

Libsmoldyn User's Manual

for Smoldyn version 2.62

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Chapter 1

About Libsmoldyn

Libsmoldyn is a C, C++, and Python interface to the Smoldyn simulator. Libsmoldyn is complementary to the stand-alone Smoldyn program in that it is a little more difficult to use, but it provides much more flexibility. In addition, Libsmoldyn provides: *(i)* an application programming interface (API) that will be relatively stable, even as Smoldyn is updated and improved, *(ii)* function names that are relatively sensible and that shouldn't collide with other function names in other software, and *(iii)* reasonably thorough error checking in every function which helps ensure that the user is using the function in a sensible way and in a way that won't crash Smoldyn.

Libsmoldyn was originally written as a C API, but one that should work with C++ as well. Then, in 2019 and 2020, Dilawar Singh added Python bindings to it, which are still quite new but generally work well.

Libsmoldyn only barely supports graphics at present due to constraints imposed by the glut code library. This will be improved in a future version (by changing to the freeglut library, which doesn't insist on controlling the main event loop, as the glut library does).

Chapter 2

Linking with C/C++

2.1 Compiling

Header files

To enable a C or C++ program to call Libsmoldyn, it has to include the Libsmoldyn header file. Libsmoldyn comes with one header file, `libsmoldyn.h`, which has function declarations for all of the Libsmoldyn functions. For most Libsmoldyn applications, this is the only header file that you will need to include. For Mac and Linux, it is typically installed to `/usr/local/include`. This is one of the standard system paths, so include it with

```
#include <libsmoldyn.h>
```

If the `libsmoldyn.h` header file is in some other directory or if your system isn't seeing its path as a system path, then include the file using double quotes rather than angle brackets and/or include more information about the path. For example, `#include "/user/local/include/libsmoldyn.h"`.

`Libsmoldyn.h` calls a second header file, `smoldyn.h`, which is also typically installed to `/usr/local/include/`. If you plan to access the Smoldyn data structure directly, then you will also need to include it with `#include <smoldyn.h>`. In general, it is safe to read from this data structure but it can be dangerous to write to it unless you really know what you are doing. Also, working with this data structure directly bypasses one of the benefits of using Libsmoldyn, which is that the interface should be relatively immune to future Smoldyn developments, because different aspects of the internal data structure get changed once in a while.

The `smoldyn.h` header calls yet another header file, `smoldynconfigure.h`, which is also installed by default in `/usr/local/include/`. That file is automatically generated by the build system. It describes what Smoldyn components are included in the build, what system the build was compiled for, etc. This might be helpful to include for some applications.

Compiling example

In the `examples/S97_libsmoldyn/testcode/` directory, you'll find the `testcode.c` program. To compile this source code to object code, enter:

```
gcc -Wall -O0 -g -c testcode.c
```

The compile flags `-O0 -g` aren't necessary but can be useful for debugging purposes. If compiling doesn't work at this stage, it's probably because you're missing the header files. Make sure that you have `libsmoldyn.h`, `smoldyn.h`, and `smoldyn_config.h` in the `/usr/local/include` directory.

2.2 Linking

Linking the different object files together to create an executable that actually runs is often one of the greatest frustrations of using software libraries. It should be easy but usually isn't.

The Libsmoldyn library can be linked statically, meaning that the Libsmoldyn code will be copied into the final result, or it can be linked dynamically, so that the final result will simply reference the Libsmoldyn code that is stored separately. Dynamic linking is somewhat more elegant in that it doesn't create unnecessary copies of the compiled code. It can also be easier. On the other hand, it's less convenient if you plan to distribute your software, because then you need to make sure that you distribute the Libsmoldyn header file and library code along with your own software. Also, I can only get the gdb debugger to help find errors within Libsmoldyn if the library is statically linked.

The Libsmoldyn static library is called `libsmoldyn_static.a` and the Libsmoldyn dynamic library is called `libsmoldyn_shared.so` (on Linux; the `.so` suffix is replaced by `.dylib` on a Mac and by `.dll` on Windows). By default, these libraries are installed to `/usr/local/lib/`.

Linking examples

Following are several example for static and dynamic linking. They are shown for C; if you use C++, then link with `g++` instead of `gcc`. The linking options for Smoldyn compiled with OpenGL are shown for Macintosh; these lines are simpler for other systems.

I have had a hard time getting static linking working on a Mac, although apparently it works fine on Ubuntu. The problem is that it doesn't find the standard C++ library. The solution is to build the Smoldyn library without NSV, so that the standard C++ library isn't needed. I also commented out a few "throw" statements from `smolsim.c` and `libsmoldyn.c` for this purpose.

Static link, no OpenGL:

```
gcc testcode.o /usr/local/lib/libsmoldyn_static.a -o testcode
```

Static link, with OpenGL:

```
gcc testcode.o /usr/local/lib/libsmoldyn_static.a -I/System/Library/Frameworks/OpenGL.framework/Headers -I/System/Library/Frameworks/GLUT.framework/Headers -framework GLUT -framework OpenGL -framework Cocoa -L/System/Library/Frameworks/OpenGL.framework/Libraries -o testcode -ltiff
```

Dynamic link, no OpenGL:

```
gcc testcode.o -o testcode -lsmoldyn_shared
```

Dynamic link, with OpenGL:

```
gcc test1.o -L/usr/local/lib -I/System/Library/Frameworks/OpenGL.framework/Headers -I/System/Library/Frameworks/GLUT.framework/Headers -framework GLUT -framework OpenGL -framework Cocoa -L/System/Library/Frameworks/OpenGL.framework/Libraries -o test1 -lsmoldyn_shared -ltiff
```


Chapter 3

Use with Python

3.1 Installing

Hopefully, you should be able to install using

```
python3 -m pip install smoldyn --user --pre
```

This should install the smoldyn nightly package, from: <https://pypi.org/project/smoldyn/>

Alternatively, the Mac distribution comes with a Python wheel, called something like smoldyn-2.62-py3-none-any.whl. You should be able to write “pip install smoldyn...whl” and that will install it for you as well.

For Windows, “pip install smoldyn” seems to work initially, but it doesn’t install the necessary compiled code, so it doesn’t actually run, yet.

3.2 Files

After building Smoldyn (assuming you were in /path/to/Smoldyn-official/build directory when you ran make command), the python module ends up in the /path/to/Smoldyn-official/build/py directory. The module can be imported into Python from this directory. Add this directory to your PYTHONPATH temporarily with `export PYTHONPATH=/path/to/Smoldyn-official/build/py:$PYTHONPATH`. With this, the module can be accessed from any directory.

Note that it’s possible to see where the library is imported from by checking `smoldyn.__file__` while in Python. For example,

```
>>> import smoldyn
>>> print(smoldyn.__file__)
/home/dilawars/Work/FORKES/Smoldyn-official/build/py/smoldyn/__init__.py
```

3.3 Python-specific functionality

Python offers a callback function that enables Smoldyn setup functions to be called repeatedly and automatically. Following is the text from Dilawar’s description.

Suppose I have the following function which generates a noisy value using the current time `t` and a list of arguments `args`.

```
def computeVm(t, args):
    x, y = args
    return math.sin(t) + x * y + random.random()
```

And I have a molecular species `ca`.

```
import smoldyn
ca = smoldyn.Species('ca', difc=1, color='blue', display_size=1)
```

Then the following connect the output of function computeVm to ca.difc parameter. This is called every 10th step.

```
smoldyn.connect(computeVm, "ca.difc", step=10, args=[1,2.1])
```

In smoldyn.connect function, both source and target must be global variables. In the example below, I made a a global variable before I used it in connect. Else there will be a runtime error.

```
import smoldyn as S
import random

a = None

def new_dif(t, args):
    global a
    x, y = args
    print(a.difc)
    return t + random.random()

def test_connect():
    global a
    S.setBounds(low=(0, 0), high=(10, 10))
    a = S.Species("a", color="red", difc=1)
    S.connect(new_dif, "a.difc", step=10, args=[0, 1])
    s = S.Simulation(100, 1)
    s.run()
    print("All done")

test_connect()
```

Also accepts a function.

```
def new_dif(t, args):
    global a, avals
    x, y = args
    # note that b.difc is not still updated.
    avals.append((t, a.difc["soln"]))
    return x * math.sin(t) + y

def update_difc(val):
    global a
    a.difc = val

def test_connect():
    global a, avals
    S.setBounds(low=(0, 0), high=(10, 10))
    a = S.Species("a", color="black", difc=0.1)
    S.connect(new_dif, update_difc, step=10, args=[1, 1])
    s = S.Simulation(100, 1)
    s.run()
    for a, b in zip(avals[1:], expected_a[1:]):
        print(a, b)
```

```
    assert math.isclose(a[1], b[1], rel_tol=1e-6, abs_tol=1e-6), (a[1], b[1])

test_connect()
```

Dilawar created some nice examples of this in use with a pre-synaptic bouton with N synaptic vesicles. These vesicles fuse with the bottom of the bouton (red surface). Upon fusion, one vesicle releases 1000 neurotransmitters which decay with time-constant τ .

