

Neuronal Spike Sorting for Multi-Day Multielectrode Array Recordings

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- 1 Method overview
- 2 Walkthrough
- 3 Some extra tips



- Neurons appear and disappear

Unique challenges of circadian spike sorting



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- Spike shape changes day-to-day

Unique challenges of circadian spike sorting



- Neurons appear and disappear
- Spike shape changes day-to-day
- Massive amounts of data



- 1 Binning into days for sorting / decision tree sorting

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- 2 Correlating spike shape between days
- 3 File formatting, creating a sorting tree based on a subsample of data (as little as 5%)

Manual

Reasoning

You need to put your files somewhere. Put them in a folder called `mcd`.

Within MCD, separate these files by days (e.g. 0day, 1day ...)
IMPORTANT: mcd files must be numbered sequentially (start with `*0000.mcd`).

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Requires: `experiment/`

Estimated time: <1 minute

Resulting directory: `experiment/mcd`

```
create_numpy_database.py
```

Reasoning

Loading .mcd files takes a long time.

Loading numpy files takes almost no time.

To take data from one electrode, you must load many .mcd files.

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Requires: `experiment/mcd`

Estimated time: 1-2 hours total (less if parallel)

Resulting directory: `experiment/numpy_database`

`subsample_mcd_array.py`

Reasoning

Clustering can take a long time if trying to cluster everything at once.

After hundreds of thousands of spikes, everything begins to look the same.

Instead, cluster using a limited number (about 1 mcd file, tuneable) of spikes.

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Instead, cluster using a limited number (about 1 mcd file, tuneable) of spikes.

Requires: `experiment/numpy_database`

Estimated time: <1 hour

Resulting directory: `experiment/subsampled_test_sets`

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sort_electrodes.py
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Using the subsampled data for an electrode, first identify the noise and remove it. Then sort resulting spikes to build a decision tree. Classify all spikes for the electrode based on this tree.

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Requires: `experiment/subsampled_test_sets`

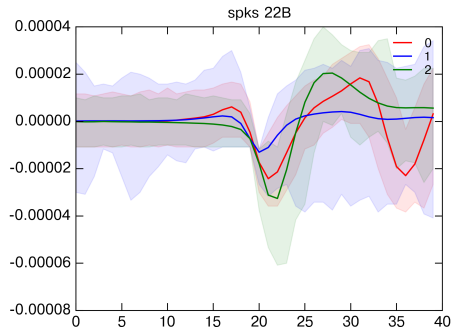
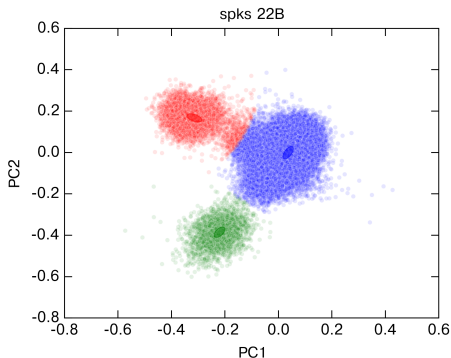
Estimated time: 1 hour per day

Resulting directory: `experiment/numpy_neurons`

Clustering: removing noise



Take subsamples, cluster using Gaussian Mixture Model (GMM), sort by expectation-maximization, delete the noise cluster.





Decision tree clustering:

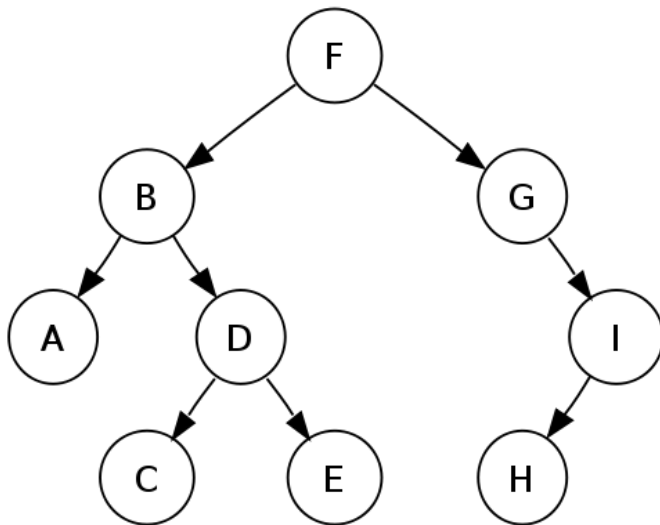
- 1 PCA, GMM cluster
- 2 EM sort
- 3 Take each cluster, PCA, GMM cluster
- 4 Take each cluster, EM sort
- 5 Continue until each dataset is a single cluster.

Record all PCA data and save it to construct a tree.

