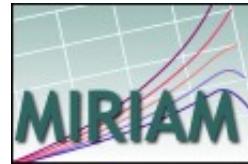
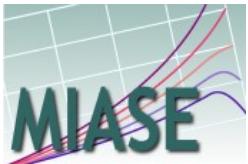
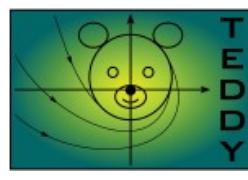


KiSAO

Kinetic Simulation Algorithm Ontology

	Models	Simulation	Results
Minimal requirements			
Data-models	 		SBRML
Ontologies			

Simulation approach

```
<listOfSimulations>
  <uniformTimeCourse id="simulation1"
    initialTime="0"
    outputStartTime="0"
    outputEndTime="140"
    numberOfPoints="1000">
    <algorithm kisaoID="KiSAO:0000030"/>
  </uniformTimeCourse>
</listOfSimulations>
```

Simulation approach

```
<listOfSimulations>
  <uniformTimeCourse id="simulation1"
    initialTime="0"
    outputStartTime="0"
    outputEndTime="140"
    numberOfPoints="1000">
    <algorithm kisaoID="KisAO:0000030"/>
  </uniformTimeCourse>
</listOfSimulations>
```

The screenshot shows a software interface for managing simulation configurations and their associated KISAO entries.

Left Panel (Simulation Configuration):

- ID:** KISAO:0000071
- Namespace:** KISAO
- Term name:** Livermore solver for ordinary differential equations
- Text:** LSODE solves explicitly given ODE systems. [and] [...] is based on the GEAR and GEARB packages. It solves ODE systems given explicitly as $dy/dt = f(t, y)$.
Hindmarsh AC. LSODE and LSODI, two new initial value ordinary differential equation solvers. SIGNUM Newsletter, Volume 15 (4), pages 10–11 (1980).
Radhakrishnan K, Hindmarsh AC. Description and Use of LSODE, the Livermore Solver for Ordinary Differential Equations. Lawrence Livermore National Laboratory Report, Vol. UCRL-ID-113855 (1993).
- Dbxrefs:** dk:16NOV2007, doi:1218052.12180

Bottom Buttons: Commit, Add, Del

DAG Viewer (Right Panel):

- Classes:**
 - kinetic simulation algorithm
 - is_a algorithm using non-spatial description
 - is_a Livermore solver
 - is_a Livermore solver for ordinary differential equations
 - is_a algorithm using continuous variables
 - is_a Livermore solver
 - is_a Livermore solver for ordinary differential equations
 - is_a algorithm using deterministic rules
 - is_a Livermore solver
 - is_a Livermore solver for ordinary differential equations
 - is_a algorithm using adaptive timesteps
 - is_a Livermore solver
 - is_a Livermore solver for ordinary differential equations

NCBO BioPortal: Kinetic Simulation Algorithm Ontology - kinetic simulation algorithm - Mozilla Firefox

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Kinetic Simulation Algorithm Ontology

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Legend 

kinetic simulation algorithm

- algorithm using adaptive timesteps
 - Bortz-Kalos-Liebowitz method
 - code value ordinary differential equation
- Gillespie-like stochastic simulation method
 - + Gillespie-like approximate simulation
 - + Gillespie-like exact stochastic simulation
 - particle-based spatial stochastic simulation
 - + sub-volume stochastic reaction-diffusion
- Green's function reaction dynamics
 - + Livermore solver
- + algorithm using continuous variables
- + algorithm using deterministic rules
- + algorithm using discrete variables
- + algorithm using fixed timesteps
- + algorithm using non-spatial description
- + algorithm using spatial description
- + algorithm using stochastic rules

ID: KISAO:0000000
Full Id: http://purl.bioontology.org/ontology/KISAO/KISAO_0000000
Definitions: Algorithm used to instantiate a simulation from a mathematical model, where the variable values evolve over time.
Xref Definition: dk:260508

     zotero 

Old KiSAO: subsumptions, multiple inheritance

The screenshot shows the Old KiSAO interface with a tree view of simulation algorithms on the left and a detailed view of a specific method on the right.

