

pharmit

virtual screening in the cloud with pharmit

David Ryan Koes

 @david_koes

Computational and Systems Biology
University of Pittsburgh



The Cloud



The Cloud



Running your stuff on other
people's computers.

The Cloud



Why Cloud?

To get access to resources.

- compute power
- software
- storage
- data

Why Cloud?

To get access to resources.

- compute power
- software
- storage
- data



Academia is well-suited to serve as a micro-cloud provider of these resources

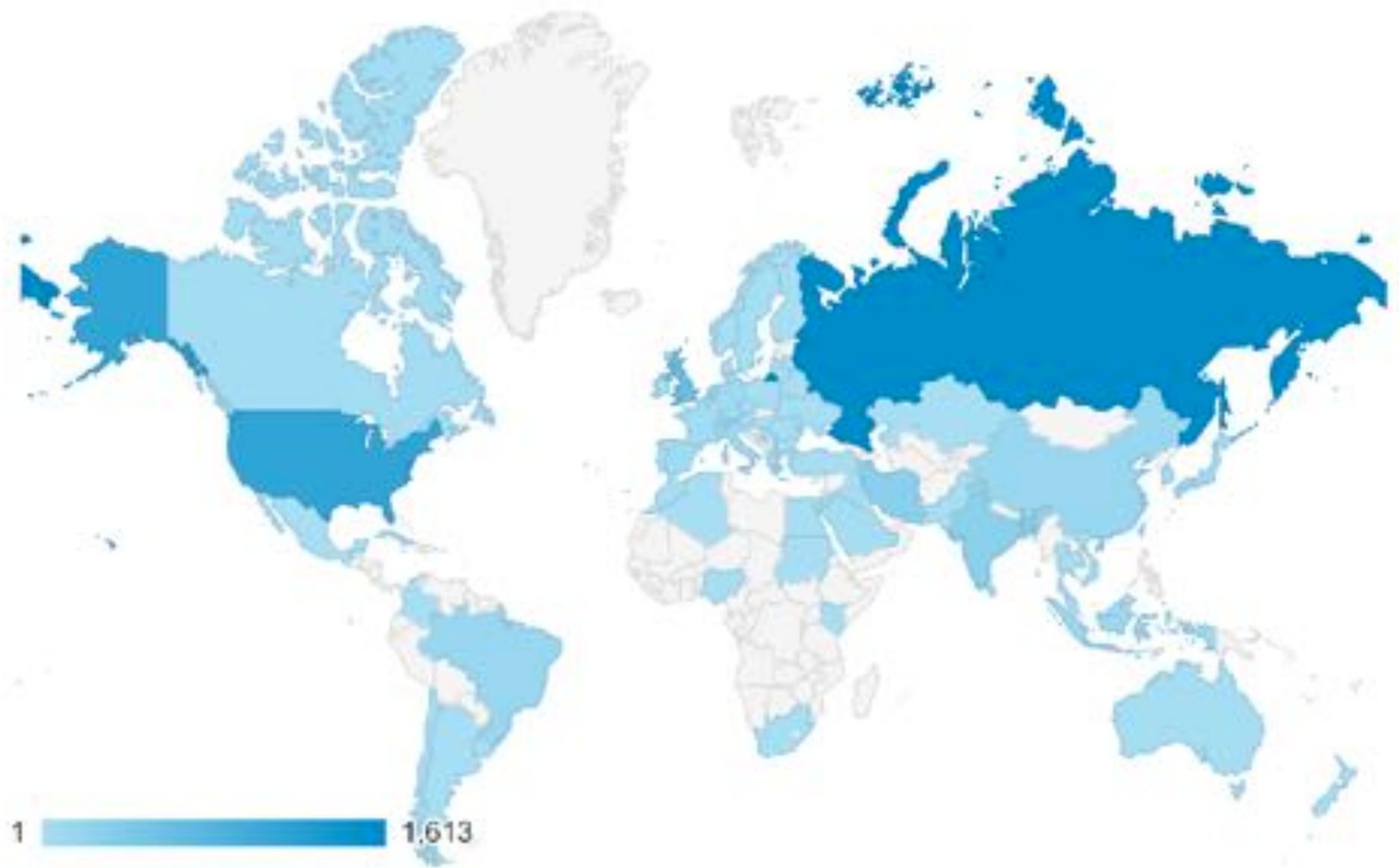


David Koes @david_koes · 23 Sep 2016

Pharmit usage by country. Online open drug discovery tools democratize access to big data resources.

pharmit.csb.pitt.edu #compchem





Country [?]	Acquisition			Behavior		
	Sessions [?]	% New Sessions [?]	New Users [?]	Bounce Rate [?]	Pages / Session [?]	Avg. Session Duration [?] 
	5,604 % of Total: 100.00% (5,604)	52.57% Avg for View: 52.48% (0.17%)	2,946 % of Total: 100.17% (2,941)	28.09% Avg for View: 28.09% (0.00%)	3.08 Avg for View: 3.08 (0.00%)	00:05:07 Avg for View: 00:05:07 (0.00%)
1.  Lithuania	1 (0.02%)	100.00%	1 (0.03%)	0.00%	7.00	00:23:01
2.  Romania	8 (0.14%)	37.50%	3 (0.10%)	25.00%	4.75	00:12:46
3.  Netherlands	156 (2.78%)	21.79%	34 (1.15%)	25.64%	7.53	00:12:43
4.  Bangladesh	77 (1.37%)	28.57%	22 (0.75%)	37.66%	3.04	00:09:52
5.  Iran	287 (5.12%)	31.71%	91 (3.09%)	23.00%	4.59	00:08:51
6.  Indonesia	119 (2.12%)	15.97%	19 (0.64%)	20.17%	3.86	00:08:07
7.  Brazil	158 (2.82%)	41.14%	65 (2.21%)	27.22%	3.85	00:07:48
8.  United Kingdom	623 (11.12%)	46.55%	290 (9.84%)	36.12%	3.91	00:06:53
9.  India	283 (5.05%)	54.42%	154 (5.23%)	41.34%	3.49	00:06:45
10.  United States	1,116 (19.91%)	42.38%	473 (16.06%)	39.61%	3.20	00:06:08

Sessions ▾ VS. Select a metric

Hourly Day Week Month

● Sessions



Sessions

5,604



Users

2,966



Pageviews

17,274



Pages / Session

3.08



Avg. Session Duration

00:05:07



Bounce Rate

28.09%



■ New Visitor ■ Returning Visitor



PharmIT 4.00 - 9/15/2014/10:00:00 AM

Search PubChem

Pharmacophore Search -> Shape Filter

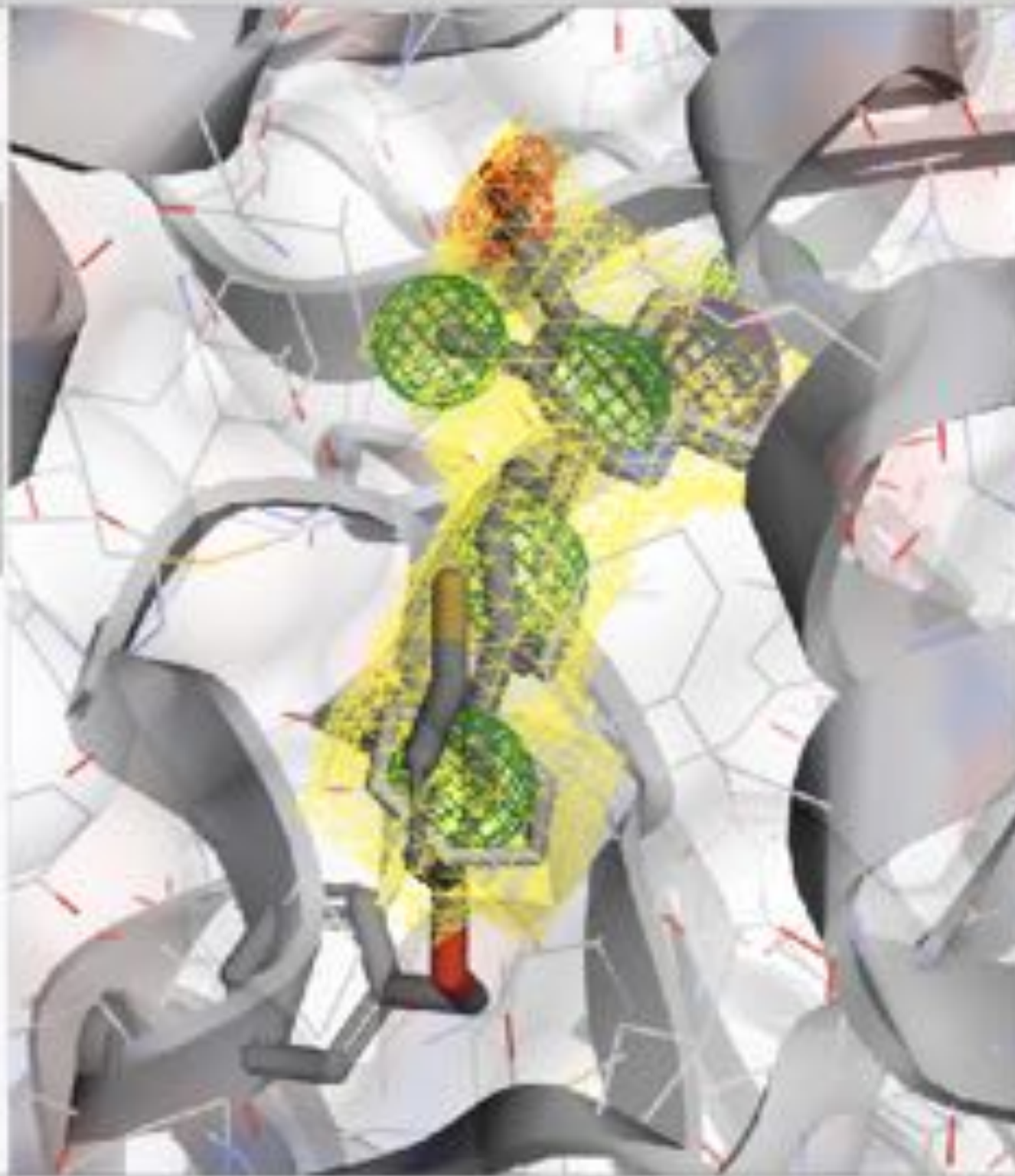
Load Receptor... Load Features...

Pharmacophore

- ☒ **Aromatic**
[0.0, 1.0, 1.0, 1.0] Radius 1.0
- ☒ **HydrogenAcceptor**
[0.0, 1.0, 1.0, 1.0] Radius 1.0
- ☒ **HydrogenAcceptor**
[0.0, 1.0, 1.0, 1.0] Radius 1.0
- ☒ **NegativIon**
[0.0, 1.0, 1.0, 1.0] Radius 1.0
- ☒ **Hydrophobic**
[0.0, 1.0, 1.0, 1.0] Radius 1.0
- ☒ **Hydrophobic**
[0.0, 1.0, 1.0, 1.0] Radius 1.0
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[0.0, 1.0, 1.0, 1.0] Radius 1.0
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- ☐ **Aromatic**
[0.0, 1.0, 1.0, 1.0] Radius 1.0
- ☐ **Aromatic**
[0.0, 1.0, 1.0, 1.0] Radius 1.0

Load Session... Save Session...

Display menu



Pharmacophore Results

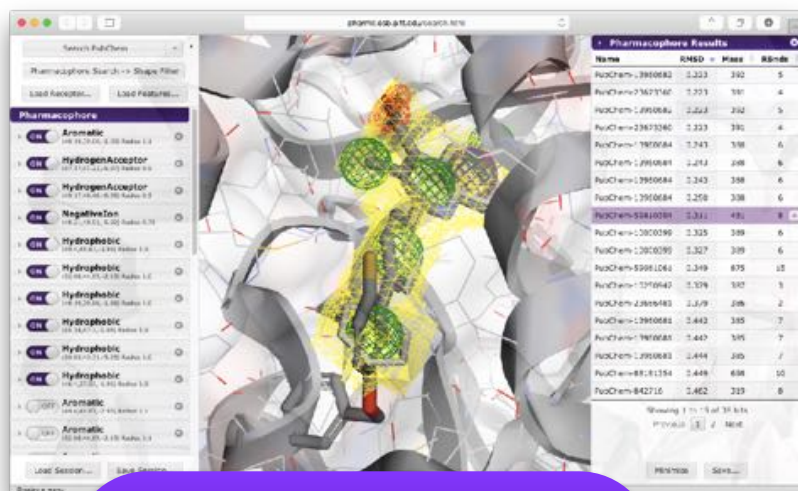
Name	RMSD	Mass	RScore
PubChem-13910082	1.223	392	5
PubChem-73673160	1.223	391	4
PubChem-13910084	1.223	392	5
PubChem-23873260	1.223	390	4
PubChem-19870784	1.243	398	6
PubChem-13910081	1.213	398	6
PubChem-13910084	1.243	398	6
PubChem-13910084	1.258	398	6
PubChem-50810084	1.311	491	8
PubChem-13910090	1.325	399	6
PubChem-13910090	1.327	399	6
PubChem-50810064	1.349	675	15
PubChem-13910040	1.379	397	3
PubChem-23873180	1.379	398	2
PubChem-13910082	1.443	395	7
PubChem-19870781	1.443	395	7
PubChem-13910082	1.444	395	7
PubChem-88111354	1.448	698	10
PubChem-842716	1.462	313	8

Showing 1 to 19 of 19 hits

Previous 1 Next

Print Table Save...

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pharmit.js

jQuery

3Dmol.js

Client

Search

User Libraries

CHEMBL

PubChem

ChemDiv

NCI

MolPort

Prebuilt
Libraries

pharmitserver

OpenBabel

ShapeDB

Pharmer

Minimization

smina

OpenBabel

Server

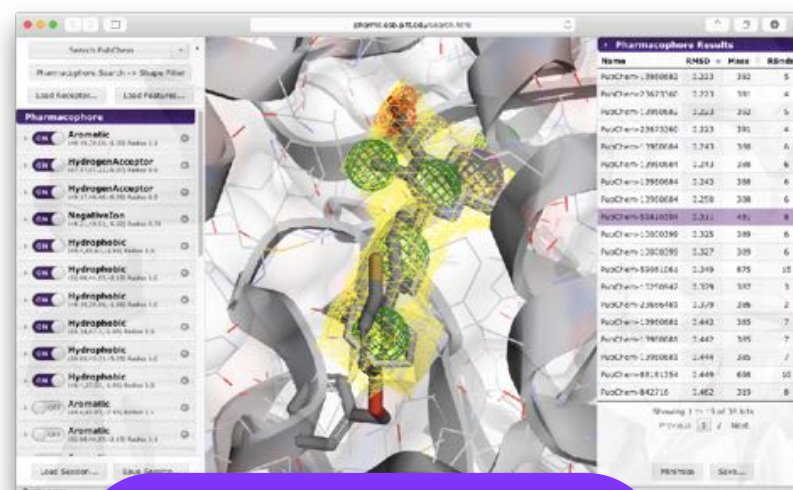
buildlibs.py

RDKit

OpenBabel

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pharmit.js

jQuery

3Dmol.js

Client

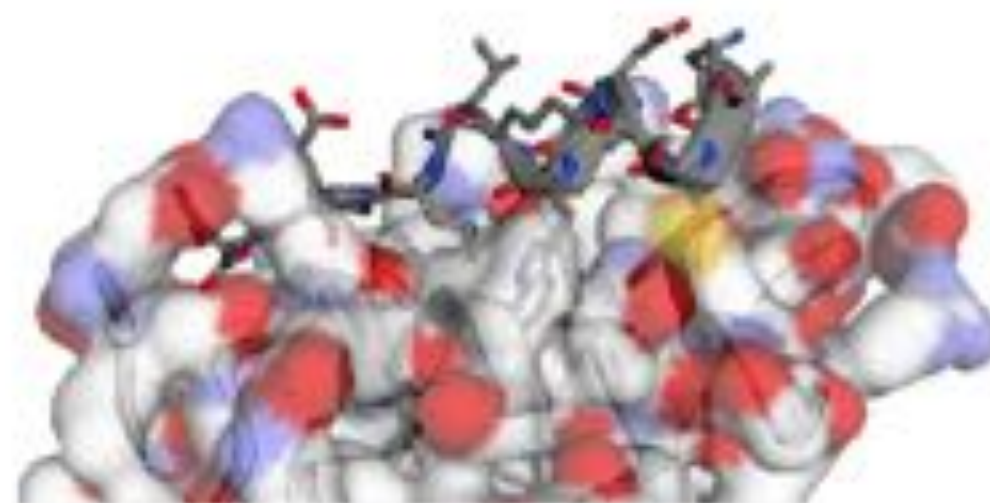
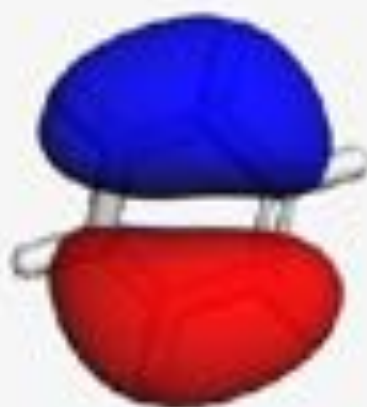
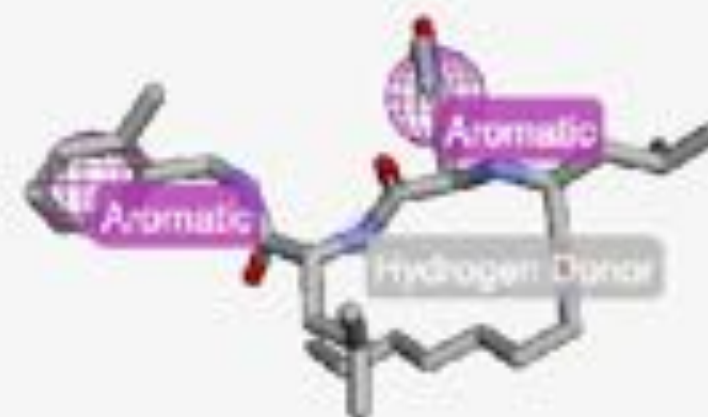
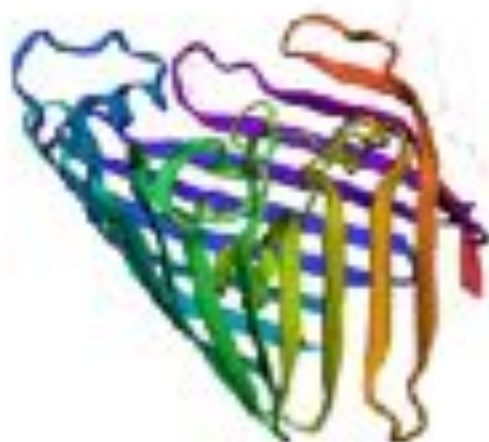
Why web application?

