

Manual parts of spike-sorting

1. Are these waveforms from a neuron or noise?
2. Are the selected waveforms sufficiently homogenous/isolated?
 - Isolated from waveforms from other units
 - Isolated from noise

Automating the Manual parts of spike-sorting

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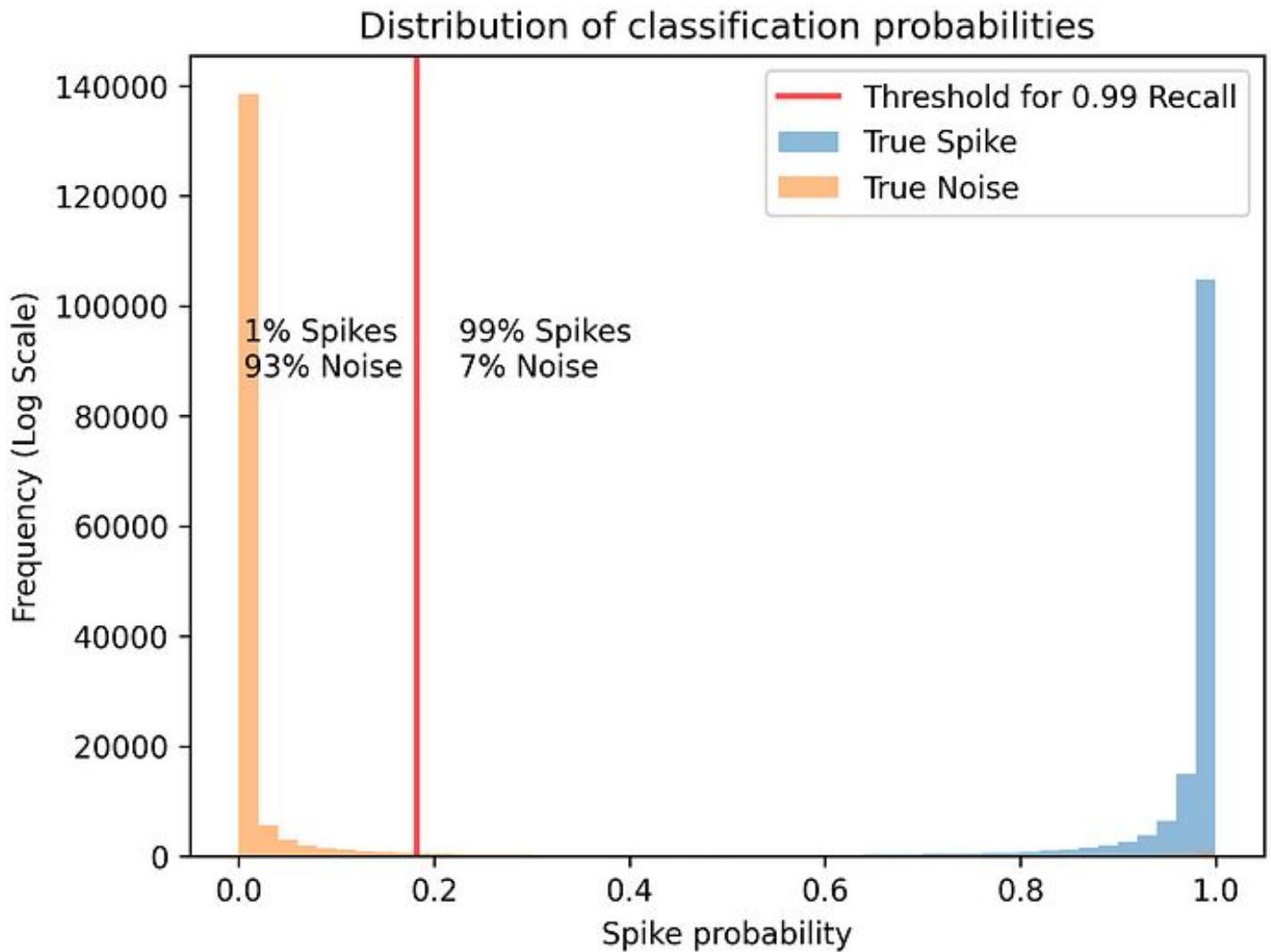
Potential Solutions

1. Train a classifier to say neuron or noise
2. Use a Bayesian Gaussian Mixture model to determine "best" number of clusters
 - o Implicitly, this will suggest a clustering configuration with the best isolation given the data
3. Select the right clusters

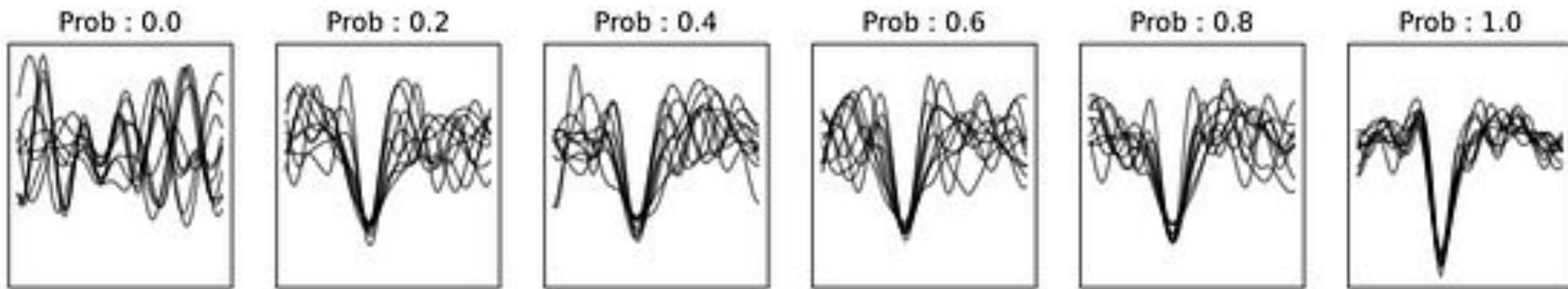
Maximizing Recall (True spikes kept)

