In [24]: import h5py import numpy as np import matplotlib.pyplot as plt import CLB.CLBXMLWriter as CLBXML import tempfile import scipy.optimize as so from display xml import XML from scipy.integrate import solve ivp First shot: spatially variable SIR model with uniform IC Zadanie E - Benchmark I: Poprawność symulacji równań typu SIR w domenie jednorodnej (OD) vs analityczne/Runge-Kutta - output jak do raportu/artykułu (nie do szkolenia) .md - dodać skrypt do examples and papers dodać ref do repo :) To evaluate, if model is working, let's start with basic case: uniform initial distribution of SIR variables, with periodic BC. This should give solution for original SIR ODE, if  $\beta$  is sufficiently large. Firstly, we prepare solution of ODE using SymPy's solve\_ivp toolbox. For details see previous workshops. In [102... #Test case settings I init = 0.1 $S_{init} = 1. - I init$ R init = 0beta = 1e20dt = 0.01R0 = 3T = 1000############################# # FD SOLUTION # def SIR OD(t, z, beta, gamma, N): # Susceptible → Infected → Removed :param t: time [days] :param z: Susceptible, Exposed, Infected, Removed :param beta: average number of contacts per day for each infected individual :param gamma: Between I and R, the transition rate is  $\gamma$ (simply the frequency of recoveries, that is, number of recovered or dead during one day divided by the total number of infected on that same day, supposing "day" is the time unit). If the duration of the infection is denoted D, then  $\gamma = 1/D$ . :return: derivatives [dS, dI, dR] S, I, R = zdSdt = -beta\*I\*S/NdIdt = beta\*I\*S/N - I\*gamma dRdt = I\*gammareturn [dSdt, dIdt, dRdt] # INITIAL CONDITIONS initial susceptible = S init # initial number of susceptible individuals in population. initial\_infections = I\_init # initial number of infected individuals in population. initial removed = R init # initial number of removed (recovered) individuals in population. IC = np.array([initial\_susceptible, initial\_infections, initial\_removed]) days to simulate = T\*dt sol = solve ivp(SIR OD, [0, days\_to\_simulate], IC, method='RK45', args=[R0, 1, 1],dense output=True) t = np.linspace(0, days to simulate, 1000)z = sol.sol(t)S rk4, I rk4, R rk4 = zparams = {'legend.fontsize': 'xx-large', 'figure.figsize': (14, 8), 'axes.labelsize': 'xx-large', 'axes.titlesize':'xx-large', 'xtick.labelsize':'xx-large', 'ytick.labelsize':'xx-large'} axes = plt.gca() plt.plot(t, S rk4, color="green", marker="", markevery=1, markersize=15, linestyle="-", linewidth=2, label='Susceptible') plt.plot(t, I rk4, color="red", marker="", markevery=1, markersize=15, linestyle="-", linewidth=2, label='Infected') plt.plot(t, R rk4, color="black", marker="", markevery=1, markersize=15, linestyle="-", linewidth=2, label='Removed') plt.xlabel('t') plt.ylabel('% of people') plt.title('SIR Epidemic Calculator') plt.legend() plt.grid() plt.show() SIR Epidemic Calculator 0.6 % of people Susceptible Infected Removed 0.4 0.2 0.0 We will use 2q9\_reaction\_diffusion\_system\_SIR\_ModifiedPeng model (or the WSIR model), aleready referenced in this workshop  $rac{\partial}{\partial t}W=eta_W\left\lceilrac{r^2}{8}W+(I-W)
