

continuumColony
15.3

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Chapter 1

Data Structure Index

1.1 Data Structures

Here are the data structures with brief descriptions:

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Chapter 2

Data Structure Documentation

2.1 blockDims Struct Reference

struct of block dimensions for CUDA kernels

Public Member Functions

- void [makeBlockDims](#) (int xl, int yl, int zl)
make up struct of block dimentions for cuda kernels

Data Fields

- dim3 **defaultNumBlocks**
- dim3 **fbaNumBlocks**
- dim3 **diffNumBlocks**
- dim3 **syncTCLwithCLNumBlocks**
- dim3 **chemReactNumBlocks**
- dim3 **expandCellsNumBlocks**
- int **BCblocksWall**
- int **BCblocksFloor**

2.1.1 Detailed Description

struct of block dimensions for CUDA kernels

2.1.2 Member Function Documentation

2.1.2.1 void **blockDims::makeBlockDims** (int *xl*, int *yl*, int *zl*) `[inline]`

make up struct of block dimentions for cuda kernels

Parameters

<i>x/</i>	lattice length in x dimensions
<i>y/</i>	lattice length in y dimensions
<i>z/</i>	lattice length in z dimensions

Returns

Void

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2.2 cellLattice Struct Reference

Cell lattice struct. This contains a lattice of the local volume fraction of a single cell type, as well as the FBA table the cell uses.

Public Member Functions

- void [readArbitraryVolumeFractionFilename](#) (float *volumeFraction, string arbitraryVolumeFractionFilename, int ly, int lz)
Initialize the starting geometry of a cell type using the an arbitrary volume fraction file.
- void [buildCellLattice](#) (fbaModel f, int x, int y, int z, float lambda, int start, string arbitraryVolumeFractionFilename, int startX, int startY, int startNumber, int agarHeight, float cellV)
Build a cell lattice struct.

Data Fields

- float * **volumeFraction**
- float * **cVolumeFraction**
- float * **tempVolumeFraction**
- float * **cTempVolumeFraction**
- float * **fbaOuts** [11]
- float * **cFbaOuts** [11]
- [fbaModel](#) **fba**
- int **starterFlag**

2.2.1 Detailed Description

Cell lattice struct. This contains a lattice of the local volume fraction of a single cell type, as well as the FBA table the cell uses.

2.2.2 Member Function Documentation

2.2.2.1 void cellLattice::buildCellLattice (fbaModel *f*, int *x*, int *y*, int *z*, float *lambda*, int *start*, string *arbitraryVolumeFractionFilename*, int *startX*, int *startY*, int *startNumber*, int *agarHeight*, float *cellV*) [inline]

Build a cell lattice struct.

Parameters

<i>f</i>	An FBA table struct
<i>x</i>	Length of the lattice in the X-direction
<i>y</i>	Length of the lattice in the Y-direction
<i>z</i>	Length of the lattice in the Z-direction
<i>lambda</i>	Lattice Spacing (in meters)
<i>start</i>	Flag (0 or 1) indicating if this cell type is present at the start of a simulation
<i>arbitrary-Volume-Fraction-Filename</i>	Name of the arbitrary volume fraction file (can be "" if none)
<i>startX</i>	Starting X location (in units of the lattice spacing) for the cell (if start = 1 and arbitraryVolumeFractionFilename = "")
<i>startY</i>	Starting Y location (in units of the lattice spacing) for the cell (if start = 1 and arbitraryVolumeFractionFilename = "")
<i>startNumber</i>	Number of cells initially in the simulation (if start = 1 and arbitraryVolumeFractionFilename = "")
<i>agarHeight</i>	Height of the agar (in units of the lattice spacing)
<i>cellV</i>	Average cell volume

Returns

Void

2.2.2.2 void cellLattice::readArbitraryVolumeFractionFilename (float * *volumeFraction*, string *arbitraryVolumeFractionFilename*, int *ly*, int *lz*) [inline]

Initialize the starting geometry of a cell type using the an arbitrary volume fraction file.