1. Making and testing the classifier

- XGBoost classifier
 - Fast
 - Effective
- ~2.5 million waveforms
 - ~48% true spikes
 - ~52% noise
 - manually labelled

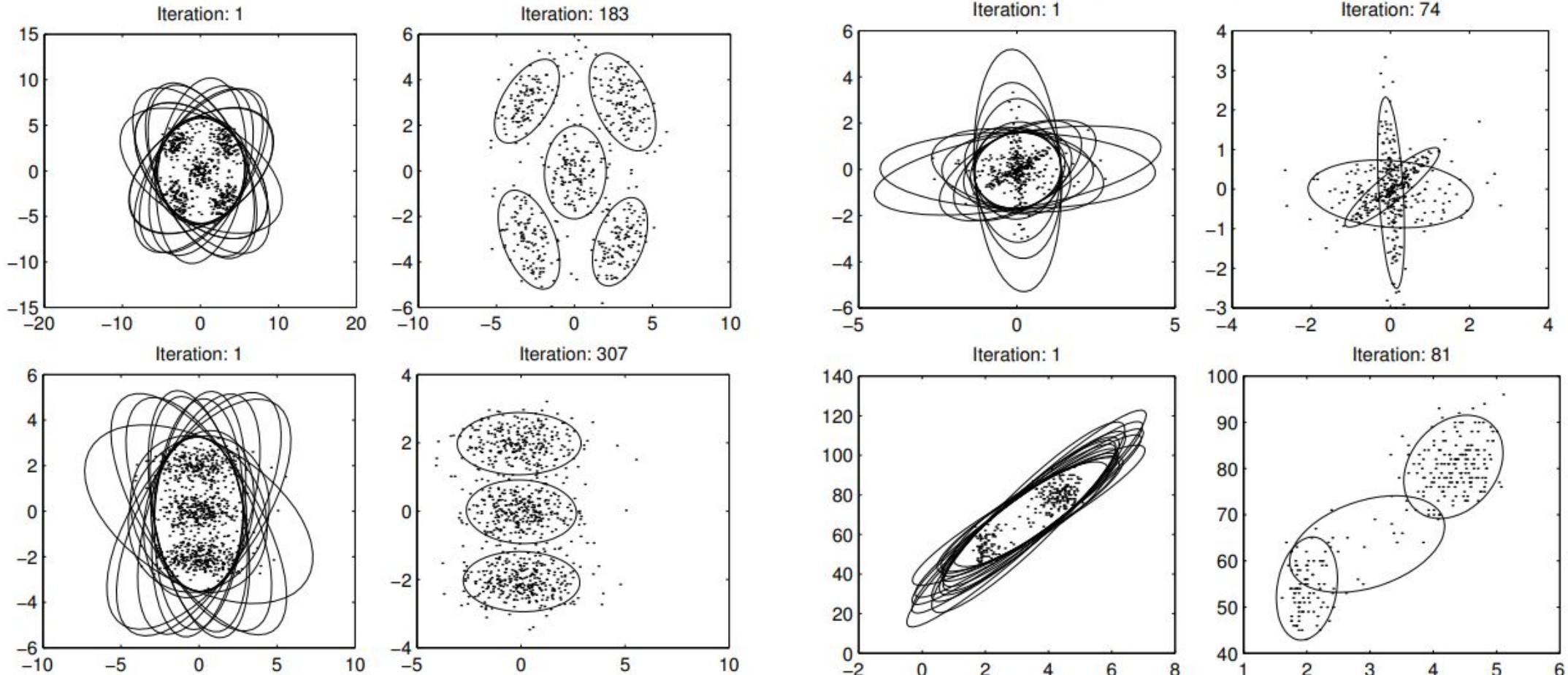


1. Making and testing the classifier



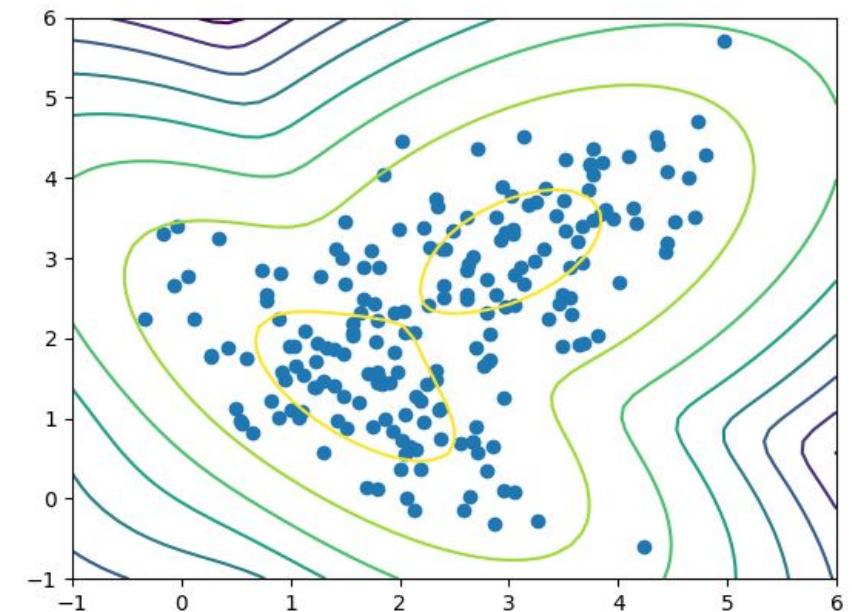
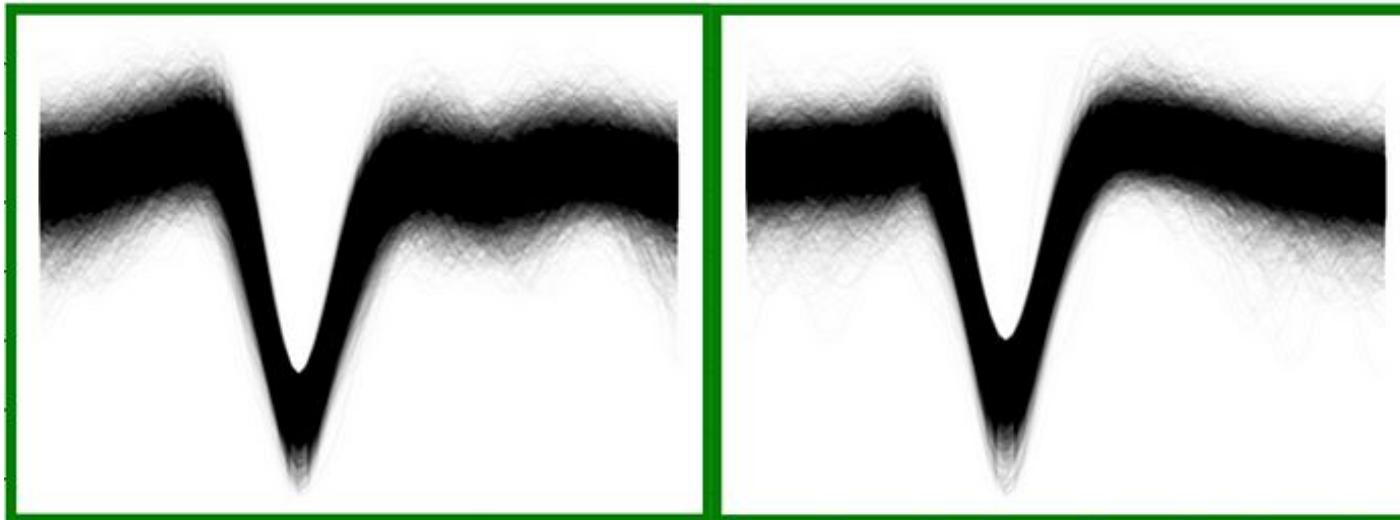
2. Automating clustering of waveforms

- Bayesian Gaussian Mixture Model
 - Only have to specify maximum allowed number of clusters
 - Sklearn.mixture.BayesianGaussianMixture



