

## utils

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</home/mpastor/soft/eTAM/src/utils.py>

```
# -*- coding: utf-8 -*-
#
# Description      eTAM utility functions
#
#
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#
# (c) PhI 2013
```

## Modules

[os](#)

[shutil](#)

## Functions

### **lastVersion(verID)**

Returns the path to the directory where the verID version of the model is located

If a value of -1 is provided, analyzes the "version0000" directories and returns the last one

### **nextVersion()**

Returns the path to the directory where the verID version of the model is located

If a value of -1 is provided, analyzes the "version0000" directories and returns the last one

### **removefile(file)**

Removes silently files or whole paths.

It checks the existence of the file in the argument and if it does not exists do not issue any error

### **splitSDF(molecules)**

Splits a SDFfile containing 1 to n molecules

Every molecule is written to a "m00000000001.sdf" file and the full list of names is returned in a list

### **writeError(error)**

Print an error message

## Data

**opt** = '/home/mpastor/soft/eTAM/opt/'

**wkd** = '/home/mpastor/soft/eTAM/src'