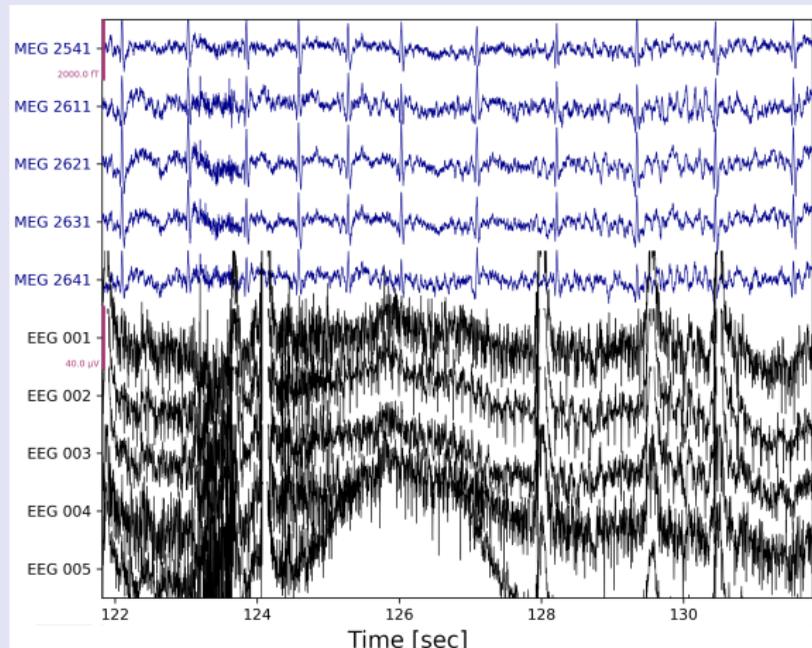


Global

Non Invasive

# M/EEG signals

## Multivariate time-series X



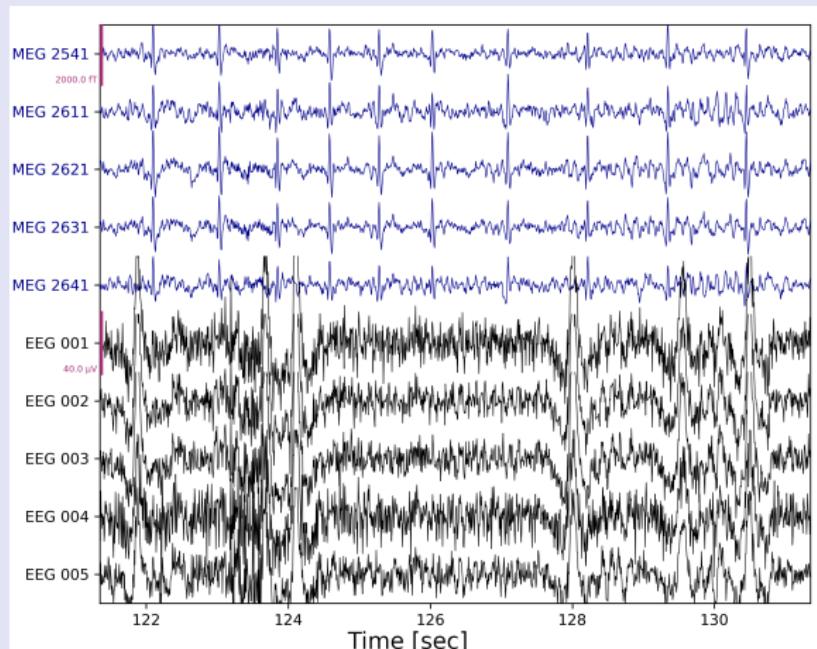
Noisy

Many artifacts

Complex

# M/EEG signals

## Multivariate time-series X



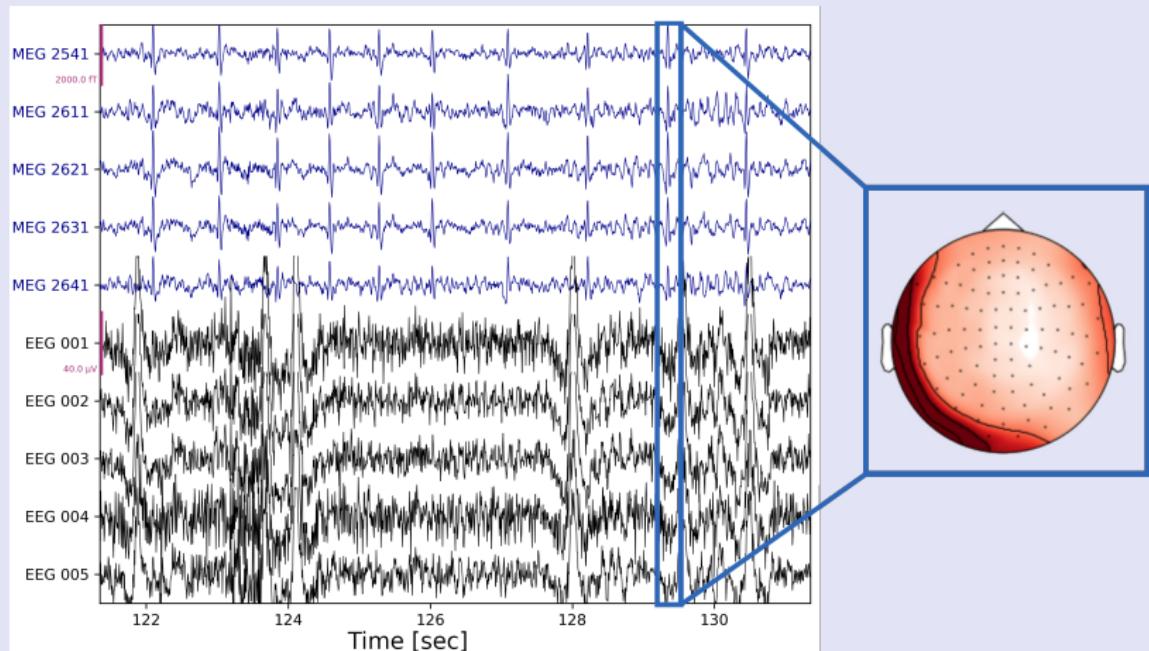
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# M/EEG signals

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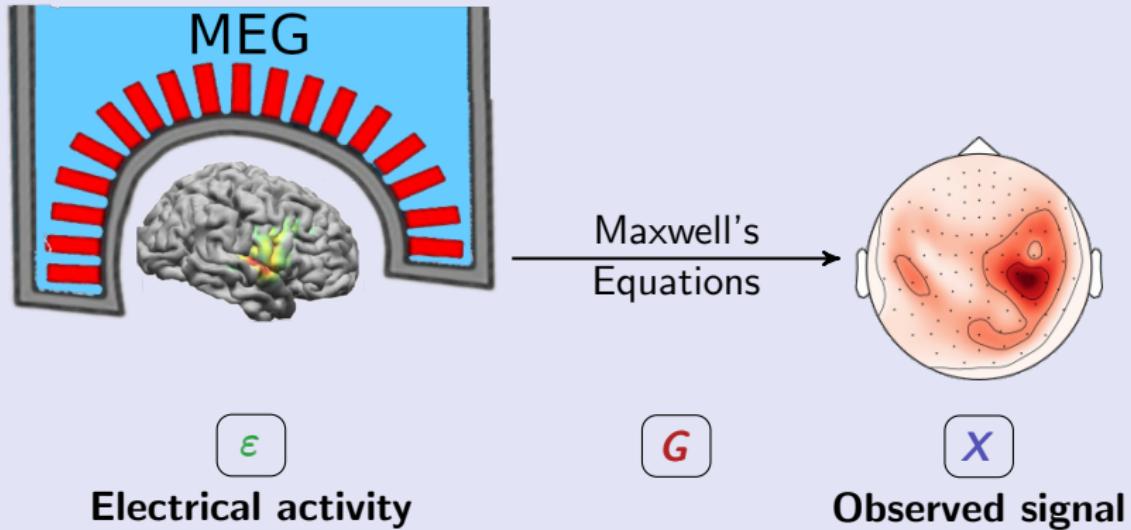


Noisy

Many artifacts

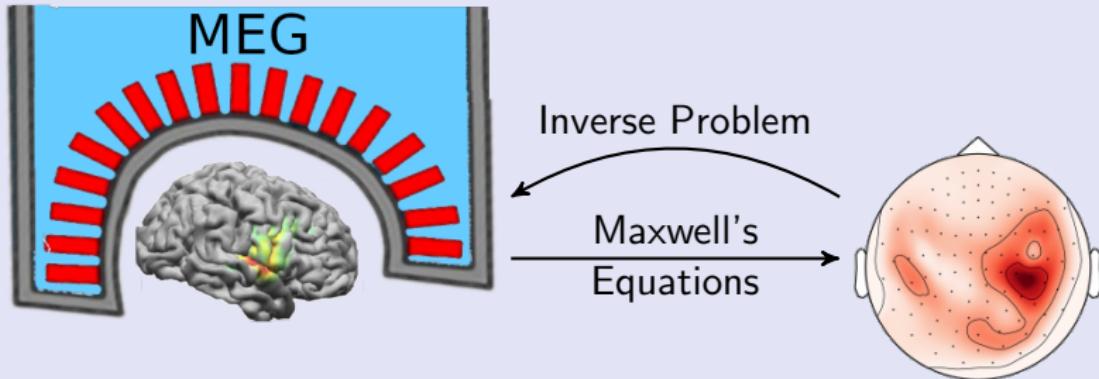
Complex

# How to get back to electrical activity?



Forward model:  $X = G\epsilon$

# How to get back to electrical activity?



$\epsilon$

Electrical activity

$G$

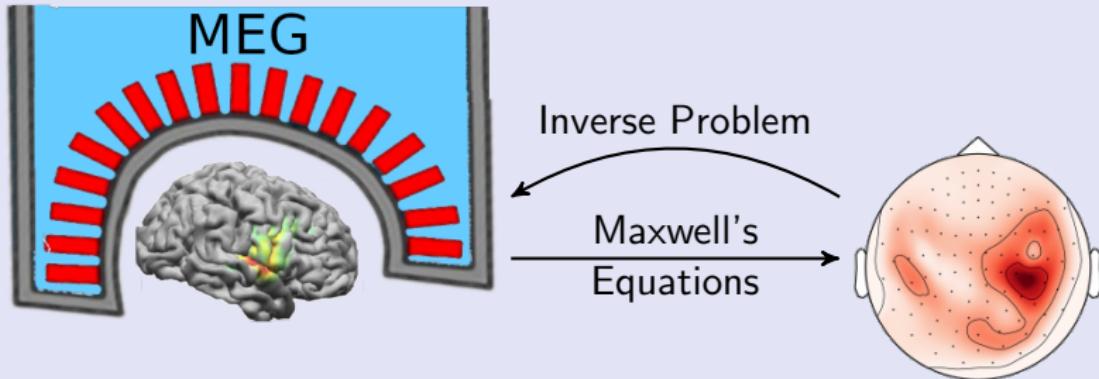
Forward model:  $X = G\epsilon$

$X$

Observed signal

Inverse problem:  $\epsilon = f(X)$  (ill-posed)

