

Setting spike boundaries: The effectiveness of three neuronal spike validation methods

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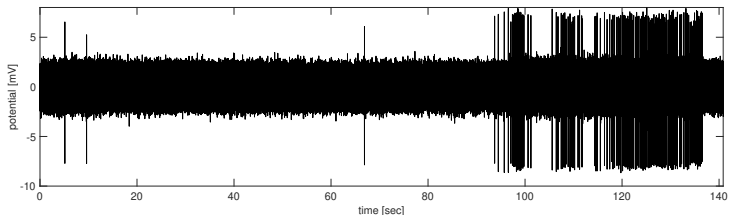
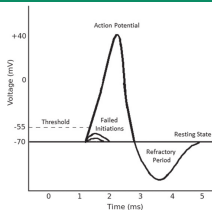
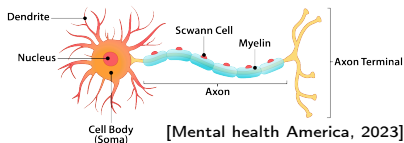
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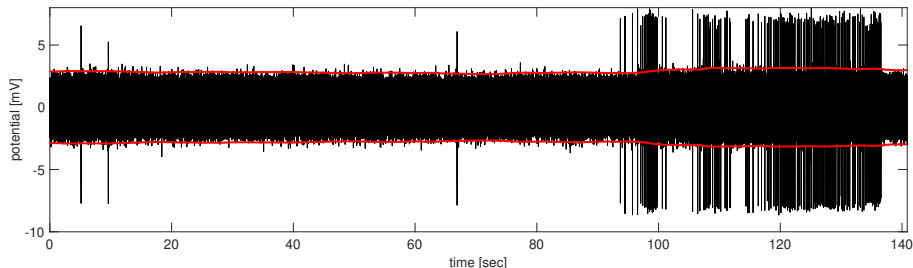
Neural spikes

NEURON ANATOMY



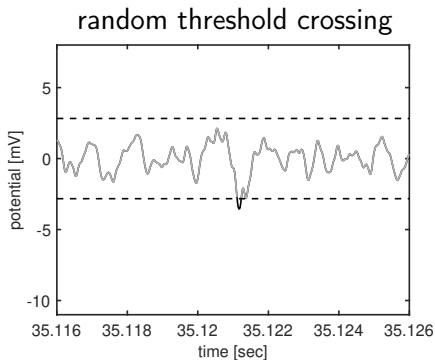
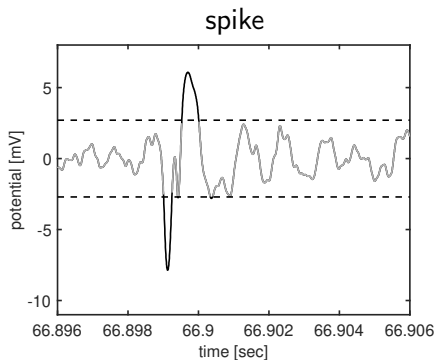
- neuron spiking activity → identification + treatment of brain diseases:
 - epilepsy, Parkinson's disease, ...
 - depression, psychiatric abnormalities, memory loss, ...

How to detect spiking activity of a neuron?



- **AdaBandFlt** [Biffi et al., 2010] → adaptive thresholding method
 - 1 = threshold crossing 0 = no threshold crossing
 - indicator vector (0,0,0,0,0,0,1,1,1,1,0,0,...)