3Dmol.js

View Embed Develop

A modern, object-oriented JavaScript library for visualizing molecular data

Feedback Download



<http://3dmol.csb.pitt.edu>

Rego, N., & Koes, D. (2014). 3Dmol.js: molecular visualization with WebGL.
Bioinformatics. doi:10.1093/bioinformatics/btu829

CHEMBL

ChemDiv

MolPort

NCI

PubChem

CHEMBL22.1 ⓘ

20,828,750 conformers of 1,581,180 molecules
Updated: 2017-Feb-24 12:37:33

ChemDiv ⓘ

21,609,296 conformers of 1,456,338 molecules
Updated: 2017-Feb-23 23:25:40

MolPort ⓘ

94,527,744 conformers of 6,544,814 molecules
Updated: 2017-Feb-26 19:55:13

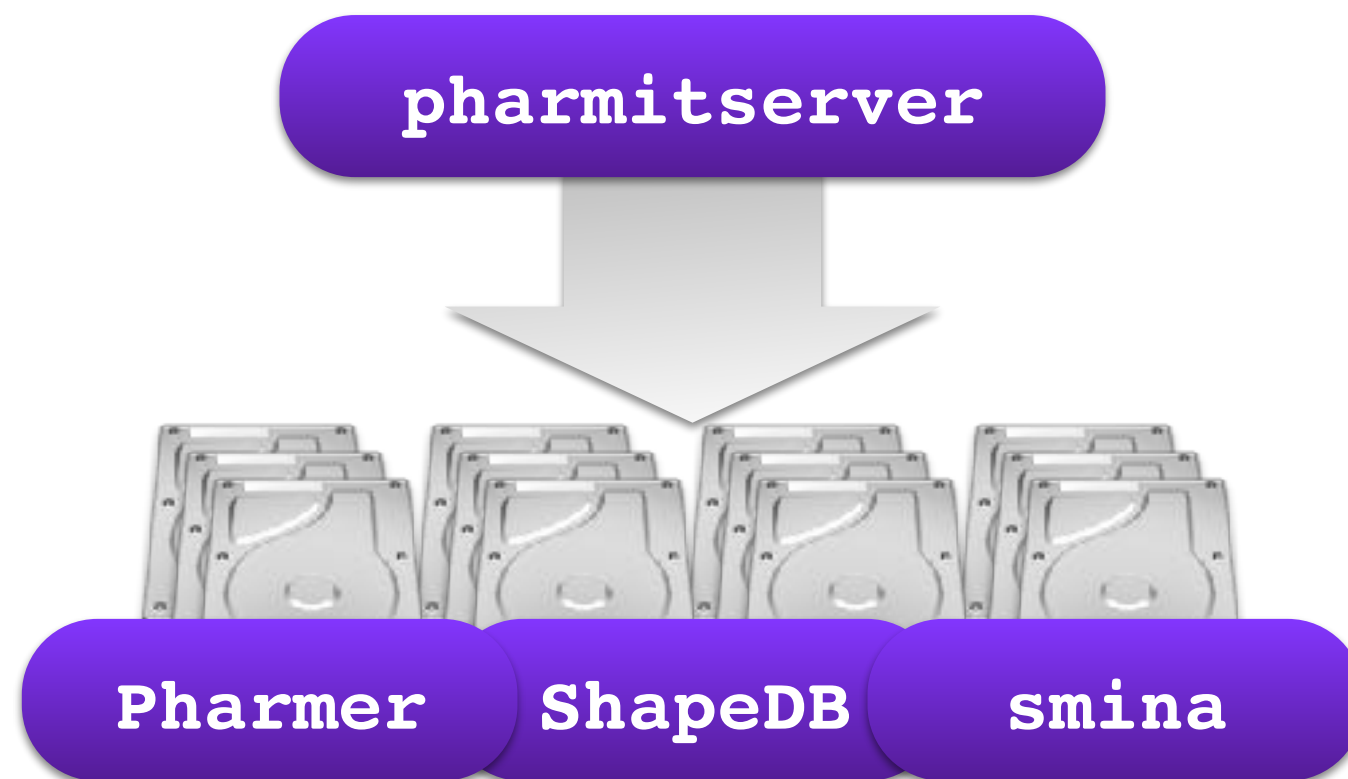
NCI Open Chemical Repository ⓘ

1,590,246 conformers of 150,388 molecules
Updated: 2017-Feb-23 10:37:26

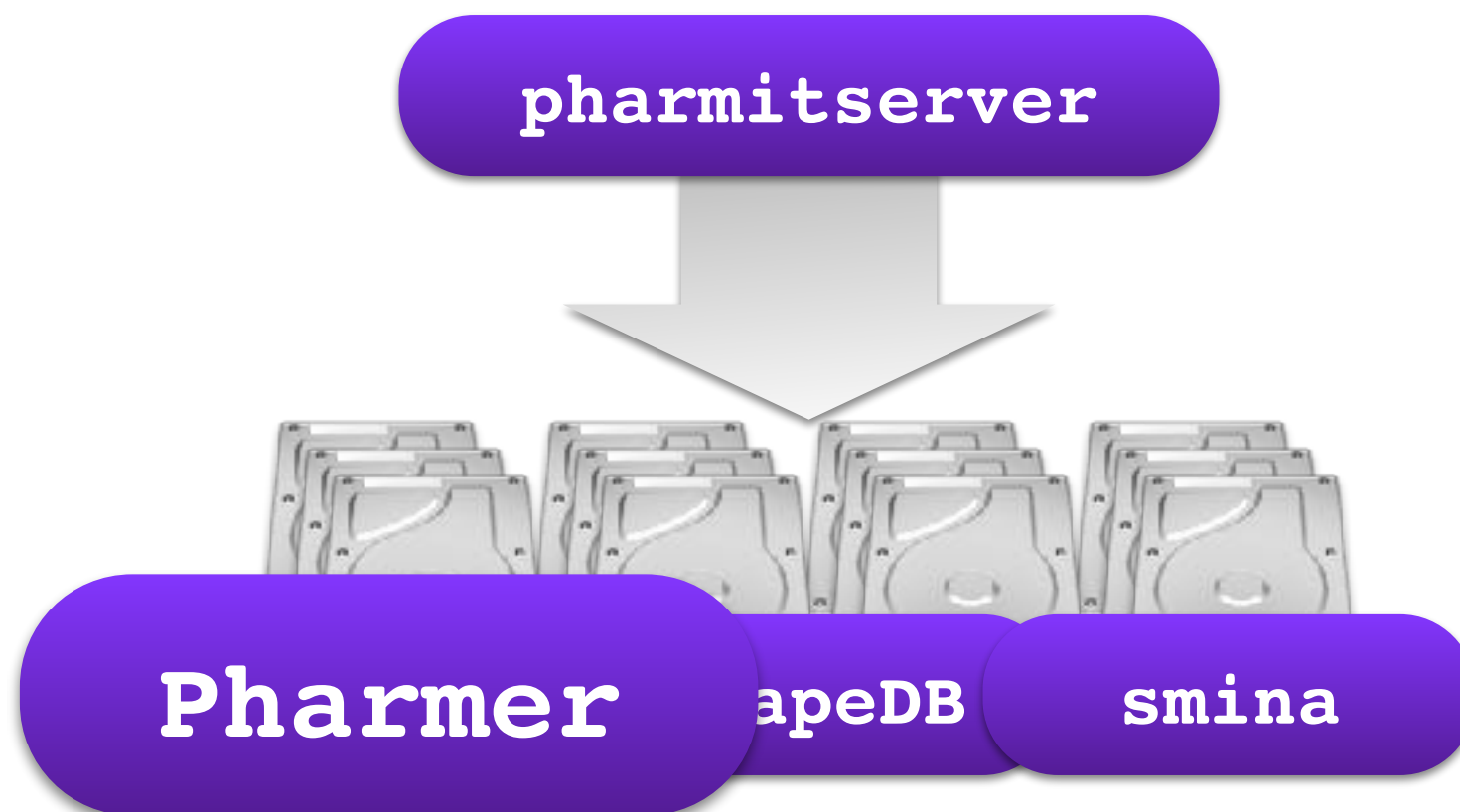
Contributed Libraries



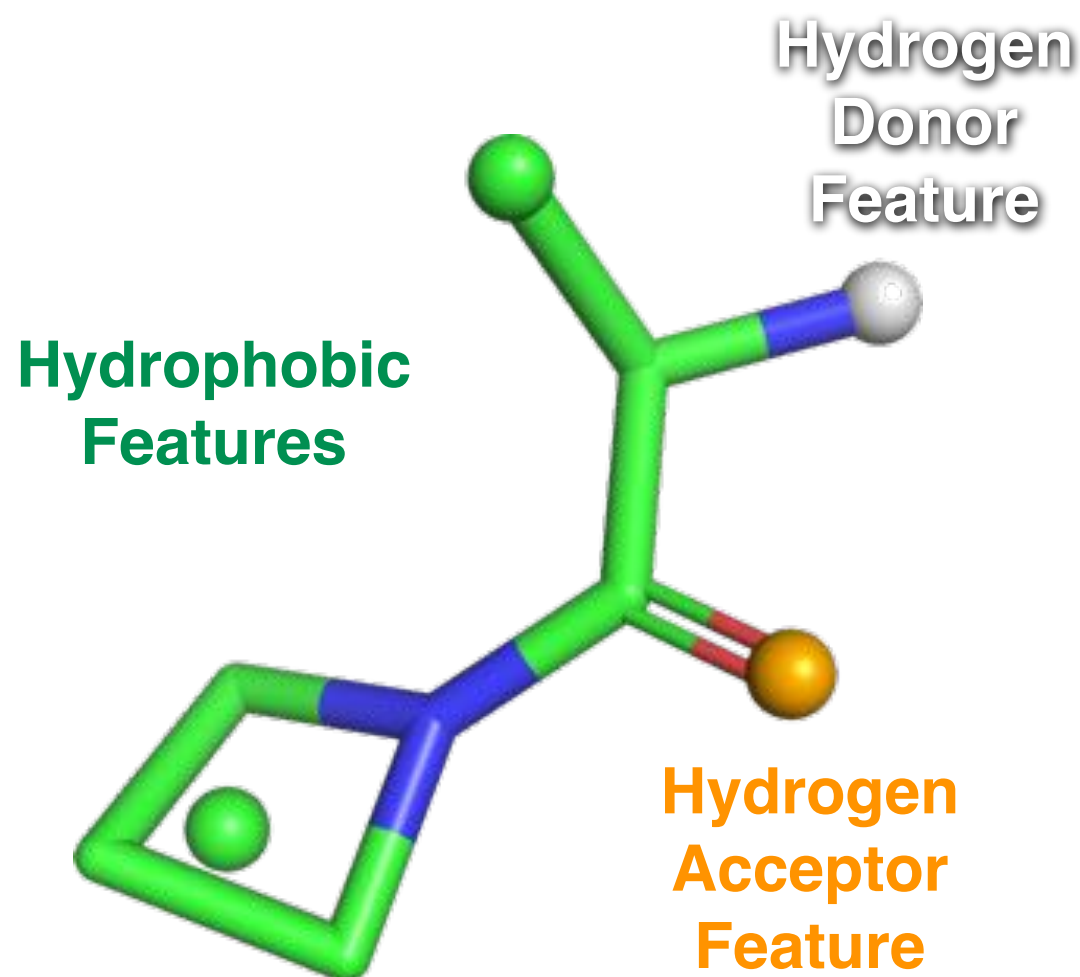
Access Private Library



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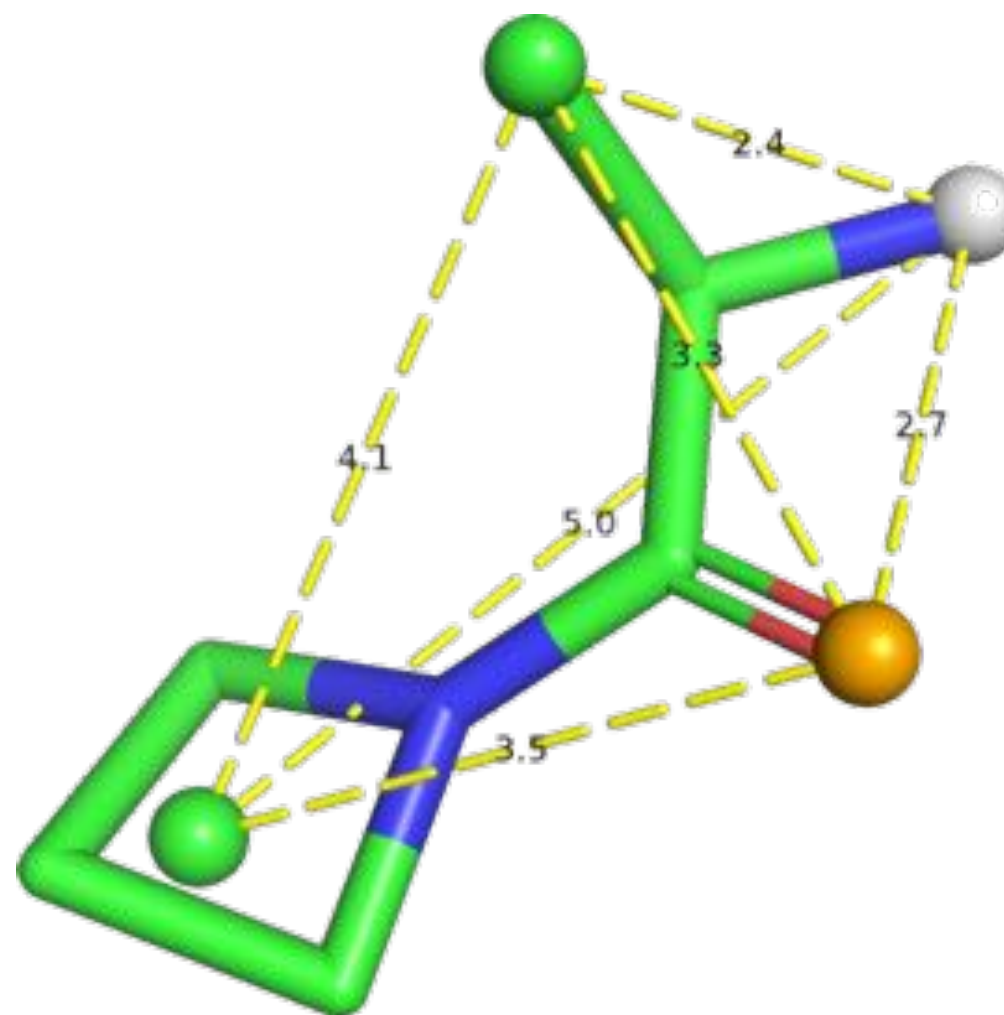
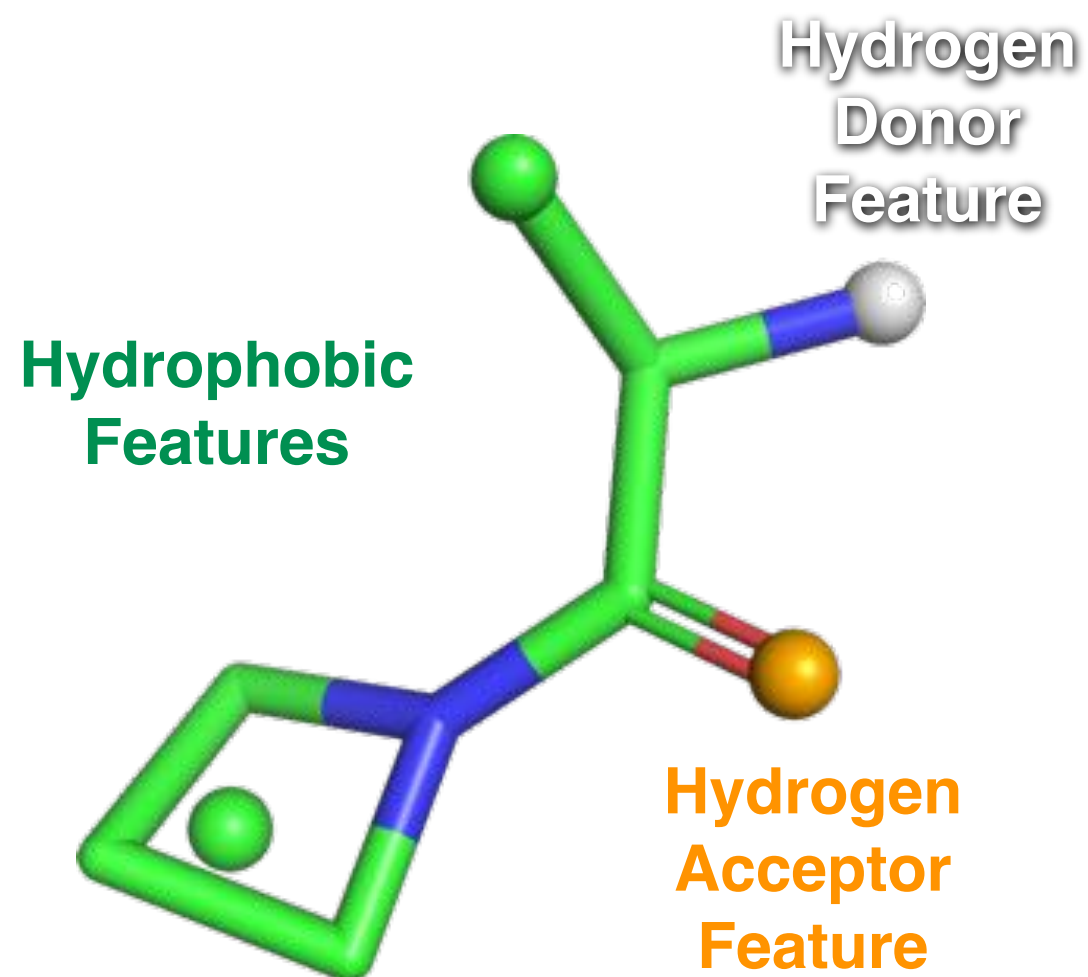