The rate of release is controlled by a function. This is set by `smoldyn.connect`. The function generates a spike 0 or 1, if the value is 1 the rate is set to 1000 else it is 0.

Chapter 4

Error trapping

Every function in Libsmoldyn checks that its input values are acceptable and also that no errors arise in the function execution. These errors are returned to the host library in a number of ways. Most Libsmoldyn functions (e.g. `smolRunSim`) return any error codes directly, which makes it easy to see if an error arose. However, a few functions (e.g. `smolNewSim`) return other types of values and so return some other indication of success or failure (e.g. `NULL`). In addition, some functions can raise warnings, which indicate that behavior is unusual but not incorrect.

For all of these errors and warnings, get the details of the problem using the function `smolGetError`, which will return the error code, the name of the function where the error arose, and a descriptive error string. This will also clear the error, if desired. If errors are not cleared, they are left until they are overwritten by subsequent errors. Warnings are also left until they are cleared or overwritten.

When writing code, it can be helpful to put Libsmoldyn into its debugging mode using the `smolSetDebugMode` function. Doing this causes any errors that arise to be displayed to `stderr`.

The possible error codes are declared in `libsmoldyn.h` with:

```
enum ErrorCode {ECok=0, ECnotify=-1, ECwarning=-2, ECnonexist=-3, ECall=-4,
    ECmissing=-5, ECbounds=-6, ECSyntax=-7, EError=-8, ECmemory=-9, Ebug=-10,
    ECsame=-11}
```

Their interpretations are:

value	code	interpretation
0	ECok	no error
-1	ECnotify	message about correct behavior
-2	ECwarning	unusual but not incorrect behavior
-3	ECnonexist	a function input specifies an item that doesn't exist
-4	ECsame	error code should be unchanged from a prior code
-5	ECall	an argument of "all" was found and may not be permitted
-6	ECmissing	a necessary function input parameter is missing
-7	ECbounds	a function input parameter is out of bounds
-8	ECSyntax	function inputs don't make syntactical sense
-9	EError	unspecified error condition
-10	ECmemory	Smoldyn was unable to allocate the necessary memory
-11	Ebug	error arose which should not have been possible

4.1 Error checking system internal to `libsmoldyn.c`

This section describes how to write Libsmoldyn functions using error checking. While it is an essential part of all Libsmoldyn functions, these details are not important for most Libsmoldyn users.

1. The first line of every Libsmoldyn function should be `const char *funcname="function_name";`. This

name will be returned with any error message to tell the user where the error arose.

2. Within the function, check for warnings or errors with either the `LCHECK` or `LCHECKNT` macros. In both cases, the macro format is `LCHECK(condition, funcname, error_code, "message");`. The macros check that the test *condition* is true, and calls either `smolSetError` or `smolSetErrorNT` to deal with it if not. The *message* should be a descriptive message that is under 256 characters in length. Use the regular version (not the “no throw” or “NT”) version for errors that arise within the function, and the “NT” version for errors that arise in subroutines of the function, so that only a single error message is displayed to the output.
3. Most functions return an “enum `ErrorCode`”. If this is the case for your function, and your function might return a notification and/or a warning, then end the main body of the function with `return libwarncode;`. If it cannot return a notification or a warning, then end it with `return ECok;`. Finally, if it does not return an “enum `ErrorCode`”, then it needs to return some other error condition that will tell the user to check for errors using `smolGetError`.
4. After the main body of the function, add a goto target called `failure:`.
5. Assuming the function returns an “enum `ErrorCode`”, end the function with `return liberrorcode;`.

The `smolSetTimeStep` function provides an excellent and simple example of how Libsmoldyn functions typically address errors. It is:

```
enum ErrorCode smolSetTimeStep(simptr sim, double timestep) {
    const char *funcname="smolSetTimeStep";

    LCHECK(sim, funcname, ECmissing, "missing sim");
    LCHECK(timestep>0, funcname, ECbounds, "timestep is not > 0");
    simsettime(sim, timestep, 3);
    return ECok;
failure:
    return liberrorcode; }
```

The `smolGet...Index` functions are worth a comment. Each of these functions returns the index of an item, such as a species or a surface, based on the name of the item. If the name is not found or other errors arise, then these functions return the error code, cast as an integer. Also, if the name is “all”, then these functions return the error code `ECall` and set the error string “species cannot be ‘all’”, or equivalent. A typical use of these functions is seen in `smolSetSpeciesMobility`, which includes the following code:

```
i=smolGetSpeciesIndex(sim, species);
if(i==(int)ECall) smolClearError();
else LCHECK(i>0, funcname, ECsame, NULL);
```

In this particular case, this function permits an input of “all”, so it clears errors that arise from this return value, and leaves `i` as a negative value for later use.

Chapter 5

Libsmoldyn quick function guide

The Libsmoldyn functions correspond relatively closely to the Smoldyn language statements, although not perfectly. However, all functionality should be available using either method. The following table lists the correspondences. Statements preceded by asterisks need to be either entered in statement blocks or preceded by the statement’s context (e.g. with **surface** *name*). Where correspondence does not apply, the table lists “N/A”. The Libsmoldyn functions are available either through the C/C++ API or through the Python API, with essentially identical input styles. The Python functions listed here use a more object-oriented approach. Here, “S” is short for smoldyn, arising for example as `import smoldyn as S`.

Statement	Libsmoldyn function	Python function
About the input		
#	N/A	
/* ... */	N/A	
read_file	LoadSimFromFile	
	ReadConfigString	
end_file	N/A	
define	N/A	
define_global	N/A	
undefine	N/A	
ifdefine	N/A	
ifundefine	N/A	
else	N/A	
endif	N/A	
display_define	N/A	
N/A	SetError	
N/A	GetError	
N/A	ClearError	
N/A	SetDebugMode	
N/A	ErrorCodeToString	
Space and time		
dim	NewSim	
boundaries	NewSim	
	SetBoundaryType	
low_wall	NewSim	
	SetBoundaryType	
high_wall	NewSim	
	SetBoundaryType	
time.start	SetSimTimes	
	SetTimeStart	
time.stop	SetSimTimes	

time_step	SetTimeStop SetSimTimes SetTimeStep	
time_now	SetTimeNow	
Molecules		
species	AddSpecies	S.Species
N/A	GetSpeciesIndex	
N/A	GetSpeciesName	
difc	SetSpeciesMobility	<i>species.difc</i>
difm	SetSpeciesMobility	
drift	SetSpeciesMobility	
mol	AddSolutionMolecules	<i>species.addToSolution</i>
surface_mol	AddSurfaceMolecules	
compartment_mol	AddCompartmentMolecules	
molecule_lists	AddMolList	
mol_list	AddSpecies SetMolList	<i>species.mol_list</i>
N/A	GetMolListIndex	
N/A	GetMolListName	
max_mol	SetMaxMolecules	
N/A	GetMoleculeCount	
Graphics		
graphics	SetGraphicsParams	
graphic_iter	SetGraphicsParams	
graphic_delay	SetGraphicsParams	
frame_thickness	SetFrameStyle	
frame_color	SetFrameStyle	
grid_thickness	SetGridStyle	
grid_color	SetGridStyle	
background_color	SetBackgroundStyle	
display_size	SetMoleculeStyle	<i>species.setStyle</i> <i>species.size</i>
color	SetMoleculeStyle	<i>species.color</i>
tiff_iter	SetTiffParams	
tiff_name	SetTiffParams	
tiff_min	SetTiffParams	
tiff_max	SetTiffParams	
light	SetLightParams	
text_color	SetTextStyle	
text_display	AddTextDisplay	
Run-time commands		
output_root	SetOutputPath	
output_files	AddOutputFile	
append_files	AddOutputFile	
output_file_number	AddOutputFile	
cmd	AddCommand AddCommandFromString	
Surfaces		
start_surface	AddSurface	
new_surface	AddSurface	
* name	AddSurface	
N/A	GetSurfaceIndex	
N/A	GetSurfaceName	
* action	SetSurfaceAction	