Which threshold crossings form the same spike?



$$\text{spike} = x \pm K$$

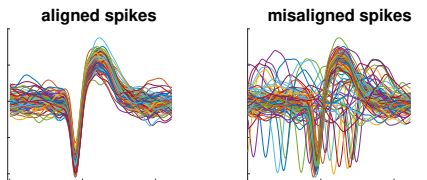
- x = “spike center”, different definitions among methods
- $K \approx 1$ ms
- refractory period \rightarrow distance between two spike centres $> \theta_{refrac}$

Spike validation methods

- method of Wagenaar [Wagenaar et al., 2005, Biffi et al., 2010]
- method of Toosi [Toosi et al., 2021]
- method of Nenadic [Nenadic and Burdick, 2005]

- comparison focused on:

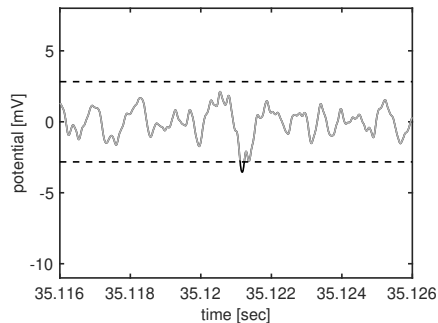
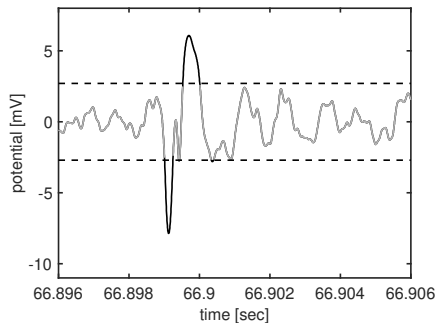
- 1 spikes alignment



- 2 ability to distinguish an actual spike from a random threshold crossing
- 3 sensitivity to presence of noise
- 4 distance between actual spike centre and detected spike centre (mD)

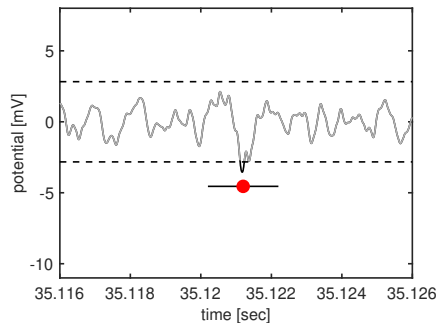
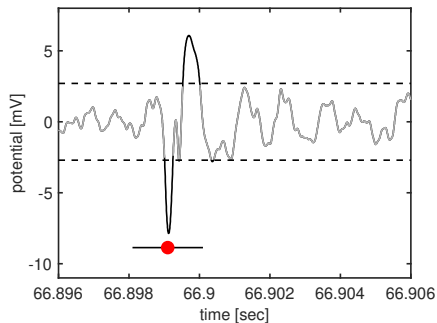
Spike validation: method of Wagenaar

- **input:** indicator vector + original electrophysiological signal
- a threshold crossing = spike center if:



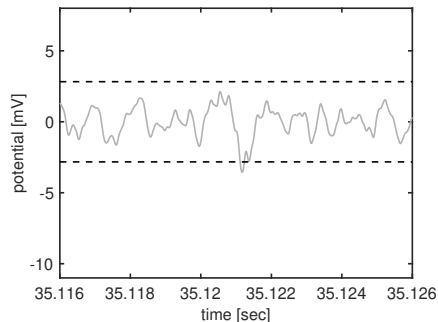
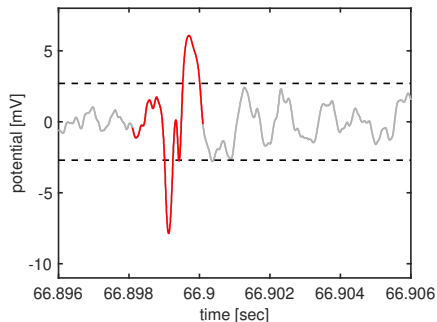
Spike validation: method of Wagenaar

- **input:** indicator vector + original electrophysiological signal
- a threshold crossing = spike center if:
 - C1: it forms the highest peak of either polarity over the ± 1 ms interval
 \Rightarrow **spikes are aligned to their dominant extremum**
 - C2: its amplitude $> 2 \times$ amplitude of the 2nd highest peak with the same polarity on this interval
 - C3: $|x_i - x_{i+1}| > \theta_{refrac}$