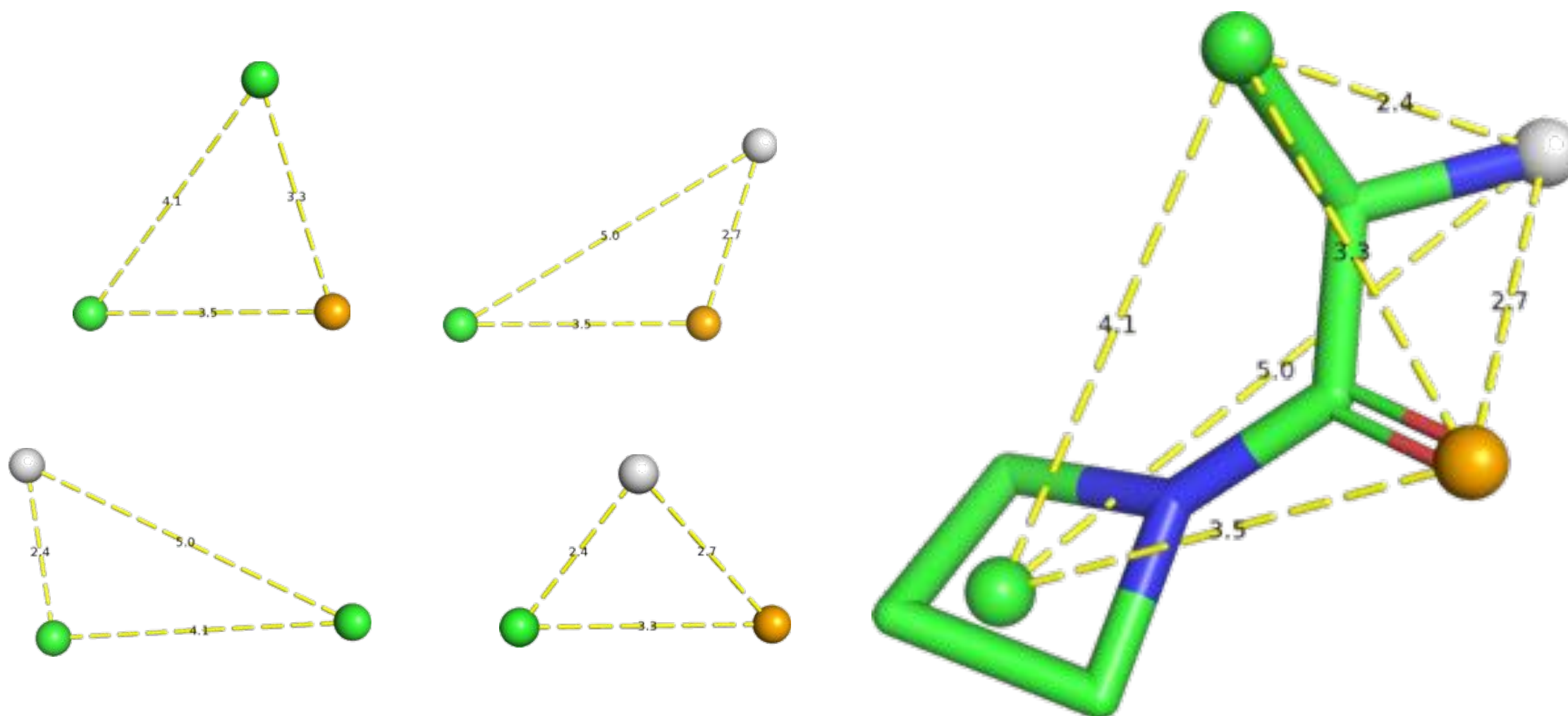
Pharmacophore

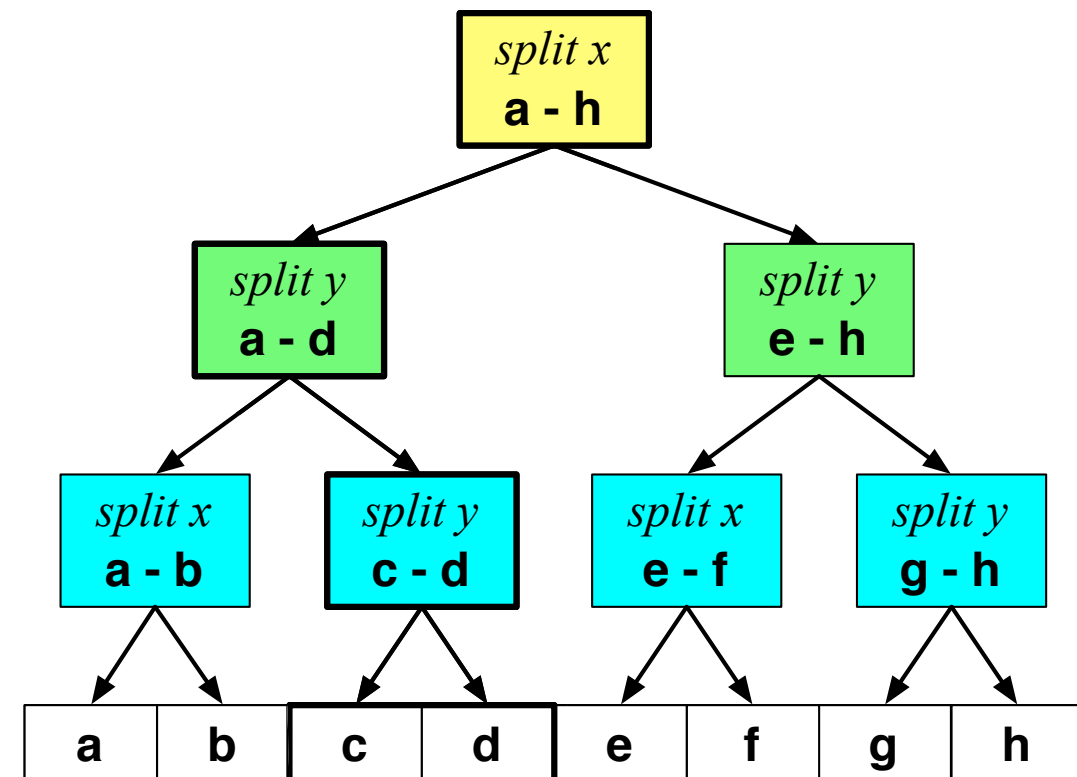
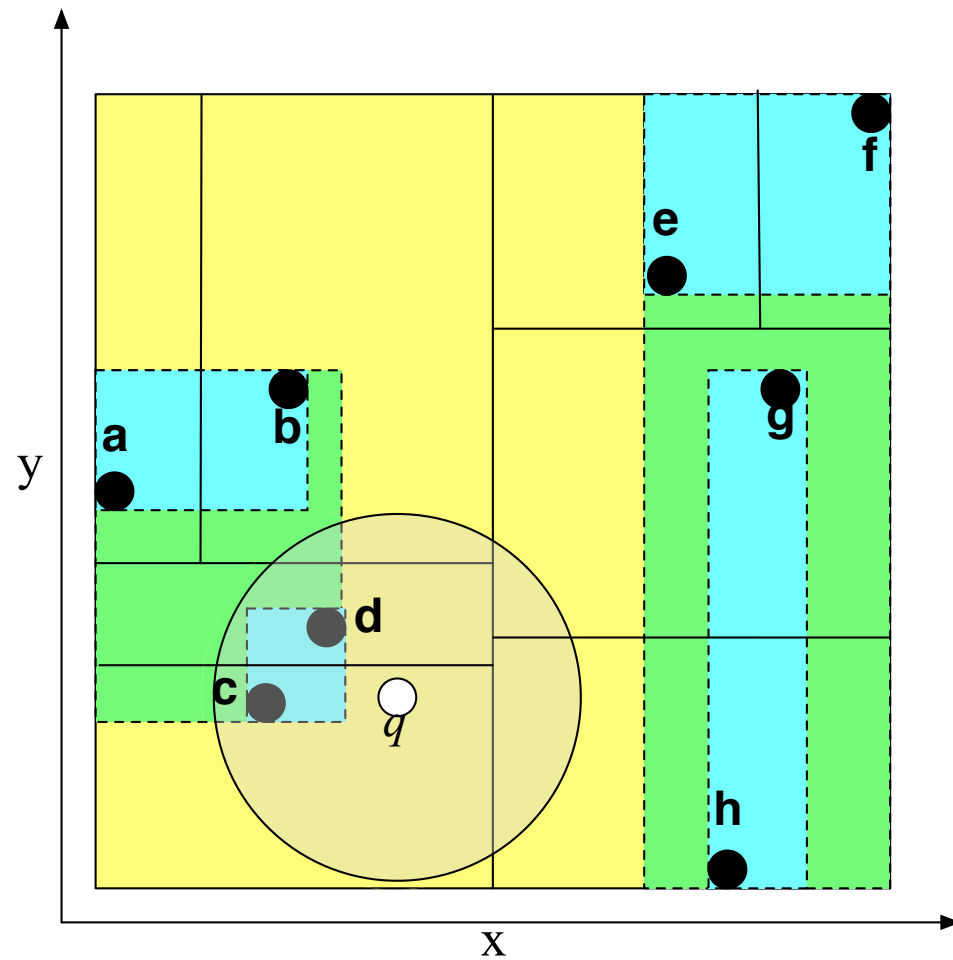
A spatial arrangement of molecular features essential for biological activity

Koes, D. R., & Camacho, C. J. (2011). Pharmer: efficient and exact pharmacophore search. *Journal of Chemical Information and Modeling*, 51(6), 1307-1314. doi:10.1021/ci200097m

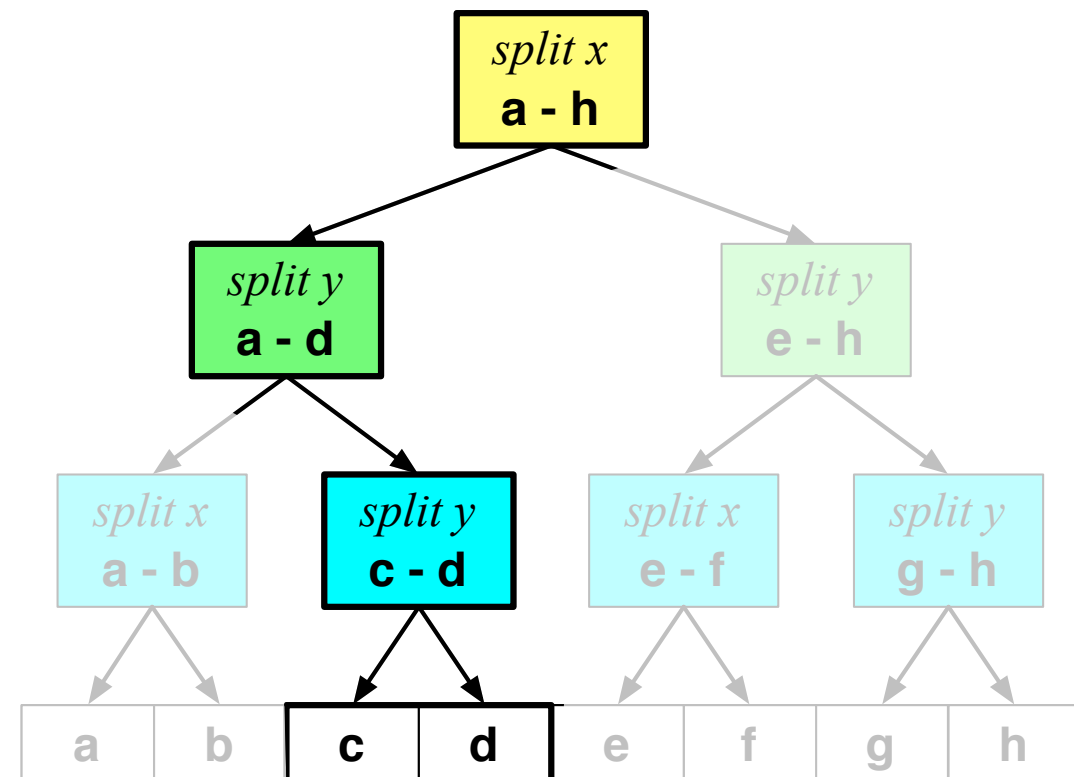
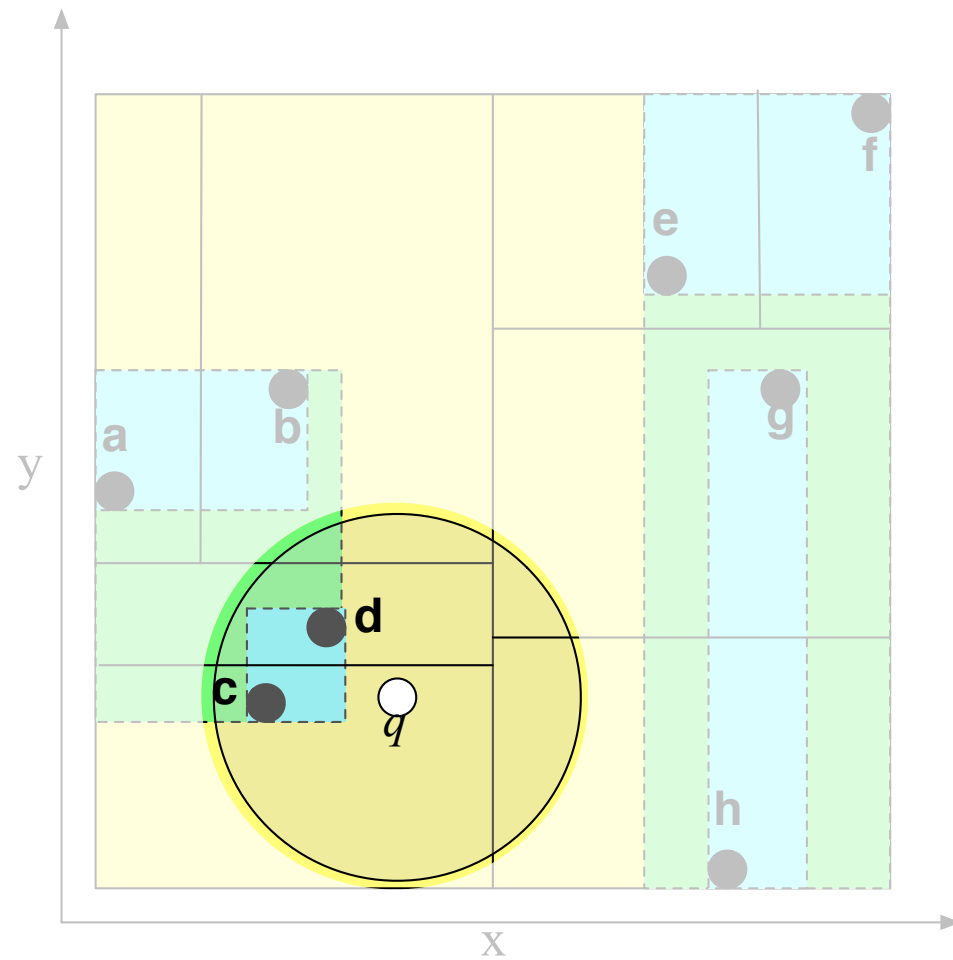
Koes, D. R., & Camacho, C. J. (2012). ZINCPharmer: pharmacophore search of the ZINC database. *Nucleic acids research*, 40(Web Server issue), W409-414. doi:10.1093/nar/gks378

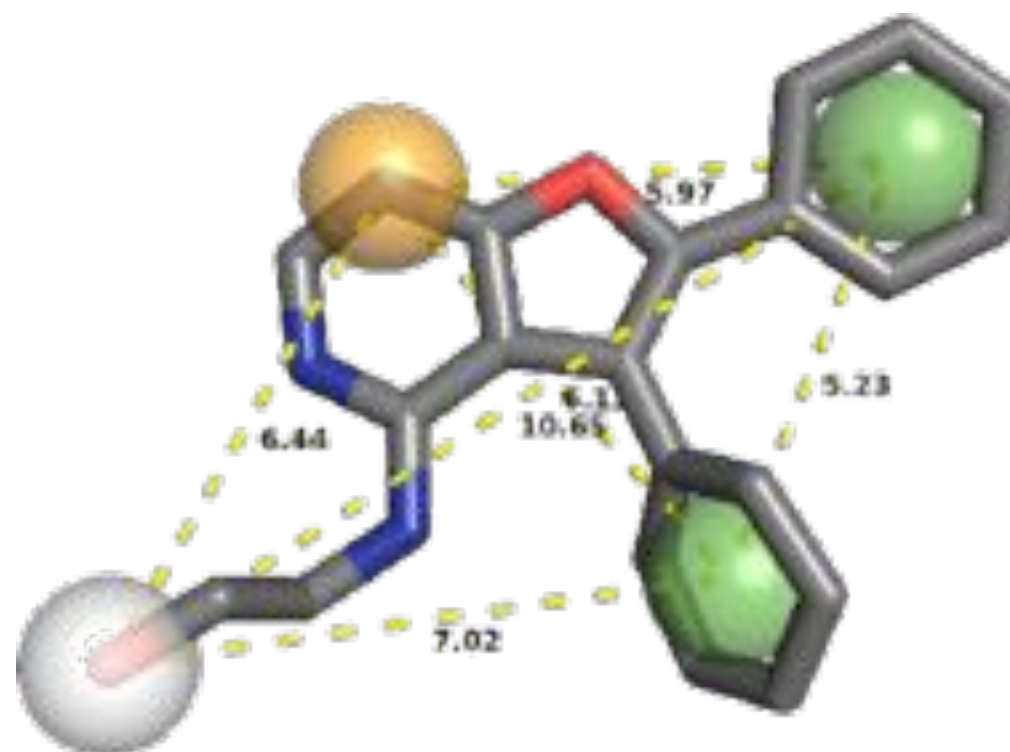
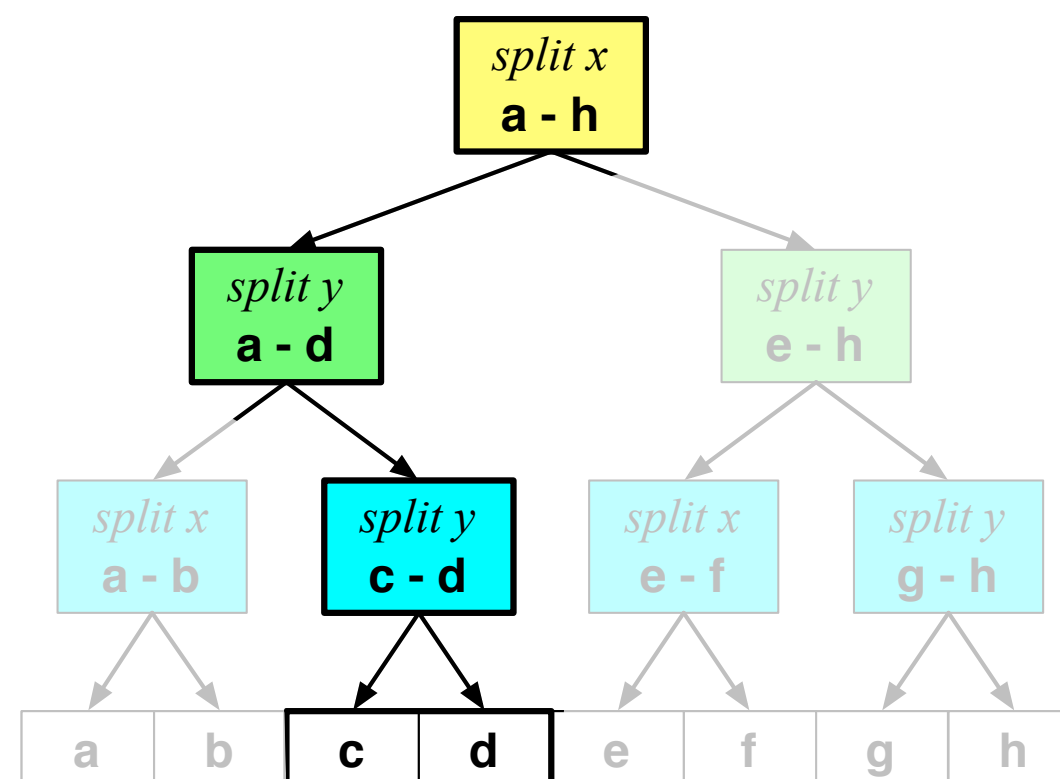
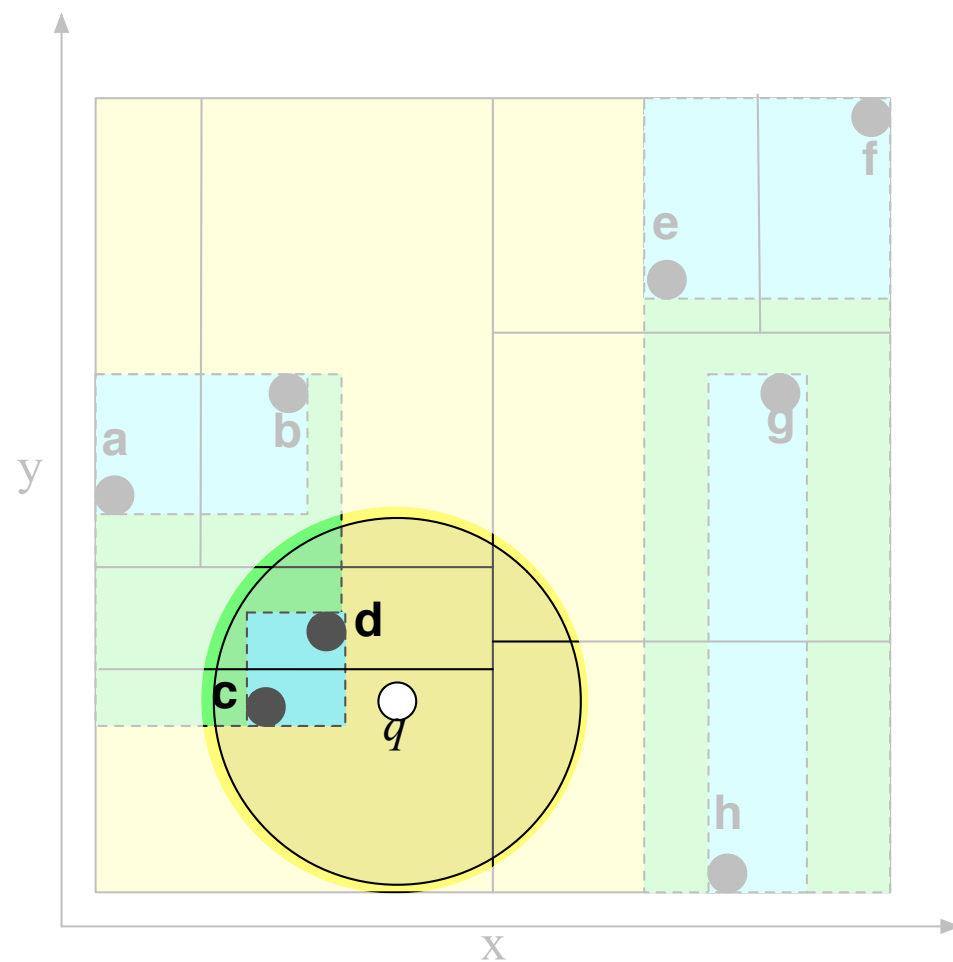


