

Approach of lazy calculations in application to topological graph indices in Sage. Introduction to a new molecular class.

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The Sage Project

Mission Statement

Create a viable free open source alternative to Magma, Maple, Mathematica, and Matlab

A “viable alternative” will have...

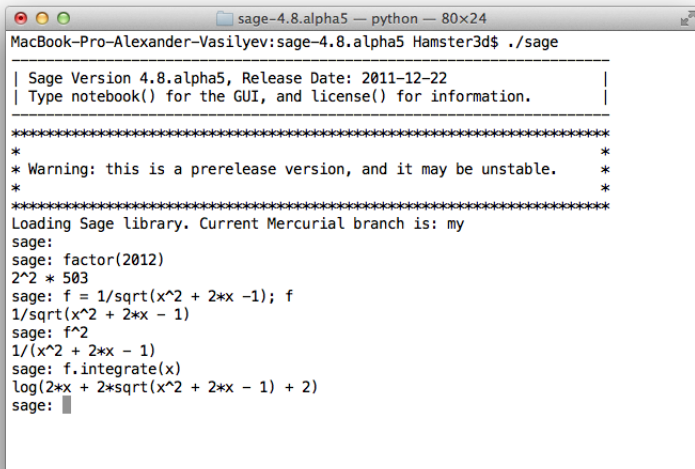
- The mathematical features of Magma, Maple, Mathematica, and Matlab with comparable speed.
- Beautiful interactive 2d and 3d graphics.
- A notebook interface and an IDE.
- Many books and Commercial support (e.g., customized notebook servers)

What is Sage

- ① **A self-contained distribution** of over 90 open source packages that is easy to build from source.
- ② **Interfaces** that smoothly tie together all these libraries and packages.
- ③ **A new library** that implements novel algorithms. About a half million lines of code written by a worldwide community of about 200 people over the last 6 years.

Sage in use

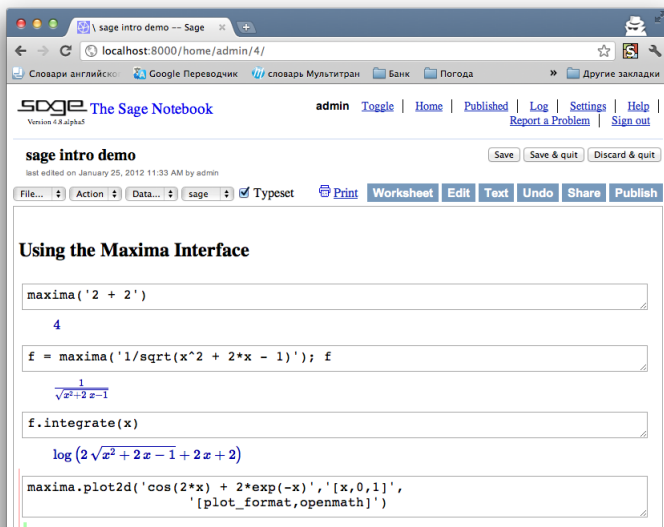
Command Line Sage



```
sage-4.8.alpha5 -- python -- 80x24
MacBook-Pro-Alexander-Vasilyev:sage-4.8.alpha5 Hamster3d$ ./sage
-----
| Sage Version 4.8.alpha5, Release Date: 2011-12-22          |
| Type notebook() for the GUI, and license() for information. |
-----
*****
*
* Warning: this is a prerelease version, and it may be unstable. *
*
*****
Loading Sage library. Current Mercurial branch is: my
sage:
sage: factor(2012)
2^2 * 503
sage: f = 1/sqrt(x^2 + 2*x -1); f
1/sqrt(x^2 + 2*x - 1)
sage: f^2
1/(x^2 + 2*x - 1)
sage: f.integrate(x)
log(2*x + 2*sqrt(x^2 + 2*x - 1) + 2)
sage: █
```

Sage in use

Sage notebook



sage intro demo

last edited on January 25, 2012 11:33 AM by admin

Save Save & quit Discard & quit

File... Action... Data... sage Typeset Print Worksheet Edit Text Undo Share Publish

Using the Maxima Interface

```
maxima('2 + 2')
```

4

```
f = maxima('1/sqrt(x^2 + 2*x - 1)'); f
```

$$\frac{1}{\sqrt{x^2+2x-1}}$$

```
f.integrate(x)
```

$$\log(2\sqrt{x^2+2x-1}+2x+2)$$

```
maxima.plot2d('cos(2*x) + 2*exp(-x)', '[x,0,1]',  
              '[plot_format,openmath]')
```

Try Sage

Sage is crossplatform. Supported platforms:

- Linux
- Apple Mac OS X
- Solaris
- Microsoft Windows (run on virtual machine with Linux)
- Live CD

The best way to try Sage is using on-line Sage notebook.