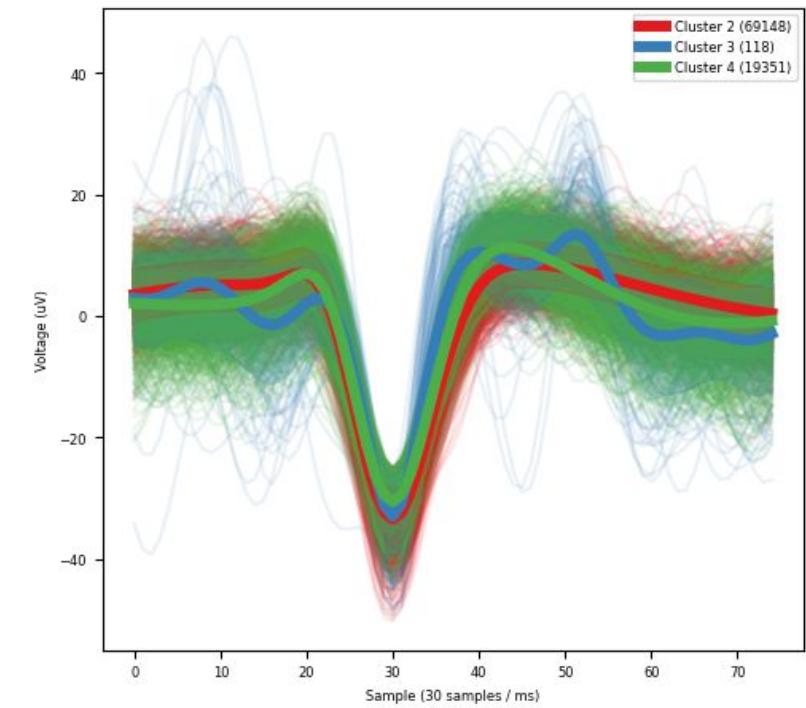
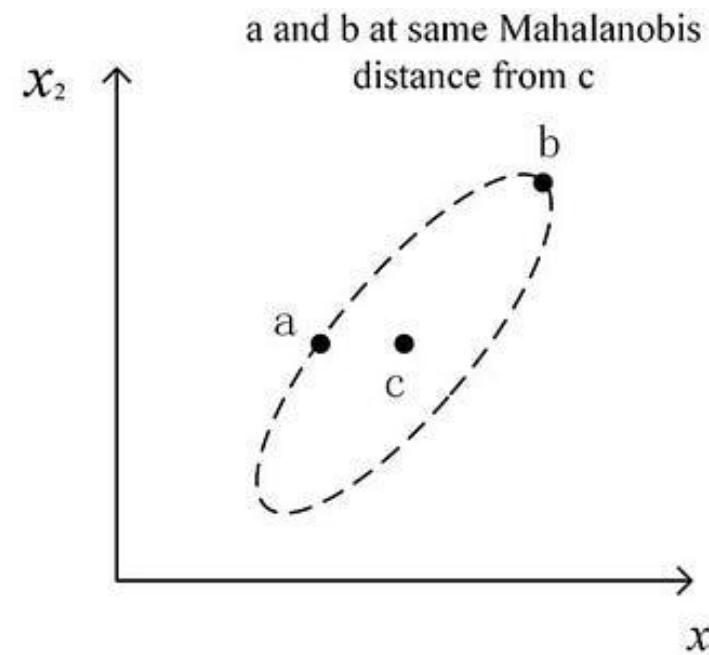
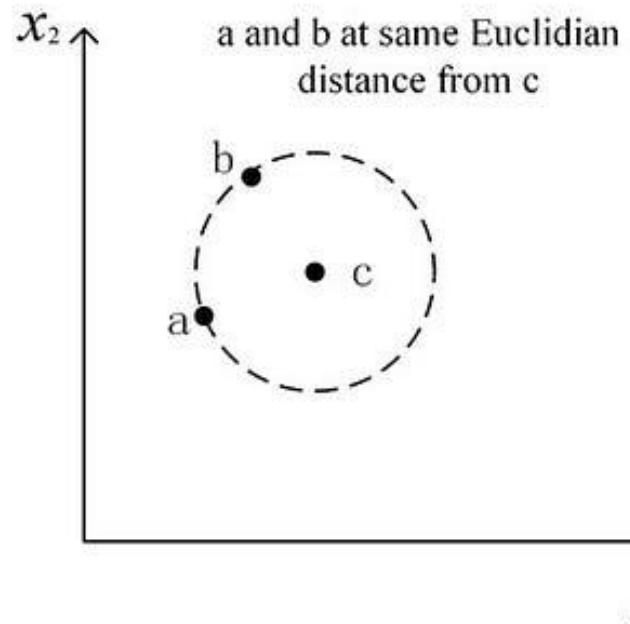
2. Automating clustering of waveforms

- BGM tends to over-split