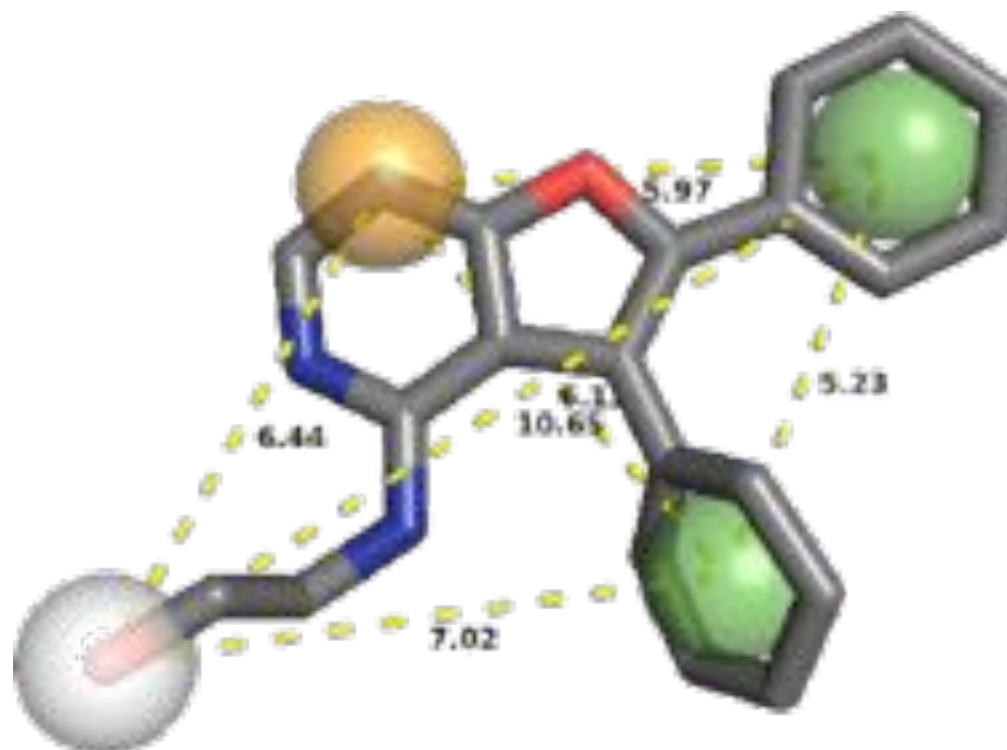
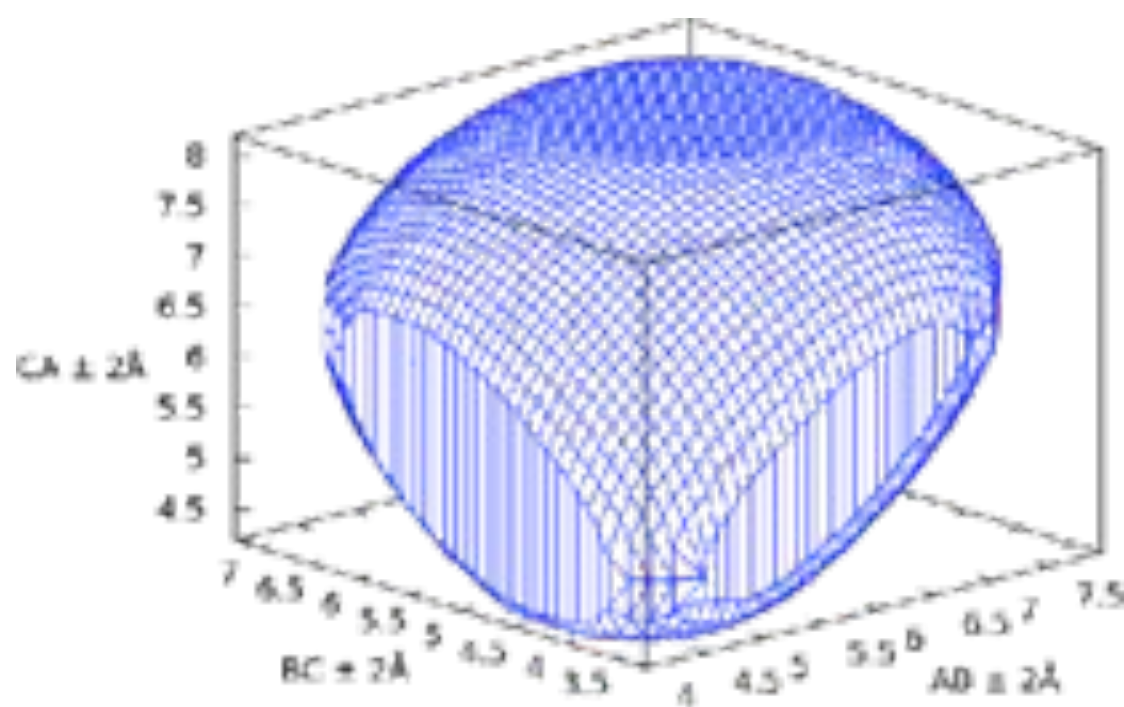
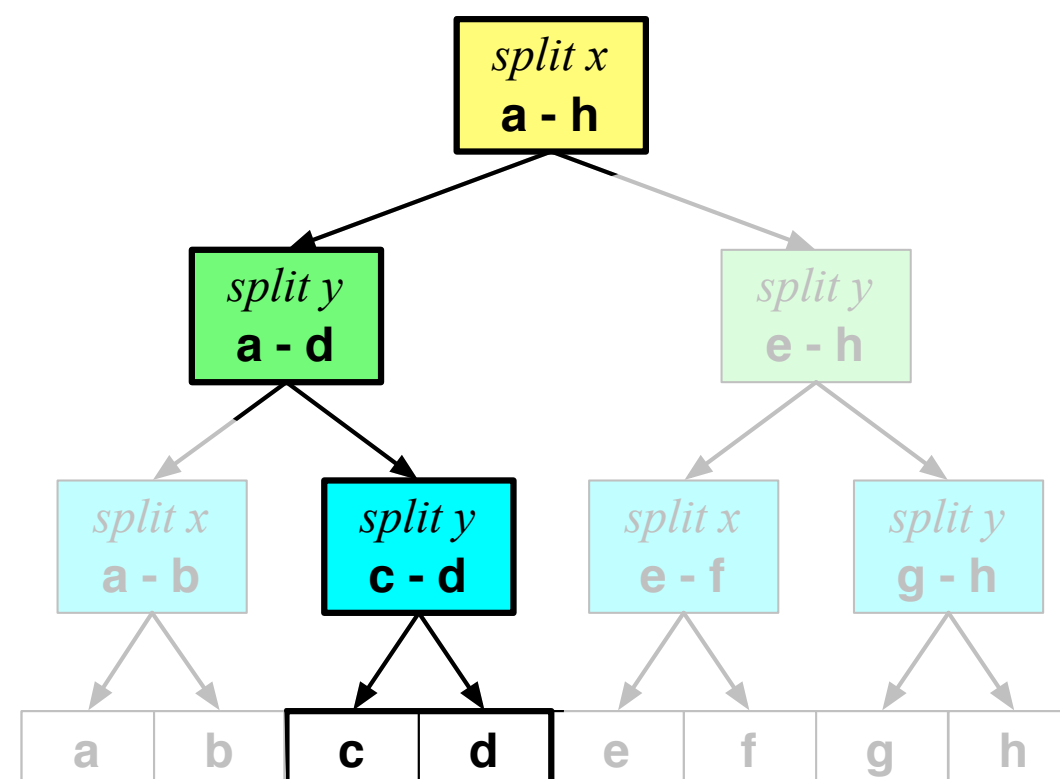
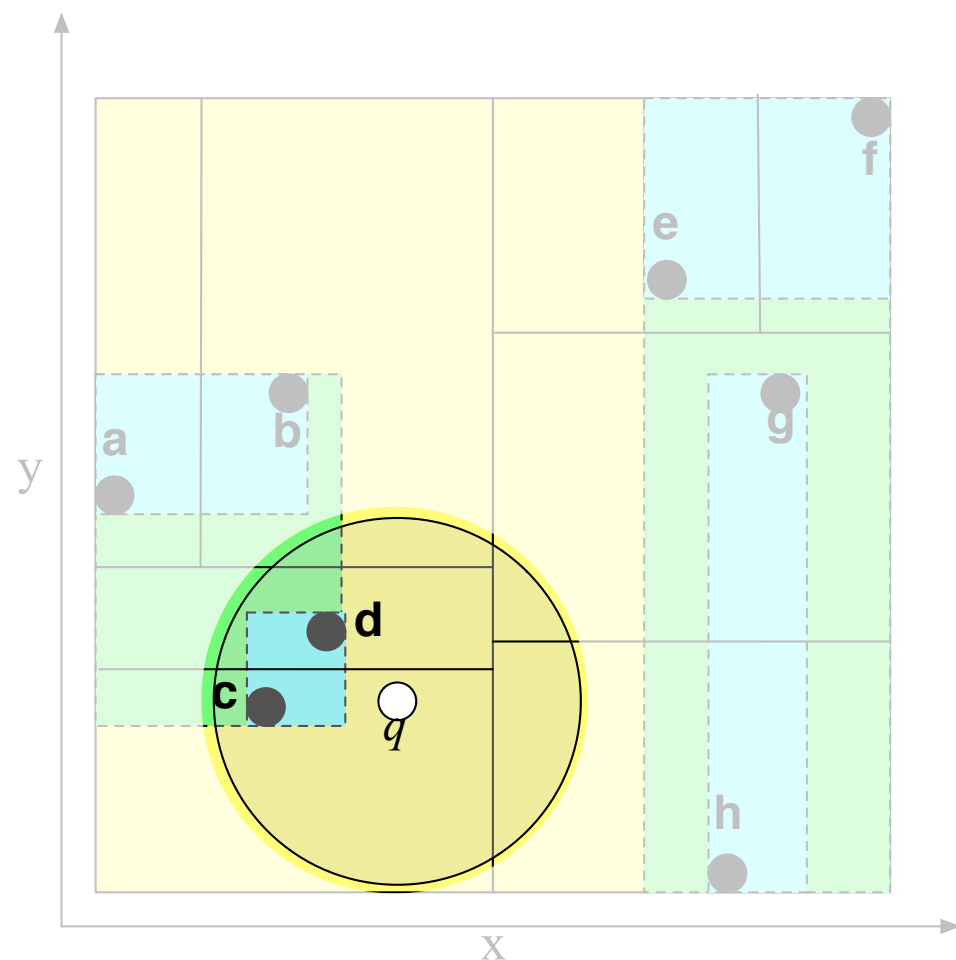


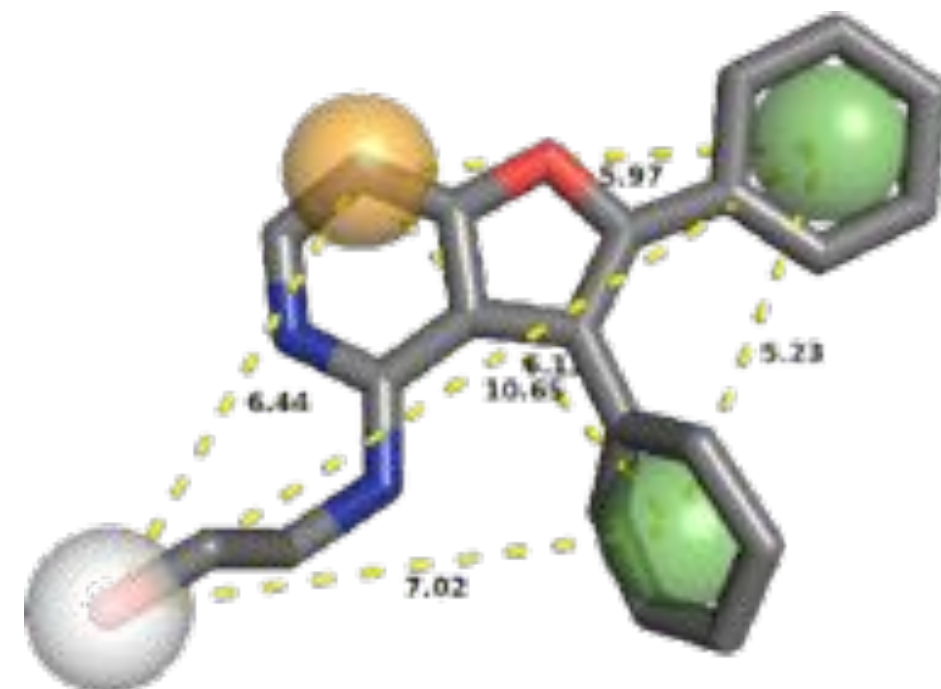
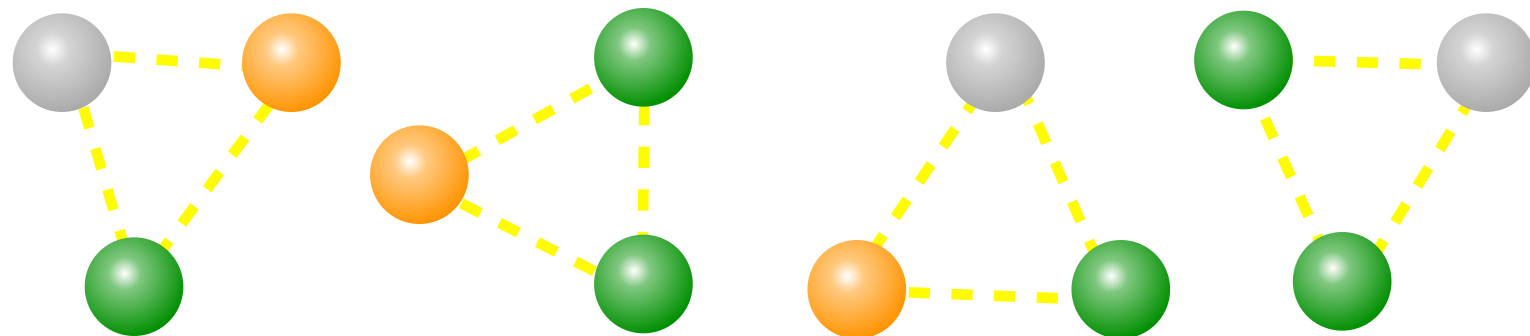


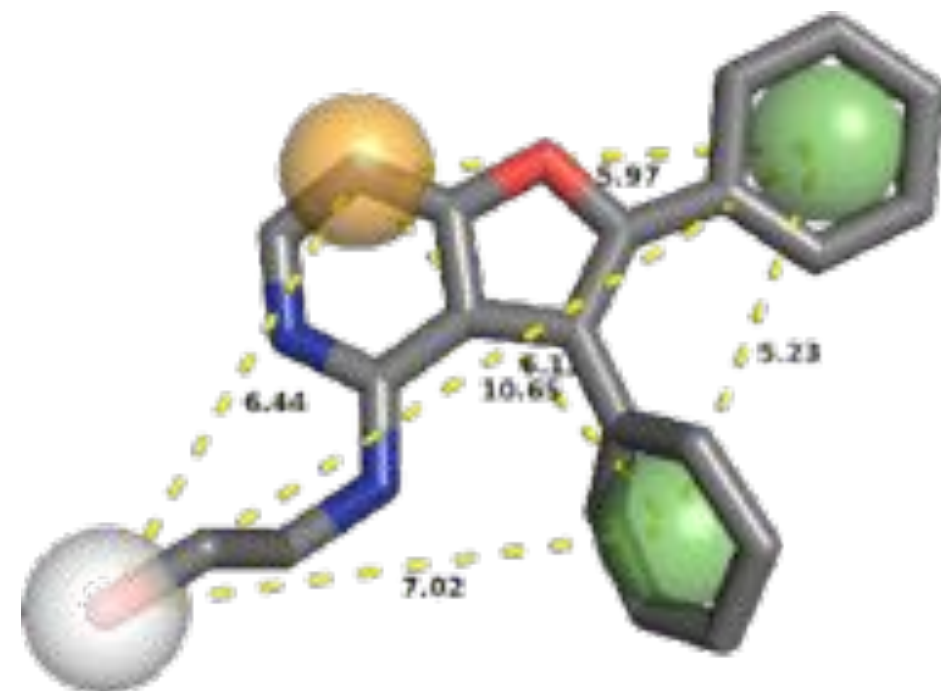
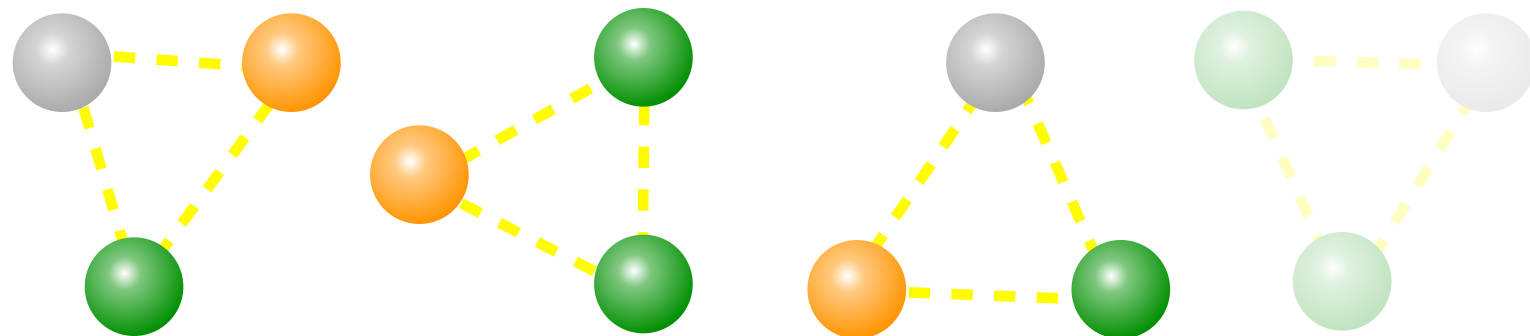
Efficient and Exact Pharmacophore Search

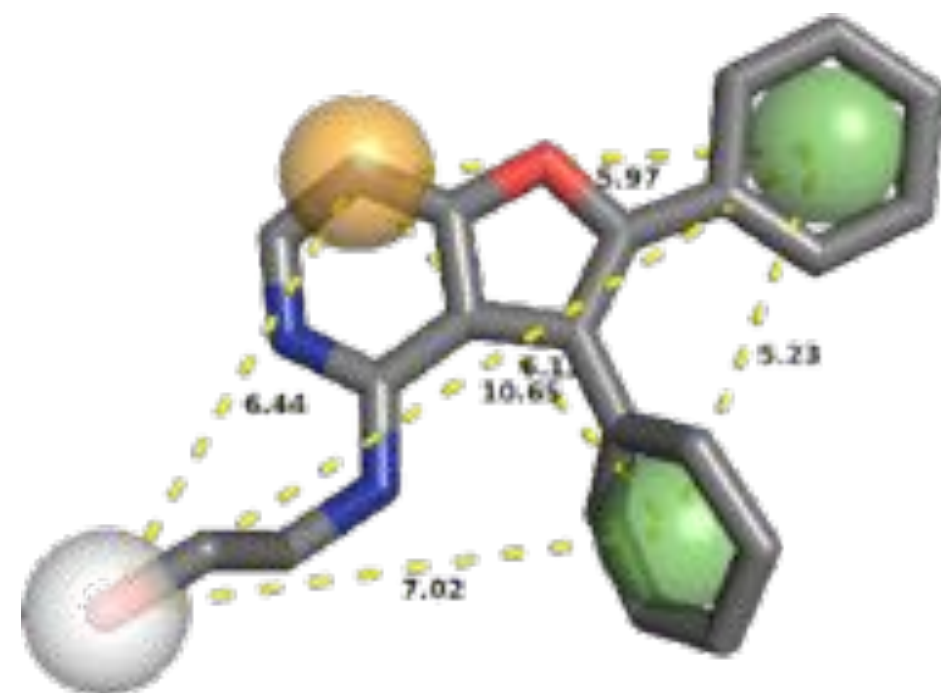
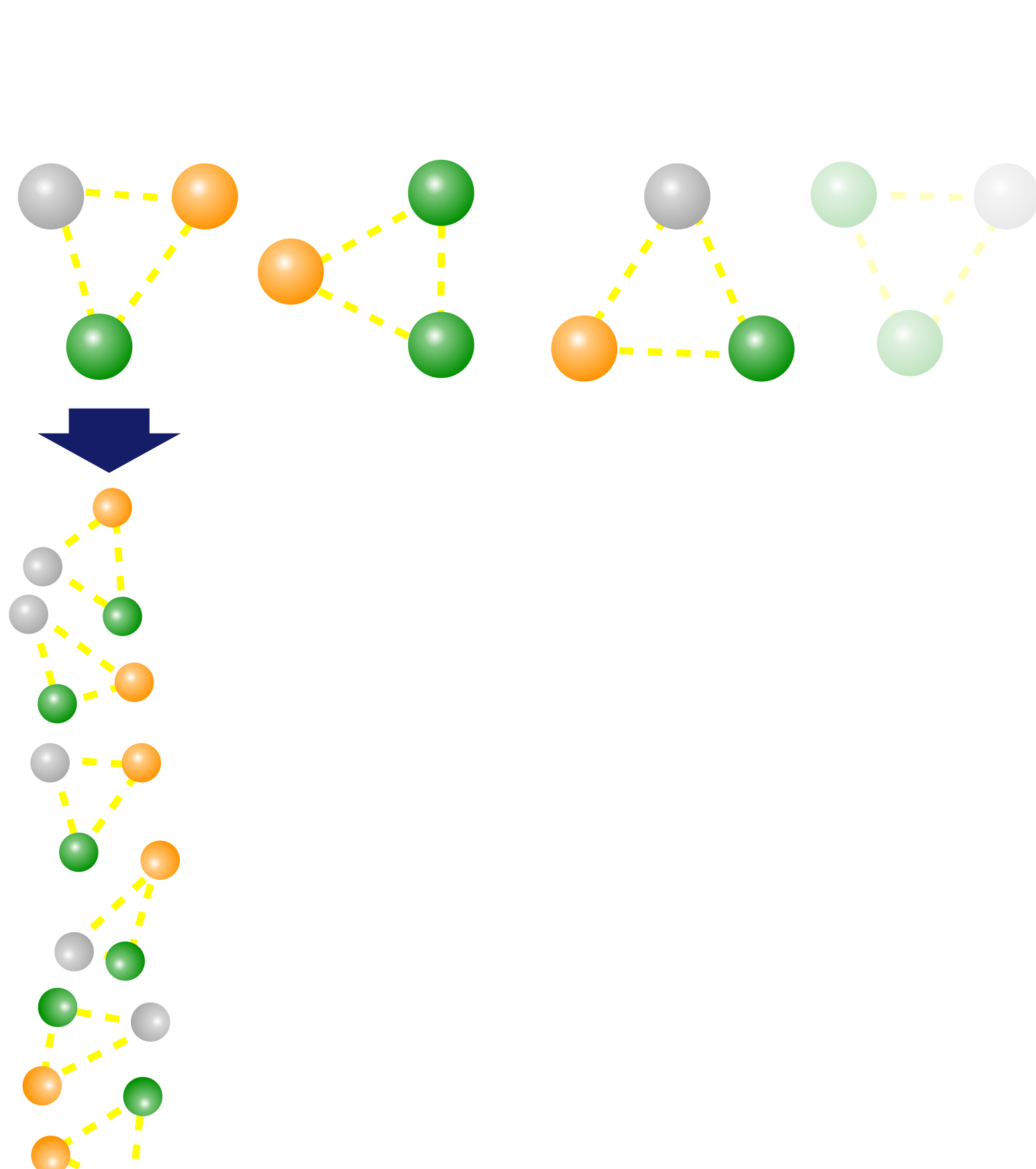