Links

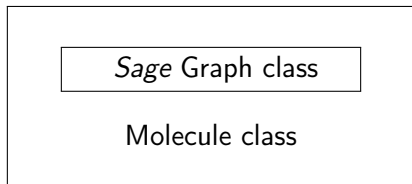
Sage homepage	www.sagemath.org
Sage Notebook	www.sagenb.org
Sage documentation	www.sagemath.org/help.html
Bug report and contribution	trac.sagemath.org/sage_trac/report

Math Chemistry package will include:

Atom-Bond Connectivity Index; Balaban J Index; Balaban-like indices; Degree Distance; Reverse Degree Distance; Detour Index; Hyper-Detour Index; Wiener Index; Hyper-Wiener Indices; Reverse Wiener Index; Modified Wiener index; Variable Wiener index; Terminal Wiener index; Edge Wiener index; y-Wiener index; Reciprocal complimentary Wiener index; Reciprocal reverse Wiener index; Estrada Index; Laplacian Estrada index; Distance Estrada index; Eccentric Connectivity Index; Eccentric Distance Sum; (Schultz) Molecular Topological Index; Reciprocal Molecular Topological Index; Randic Index; (Zeroth-order) General Randic Index; Variable Randic Index; Augmented Zagreb Index; M1, M2 Zagreb index; Path-Zagreb index; Zagreb Coindices; Reformulated Zagreb indices; Edge-Szeged index; Szeged Index; Revised Szeged Index; (Padmakar-Ivan) PI Index; Vertex PI Index; Connectivity Index; Matching number; Number of perfect matchings; 1th,2th,3th Geometric-Arithmetic index; Harary Index; Kirchhoff Index; Hosoya index; Merrifield-Simmons Indices; **Spectrum, spread, spectral moments, characteristic polynomial, Energy** (Adjacency, Laplacian, Detour, Distance, Reciprocal distance, Normalized laplacian, Signless laplacian, Resistance distance, Harary); **Polynomials** (Hosoya, Omega, Eccentric connectivity, Wiener, PI, Szeged, Edge-Szeged); Anti-Kekule number; Anti-Forcing Number;

First attempt: classical object-oriented approach

Molecule class inherited from the Sage Graph class

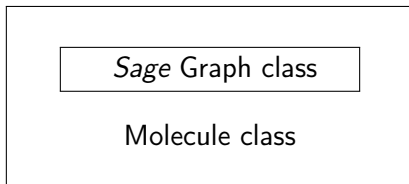


+ Advantages

- Easy to develop
- Inherited methods such as
 - `Graph.show()` → `Molecule.show()`
 - `Graph.spectrum()` → `Molecule.spectrum()`
 - `Graph.min_cut()` → `Molecule.min_cut()`
 - ...
- The package could be included into main Sage repository

First attempt: classical object-oriented approach

Molecule class inherited from the Sage Graph class

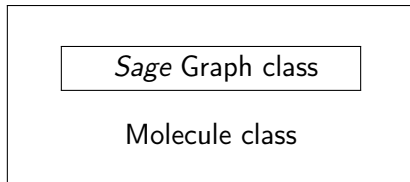


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First attempt: classical object-oriented approach

Molecule class inherited from the Sage Graph class



– Disadvantages

- Performance
- Dependence on the framework

Second attempt: lazy calculations

We calculate data only when it is needed and then we save the results.

Sage Graph class

Mol class

+ Advantages

- Performance
- Independence on the framework (the package can be used in any Python-based project)

– Disadvantages

- Hard to code, easy to get bugs
- The package will not be included into main Sage bundle

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Tests

Graphs for test: all connected simple graphs of 7, 8, 9 vertices

	7c	8c	9c
number of graphs	853	11117	261080

1. Zagreb M1 and M2 indices:

$$M1 = \sum_{v \in V(G)} d_v^2 \quad M2 = \sum_{u \sim v \in V(G)} d_v d_u \quad (1)$$

2. Molecular Topological Index MTI, Degree Distance DD, Reverse Degree Distance rDD:

$$MTI = \sum_{v \in V(G)} (A + D) d_v \quad DD = \sum_{v \in V(G)} d_v \sum_{u \in V(G)} d_{u,v} \quad (2)$$
$$rDD = 2(n - 1)md_G - DD$$

Where d_v is degree of v , A is adjacency matrix, D is distance matrix, $d_{u,v}$ is distance between u and v , d_G is diameter of G , n is number of vertices, m is number of edges

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Tests. First test.

Performance test of *Molecule* and *Mol* classes.

Time (in seconds) of calculation Zagreb M1 and M2 indices for all connected graphs of 7, 8, 9 vertices.

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Molecule	0.48	7.36	206.7
Mol	0.07	1.17	33.0
multiplier	6.85	6.29	6.26

The new *Mol* class is ≈ 6 times faster

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Molecule	0.86	12.72	336.34
Mol	0.75	10.33	263.07
multiplier	1.14	1.23	1.27

The new *Mol* class is ≈ 1.2 times faster

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Easy link to Sage and NetworkX

```
Mol().sage_graph()
```

```
Mol().NX_graph()
```

Thank you!