* rate	SetSurfaceRate	
* rate_internal	SetSurfaceRate	
* color	SetSurfaceFaceStyle	
	SetSurfaceEdgeStyle	
* thickness	SetSurfaceEdgeStyle	
* stipple	SetSurfaceEdgeStyle	
* polygon	SetSurfaceFaceStyle	
* shininess	SetSurfaceFaceStyle	
* panel	AddPanel	
N/A	GetPanelIndex	
N/A	GetPanelName	
* jump	SetPanelJump	
* neighbors	AddPanelNeighbor	
* unbounded_emitter	AddSurfaceUnboundedEmitter	
* end_surface	N/A	
epsilon	SetSurfaceSimParams	
margin	SetSurfaceSimParams	
neighbor_dist	SetSurfaceSimParams	
Compartments		
start_compartment	AddCompartment	
new_compartment	AddCompartment	
* name	AddCompartment	
N/A	GetCompartmentIndex	
N/A	GetCompartmentName	
* surface	AddCompartmentSurface	
* point	AddCompartmentPoint	
* compartment	AddCompartmentLogic	
* end_compartment	N/A	
Reactions		
reaction	AddReaction	S.Reaction
N/A	GetReactionIndex	
N/A	GetReactionName	
reaction_cmpt	SetReactionRegion	
reaction_surface	SetReactionRegion	
reaction_rate	AddReaction	
	SetReactionRate	
confspread_radius	SetReactionRate	
binding_radius	SetReactionRate	
reaction_probability	SetReactionRate	
reaction_production	SetReactionRate	
reaction_permit	not supported	
reaction_forbid	not supported	
product_placement	SetReactionProducts	
Ports		
start_port	AddPort	
new_port	AddPort	
* name	AddPort	
N/A	GetPortIndex	
N/A	GetPortName	
* surface	AddPort	
* face	AddPort	
* end_port	N/A	
N/A	AddPortMolecules	

N/A	GetPortMolecules
Simulation settings	
rand_seed	SetRandomSeed
accuracy	not supported
molperbox	SetPartitions
boxsize	SetPartitions
gauss_table_size	not supported
epsilon	SetSurfaceSimParams
margin	SetSurfaceSimParams
neighbor_dist	SetSurfaceSimParams
pthreads	not supported
Libdyn actions	
N/A	UpdateSim
N/A	RunTimeStep
N/A	RunSim
N/A	RunSimUntil
N/A	FreeSim
N/A	DisplaySim
N/A	PrepareSimFromFile

Chapter 6

Data structures and declarations

6.1 Enumerations

In C, enumerations are already defined, so they can be used as is. Here is an example of using an enumerated error code as an argument,

```
smolErrorCodeToString(ECwarning, mystring)
```

In Python, enumerations are most easily dealt with by defining a variable for the enumerated list and then choosing from it. Here is an example,

```
import smoldyn._smoldyn as S
EC=S.ErrorCode
S.errorCodeToString(EC.warning, mystring)
```

Surface actions (SrfAction)

Statement	Libsmoldyn	Python	Notes
reflect	SAreffect	reflect	
transmit	SAttrans	trans	
absorb	SAabsorb	absorb	
jump	SAjump	jump	
port	SAsport	port	
multiple	SAmult	mult	multiple actions
N/A	SAno	no	static surface-bound molecules
N/A	SAnone	none	none of the other options
N/A	SAadsorb	adsorb	internal use only
N/A	SArevdes	revdes	internal use only
N/A	SAirrevdes	irrevdes	internal use only
N/A	SAflip	flip	internal use only

Molecule state (MolecState)

Statement	Libsmoldyn	Python	Notes
soln	MSsoln	soln	
front	MSfront	front	
back	MSback	back	
up	MSup	up	
down	MSdown	down	
bsoln	MSbsoln	bsoln	pseudo-state for surface interactions
all	MSall	all	for model creation by user
N/A	MSnone	none	internal use only

Panel face (PanelFace)

Statement	Libsmoldyn	Python	Notes
front	PFfront	front	
back	PFback	back	
N/A	PFnone	none	internal use only
both	PFboth	both	for model creation by user

Panel shape (PanelShape)

Statement	Libsmoldyn	Python	Notes
rect	PSrect	rect	rectangle
tri	PStri	tri	triangle
sph	PSsph	sph	sphere
cyl	PScyl	cyl	cylinder
hemi	PShemi	hemi	hemisphere
disk	PSdisk	disk	disk
all	PSall	all	for model creation by user
N/A	PSnone	none	internal use only

Libsmoldyn error code (ErrorCode)

Statement	Libsmoldyn	Python	Notes
N/A	ECok	ok	value 0
N/A	ECnotify	notify	value -1
N/A	ECwarning	warning	value -2
N/A	ECnonexist	nonexist	value -3
N/A	ECall	all	value -4
N/A	ECmissing	missing	value -5
N/A	ECbounds	bounds	value -6
N/A	ECsyntax	syntax	value -7
N/A	ECerror	error	value -8
N/A	ECmemory	memory	value -9
N/A	ECbug	bug	value -10
N/A	ECsame	same	value -11
N/A	ECwildcard	wildcard	value -12

6.2 Libsmoldyn header file

The Libsmoldyn header file is libsmoldyn.h. It lists all of the function declarations. This file references smoldyn.h, which lists all of the data structure declarations and enumerated type definitions.

If you compiled and installed Smoldyn using the default configuration, both files should be in your /usr/local/include/smoldyn directory. Also in this directory is the smoldyn_config.h file. This file was used for compiling Smoldyn and Libsmoldyn but is not needed afterwards. Nevertheless, it's copied to the /usr/local/include/smoldyn directory so that programs that call Libsmoldyn can know what options Libsmoldyn was built with.

Chapter 7

Libsmoldyn functions

7.1 General comments

None of the functions allocate memory, except within the simulation data structure. This means, for example, that all functions that return strings do not allocate these strings themselves, but instead write the string text to memory that the library user allocated and gave to the function. All strings are fixed at `STRCHAR` characters, where this constant is defined in `string2.h` to 256 characters.

7.2 Miscellaneous

GetVersion

C: `double smolGetVersion(void)`
Python: `float getVersion()`
Returns the Smoldyn version number.

7.3 Errors

SetError

C: `void smolSetError(const char *errorfunction, enum ErrorCode errorcode, const char *errorstring)`
C: `void smolSetErrorNT(const char *errorfunction, enum ErrorCode errorcode, const char *errorstring)`
Python: N/A

These functions are probably not useful for most users. Sets the Libsmoldyn error code to `errorcode`, error function to `errorfunction`, and error string to `errorstring`. The sole exception is if `errorcode` is `ECsame` then this does nothing and simply returns. Back to it's normal operation, this also either sets or clears the Libsmoldyn warning code, as appropriate. If `errorstring` is entered as `NULL`, this clears the current error string, and similarly for `errorfunction`. For the regular version without the "NT", if the library is being run in debug mode, then this function prints the notification, warning, or error out to `stderr`. It would also ideally throw exceptions if the error code is more severe than the `LibThrowThreshold` value, but this throwing doesn't work at present because throwing exceptions to the host code is incompatible with static linking.

The "NT" version is the "no throw" version, which is the same as the regular version but doesn't display messages to `stderr` and doesn't throw exceptions. In general, library functions should call `smolSetError` for errors caught by that function itself, and `smolSetErrorNT` for errors caught by subroutines of that function, so that each error only leads to a single call of `smolSetError`.

GetError

C: `enum ErrorCode smolGetError(char *errorfunction, char *errorstring, int`

`clearerror()`

Python: N/A

Returns the current LibSmoldyn error code directly, returns the function where the error occurred in `errorfunction` if it is not NULL, and returns the error string in `errorstring` if it is not NULL. Set `clearerror` to 1 to clear the error and 0 to leave any error condition unchanged.

ClearError

C: `void smolClearError(void)`

Python: N/A

Clears any error condition.

SetDebugMode

C: `void smolSetDebugMode(int debugmode)`

Python: `setDebugMode(int debugmode)`

Enter `debugmode` as 1 to enable debugging and 0 to disable debugging. When debug mode is turned on, all errors are displayed to `stderr`, as are all cleared errors. By turning on debug mode, you can often avoid checking for errors with additional code and you also typically don't need to call `smolGetError`.

ErrorCodeToString

C: `char* smolErrorCodeToString(enum ErrorCode ERC, char *string)`

Python: `str errorCodeToString(enum ErrorCode ERC, str string)`

Returns a string both directly and in `string` that corresponds to the error code in `ERC`. For example, if `ERC` is `ECmemory`, this returns the string "memory". To do: The string is not needed or used in the Python version.

7.4 Sim structure

NewSim

C: `simptr smolNewSim(int dim, double *lowbounds, double *highbounds)`

Python: `simptr newSim(int dim, List[float] lowbounds, List[float] highbounds)`

Creates and returns a new sim structure. The structure is initialized for a `dim` dimensional system that has boundaries defined by the points `lowbounds` and `highbounds`. Boundaries are transmitting (modify them with `smolSetBoundaryType`). Returns NULL upon failure.

Python to do: there's no need for the `dim` parameter because it's redundant with vector lengths.

