Multiple sequence Alignment

```
import java.io.*;
import java.util.*;
public class Consensus
public static void main(String str[]) throws IOException
{
int n, i,j,k,count;
String seq[],cons[];
ArrayList<Integer> a = new ArrayList<Integer>();
ArrayList s = new ArrayList();
BufferedReader br=new BufferedReader(new InputStreamReader(System.in));
System.out.println("Enter the no of Sequences");
n=Integer.parseInt(br.readLine());
seq=new String[n];
System.out.println("Enter sequences");
for(i=0;i<n;i++)
seq[i]=br.readLine();
cons=new String[seq[0].length()];
for(j=0;j<seq[0].length();j++)</pre>
cons[j]=" ";
for(j=0;j < seq[0].length();j++)
{
a.clear();
s.clear();
for(i=0;i<n;i++)
{
count=1;
```

```
for(k=i+1;k< n;k++)
{
if(seq[i].charAt(j)==seq[k].charAt(j))
count++;
}
System.out.println("count="+count);
a.add(count);
s.add(seq[i].charAt(j));
}
/**Updated Snippet 1**/
Set<String> set = new HashSet<>(s);
ArrayList setlist = new ArrayList(set);
Collections.sort(setlist);
if (setlist.contains('-') &&setlist.size()==2){
cons[j]+="-"+setlist.get(1);
}
else if (setlist.size()==1){
cons[j]+="-"+setlist.get(0);
}
else{
int m = Collections.max(a);
int index=a.indexOf(m);
System.out.println("Max="+m);
cons[j]+=s.get(index);
System.out.println("index="+index);
for(i=index+1;i<a.size();i++)</pre>
{
```

```
if(a.get(i)==m)
cons[j]+="/"+s.get(i);
}
}
}
System.out.println("Consensus=");
for(j=0;j \le eq[0].length();j++){
/**Updated Snippet 2**/
if(cons[j].length()==2)
System.out.print(cons[j].toLowerCase());
else if(cons[j].length()==3)
System.out.print(cons[j].replace("-",""));
else
System.out.print(cons[j]);
}
}
OUTPUT:
```

```
Enter the no of Sequences

Sequences

Enter sequences

ACTG

ICGA

TATG

TATG

Count=1

count=2

count=1

fax=3

count=1

count=1

fax=3

count=1

count=1

fax=3

count=1

count=1

count=1

fax=3

count=1

count=1

count=1

count=1

count=1

count=1

fax=3

count=1

count=
```