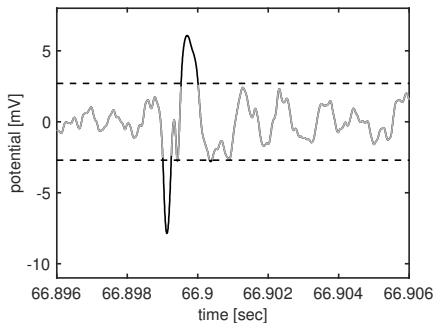
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Spike validation: method of Toosi

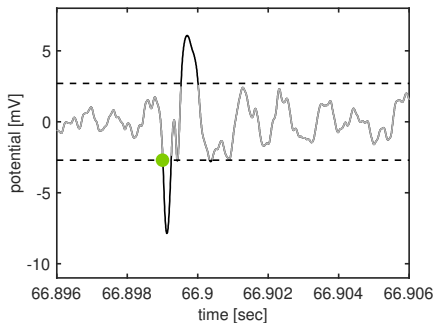
- [Toosi et al., 2021]



Spike validation: method of Toosi

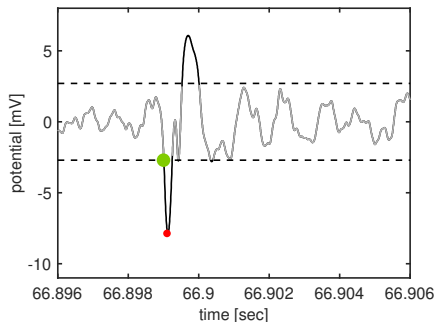
- [Toosi et al., 2021]

1. find the first threshold crossing ●



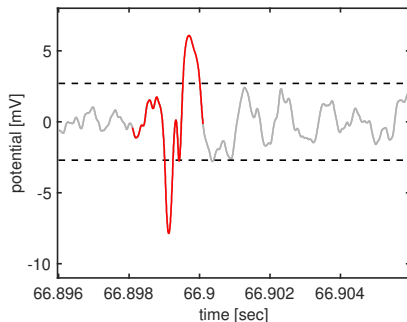
Spike validation: method of Toosi

- [Toosi et al., 2021]
1. find the first threshold crossing ●
 2. spike center x ● = local minimum to the right of ●



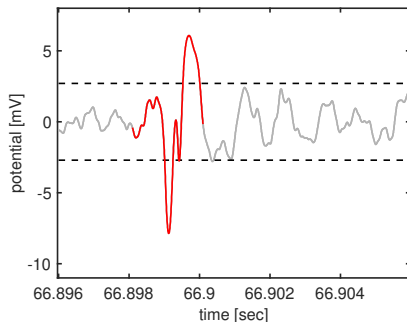
Spike validation: method of Toosi

- [Toosi et al., 2021]
1. find the first threshold crossing ●
 2. spike center x ● = local minimum to the right of ●
 3. spike: $x \pm K$ ms



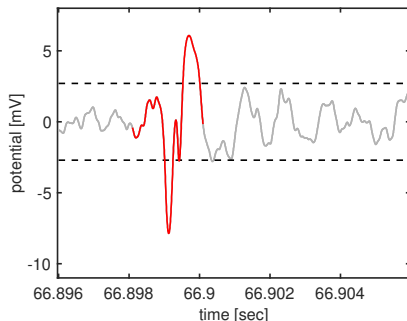
Spike validation: method of Toosi

- [Toosi et al., 2021]
1. find the first threshold crossing ●
 2. spike center x ● = local minimum to the right of ●
 3. spike: $x \pm K$ ms
 4. discard threshold crossings within the refractory period of the spike