Left Panel (Tree View):

- Thing
 - 'kinetic simulation algorithm'
 - 'algorithm using adaptive timesteps'
 - 'algorithm using continuous variables'
 - 'algorithm using deterministic rules'
 - 'algorithm using discrete variables'
 - 'algorithm using fixed timesteps'
 - 'Euler backward method'
 - 'Euler forward method'
 - 'Runge-Kutta based method'
 - 'StochSim nearest-neighbour algorithm'
 - 'brownian diffusion Smoluchovski method'**
 - 'deterministic cellular automata update algorithm'
 - 'multi-state agent-based simulation method'
 - 'algorithm using non-spatial description'
 - 'algorithm using spatial description'
 - 'Green's function reaction dynamics'
 - 'StochSim nearest-neighbour algorithm'
 - 'binomial tau-leap spatial stochastic simulation algorithm'
 - 'brownian diffusion Smoluchovski method'
 - 'deterministic cellular automata update algorithm'
 - 'partial differential equation method'
 - 'particle-based spatial stochastic method'
 - 'sub-volume stochastic reaction-diffusion algorithm'
 - 'algorithm using stochastic rules'
 - 'Bortz-Kalos-Liebowitz method'
 - 'Gillespie-like stochastic simulation method'
 - 'Smoluchowski equation based method'
 - 'Green's function reaction dynamics'
 - 'brownian diffusion Smoluchovski method'**

Old KiSAO: subsumptions, multiple inheritance

The screenshot shows the Old KiSAO interface with a tree view on the left and a detailed view on the right.

Tree View (Left):

- Thing
 - 'kinetic simulation algorithm'
 - 'algorithm using adaptive timesteps'
 - 'algorithm using continuous variables'
 - 'algorithm using deterministic rules'
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 - 'algorithm using fixed timesteps'
 - 'Euler backward method'
 - 'Euler forward method'
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 - 'brownian diffusion Smoluchovski method'**
 - 'deterministic cellular automata update algorithm'
 - 'multi-state agent-based simulation method'
 - 'algorithm using non-spatial description'
 - 'algorithm using spatial description'
 - 'Green's function reaction dynamics'
 - 'StochSim nearest-neighbour algorithm'
 - 'binomial tau-leap spatial stochastic simulation algorithm'
 - 'brownian diffusion Smoluchovski method'**
 - 'deterministic cellular automata update algorithm'
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 - 'Gillespie-like stochastic simulation method'
 - 'Smoluchowski equation based method'**
 - 'Green's function reaction dynamics'
 - 'brownian diffusion Smoluchovski method'**

def

"\"In the Brownian diffusion Smoluchowski method, each molecule is treated as a point-like particle that diffuses freely in three-dimensional space. When a pair of reactive molecules collide, such as an enzyme and its substrate, a reaction occurs and the simulated reactants are replaced by products. [...] Analytic solutions are presented for some simulation parameters while others are calculated using look-up tables.\""
Supported chemical processes include molecular diffusion, treatment of surfaces, zeroth-order-, unimolecular-,

Description: 'brownian diffusion Smoluchovski method'

Equivalent classes +

Superclasses +

- 'Smoluchowski equation based method'**
- 'algorithm using discrete variables'
- 'algorithm using fixed timesteps'
- 'algorithm using spatial description'**

Inherited anonymous classes

Members +

Keys +

New KiSAO: 'algorithm characteristic' branch, 'hasProperty' relation

Class hierarchy: 'Brownian diffusion Smoluchowski method'

comment

"In the Brownian diffusion Smoluchowski method, "each molecule is treated as a point-like particle that diffuses freely in three-dimensional space. When a pair of reactive molecules collide, such as an enzyme and its substrate, a reaction occurs and the simulated reactants are replaced by products. [...] Analytic solutions are presented for some simulation parameters while others are calculated using look-up tables." Supported chemical processes include molecular diffusion, treatment of surfaces, zeroth-order-, unimolecular-, and bimolecular reactions."