# How to get back to electrical activity?



$\epsilon$

Electrical activity

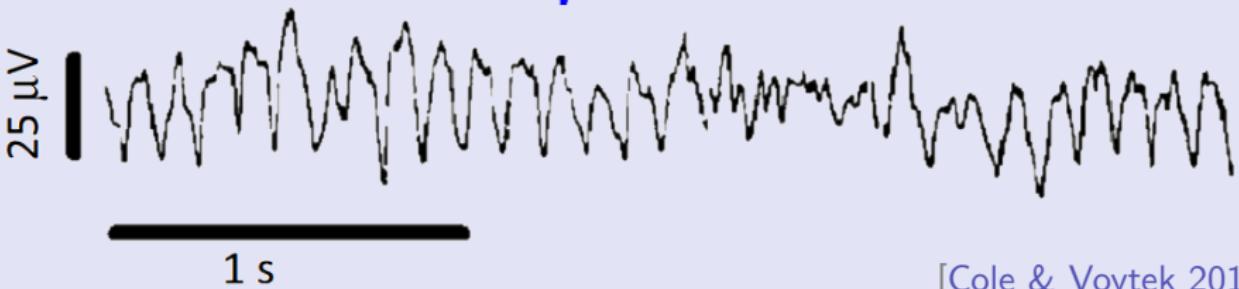
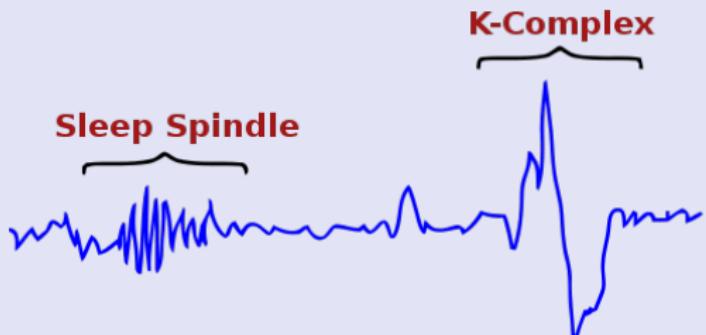
$G$

Observed signal

Forward model:  $X = G\epsilon$

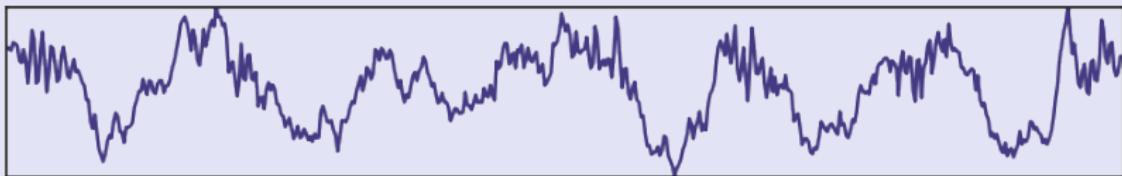
Inverse problem:  $\epsilon = f(X)$  (ill-posed)

- ▶ Dipole fit [Sarvas, 1987]
- ▶ Regularized optimization [Gramfort et al., 2012]
- ▶ Deep-learning [Hecker et al., 2021]

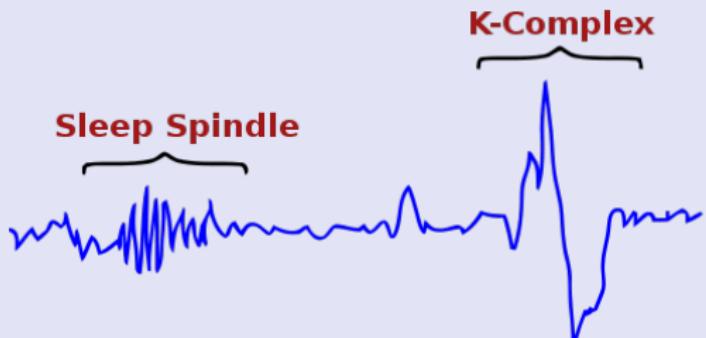


Neural signals exhibit diverse and complex morphologies

[Cole & Voytek 2017]



[Dupré la Tour et al. 2017]



Neural signals exhibit diverse and complex morphologies

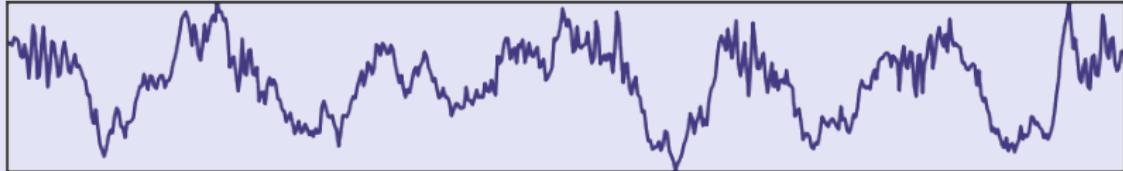
25  $\mu$ V

Waveform shape can be related to diseases  
e.g. Parkinson

[Jackson et al. 2019]

1 s

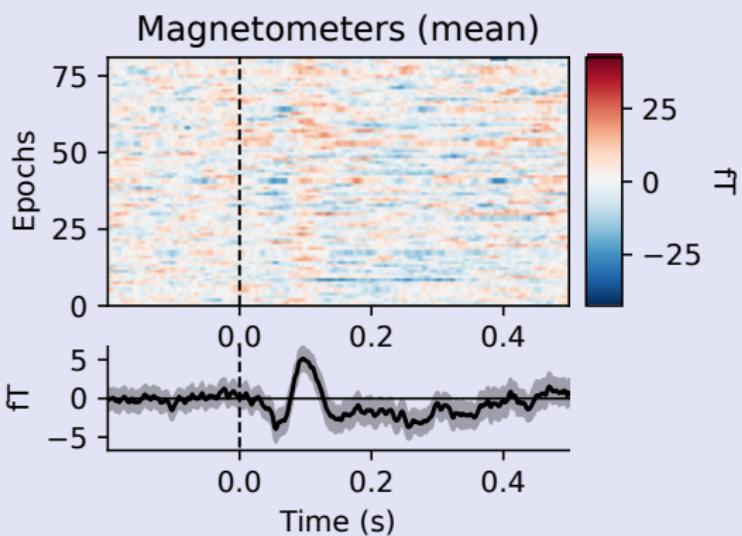
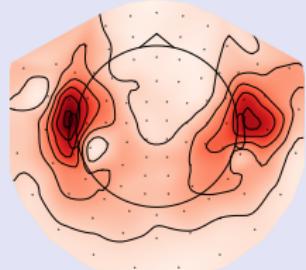
[Cole & Voytek 2017]



[Dupré la Tour et al. 2017]

- ▶ Subject is presented some stimuli – Audio, Visual, Motor, ...
- ▶ Record onset of the stimuli
- ▶ Average signal on window aligned around the stimulus

Evoked response to an auditory stimuli



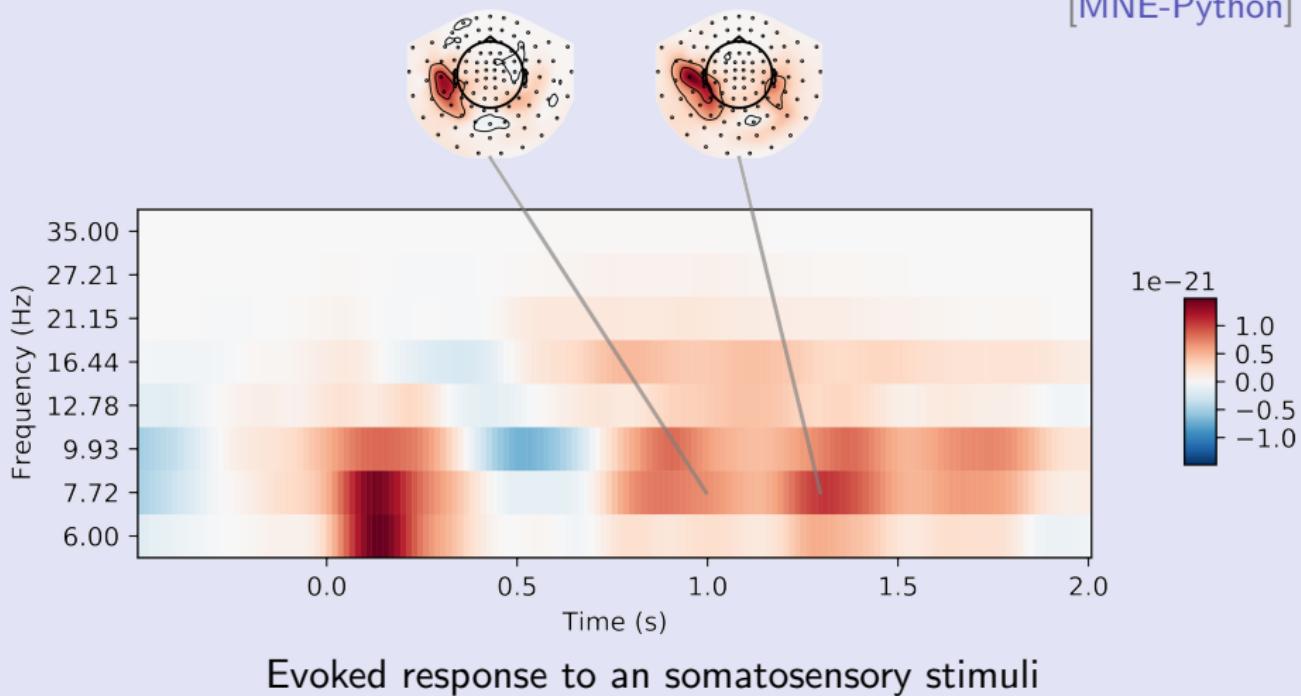
[MNE-Python]