UpdateSim

C: `enum ErrorCode smolUpdateSim(simptr sim)`

Python: `ErrorCode updateSim()`

Updates the simulation structure. This calculates all simulation parameters from physical parameters, sorts lists, and generally does everything required to make a simulation ready to run. It may be called multiple times.

Python to do: doesn't work. Python wants no argument, but Libsmoldyn then complains about no argument.

RunTimeStep

C: `enum ErrorCode smolRunTimeStep(simptr sim)`

Python: `ErrorCode runTimeStep()`

Runs one time step of the simulation. Returns an error if the simulation terminates unexpectedly during this time step or a warning if it terminates normally.

Python to do: doesn't work. Python wants no argument, but Libsmoldyn then complains about no argument.

RunSim

C: `enum ErrorCode smolRunSim(simptr sim)`

Python: `ErrorCode runSim()`

Python: `run(stop=None, start=None, step=None)`

Runs the simulation until it terminates. Returns an error if the simulation terminates unexpectedly during this time step or a warning if it terminates normally.

Python to do: doesn't work. Python wants no argument, but Libsmoldyn then complains about no argument.

RunSimUntil

C: `enum ErrorCode smolRunSimUntil(simptr sim, double breaktime)`

Python: `ErrorCode runSimUntil(float breaktime)`

Python: `runUntil(t, dt)`

Runs the simulation either until it terminates or until the simulation time equals or exceeds **breaktime**.

Python to do: doesn't work. Python wants no argument, but Libsmoldyn then complains about no argument.

FreeSim

C: `enum ErrorCode smolFreeSim(simptr sim)`

Python: `ErrorCode freeSim()`

Frees the simulation data structure.

DisplaySim

C: `enum ErrorCode smolDisplaySim(simptr sim)`

Python: `ErrorCode displaySim()`

Displays all relevant information about the simulation system to stdout.

7.5 Read configuration file

PrepareSimFromFile

C: `simptr smolPrepareSimFromFile(char *filepath, char *filename, char *flags)`

Python: `simptr prepareSimFromFile(str filename, str flags)`

Reads the Smoldyn configuration file that is at **filepath** and has file name **filename**, sets it up, and outputs simulation diagnostics to stdout. Returns the sim structure, or NULL if an error occurred. **flags** are the command line flags that are entered for normal Smoldyn use. Either or both of **filepath** and **flags** can be sent in as NULL if there is nothing to report. After this function runs successfully, it should be possible to call **smolRunSim** or **smolRunTimeStep**.

LoadSimFromFile

C: `enum ErrorCode smolLoadSimFromFile(char *filepath, char *filename, simptr *simpointer, char *flags)`

Python: `ErrorCode loadSimFromFile(str filename, str flags)`

Loads part or all of a sim structure from the file that is at **filepath** and has file name **filename**. Send in **simpointer** as a pointer to sim, where sim may be an existing simulation structure that this function will append or NULL if it is to be created by this function. **flags** are the command line flags that are entered for normal Smoldyn use. Either or both of **filepath** and **flags** can be sent in as NULL if there is nothing to report. After this function runs successfully, call **smolUpdateSim** to calculate simulation parameters.

ReadConfigString

C: `enum ErrorCode smolReadConfigString(simptr sim, char *statement, char *parameters)`

Python: `ErrorCode readConfigString(str statement, str parameters)`

Reads and processes what would normally be a single line of a configuration file. The first word of the line is the statement name, entered here as **statement**, while the rest of the line is entered as **parameters**. Separate different parameters with spaces. The same parser is used as for normal Smoldyn configuration files. This function does not make use of block style input formatting, such as for surface definitions. This means that a new surface needs to be declared with “**new_surface name**” and all subsequent surface definitions need to start with “**surface name**”. Analogous rules apply to compartments and port.

7.6 Simulation settings

SetSimTimes

C: `enum ErrorCode smolSetSimTimes(simptr sim, double timestart, double timestep, double timestep)`

Python: `ErrorCode setSimTimes(float timestart, float timestep, float timestep)`

Sets all of the simulation time parameters to the values entered here. In addition the simulation “time now” is set to `timestart`.

SetTimeStart

C: `enum ErrorCode smolSetTimeStart(simptr sim, double timestart)`

Python: `ErrorCode setTimeStart(float timestart)`

Sets the simulation starting time.

SetTimeStop

C: `enum ErrorCode smolSetTimeStop(simptr sim, double timestep)`

Python: `ErrorCode setTimeStop(float timestep)`

Sets the simulation stopping time.

SetTimeNow

C: `enum ErrorCode smolSetTimeNow(simptr sim, double timenow)`

Python: `ErrorCode setTimeNow(float timenow)`

Sets the simulation current time.

SetTimeStep

C: `enum ErrorCode smolSetTimeStep(simptr sim, double timestep)`

Python: `ErrorCode setTimeStep(float timestep)`

Sets the simulation time step, which must be greater than 0.

SetRandomSeed

C: `enum ErrorCode smolSetRandomSeed(simptr sim, double seed)`

Python: `ErrorCode setRandomSeed(int seed)`

Sets the random number generator seed to `seed` if `seed` is at least 0, and sets it to the current time value if `seed` is less than 0.

SetAccuracy

C: not supported

Python: `accuracy(accuracy: float)`

Sets or gets the simulation accuracy.

SetPartitions

C: `enum ErrorCode smolSetPartitions(simptr sim, char *method, double value)`

Python: `ErrorCode setPartitions(str method, float value)`

Python: `Partition(name: str, value: float)`

Sets the virtual partitions in the simulation volume. Enter `method` as “molperbox” and then enter `value` with the requested number of molecules per partition volume; the default, which is used if this function is not called at all, is a target of 4 molecules per box. Or, enter `method` as “boxsize” and enter `value` with the requested partition spacing. In this latter case, the actual partition spacing may be larger or smaller than the requested value in order to fit an integer number of partitions into each coordinate of the simulation volume.

The second Python option is its own class. I think this should be removed because partitions aren’t physical objects and so don’t really make sense here.

MoleculePerBox

Python: `MoleculePerBox(size: float)`

This is only available in Python. Again, I think this should be removed because partitions aren’t physical objects.

Box

Python: `Box(size: float)`

This is only available in Python. Again, I think this should be removed because partitions aren't physical objects.

7.7 Graphics

SetGraphicsParams

C: `enum ErrorCode smolSetGraphicsParams(simptr sim, char *method, int timesteps, double delay)`

Python: `ErrorCode setGraphicsParams(str method, int timesteps, int delay)`

Python: `setGraphics(method: str, timestep: int, delay: int = 0)`

Sets basic simulation graphics parameters. Enter `method` as “none” for no graphics (the default), “opengl” for fast but minimal OpenGL graphics, “opengl_good” for improved OpenGL graphics, “opengl_better” for fairly good OpenGL graphics, or as `NULL` to not set this parameter currently. Enter `timesteps` with a positive integer to set the number of simulation time steps between graphics renderings (1 is the default) or with a negative number to not set this parameter currently. Enter `delay` as a non-negative number to set the minimum number of milliseconds that must elapse between subsequent graphics renderings in order to improve visualization (0 is the default) or as a negative number to not set this parameter currently.

SetTiffParams

C: `enum ErrorCode smolSetTiffParams(simptr sim, int timesteps, char *tiffname, int lowcount, int highcount)`

Python: `ErrorCode setTiffParams(int timesteps, str tiffname, int lowcount, int highcount)`

Sets parameters for the automatic collection of TIFF format snapshots of the graphics window. `timesteps` is the number of simulation timesteps that should elapse between subsequent snapshots, `tiffname` is the root filename of the output TIFF files, `lowcount` is a number that is appended to the filename of the first snapshot and which is then incremented for subsequent snapshots, and `highcount` is the last numbered file that will be collected. Enter negative numbers for `timesteps`, `lowcount`, and/or `highcount` to not set these parameters, and enter `NULL` for `tiffname` to not set the file name.

SetLightParams

C: `enum ErrorCode smolSetLightParams(simptr sim, int lightindex, double *ambient, double *diffuse, double *specular, double *position)`

Python: `ErrorCode smolSetLightParams(int lightindex, List[float] ambient, List[float] diffuse, List[float] specular, List[float] position)`

Sets the lighting parameters that are used for the rendering method “opengl_better”. Enter `lightindex` as -1 for the global ambient light (in which case `diffuse`, `specular`, and `position` should all be `NULL`) or as 0 to 8 for one of the 8 light sources. For each light source, you can specify the 4-value color vector for the light's ambient, diffuse, and specular properties (all values should be between 0 and 1). You can also specify the 3-dimensional position for the light. To not set a property, just enter the respective vector as `NULL`.