ight
ceil$  $\frac{\partial}{\partial t}S = -\beta \frac{S}{N}W$  $\frac{\partial}{\partial t}I = \beta \frac{S}{N}W - \gamma I$  $\frac{\partial}{\partial t}R = \gamma I$ where  $S = ODE_1$  $I = ODE_2$  $R = ODE_3$  $C_1 = R_0$  $C_2 = \beta$  $C_3 = dt$  $S = ODE_1$ (1) $I = ODE_2$ (2) $R = ODE_3$ (3) $C_1 = R_0$  $C_2 = \beta$  $C_3 = dt$  $S = ODE_1$ In [103... CLBc = CLBXML.CLBConfigWriter() CLBc.addGeomParam('nx', 100) CLBc.addGeomParam('ny', 2) params = { "Diffusivity\_DRE\_1" : 1./6., "C 1":R0, "C 2":beta, "C 3":dt, "Init DRE 1":I init, #This is W equation, initially equal to I. large Beta reduces Relaxation-To-SIR "Init ODE 1":S init, "Init ODE 2":I init, "Init\_ODE\_3":R\_init CLBc.addModelParams(params) CLBc.addHDF5() solve = CLBc.addSolve(iterations=T) CLBc.addHDF5(Iterations=int(T / 50), parent=solve) CLBc.write('run.xml') In [107... ! rm -rf output/\* && tclb d2q9 reaction diffusion system SIR ModifiedPeng run.xml MPMD: TCLB: local:0/1 work:0/1 --- connected to: ] - CLB version: v6.0-beta-1713-g0861a50 Model: d2q9\_reaction\_diffusion\_system\_SIR\_ModifiedPeng [ ] Setting output path to: run ] Discarding 1 comments [ 0] Running on CPU [ 0] WARNING: No "Units" element in config file [ ] Mesh size in config file: 100x2x1 ] Global lattice size: 100x2x1 ] Max region size: 200. Mesh size 200. Overhead: 0% [ ] Local lattice size: 100x2x1 Hello allocator! Threads | [ ] Action | Primal , NoGlobals , BaseIteration 1x1| Tangent , NoGlobals , BaseIteration 1x1| Optimize , NoGlobals , BaseIteration 1x1| SteadyAdjoint , NoGlobals , BaseIteration 1x1| Primal , IntegrateGlobals , BaseIteration 1x1| Tangent , IntegrateGlobals , BaseIteration 1x1| Optimize , IntegrateGlobals , BaseIteration 1x11x1| SteadyAdjoint , IntegrateGlobals , BaseIteration | Primal , OnlyObjective , BaseIteration 1x1| Tangent , OnlyObjective , BaseIteration ] 1x1| Optimize , OnlyObjective , BaseIteration 1x1| SteadyAdjoint , OnlyObjective , BaseIteration 1x11x1| Primal , NoGlobals , BaseInit 1x1| Tangent , NoGlobals , BaseInit | Optimize , NoGlobals , BaseInit 1x1] 1x1| SteadyAdjoint , NoGlobals , BaseInit ] 1x1| Primal , IntegrateGlobals , BaseInit | Tangent , IntegrateGlobals , BaseInit ] 1x11x1| Optimize , IntegrateGlobals , BaseInit 1x1| SteadyAdjoint , IntegrateGlobals , BaseInit 1x1| Primal , OnlyObjective , BaseInit | Tangent , OnlyObjective , BaseInit [ ] 1x1| Optimize , OnlyObjective , BaseInit [ 1x11x1| SteadyAdjoint , OnlyObjective , BaseInit [0] Cumulative allocation of 71424 b (71.4 kB) ] Creating geom size:200 ] Setting output path to: run ] Setting output path to: output/run ] loading geometry ... ] Setting number of zones to 1 ] Setting Diffusivity\_DRE\_1 to 0.16666666666666 (0.166667) [0] Settings [Diffusivity for DRE\_1] to 0.166667 ] Setting C\_1 to 3.0000000000000 (3.000000) [0] Settings [Model parameter C\_1] to 3.000000 ] 000) ] Setting C\_3 to 0.01000000000000 (0.010000) [0] Settings [Model parameter C\_3] to 0.010000 Setting Init\_DRE\_1 in zone (-1) to 0.10000000000000 (0.100000) ] Setting Init\_ODE\_1 in zone (-1) to 0.90000000000000 (0.900000) Setting Init\_ODE\_2 in zone (-1) to 0.10000000000000 (0.100000) ] Setting Init\_ODE\_3 in zone (-1) to 0.00000000000000 (0.000000) ] Initializing Lattice ... Callback HDF5 with no Iterations attribute==] 0s ] Negotiated HDF5 chunks: 1x2x100[x3] [ 0 it writing hdf5 ] Setting action Solve at 1000.000000 iterations ] Setting callback HDF5 at 20.000000 iterations ] Negotiated HDF5 chunks: 1x2x100[x3] ] Adding HDF5 to the solver hands [ 0.28 GB/s [========] 1.3 MLBUps [ ] 20 it writing hdf5 [ ] [ ] 0.5 MLBUps 0.10 GB/s [==========] 40 it writing hdf5 [ ] 0.6 MLBUps 0.13 GB/s [==========] ] 60 it writing hdf5 ] 0.6 MLBUps ] ] 80 it writing hdf5 0.6 MLBUps 0.13 GB/s [=========] ] 100 it writing hdf5 ] 0.6 MLBUps ] 120 it writing hdf5 [ ] ] 0.6 MLBUps 0.13 GB/s [=========] 140 it writing hdf5 [ ] 0.13 GB/s [=========] ] 0.6 MLBUps 160 it writing hdf5 ] ] 0.6 MLBUps ] 180 it writing hdf5 ] 0.6 MLBUps 200 it writing hdf5 ] 0.13 GB/s [==========] ] 0.6 MLBUps 220 it writing hdf5 ] ] 0.6 MLBUps 0.13 GB/s [==========] 240 it writing hdf5 ] 0.13 GB/s [==========] ] 0.6 MLBUps 260 it writing hdf5 ] 0.6 MLBUps 0.13 GB/s [==========] ] ] 280 it writing hdf5 0.6 MLBUps 0.13 GB/s [==========] ] ] 300 it writing hdf5 0.13 GB/s [==========] 0.6 MLBUps 320 it writing hdf5 0.6 MLBUps 340 it writing hdf5 ] 0.13 GB/s [========= 0.6 MLBUps 360 it writing hdf5 0.6 MLBUps 380 it writing hdf5 [ 0.6 MLBUps [ ] 400 it writing hdf5 0.13 GB/s [=========] [ ] 0.6 MLBUps [ ] 420 it writing hdf5 0.14 GB/s [========] [ ] 0.7 MLBUps 440 it writing hdf5 [ ] 0.16 GB/s [=========] [ ] 0.8 MLBUps [ ] 460 it writing hdf5 0.16 GB/s [=========] [ ] 0.8 MLBUps [ ] 480 it writing hdf5 0.16 GB/s [=========] [ ] 0.8 MLBUps [ ] 500 it writing hdf5 0.16 GB/s [========] 0.8 MLBUps [ ] [ ] 520 it writing hdf5 0.16 GB/s [=========] [ ] 0.8 MLBUps [ ] 540 it writing hdf5 0.16 GB/s [==========] 0.8 MLBUps [ ] [ ] 560 it writing hdf5 0.16 GB/s [==========] 0.8 MLBUps [ ] [ ] 580 it writing hdf5 0.16 GB/s [=========] 0.8 MLBUps [ ] [ ] 600 it writing hdf5 0.16 GB/s [=========] [ ] 0.8 MLBUps [ ] 620 it writing hdf5 0.16 