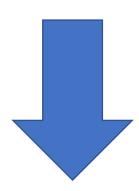


Choose a few files as sample to run normal CNMF-E individually. Each file will get some neurons that are common to all sample files as well as some unique neurons that file only in that file.



Generate a common spatial footprint of neuron (A): merge common ones and keep unique ones.



Use this A to extract neuron temporal C in all data you are interested in.

data folder