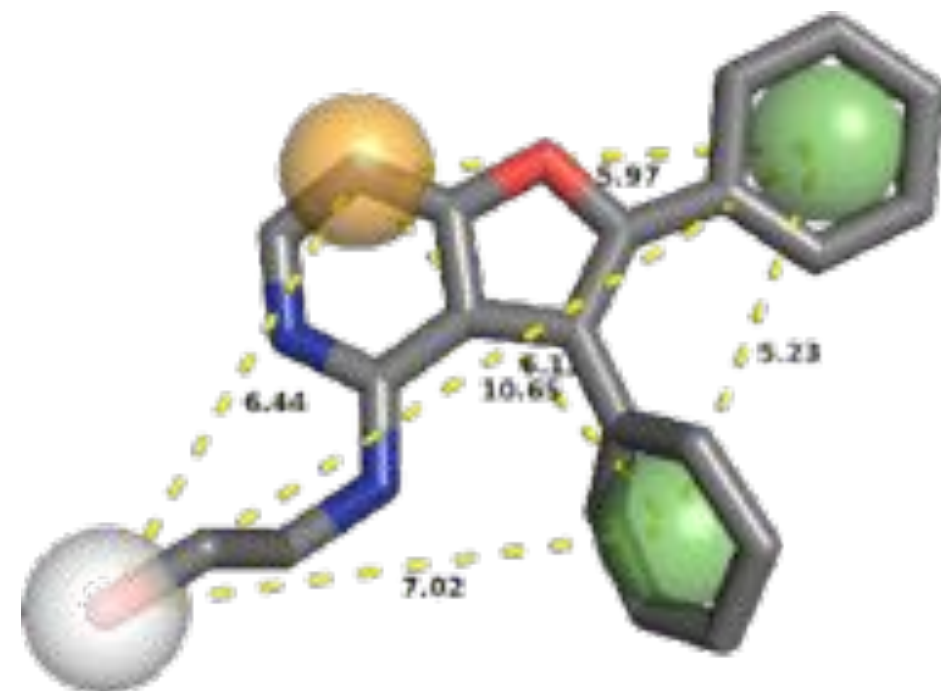
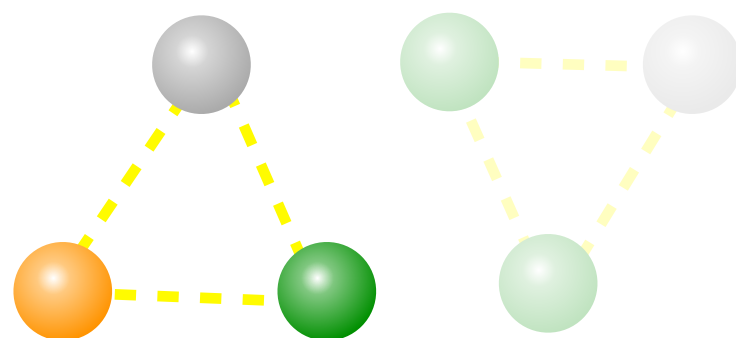
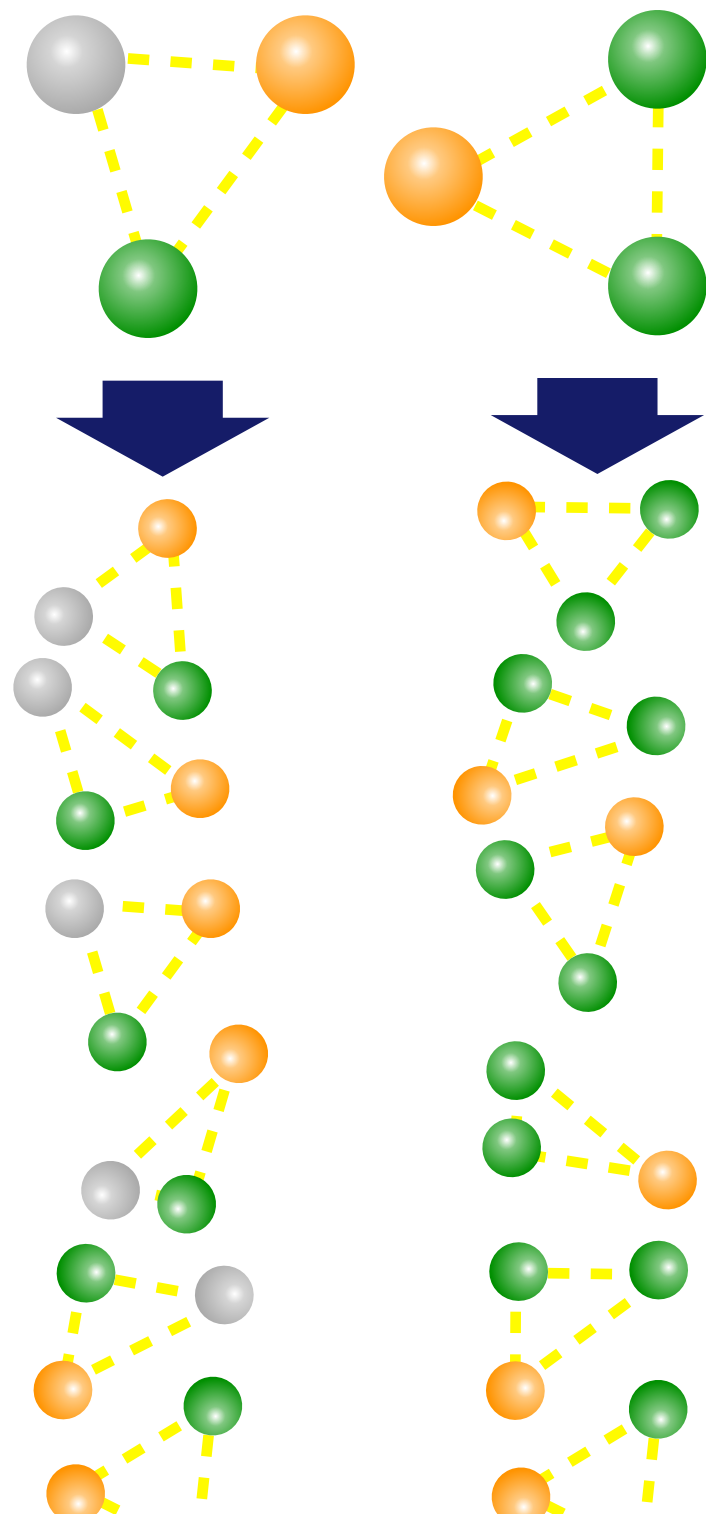


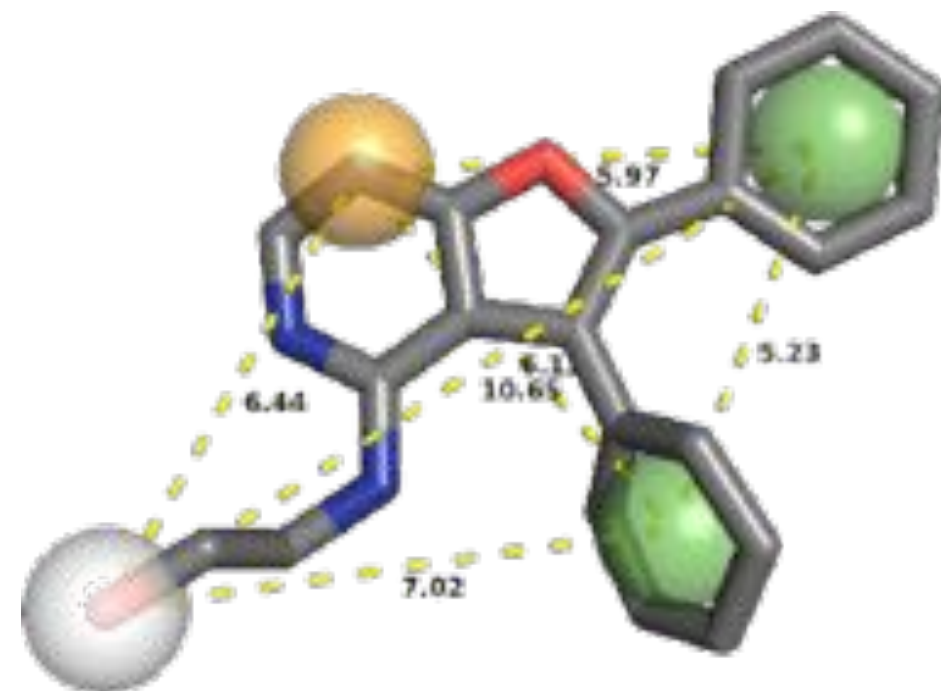
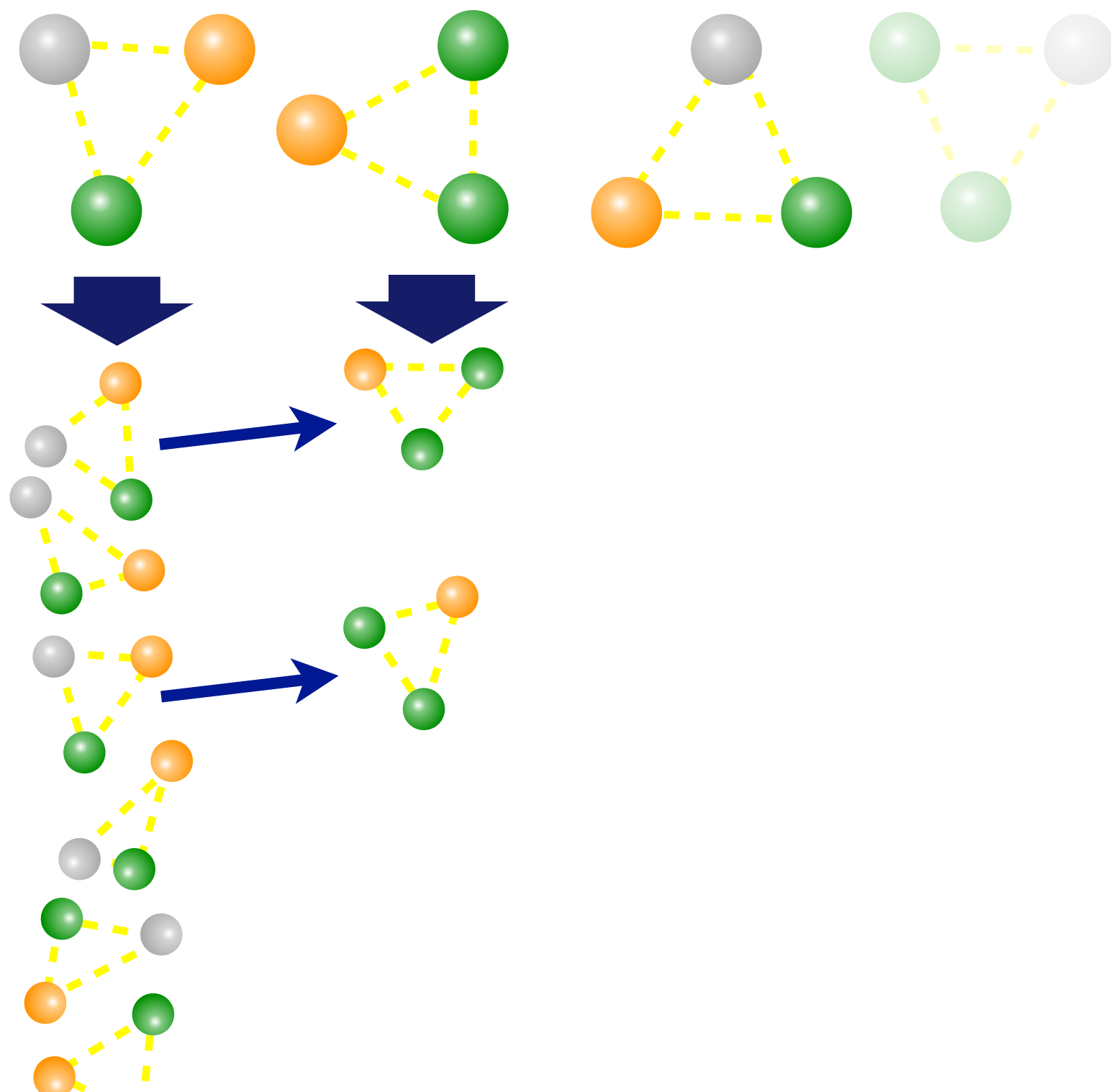


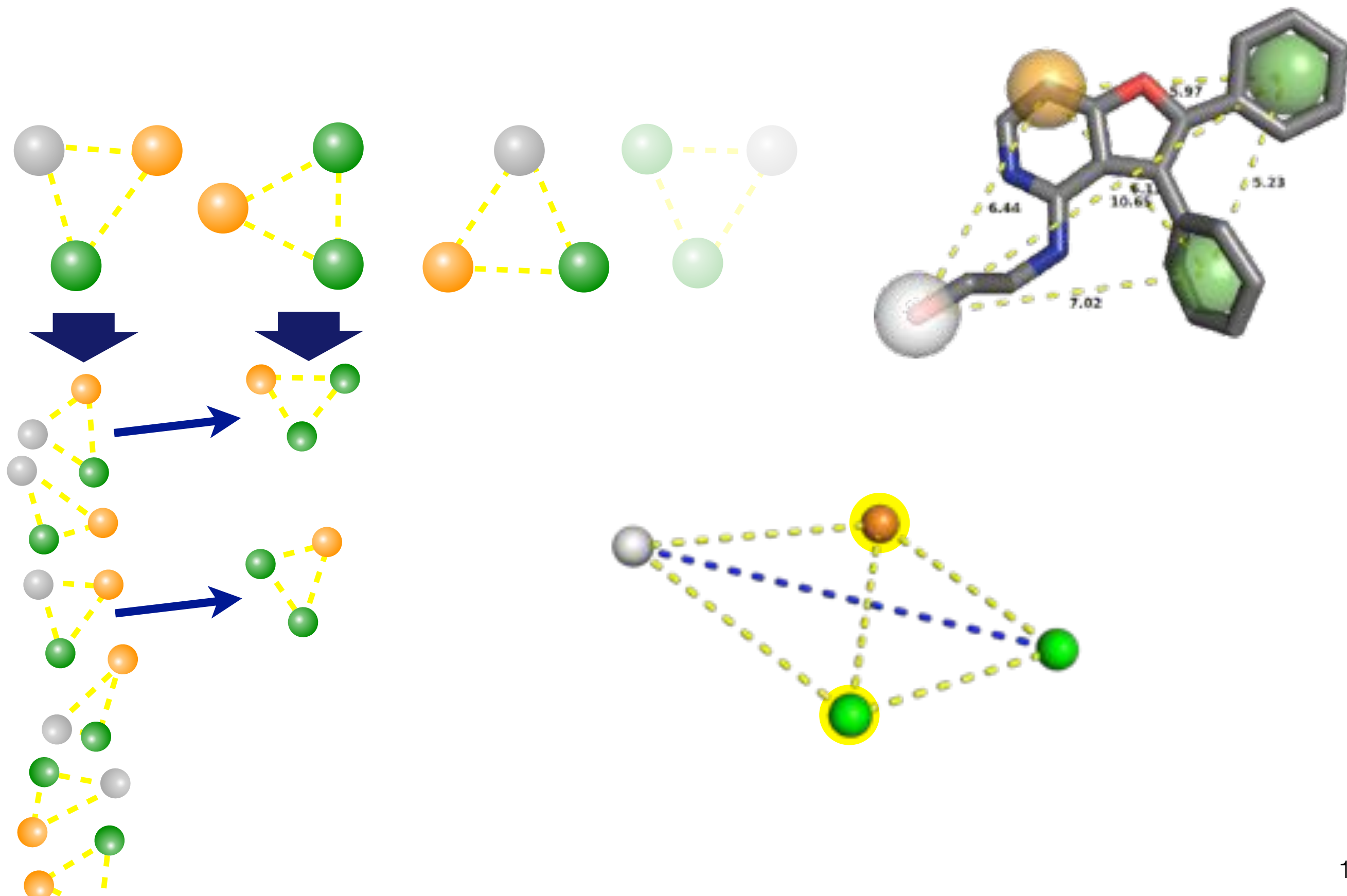


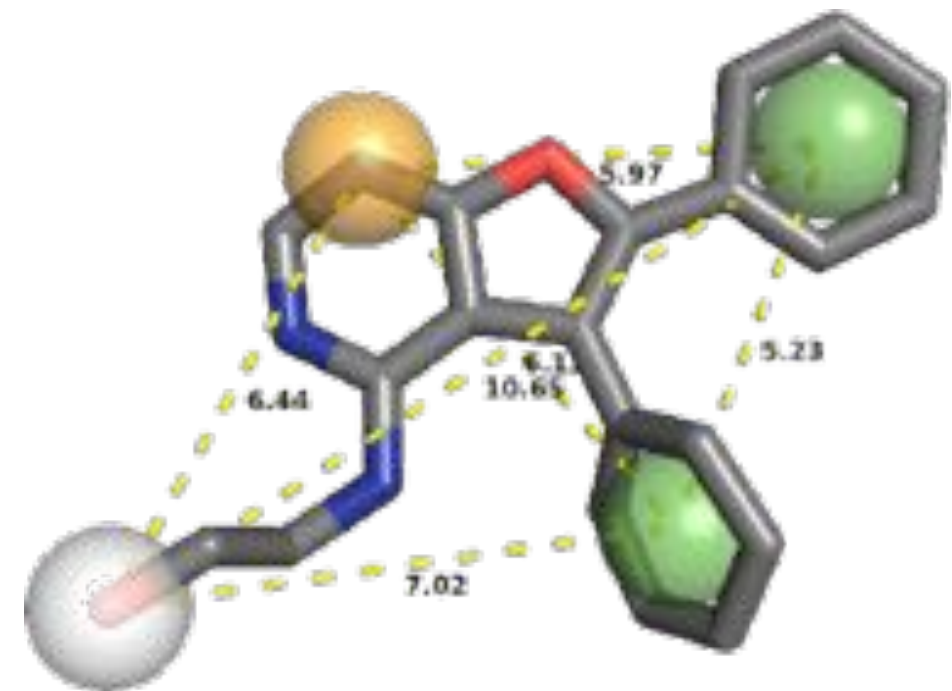
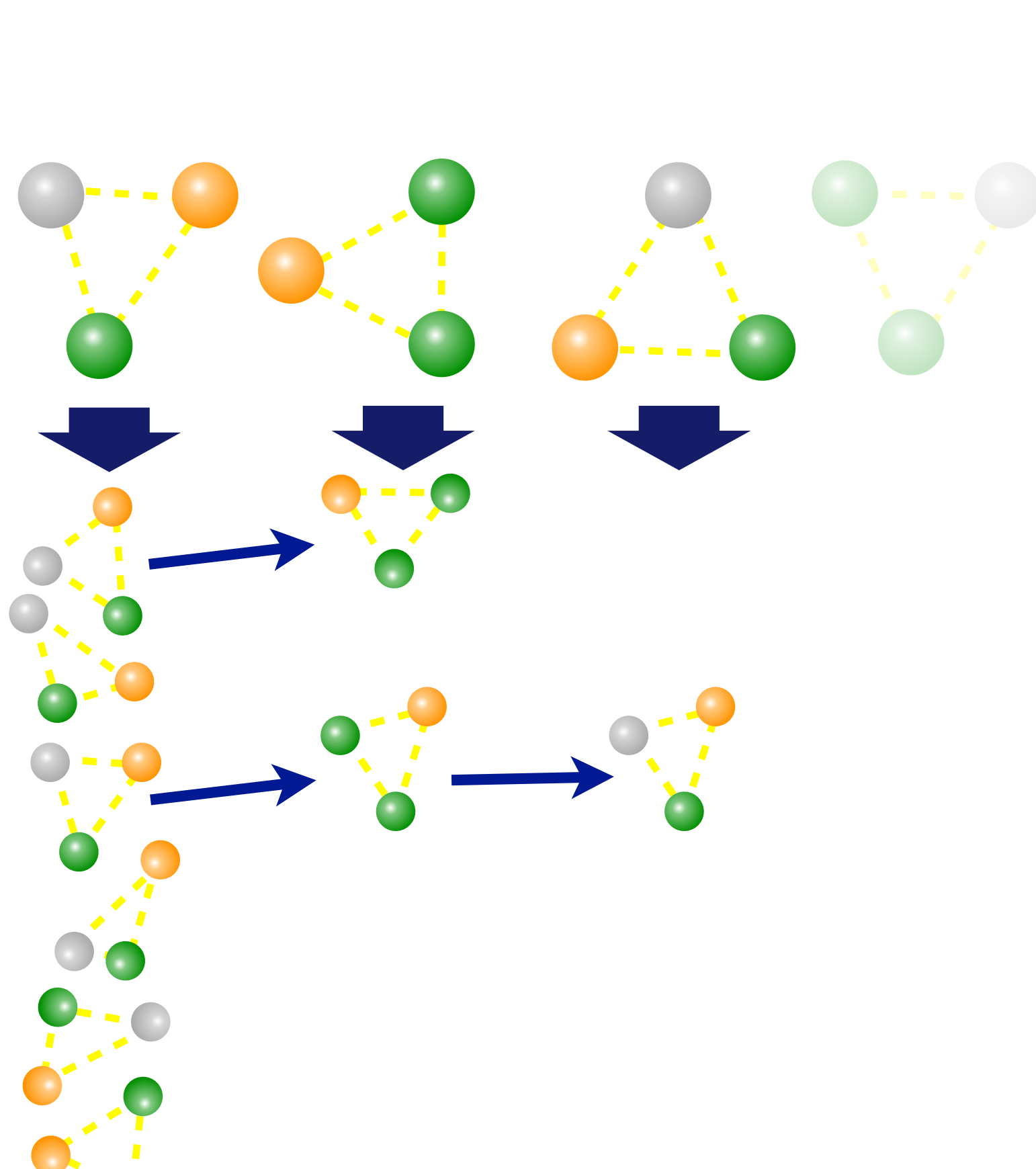


