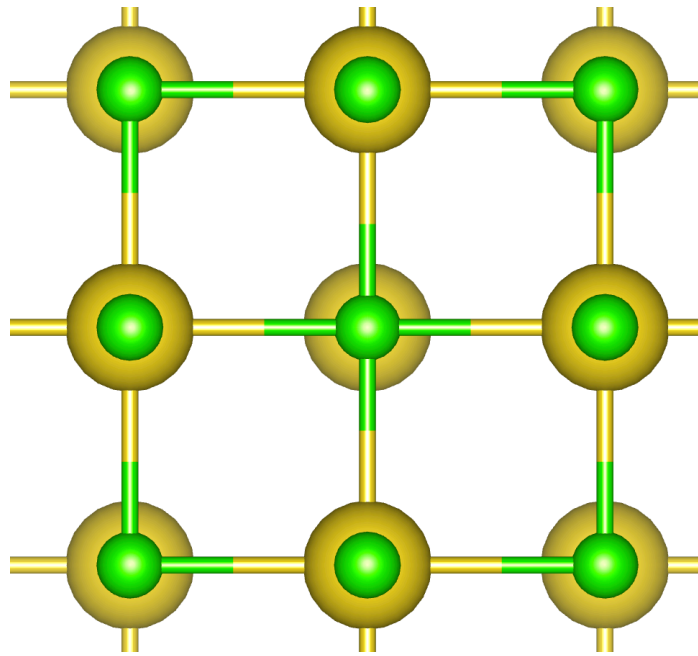


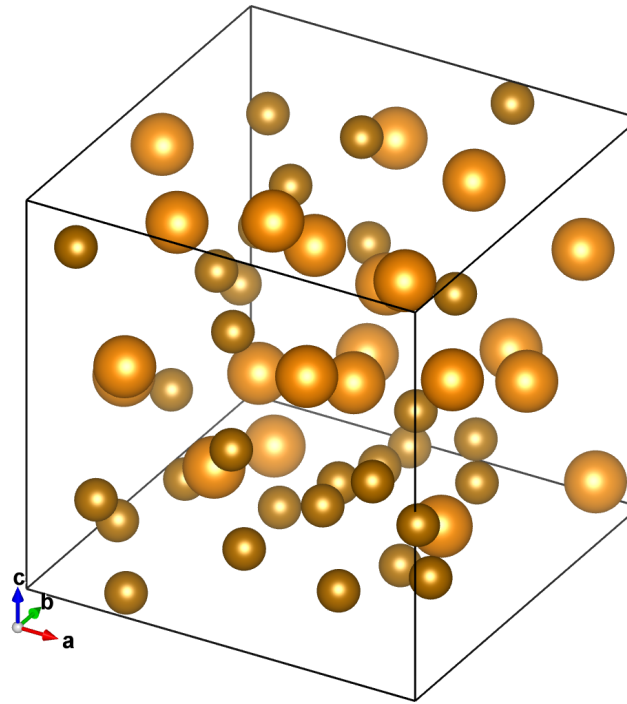
Atomic coordinates to atomic bonding graphs: exploration of data-scientific treatment

Materials Design Group
2nd Innovation Camp of
Computational Materials Science
Oct. 4th, 2018

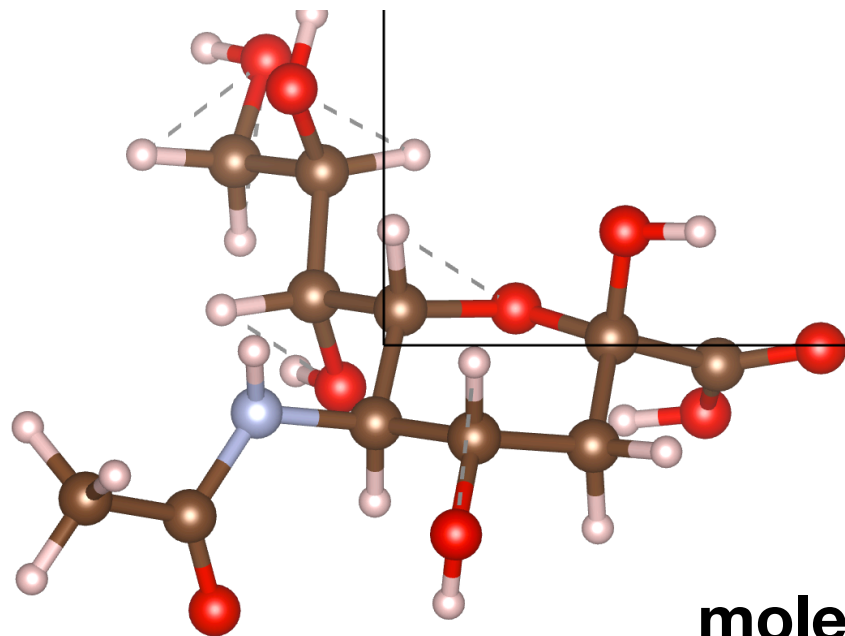
Extraction of information from atomic coordinates