Parameters

<i>volume-Fraction</i>	Pointer to a cell's volume fraction lattice
<i>arbitrary-Volume-Fraction-Filename</i>	Name of the arbitrary volume fraction file
<i>ly</i>	Length of the lattice in the Y-direction
<i>lz</i>	Length of the lattice in the Z-direction

Returns

Void

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2.3 cellLattices Struct Reference

Cell lattices struct. This is a container struct for all the cell lattice structs in the simulation. It also stores to total volume fraction lattice.

Public Member Functions

- void [buildCellLattices](#) (float mvf, int x, int y, int z, float l, float t, int ah, float cellV, float cellM)
Build the [cellLattices](#) struct.
- void **add** ([cellLattice](#) cL)

Data Fields

- int * **typeFlags**
- int * **cTypeFlags**
- float * **totalVolumeFraction**
- float * **cTotalVolumeFraction**
- int **numberCells**
- [cellLattice](#) **lattices** [10]
- float **maxVolFraction**
- float **conversionFactorConcPerSecondToMmolPerGDwtPerHr**
- int **xl**
- int **yl**
- int **zl**
- float **lambda**
- float **fbaTau**
- int **agarHeight**
- int **N**
- int **isRelaxed**

2.3.1 Detailed Description

Cell lattices struct. This is a container struct for all the cell lattice structs in the simulation. It also stores to total volume fraction lattice.

2.3.2 Member Function Documentation

2.3.2.1 `void cellLattices::buildCellLattices (float mvf, int x, int y, int z, float l, float t, int ah, float cellV, float cellM) [inline]`

Build the `cellLattices` struct.

Parameters

<i>mvf</i>	Maximum volume fraction parameter
<i>x</i>	Length of the lattice in the X-direction
<i>y</i>	Length of the lattice in the Y-direction
<i>z</i>	Length of the lattice in the Z-direction
<i>l</i>	Lattice Spacing (in meters)
<i>t</i>	FBA timestep
<i>ah</i>	Agar height in units of the lattice spacing
<i>cellV</i>	Average cell volume
<i>cellM</i>	Average cell mass

Returns

Void

The documentation for this struct was generated from the following file:

- `continuumColony_15.3.cu`

2.4 chemCognate Struct Reference

Chem cognate struct.

Data Fields

- int `reactChemIdx`
- int `prodChemIdx`

2.4.1 Detailed Description

Chem cognate struct.

The documentation for this struct was generated from the following file:

- `continuumColony_15.3.cu`

2.5 chemLattice Struct Reference

Cell lattice struct. This contains a lattice of chem concentrations, a lattice of diffusion rates, and info about time steps, boundary conditions, and other such things.

Public Member Functions

- void [buildChemLattice](#) (float *dw*, float *da*, float *l*, float *t*, int *x*, int *y*, int *z*, int *agar*, float *initAg*, float *initAir*, int *isInt*, int *cicli*, int *isHin*, float *cellVol*, int *wallBcFlag*, int *floorBcFlag*)

Build a chem lattice struct.

Data Fields

- float * **lattice**
- float * **tempLattice**
- float * **cLattice**
- float * **cTempLattice**
- float * **Dlattice**
- float * **cDlattice**
- float **Dwater**
- float **Dagar**
- float **initialAgar**
- float **initialAir**
- float **tau**
- int **isIntracellular**
- int **cellIdxChemLivesIn**
- int **isHindered**
- int **refWallBcFlag**
- int **refFloorBcFlag**

2.5.1 Detailed Description

Cell lattice struct. This contains a lattice of chem concentrations, a lattice of diffusion rates, and info about time steps, boundary conditions, and other such things.

2.5.2 Member Function Documentation

- 2.5.2.1 void **chemLattice::buildChemLattice** (float *dw*, float *da*, float *l*, float *t*, int *x*, int *y*, int *z*, int *agar*, float *initAg*, float *initAir*, int *isInt*, int *cicli*, int *isHin*, float *cellVol*, int *wallBcFlag*, int *floorBcFlag*) `[inline]`

Build a chem lattice struct.