SetBackgroundStyle

C: `enum ErrorCode smolSetBackgroundStyle(simptr sim, double *color)`

Python: `ErrorCode setBackgroundStyle(string color)`

Sets the color of the graphics display background. `color` is a 4-value vector with red, green, blue, and alpha values (or, in Python, a color word).

SetFrameStyle

C: `enum ErrorCode smolSetFrameStyle(simptr sim, double thickness, double *color)`

Python: `ErrorCode setFrameStyle(float thickness, string color)`

Sets the thickness and the color of the wire frame that outlines the simulation system in the graphics

window. Enter `thickness` as 0 for no frame, as a positive number for the number of points in thickness, or as a negative number to not set this parameter. Enter `color` as a 4-value vector with the frame color, or as NULL to not set it (or, in Python, a color word).

SetGridStyle

C: `enum ErrorCode smolSetGridStyle(simptr sim, double thickness, double *color)`

Python: `ErrorCode setGridStyle(float thickness, string color)`

Sets the thickness and the color of a grid that shows where the partitions are that separate Smoldyn's virtual boxes. Enter `thickness` as 0 for no grid, as a positive number for the number of points in thickness, or as a negative number to not set this parameter. Enter `color` as a 4-value vector with the grid color, or as NULL to not set it (or, in Python, a color word).

SetTextStyle

C: `enum ErrorCode smolSetTextStyle(simptr sim, double *color)`

Python: `ErrorCode setTextStyle(string color)`

Sets the color of any text that is displayed to the graphics window. `color` is a 4-value vector with red, green, blue, and alpha values (or, in Python, a color word).

AddTextDisplay

C: `enum ErrorCode smolAddTextDisplay(simptr sim, char *item)`

Python: `ErrorCode addTextDisplay(string item)`

Adds `item` to the list of things that Smoldyn should display as text to the graphics window. Currently supported options are "time" and the names of species and, optionally, their states. For species and states, the graphics window shows the number of molecules.

7.8 Runtime commands

SetOutputPath

C: `enum ErrorCode smolSetOutputPath(simptr sim, char *path)`

Python: `ErrorCode setOutputPath(string path)`

Sets the file path for text output files to `path`.

AddOutputFile

C: `enum ErrorCode smolAddOutputFile(simptr sim, char *filename, int suffix, int append)`

Python: `ErrorCode addOutputFile(string filename, int suffix, int append)`

Declares the file called `filename` as a file for output by one or more runtime commands. Note that spaces are not permitted in the file name. If `suffix` is non-negative, then the file name is suffixed by this integer, which can be helpful for creating output file stacks. Enter `append` as 1 if any current file should simply be appended, or to 0 if any current file should be overwritten.

OpenOutputFiles

C: `enum ErrorCode smolOpenOutputFiles(simptr sim, int overwrite = 0)`

Opens output files for writing. Enter `overwrite` as 1 if any existing file should be overwritten. If `overwrite` is 0 and a file with this name already exists, then Smoldyn asks the user if it should be overwritten. If the user replies no, then this function ends with an error of `ECerror`.

AddCommand

C: `enum ErrorCode smolAddCommand(simptr sim, char type, double on, double off, double step, double multiplier, char *commandstring)`

Python: `ErrorCode addCommand(string type, float on, float off, float step, float multiplier, string commandstring)`

Adds a run-time command to the simulation, including its timing instructions. This function should generally be called after `smolSetSimTimes` to make sure that command times get set correctly. The following table lists the command type options along with the other parameters that are used for each

type. Parameters that are not required are simply ignored. The `commandstring` is the command name followed by any command parameters.

type	meaning	on	off	step	multiplier
Continuous time queue					
b	before simulation	-	-	-	-
a	after simulation	-	-	-	-
@	at fixed time	time	-	-	-
i	fixed intervals	time on	time off	time step	-
x	exponential intervals	time on	time off	min. time step	multiplier
Integer time queue					
B	before simulation	-	-	-	-
A	after simulation	-	-	-	-
&	at fixed iteration	iteration	-	-	-
I	fixed iteration intervals	iter. on	iter. off	iter. step	-
E	every time step	-	-	-	-
N	every n'th time step	-	-	iter. step	-

AddCommandFromString

C: `enum ErrorCode smolAddCommandFromString(simptr sim, char *string)`

Python: `ErrorCode addCommandFromString(str string)`

Defines a runtime command, including its execution timing parameters, from the string `string`. This string should be identical to ones used in configuration files, except that they do not include the “cmd” statement.

7.9 Molecules

AddSpecies

C: `enum ErrorCode smolAddSpecies(simptr sim, char *species, char *molllist)`

Python: `ErrorCode addSpecies(str species, str molllist)`

Python: `Species name: str, state: str = "soln", color: T.Color = "", difc: float = 0.0, display_size: int = 2, molllist: str = ""`

Adds a molecular species named `species` to the system. If you have already created species lists and want all states of this species to live in a specific list, then enter it in `molllist`; otherwise, enter `molllist` as `NULL` or an empty string to request default behavior.

GetSpeciesIndex

C: `int smolGetSpeciesIndex(simptr sim, char *species)`

C: `int smolGetSpeciesIndexNT(simptr sim, char *species)`

Python: `int getSpeciesIndex(str species)`

Returns the species index that corresponds to the species named `species`. Upon failure, this function returns an error code cast as an integer. The “NT” version is identical, but doesn’t throw exceptions or print errors to stderr.

GetSpeciesName

C: `char* smolGetSpeciesName(simptr sim, int speciesindex, char *species)`

Python: `str getSpeciesName(int speciesindex, str species)`

Returns the species name that corresponds to the species index in `speciesindex`. The name is returned both in `species` and directly, where the latter simplifies function use. Upon failure, this function returns `NULL`.

SetSpeciesMobility

C: `enum ErrorCode smolSetSpeciesMobility(simptr sim, char *species, enum MolecState state, double difc, double *drift, double *difmatrix)`

Python: `ErrorCode setSpeciesMobility(str species, MolecState state, float difc, List[float] drift, List[float] difmatrix)`

Python: `species.difc`

Sets any or all of the mobility coefficients for species `species` (which may be “all”) and state `state` (which may be `MSall`). `difc` is the isotropic diffusion coefficient, `drift` is the drift vector, and `difmatrix` is the square of the anisotropic diffusion matrix (see the User’s manual). To not set coefficients, enter a negative number in `difc` and/or enter a `NULL` pointer in the other inputs, respectively.

The last version shown can also be used to get the diffusion coefficient for a species.

AddMolList

C: `int smolAddMolList(simptr sim, char *mollist)`

Python: `int addMolList(str mollist)`

Adds a new molecule list, named `mollist`, to the system.

GetMolListIndex

C: `int smolGetMolListIndex(simptr sim, char *mollist)`

C: `int smolGetMolListIndexNT(simptr sim, char *mollist)`

Python: `int getMolListIndex(str mollist)`

Returns the list index that corresponds to the list named `mollist`. The “NT” version is identical but doesn’t throw exceptions or print errors to the `stderr` output.

GetMolListName

C: `char* smolGetMolListName(simptr sim, int mollistindex, char *mollist)`

Python: `str getMolListName(int mollistindex, str mollist)`

Returns the molecule list name that corresponds to the molecule list with index `mollistindex`. The name is returned both in `mollist` and directly. On error, this function `NULL`.

SetMolList

C: `enum ErrorCode smolSetMolList(simptr sim, char *species, enum MolecState state, char *mollist)`

Python: `ErrorCode setMolList(str species, MolecState state, str mollist)`

Python: `species.mol_list`

Sets the molecule list for species `species` (which may be “all”) and state `state` (which may be `MSall`) to molecule list `mollist`.

The last version can either set or retrieve the molecule list.

SetMaxMolecules

C: `smolSetMaxMolecules(simptr sim, int maxmolecules)`

Python: `setMaxMolecules(int maxmolecules)`

Sets the maximum number of molecules that can simultaneously exist in a system to `maxmolecules`. At present, this function needs to be called for a simulation to run, although it will become optional once dynamic molecule memory allocation has been written.

AddSolutionMolecules

C: `enum ErrorCode smolAddSolutionMolecules(simptr sim, char *species, int number, double *lowposition, double *highposition)`

Python: `ErrorCode addSolutionMolecules(str species, int number, List[float] lowposition, List[float] highposition)`

Python: `species.addToSolution(mol: float, highpos: List[float] = [], lowpos: List[float] = [])`

Adds `number` solution state molecules of species `species` to the system. They are randomly distributed within the box that has its opposite corners defined by `lowposition` and `highposition`. Any or all of these coordinates can equal each other to place the molecules along a plane or at a point. Enter `lowposition` and/or `highposition` as `NULL` to indicate that the respective corner is equal to that corner of the entire system volume.