# Repeated Stimuli – Induced Response

[Gramfort et al. 2013]

- ▶ Subject is presented some stimuli – Audio, Visual, Motor, ...
- ▶ Average PSD on window aligned around the stimulus

[MNE-Python]

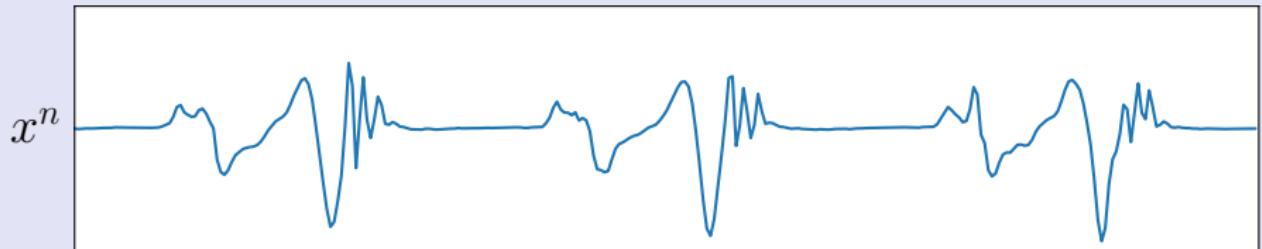


## Learning the waveform: Convolutional Dictionary Learning

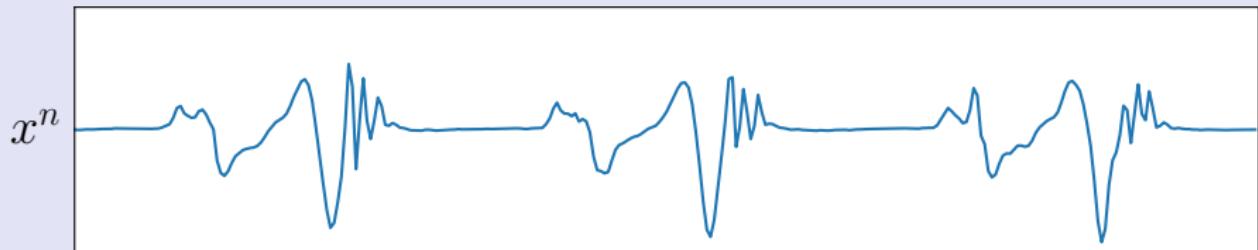
### References

- ▶ Grosse, R., Raina, R., Kwong, H., and Ng, A. Y. (2007). Shift-Invariant Sparse Coding for Audio Classification.  
*Cortex*, 8:9

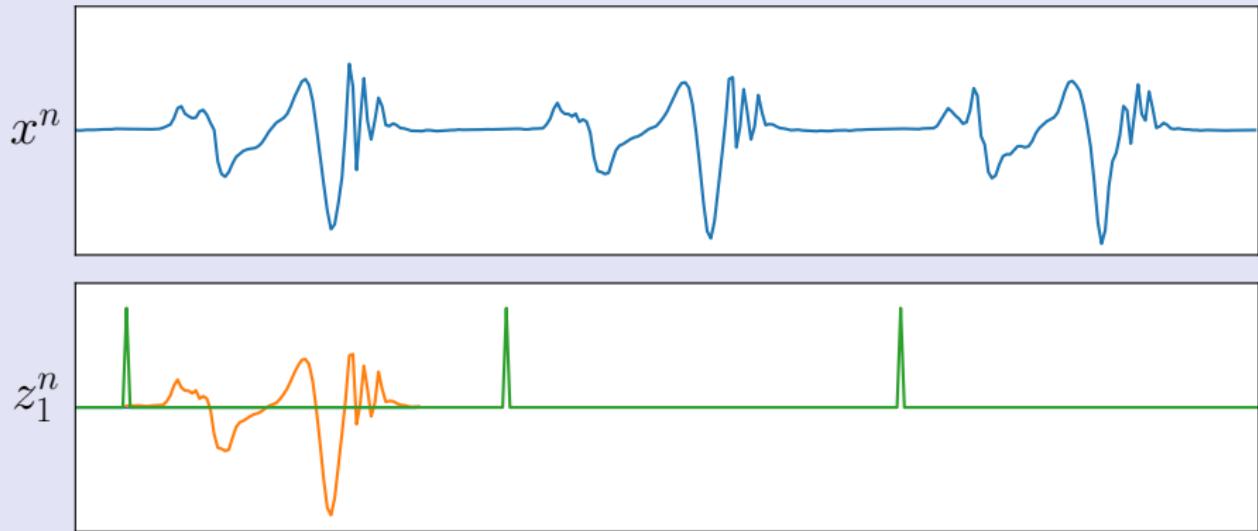
## Local structure in signals



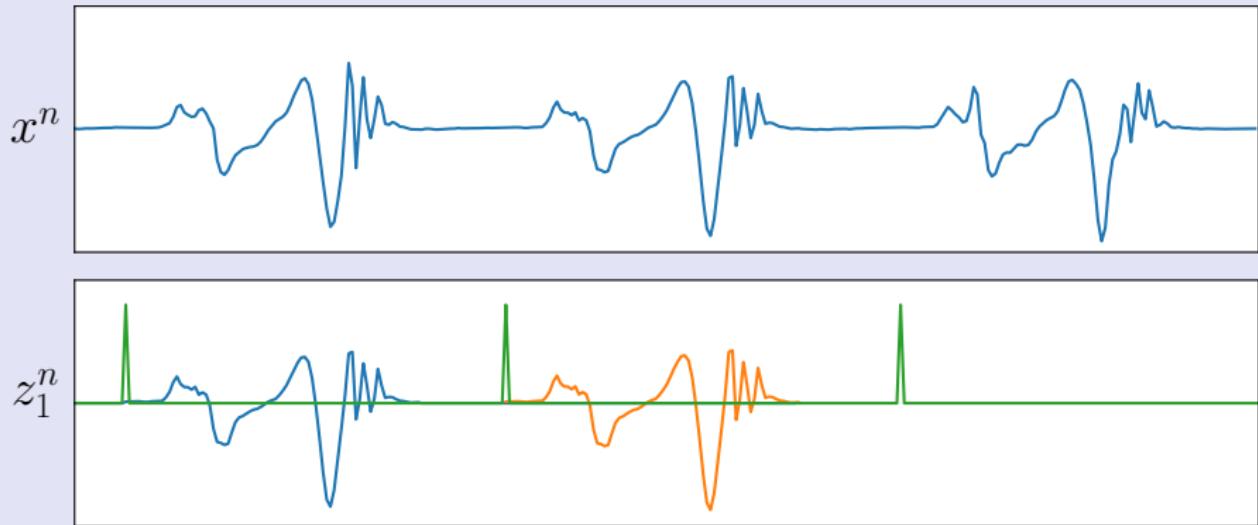
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**Key idea:** decouple the localization of the patterns and their shape



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$$x^n[t] = \sum_{k=1}^K (z_k^n * d_k)[t] + \varepsilon[t]$$

For a set of  $N$  univariate signals  $x^n$ , solve

$$\begin{aligned} \min_{d,z} \sum_{n=1}^N \frac{1}{2} \left\| \boxed{x^n} - \sum_{k=1}^K \boxed{z_k} * \boxed{d_k} \right\|_2^2 + \lambda \sum_{k=1}^K \|\boxed{z_k}\|_1, \\ \text{s.t. } \|\boxed{d_k}\|_2^2 \leq 1 \end{aligned}$$

**Hypothesis:** patterns  $d_k$  are not present everywhere in the signal. They are localized in time.

⇒ Sparse activation signals  $z$

**Technical hypothesis:** the patterns are in the  $\ell_2$ -ball:  $\|d_k\|_2^2 \leq 1$ .

## Optimization strategy

**Bi-convex:** The problem is not jointly convex in  $z_k^n$ , and  $d_k$  but it is convex in each block of coordinate.

**Alternate minimization** (a.k.a. Bloc Coordinate Descent):

- ▶ **Z-step:** given a fixed estimate of the atom, compute the activation signal  $z_k^n$  associated to each signal  $x^n$ .
- ▶ **D-step:** given a fixed estimate of the activation, update the atoms in the dictionary  $d_k$ .

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**Unrolled optimization:**

- ▶ **Z-step:** use an fixed differentiable procedure  $f(x^n, D)$ .
- ▶ **D-step:** learn  $D$  through back-propagation.

[Malezieux et al. 2022]

## How to extend CSC to multivariate signals?

We can just use multivariate convolution,

$$\underbrace{X[t]}_{\in \mathbb{R}^P} = \sum_{k=1}^K (z_k * D_k)[t] = \sum_{k=1}^K \sum_{\tau=1}^L z_k[t - \tau] \underbrace{D_k[\tau]}_{\in \mathbb{R}^P}$$

with:

- ▶  $X$  a multivariate signal of length  $T$  in  $\mathbb{R}^P$
- ▶  $D_k$  a multivariate signal of length  $L$  in  $\mathbb{R}^P$
- ▶  $z_k$  a univariate activation signal of length  $\tilde{T} = T - L + 1$

However, this model does not account for the physics of the problem.