Parameters

<i>dw</i>	The diffusion rate in water
<i>da</i>	The diffusion rate in agar
<i>l</i>	The lattice spacing
<i>t</i>	The timestep
<i>x</i>	Number of lattice sites in the x direction
<i>y</i>	Number of lattice sites in the y direction
<i>z</i>	Number of lattice sites in the z direction
<i>agar</i>	The height (in lattice sites) of the agar
<i>initAg</i>	The initial concentration of the chemical in the agar
<i>initAir</i>	The initial concentration of the chemical in the air
<i>isInt</i>	Flag determining if the chemical species is intracellular (1) or extracellular (0)
<i>cicli</i>	Cell index the chemical lives is--if a chemical species is intracellular, this int denotes the index of the cell type where the chemical resides
<i>isHin</i>	Flag determining if a chemical experiences hindered diffusion around cells (1) or does not experience hindered diffusion (0)
<i>cellVol</i>	Volume of the cell (in meters ³)
<i>wallBcFlag</i>	Flag determining if a chemical will experience reflective boundary conditions at the 4 walls of the simulation box (1) or will experience fixed-concentration boundary conditions (0)
<i>floorBcFlag</i>	Flag determining if a chemical will experience reflective boundary conditions at the floor of the simulation box (1) or will experience fixed-concentration boundary conditions (0)

Returns

Void

The documentation for this struct was generated from the following file:

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2.6 chemLattices Struct Reference

Chem lattices struct. This is a container struct for all the chem lattice structs in the simulation.

Public Member Functions

- void [buildChemLattices](#) (int x, int y, int z, float l, float t, int ah)
Build the chem lattices struct.
- void [add](#) ([chemLattice](#) cl)
Add a chem lattice to this container struct of chem lattices.

Data Fields

- int **xl**
- int **yl**
- int **zl**
- float **lambda**
- float **fbaTau**
- int **agarHeight**
- int **N**
- int **numberChems**
- [chemLattice](#) **lattices** [10]

2.6.1 Detailed Description

Chem lattices struct. This is a container struct for all the chem lattice structs in the simulation.

2.6.2 Member Function Documentation

2.6.2.1 void chemLattices::add (chemLattice *cl*) [inline]

Add a chem lattice to this container struct of chem lattices.

Parameters

<i>cl</i>	chem lattice
-----------	--------------

Returns

Void

2.6.2.2 void chemLattices::buildChemLattices (int *x*, int *y*, int *z*, float *l*, float *t*, int *ah*) [inline]

Build the chem lattices struct.

Parameters

<i>x</i>	Number of lattice sites in the x direction
<i>y</i>	Number of lattice sites in the y direction
<i>z</i>	Number of lattice sites in the z direction
<i>l</i>	The lattice spacing in meters
<i>t</i>	The simulation timestep
<i>ah</i>	The agar height in lattice sites

Returns

Void

The documentation for this struct was generated from the following file:

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2.7 chemReaction Struct Reference

Chem uptake reaction struct.

Public Member Functions

- void [buildChemReaction](#) (float *m*, float *cat*, float *e*, int *type*, int *react*, int *prod*, float *cellV*)

Build an uptake reaction struct.

Data Fields

- float **km**
- float **kcat**
- float **E**
- int **cellType**
- int **reactant**
- int **product**

2.7.1 Detailed Description

Chem uptake reaction struct.

2.7.2 Member Function Documentation

2.7.2.1 void `chemReaction::buildChemReaction` (float *m*, float *cat*, float *e*, int *type*, int *react*, int *prod*, float *cellV*) `[inline]`

Build an uptake reaction struct.