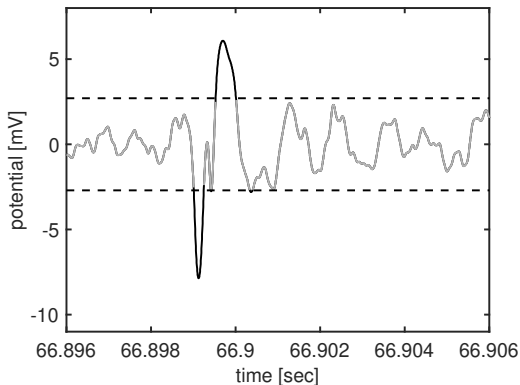
Spike validation: method of Toosi

- [Toosi et al., 2021]
- 1. find the first threshold crossing ●
- 2. spike center x ● = local minimum to the right of ●
- 3. spike: $x \pm K$ ms
- 4. discard threshold crossings within the refractory period of the spike
- repeat steps 1.-4. with the next (not discarded) threshold crossing
- **spikes are aligned to their global minimum**



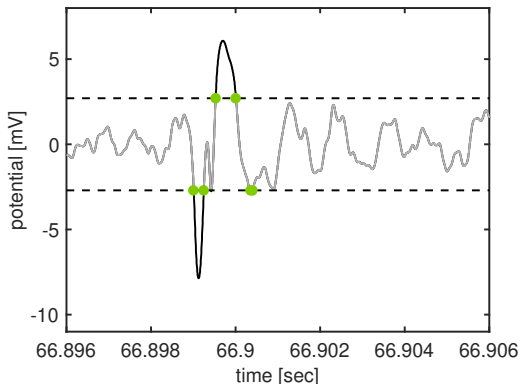
Spike validation: method of Nenadic

- [Nenadic and Burdick, 2005]



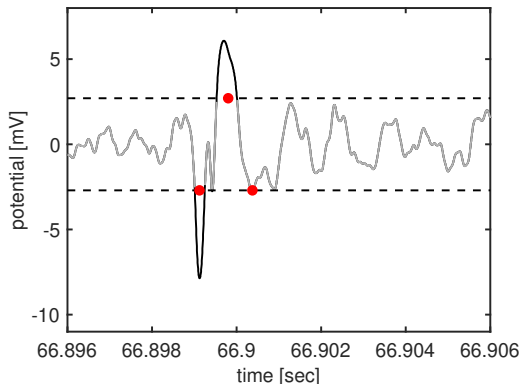
Spike validation: method of Nenadic

- [Nenadic and Burdick, 2005]
- indicator vector - 1 = threshold crossing, 0 = no threshold crossing
- $0 \rightarrow 1 \Rightarrow$ possible spike starting point
- $1 \rightarrow 0 \Rightarrow$ possible spike ending point



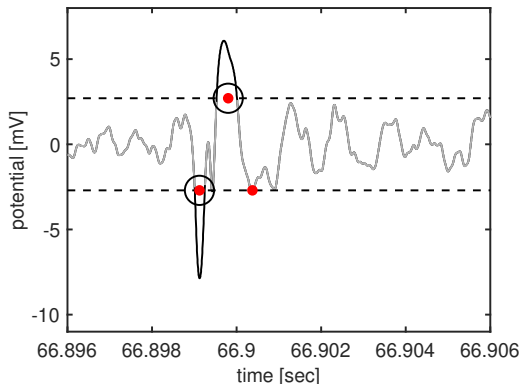
Spike validation: method of Nenadic

- [Nenadic and Burdick, 2005]
- candidate spike centre x
= mean between consecutive starting and ending points



Spike validation: method of Nenadic

- [Nenadic and Burdick, 2005]
- consecutive candidate spike centres x_i and x_{i+1} are analysed sequentially