label

"Brownian diffusion Smoluchowski method"^^string

Description: 'Brownian diffusion Smoluchowski method'

Equivalent classes +

Superclasses +

- 'Smoluchowski equation based method'
- hasProperty some 'discrete variables'
- hasProperty some 'progression with fixed timesteps'
- hasProperty some 'spatial description'

Inherited anonymous classes

- hasProperty some 'stochastic system behaviour'

The screenshot shows the KiSAO class hierarchy interface. At the top, there's a navigation bar with icons for search, refresh, and other functions. Below it, the 'Class hierarchy' section shows the path: 'Brownian diffusion Smoluchowski method' > 'Thing'. The tree view on the left lists various algorithm types and characteristics. The 'Brownian diffusion Smoluchowski method' node is highlighted in blue. To the right of the tree, detailed information is provided for this node, including its definition (comment), label ('Brownian diffusion Smoluchowski method'), and superclasses. Below these, four 'hasProperty' relations are listed: 'discrete variables', 'progression with fixed timesteps', 'spatial description', and 'stochastic system behaviour'. The 'stochastic system behaviour' entry is also highlighted in blue.

New KiSAO: 'algorithm characteristic' branch, 'hasProperty' relation

Class hierarchy: 'Brownian diffusion Smoluchowski method'

comment

In the Brownian diffusion Smoluchowski method, "each molecule is treated as a point-like particle that diffuses freely in three-dimensional space. When a pair of reactive molecules collide, such as an enzyme and its substrate, a reaction occurs and the simulated reactants are replaced by products. [...] Analytic solutions are presented for some simulation parameters while others are calculated using look-up tables." Supported chemical processes include molecular diffusion, treatment of surfaces, zeroth-order-, unimolecular-, and bimolecular reactions."

label

"Brownian diffusion Smoluchowski method"^^string

Description: 'Brownian diffusion Smoluchowski method'

Equivalent classes +

Superclasses +

- 'Smoluchowski equation based method'
- hasProperty some 'discrete variables'
- hasProperty some 'progression with fixed timesteps'
- hasProperty some 'spatial description'