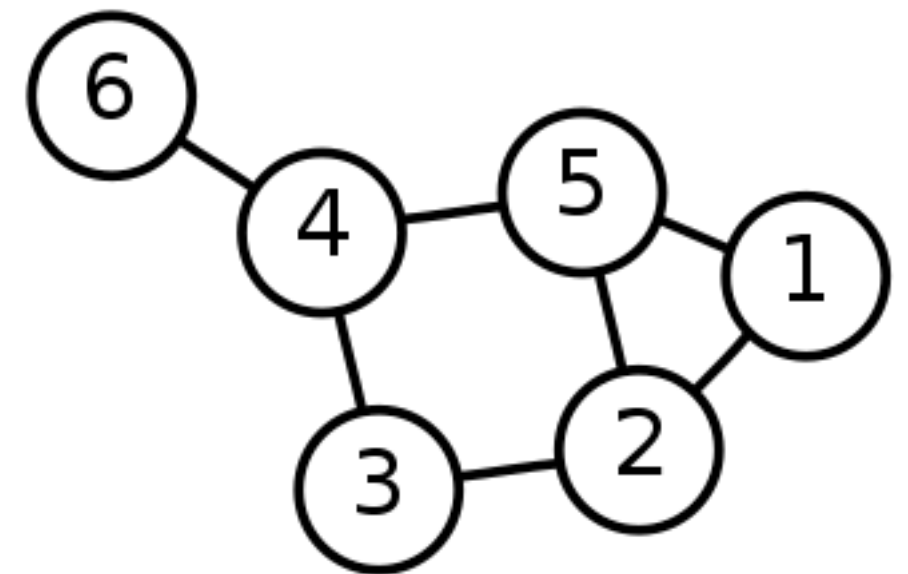
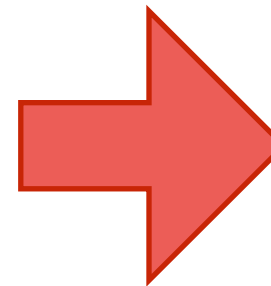
crystal?



amorphous?

