



#### 4. Reciprocate and Divide

```
void Fraction::reciprocal() {
    int tempDenom = denom;
    denom = num;
    num = tempDenom;
}

void Fraction::divide(Fraction other) {
    mult(Fraction(other.getDenom(), other.getNum()));
}
```

#### 5. First Date (Class)

```
// .h file
class Date {
public:
    Date(int m, int d);

    int getDay();
    int getMonth();
    int daysInMonth();
    void nextDay();
private:
    int m;
    int d;
};

// .cpp file
Date::Date(int month, int day) {
    m = month;
    d = day;
}

int Date::getDay() {
    return d;
}

int Date::getMonth() {
    return m;
}

int Date::daysInMonth() {
    switch (m) {
        case 2:
            return 28;
        case 1:
        case 3:
        case 5:
        case 7:
        case 8:
        case 10:
        case 12:
            return 31;
        default:
            return 30;
    }
}

void Date::nextDay() {
    d += 1;
    if (d > daysInMonth()) {
        m = ((m + 1) % 12);
        d = 1;
    }
}
```

## 6. Circle (Class) of Life

```
// .h file
class Circle {
public:
    Circle(double radius);

    double area();
    double circumference();
    double getRadius();
    string toString();
private:
    double r;
};

// .cpp file
Circle::Circle(double radius) {
    r = radius;
}

double Circle::area() {
    return PI * r * r;
}

double Circle::circumference() {
    return 2 * PI * r;
}

double Circle::getRadius() {
    return r;
}

string Circle::toString() {
    return string("Circle{radius=") + realToString(r) +
        string("}");
}
```

## 7. Align DNA Strands

```
int alignStrands(string &one, string &two, Map<string, int> &cache) {
    if (one.empty()) return -2 * two.length();
    if (two.empty()) return -2 * one.length();

    string key = one + ":" + two;
    if (cache.containsKey(key)) return cache[key];

    if (one[0] == two[0]) { // two leading bases match
        int score = alignStrands(one.substr(1), two.substr(1), cache) + 1;
        cache[key] = score;
        return cache[key];
    }

    int first = alignStrands(one, two.substr(1), cache) - 2; // insert the space in one
    int second = alignStrands(one.substr(1), two, cache) - 2; // insert the space in two
    int third = alignStrands(one.substr(1), two.substr(1), cache) - 1; // don't insert space
    cache[key] = max(first, max(second, third));
    return cache[key];
}

int alignStrands(const string& one, const string& two) {
    Map<string, int> cache;
    return alignStrands(one, two, cache);
}
```

## 8. Sorting in $O(n)$ Time

```
void linearSort(Vector<int> &vec, int k) {
    Vector<int> sorted(k + 1, 0); // Vector with 0-k (inclusive) elements, initialized to 0

    // create a histogram of the values in vec
    for (int i = 0; i < vec.size(); i++) {
        int elem = vec[i];
        sorted[elem] += 1;
    }

    // transform the histogram back into the individual values
    int counter = 0; // index at which to insert next element
    for (int i = 0; i < sorted.size(); i++) {
        for (int j = 0; j < sorted[i]; j++) {
            vec[counter++] = i;
        }
    }
}
```

This is a variation of a sorting algorithm called counting sort. Unlike many of the algorithms we looked at in class, it doesn't require comparing two or more elements, which is why it runs faster than  $O(n \log n)$  – you'll learn more about this if you go on to study algorithms. Why might you not want to use counting sort? Think of circumstances where the limitations of counting sort would prevent its use.