Name:

Instructions

- 1. Write your name at the top of the *first* page and your initials at the bottom of *every* page.
- 2. Do not staple the exam.
- 3. Return the exam with all the pages, arranged in ascending order.
- 4. This is a closed-book exam.
- 5. No electronic devices are permitted.
- 6. You may use the blank spaces for any scratch work.
- 7. Discussing the exam before the solutions have been posted is a violation of the Honor Code.
- 8. There are 11 problems on this exam and you have 75 minutes to answer them.
- 9. Problems 1-12 involve 19 multiple-choice questions, each worth 4 points. Each question must have *exactly one* response clearly marked in the circle provided or else your answer will be considered incorrect.
- 10. Problems 13 (worth 16 points) and 14 (worth 8 points), must be answered clearly in the boxed space provided for those problems.

Problem 1. Consider the following method:

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Problem 2. Consider the following recursive method:
<pre>public static int mystery(int a, int b) { return (b == 0) ? a : mystery(b, a % b); }</pre>
a. What does mystery(27, 72) return?
<u> </u>
9
45
1
72
b. What does mystery(48, 15) return?
48
<u> </u>
1
33
3
c. What does mystery() compute and return in general?
\bigcirc b
\bigcirc Largest number that divides both a and b
\bigcirc Smallest number that divides both a and b
$\bigcap a-b $
\bigcirc a
Problem 3. Consider the following recursive methods::

return (x == null) ? 0 : 1 + f(x.next);

return (x == null) ? 0 : g.item + g(x.next);

public static int f(Node x) {

public static int g(Node x) {

}

}

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a. What	does f(a) return, where a is a linked list containing the items 1, 1, 2, 3, 5, 8, and 13 and ir order?
	7
	33
	1
	13
	0
b. What	does g(a) return, where a is a linked list containing the items 1, 1, 2, 3, 5, 8, and 13 and ir order?
	7
	1
	0
	33
	13
Probler	m 4. Consider the following program Mystery.java:
pub	<pre>class Mystery { lic static void main(String[] args) { String x = StdIn.readString(); String y = StdIn.readString(); StdOut.print(x + y); StdOut.print(" "); StdOut.print(y + x); StdOut.println();</pre>
}	
·	ppose that the file input.txt contains the two strings AB and BA separated by a space. What following command output?
\$ java	Mystery < input.txt java Mystery java Mystery
	AB BA
	ABBA BAAB
	ABBABAABBAABBA BAABABBAABBABAAB
	ABBABAABBAABABABAABBAABBAABB BAABABBAABBAABBABAABBAABBAABBA
	ABBABAAB BAABABBA

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Problem 5. Consider the following table, which gives the running time T(N) in seconds for a program for various values of the input size N:

N	T(N)
1000	3
2000	12
4000	48
8000	192

What	is	the	order	$\circ f$	growth	$\circ f$	T	(N)	17
vvnat	19	une	oraer	OI	growth	OΙ	1	. ∠ V .	<i>)</i> :

	Quadratic
	Quadranc

· \	т •
)	Linear

Exponential

Linearithmic

Cubic

Problem 6. What is the order of growth (T(N)) of the following code fragment?

```
int sum = 0;
for (int i = 0; i < N; i++) {
    for (int j = 0; j < 100; j++) {
        for (int k = 0; k < 1000; k++) {
            sum++;
        }
    }
}</pre>
```

() Cubic

() Linear

Quadratic

Exponential

Linearithmic

Problem 7. Consider a data type T with two instance variables: int x and double y. Ignoring array and object overheads, what is the memory footprint (in bytes) of the array a[] created and initialized as follows?

```
T[] a = new T[100];
for (int i = 0; i < 100; i++) {
    T[i] = new T();
}</pre>
```

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		100
		1200
		12
		400
		800
Probl	en	8. Consider the following methods:
Itera Q f	tor uei	<pre>c<character> f(String s) { ne<character> Q = new Queue<character>(); (int i = 0; i < s.length(); i++) { Q.enqueue(s.charAt(i)); } nr Q.iterator();</character></character></character></pre>
S f	tao or	<pre>c<character> g(String s) { ck<character> S = new Stack<character>(); (int i = 0; i < s.length(); i++) { S.push(s.charAt(i)); } urn S.iterator();</character></character></character></pre>
a. Wh	at	is the value returned by the method call f("alice").next()?
)	'e'
		'1'
		'c'
		'a'
)	'i'
b. Wh	at	is the value returned by the method call g("alice").next()?
)	'i'
		'1'
		'c'
		'a'
)	'e'
Probl	en	9. Suppose we use the QuickUnionUF data structure to solve the dynamic con

Problem 9. Suppose we use the QuickUnionUF data structure to solve the dynamic connectivity problem with 10 sites and input pairs (8,1), (7,6), (9,2), (7,8), (4,6), (6,0), and (4,1), arriving in that order; the code for the union() method in QuickUnionUF is shown below.

```
public void union(int p, int q) {
   int rootP = find(p);
   int rootQ = find(q);
   if (rootP == rootQ) { return; }
   parent[rootP] = rootQ;
   count --;
}
```