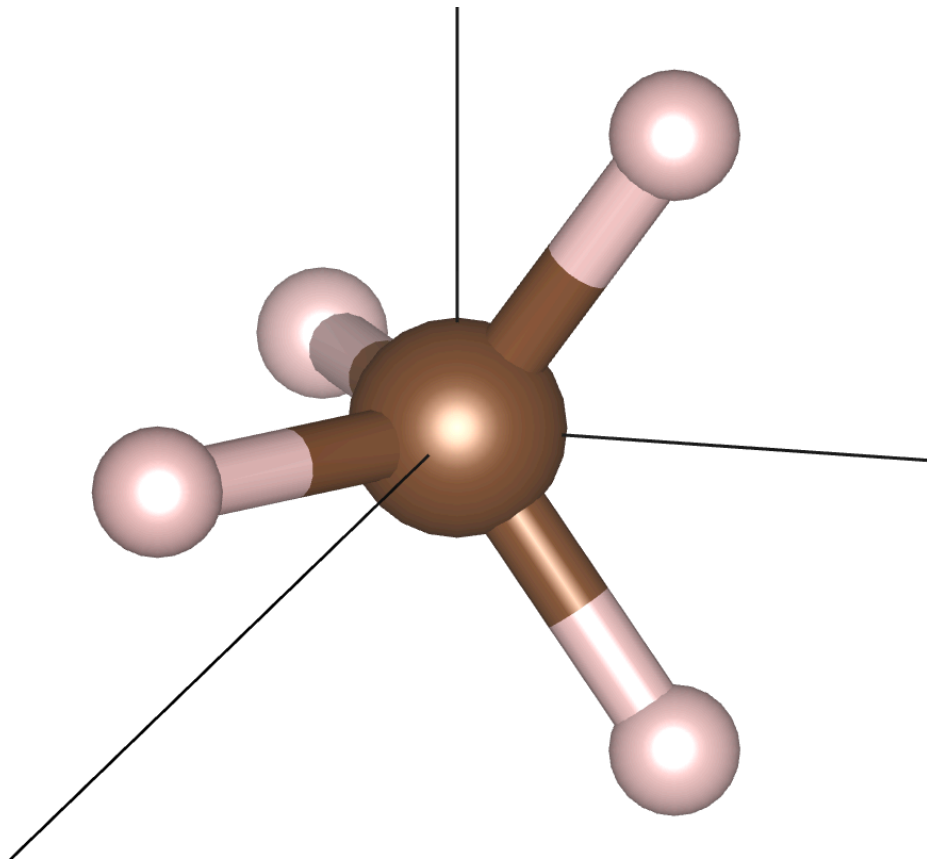


molecule?

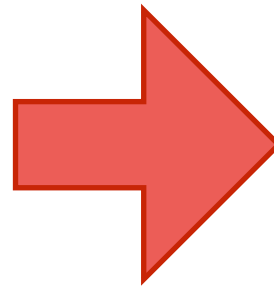


Conversion to graphs!

Atomic bonding graph



**Atomic coordinates
(sample input of OpenMX)**



1	C	2	H	1.09
1	C	3	H	1.09
1	C	4	H	1.09
1	C	5	H	1.09
.....				

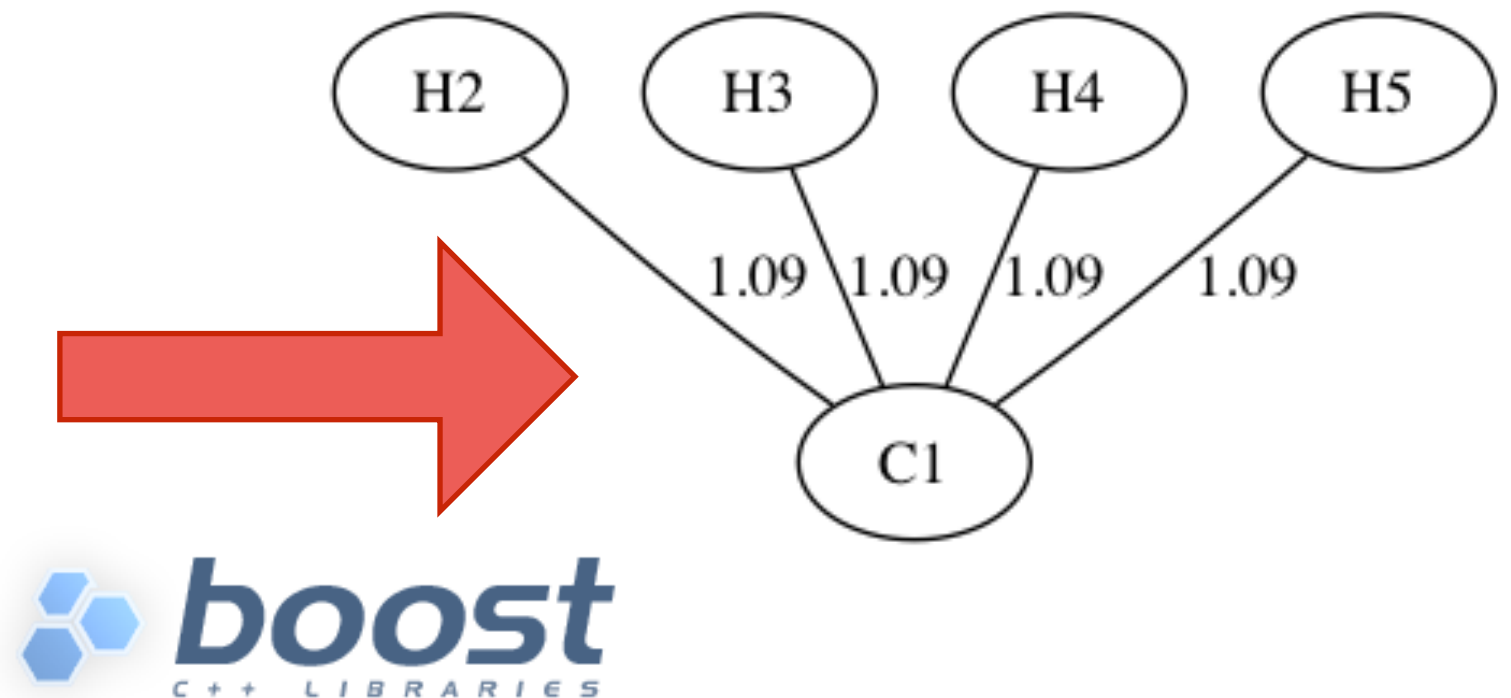
**Corresponding graphs
treatable by libraries**