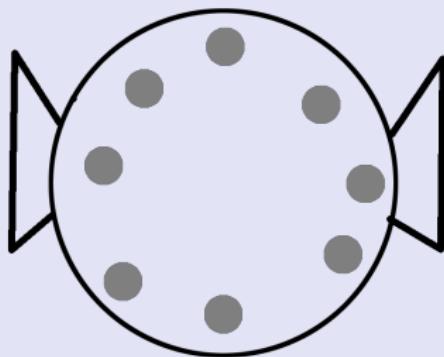
## Rank-1 constrained dictionary learning

### References

- ▶ Dupré la Tour, T., **Moreau, T.**, Jas, M., and Gramfort, A. (2018).  
Multivariate Convolutional Sparse Coding for Electromagnetic Brain Signals.  
In *Advances in Neural Information Processing Systems (NeurIPS)*, pages  
3296–3306, Montreal, Canada

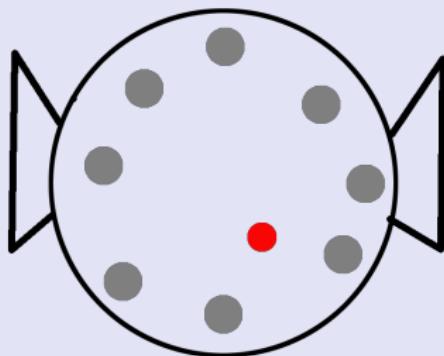
## EM wave diffusion

- ▶ Recording here with 8 sensors



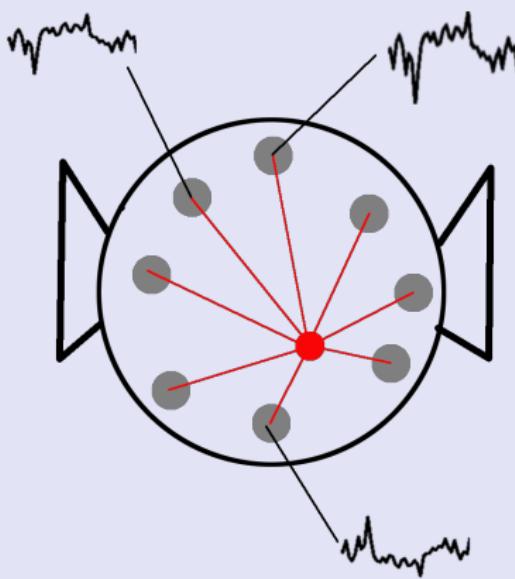
## EM wave diffusion

- ▶ Recording here with 8 sensors
- ▶ EM activity in the brain



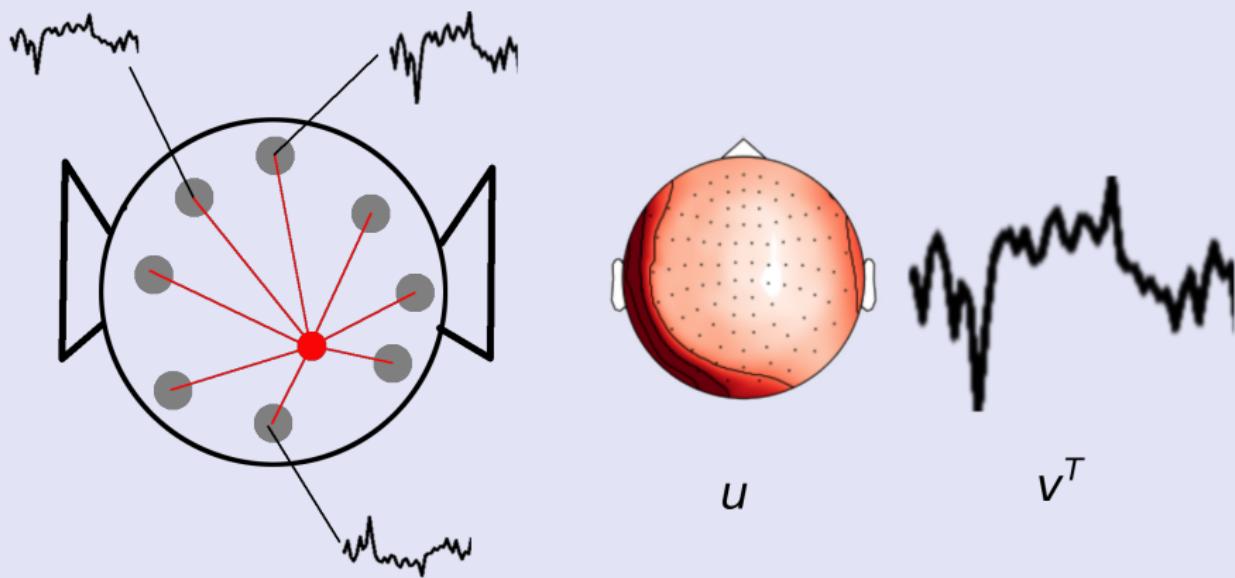
## EM wave diffusion

- ▶ Recording here with 8 sensors
- ▶ EM activity in the brain
- ▶ The electric field is spread **linearly** and **instantaneously** over all sensors (Maxwell equations)



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## Multivariate CSC with rank-1 constraint

**Idea:** Impose a rank-1 constraint on each dictionary atom  $D_k$

To make the problem tractable, use  $u_k$  and  $v_k$  s.t.  $D_k = u_k v_k^\top$ .

$$\begin{aligned} \min_{u_k, v_k, z_k^n} & \sum_{n=1}^N \frac{1}{2} \left\| X^n - \sum_{k=1}^K z_k^n * (u_k v_k^\top) \right\|_2^2 + \lambda \sum_{k=1}^K \|z_k^n\|_1, \\ \text{s.t. } & \|u_k\|_2^2 \leq 1, \|v_k\|_2^2 \leq 1 \text{ and } z_k^n \geq 0. \end{aligned} \quad (1)$$

Here,

- ▶  $u_k \in \mathbb{R}^P$  is a spatial pattern
- ▶  $v_k \in \mathbb{R}^L$  is a temporal pattern

⇒ This is a tri-convex problem

## Optimization strategy

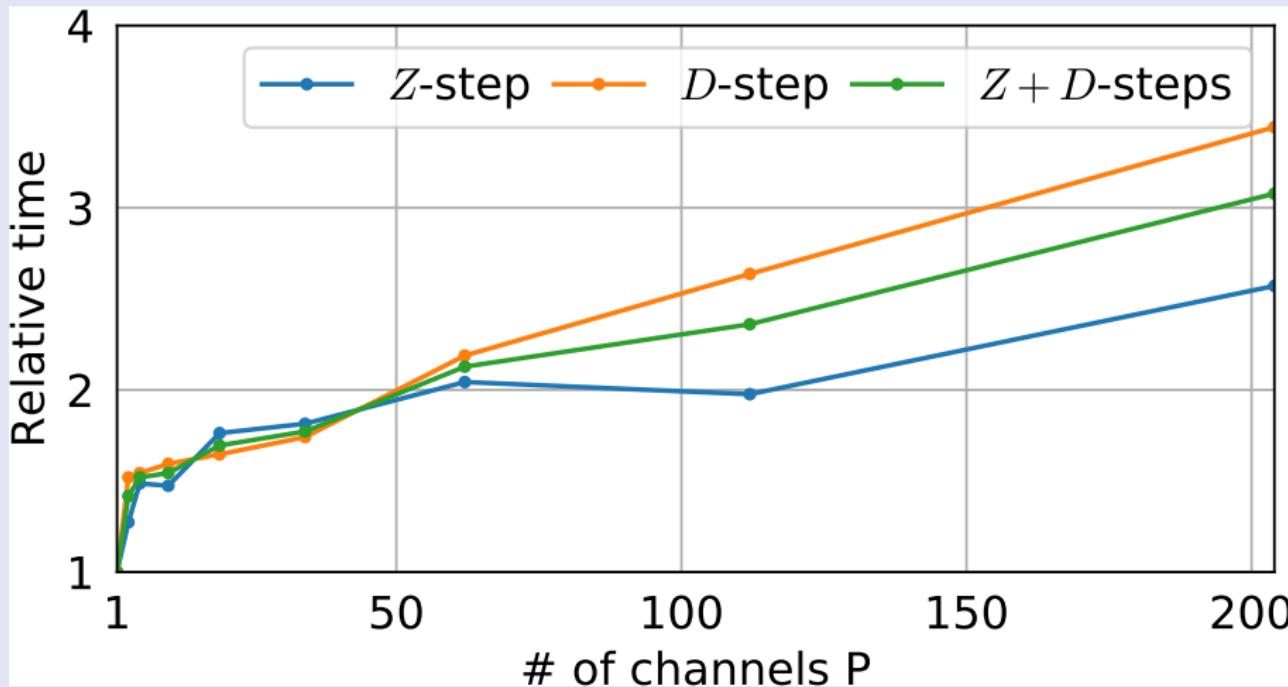
**Tri-convex:** The problem is not jointly convex in  $z_k^n$ ,  $u_k$  and  $v_k$  but it is convex in each block of coordinate.

We can use a block coordinate descent, aka alternate minimization, to converge to a local minima of this problem. The 3 following steps are applied alternatively:

- ▶ **Z-step:** given a fixed estimate of the atom, compute the activation signal  $z_k^n$  associated to each signal  $X^n$ . (LGCD)
- ▶ **u-step:** given a fixed estimate of the activation and temporal pattern, update the spatial pattern  $u_k$ . (PGD)
- ▶ **v-step:** given a fixed estimate of the activation and spatial pattern, update the temporal pattern  $v_k$ . (PGD)

## Good scaling in the number of channels $P$

Scaling relative to  $P$  on somato dataset with  $T = 134,700$ ,  $K = 2$ , and  $L = 128$



## Pattern recovery

Test the pattern recovery capabilities of our method on simulated data,

$$X^n = \sum_{k=1}^2 z_k * (u_k v_k^\top) + \mathcal{E}$$

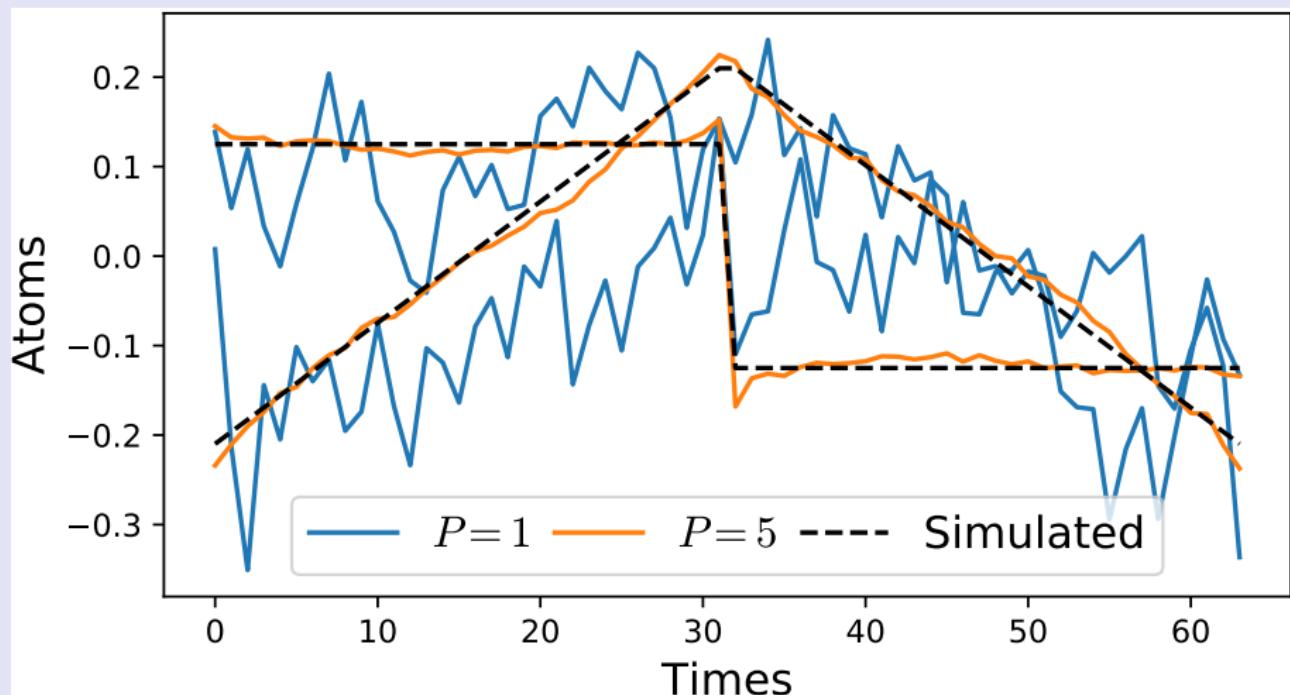
where  $(u_k, v_k)$  are chosen patterns of rank-1 and the activated coefficient  $z_k^n[t]$  are drawn uniformly and their value are uniform in  $[0, 1]$ .