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a. What are the values in the parent array after all the pairs are processed?
<pre>parent = {0, 0, 2, 0, 0, 2, 6, 0, 8, 0}</pre>
<pre>parent = {0, 0, 2, 3, 1, 5, 1, 6, 1, 2}</pre>
<pre>parent = {0, 0, 0, 3, 2, 2, 0, 0, 0, 9}</pre>
parent = {2, 0, 2, 0, 0, 6, 0, 8, 0}
parent = {0, 1, 0, 2, 0, 0, 0, 2, 8, 0}
b. What is the size of the largest component?
\bigcirc 2
\bigcirc 5
\bigcirc 3
\bigcirc 4
\bigcirc 6
c. What is the identifier of the largest component?
\bigcirc 2
\bigcirc 3
\bigcirc 1
\bigcirc 0
\bigcirc 4
Problem 10. Consider sorting an array a[] containing the following strings, using selection sort (show below):

S M

Z

N

H Y V T

```
public static void sort(Comparable[] a) {
    int N = a.length;
    for (int i = 0; i < N; i++) {
         int min = i;
         for (int j = i + 1; j < N; j++) { if (less(a[j], a[min])) {
                  min = j;
              }
         }
         exch(a, i, min);
    }
}
```

C

G

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What is the value that Y is exchanged with?
Н
○ s
\bigcirc M
O z
O N
Problem 11. Consider sorting an array a[] containing the following strings, using insertion sort (shown below):
D J M R S T Y Z O F
<pre>public static void sort(Comparable[] a) { int N = a.length; for (int i = 1; i < N; i++) { for (int j = i; j > 0 && less(a[j], a[j - 1]); j) { exch(a, j, j - 1); } } } Where is the item 0 sorted (ie, what is its index) relative to the items before?</pre>
Problem 12. Consider sorting an array a[] containing the following strings, using quick sort (shown below):
V U Z L S Y R E I J
<pre>public static void sort(Comparable[] a) { sort(a, 0, a.length - 1); }</pre>
<pre>private static void sort(Comparable[] a, int lo, int hi) { if (hi <= lo) return; int j = partition(a, lo, hi); sort(a, lo, j - 1); sort(a, j + 1, hi); }</pre>

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private static int partition(Comparable[] a, int lo, int hi) {

int i = lo;

```
int j = hi + 1;
Comparable v = a[lo];
while (true) {
    while (less(a[++i], v)) { if (i == hi) { break; } }
    while (less(v, a[--j])) { if (j == lo) { break; } }
    if (i >= j) { break; }
    exch(a, i, j);
}
exch(a, lo, j);
return j;
}
```

a. What is the state of the array a after the first call to partition()?

\bigcirc	E	U	J	L	S	I	R	V	Y	Z
	I	E	J	R	U	S	L	V	Z	Y
	Ι	S	U	R	J	L	E	V	Y	Z
	E	J	I	S	R	U	L	V	Y	Z
	R	L	U	J	E	I	S	V	Z	Y

b. What is pivot element in the next call to partition()?

	Z
	E
	R
	Ι
_	

) Y

Problem 13. Implement a comparable and iterable data type Genome that represents a genome sequence (a string of letters A, T, G, or C denoting nucleotides), and supports the following API:

	method	description
	Genome(String s)	construct a genome from s
a. (2 points)	double gcContent()	$\frac{G+C}{A+T+G+C} \times 100$; for example, GC content of the genome sequence "ACTGCG" is 67%
b. (2 points)	<pre>int compareTo(Genome that)</pre>	a comparison of this and that genome by their lengths
c. (2 points)	static GCContentOrder	for comparing genome sequences by their GC content
d-f (6 points)	<pre>Iterator iterator()</pre>	for iterating over the genome in <i>reverse</i> order

If s is a String object, you may use s.charAt(i) to obtain the ith character in the string.

```
import java.util.Comparator;
import java.util.Iterator;

public class Genome implements Comparable < Genome > , Iterable < Character > {
    private final String s; // the genome sequence

    public Genome (String s) { this.s = s; }
```

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```
public double gcContent() {
}
public int compareTo(Genome that) {
}
\verb"public static class GCC ontentOrder implements Comparator < \texttt{Genome} > \{
    public int compare(Genome g1, Genome g2) {
    }
}
public Iterator < Character > iterator() { return new ReverseIterator(); }
private class ReverseIterator implements Iterator {
    private int i; // index of current letter
    public ReverseIterator() {
    }
    public boolean hasNext() {
    }
    public Character next() {
    }
}
```

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Initials:

g. (2 points) Suppose sequences is an array of Genome objects. Write down a statement that uses Arrays.sort() to sort sequences by length.
h. (2 points) Write down a statement that uses Arrays.sort() to sort sequences by GC content.
Problem 14. a. (6 points) Given an array a containing N integers, provide a crisp and concise English description of an algorithm for finding the <i>closest</i> pair of integers. For example, (3, 4) is the closest pair in the array $a = \{4, 9, 3, -1, 6\}$.
b. (2 points) What is the order of growth of the worst case running time of your algorithm?

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