- Have to figure out how to automatically merge clusters



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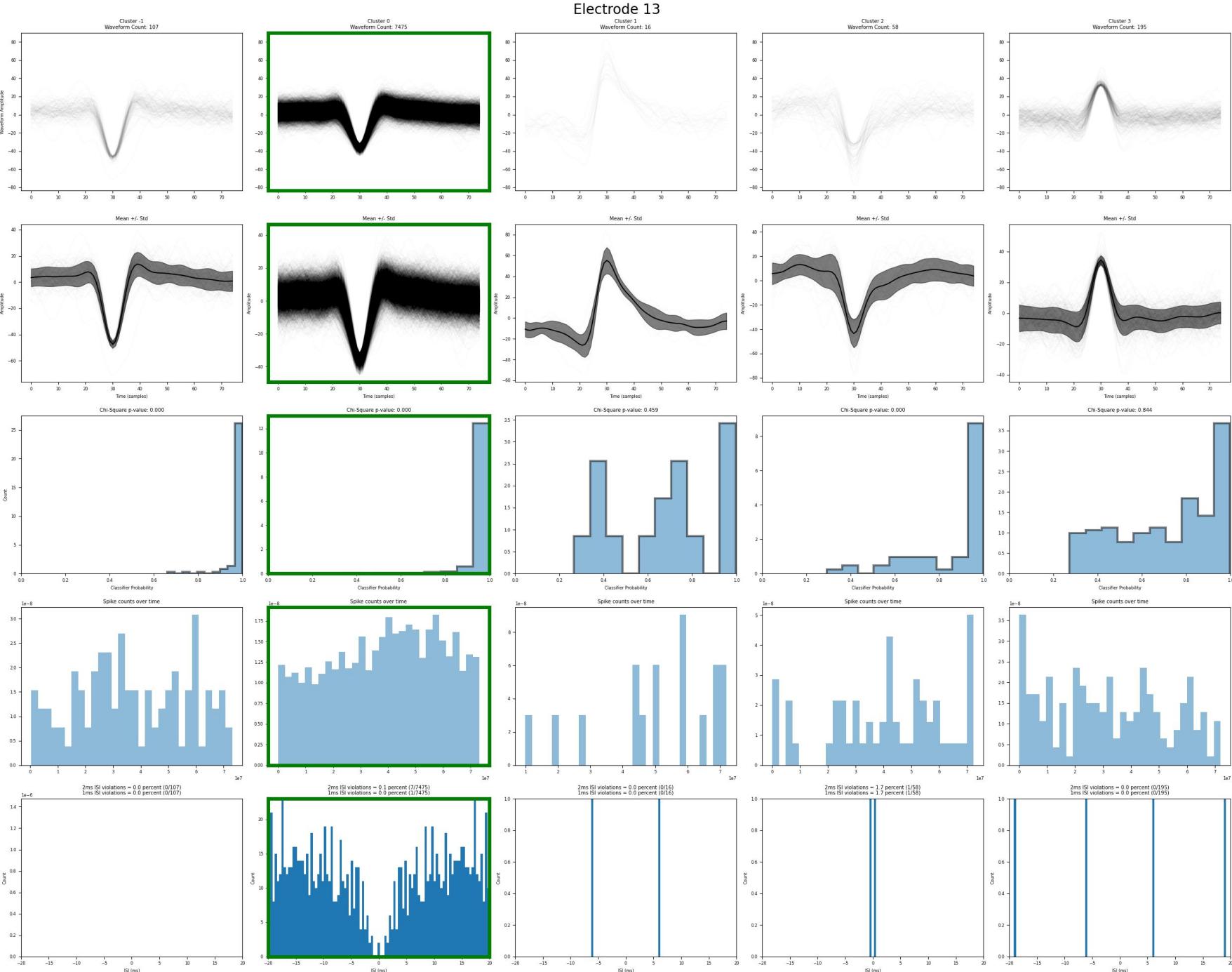
3. Selecting the right clusters

