

# Module 6: Dictyostelium cluster formation & slug movement

Author: Fabian Rost

## Aim

- this was planned as a one week project, students should develop a complex model
- teach them about biology: morphogenesis, cellular behavior, ...
- teach them about mathematics: CPM, PDE, coupling

## Description

- good description on what has been done can be found in the report by the students (see section Papers)
- it's been two days of teaching:
  - first explain CPM and let them play with that a little (e.g. cell sorting)
  - explain PDE system we are going to use
    - I had to explain about phase plane analysis quite a bit
- then on day 3 start to couple PDE & CPM
  - start with coupling in one direction only
    - e.g. first give a fixed gradient and let CPM cells perform chemotaxis
    - then formulate a PDE where a substance is only produced where CPM cells are
  - then couple in both directions (implement the Savill model)
    - perform clustering and slug movement experiment

## Paper

- Savill, N. J., & Hogeweg, P. (1997). Modelling morphogenesis: from single cells to crawling slugs. *J. Theor. Biol*, 184, 229-235. [link](#)
- Stan Marée (2002), From pattern formation to morphogenesis: multicellular coordination in *Dictyostelium discoideum* [link](#)
- Alberto Quintero, Mirko Myllykoski, Anna Igolkina, Alexandra Rohde O'Sullivan Freltoft, Nitya Dixit, Fabian Rost (2012), Morphogenesis and Dynamics of Multicellular Systems, [link](#)

## Morpheus models

h ClusterFormation.xml |h

```
<?xml version='1.0' encoding='UTF-8'?>
```

```
<MorpheusModel version="1">
  <Description>
    <Title>Cluster Formation</Title>
  </Description>
  <Space>
    <Lattice class="square">
      <Size symbol="size" value="200 200 0"/>
      <BoundaryConditions>
        <Condition boundary="x" type="constant"/>
        <Condition boundary="y" type="constant"/>
        <Condition boundary="-x" type="constant"/>
        <Condition boundary="-y" type="constant"/>
      </BoundaryConditions>
      <NodeLength value="0.37"/>
    </Lattice>
    <SpaceSymbol symbol="space"/>
  </Space>
  <Time>
    <StartTime value="0"/>
    <StopTime symbol="end" value="5000"/>
    <TimeSymbol symbol="time"/>
  </Time>
  <CellTypes>
    <CellType class="medium" name="medium">
      <Property symbol="b" value="0"/>
      <Property symbol="max_c" value="0"/>
      <Property symbol="phase" value="0"/>
    </CellType>
    <CellType class="biological" name="ameba">
      <VolumeConstraint>
        <Strength value="1.0"/>
        <Target value="20"/>
      </VolumeConstraint>
      <Chemotaxis>
        <Layer symbol-ref="c"/>
        <Strength symbol-ref="mu"/>
      </Chemotaxis>
      <Property symbol="b" value="0"/>
      <Property symbol="max_c" value="0"/>
      <PDEReporter>
        <Input symbol-ref="c"/>
        <Maximum symbol-ref="max_c"/>
      </PDEReporter>
      <Property symbol="phase" value="1"/>
      <Property symbol="phaseTime" value="0"/>
      <Equation symbol-ref="phaseTime">
        <Expression>if(phase == 2, phaseTime+MCStime,
0)</Expression>
      </Equation>
      <Equation symbol-ref="phase" name="Rule_2_3">
```

```

        <Expression>if(phase == 1 and max_c > 0.1, 2,
if(phase == 2 and phaseTime>phase2duration, 3,
if(phase == 3 and max_c < 0.05, 1,
phase
))) </Expression>
    </Equation>
    <Property symbol="phase2duration" value="2"/>
    <Function symbol="mu">
        <Expression>if(phase == 2, 300, 0)</Expression>
    </Function>
</CellType>
<CellType class="biological" name="autoAmeba">
    <VolumeConstraint>
        <Strength value="1.0"/>
        <Target value="20"/>
    </VolumeConstraint>
    <!--
<Disabled>
    <Chemotaxis>
        <Layer symbol-ref="c"/>
        <Strength symbol-ref="mu"/>
    </Chemotaxis>
</Disabled> -->
    <Property symbol="b" value="0.5"/>
    <Property symbol="max_c" value="0"/>
    <Property symbol="phase" value="0"/>
</CellType>
</CellTypes>
<CPM>
    <Interaction>
        <Contact type1="ameba" type2="ameba" value="4"/>
        <Contact type1="ameba" type2="medium" value="2"/>
        <Contact type1="ameba" type2="autoAmeba" value="4"/>
        <Contact type1="autoAmeba" type2="medium" value="3"/>
        <Contact type1="autoAmeba" type2="autoAmeba" value="-15"/>
    </Interaction>
    <MCSDuration symbol="MCStime" value="0.1"/>
    <MetropolisKinetics temperature="1.0" yield="0.1"
stepper="edgelist">
    <Neighborhood>
        <Order>2</Order>
    </Neighborhood>
</MetropolisKinetics>
</CPM>
<PDE>
    <Layer symbol="c">
        <Diffusion rate="1"/>
        <Initial>
            <InitPDEExpression>
                <Expression>0</Expression>
            </InitPDEExpression>