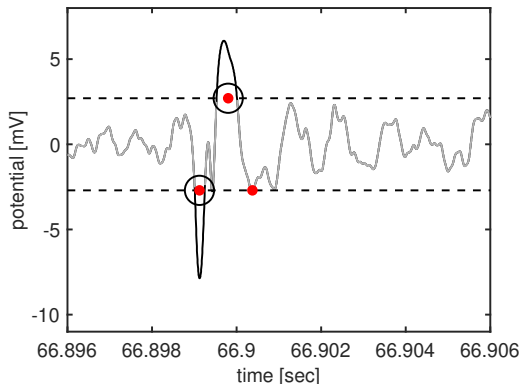
Spike validation: method of Nenadic

- [Nenadic and Burdick, 2005]

1. $|x_i - x_{i+1}| \leq \theta_{merge}$

⇒ x_i and x_{i+1} represent one spike

⇒ new candidate spike center $x_i^{new} = \lceil \frac{x_i + x_{i+1}}{2} \rceil$



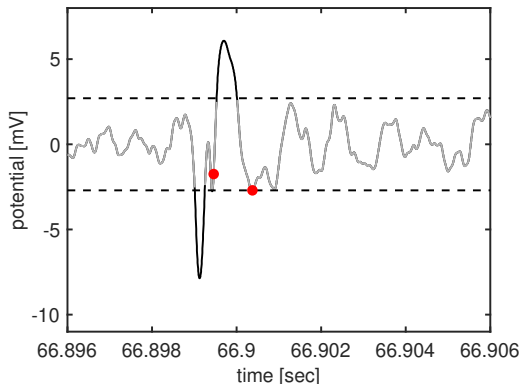
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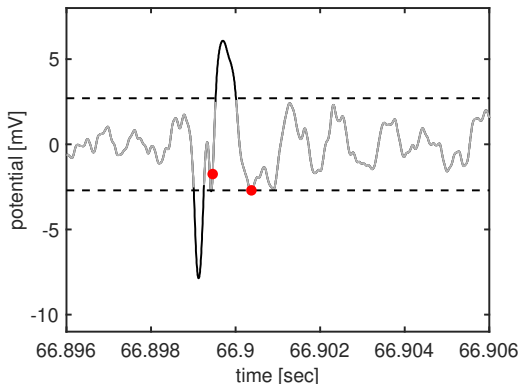


Spike validation: method of Nenadic

- [Nenadic and Burdick, 2005]

2. $|x_i - x_{i+1}| > \theta_{merge}$ and $|x_i - x_{i+1}| \leq \theta_{refrac}$

⇒ x_{i+1} is discarded

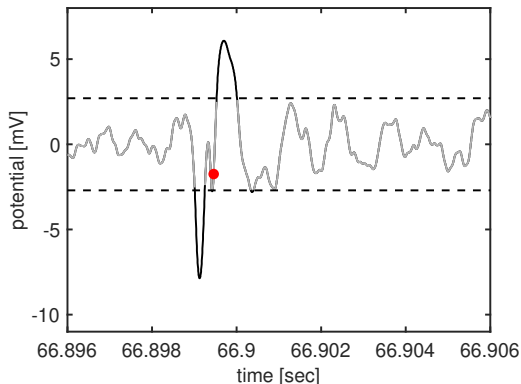


Spike validation: method of Nenadic

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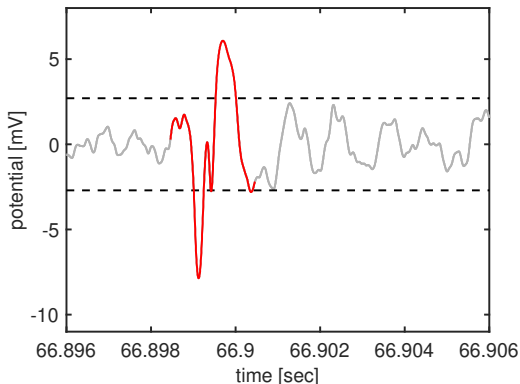


Spike validation: method of Nenadic

- [Nenadic and Burdick, 2005]

2. $|x_i - x_{i+1}| > \theta_{merge}$ and $|x_i - x_{i+1}| \leq \theta_{refrac}$

⇒ x_{i+1} is discarded



Spike validation: theoretical comparison

	Wagenaar	Toosi	Nenadic
spikes alignment	partially	yes	no
rejecting random threshold crossings	yes (C2)	no	no
presence of noise	?	?	?
$ X_{true} - X_{detected} $?	?	?

