## Performance comparison of FMU vs Matlab

Model by Smith [1], implemented in Modelica by [2], source code available at [3], model name Cardiovascular.Model.Smith2004.HemodynamicsSmith\_shallow has been simulated using several methods and compared to Matlab implementation by Brian Carlson.

- using Modelon FMIToolbox code
  - Export FMI2 from Dymola
    - Model exchange (ME) not containing solvers takes longest time. Not recommended.
    - Co-simulation (CS) using Dymola solvers performs probably the best
    - Co-simulation (CS) using CVODE solvers the Dymola 2019 suffers from a CVODE related bug. Should be fixed in FD01
  - Export from OpenModelica,
    - FMI1 ME provides rubbish output
    - FMI2 ME does not work
    - FMI2 CS crashes matlab (2018a)
- using Modelon FMIToolbox Simulink
  - Export FMI2 from Dymola
    - CS provides similar results as CS through code. The fastest solution!
  - Export from OpenModelica
    - Not being able to run simulation
- Load into Simulink's own FMIBlock
  - Export FMI2 from Dymola
    - Strange error:

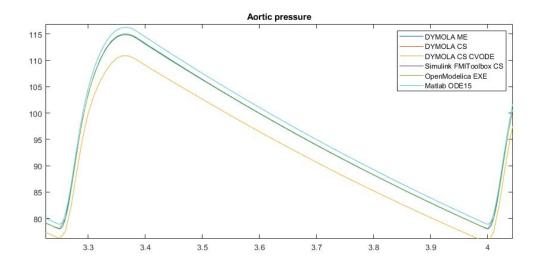
Value type mismatch for parameter 'aorta' in 'SmithCS/FMU'. Caused by:

Value type does not match the structure of variable 'aorta' defined in the modelDescription.xml file.

Component:Simulink | Category:Block error

- Export from OpenModelica
  - no combination of FMI export proved working
- OpenModelica exe simulator the performance is somewhat lower than Dymola CS FMIs, but still acceptable.
- Matlab reference implementation by Brian Carlson

The working solutions are presented in figures below. The tests were run at reference machine with i7-3610QM 2.3Ghz with 32GB RAM on windows 10, with matlab 2018a, Dymola 2019 and OpenModelica 1.12. The model has been run in Matlab for {1, 2, 3, 5, 8, 13, 21, 34, 55, 89} seconds and the the elapsed time has been measured. Each measurement has been repeated 10x. The measurement could be repeated using *FMUs.m.* All required sources are made public at



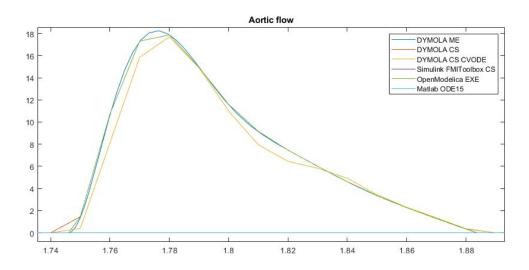


Figure 1: Comparison of output - The matlab solution provides slightly different, yet still acceptable result, probably caused by different numerics and /or unknown parametrization mismatch. All Modelica based results behave the same, except for Dymola CVODE, which suffers from a known bug in Dymola 2019 (waiting for a fix in the following version)

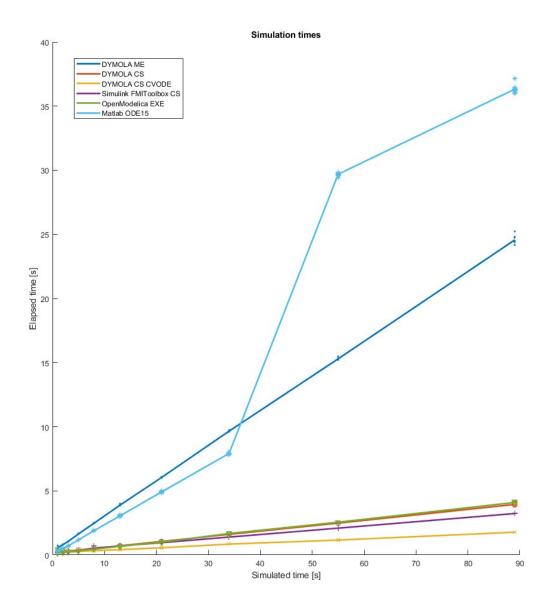


Figure 2: Simulation time required per simulated time. Each method and simulation time is run 10x and averaged. The vector-format figure *SimulationTimes.fig* might be downloaded from [4].

## References

- 1. Smith BW, Chase JG, Nokes RI, Shaw GM, Wake G. Minimal haemodynamic system model including ventricular interaction and valve dynamics. Med Eng Phys. Elsevier; 2004;26: 131–139. doi:10.1016/j.medengphy.2003.10.001
- 2. Ježek F, Kulhánek T, Kalecký K, Kofránek J. Lumped models of the cardiovascular system of various complexity. Biocybernetics and Biomedical Engineering. 2017;37:

666-678. doi:10.1016/j.bbe.2017.08.001

- 3. Ježek F. Physiolibrary.models. In: GitHub [Internet]. [cited 20 Jul 2017]. Available: https://github.com/filip-jezek/Physiolibrary.models
- 4. Ježek F. FMIComparison [Internet]. Github; Available: https://github.com/filip-jezek/FMIComparison