The noise  $\mathcal{E}$  is generated as a gaussian white noise with variance  $\sigma$ .

We set  $N = 100$ ,  $L = 64$  and  $\tilde{T} = 640$

## Pattern recovery

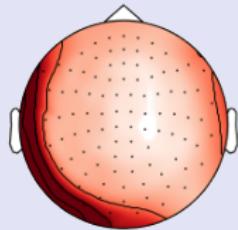
Patterns recovered with  $P = 1$  and  $P = 5$ . The signals were generated with the two simulated temporal patterns and with  $\sigma = 10^{-3}$ .



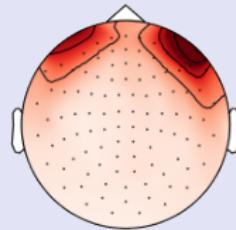
## MNE sample data

A selection of temporal waveforms of the atoms learned on the MNE sample dataset.

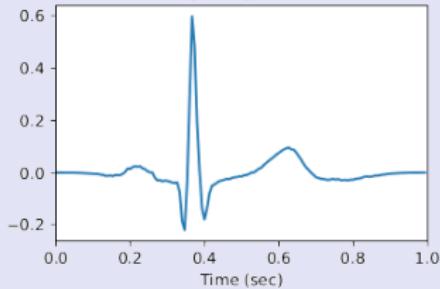
Spatial pattern 0  
Explained variance 5.62 %



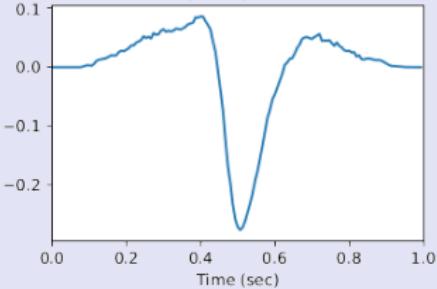
Spatial pattern 1  
Explained variance 2.38 %



Temporal pattern 0

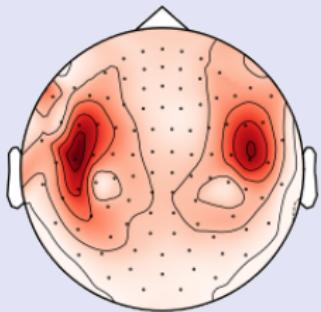


Temporal pattern 1

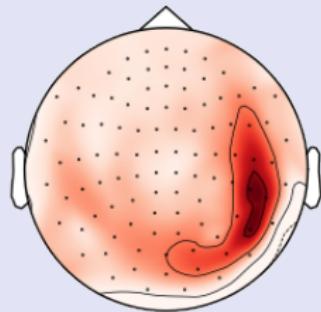


## Learned atoms – Evoked response

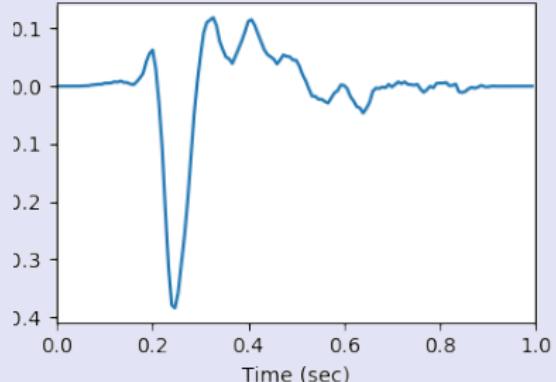
Spatial pattern 3



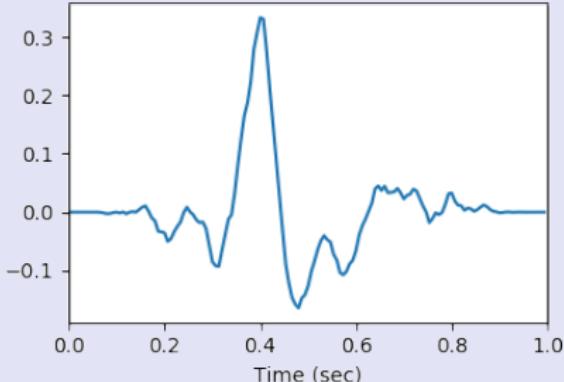
Spatial pattern 15



Temporal pattern 3

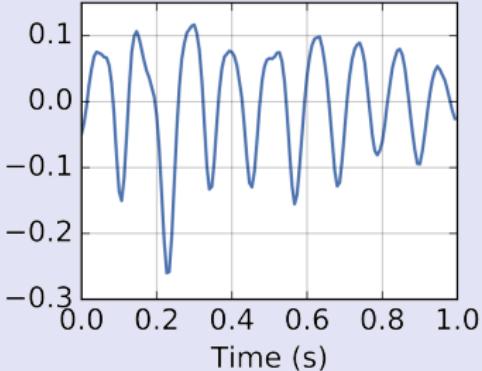


Temporal pattern 15

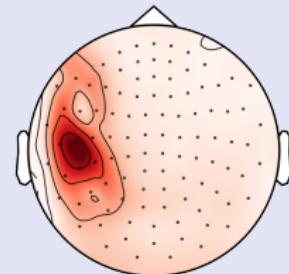


## Learned atoms – Induced responses

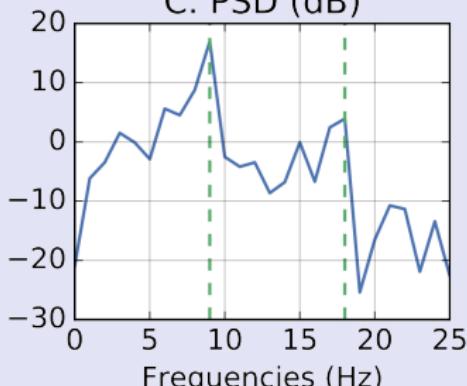
A. Temporal waveform



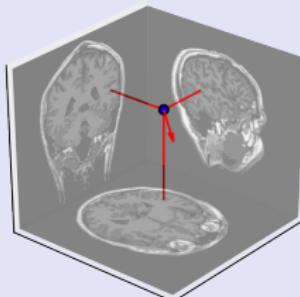
B. Spatial pattern



C. PSD (dB)



D. Dipole fit



# alphaCSC: Convolution sparse coding for time-series



This is a library to perform shift-invariant sparse dictionary learning (CSC), on time-series data. It includes a number of different r

1. univariate CSC
2. multivariate CSC
3. multivariate CSC with a rank-1 constraint [1]
4. univariate CSC with an alpha-stable distribution [2]

A mathematical descriptions of these models is available [in the documentation](#).

Python code online:  
<https://alphacsc.github.io>

`pip install alphacsc`

## Installation

To install this package, the easiest way is using [pip](#). It will install this package and its dependencies. The `setup.py` depends on `numpy` and `cython` for the installation so it is advised to install them beforehand. To install this package, please run one of the two commands:

(Latest stable version)

```
pip install alphacsc
```

(Development version)

```
pip install git+https://github.com/alphacsc/alphacsc.git#egg=alphacsc
```

(Dicodile backend)

```
pip install numpy cython  
pip install alphacsc[dicodile]
```

Examples reproduce figures from this talk!

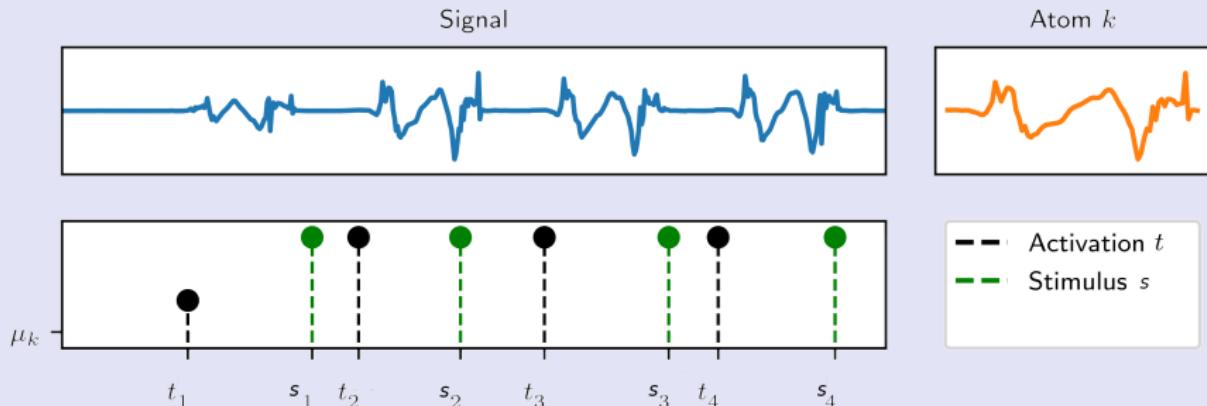
## Modeling stimuli induced patterns with Point Processes

### References

- ▶ Allain, C., Gramfort, A., and **Moreau, T.** (2022). DriPP: Driven Point Process to Model Stimuli Induced Patterns in M/EEG Signals.  
In *International Conference on Learning Representations (ICLR)*
- ▶ Staerman, G., Allain, C., Gramfort, A., and **Moreau, T.** (2023). FaDIIn: Fast Discretized Inference for Hawkes Processes with General Parametric Kernels.  
In *International Conference on Machine Learning (ICML)*, Honolulu, HI, USA. PMLR

## Stimuli Induced Patterns

- ▶ Manual pattern identification
- ▶ No quantification of how stimuli influence patterns activation.