1. We know whether a single waveform is noise or spike
2. How do we select whether a cluster is noise or spiking

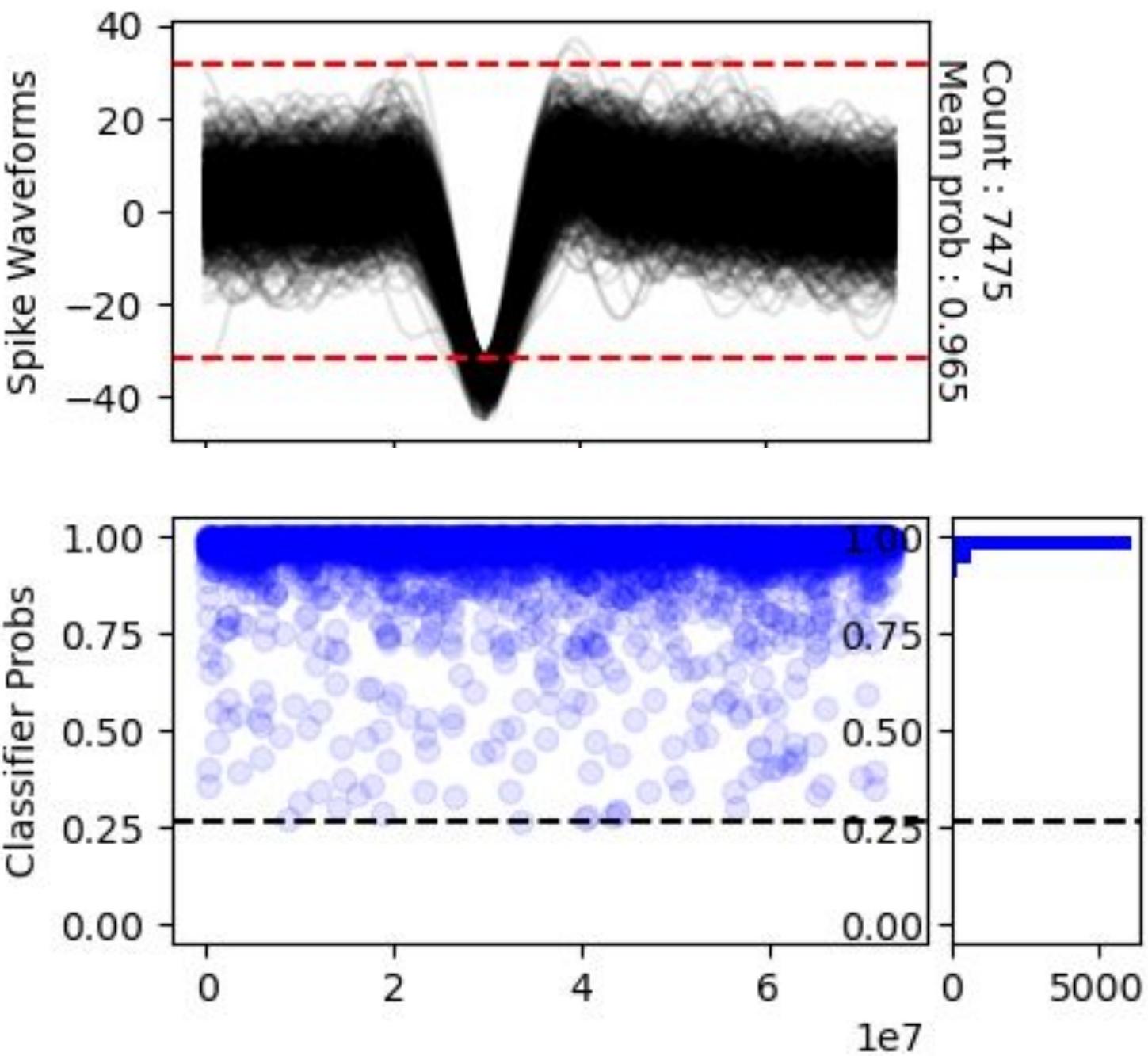
Criteria

1. Minimum number of waveforms (~2000)
2. Inter-spike interval violations
3. Distribution of classification probability (chi-square test for non-uniformity)