⇒ simulated data with different levels of signal-to-noise ratio (SNR)

Simulated data

- inspired by the work of [Smith and Mtetwa, 2007]
- duration: 5 seconds, sampling rate: 100 kHz
- **target neuron**
 - Naundorf model [Naundorf et al., 2006] , Poisson distribution of spike times
 - spike duration: 4,44 ms, refractory period: 10 ms
- **background electrophysiological activity**
 - 7 neurons - firing in line with the target neuron
 - 100 neurons - firing independently of target neuron
- **Gaussian noise** \rightarrow $SNR \in \{50, 35, 25, 10, 0, -5\}$ dB

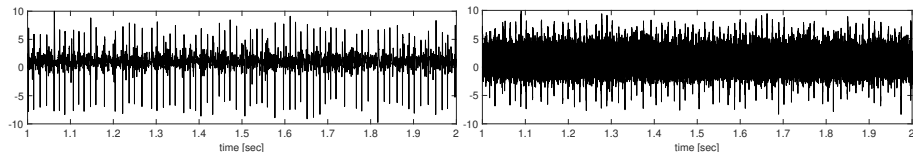
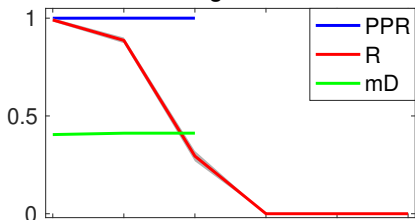


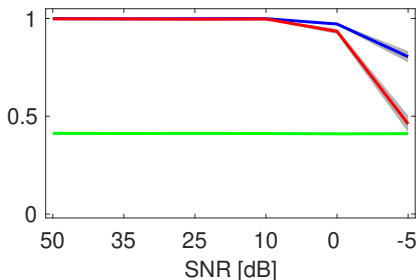
Figure: Simulated data with $SNR = 50$ (left) and $SNR = 0$ (right).

Results

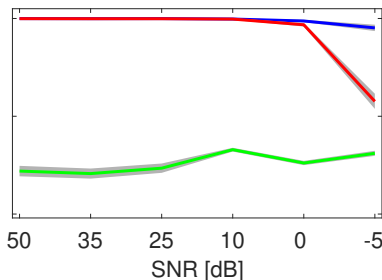
Wagenaar



Toosi

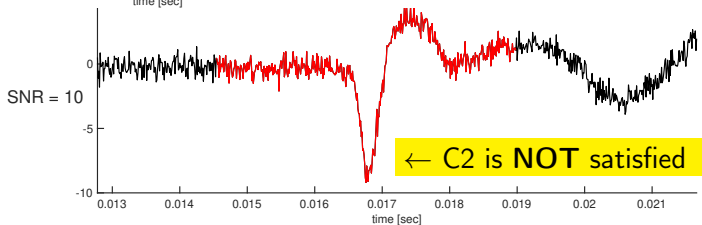
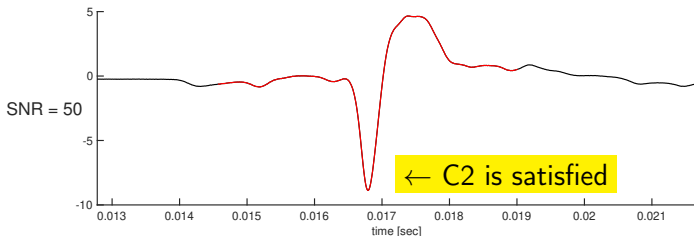