- Determine atomic bonding by comparing with cut-off distances

$$d_{AB} < r_{\max}(A) + r_{\max}(B)$$

Boost Graph Library (1)

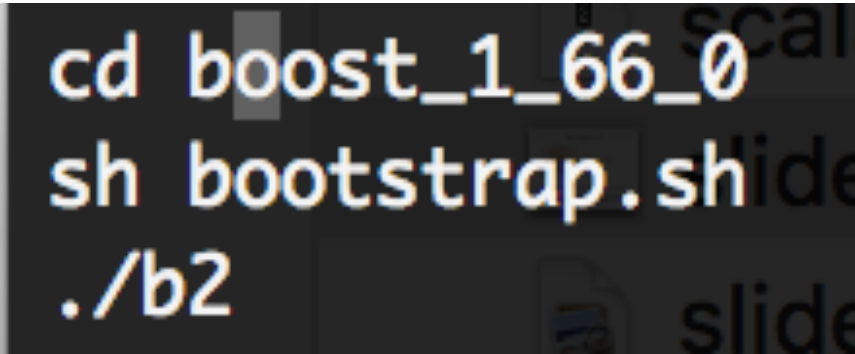
1	C	2	H	1.09
1	C	3	H	1.09
1	C	4	H	1.09
1	C	5	H	1.09
.....				



- Generate graphs from atomic coordinates and/or any physical systems
- Analyze those graphs using Boost Graph Library
- <https://www.boost.org/>
- Detailed Japanese instructions in <https://boostjp.github.io/>

Boost graph Library (2)