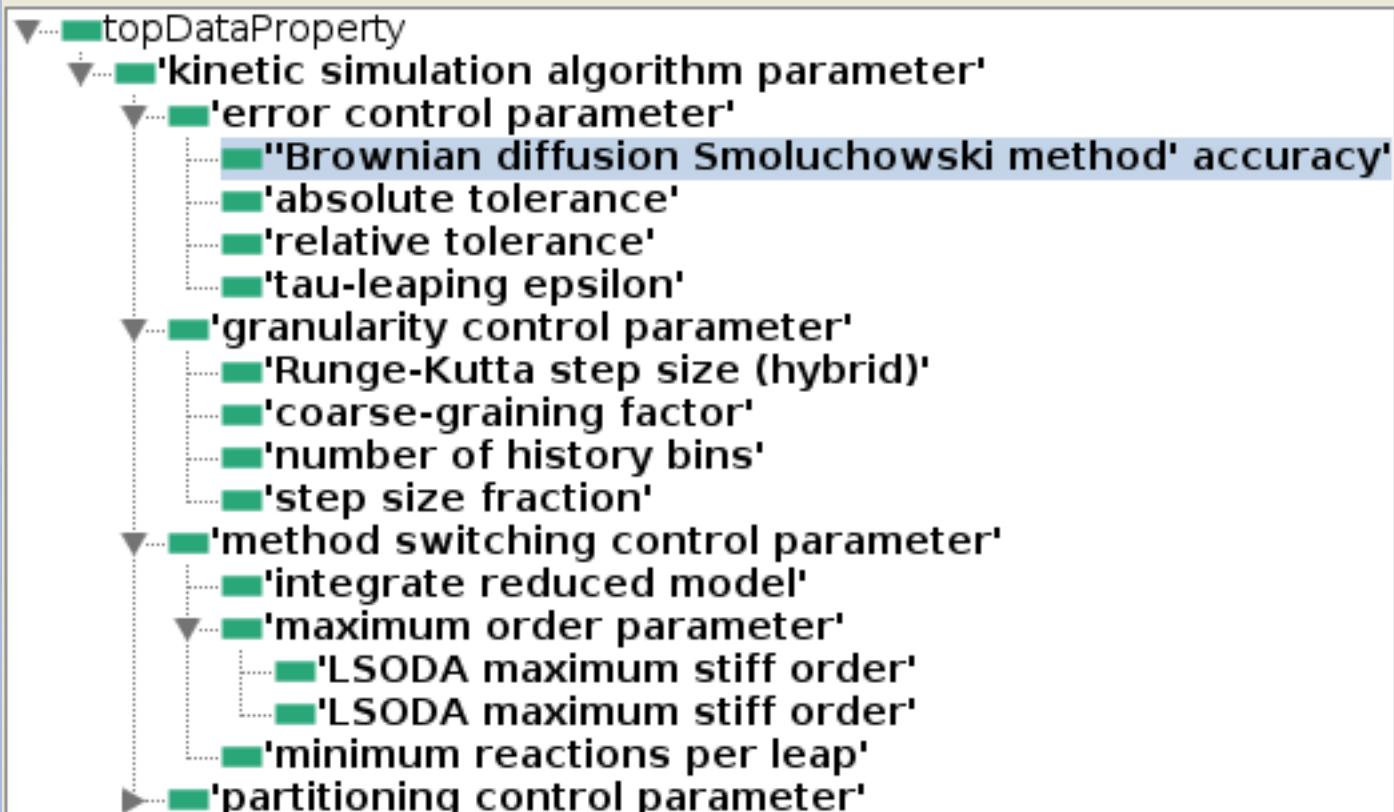
Inherited anonymous classes

- hasProperty some 'stochastic system behaviour'

hasProperty

- 'kinetic simulation algorithm'
- CVODE
- Euler method
- Gillespie-like method
- Livermore solver
- Pahle hybrid method
- Runge-Kutta based method
- 'Smoluchowski equation based method'
- 'Brownian diffusion Smoluchowski method'
- GFRD
- cellular automata update method
- multi-state agent-based simulation method
- 'kinetic simulation algorithm characteristic'
- implicit or explicit method type
- solution type
- spatial description
- system behaviour
 - deterministic system behaviour
 - stochastic system behaviour
- type of progression timesteps
 - progression with adaptive timesteps
 - progression with fixed timesteps
- type of variables
 - continuous variables
 - discrete variables

New KiSAO: algorithm parameters



Consider the reaction $A + B \rightarrow C$. Suppose that A and B are within the binding radius of each other. Then the reaction will always be performed when A and B are in the same virtual bin. If the accuracy is set to at least 3, then no reaction will occur if A and B are in different bins.

Description: "Brownian diffusion Smoluchowski method"

