

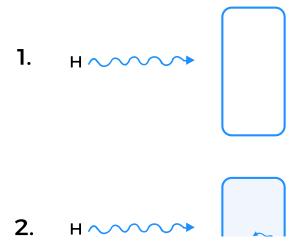
This problem simulated the creation of water molecule by grouping three threads representing Hydrogen and Oxygen atoms.

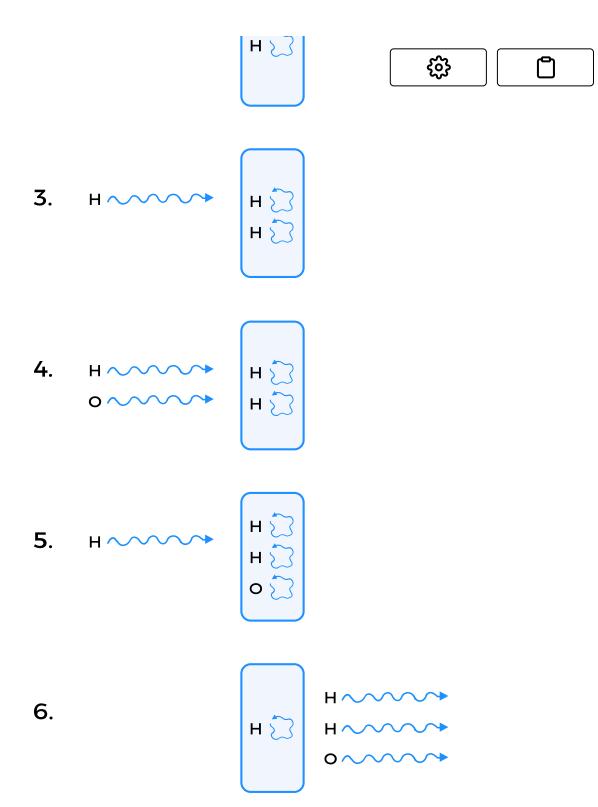
We'll cover the following ^

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Problem Statement#

Suppose we have a machine that creates molecules by combining atoms. We are creating water molecules by joining one Oxygen and two Hydrogen atoms. The atoms are represented by threads. The machine will wait for the required atoms (threads), then group one Oxygen and two Hydrogen threads to simulate the creation of a molecule. The molecule then exists the machine. You have to ensure that one molecule is completed before moving onto the next molecule. If more than the required number of threads arrive, they will have to wait. The figure below explains the working of our machine:





Two Hydrogen threads are admitted in the machine as they arrive but when the third thread arrives in step 3, it is made to wait. When an Oxygen thread arrives in step 4, it is allowed to enter the machine. A water molecule is formed in step 5 which exists the machine in step 6. That is when the waiting Hydrogen thread is notified and the process of creating more molecules continues. The threads can arrive in any order which means that HHO, OHH and HOH are all valid outputs.

The code for the class is as follows:





```
class H20Machine {
   public H20Machine() {
   }
   public void HydrogenAtom() {
   }
   public void OxygenAtom() {
   }
}
```

The input to the machine can be in any order. Your program should enforce a 2:1 ratio for Hydrogen and Oxygen threads, and stop more than the required number of threads from entering the machine.

Solution#

Our molecule making machine is represented by the class H20Machine, which contains two main methods; HydrogenAtom() and OxygenAtom().

The problem is solved by using basic utility functions like notify() and wait(). The class consits of 3 private members: sync for synchronization, molecule which is a string array with a capacity of 3 elements (atoms) and count to store the current index of the molecule array.





```
class H20Machine {
   Object sync;
   String[] molecule;
   int count;

public H20Machine() {
   }

public void HydrogenAtom() {
   }

public void 0xygenAtom() {
   }
}
```

The constructor initializes the molecule array with a capacity of 3 atoms and the integer count is initialized with 0.

```
public H20Machine() {
    molecule = new String[3];
    count = 0;
    sync = new Object();
}
```

For synchronization purpose, the entire logic of HydrogenAtom() is wrapped in sync. First of all, we check the frequency of Hydrogen atom in the molecule array by using frequency() function found in the **Collections** library of Java. The function deals the array molecule as an ArrayList and checks the count of Hydrogen atoms in it. If the array has reached its capacity of 2 Hydrogen atoms, then the thread should wait for space in a new molecule. If the frequency of Hydrogen is less than 2 it means space is available in the current molecule. Hence, H is placed in the array and count is incremented. So far, the code of HydrogenAtom() is as follows:





```
public void HydrogenAtom() {
    synchronized (sync) {

        // if 2 hydrogen atoms already exist
        while (Collections.frequency(Arrays.asList(molecule), "H") =
        = 2) {
            sync.wait();
        }

        molecule[count] = "H";
        count++;
    }
}
```

In case molecule is full and count is 3, then print the molecule and exit the machine. The array molecule is reset (initialized with null) and count goes back to 0 for a new molecule to be built. At the end of the method, the waiting threads (atoms) are notified using notifyAll(). The complete code for HydrogenAtom() is given below:





```
public void HydrogenAtom() {
    synchronized (sync) {
        // if 2 hydrogen atoms already exist
        while (Collections.frequency(Arrays.asList(molecule),"H")
== 2) {
            sync.wait();
        }
        molecule[count] = "H";
        count++;
        // if molecule is full, then exit.
        if(count == 3) {
            for (String element: molecule) {
                System.out.print(element);
            }
            Arrays.fill(molecule, null);
            count = 0;
        }
        sync.notifyAll();
    }
}
```

The second method <code>OxygenAtom()</code> is the same as <code>HydrogenAtom()</code> with the only difference of the atom frequency check in the array <code>molecule</code>. If it contains one Oxygen atom, then the calling thread goes into <code>wait()</code>. If the count of Oxygen atom is not equal to 1 in the <code>molecule</code>, then an Oxygen atom "O" is placed in the next available space. The complete code of <code>OxygenAtom()</code> is shown below:





```
public void oxygen() {
    synchronized (sync) {
        // if 1 oxygen atom already exists
        while (Collections.frequency(Arrays.asList(molecule),"0")
== 1) {
            sync.wait();
        }
        molecule[count] = "0";
        count++;
        // if molecule is full, then exit.
        if(count == 3) {
            for (String element: molecule) {
                System.out.print(element);
            Arrays.fill(molecule, null);
            count = 0;
        }
        sync.notifyAll();
    }
}
```

The complete code for the solution is as follows:

```
class H20Machine {
    Object sync;
    String[] molecule;
    int count;
    public H20Machine() {
        molecule = new String[3];
        count = 0;
        sync = new Object();
    }
    public void HydrogenAtom() {
        synchronized (sync) {
            // if 2 hydrogen atoms already exist
            while (Collections.frequency(Arrays.asList(molecule),
"H") == 2) {
                sync.wait();
            }
            molecule[count] = "H";
            count++;
            // if molecule is complete, then exit.
            if(count == 3) {
                for (String element: molecule) {
                    System.out.print(element);
                }
                Arrays.fill(molecule, null);
                count = 0;
            sync.notifyAll();
        }
    }
    public void OxygenAtom() throws InterruptedException {
        synchronized (sync) {
```

```
(3)
            // if 1 oxygen atom already exists
            while (Collections.frequency(Arrays.asList(molecule),
"0") == 1) {
                sync.wait();
            }
            molecule[count] = "0";
            count++;
            // if molecule is complete, then exit.
            if(count == 3) {
                for (String element: molecule) {
                    System.out.print(element);
                }
                Arrays.fill(molecule,null);
                count = 0;
            }
            sync.notifyAll();
        }
    }
}
```

We will be creating another class H20MachineThread for multi-threading purpose. It takes an object of H20Machine and calls the relevant method from the string passed to it.





```
class H20MachineThread extends Thread {
    H20Machine molecule;
    String atom;
    public H20MachineThread(H20Machine molecule, String atom){
        this.molecule = molecule;
        this.atom = atom;
    }
    public void run() {
        if ("H".equals(atom)) {
            try {
                molecule.HydrogenAtom();
            catch (Exception e) {
        else if ("0".equals(atom)) {
            try {
                molecule.OxygenAtom();
            catch (Exception e) {
        }
    }
}
```

We will now be creating 4 threads in order to test our proposed solution. Same object of H20Machine is passed to the 4 threads: **t1**, **t2**, **t3** and **t4**.

t1 and t3 act as Hydrogen atoms trying to enter the machine where as t2 and t4 act as Oxygen atoms. It can be seen from the output that only 1 molecule of H2O exits the machine while the extra Oxygen atom is not utilized.





```
import java.util.Arrays;
import java.util.Collections;
class H20Machine {
    Object sync;
    String[] molecule;
    int count;
    public H20Machine() {
        molecule = new String[3];
        count = 0;
        sync = new Object();
   }
    public void HydrogenAtom() {
        synchronized (sync) {
            // if 2 hydrogen atoms already exist
            while (Collections.frequency(Arrays.asList(molecule),"H") == 2) {
                try {
                    sync.wait();
                catch (Exception e) {
            }
            molecule[count] = "H";
            count++;
            // if molecule is complete, then exit.
            if(count == 3) {
                for (String element: molecule) {
                    System.out.print(element);
                Arrays.fill(molecule, null);
                count = 0;
            sync.notifyAll();
        }
    }
   public void OxygenAtom() throws InterruptedException {
        synchronized (sync) {
            // if 1 oxygen atom already exists
            while (Collections.frequency(Arrays.asList(molecule),"0") == 1) {
                try {
                    sync.wait();
                catch (Exception e) {
```

```
\mathcal{E}_{\mathcal{E}}
```



```
molecule[count] = "0";
            count++;
            // if molecule is complete, then exit.
            if(count == 3) {
                for (String element: molecule) {
                    System.out.print(element);
                Arrays.fill(molecule, null);
                count = 0;
            sync.notifyAll();
        }
    }
}
class H20MachineThread extends Thread {
    H20Machine molecule;
    String atom;
    public H20MachineThread(H20Machine molecule, String atom){
        this.molecule = molecule;
        this.atom = atom;
    }
    public void run() {
        if ("H".equals(atom)) {
            try {
                molecule.HydrogenAtom();
            catch (Exception e) {
            }
        else if ("0".equals(atom)) {
            try {
                molecule.OxygenAtom();
            catch (Exception e) {
        }
    }
}
public class Main
        public static void main(String[] args) {
                H20Machine molecule = new H20Machine();
                Thread t1 = new H20MachineThread(molecule,"H");
            Throad +2 - now U20MachinaThroad/malacula U0U).
```

}

```
Thread t3 = new H20MachineThread(molecule, "O");
Thread t4 = new H20MachineThread(molecule, "O");

t2.start();
t1.start();
t4.start();
t3.start();
}

}
```

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