- Installation
- Test program (from boostjp)



```
cd boost_1_66_0
sh bootstrap.sh
./b2
```

```
#include <utility>
#include <string>
#include <boost/graph/adjacency_list.hpp>
#include <boost/graph/graph_utility.hpp>

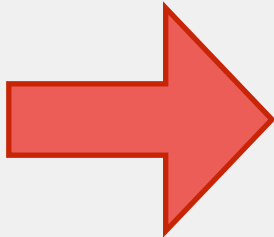
typedef boost::adjacency_list<boost::listS, boost::vecS, boost::directedS> Graph;
typedef std::pair<int, int> Edge;

enum { A, B, C, D, E, N };
const std::string name = "ABCDE";

int main()
{
    const std::vector<Edge> edges = {
        {A, B}, {A, C}, {A, D},
        {B, E}, {C, E}, {D, E}
    };

    const Graph g(edges.begin(), edges.end(), N);

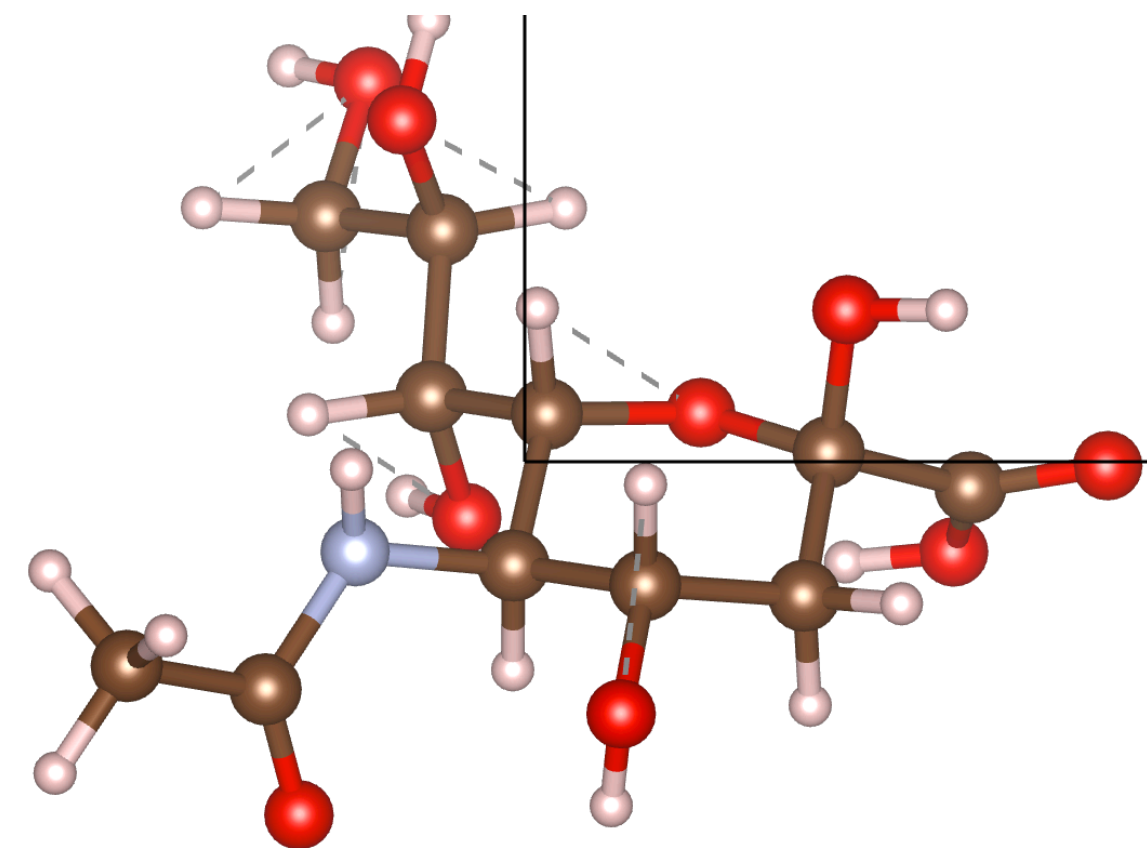
    boost::print_graph(g, name.c_str());
}
```



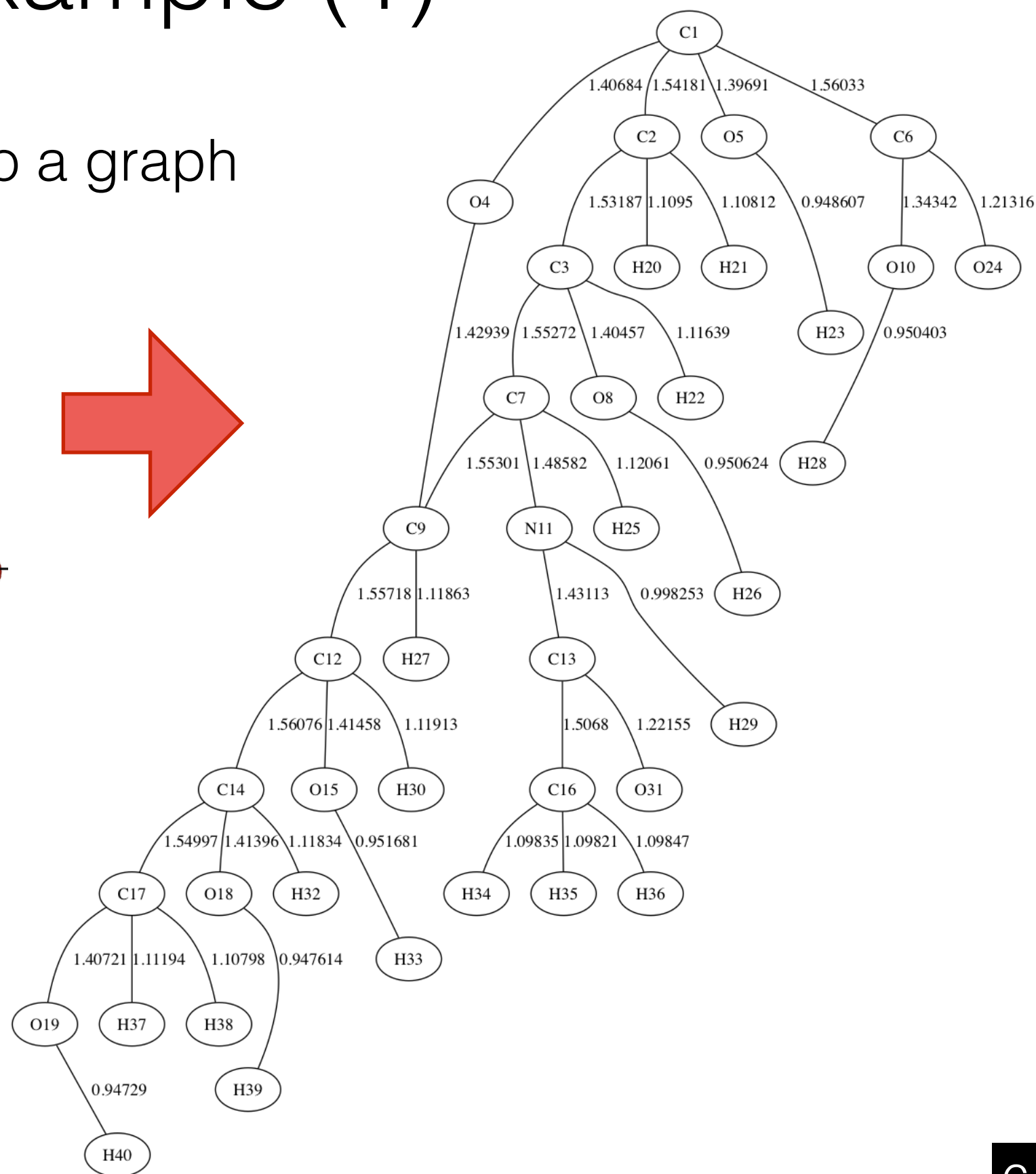
	A	B	C	D
A		-->		
B				
C				
D				
E				

Example (1)