```

```
</Initial>
<BoundaryConditions>
  <Condition boundary="x" value="0"/>
  <Condition boundary="-x" value="0"/>
  <Condition boundary="y" value="0"/>
  <Condition boundary="-y" value="0"/>
</BoundaryConditions>
</Layer>
<Layer symbol="r">
  <Diffusion rate="0"/>
  <Initial>
    <InitPDEExpression>
      <Expression>0</Expression>
    </InitPDEExpression>
  </Initial>
  <BoundaryConditions>
    <Condition boundary="x" value="0"/>
    <Condition boundary="-x" value="0"/>
    <Condition boundary="y" value="0"/>
    <Condition boundary="-y" value="0"/>
  </BoundaryConditions>
</Layer>
<System solver="euler" time-step="0.1">
  <DiffEqn symbol-ref="c">
    <Expression>alpha*(-f -r)</Expression>
  </DiffEqn>
  <DiffEqn symbol-ref="r">
    <Expression>alpha*epsilon*(3.5*c-b-r)</Expression>
  </DiffEqn>
  <Function symbol="epsilon">
    <Expression>if(c < 0.0065, 0.5,
if(c < 0.841, 0.0589,
0.5))</Expression>
  </Function>
  <Function symbol="f">
    <Expression>if(c < 0.0065, 20*c,
if(c < 0.841, -3*c+0.15,
15*(c-1))</Expression>
  </Function>
  <Function symbol="alpha">
    <Expression>if(cell.id != 0, 1, 0)</Expression>
  </Function>
</System>
</PDE>
<CellPopulations>
  <Population size="1" type="autoAmeba">
    <InitRectangle cells="1" type="regular">
      <Dimensions size="10 10 0" center="100 100 0"/>
    </InitRectangle>
  </Population>
```

```

    <Population size="1" type="ameba">
      <InitRectangle cells="500" type="random">
        <Dimensions size="200 200 0" origin="0 0 0"/>
      </InitRectangle>
    </Population>
  </CellPopulations>
  <Analysis>
    <Gnuplotter interval="10">
      <Terminal opacity="0.7" name="png"/>
      <PDE symbol-ref="c" min="-0.5" max="1.0">
        <ColorMap>
          <Color value="-0.5" color="dark-violet"/>
          <Color value="0" color="white"/>
          <Color value="1.0" color="red"/>
        </ColorMap>
      </PDE>
      <Cells symbol-ref="phase" min="0" max="3"/>
      <Cells symbol-ref="cell.volume" min="15" max="25"/>
      <PDE symbol-ref="c" min="-0.5" max="1.0">
        <ColorMap>
          <Color value="-0.5" color="dark-violet"/>
          <Color value="0" color="white"/>
          <Color value="1.0" color="red"/>
        </ColorMap>
      </PDE>
    </Gnuplotter>
  </Analysis>
</MorpheusModel>