GB/s [=========] [ ] 0.8 MLBUps [ ] 640 it writing hdf5 0.16 GB/s [=========] [ ] 0.8 MLBUps [ ] 660 it writing hdf5 0.16 GB/s [=========] [ ] 0.8 MLBUps [ ] 680 it writing hdf5 0.16 GB/s [=========] [ ] 0.8 MLBUps [ ] 700 it writing hdf5 0.16 GB/s [=========] 0.8 MLBUps [ ] [ ] 720 it writing hdf5 0.16 GB/s [=========] [ ] 0.8 MLBUps 740 it writing hdf5 [ ] 0.16 GB/s [=========] 0.8 MLBUps [ ] [ ] 760 it writing hdf5 0.16 GB/s [=========] 0.8 MLBUps [ ] [ ] 780 it writing hdf5 0.16 GB/s [=========] 0.8 MLBUps [ ] [ ] 800 it writing hdf5 0.16 GB/s [=========] [ ] 0.8 MLBUps [ ] 820 it writing hdf5 0.16 GB/s [=========] [ ] 0.8 MLBUps [ ] 840 it writing hdf5 0.16 GB/s [=========] 0.8 MLBUps [ ] [ ] 860 it writing hdf5 0.16 GB/s [=========] 0.8 MLBUps [ ] [ ] 880 it writing hdf5 0.16 GB/s [=========] 0.8 MLBUps [ ] [ ] 900 it writing hdf5 0.16 GB/s [=========] 0.8 MLBUps [ ] [ ] 920 it writing hdf5 0.16 GB/s [=========] [ ] 0.8 MLBUps 940 it writing hdf5 [ ] 0.8 MLBUps 0.16 GB/s [==========] [ ] [ ] 960 it writing hdf5 0.8 MLBUps [ ] [ 980 it writing hdf5 0.16 GB/s [==========] 0.8 MLBUps 1000 it writing hdf5 ] Total duration: 0.338237 s = 0.005637 min = 0.000094 hIn [108... S = list()I = list()R = list()for i in range (0, T, int(T / 50)): f = h5py.File('output/run\_HDF5\_%08d.h5'%i) #plt.plot(f['DRE 1'][0,25,:]) S.append( f['ODE\_1'][0,0,0] I.append( f['ODE\_2'][0,0,0] ) R.append( f['ODE 3'][0,0,0]) t lb = np.linspace(0,T,len(S))\*dtplt.figure(figsize=(8,8)) plt.plot(t lb,S, 'gx', label='S - TCLB') plt.plot(t lb,I, 'rx', label='I - TCLB') plt.plot(t lb,R, 'kx', label='R - TCLB') plt.plot(t, S rk4, color="green", marker="", markevery=1, markersize=15, linestyle="-", linewidth=2, label='S - RK45') plt.plot(t, I rk4, color="red", marker="", markevery=1, markersize=15, linestyle="-", linewidth=2, label='I - RK45') plt.plot(t, R rk4, color="black", marker="", markevery=1, markersize=15, linestyle="-", linewidth=2, label='R - RK45') plt.legend() plt.grid(which='both') 0.8 0.6 × S - TCLB X I - TCLB × R - TCLB S - RK45 I - RK45 R - RK45 0.2 We could use TCLB shortcuts in a loop, to see effects of  $\beta$ In [109... plt.figure(figsize=(8,8)) for beta in np.logspace(-5,6,3): CLBc = CLBXML.CLBConfigWriter() CLBc.addGeomParam('nx', 5) CLBc.addGeomParam('ny', 5) params = { "Diffusivity DRE 1" : 1./6., "C 1":R0, "C 2": beta, "C 3":dt, "Init\_DRE\_1":I\_init, #This is W equation, initlally equall to I. large Beta reduces Relaxation-To-S "Init ODE 1":S init, "Init ODE 2":I init, "Init ODE 3":R init CLBc.addModelParams(params) CLBc.addHDF5() solve = CLBc.addSolve(iterations=T) CLBc.addHDF5(Iterations=int(T / 50), parent=solve) CLBc.write('run.xml') ! rm -rf output/\* && tclb d2q9\_reaction\_diffusion\_system\_SIR\_ModifiedPeng run.xml > /dev/null && echo "DON" S = list()I = list()R = list()for i in range (0,T,int(T / 50)): f = h5py.File('output/run HDF5 %08d.h5'%i) #plt.plot(f['DRE\_1'][0,25,:]) S.append( f['ODE 1'][0,0,0]) I.append( f['ODE 2'][0,0,0] ) R.append( f['ODE\_3'][0,0,0])  $t_{b} = np.linspace(0, T, len(S))*dt$  $plt.plot(t lb,S, 'g-', label=r'S - TCLB ($\beta=$)')$ plt.plot(t lb,I, '--', label=r'Infected - TCLB (\$\beta=%e\$)'%\_beta ) plt.plot(t lb,R, 'k-', label='R - TCLB') #plt.plot(t, S rk4, color="green", marker="", markevery=1, markersize=15, linestyle="-", linewidth=2, label='S - RK45') plt.plot(t, I rk4, color="red", marker="", markevery=1, markersize=15, linestyle="-", linewidth=2, label='Infected - RK45') #plt.plot(t, R rk4, color="black", marker="", markevery=1, markersize=15, linestyle="-", linewidth=2, label='R - RK45') plt.legend() plt.grid(which='both') plt.xlabel('Time') plt.ylabel('% of population') Hello allocator! DONE! Hello allocator! DONE! Hello allocator! Text(0, 0.5, '% of population') Out[109... 0.35 --- Infected - TCLB (β = 1.000000e - 05) --- Infected - TCLB (β = 3.162278e + 00) --- Infected - TCLB (β = 1.000000e + 06) 0.30 Infected - RK45 0.25 0.20 of population 0.15 0.10 0.05 0.00 Time SIR - Simple Laplace The same could be done with SIR\_SimpleLaplace model - for large eta results are identical In [110... CLBc = CLBXML.CLBConfigWriter() CLBc.addGeomParam('nx', 5) CLBc.addGeomParam('ny', 5) params = { "Diffusivity\_DRE\_1" : 1./6., "Diffusivity\_DRE\_2" : 1./6., "Diffusivity\_DRE\_3" : 1./6., "C 1":R0, "C 2":dt, "Init DRE 1":S init, "Init DRE 2":I init, "Init DRE 3":R init CLBc.addModelParams(params) CLBc.addHDF5() solve = CLBc.addSolve(iterations=T) CLBc.addHDF5(Iterations=int(T / 50), parent=solve) CLBc.write('run.xml') ! rm -rf output/\* && tclb d2q9\_reaction\_diffusion\_system\_SIR\_SimpleLaplace run.xml > /dev/null S = list()I = list()R = list()for i in range(0,T,int(T / 50)): f = h5py.File('output/run\_HDF5\_%08d.h5'%i) #plt.plot(f['DRE 1'][0,25,:]) S.append( f['DRE\_1'][0,0,0] ) I.append( f['DRE 2'][0,0,0] ) R.append( f['DRE\_3'][0,0,0])  $t_{b} = np.linspace(0, T, len(S))*dt$ plt.figure(figsize=(8,8)) plt.plot(t\_lb,S, 'gx', label='S - TCLB') plt.plot(t lb,I, 'rx', label='I - TCLB') plt.plot(t\_lb,R, 'kx', label='R - TCLB') plt.plot(t, S rk4, color="green", marker="", markevery=1, markersize=15, linestyle="-", linewidth=2, label='S - RK45') plt.plot(t, I rk4, color="red", marker="", markevery=1, markersize=15, linestyle="-", linewidth=2, label='I - RK45') plt.plot(t, R rk4, color="black", marker="", markevery=1, markersize=15, linestyle="-", linewidth=2, label='R - RK45') plt.legend() plt.grid(which='both') Hello allocator! 0.6 S - TCLB I - TCLB R - TCLB S - RK45 I - RK45 R - RK45 0.0 Simple spatially variable SIR model with non-uniform IC In [ ]: