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# Data Structures and Algorithms

**COMP 103**

**2019-20**

**Semester 2**

**Lecture 08b**

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**Victoria University of Wellington**

## Assignment 2b - Core

[https://ecs.wgtn.ac.nz/Courses/XMUT103\\_2020T1/Assignment2PartB](https://ecs.wgtn.ac.nz/Courses/XMUT103_2020T1/Assignment2PartB)

### Core:

Complete the `MoleculeRender` class so that it

- Reads the data from the element-info file into a Map of Elements
- Reads the data from a molecule file into a List of `AtomS`.
- Renders the molecule from the front

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```
69 /**
70  * Reads the file "element-info.txt" which contains information about each type of element:
71  *   element name, atomic number, atomic weight, a radius, and red, green, blue, components of the color (integers)
72  * Stores the info in a Map in the elements field.
73  * The element name is the key.
74  */
75 private void readElements() {
76     /*# YOUR CODE HERE */
77 }
78
```

```
79
80 /**
81  * Asks the user to choose a file that contains info about a molecule,
82  * loads the information, resets the view angle to 0, and
83  * renders the molecule on the graphics pane.
84  * Should call other methods to do the main steps
85  */
86 public void loadAndRenderMolecule(){
87     /*# YOUR CODE HERE */
88 }
89
```

## Assignment 2b - Core

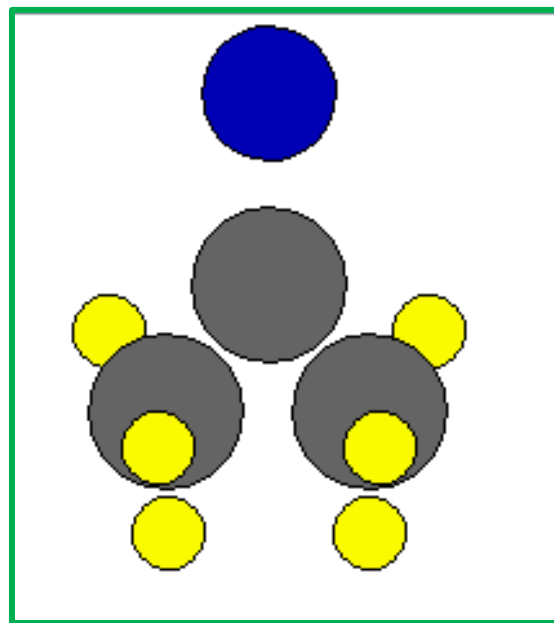
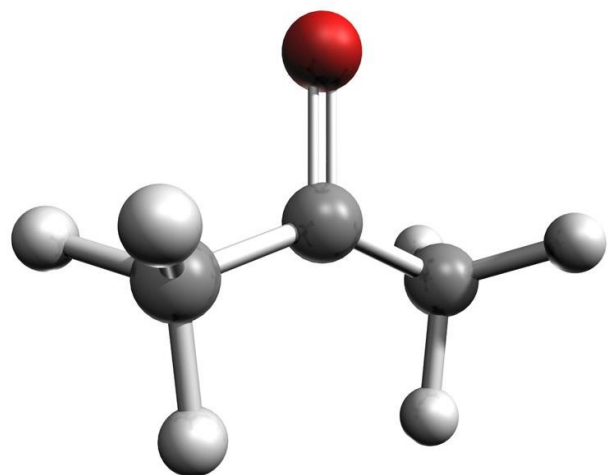
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Element name, atomic number, atomic weight, a radius, (red, green & blue values for the colour)

Example:

H	1	1.008	14	250	250	0
C	6	12.011	30	100	100	100
O	8	15.999	26	0	0	180

Acetone molecule:  $C_3H_6O$



element-info.txt - Notepad

File	Edit	Format	View	Help			
H	1	1.008	14	250	250	0	
F	9	18.998	25	60	0	140	
O	8	15.999	26	0	0	180	
N	7	14.007	27	0	180	0	
C	6	12.011	30	100	100	100	
B	5	10.811	34	40	0	160	
Be	4	9.012	35	20	0	180	
Cl	17	35.453	39	200	0	0	
I	53	126.904	80	80	0	80	
S	16	32.065	41	180	0	20	
P	15	30.974	43	160	0	40	
Br	35	79.904	44	0	60	140	
Co	27	58.933	45	20	180	0	
Ni	28	58.693	45	0	200	0	
Cr	24	51.996	46	80	120	0	
Cu	29	63.546	46	0	180	20	
Fe	26	55.845	46	40	160	0	
Mn	25	54.938	46	60	140	0	
Se	34	78.960	46	0	80	120	
Si	14	28.086	46	140	0	60	
As	33	74.922	47	0	100	100	
Ge	32	72.640	48	0	120	80	
V	23	50.942	48	100	100	0	
Al	13	26.982	49	120	0	80	
Ga	31	69.723	49	0	140	60	
Ru	44	101.070	49	0	230	0	
Zn	30	65.390	49	0	160	40	
Ti	22	47.867	51	120	80	0	
Mg	12	24.305	53	100	0	100	
Sc	21	44.956	56	140	60	0	
Na	11	22.990	61	80	0	120	
Ca	20	40.078	68	160	40	0	
Ba	56	137.327	77	0	20	180	
K	19	39.098	39	180	20	0	
Rb	37	85.468	84	0	40	160	

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```
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70  * Reads the file "element-info.txt" which contains information about each type of element:
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72  * Stores the info in a Map in the elements field.
73  * The element name is the key.
74  */
75  private void readElements() {
76      /*# YOUR CODE HERE */
77  }
78  }
```

## Assignment 2b - Core

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74  */
75  private void readElements() {
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77  }
78  
```

Code line 77 has sample Java code to read the contents of the `element-info.txt` file and store it into the `allLines` `ArrayList`

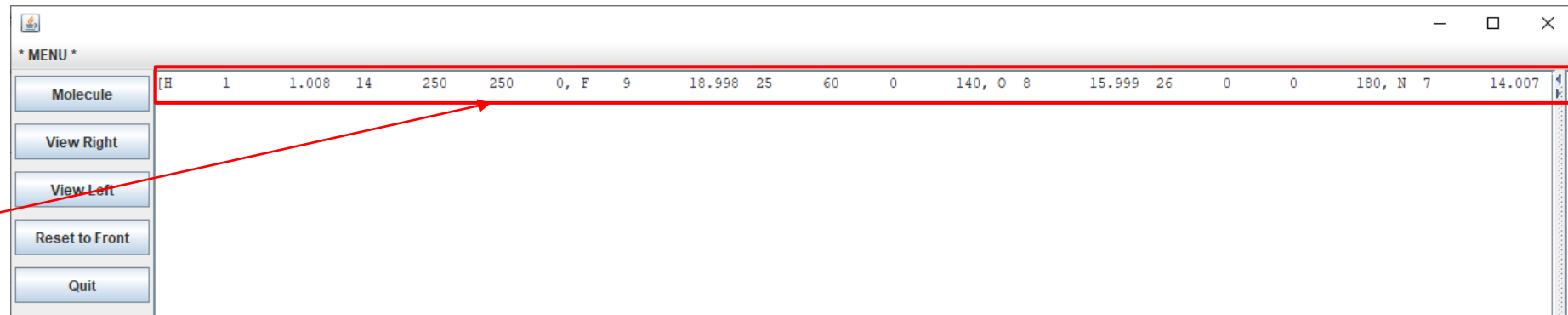
```

75  private void readElements() {
76      /*# YOUR CODE HERE */
77      ArrayList<String> allLines = readAllLines("element-info.txt");
78      UI.println(allLines);

```

Compiling and running the program, produces the shown output.

Note that the contents of the `element-info.txt` file is now stored in the `allLines` `ArrayList`. All printed on one line.



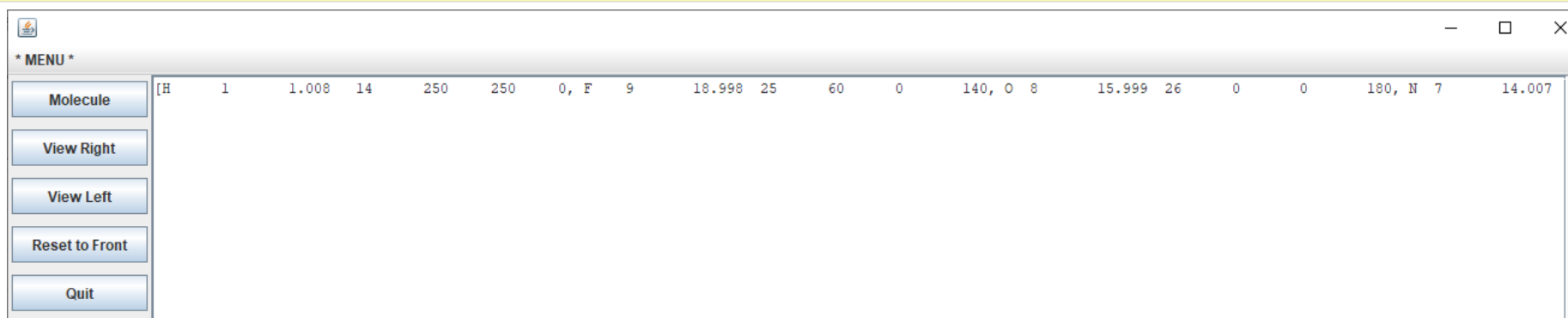
## Assignment 2b - Core

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75  private void readElements() {
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```



Element name, atomic number, atomic weight, a radius, (red, green & blue values for the colour)

Example:

H	1	1.008	14	250	250	0
C	6	12.011	30	100	100	100
O	8	15.999	26	0	0	180

## Assignment 2b - Core

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```
75 private void readElements() {  
76     /*# YOUR CODE HERE */  
77     ArrayList<String> allLines = readAllLines("element-info.txt");  
78     UI.println(allLines);  
79  
80     for (String line : allLines) {  
81         Scanner scan = new Scanner(line);  
82         String elementName = scan.next();  
83         int number = scan.nextInt();  
84         double weight = scan.nextDouble();  
85         double radius = scan.nextDouble();  
86         UI.println(elementName+" "+number+" "+weight+" "+radius);  
    }
```