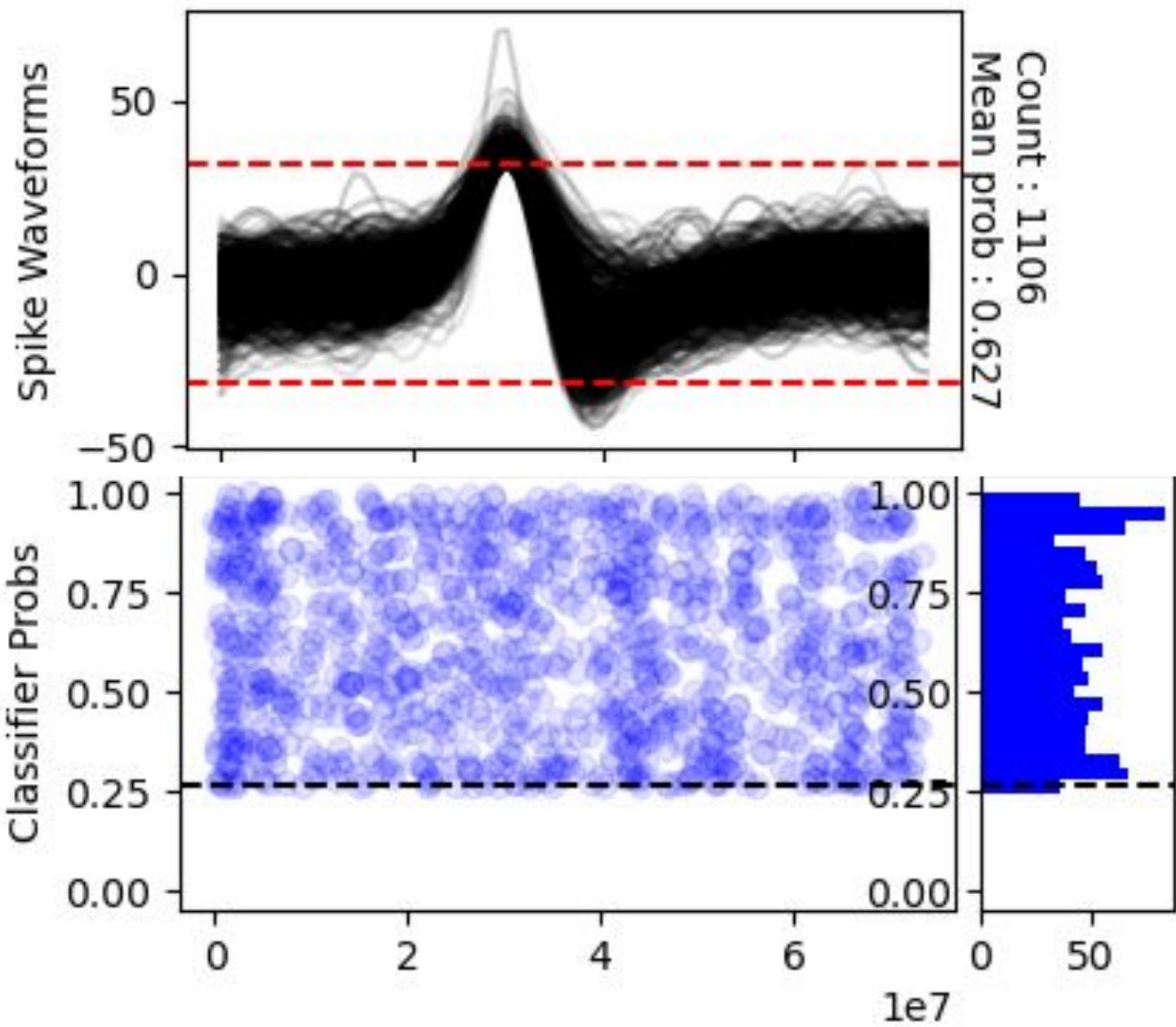
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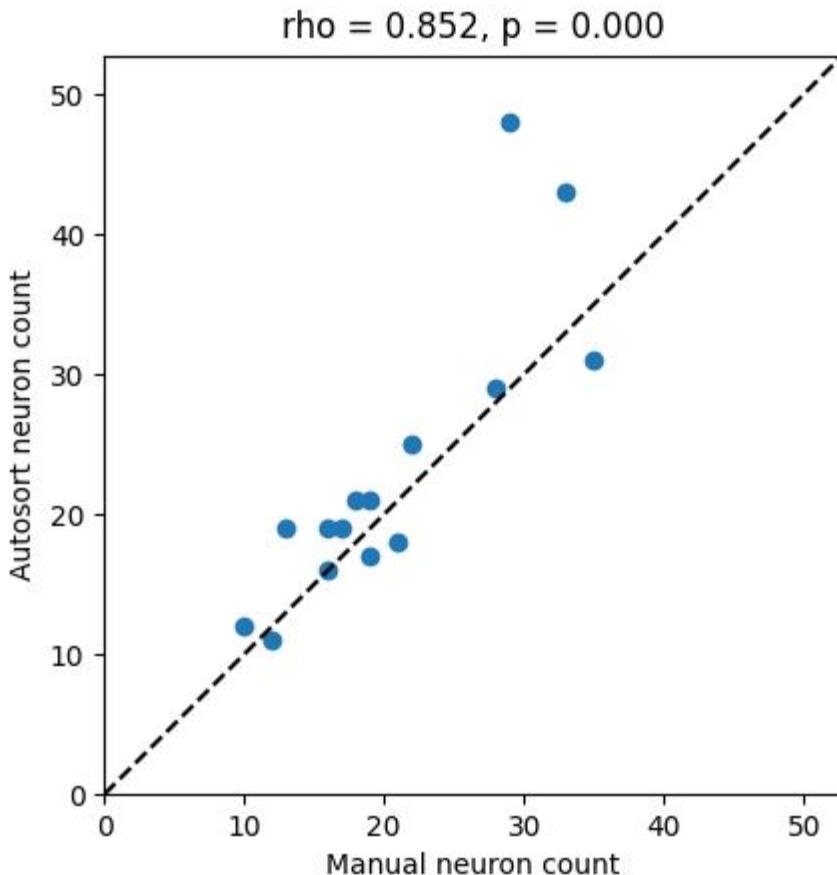


How well does it work?