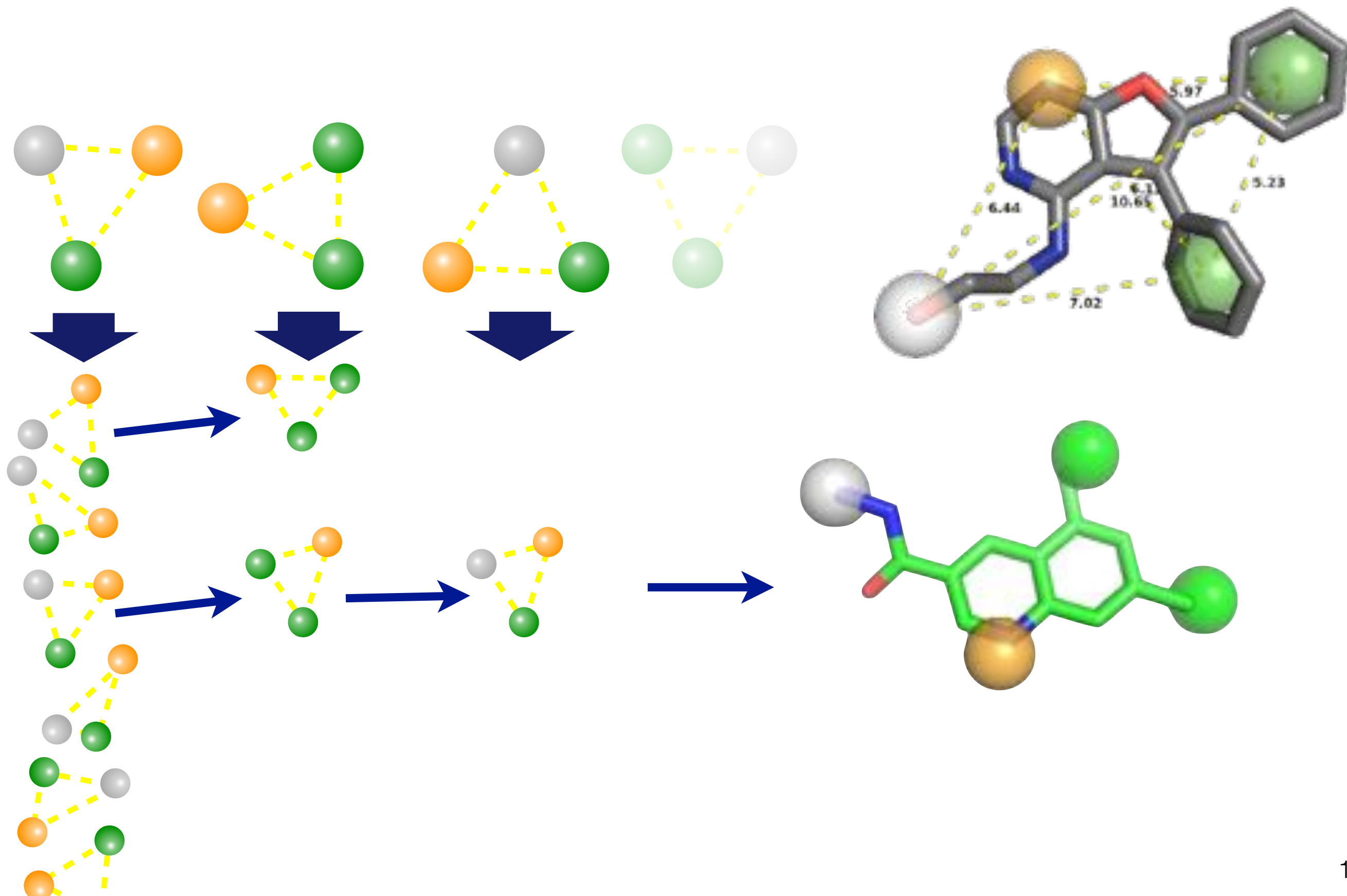




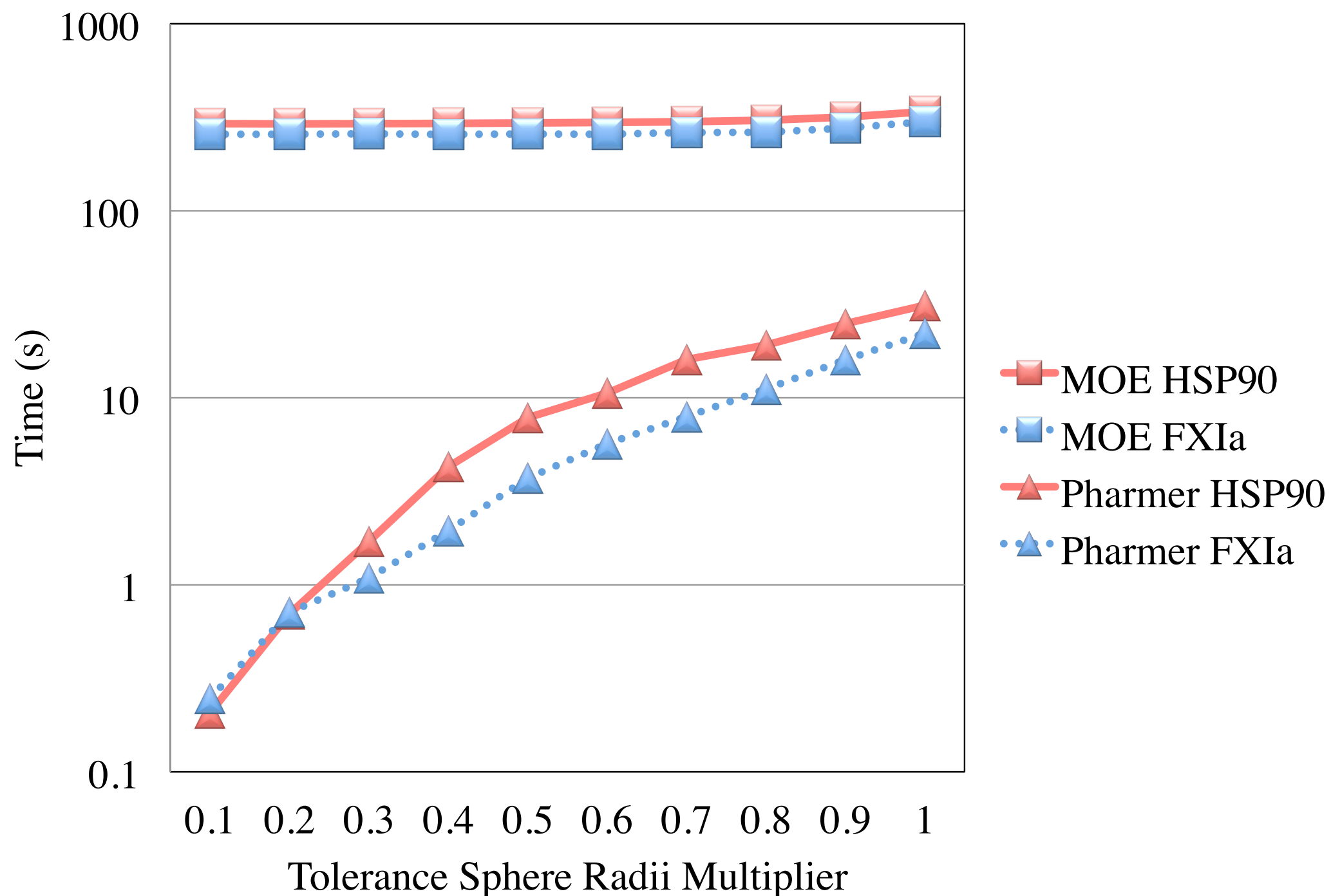


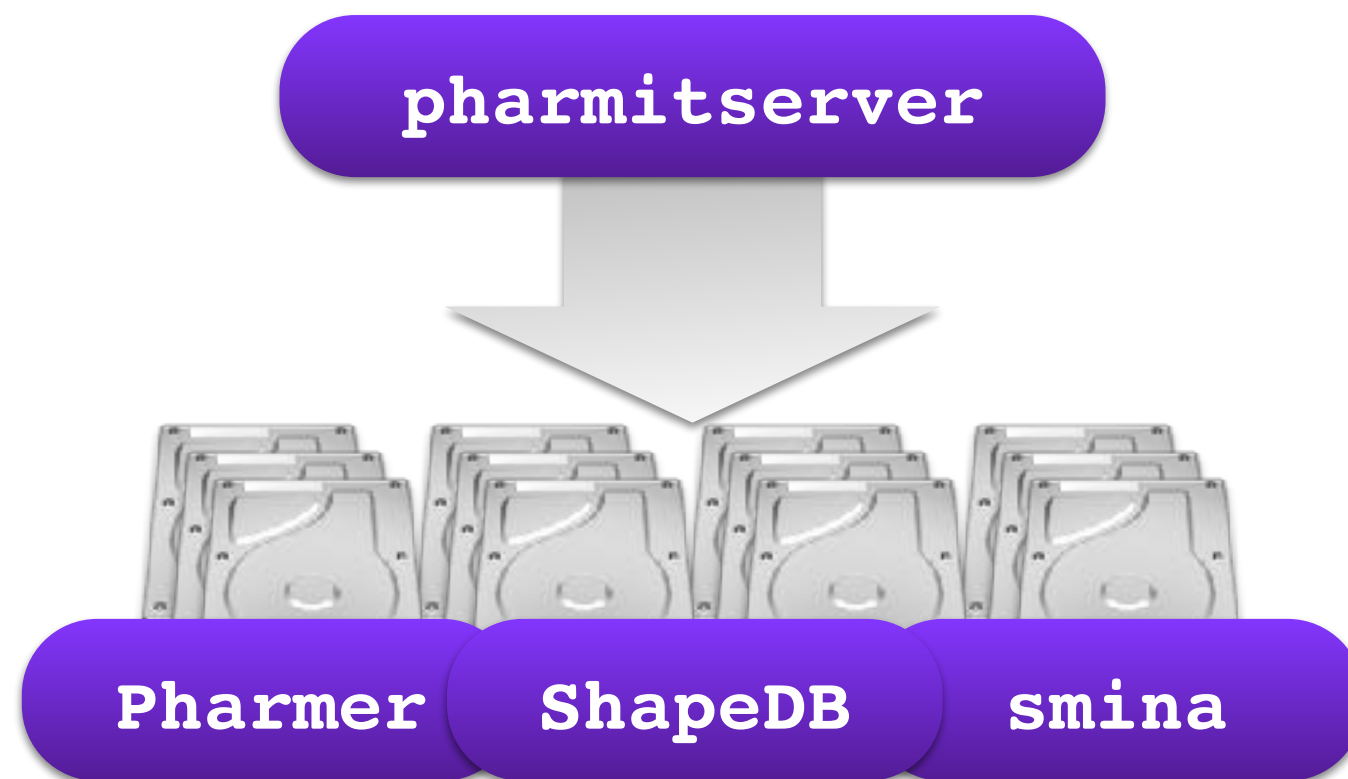




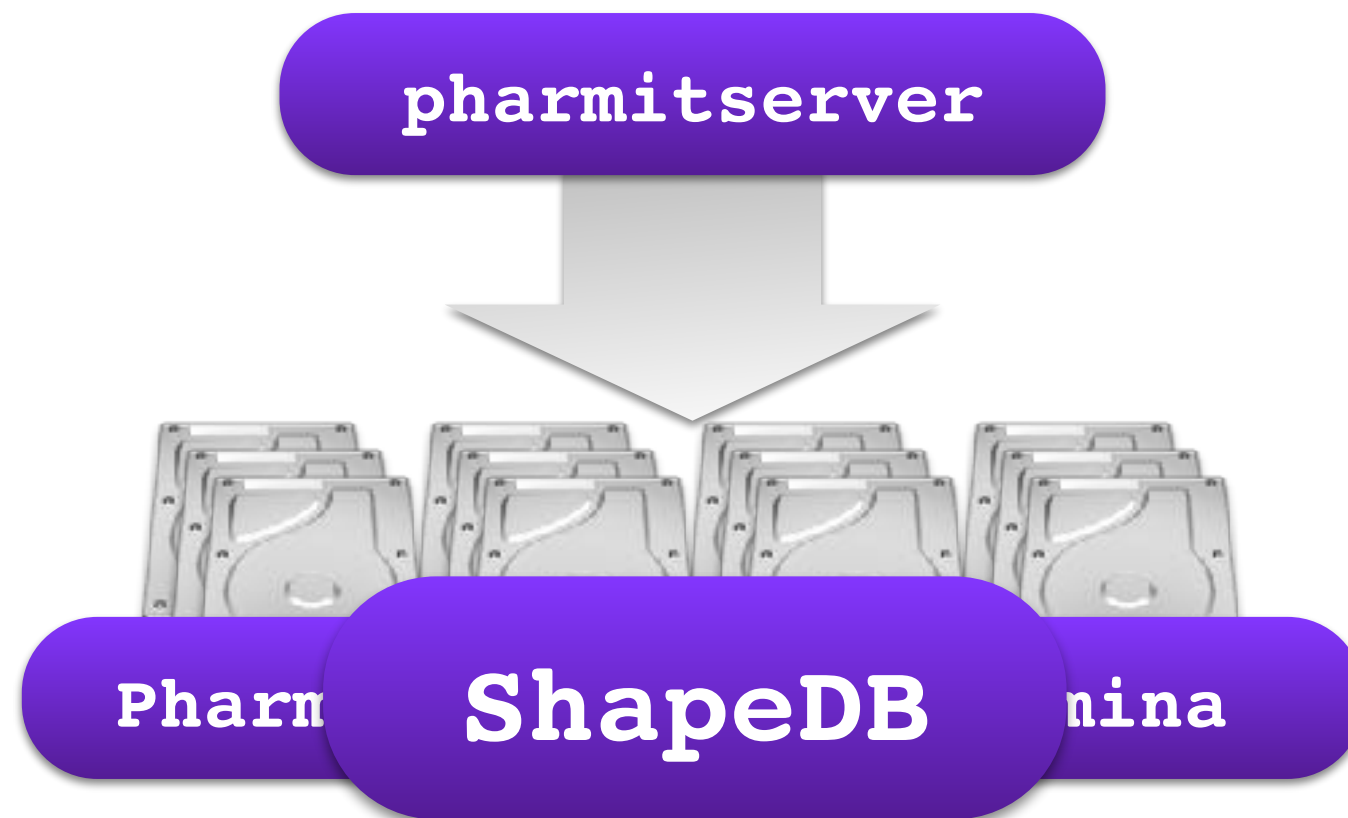


Efficient and Exact Pharmacophore Search



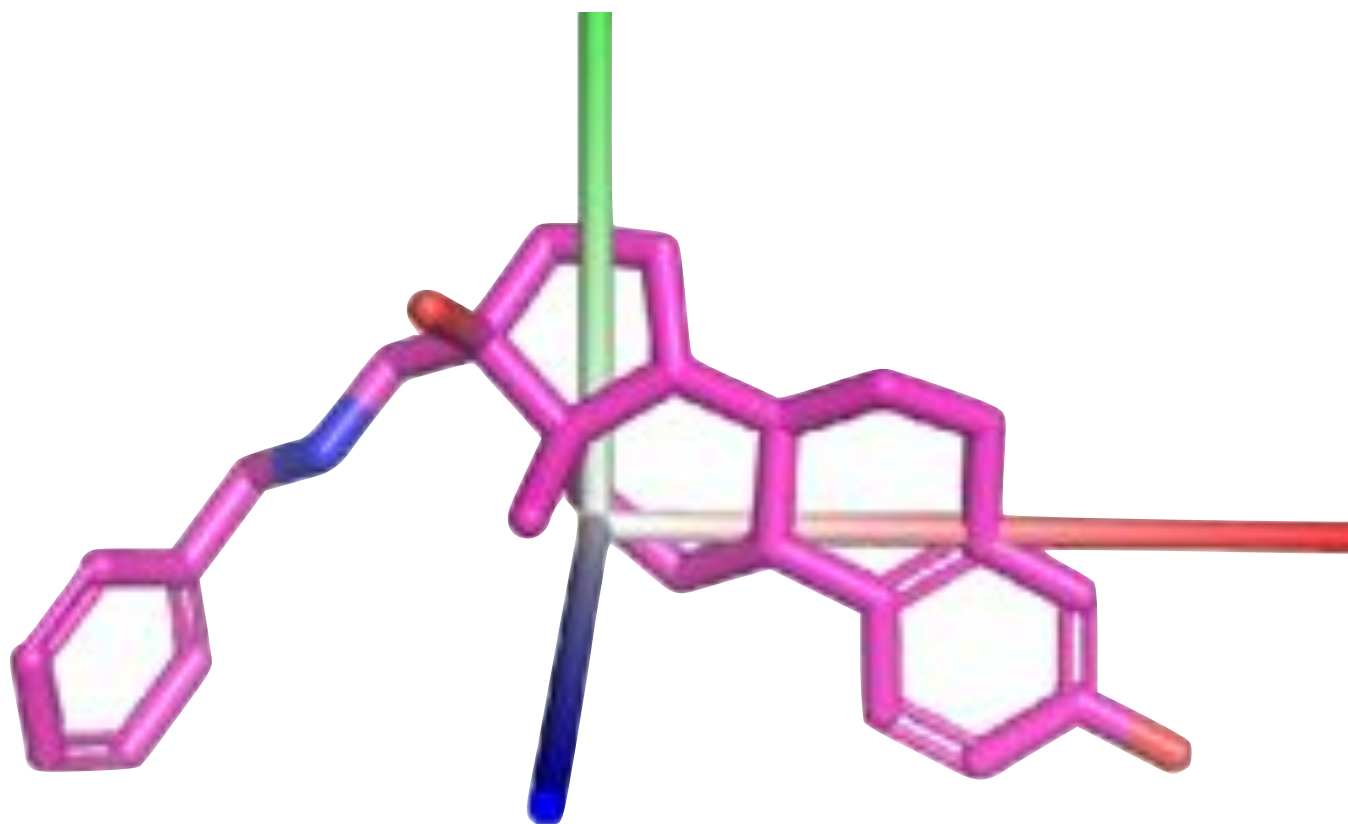


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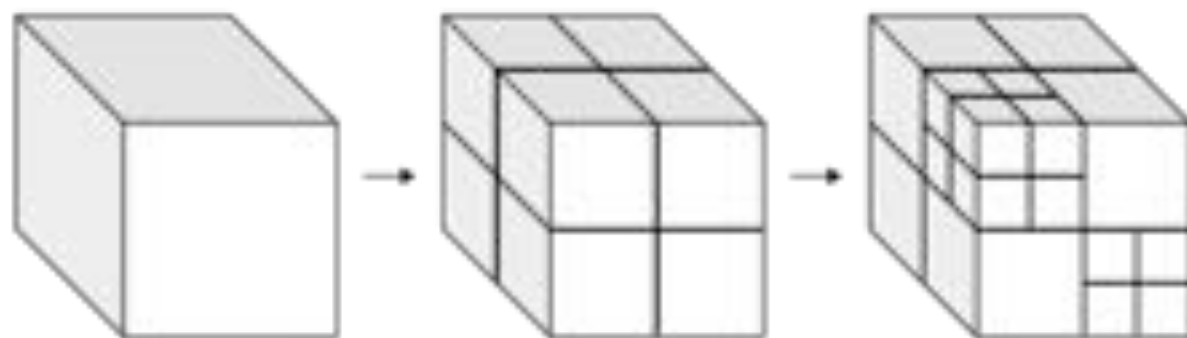
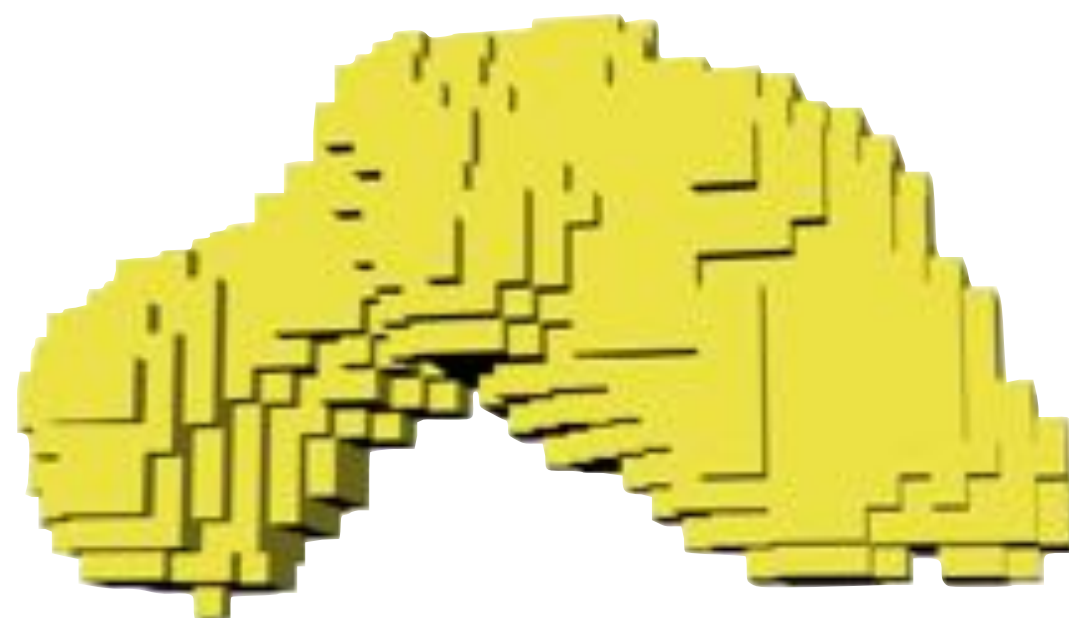


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Align to Moments of Inertia



