

# Dynamic Mode Decomposition for the Monodomain Sub-Model in the Neuro-Muscular System

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## ABSTRACT

The neuro-muscular system is a complex multiscale coupled system, for which detailed biophysics-based chemo-electro-mechanical models are available. Realistic simulations of such models are computationally extremely demanding, see, e.g., [1] and references therein. The 3D continuum-mechanical problem, which describes the deformation of the muscle, is coupled with a 1D problem, which describes the propagation of the action potential along muscle fibers. The propagation of action potentials can be described with the monodomain model, consisting of a partial differential equation (PDE) and associated ordinary differential equations (ODEs), representing the ionic currents across the muscle cell membranes. The ODEs derived from a Hodgkin-Huxley type [2, 3] modeling approach are stiff and thus require very small time steps.

To accelerate the simulations, we have recently applied a POD-DEIM model reduction technique to the above mentioned problem [4]. Considering a total reduction of the action potential, together with the ODE state variables, there is a high potential for reducing the coupled system. However, the system is solved for large physical time intervals with repeated stimulation of the muscle fibers. This motivates extrapolation of the solution of the monodomain equation in time instead of solving it in each activation period. For this purpose, the higher order dynamic mode decomposition (HODMD) [5, 6] seems to be an appropriate method, which uses time-lagged snapshots. A tunable parameter  $d \geq 1$  (number of related time-lagged snapshots) has to be adopted for accuracy. With  $d = 1$  one obtains the classical dynamic mode decomposition (DMD). To the best of our knowledge, the DMD/HODMD is not yet studied in context of neuro-muscular problems. We consider a 1D muscle fiber and stimulate it in the middle with different frequencies. The simulations are performed for different spatial and temporal resolutions. We use these full-order solutions and consider several values of  $d$  to find out an appropriate basis for the reduced solution. The first results show that the HODMD is more accurate than DMD in capturing the underlying physical modes of solution and basically applicable to neuro-muscular problems.

## REFERENCES

- [1] Bradley, C. P., Emamy, N., Ertl, T., Göddeke, D., Hessenthaler, A., Klotz, T., Krämer, A., Krone, M., Maier, B., Mehl, M., Rau, T. and Röhrle, O. Enabling detailed, biophysics-based skeletal muscle models on HPC systems. *Front. Physiol.* (2018) **9**.
- [2] Hodgkin, A. L. and Huxley, A. F. A quantitative description of membrane current and its application to conduction and excitation in nerve. *The Journal of Physiology* (1952) **117**(4):500–544.

- [3] Shorten, P. R., O'Callaghan, P., Davidson, J. B. and Soboleva, T. K. A mathematical model of fatigue in skeletal muscle force contraction. *Journal of Muscle Research and Cell Motility* (2007) **28(6)**:293–313.
- [4] N. Emamy, P. Litty, T. Klotz, M. Mehl and O. Röhrle. POD-DEIM Model Order Reduction for the Monodomain Reaction-Diffusion Sub-Model of the Neuro-Muscular System. (accepted) *MORCOS 2018 Proceedings*.
- [5] Le Clainche, S. and Vega, J.M. Higher Order Dynamic Mode Decomposition. *SIAM J. Appl. Dyn. Syst.* (2017) **16**:882–925.
- [6] Le Clainche, S. and Vega, J.M. Higher order dynamic mode decomposition to identify and extrapolate flow patterns. *Phys. Fluids* (2017) **29**:084102.