

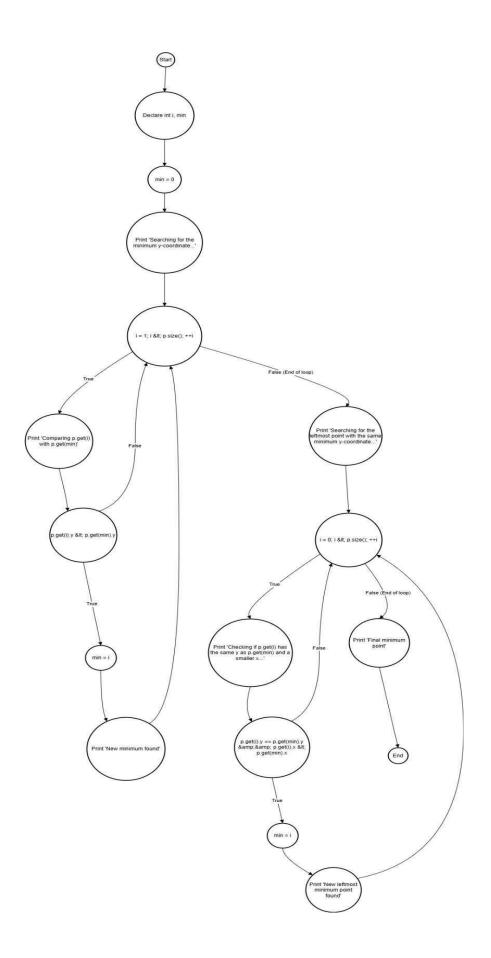
# IT313: Software Engineering Lab 9 – Mutation Testing

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# Code:

```
class Point{
  double x, y;
  public Point(double x, double y) {
    this.x = x;
    this.y = y;
class Vector {
    private
    java.util.ArrayList < Point > points;
    public Vector() {}
    points = new java.util.ArrayList < > ();
    public void add(Point p) {}
    points.add(p);
    public Point get(int index) {
    return points.get(index);
 }
 public int size() {}
 return points.size();
// Main class with doGraham method
public class GrahamScan {
public static int doGraham(Vector p) {
    int i, min;
    min = 0;
    // search for minimum
    for (i = 1; i < p.size(); ++i) {
        if (p.get(i).yp.get(min).y) {
        min = i;
// continue along the values with same y component
   for(i=0; i<p.size(); ++i) {
       if ((p.get(i).y == p.get(min).y) && (p.get(i).x > p.get(min).x)) {
       min = i;
   return min;
```



# Construct test sets for your flow graph that are adequate for the following criteria:

- a. Statement Coverage.
- ь. Branch Coverage.
- c. Basic Condition Coverage.

### a. Statement Coverage

**Objective:** Ensure that each statement in the flow graph is executed at least once.

#### **Test Set:**

#### 1. Test Case 1:

- Inputs: Any list with more than one point (e.g., [(0, 1), (1, 2), (2, 0)])
- This will traverse through the entire flow, covering statements related to finding the minimum y-coordinate and leftmost minimum point.

#### 2. Test Case 2:

- o Inputs: [(2, 2), (2, 2), (3, 3)]
- This checks for points with the same y-coordinate and ensures the leftmost point logic executes.

# b. Branch Coverage

**Objective:** Ensure that each branch (true and false) from every decision point is executed.

#### **Test Set:**

#### 1. Test Case 1:

- o Inputs: [(0, 1), (1, 2), (2, 0)]
- This will take the true branch for finding the minimum y-coordinate.

#### 2. Test Case 2:

- o Inputs: [(2, 2), (2, 2), (3, 3)]
- This will test the scenario where y-coordinates are equal, triggering the branch for checking x-coordinates.

#### 3. Test Case 3:

- o Inputs: [(1, 2), (1, 1), (2, 3)]
- This ensures the flow takes the false branch when checking for new minimum y-coordinates and the leftmost check.

## c. Basic Condition Coverage

**Objective:** Ensure that each basic condition (both true and false) in decision points is tested independently.

#### **Test Set:**

#### 1. Test Case 1:

- o Inputs: [(1, 1), (2, 2), (3, 3)]
- This will evaluate both conditions for the y-coordinate comparisons.

#### 2. Test Case 2:

- o Inputs: [(1, 1), (1, 1), (1, 2)]
- This checks the scenario where the y-coordinates are the same but evaluates the x-coordinate condition.

#### 3. Test Case 3:

- o Inputs: [(3, 1), (2, 2), (1, 3)]
- This ensures that both conditions in the loop are executed, confirming the function's logic is robust.