Clustering: building a classifier



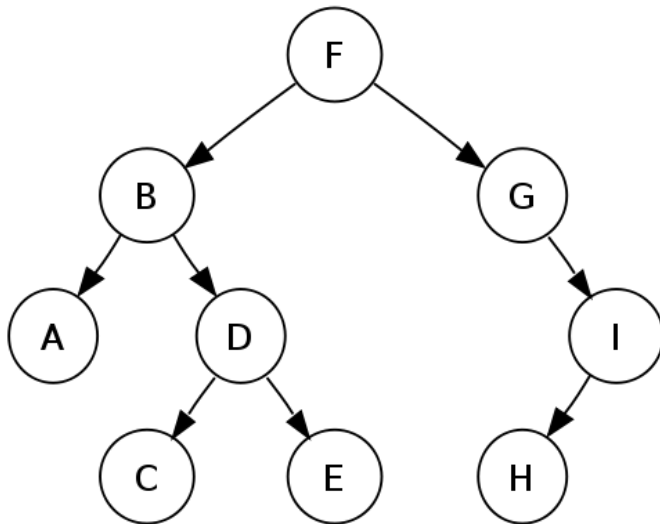
Decision tree clustering:



Clustering: Classification



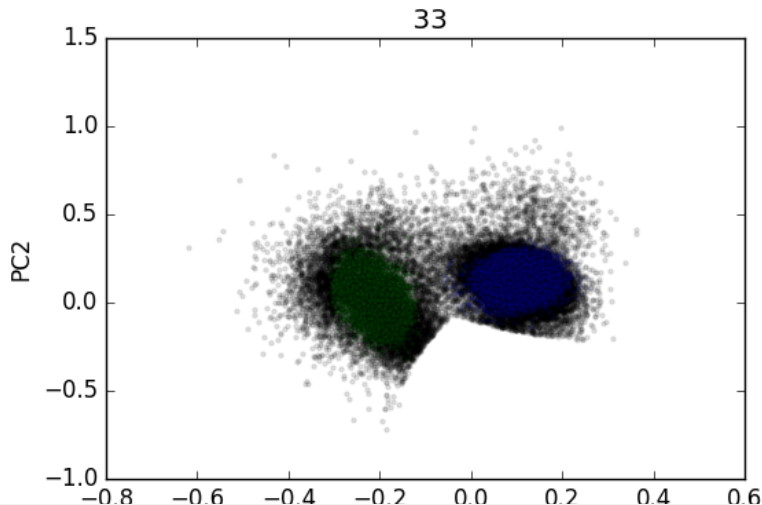
Take all data and sort based on the tree.



Clustering: Classification



Use mahalanobis distance to classify spikes in terminal clusters.
“False negatives are better than false positives.”



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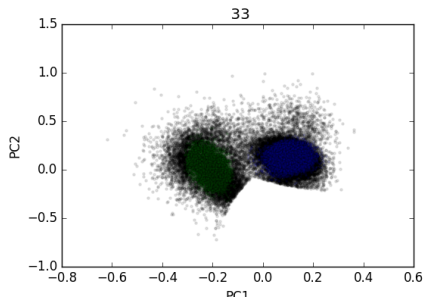
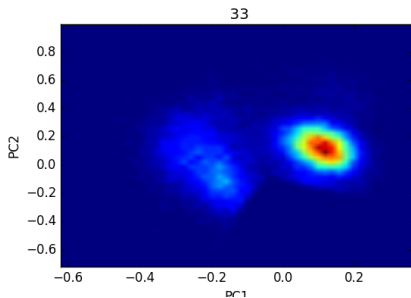
Requires: `experiment/numpy_neurons`

Estimated time: 1-2 minutes

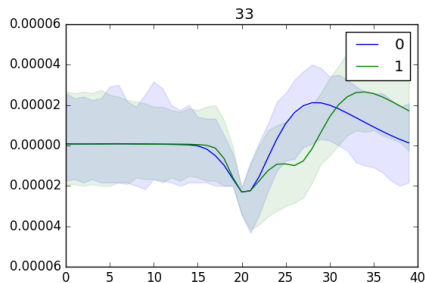
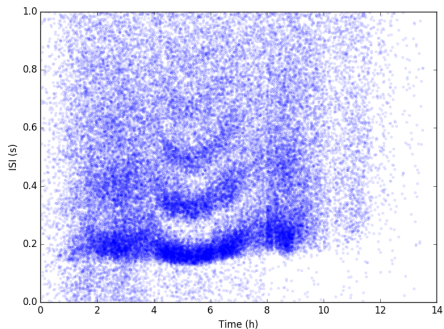
Resulting directory:

`experiment/numpy_neurons/combined_neurons`

Piecing it back together



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github.com/JohnAbel/spikesort



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- Don't fly blind! Check the sorting once it's finished and email me for help if you need. A variety of plots are generated.
- This isn't magic, and even though it works more often than not, it sometimes sorts badly.