Domains (intersection) +

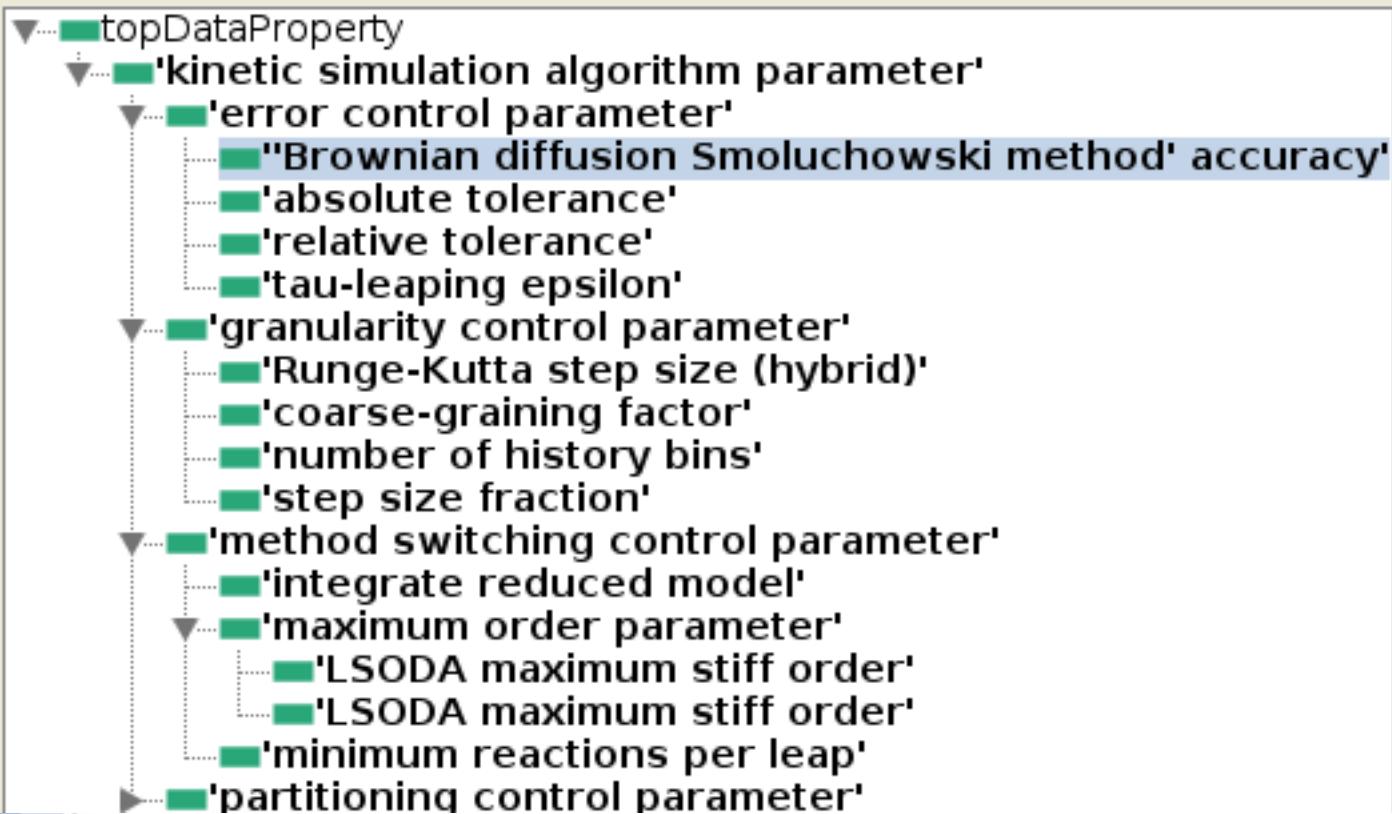
● 'Brownian diffusion Smoluchowski method'

Ranges +

● int

Equivalent properties +

New KiSAO: algorithm parameters



Consider the reaction $A + B \rightarrow C$. Suppose that A and B are within the binding radius of each other. The reaction will always be performed if A and B are in the same virtual bin. Accuracy is set to at least 3, though it also occurs if A and B are in different bins.

Description: "Brownian diffusion Smoluchowski method"

Domains (intersection) +

'Brownian diffusion Smoluchowski method'

Ranges +

int

Equivalent properties +

Algorithm Parameter Survey

[https://www.surveymonkey.com/
s/kisao_parameters](https://www.surveymonkey.com/s/kisao_parameters)

Acknowledgements



Dagmar Waltemath



Anna Zhukova