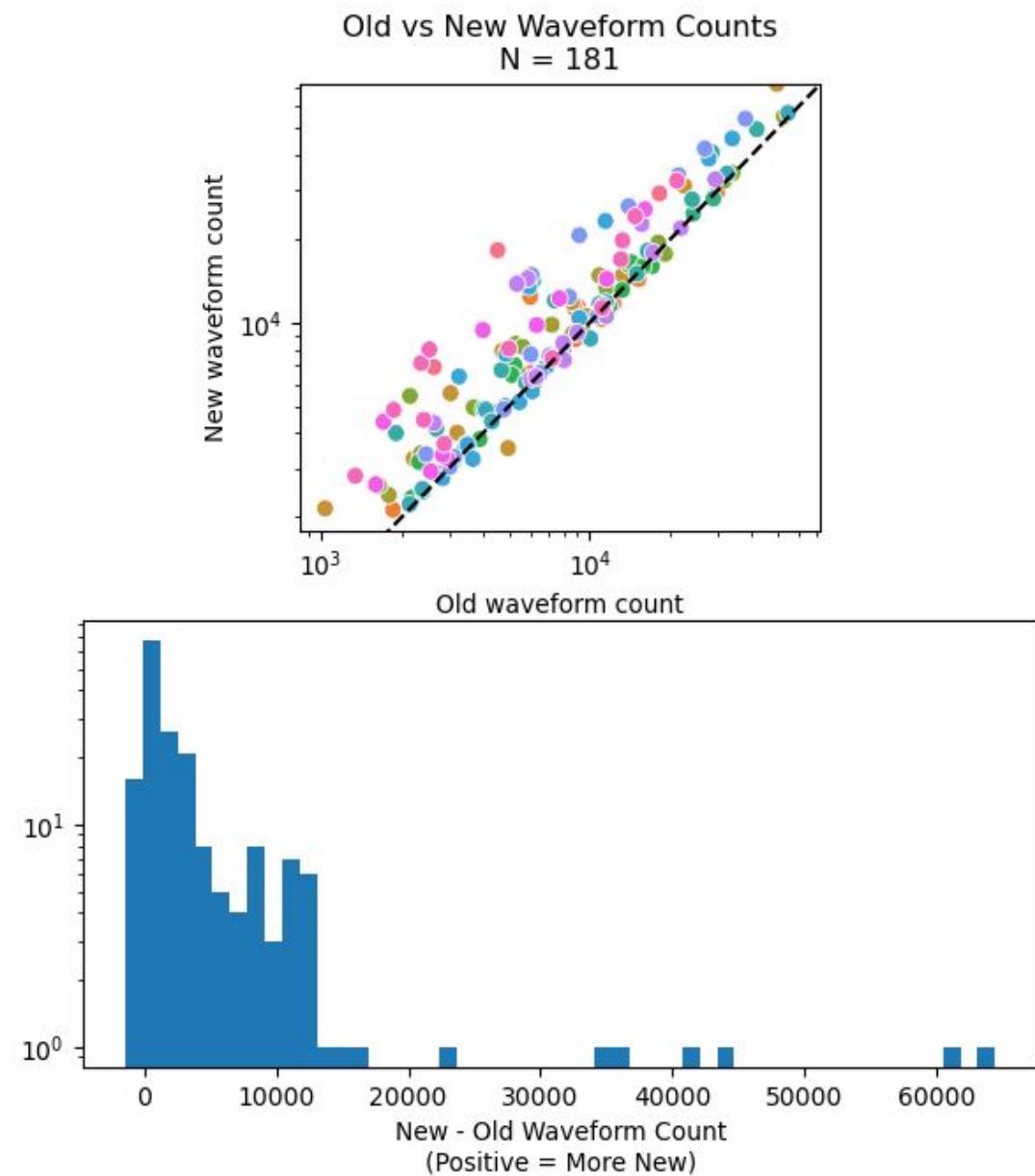
1. At the unit level
 - Comparing number of detected neurons
 - Comparing waveforms detected per unit
2. At the activity level
 - Single-trial activity
 - Mean-trial activity
 - Changes in similarity of activity with neuron counts

How well does it work?