```

- [Movie: Cluster Formation & cAMP Spirals](#)
- for the movie the system size was 500×500 with 4000 cells randomly placed, everything else was like in the example above

h CrawlingSlug.xml |h

```

<?xml version='1.0' encoding='UTF-8'?>
<MorpheusModel version="1">
  <Description>
    <Title>test</Title>
  </Description>
  <Space>
    <Lattice class="square">
      <Size symbol="size" value="100 200 0"/>
      <BoundaryConditions>
        <Condition boundary="x" type="constant"/>
        <Condition boundary="y" type="constant"/>
        <Condition boundary="-x" type="constant"/>
        <Condition boundary="-y" type="constant"/>
      </BoundaryConditions>
    </Lattice>
  </Space>
</MorpheusModel>

```

```
        <NodeLength value="0.37"/>
    </Lattice>
    <SpaceSymbol symbol="space"/>
</Space>
<Time>
    <StartTime value="1.0"/>
    <StopTime symbol="end" value="5000"/>
    <TimeSymbol symbol="time"/>
</Time>
<CellTypes>
    <CellType class="medium" name="medium">
        <Property symbol="b" value="0"/>
        <Property symbol="max_c" value="0"/>
        <Property symbol="phase" value="0"/>
    </CellType>
    <CellType class="biological" name="ameba">
        <VolumeConstraint>
            <Strength value="1.0"/>
            <Target value="20"/>
        </VolumeConstraint>
        <Chemotaxis>
            <Layer symbol-ref="c"/>
            <Strength symbol-ref="mu"/>
        </Chemotaxis>
        <Property symbol="b" value="0"/>
        <Property symbol="max_c" value="0"/>
        <PDEReporter>
            <Input symbol-ref="c"/>
            <Maximum symbol-ref="max_c"/>
        </PDEReporter>
        <Property symbol="phase" value="1"/>
        <Property symbol="phaseTime" value="0"/>
        <Equation symbol-ref="phaseTime">
            <Expression>if(phase == 2, phaseTime+MCStime,
0)</Expression>
        </Equation>
        <Equation symbol-ref="phase" name="Rule_2_3">
            <Expression>if(phase == 1 and max_c > 0.1, 2,
if(phase == 2 and phaseTime>phase2duration, 3,
if(phase == 3 and max_c < 0.05, 1,
phase
))) </Expression>
        </Equation>
        <Property symbol="phase2duration" value="0.5"/>
        <Function symbol="mu">
            <Expression>if(phase == 2, 50, 0)</Expression>
        </Function>
    </CellType>
    <CellType class="biological" name="autoAmeba">
        <VolumeConstraint>
```

```

        <Strength value="1.0"/>
        <Target value="20"/>
    </VolumeConstraint>
    <!--
<Disabled>
  <Chemotaxis>
    <Layer symbol-ref="c"/>
    <Strength symbol-ref="mu"/>
  </Chemotaxis>
</Disabled> -->
  <Property symbol="b" value="0.5"/>
  <Property symbol="max_c" value="0"/>
  <Property symbol="phase" value="0"/>
</CellType>
</CellTypes>
<CPM>
  <Interaction>
    <Contact type1="ameba" type2="ameba" value="3"/>
    <Contact type1="ameba" type2="medium" value="2"/>
    <Contact type1="ameba" type2="autoAmeba" value="4"/>
    <Contact type1="autoAmeba" type2="medium" value="3"/>
    <Contact type1="autoAmeba" type2="autoAmeba" value="-2"/>
  </Interaction>
  <MCSDuration symbol="MCStime" value="0.01"/>
  <MetropolisKinetics temperature="1.0" yield="2" stepper="edgelist">
    <Neighborhood>
      <Order>2</Order>
    </Neighborhood>
  </MetropolisKinetics>
</CPM>
<PDE>
  <Layer symbol="c">
    <Diffusion rate="1"/>
    <Initial>
      <InitPDEExpression>
        <Expression>0</Expression>
      </InitPDEExpression>
    </Initial>
    <BoundaryConditions>
      <Condition boundary="x" value="0"/>
      <Condition boundary="-x" value="0"/>
      <Condition boundary="y" value="0"/>
      <Condition boundary="-y" value="0"/>
    </BoundaryConditions>
  </Layer>
  <Layer symbol="r">
    <Diffusion rate="0"/>
    <Initial>
      <InitPDEExpression>
        <Expression>0</Expression>
      </InitPDEExpression>
    </Initial>
  </Layer>
</PDE>