\* MENU \*

Molecule

View Right

View Left

Reset to Front

Quit

[H	1	1.008	14	250	250	0	F	9	18.998	25	60	0	140, 0	8	15.999	26	0	0	180, N	7	14.007
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## Assignment 2b - Core

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```
27 // Map containing info about each type of element.  
28 private Map<String, Element> elements;  
29
```

```
75 private void readElements() {  
76     /*# YOUR CODE HERE */  
77     ArrayList<String> allLines = readAllLines("element-info.txt");  
78     UI.println(allLines);  
79     elements = new HashMap<String, Element>();  
80     for (String line : allLines) {  
81         Scanner scan = new Scanner(line);  
82         String elementName = scan.next();  
83         int number = scan.nextInt();  
84         double weight = scan.nextDouble();  
85         double radius = scan.nextDouble();  
86         UI.println(elementName+" "+number+" "+weight+" "+radius);  
87     }  
88 }
```

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75  private void readElements() {
76      /*# YOUR CODE HERE */
77  }
78  }
```

```
80  /**
81  * Asks the user to choose a file that contains info about a molecule,
82  * loads the information, resets the view angle to 0, and
83  * renders the molecule on the graphics pane.
84  * Should call other methods to do the main steps
85  */
86  public void loadAndRenderMolecule(){
87      /*# YOUR CODE HERE */
88  }
89  }
```

## Assignment 2b - Core

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```
/**
 * Asks the user to choose a file that contains info about a molecule,
 * loads the information, resets the view angle to 0, and
 * renders the molecule on the graphics pane.
 * Should call other methods to do the main steps
 */
public void loadAndRenderMolecule(){
    /*# YOUR CODE HERE */
    String fn = (UIFileChooser.open());
    ArrayList<String> allLines = readAllLines(fn);
    UI.println(allLines);
}
```

Ask user to choose a file containing info about a molecule from the list of files shown

Name

- aa.txt
- Atom.java
- Element.java
- element-info.txt
- molecule0.txt
- molecule1.txt
- molecule2.txt
- molecule3.txt
- molecule4.txt
- molecule5.txt
- molecule6.txt
- molecule7.txt
- molecule8.txt
- molecule8-with-bonds.txt
- molecule9.txt
- molecule9-with-bonds.txt
- molecule10.txt
- molecule10-with-bonds.txt
- MoleculeRenderer.java
- moleculeTest.txt
- package.bluej

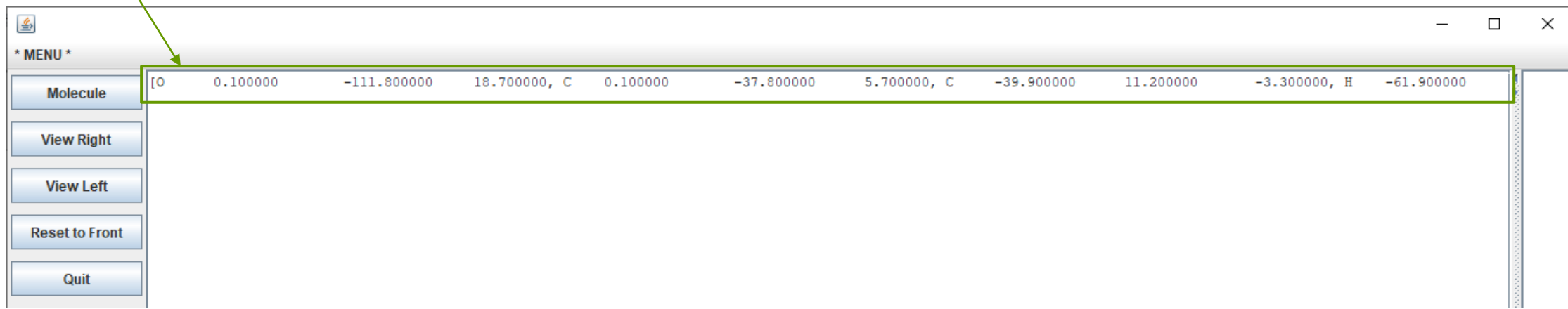
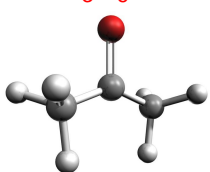
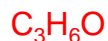
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```
/**
 * Asks the user to choose a file that contains info about a molecule,
 * loads the information, resets the view angle to 0, and
 * renders the molecule on the graphics pane.
 * Should call other methods to do the main steps
 */
public void loadAndRenderMolecule(){
    /*# YOUR CODE HERE */
    String fn = (UIFileChooser.open());
    ArrayList<String> allLines = readAllLines(fn);
    UI.println(allLines);
}
```