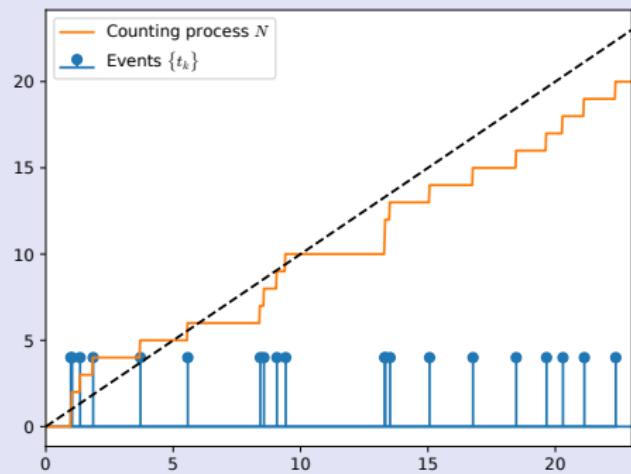
Activations and stimuli can be seen as *Point Processes*.

- ▶ Stochastic model for stream of events
- ▶ Time of arrival  $\{t_k\}$  associated with counting process  $N(t)$
- ▶ Characterized by the intensity:

$$\lambda(t|\mathcal{F}_t) = \lim_{dt \rightarrow 0} \frac{P(N(t+dt) - N(t) = 1 | \mathcal{F}_t)}{dt}$$

Poisson process with constant probability of arrival

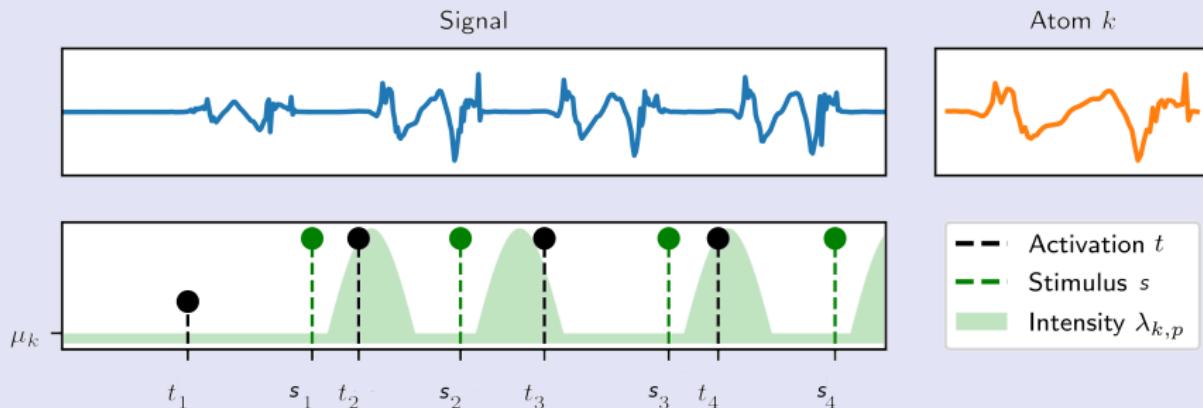
$$\lambda(t) = \mu_0$$



# DriPP – Driven Point Process

**Idea:** Model the probability of activation  $\{t_k\}$  depending on the PP from the stimuli  $\{s_p\}$ .

$$\lambda(t|\mathcal{F}_t) = \lambda(t|\{s_p; s_p < t\}) = \mu_0 + \sum_{s_p < t} \kappa(t - s_p)$$

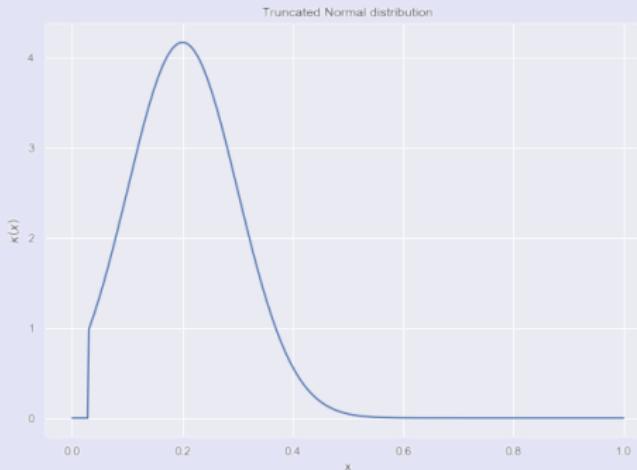


# Modeling latency

Choosing a model for stimuli based modeling:

$$\lambda(t|\mathcal{F}_t) = \mu_0 + \sum_{s_p < t} \alpha \kappa(t - s_p)$$

- ▶  $\mu_0 \geq 0$ : spontaneous activity.
- ▶  $\alpha \geq 0$ : allow for stimuli to have no effect.
- ▶  $\kappa(\tau)$ : pdf of a truncated Gaussian  $\mathcal{N}(m, \sigma^2)$  to model latency.



## Parameters estimation

The negative log-likelihood of the model can be computed using the intensity  $\lambda$ :

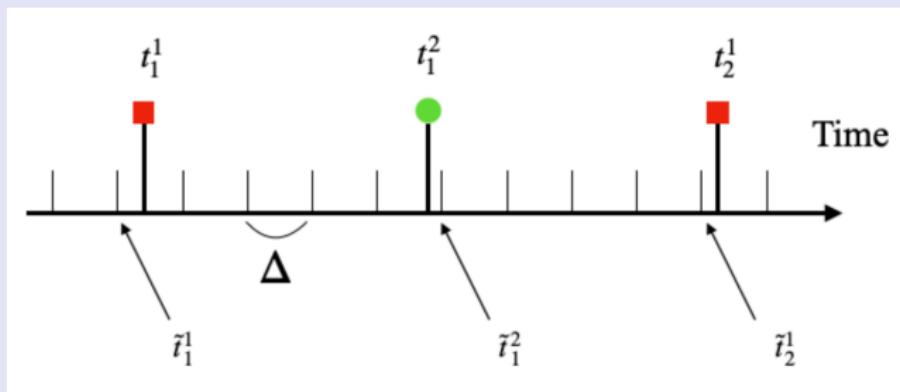
$$\begin{aligned}\mathcal{L}(\{t_k\}; \Theta) &= \int_0^T \lambda(t) dt - \sum_{t_k} \log \lambda(t_k) \\ &= \mu_0 T + \alpha |\{t_k\}| - \sum_{t_k} \log(\mu_0 \sum_{s_p < t_k} \alpha \kappa(t_k - s_p))\end{aligned}$$

with  $\Theta = (\mu_0, \alpha, m, \sigma^2)$

⇒ Parameter estimation with an EM algorithm.

- ▶ Slow EM algorithm
- ▶ Not general for parametric kernels.

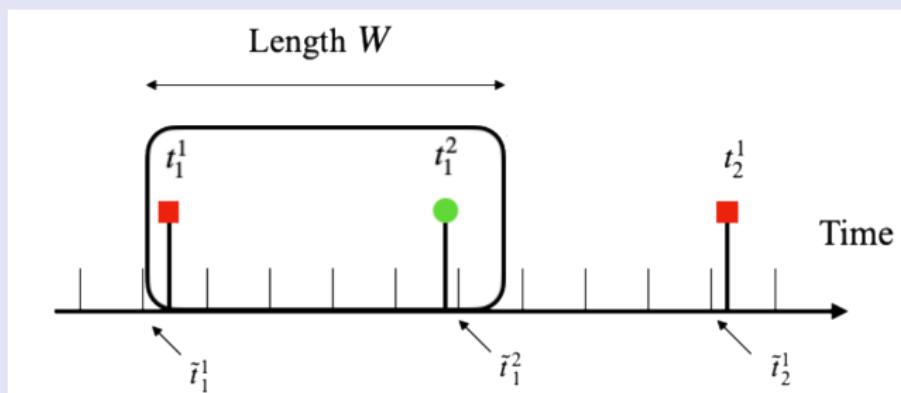
**FaDIn** Inference method for general parametric kernels



**Discretization**

**FaDIn** Inference method for general parametric kernels

## ► Discretization



**Finite support**

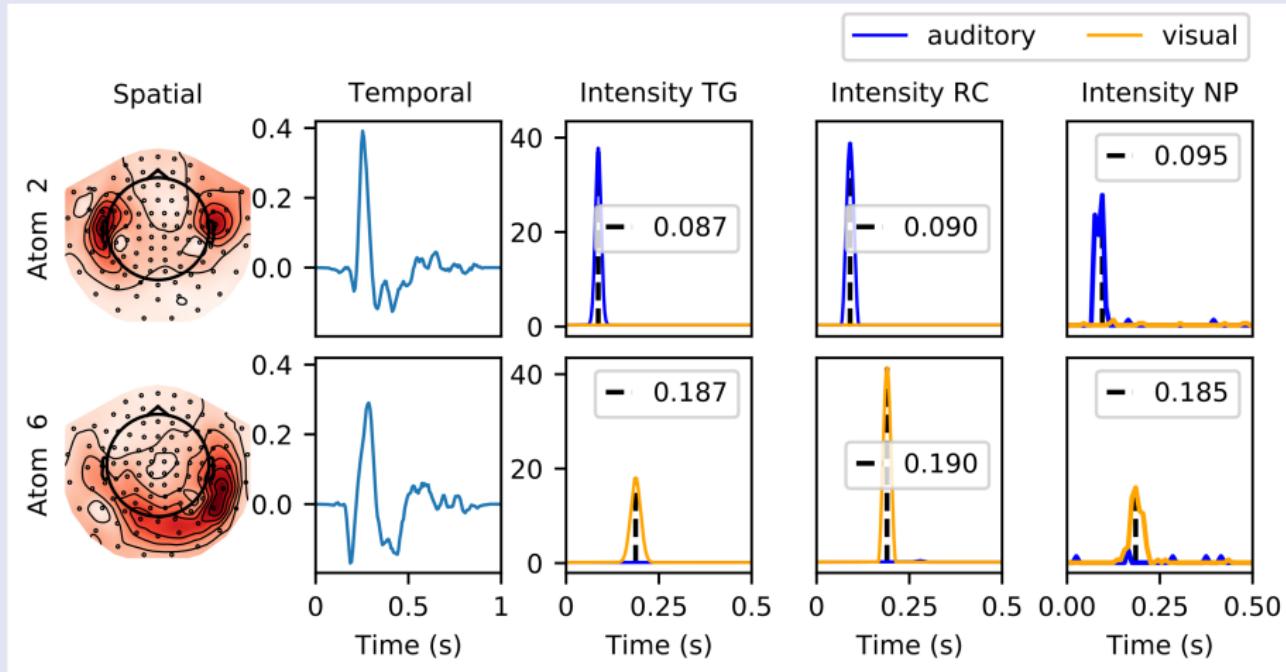
## FaDIn Inference method for general parametric kernels

- ▶ Discretization
- ▶ Finite support
- ▶  $\ell_2$  loss

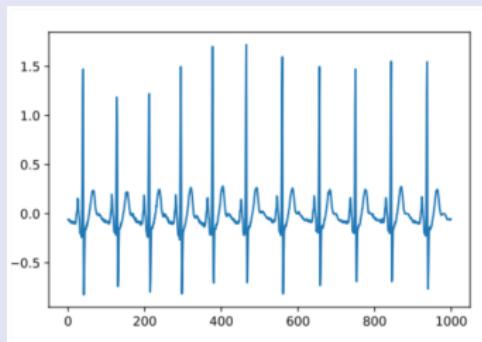
$$\mathcal{L}(\{t_k\}; \theta) = \sum_{t=0}^T \frac{1}{2} \|z[t] - (z * k)[t]\|_2^2$$

with  $z[t] = 1$  if  $t \in \{t_k\}$ , 0 otherwise.