- Convert a molecule into a graph

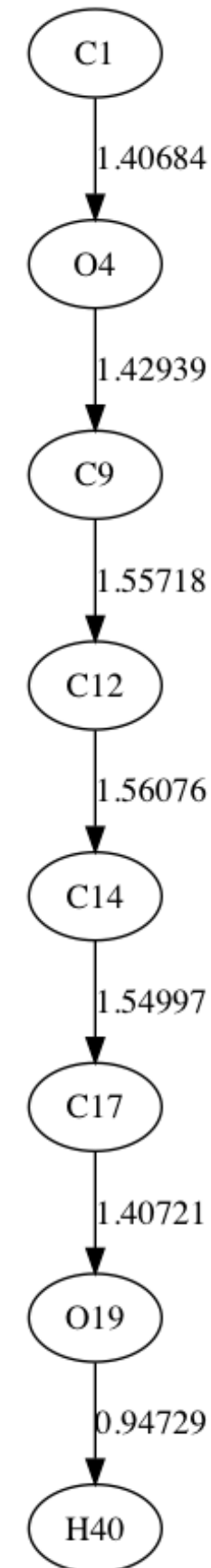
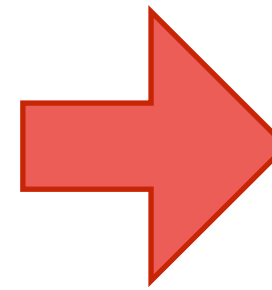
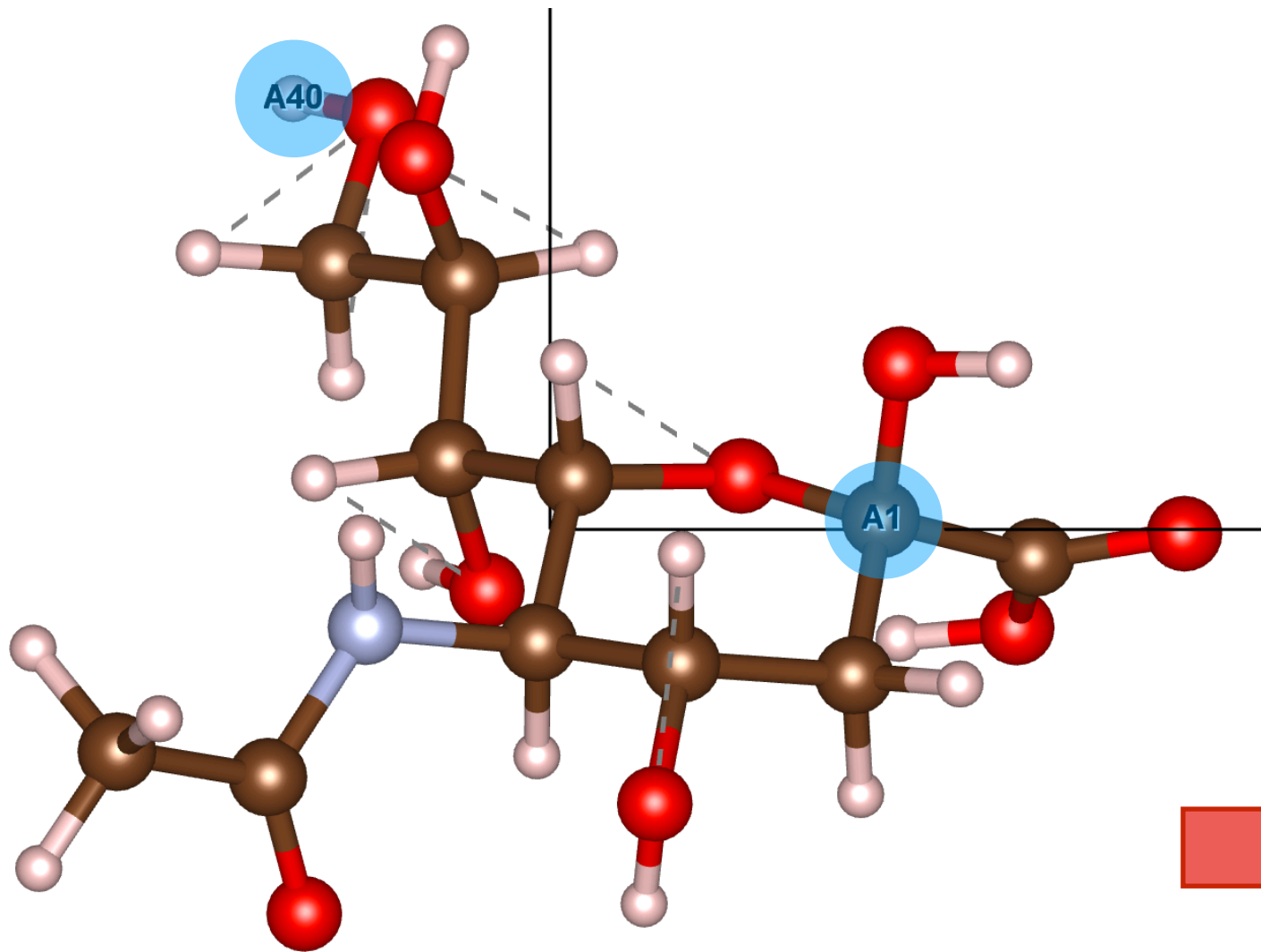


Sialic acid
(from sample input of OpenMX)



Example (2)