Use the readAllLines method (utility method given at the end of the original moleculeRenderer.java file) to open the chosen file fn and store contents of the file into an ArrayList named allLines

Acetone molecule



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```
/**
 * Asks the user to choose a file that contains info about a molecule,
 * loads the information, resets the view angle to 0, and
 * renders the molecule on the graphics pane.
 * Should call other methods to do the main steps
 */
public void loadAndRenderMolecule(){
    /*# YOUR CODE HERE */
    String fn = (UIFileChooser.open());
    ArrayList<String> allLines = readAllLines(fn);
    UI.println(allLines);
}
```

Reset the view angle to 0

```
33 private double horizViewAngle = 0; // The current horizontal viewing angle ("Yaw")
34 // reset it to 0 when loading a new molecule!
```

## Assignment 2b - Core

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```
/**
 * Asks the user to choose a file that contains info about a molecule,
 * loads the information, resets the view angle to 0, and
 * renders the molecule on the graphics pane.
 * Should call other methods to do the main steps
 */
public void loadAndRenderMolecule(){
    /*# YOUR CODE HERE */
    String fn = (UIFileChooser.open());
    ArrayList<String> allLines = readAllLines(fn);
    UI.println(allLines);
    this.horizViewAngle = 0; // reset to 0 before drawing
    this.vertViewAngle = 0;
```

Reset the view angle to 0

```
for (String line : allLines) {  
    Scanner scan = new Scanner(line);  
    String atomType = scan.next();  
    UI.println(atomType);  
    Element element = elements.get(atomType);  
  
    double x = scan.nextDouble();  
    double y = scan.nextDouble();  
    double z = scan.nextDouble();  
  
    Atom atom = new Atom(x, y, z, element);  
    molecule.add(atom);  
}  
showFromFront();
```

Type the given Java code after the reset code lines for the `horizViewAngle` and the `vertViewAngle` as shown in the previous slide.



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```
for (String line : allLines) {
    Scanner scan = new Scanner(line);
    String atomType = scan.next();
    UI.println(atomType);
    Element element = elements.get(atomType);

    double x = scan.nextDouble();
    double y = scan.nextDouble();
    double z = scan.nextDouble();

    Atom atom = new Atom(x, y, z, element);
    molecule.add(atom);
}
showFromFront();
```

You need to add the Java code to ensure that the showFromFront method renders the molecule from the front

```
/**
 * Renders the molecule from the front.
 * Sorts the Atoms in the List by their z value, back to front
 * Uses the default ordering of the Atoms
 * Then renders each atom
 */
public void showFromFront() {
    /*# YOUR CODE HERE */

    UI.clearGraphics();
    for(Atom atom : molecule) {
        atom.render(MIDX, MIDY, 0, 0);
    }
}
```

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Only 1 line of Java code  
is required in the  
showFromFront method.  
**Hint**

```
/**
 * Renders the molecule from the front.
 * Sorts the Atoms in the List by their z value, back to front
 * Uses the default ordering of the Atoms
 * Then renders each atom
 */
public void showFromFront() {
    /*# YOUR CODE HERE */

    UI.clearGraphics();
    for(Atom atom : molecule) {
        atom.render(MIDX, MIDY, 0, 0);
    }
}
```

**Outcome** for the  
molecule0.txt file