AddCompartmentMolecules

C: `enum ErrorCode smolAddCompartmentMolecules(simptr sim, char *species, int number, char *compartment)`

Python: `ErrorCode addCompartmentMolecules(str species, int number, str compartment)`

Adds `number` solution state molecules of species `species` to the compartment `compartment`. Molecules are randomly distributed within the compartment.

AddSurfaceMolecules

C: `enum ErrorCode smolAddSurfaceMolecules(simptr sim, int speciesindex, enum MolecState state, int number, int surface, enum PanelShape panelshape, int panel, double *position)`

Python: `ErrorCode addSurfaceMolecules(int speciesindex, MolecState state, int number, int surface, PanelShape panelshape, int panel, List[float] position)`

Adds `number` molecules of species `species` and state `state` to surface(s) in the system. It is permissible for `surface` to be “all”, `panelshape` to be `PSall`, and/or `panel` to be “all”. If you want molecules at a specific position, then you need to enter a specific surface, panel shape, and panel, and then enter the position in `position`.

GetMoleculeCount

C: `int smolGetMoleculeCount(simptr sim, char *species, enum MolecState state)`

Python: `int getMoleculeCount(str species, MolecState state)`

Returns the total number of molecules in the system that have species `species` (“all” is permitted) and state `state` (`MSall` is permitted). Any error is returned as the error code cast as an integer.

SetMoleculeColor

C: `enum ErrorCode smolSetMoleculeStyle(simptr sim, const char *species, enum MolecState state, double *color)`

SetMoleculeSize

C: `enum ErrorCode smolSetMoleculeStyle(simptr sim, const char *species, enum MolecState state, double size)`

SetMoleculeStyle

C: `enum ErrorCode smolSetMoleculeStyle(simptr sim, const char *species, enum MolecState state, double size, double *color)`

Python: `ErrorCode setMoleculeStyle(str species, MolecState state, float size, List[float] color)`

Python: `species.setStyle`

Python: `species.color`

Python: `species.size`

Sets the graphical display parameters for molecules of species `species` (“all” is permitted) and state `state` (`MSall` is permitted). Enter `size` with the drawing size (in pixels if graphics method is “opengl” and in simulation system length units for better drawing methods) or with a negative number to not set the size. Enter `color` with the 3-value color vector or with `NULL` to not set the color (or, in Python, a color word).

7.10 Surfaces

Boundaries

Python: `Boundaries(low: List[float], high: List[float], types: List[str] = field(default_factory=lambda: ["r"]), dim: field(init=False) = 0)`

Python: `setBounds()`

This functionality is only available in Python. There is a class called `Boundaries` and a function called `setBounds`. They do basically the same thing.

SetBoundaryType

C: `enum ErrorCode smolSetBoundaryType(simptr sim, int dimension, int highside, char type)`

Python: `ErrorCode setBoundaryType(int dimension, int highside, str type)`

Sets the molecule interaction properties for a system boundary that bounds the `dimension` axis. Enter `dimension` as -1 to indicate all dimensions. Set `highside` to 0 for the lower boundary, to 1 for the upper boundary, and to -1 for both boundaries. The boundary type is entered in `type` as ‘r’ for reflecting, ‘p’ for periodic, ‘a’ for absorbing, or ‘t’ for transmitting. Note that Smoldyn only observes these properties if no surfaces are declared; otherwise all boundaries are transmitting regardless of what’s entered here.

AddSurface

C: `int smolAddSurface(simptr sim, char *surface)`

Python: `int addSurface(str surface)`

Adds a surface called `surface` to the system.

GetSurfaceIndex

C: `int smolGetSurfaceIndex(simptr sim, char *surface)`

C: `int smolGetSurfaceIndexNT(simptr sim, char *surface)`

Python: `int getSurfaceIndex(str surface)`

Returns the surface index that corresponds to the surface named `surface`. The index is non-negative. On failure, this returns an error code cast as an integer. The “NT” version is identical but errors aren’t printed to the stderr output and don’t throw exceptions.

GetSurfaceName

C: `char* smolGetSurfaceName(simptr sim, int surfaceindex, char *surface)`

Python: `str getSurfaceName(int surfaceindex, str surface)`

Returns the surface name for surface number `surfaceindex` both directly and in the `surface` string. On failure, this returns NULL.

SetSurfaceAction

C: `enum ErrorCode smolSetSurfaceAction(simptr sim, char *surface, enum PanelFace face, char *species, enum MolecState state, enum SrfAction action, char *newspecies)`

Python: `ErrorCode setSurfaceAction(str surface, PanelFace face, str species, MolecState state, SrfAction action)`

Python: `surface.addAction(face, species: Union[Species, str], action: str, new_spec=None)`

Sets the action that should happen when a molecule of species `species` (may be “all”) and state `state` (may be MSall) diffuses into face `face` (may be PFboth) of surface `surface`. The action is set to `action`. Enter `newspecies` to the name of a new species if the molecule should change species, or as either NULL or an empty string if it should not change species.

SetSurfaceRate

C: `enum ErrorCode smolSetSurfaceRate(simptr sim, char *surface, char *species, enum MolecState state, enum MolecState state1, enum MolecState state2, double rate, char *newspecies, int isinternal)`

Python: `ErrorCode setSurfaceRate(str surface, str species, MolecState state, MolecState state1, MolecState state2, float rate, str newspecies, int isinternal)`

Sets the surface interaction rate(s) for surface `surface` (may be “all”) and species `species` (may be “all”) and state `state`. The transition being considered is from `state1` to `state2` (this function uses the tri-state format for describing surface interactions, shown below). The interaction rate is set to `rate`, which is interpreted as a probability value for internal use if `isinternal` is 1 and as a physical interaction coefficient if `isinternal` is 0. If the molecule ends up interacting with the surface, it changes to new species `newspecies`. Enter `newspecies` as either NULL or an empty string to indicate that molecules should not change species upon interactions. The molecule states are most easily understood with the following table. If the action listed in the table is in *italics*, then the corresponding combination of states is not a permitted input.

interaction class	tristate format			action
	state	state1	state2	
collision from solution state	soln	soln	soln	<i>reflect</i>
	"	"	bsoln	transmit
	"	"	bound	adsorb
	"	bsoln	soln	transmit
	"	"	bsoln	<i>reflect</i>
action from bound state	"	"	bound	adsorb
	"	bound	soln	desorb
	"	"	bsoln	desorb
	"	"	bound	<i>no change</i>
collision from bound state	"	"	bound'	flip
	bound	soln	soln	<i>reflect</i>
	"	"	bsoln	transmit
	"	"	bound	hop
	"	"	bound'	hop
	"	bsoln	soln	transmit
	"	"	bsoln	<i>reflect</i>
	"	"	bound	hop
action from bound state	"	"	bound'	hop
	"	bound	soln	desorb
	"	"	bsoln	desorb
	"	"	bound	<i>no change</i>
impossible	"	"	bound'	flip
	"	bound'	any	<i>nonsense</i>

AddPanel

C: `int smolAddPanel(simptr sim, char *surface, enum PanelShape panelshape, char *panel, char *axisstring, double *params)`

Python: `int addPanel(str surface, PanelShape panelshape, str panel, str axisstring, List[float] params)`

Adds or modifies a panel of shape `panelshape` of surface `surface`. `axisstring` lists any text parameters for the panel, which in practice is only a single word that gives the orientation of a rectangle panel (e.g. "+0" or "-y"). `params` lists the numerical parameters for the panel location, size, and drawing characteristics. These are exactly the same parameters that are listed for the "panel" statement in Smoldyn configuration files, with the sole exception that the first rectangle "parameter" is actually a string that is entered in `axisstring`. `panelname` is an optional parameter for naming the panel; if it is included and is not an empty string, the panel is named `panelname`. If this panel name was already used by a panel of the same shape, then this function overwrites that panel's data with the new data. If the name was already used by a panel with a different shape, then this creates an error, and if the name was not used before, then a new panel is created. To use default panel naming, send in `panelname` as either NULL or as an empty string. In the latter case, `panelname` is returned with the newly assigned default name.

In Python, each panel shape is a separate class. These classes are:

- Rectangle: corner: List[float], dimensions: List[float], axis: str, name=""
- Triangle: vertices: List[List[float]] = [], name=""
- Sphere: center: List[float], radius: float, slices: int, stacks: int, name=""
- Hemisphere: center: List[float], radius: float, vector: List[float], slices: int, stacks: int, name: str = ""
- Cylinder: start: List[float], end: List[float], radius: float, slices: int, stacks: int, name=""
- Disk: center: List[float], radius: float, vector: List[float], name=""

GetPanelIndex

C: `int smolGetPanelIndex(simptr sim, char *surface, enum PanelShape *panelshapeptr, char *panel)`

C: `int smolGetPanelIndexNT(simptr sim, char *surface, enum PanelShape *panelshapeptr, char *panel)`

Python: `int getPanelIndex(str surface, PanelShape *panelshapeptr, str panel)`

Returns the panel index for the panel called `panel` on surface `surface`. If `panelshapeptr` is not NULL, this also returns the panel shape in `panelshapeptr`. On failure, this returns the error code cast as an integer. The “NT” version is identical but errors aren’t printed to the stderr output and don’t cause exceptions to be thrown.