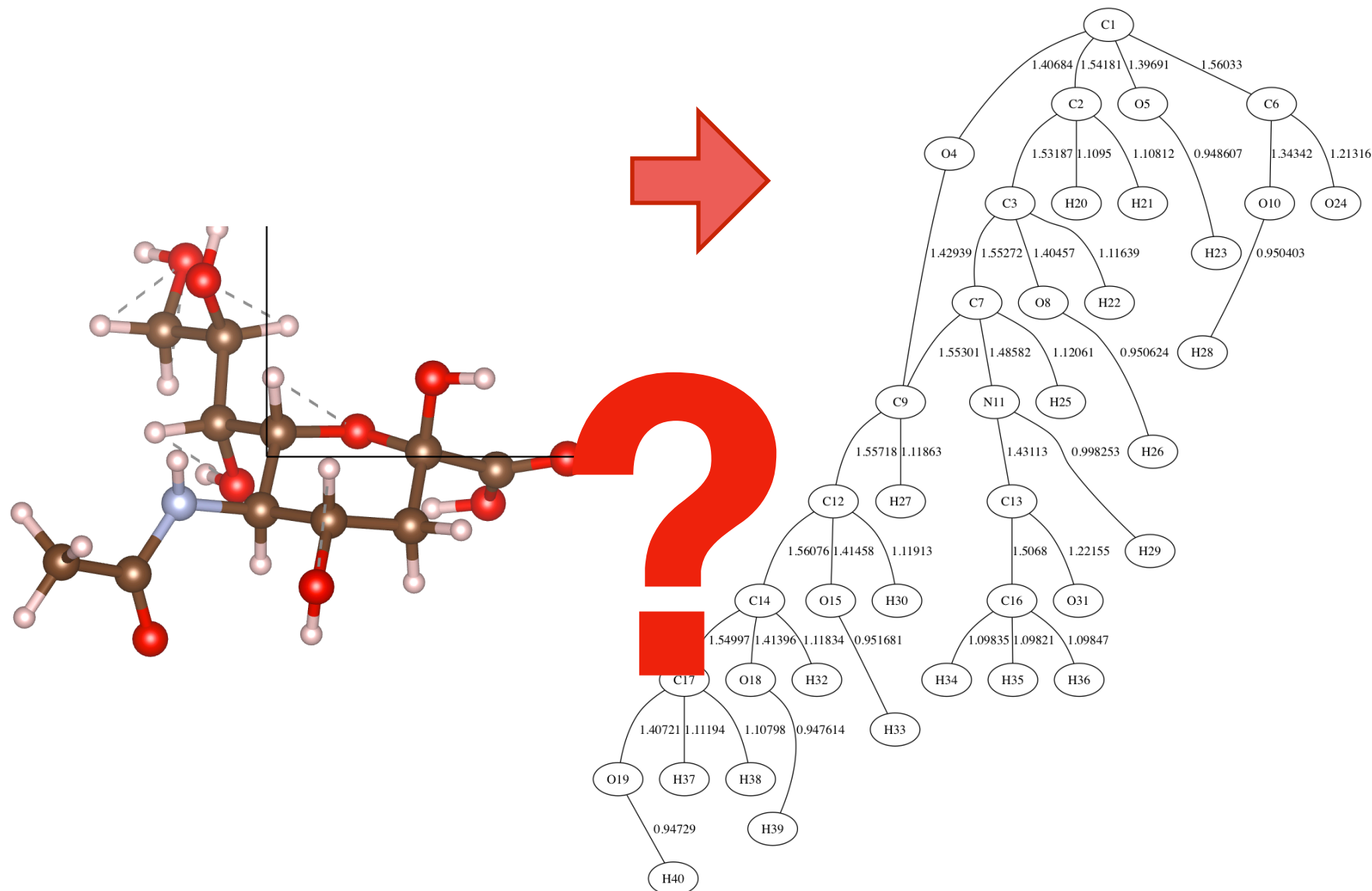
- Extraction of shortest path



**route length:
9.85865**

Our proposal...

- Prepare graph data by text file from atomic coordinates!
- Then read it by c++ program!
- Use Boost library, then we can do something... (?)



Possible application to materials design

- Analysis of **molecular conduction**?
 - Effective electron paths should be important!