Voxelize

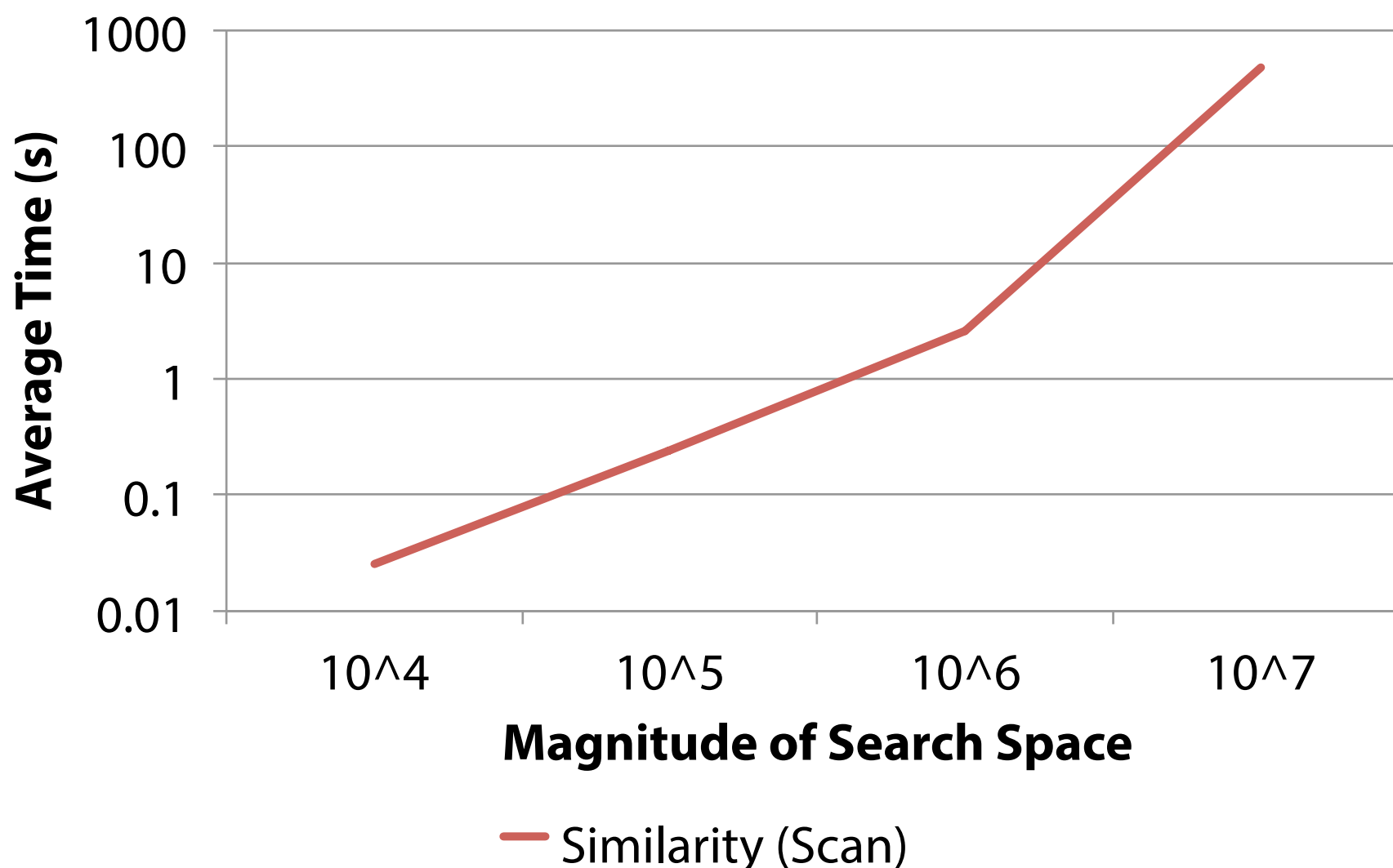


Oct-tree

- Scales with Surface Area, not Volume
- Fast Intersection/Union Operations

$$\delta(A, B) = \frac{A \cap B}{A \cup B}$$

Performance of Shape Similarity Search



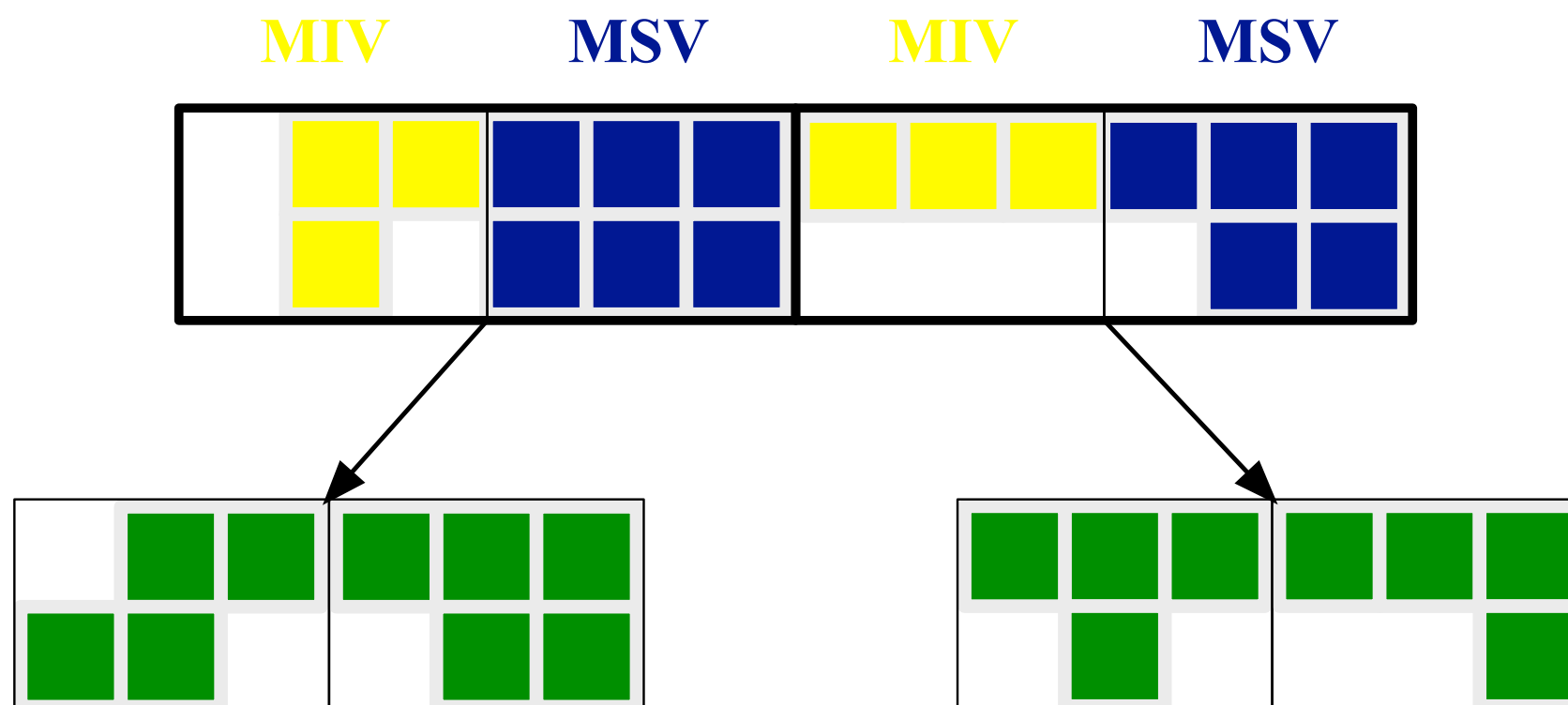
Indexed Search of Molecular Shapes

MIV

Maximum Included Volume

Intersection**MSV**

Minimum Surrounding Volume

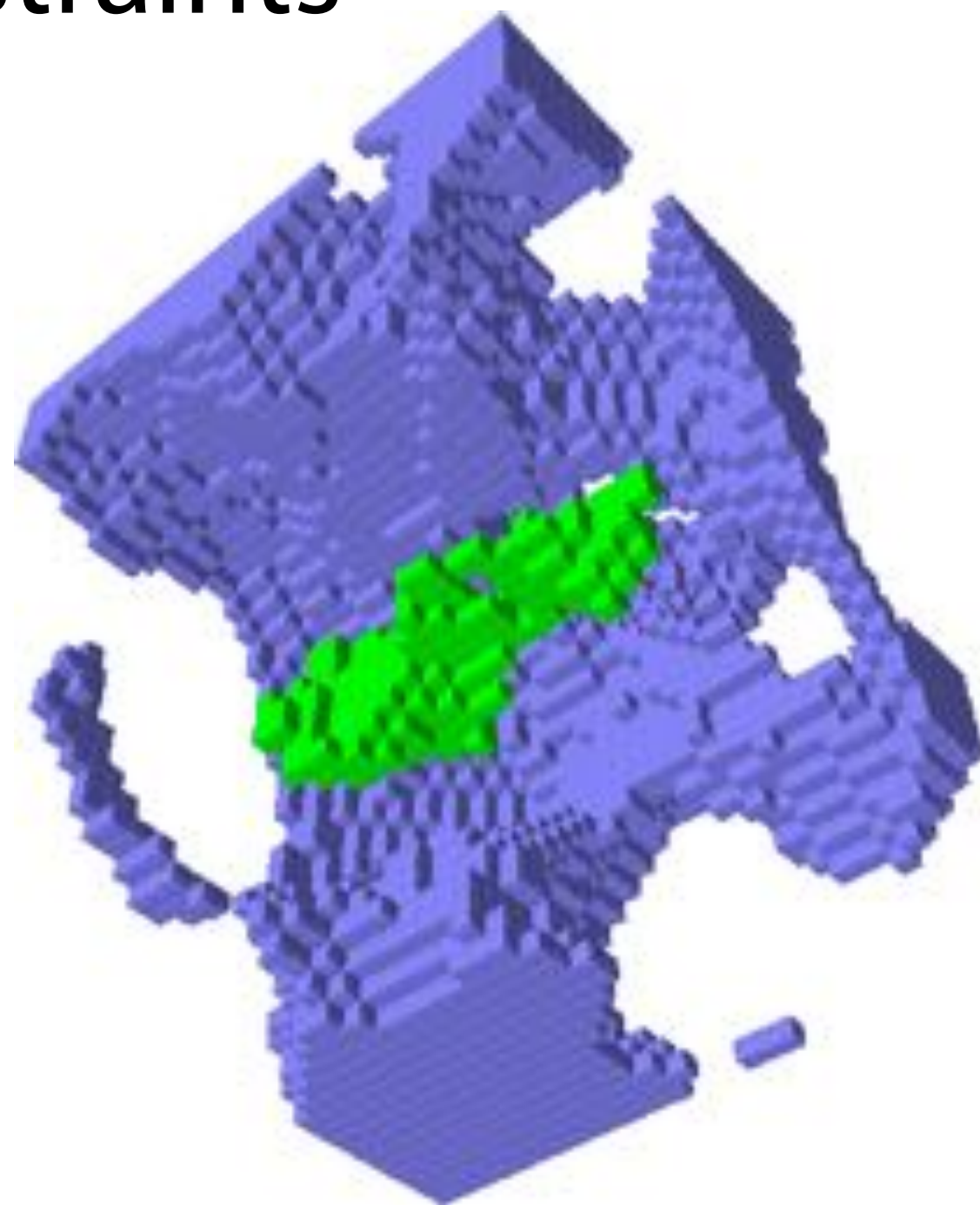
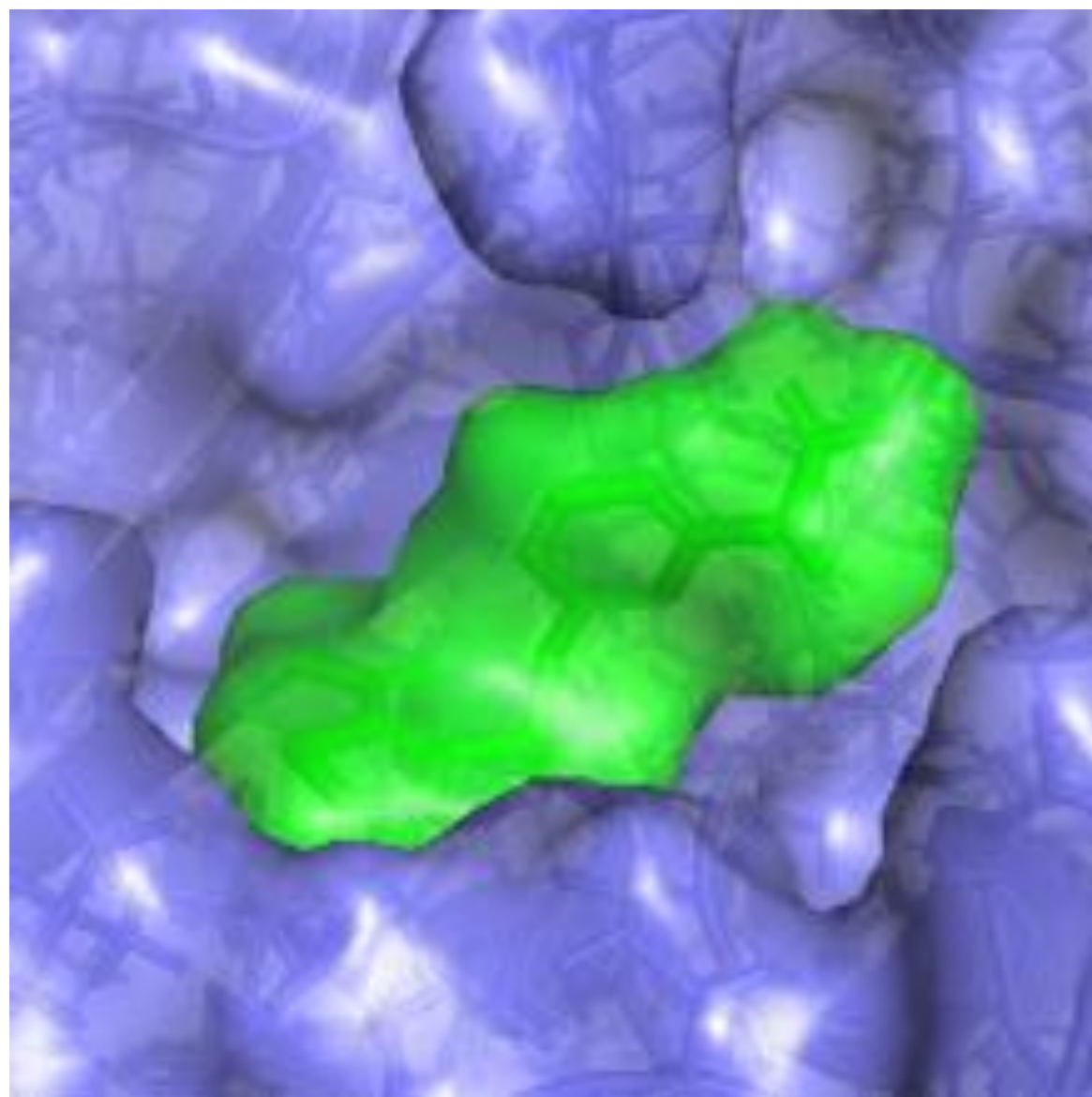
Union