GetPanelName

C: `char* smolGetPanelName(simptr sim, char *surface, enum PanelShape panelshape, int panelindex, char *panel)`

Python: `str getPanelName(str surface, PanelShape panelshape, int panelindex, str panel)`

Returns the name of the panel that is in surface `surface`, has shape `panelshape`, and has index `panelindex`, both directly and in the string `panel`. On failure, this returns NULL.

SetPanelJump

C: `enum ErrorCode smolSetPanelJump(simptr sim, const char *surface, const char *panel1, enum PanelFace face1, const char *panel2, enum PanelFace face2, int isbidirectional)`

Python: `ErrorCode setPanelJump(str surface, str panel1, PanelFace face1, str panel2, PanelFace face2, int isbidirectional)`

Sets a jumping link between face `face1` of panel `panel1` and face `face2` of panel `panel2` of surface `surface`. The link goes from `panel1` to `panel2` if `bidirectional` is entered as 0 and goes in both directions if `bidirectional` is entered as 1. None of the surface, panel, or face entries is allowed to be “all”. This does not set the actions of any species to “jump”, which has to be done using the `smolSetSurfaceAction` function.

AddSurfaceUnboundedEmitter

C: `enum ErrorCode smolAddSurfaceUnboundedEmitter(simptr sim, const char *surface, enum PanelFace face, const char *species, double emitamount, double *emitposition)`

Python: `ErrorCode addSurfaceUnboundedEmitter(str surface, PanelFace face, str species, float emitamount, List[float] emitposition)`

Adds information about a point molecular source so that face `face` of surface `surface` can have its absorption properties calculated so that the molecular concentrations will become the same as they would be if the surface weren’t there at all. The point molecular source emits molecules of species `species`, with a rate of `emitamount` and is at location `emitposition`. The emission rate does not need to be in absolute units, but only has to be correct relative to other unbounded emitters. None of the inputs to this function are allowed to be “all”.

SetSurfaceSimParams

C: `enum ErrorCode smolSetSurfaceSimParams(simptr sim, const char *parameter, double value)`

Python: `ErrorCode setSurfaceSimParams(str parameter, float value)`

Sets the surface simulation parameter named with `parameter` to value `value`. The possible parameters are “epsilon”, “margin”, and “neighbordist”. In all cases, the defaults are nearly always good, although this function allows them to be modified if desired. Epsilon is the maximum distance away from a surface that Smoldyn is allowed to place a surface-bound molecule. Margin is the distance inside from the edge of a surface panel that Smoldyn will place surface-bound molecules that hop onto this panel. Neighbor distance is the maximum distance over which surface-bound molecules are allowed to hop to transition from one panel to a neighboring panel.

AddPanelNeighbor

C: `enum ErrorCode smolAddPanelNeighbor(simptr sim, const char *surface1, const char *panel1, const char *surface2, const char *panel2, int reciprocal)`

Python: `ErrorCode addPanelNeighbor(str surface1, str panel1, str surface2, str panel2, int reciprocal)`

Adds panel `panel2` of surface `surface2` as a neighbor of panel `panel1` or surface `surface1`, meaning that surface-bound molecules will be allowed to diffuse from `panel1` to `panel2`. These are not allowed to be the same panel. Also, “all” values are not permitted. Otherwise, essentially any possible entries are legitimate. If surface-bound molecules should also be allowed to diffuse from `panel2` to `panel1`, enter `reciprocal` as 1; if not, enter `reciprocal` as 0.

SetSurfaceStyle

C: `enum ErrorCode smolSetSurfaceStyle(simptr sim, const char *surface, enum PanelFace face, enum DrawMode mode, double thickness, double *color, int stipplefactor, int stipplepattern, double shininess)`

Python: `ErrorCode setSurfaceStyle(str surface, PanelFace face, DrawMode mode, float thickness, List[float] color, int stipplefactor, int stipplepattern, float shininess)`

Python: `surface.setStyle(face, drawmode: str, color: T.Color = "", thickness: float = 1, stipplefactor: int = -1, stipplepattern: int = -1, shininess: int = -1,)`

Sets the graphics output style for face `face` of surface `surface`. `mode` is the drawing mode; enter it as `DMnone` to not set this parameter and otherwise enter it as `DMno` to not draw the surface, `DMvert` for vertices, `DMedge` for edges, or `DMface` for faces. The `thickness` parameter gives the point size or line width for drawing vertices or edges, or can be entered as a negative number to not set this parameter. `color` is the 4-value color vector for the surface, or can be entered as `NULL` to not set this parameter (or, in Python, a color word). `stipplefactor` is the repeat distance for the entire edge stippling pattern, or can be entered as a negative number to not set it. `stipplepattern` is the edge stippling pattern, which needs to be between 0 and 0xFFFF, or can be entered as -1 to not set this parameter. And `shininess` is the surface shininess, for use with lighting in the “opengl.better” graphics display option, or can be entered as -1 to not set this parameter. The parameters `thickness`, `stipplefactor`, and `stipplepattern` only apply to edge style drawing modes and ignore any input in the `face` entry. The `shininess` parameter only applies to the face style drawing modes.

7.11 Compartments

AddCompartment

C: `int smolAddCompartment(simptr sim, char *compartment)`

Python: `int addCompartment(str compartment)`

Adds a compartment called `compartment` to the system.

GetCompartmentIndex

C: `int smolGetCompartmentIndex(simptr sim, char *compartment)`

C: `int smolGetCompartmentIndexNT(simptr sim, char *compartment)`

Python: `int getCompartmentIndex(str compartment)`

Returns the index of the compartment named `compartment`. On failure, this returns an error code cast as an integer. The “NT” version is identical but errors aren’t printed to the `stderr` output or cause exceptions to be thrown.

GetCompartmentName

C: `char* smolGetCompartmentName(simptr sim, int compartmentindex, char *compartment)`

Python: `str getCompartmentName(int compartmentindex, str compartment)`

Returns the name of the compartment that has index `compartmentindex` both directly and in the string `compartment`. Returns `NULL` if an error arises.

AddCompartmentSurface

C: `enum ErrorCode smolAddCompartmentSurface(simptr sim, char *compartment, char`

`*surface)`
 Python: `ErrorCode addCompartmentSurface(str compartment, str surface)`
 Adds surface `surface` as one of the bounding surfaces of compartment `compartment`.

AddCompartmentPoint

C: `enum ErrorCode smolAddCompartmentPoint(simptr sim, char *compartment, double *point)`
 Python: `ErrorCode addCompartmentPoint(str compartment, List[float] point)`
 Adds `point` as one of the interior-defining points of compartment `compartment`.

AddCompartmentLogic

C: `enum ErrorCode smolAddCompartmentLogic(simptr sim, char *compartment, enum CmptrLogic logic, char *compartment2)`
 Python: `ErrorCode addCompartmentLogic(str compartment, CmptrLogic logic, str compartment2)`
 Modifies the current definition of compartment `compartment` using a logical rule specified in `logic` and the definition of `compartment2`.

7.12 Reactions

AddReaction

C: `enum ErrorCode smolAddReaction(simptr sim, const char *reaction, const char *reactant1, enum MolecState rstate1, const char *reactant2, enum MolecState rstate2, int nproduct, const char **productspecies, enum MolecState *productstates, double rate)`
 Python: `ErrorCode addReaction(str reaction, str reactant1, MolecState rstate1, str reactant2, MolecState rstate2, int nproduct, List[str] productspecies, List[MolecState] productstates, float rate)`
 Python: `Reaction(subs: List[Species], prds: List[Species], kf, kb=0.0)`
 Adds reaction named `reaction` to the system. This reaction can have up to two reactants, whose species are listed in `reactant1` and `reactant2` and whose states are listed in `rstate1` and `rstate2`. If the reaction has fewer than two reactants, set either or both of `reactant1` and `reactant2` to either `NULL` or an empty string. State the number of reaction products in `nproduct`, list their species in `productspecies`, and list their states in `productstates`. To set the reaction rate, enter it in `rate`; otherwise, enter `rate` as a negative number.