```

```
</Initial>
<BoundaryConditions>
  <Condition boundary="x" value="0"/>
  <Condition boundary="-x" value="0"/>
  <Condition boundary="y" value="0"/>
  <Condition boundary="-y" value="0"/>
</BoundaryConditions>
</Layer>
<!--
<Disabled>
  <Function symbol="b">
    <Expression>0</Expression>
  </Function>
</Disabled> -->
  <System solver="euler" time-step="0.1">
    <DiffEqn symbol-ref="c">
      <Expression>alpha*(-f -r)</Expression>
    </DiffEqn>
    <DiffEqn symbol-ref="r">
      <Expression>alpha*epsilon*(3.5*c-b-r)</Expression>
    </DiffEqn>
    <Function symbol="f">
      <Expression>if(c < 0.0065, 20*c,
if(c < 0.841, -3*c+0.15,
15*(c-1))</Expression>
    </Function>
    <Function symbol="alpha">
      <Expression>if(cell.id != 0, 1, 0)</Expression>
    </Function>
    <Function symbol="epsilon">
      <Expression>if(c < 0.0065, 0.5,
if(c < 0.841, 0.0589,
0.5))</Expression>
    </Function>
  </System>
<!--
<Disabled>
  <Equation symbol-ref="r">
    <Expression>if(cell.id != 0, r, 0)</Expression>
  </Equation>
</Disabled> -->
</PDE>
<CellPopulations>
  <Population size="2" type="autoAmeba">
    <InitRectangle cells="2" type="regular">
      <Dimensions size="5 5 0" center="50 50 0"/>
    </InitRectangle>
  </Population>
  <Population size="1" type="ameba">
    <InitRectangle cells="50" type="regular">
```



```

        <Dimensions size="35 50 0" center="50 25 0"/>
    </InitRectangle>
</Population>
<!--
<Disabled>
    <Population size="2" type="autoAmeba">
        <InitRectangle cells="2" type="regular">
            <Dimensions size="5 5 0" center="55 50 0"/>
        </InitRectangle>
    </Population>
</Disabled> -->
</CellPopulations>
<Analysis>
    <Gnuplotter interval="10">
        <Terminal opacity="0.7" name="png"/>
        <PDE symbol-ref="c" min="-0.5" max="1.0">
            <ColorMap>
                <Color value="-0.5" color="dark-violet"/>
                <Color value="0" color="white"/>
                <Color value="1.0" color="red"/>
            </ColorMap>
        </PDE>
        <Cells symbol-ref="phase" min="1" max="3">
            <ColorMap>
                <Color value="1" color="yellow"/>
                <Color value="2" color="green"/>
                <Color value="3" color="red"/>
            </ColorMap>
        </Cells>
    <!--
<Disabled>
    <Cells symbol-ref="cell.volume" min="15" max="25"/>
</Disabled> -->
    <PDE symbol-ref="c" min="-0.5" max="1.0">
        <ColorMap>
            <Color value="-0.5" color="dark-violet"/>
            <Color value="0" color="white"/>
            <Color value="1.0" color="red"/>
        </ColorMap>
    </PDE>
    <!--
<Disabled>
    <PDE symbol-ref="r" min="-0.5" max="3.0">
        <ColorMap>
            <Color value="-0.5" color="dark-violet"/>
            <Color value="0" color="white"/>
            <Color value="3.0" color="red"/>
        </ColorMap>
    </PDE>
</Disabled> -->
    </Gnuplotter>

```

```
<!--  
<Disabled>  
  <Logger interval="0.1">  
    <Filename name="log"/>  
    <Format string="c r"/>  
    <Input>  
      <PDE mapping="all" slice="1"/>  
    </Input>  
    <Plot interval="0.1" terminal="png">  
      <X-axis column="1"/>  
      <Y-axis columns="2 3"/>  
    </Plot>  
  </Logger>  
</Disabled> -->  
</Analysis>  
</MorpheusModel>
```

- [Movie: Crawling Slug](#)

From:  
<https://imc.zih.tu-dresden.de/wiki/morpheus/> - **Morpheus**

Permanent link:  
<https://imc.zih.tu-dresden.de/wiki/morpheus/doku.php?id=documentation:course:module6>

Last update: **10:06 17.11.2017**