## Results for evoked atoms - samples



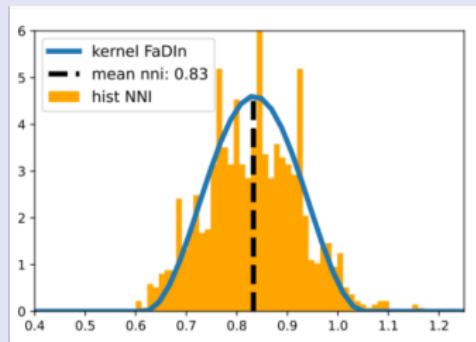
# Results for heart rate variability



CDL + FaDIn



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## Conclusion

---

- ▶ CDL can learn recurring patterns in multivariate signals.
- ▶ Converts the signal into a stream of events.
- ▶ PP framework can model the activation distribution.

### Limitations and on-going work:

- ▶ Not easy to apply to population level.
- ▶ DriPP does not model inhibition.
- ▶ CDL and PP are separated.

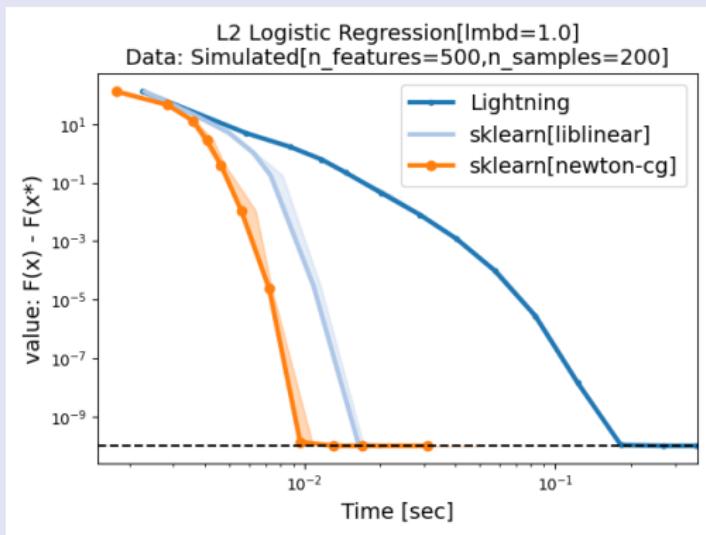
## Benchopt

### References

- ▶ **Moreau, T.**, Massias, M., Gramfort, A., Ablin, P., Bannier, P.-A., Charlier, B., Dagréou, M., la Tour, T. D., Durif, G., Dantas, C. F., Klopfenstein, Q., Larsson, J., Lai, E., Lefort, T., Malézieux, B., Moufad, B., Nguyen, B. T., Rakotomamonjy, A., Ramzi, Z., Salmon, J., and Vaiter, S. (2022). [Benchopt: Reproducible, efficient and collaborative optimization benchmarks](#). In *Advances in Neural Information Processing Systems (NeurIPS)*, volume 36, New-Orlean, LA, USA. Curran Associates, Inc.

Doing a benchmark for the  $\ell_2$  regularized logistic regression with multiple solvers and datasets is now easy as calling:

```
git clone https://github.com/benchopt/benchmark_logreg_l2  
benchopt run ./benchmark_logreg_l2
```



## Benchmark: principle

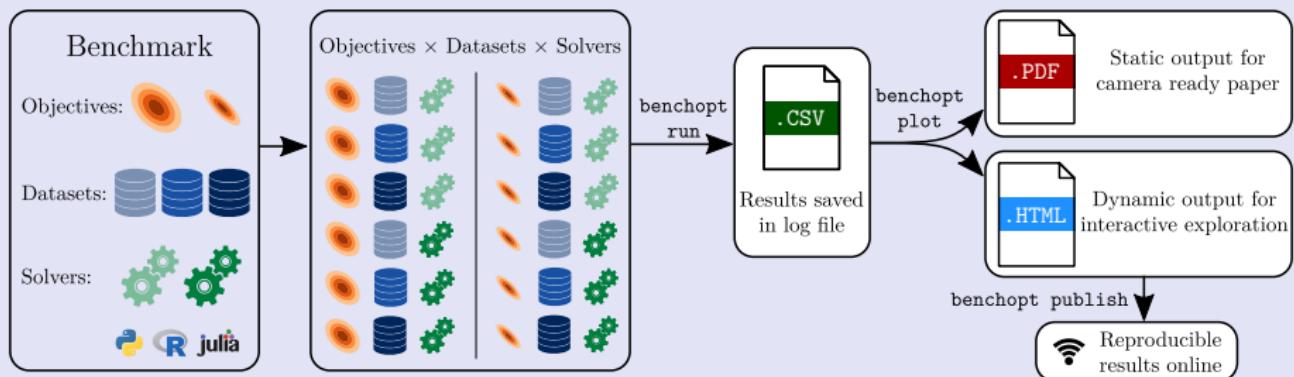
A benchmark is a directory with:

- ▶ An `objective.py` file with an `Objective`
- ▶ A directory `solvers` with one file per `Solver`
- ▶ A directory `datasets` with `Dataset` generators/fetchers

```
my_benchmark/
├── README.rst
├── datasets
│   ├── simulated.py  # some dataset
│   └── real.py  # some dataset
└── objective.py  # contains the definition of the objective
└── solvers
    ├── solver1.py  # some solver
    └── solver2.py  # some solver
```

The `benchopt` client runs a cross product and generates a csv file + convergence plots like above.

# Benchopt: principle



⇒ Each object can be parametrized so multiple scenario can be tested.

## Automatizing tasks:

- ▶ Automatic installation of competitors solvers.
- ▶ Parametrized datasets, objectives and solvers and run on cross products.
- ▶ Make sure to quantify the variance.
- ▶ Automatic caching.
- ▶ Interactive visualization of the results
- ▶ Automatic parallelization, run on SLURM,
- ▶ ...?

# Thanks for your attention!

Code available online:

⌚ **alphacsc** : [alphacsc.github.io](https://alphacsc.github.io)

⌚ **DriPP** : [github.com/CedricAllain/dripp](https://github.com/CedricAllain/dripp)

⌚ **benchopt** : [benchopt.github.io](https://benchopt.github.io)

Slides are on my web page:

🌐 [tommoral.github.io](https://tommoral.github.io)

⌚ [@tomamoral](#)

## Z-step: Locally greedy coordinate descent (LGCD)

$N$  independent problem such that

$$\min_{z_k^n \geq 0} \frac{1}{2} \left\| X^n - \sum_{k=1}^K z_k^n * D_k \right\|_2^2 + \lambda \sum_{k=1}^K \|z_k^n\|_1.$$

This problem is convex in  $z_k$  and can be solved with different techniques:

- ▶ FISTA [Chalasani et al., 2013]
- ▶ ADMM [Bristow et al., 2013]
- ▶ L-BFGS [Jas et al., 2017]
- ▶ Greedy CD [Kavukcuoglu et al., 2010]

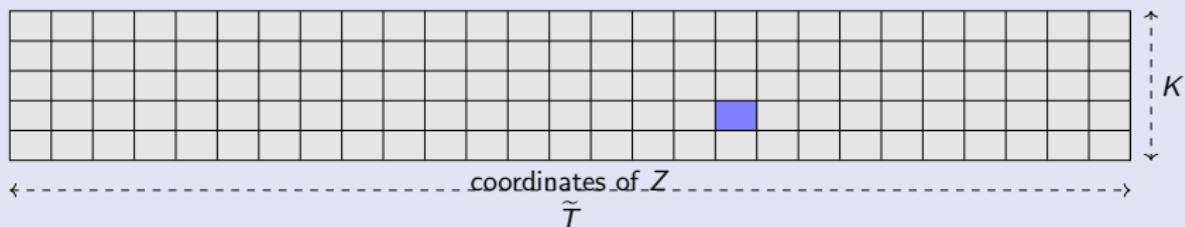
⇒ These methods can be slow for long signals as the complexity of each iteration is at least linear in the length of the signal.

## Z-step: Locally greedy coordinate descent (LGCD)

**Coordinate Descent:** only 1 coordinate is updated at each iteration:

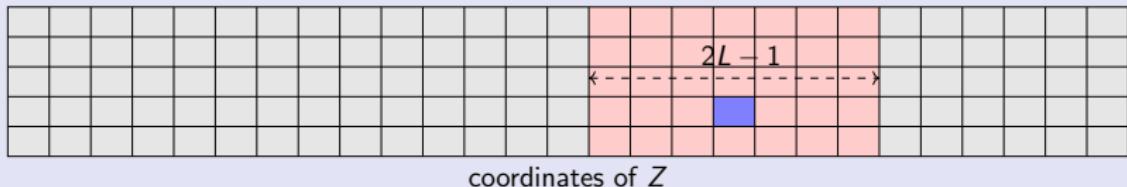
1. The coordinate  $z_{k_0}[t_0]$  is updated to its optimal value  $z'_{k_0}[t_0]$  when all other coordinate are fixed.
2. The updated coordinate is chosen
  - ▶ Cyclic:  $\mathcal{O}(1)$  [Friedman et al., 2007]
  - ▶ Randomized:  $\mathcal{O}(1)$  [Nesterov, 2010]
  - ▶ Greedy:  $\mathcal{O}(K\tilde{T})$  by maximizing  $|z_k[t] - z'_k[t]|$  [Osher and Li, 2009]
  - ▶ Locally Greedy:  $\mathcal{O}(KL)$  by maximizing  $|z_k[t] - z'_k[t]|$  on a window [Moreau et al., 2018]

We introduced the LGCD method which is an extension of GCD.



GCD has  $\mathcal{O}(K\tilde{T})$  computational complexity.

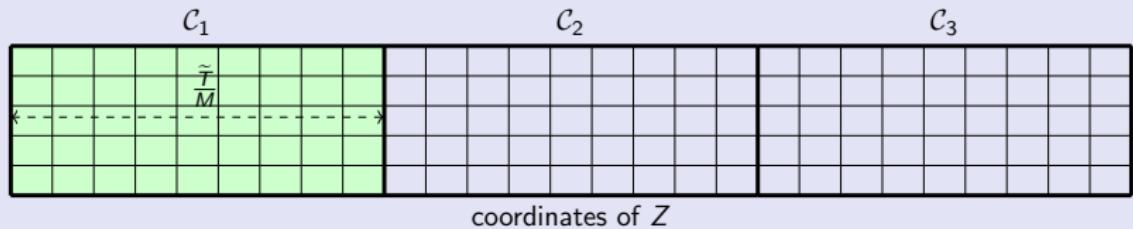
We introduced the LGCD method which is an extension of GCD.



GCD has  $\mathcal{O}(K\tilde{T})$  computational complexity.

But the update itself has complexity  $\mathcal{O}(KL)$

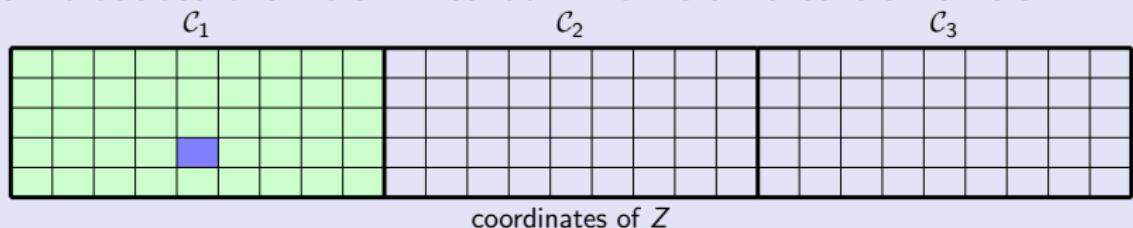
We introduced the LGCD method which is an extension of GCD.



With a partition  $\mathcal{C}_m$  of the signal domain  $[1, K] \times [0, \tilde{T}]$ ,

$$\mathcal{C}_m = [1, K] \times \left[ \frac{(m-1)\tilde{T}}{M}, \frac{m\tilde{T}}{M} \right]$$

We introduced the LGCD method which is an extension of GCD.



With a partition  $\mathcal{C}_m$  of the signal domain  $[1, K] \times [0, \tilde{T}]$ ,

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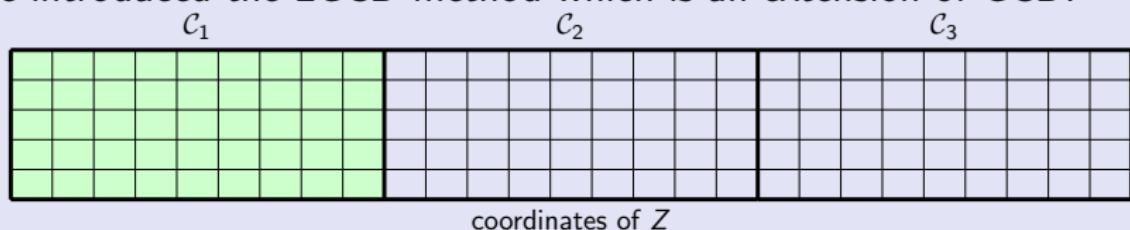
The coordinate to update is chosen greedily on a sub-domain  $\mathcal{C}_m$

$$\frac{\tilde{T}}{M} = 2L - 1 \Rightarrow \mathcal{O}(\text{Coordinate selection}) = \mathcal{O}(\text{Coordinate Update})$$

The overall iteration complexity is  $\mathcal{O}(KL)$  instead of  $\mathcal{O}(K\tilde{T})$ .

$\Rightarrow$  Efficient for sparse  $Z$

We introduced the LGCD method which is an extension of GCD.



With a partition  $\mathcal{C}_m$  of the signal domain  $[1, K] \times [0, \tilde{T}]$ ,

$$\mathcal{C}_m = [1, K] \times \left[ \frac{(m-1)\tilde{T}}{M}, \frac{m\tilde{T}}{M} \right]$$

The coordinate to update is chosen greedily on a sub-domain  $\mathcal{C}_m$

$$\frac{\tilde{T}}{M} = 2L - 1 \Rightarrow \mathcal{O}(\text{Coordinate selection}) = \mathcal{O}(\text{Coordinate Update})$$

The overall iteration complexity is  $\mathcal{O}(KL)$  instead of  $\mathcal{O}(K\tilde{T})$ .

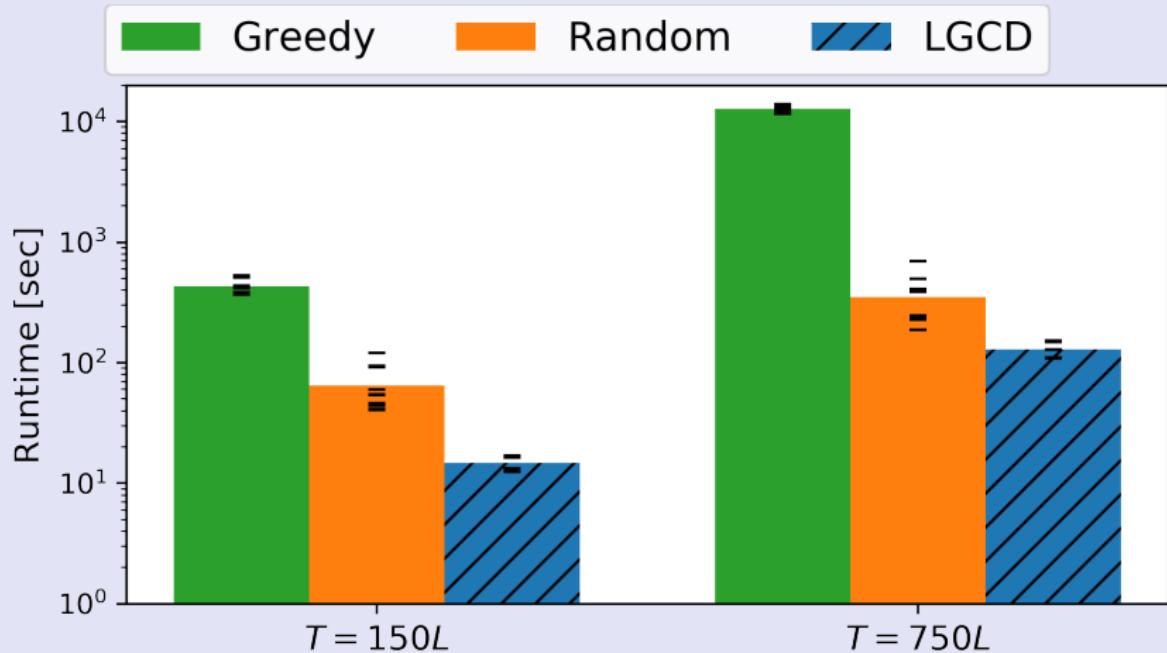
$\Rightarrow$  Efficient for sparse  $Z$

$\Rightarrow$  Can be efficiently parallelized.

## Fast optimization

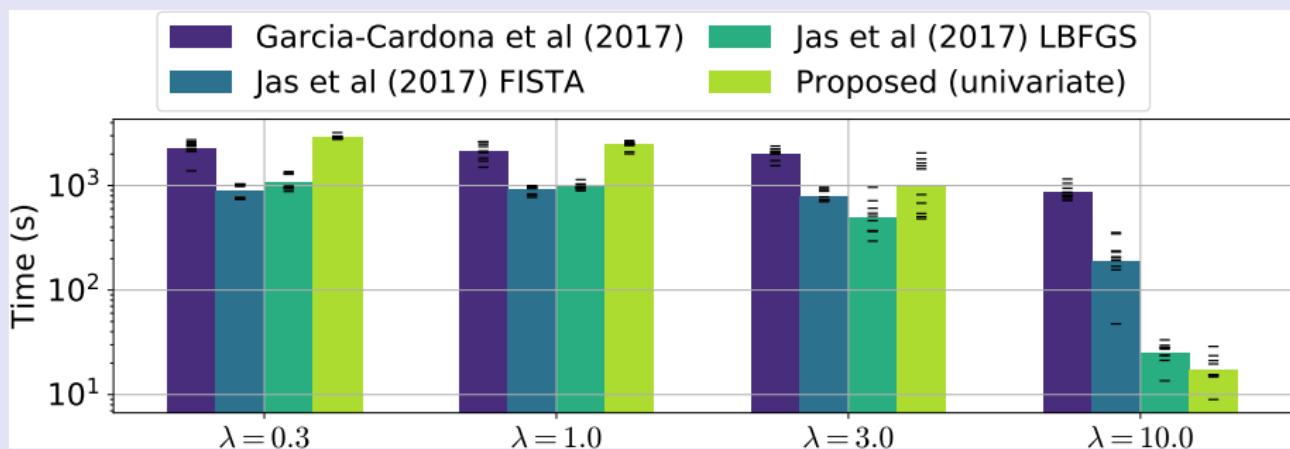
Comparison of the coordinate selection strategy for CD on simulated signals

We set  $K = 10$ ,  $L = 150$ ,  $\lambda = 0.1\lambda_{\max}$



# Fast optimization

Comparison with univariate methods on somato dataset with  
 $T = 134,700$ ,  $K = 8$  and  $L = 128$



# Fast optimization

Comparison with multivariate methods on somato dataset with  
 $T = 134,700$ ,  $K = 8$ ,  $P = 5$  and  $L = 128$