GetReactionIndex C: `int smolGetReactionIndex(simptr sim, int *orderptr, char *reaction)`

C: `int smolGetReactionIndexNT(simptr sim, int *orderptr, char *reaction)`

Python: `int getReactionIndex(List[int] orderptr, str reaction)`

Returns the index and order for the reaction that is named `reaction`. If the order is known, send in `orderptr` pointing to this value. If it is not known, send in `orderptr` equal to either `NULL` or pointing to a negative number; in this case, it will be returned pointing to the reaction order, if the reaction was found. On failure, this returns the error code, cast as an integer. The “NT” version is identical but errors don’t get displayed to the stderr output or cause exceptions to be thrown.

GetReactionName

C: `char* smolGetReactionName(simptr sim, int order, int reactionindex, char *reaction)`

Python: `str getReactionName(int order, int reactionindex, str reaction)`

Returns the name of the reaction that has reaction order `order` and index `reactionindex` in the string `reaction`. Also returns the result directly. Returns `NULL` if an error arises.

SetReactionRate

C: `enum ErrorCode smolSetReactionRate(simptr sim, int order, char *reaction, double rate, int isinternal)`

Python: `ErrorCode setReactionRate(int order, str reaction, float rate, int isinternal)`

Set the reaction rate to `rate`. If this value is to be interpreted as an internal reaction rate parameter, meaning the production rate for zeroth order reactions, the reaction probability for first order reactions, or the binding radius for second order reactions, then set `isinternal` to 1. Rather than calling this function at all, it's usually easier to use the `rate` parameter of the `smolAddReaction` function, although that doesn't cope with internal rate values.

SetReactionRegion

C: `enum ErrorCode smolSetReactionRegion(simptr sim, const char *reaction, const char *compartment, const char *surface)`

Python: `ErrorCode setReactionRegion(str reaction, str compartment, str surface)`

Limits the spatial region where a reaction can take place to the compartment `compartment` and/or the surface `surface`. To not set one of these limits, enter `compartment` and/or `surface` as NULL. To remove a previously set limit, enter `compartment` and/or `surface` as the empty string, "".

SetReactionIntersurface

C: `enum ErrorCode smolSetReactionIntersurface(simptr sim, const char *reaction, int *rulelist)`

Set the intersurface reaction rules for the bimolecular reaction called `reaction`. Intersurface reactions are reactions between two surface-bound molecules that are on two different surfaces. If `rulelist` is NULL, then this returns the reaction to the default state, which is that intersurface reactions are not allowed for this reaction. Otherwise, `rulelist` should have one entry for each product. If the entry is 1, then that product is placed on the surface with the first reactant; if it is 2, then that product is placed on the surface with the second reactant. If a reaction has no products, then create a single element in `rulelist` equal to 0 to indicate that intersurface reactions are permitted.

SetReactionProducts

C: `enum ErrorCode smolSetReactionProducts(simptr sim, const char *reaction, enum RevParam method, double parameter, const char *product, double *position)`

Python: `ErrorCode setReactionProducts(str reaction, RevParam method, float parameter, str product, List[float] position)`

Sets the reaction product parameters for reaction `reaction`. At a minimum, the `method` reversible parameter is required. Most of these methods require a single parameter, entered in `parameter`. A few methods also require a product, in `product` and the relative position of this product in `position`.

method	parameter	product	position
RPnone	-	-	-
RPirrev	-	-	-
RPconfspread	-	-	-
RPbounce	σ_u	-	-
RPpgem	ϕ	-	-
RPpgemmax	ϕ_{max}	-	-
RPpgemmaxw	ϕ_{max}	-	-
RPratio	σ_u/σ_b	-	-
RPunbindrad	σ_u	-	-
RPpgem2	ϕ	-	-
RPpgemmax2	ϕ_{max}	-	-
RPratio2	σ_u/σ_b	-	-
RPOffset	-	product number	relative position
RPfixed	-	product number	relative position

If `method` is `RPbounce`, then a negative number for the `parameter` indicates default bounce behavior, which is that molecules are separated by an amount that is equal to their previous overlap.

7.13 Ports

AddPort

C: `enum ErrorCode smolAddPort(simptr sim, const char *port, const char *surface, enum PanelFace face)`

Python: `ErrorCode addPort(str port, str surface, PanelFace face)`

Python: `Port(name: str, surface: Union[Surface, str], panel: str)`

Adds a port to the simulation. The port will be named `port` and will port at the `face` face of surface `surface`.

GetPortIndex

C: `int smolGetPortIndex(simptr sim, const char *port)`

C: `int smolGetPortIndexNT(simptr sim, const char *port)`

Python: `int getPortIndex(str port)`

Returns the index of the port named `port`. The “NT” version is identical but errors don’t get displayed to the stderr output or cause exceptions to be thrown.

GetPortName

C: `char* smolGetPortName(simptr sim, int portindex, char *port)`

Python: `str getPortName(int portindex, str port)`

Returns the name of the port with index `portindex`, both directly and in `port`.

AddPortMolecules

C: `enum ErrorCode smolAddPortMolecules(simptr sim, const char *port, int nmolec, const char *species, double **positions)`

Python: `ErrorCode addPortMolecules(str port, int nmolec, str species, List[float] positions)`

Adds `nmolec` molecules to Smoldyn’s import buffer of port `port`. These molecules will all have species `species` and state `MSsoln`. Enter `positions` as NULL to have the molecules positioned randomly over the porting surface and as an `nmolec` length list of position vectors to have them located at those specific initial positions. These initial positions should be close to the porting surface, and on the Smoldyn system side of it.

GetPortMolecules

C: `int smolGetPortMolecules(simptr sim, const char *port, const char *species, enum MolecState state, int remove)`

Python: `int getPortMolecules(str port, str species, MolecState state, int remove)`

Returns the number of molecules that are in Smoldyn’s export buffer of port `port`. Enter `species` with the species of the molecules that should be retrieved, or “all” for all species. Enter `state` with the states of the molecules that should be retrieved, or `MSall` for all states. Enter `remove` with 1 to remove molecules from the export buffer after they are retrieved or with 0 to leave them in the buffer. If an error arises, this returns the error code cast as an integer.

7.14 Lattices

AddLattice

C: `enum ErrorCode smolAddLattice(simptr sim, const char *lattice, const double *min, const double *max, const double *dx, const char *btype)`

Python: `ErrorCode AddLattice(str lattice, List[float] min, List[float] max, List[float] dx, str btype)`

Adds a lattice to the simulation for hybrid operation. The lattice is named `lattice` and extends from `min` to `max`, with lattice spacing `dx`. The boundary types are given with `btype`, which is a string of either ‘r’ characters for reflective boundary or ‘p’ characters for periodic boundary, with one character for each dimension and the dimensions listed in order (e.g. “rrp” is reflective on the *x* and *y* axes and periodic on the *z* axis).

GetLatticeIndex

C: `int smolGetLatticeIndex(simptr sim, const char *lattice)`

C: `int smolGetLatticeIndexNT(simptr sim, const char *lattice)`

Python: `int GetLatticeIndex(str lattice)`

Returns the index of the lattice named `lattice`. The “NT” version is identical but errors don’t get displayed to the stderr output or cause exceptions to be thrown.

GetLatticeName

C: `char *smolGetLatticeName(simptr sim, int latticeindex, char *lattice)`

Python: `str getLatticeName(int latticeindex, str lattice)`

Returns the name of the lattice with index `latticeindex`, both directly and in `lattice`.

AddLatticeMolecules

C: `enum ErrorCode smolAddLatticeMolecules(simptr sim, const char *lattice, const char *species, int number, double *lowposition, double *highposition)`

Python: `ErrorCode AddLatticeMolecules(str lattice, str species, int number, List[float] lowposition, List[float] highposition)`

Adds `number` molecules of species `species` to the lattice named `lattice`, randomly positioned over the volume extending from `lowposition` to `highposition`. These molecules will all have state `MSsoln`.

AddLatticePort

C: `enum ErrorCode smolAddLatticePort(simptr sim, const char *lattice, const char *port)`

Python: `ErrorCode AddLatticePort(str lattice, str port)`

Connects port `port` with lattice `lattice`, so that molecules can transition across this port between the particle-based simulation region and the lattice-based simulation region.

AddLatticeSpecies

C: `enum ErrorCode smolAddLatticeSpecies(simptr sim, const char *lattice, const char *species)`

Python: `ErrorCode AddLatticeSpecies(str lattice, str species)`

Not all particle-based Smoldyn species are necessarily listed in a lattice portion of space, so use this function to add species to the lattice region of space. Clearly, the lattice is `lattice` and the species being added is `species`.

AddLatticeReaction

C: `enum ErrorCode smolAddLatticeReaction(simptr sim, const char *lattice, const char *reaction, const int move)`

Python: `ErrorCode AddLatticeReaction(str lattice, str reaction, int move)`

Not all particle-based Smoldyn reactions are necessarily listed in a lattice portion of space, so use this function to add reactions to the lattice region of space. Clearly, the lattice is `lattice` and the reaction being added is `reaction`.