

LINKING BIOPHYSICAL MUSCLE MODELS WITH FINITE ELEMENT SOLVER

BOGDAN MILICEVIC^{1,2*}, MOMCILO PRODANOVIC^{1,3}, DANICA PRODANOVIC^{1,4}, MILJAN MILOSEVIC^{1,3,5}, BOBAN STOJANOVIC^{1,4}, SRBOLJUB M. MIJAILOVICH⁶, MILOS KOJIC^{1,7,8}, NENAD FILIPOVIC^{1,2}

¹Bioengineering Research and Development Center (BioIRC), Kragujevac, Serbia

²University of Kragujevac, Faculty of Engineering, Kragujevac, Serbia

³Institute for Information Technologies, University of Kragujevac, Kragujevac, Serbia

⁴University of Kragujevac, Faculty of Science, Kragujevac, Serbia

⁵Belgrade Metropolitan University, Belgrade, Serbia

⁶Illinois Institute of Technology, Department of Biology, Chicago, IL 60616, USA

⁷Serbian Academy of Sciences and Arts, Belgrade, Serbia

⁸Houston Methodist Research Institute, Houston, USA

e-mails: bogdan.milicevic@uni.kg.ac.rs; momcilo.prodanovic@kg.ac.rs;
danica.prodanovic@pmf.kg.ac.rs; miljan.m@kg.ac.rs;
boban.stojanovic@pmf.kg.ac.rs; smijailo@gmail.com; mkojic42@gmail.com;
fica@kg.ac.rs

Key words: muscle micromodel, biophysical models, multi-scale finite element analysis, software integration, drug effects.

ABSTRACT

Understanding the effects of structural and kinetics changes of sarcomeric proteins at the level on muscle cells or tissue requires detailed and comprehensive biophysical muscle models, such as MUSICO [1]. However, these models require large computational resources to track all molecular interactions. Thus, they are, in most cases, impractical for use in finite element solvers. To resolve this problem we have developed Mijailovich-Prodanovich (MP) surrogate model that is reduced to a system of ordinary differential equations and uses parameters that replicate transient responses obtained by detailed comprehensive models. This approach is computationally effective for use in finite element solvers and provides tools to translate changes across the length scales, i.e. from modulated molecular interactions to changes in functional behavior at the level of muscle cells and tissues. To demonstrate ability of this novel approach we have implemented the MP surrogate model into finite element solver PAK [2,3] to provide calculations of active tension and changes in muscle stiffness during (cardiac) muscle transient contractions. At a macro level, finite element solver PAK provides stretches along muscle fibers as a feedback loop to the MP model. Passive stresses, obtained from the recently implemented Holzapfel experimental material model [3], are also included in the calculation of total stress. Moreover, openMP library is incorporated in the multiscale model, which is typically used for loop-level parallelism, to speed up the calculations of integrated PAK-MP simulations. With the MP model integrated into the finite element solver PAK, we tracked the effects of several drugs, i.e., digoxin, disopyramide and mavacamten, on the cardiac muscle contraction of left ventricle. Simulations showed increase of ejection fraction, changes in outlet velocities and improvements in pressures and volumes. With pressure-volume diagrams, produced by our coupled simulations, we can help evaluate any realistic or

hypothetic clinical scenario, and potentially guide the treatment of patients with cardiac muscle diseases and disorders.

Acknowledgments: This work is supported by the SILICOFCM project that has received funding from the European Union’s Horizon 2020 research and innovation programme under grant agreement No 777204. The research was also funded by Serbian Ministry of Education, Science, and Technological Development, grants [451-03-9/2021-14/200378 (Institute for Information Technologies, University of Kragujevac)] and [451-03-9/2021-14/200107 (Faculty of Engineering, University of Kragujevac)].

REFERENCES

- [1] S. Mijailovich, O. Kayser-Herold, B. Stojanovic, Dj. Nedic, T. Irving, M. Geeves. Three-dimensional stochastic model of actin–myosin binding in the sarcomere lattice. *The Journal of General Physiology* 2016, 148(6): 459-488.
- [2] M. Kojić, R. Slavković, M. Živković and N. Grujović, “PAK - Finite Element Program for Linear and Nonlinear Analysis,” Kragujevac: University of Kragujevac and R&D Center for Bioengineering, 20
- [3] M. Kojic, M. Milosevic, B. Milicevic, and V. Simic, “Heart Mechanical Model Based on Holzapfel Experiments,” in *Learning and Analytics in Intelligent Systems*, Springer International Publishing, 2020, pp. 12–21. doi: 10.1007/978-3-030-43658-2_2.