Parameters

<i>m</i>	Michaelis constant for a michaelis-menten uptake reaction
<i>cat</i>	Kcat for a michaelis-menten uptake reaction
<i>e</i>	Number of uptake enzymes on an average cell
<i>type</i>	The index of the cell type that does this uptake reaction
<i>react</i>	The index of the chem type that is uptaken
<i>prod</i>	The index of the chem type that is produced (intercellular)
Generated on Thu Nov 18 2016 14:53 from the cell type	

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2.8 chemReactions Struct Reference

container struct for all the uptake reaction structs

Public Member Functions

- void [add](#) ([chemReaction](#) cR)
add an uptake reaction to the container struct

Data Fields

- int **numberReactions**
- [chemReaction](#) **reactions** [10]

2.8.1 Detailed Description

container struct for all the uptake reaction structs

2.8.2 Member Function Documentation

2.8.2.1 void chemReactions::add (chemReaction cR) [inline]

add an uptake reaction to the container struct

Parameters

<i>cR</i>	The uptake reaction struct to be added
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Returns

Void

The documentation for this struct was generated from the following file:

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2.9 fbaModel Struct Reference

fbaModel struct. This is a struct that contains an FBA table and the parameters needed to index into it.

Public Member Functions

- void **buildFbaModel** (string inputFilename)
build the FBA table using an FBA table file.

Data Fields

- int **nInputs**
- int **nOutputs**
- int **inputIndexes** [10]
- int **outputIndexes** [11]
- int **numberOfTabledValues** [10]
- float **inputStride** [10]
- int **indexMultipliers** [10]
- float * **fbaTable**
- float * **cFbaTable**

2.9.1 Detailed Description

fbaModel struct. This is a struct that contains an FBA table and the parameters needed to index into it.

2.9.2 Member Function Documentation

2.9.2.1 void **fbaModel::buildFbaModel** (string *inputFilename*) [inline]

build the FBA table using an FBA table file.

Parameters

<i>filename</i>	The name of the FBA table file
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Returns

Void

The documentation for this struct was generated from the following file:

- continuumColony_15.3.cu

2.10 regReaction Struct Reference

Regulation reaction struct.

Public Member Functions

- void [buildRegReaction](#) (int chem1, int chem2, int reactCell, int prodCell, string mults)
Build a regulation reaction.
- void [fillChemCognate](#) (string cogs)
Regulation reactions turn one cell type into another; the intracellular species within the reactant cell type also need to transform into the intracellular species of the product cell type.

Data Fields

- float **multipliers** [10]
- [chemCognate](#) **cognates** [10]
- int **numChemCognates**
- int **chemType1**
- int **chemType2**
- int **reactantCell**
- int **productCell**

2.10.1 Detailed Description

Regulation reaction struct.

2.10.2 Member Function Documentation

2.10.2.1 void `regReaction::buildRegReaction` (int *chem1*, int *chem2*, int *reactCell*, int *prodCell*, string *mults*) `[inline]`

Build a regulation reaction.

Parameters

<i>chem1</i>	Index of the first chemical species a cell is assumed to change regulatory state in response to
<i>chem2</i>	Index of the second chemical species a cell is assumed to change regulatory state in response to
<i>reactCell</i>	Index of the cell type transformed by this regulation reaction
<i>prodCell</i>	Index of the cell type produced by this regulation reaction
<i>mults</i>	Coefficients for the polynomial rate law that governs this reaction

Returns

Void

2.10.2.2 void regReaction::fillChemCognate (string *cogs*) [inline]

Regulation reactions turn one cell type into another; the intracellular species within the reactant cell type also need to transform into the intracellular species of the product cell type.

Parameters

<i>cogs</i>	Tab delimited set of indexes of the intracellular chem index in the reactant cell (first) and product cell (second)
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Returns

Void

The documentation for this struct was generated from the following file:

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2.11 regReactions Struct Reference

container struct for the regulation reactions

Public Member Functions

- void [add](#) ([regReaction](#) reg)
add a regulation reaction to this container struct

Data Fields

- int **numberReactions**
- [regReaction](#) **reactions** [10]

2.11.1 Detailed Description

container struct for the regulation reactions

2.11.2 Member Function Documentation

2.11.2.1 void regReactions::add (regReaction *reg*) [inline]

add a regulation reaction to this container struct

Parameters

<i>reg</i>	The regulation reaction struct to be added
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Returns

Void

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