Matching and packing algorithm for
efficient and effective initialization

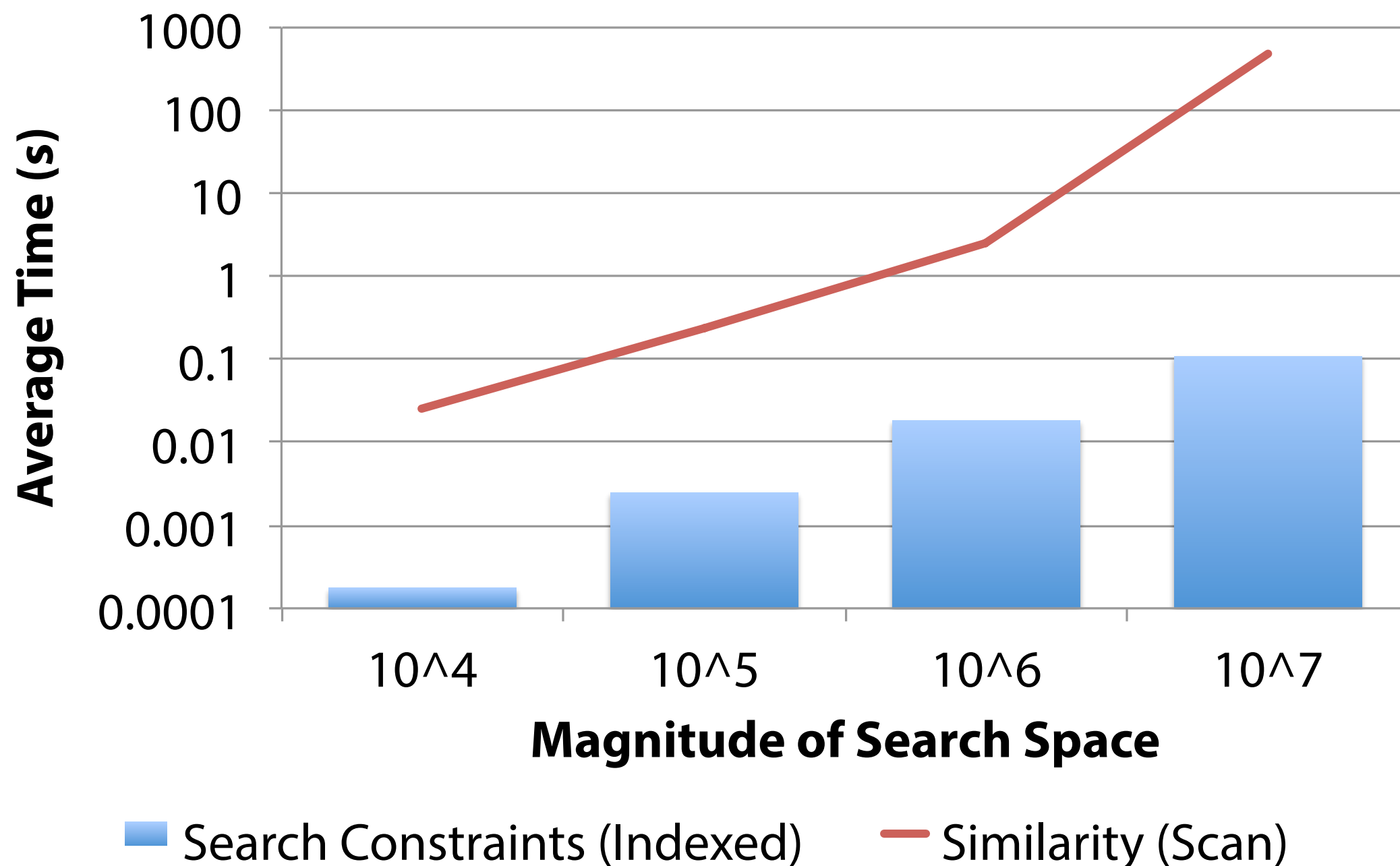
Koes, D. R., & Camacho, C. J. (2014). Shape-based virtual screening with volumetric aligned molecular shapes. *J Comput Chem*, 35(25), 1824-1834. doi:10.1002/jcc.23690

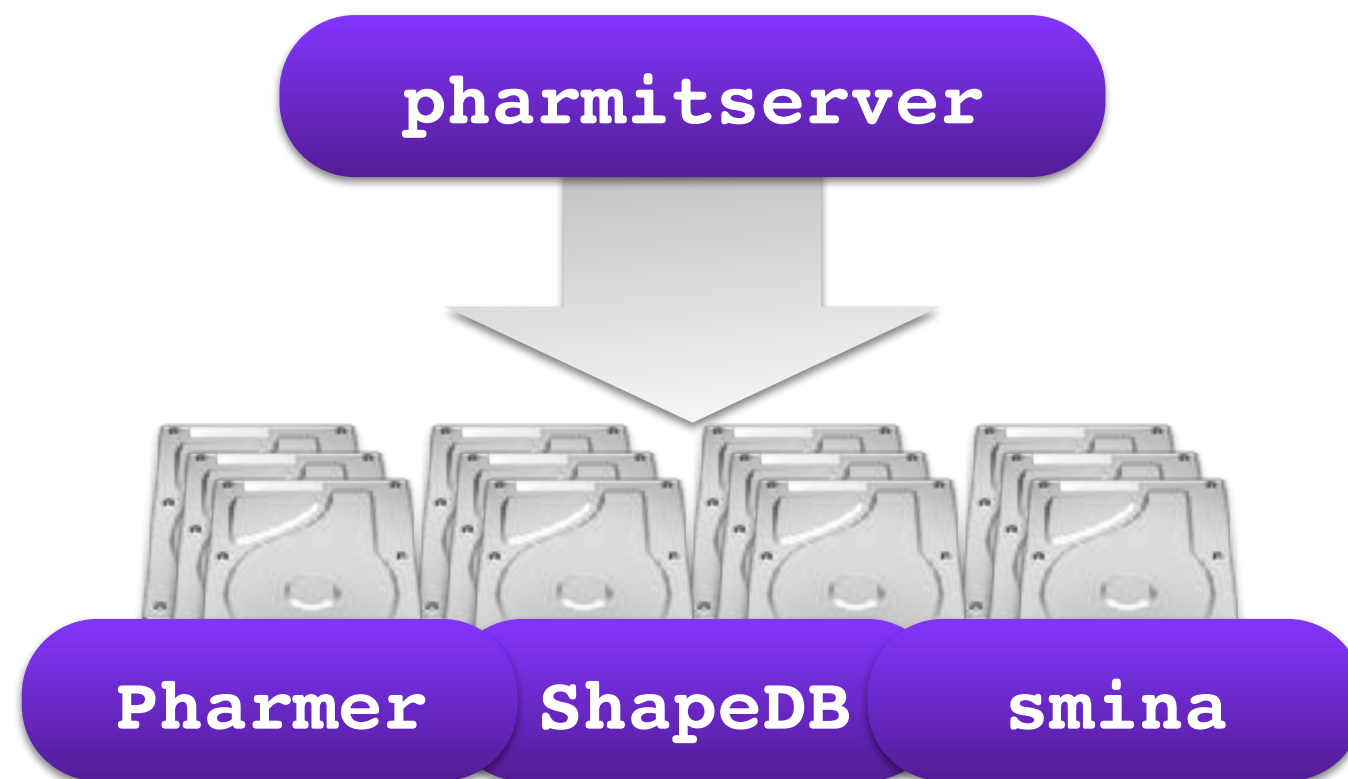
Koes, D., & Camacho, C. (2014). Indexing volumetric shapes with matching and packing. *Knowledge and Information Systems*, 1-24. doi:10.1007/s10115-014-0729-z

Shape Constraints

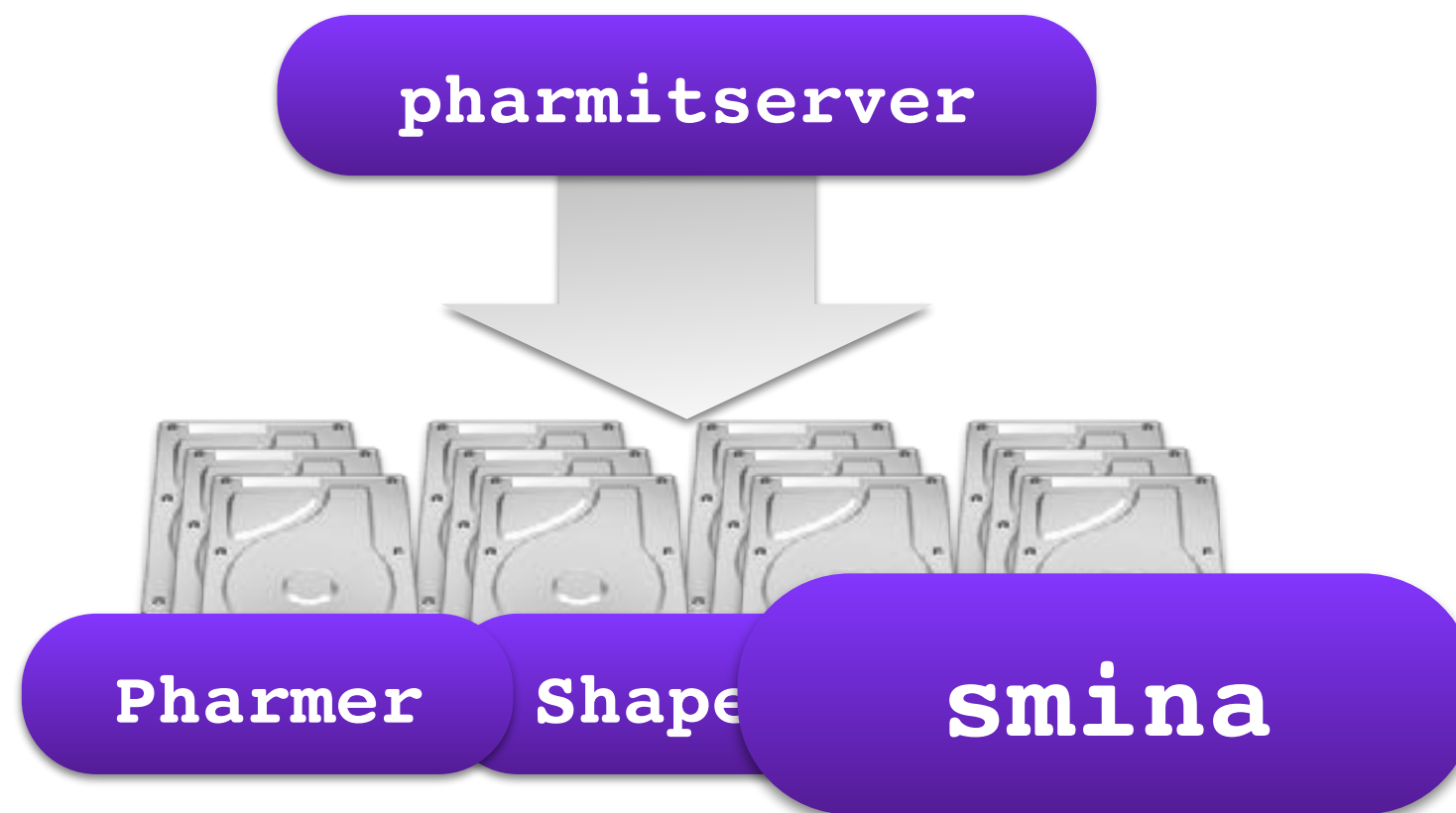


Performance of Shape Constraint Search

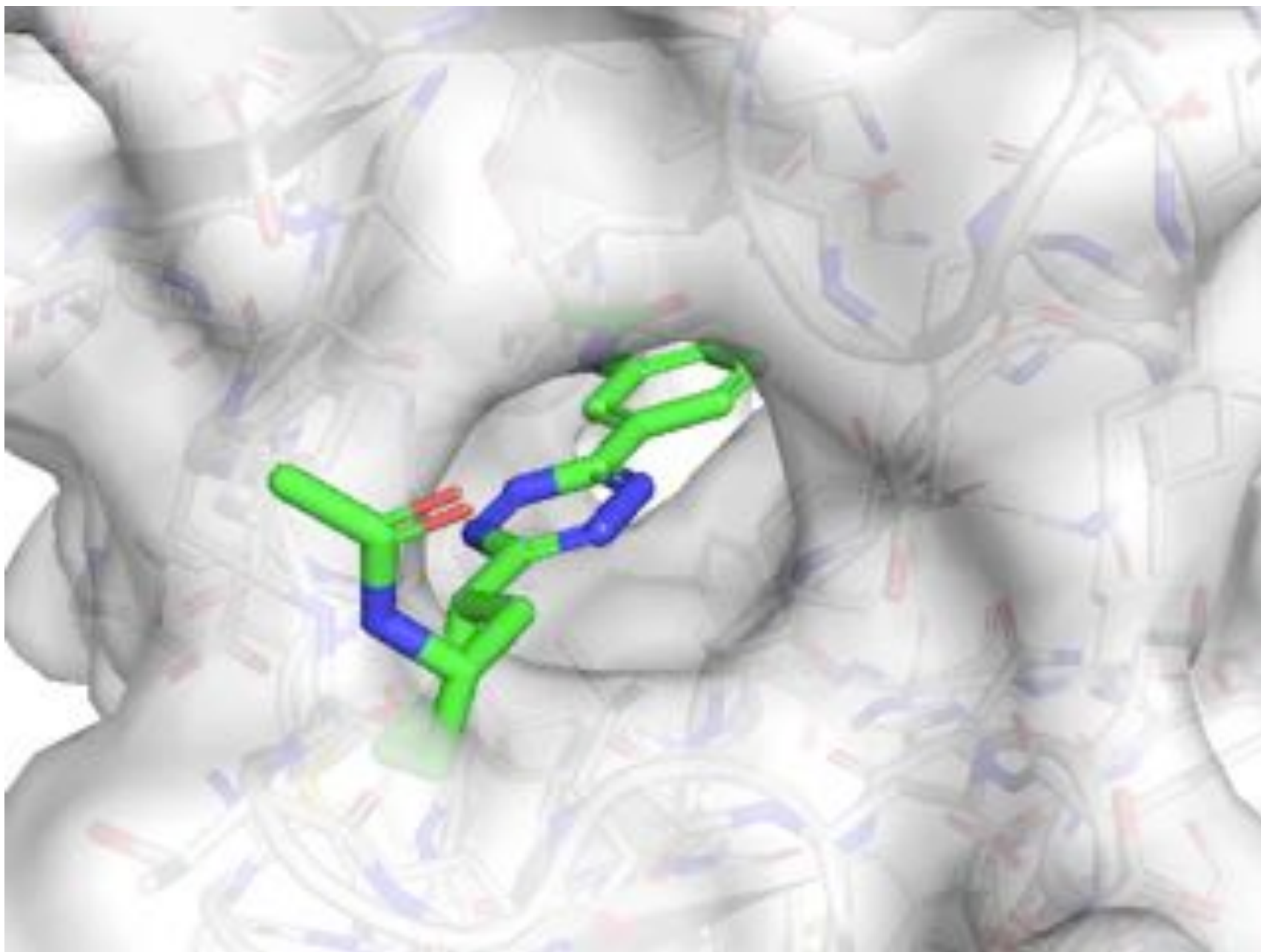




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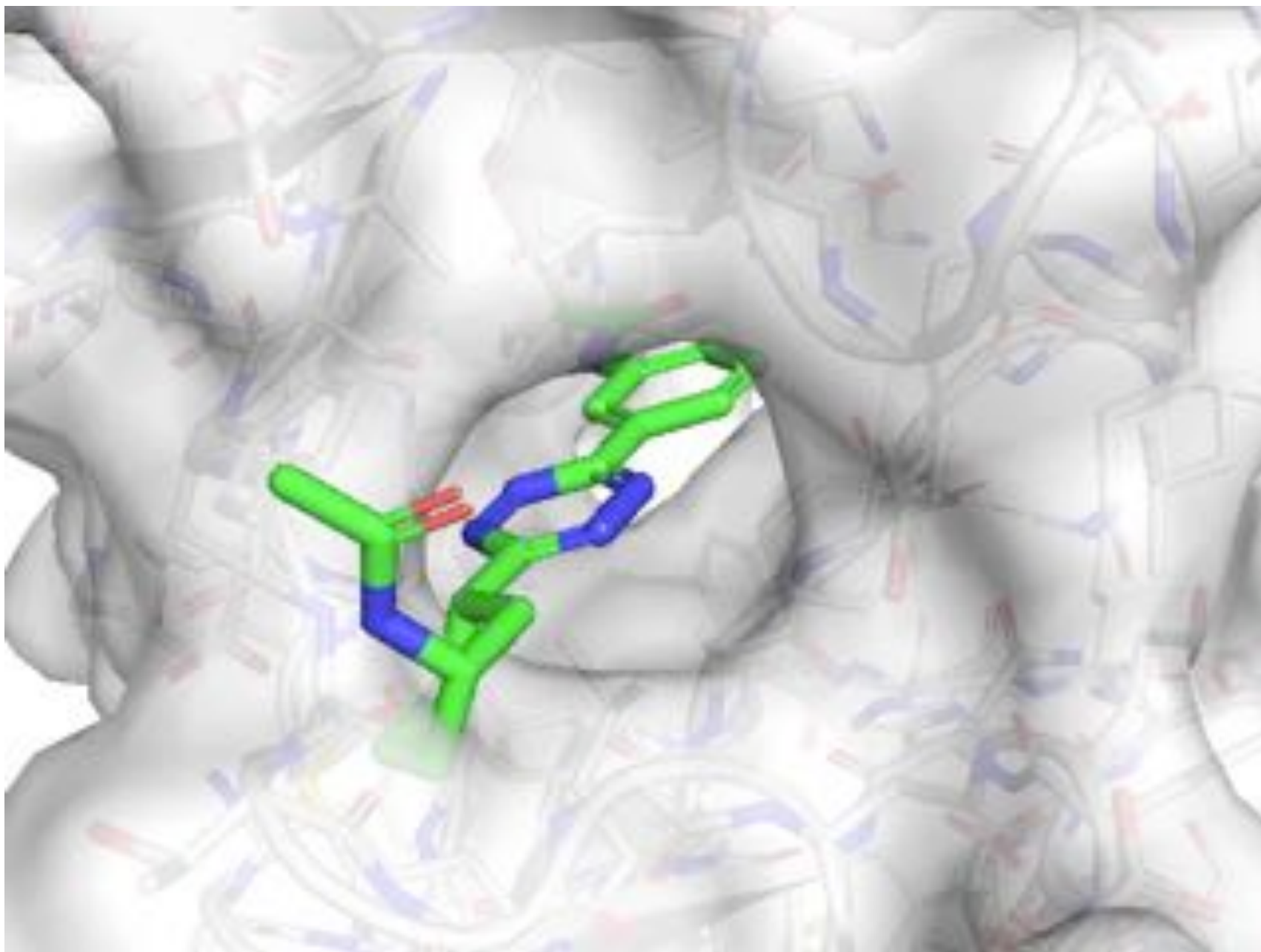


Improved energy minimization

Facilities for scoring function development

Custom format with precomputed torsion tree

Implementation of server protocol



Improved energy minimization

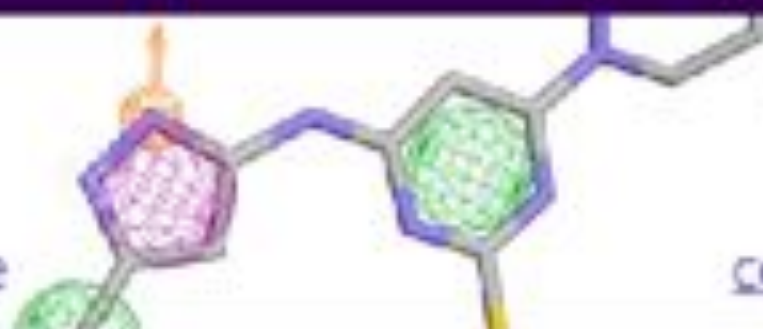
Facilities for scoring function development

Custom format with precomputed torsion tree

Implementation of server protocol

pharmit^{beta}

interactive exploration of chemical space

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binding site waters:

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create

submit your own chemical libraries

[log in to manage libraries](#)

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password:

[log in](#)

[register new account](#)

[log in as guest](#)

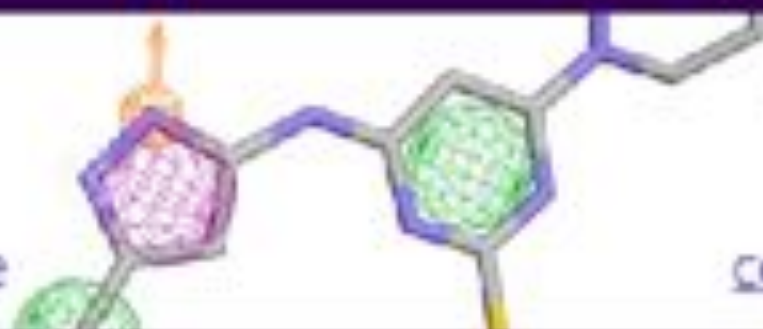
pharmit currently has

5 built-in libraries containing 1,039,705,461 conformations of 75,076,775 compounds and

7 publicly accessible user-contributed libraries containing

pharmit^{beta}

interactive exploration of chemical space

[code](#)[help](#)[contact](#)

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virtual screening in your browser

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binding site waters:

[submit](#)

[examples](#)

create

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pharmit currently has

5 built-in libraries containing 1,039,705,461 conformations of 75,076,775 compounds and

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Acknowledgements

Group Members

Jocelyn Sunseri

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Alec Helbling

Lily Turner

Aaron Zheng

Sara Amato

Lily Turner

Aaron Zheng

Gibran Biswas



Department of
Computational and
Systems Biology



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R01GM108340

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