Nenadic



- spike = $x \pm 2, 22$ ms
- $mD = \|x_{detected} - x_{actual}\|$
- $PPR = \frac{\# \text{ correctly identified spikes}}{\# \text{ detected spikes}}$
- $R = \frac{\# \text{ correctly identified spikes}}{\# \text{ actual spikes}}$

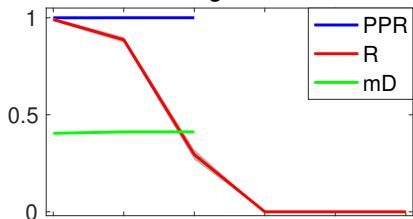
Results - problem in Wagenaar's method



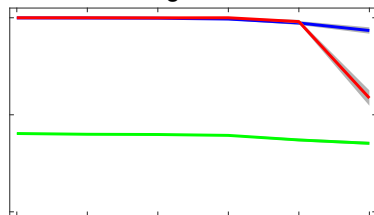
- **solution 1** → zero-phase bandpass filter between 300 Hz and 3000 Hz
- **solution 2** → removing condition C2 → Wagenaar2

Results

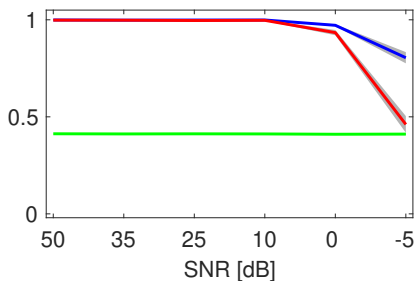
Wagenaar



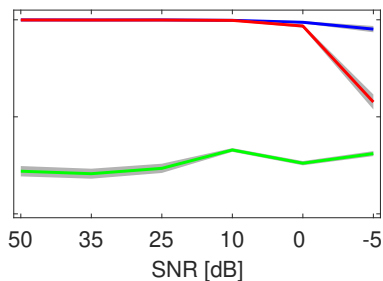
Wagenaar2



Toosi

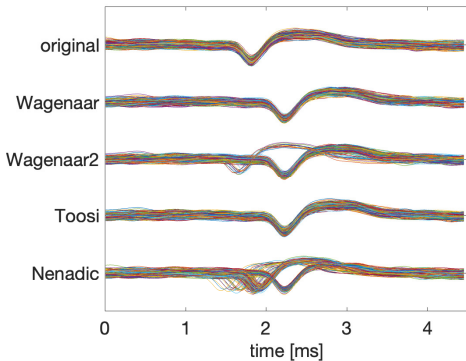


Nenadic

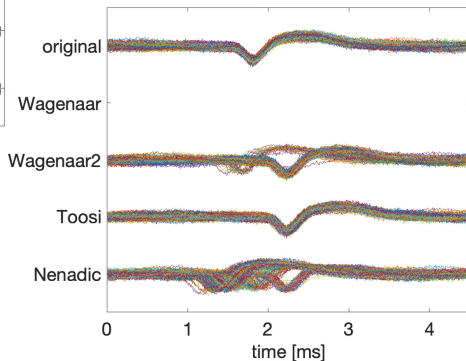


Results spikes alignment

SNR = 35 dB



SNR = 10 dB



Conclusion

	Wagenaar	Wagenaar2	Toosi	Nenadic
spikes alignment	partially	partially	yes	no
rejecting random threshold crossings	yes (C2)	no	no	no
presence of noise	problem	deteriorates alignment	ok	ok
$ x_{true} - x_{detected} $	\approx const.	\approx const.	const.	non-const.

- **future work** → methods' comparison on simulated data with:
 - spikes with varying amplitude
 - lower refractory period or overlapping spikes
 - ...

Literature I



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