Comparing selected neuron counts



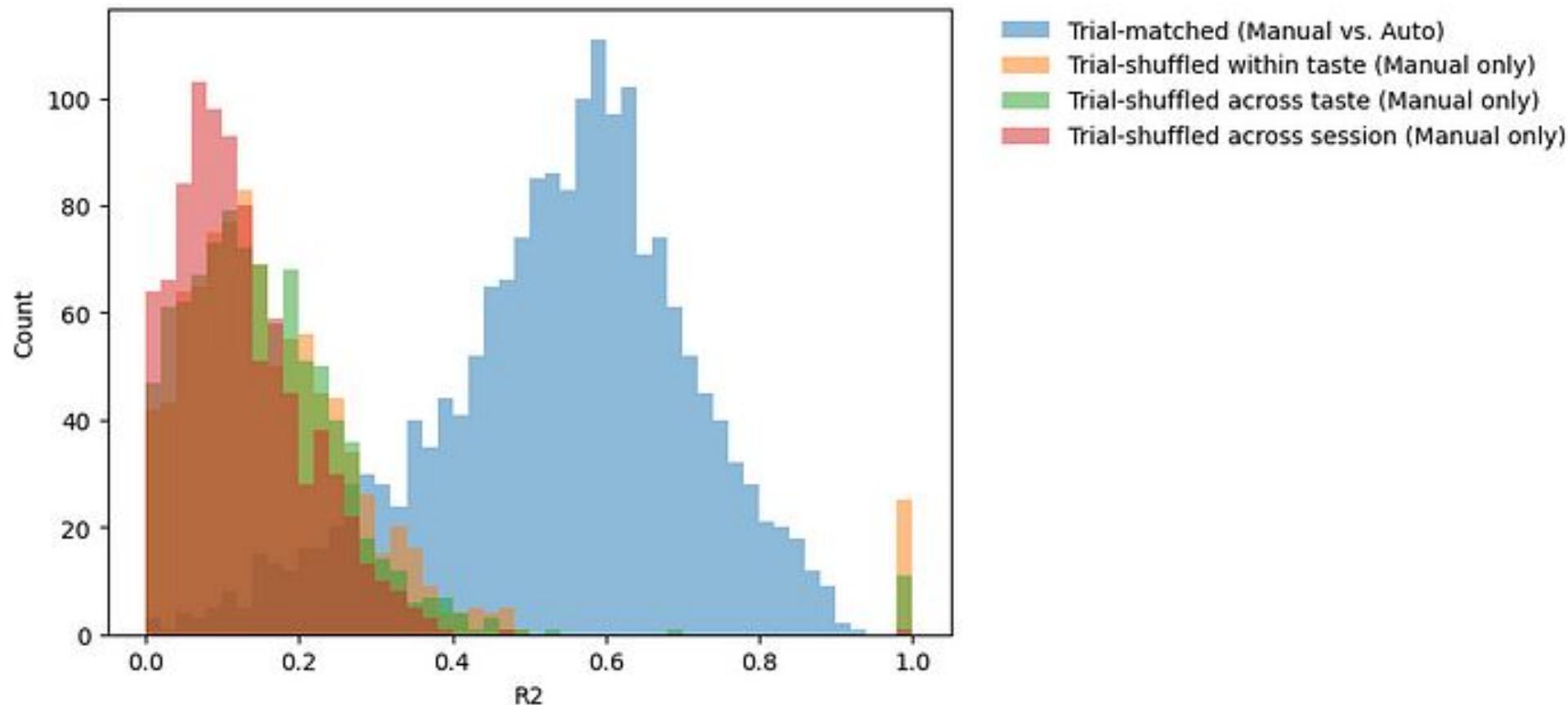
Comparing waveform counts



How well does it work?

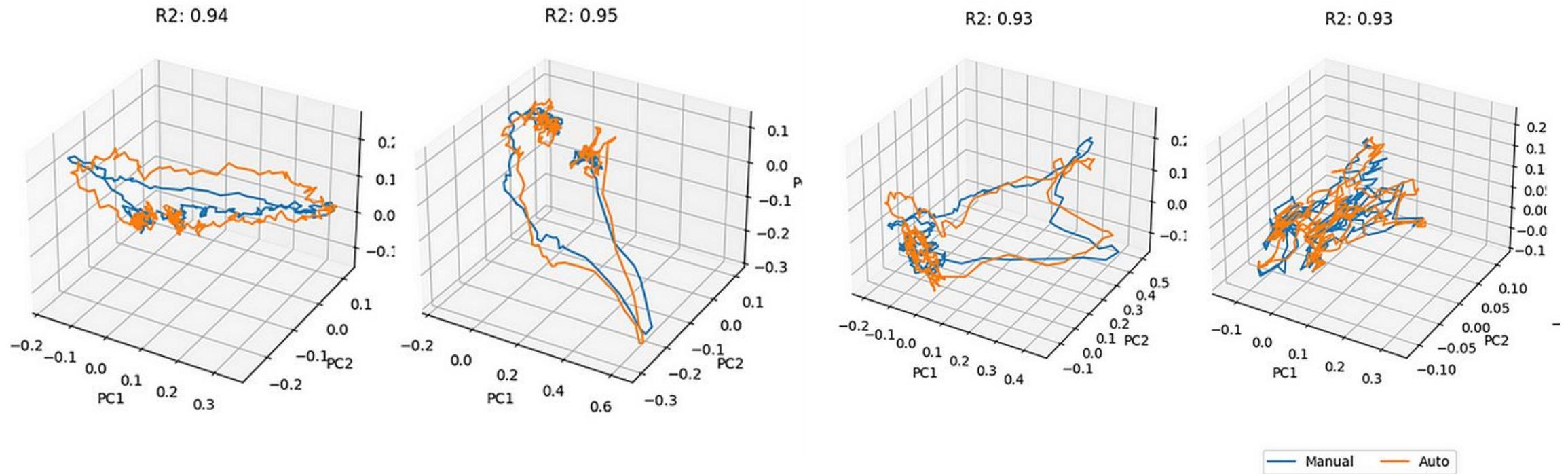
Comparison of single-trial firing rates (population PCA)

Single-Trial R2 value comparison



How well does it work?

Trial-averaged population PCA



How well does it work?

Changes in similarity of mean neural activity vs. Neuron Counts