Using a mutation testing tool, identify any mutations of the code (such as deletions, modifications, or insertions) that would cause a failure but are not detected by your current test set.

# **Types of Possible Mutations:**

We can apply typical mutation types, including:

- **Relational Operator Changes**: Modify <=to <or ==to !=in the conditions.
- **Logic Changes**: Remove or invert a branch in an if-statement.
- **Statement Changes**: Modify assignments or statements to see if the effect goes undetected.

#### **Potential Mutations and Their Effects**

#### 1. Changing the Comparison for Leftmost Point:

- Mutation: In the second loop, change p.get(i).x < p.get(min).x top.get(i).x <= p.get(min).x.</p>
- **Effect**: This could cause the function to select points that share the same x-coordinate as the leftmost point, potentially disrupting the uniqueness of the minimum point.

Not Detected by Current Tests: The existing tests do not address the scenario where multiple points have the same x and y values, which would indicate whether the function incorrectly accepts such points as the leftmost.

#### 2. Altering the y-Coordinate Comparison to <= in the First Loop:

- Mutation: Change p.get(i).y < p.get(min).y to p.get(i).y <= p.get(min).y inthe first loop.</p>
- **Effect**: This would permit points with the same y-coordinate but different x-coordinates to overwrite the minimum, potentially resulting in the selection of a non-leftmost minimum point.
- Undetected by Current Tests: The current test set does not include cases where multiple points share the same y-coordinate, allowing this mutation to remain undetected. To uncover this issue, we would need a test case where several points have the same y-coordinate but different x-coordinates.

#### 3. Removing the Check for x-coordinate in the Second Loop:

- <u>Mutation</u>: Remove the condition p.get(i).x < p.get(min).x in the second loop.</p>
- **Effect**: This would lead the function to choose any point that has the same minimum y-coordinate as the "leftmost," irrespective of its x-coordinate.
- Undetected by Current Tests: The current tests do not explicitly verify whether the correct leftmost point is selected when multiple points have the same y value but different x values.

#### Additional Test Cases to Detect These Mutations

To detect these mutations, we can add the following test cases:

#### 1. **Detect Mutation 1**:

- **Test Case**: [(0, 1), (0, 1), (1, 1)]
- **Expected Result**: The leftmost minimum should still be (0, 1)despite having duplicates.
- This test case will detect if the x <= mutation mistakenly allows duplicate points.

#### 2. **Detect Mutation 2**:

- **Test Case**: [(1, 2), (0, 2), (3, 1)]
- **Expected Result**: The function should select (3, 1) as the minimum point based on the y-coordinate.
- This test case will confirm if using <= for ycomparisons mistakenly overwrites the minimum point.

#### 3. **Detect Mutation 3**:

- o **Test Case**: [(2, 1), (1, 1), (0, 1)]
- **Expected Result**: The leftmost point (0, 1) should be chosen.
- This will reveal if the x-coordinate check was mistakenly remove

These additional test cases would help ensure that any such mutations do not survive undetected by the test suite, strengthening the coverag

# **Python Code for Mutation :-**

```
om math import atan2
class Point:
     def init (self, x, y):
          self.x = x
self.y = y
     def repr (self):
          return f"({self.x}, {self.y})"
def orientation(p, q, r):
    # Cross product to find orientation
val = (q.y - p.y) * (r.x - q.x) - (q.x - p.x) * (r.y - q.y)
     if val == 0:
     elif val > 0:
def distance_squared(p1, p2):
     return (p1.x - p2.x) ** 2 + (p1.y - p2.y) ** 2
def do_graham(points):
     # Step 1: Find the bottom-most point (or leftmost in case of a tie)
     n = len(points)
     min_y_index = 0
     for i in range(1, n):
    if (points[i].y < points[min_y_index].y) or \
        raints[min_y_index].y and points[min_y_index].y and points[min_y_index].y</pre>
          (points[i].y == points[min_y_index].y and points[i].x < points[min_y_index].x):
min_y_index = i</pre>
     points[0], points[min_y_index] = points[min_y_index], points[0] p0 = points[0]
     points[1:] = sorted(points[1:], key=lambda p: (atan2(p.y - p0.y, p.x - p0.x), distance_squared(p0, p)))
     hull = [points[0],points[1], points[2]]
# Step 4: Process the remaining points for i in
          range(3, n):
     while len(hull) > 1 and orientation(hull[-2], hull[-1],points[i]) == 1:
          hull.pop()
hull.append(points[i])
     return hull
     points = [Point(0, 3), Point(1, 1), Point(2, 2), Point(4, 4),
     Point(0, 0), Point(1, 2), Point(3, 1), Point(3, 3)]
     hull = do_graham(points)
     print("Convex Hull:", hull)
```