1. About task 1 (15%)

1.1 Briefly explain how you implement your_fk() function (3%) (You can paste the screenshot of your code and explain it)

Initialize Base Position:

- Set up the transformation matrix A based on the initial base_pos, which initializes the robot arm's base position. This matrix will be used to iteratively apply each joint transformation.
- Initialize a 6x6 Jacobian matrix jacobian to later compute the linear and angular velocity mappings.

```
# Initialize the transformation matrix of the base position
base_pose = list(base_pos) + [0, 0, 0] # Initial base position with zero rotation
A = get_matrix_from_pose(base_pose) # 4x4 transformation matrix
jacobian = np.zeros((6, 6)) # 6x6 Jacobian matrix
```

Verify Input Length:

• Ensure that the DH parameters and joint angles q each contain 6 values, matching the robot arm's configuration.

```
# Ensure that both DH_params and q contain 6 values
assert len(DH_params) == 6 and len(q) == 6, f'Both DH_params and q should contain 6 values,\n' \
f'but received len(DH_params) = {len(DH_params)}, len(q) = {len(q)}'
```

Initialize Position and Orientation Storage:

• Store the base position and z-axis orientation, which will serve as the starting point for calculating the Jacobian matrix later.

```
# Store each joint's position and orientation (z-axis direction)
positions = [A[:3, 3]] # Store the base position
orientations = [A[:3, 2]] # Store the base z-axis direction
```

Define DH Transformation Matrix Generator:

• Use the dh_transform() function to generate the transformation matrix for each joint. This function calculates the transformation matrix based on each joint's DH parameters, linking each joint's position and orientation.

Apply Each Joint's DH Transformation Matrix Iteratively:

- For each joint, generate the transformation matrix T using its DH parameters and joint angle q[i], and update the cumulative matrix A by multiplying it with T.
- Simultaneously, store the position and z-axis orientation of each joint, which will be used to compute the Jacobian matrix later.

```
# Iteratively apply the DH transformation matrix for each joint
for i, params in enumerate(DH_params):
    # Generate the transformation matrix using the current joint angle q[i] and corresponding DH parameters
    T = dh_transform(params['a'], params['d'], params['alpha'], q[i])
    A = A @ T # Update the cumulative transformation matrix

# Store the current joint's position and z-axis direction for Jacobian computation
    positions.append(A[:3, 3])
    orientations.append(A[:3, 2])
```

Calculate the Jacobian Matrix:

- For calculating the Jacobian matrix, retrieve the end-effector position end effector pos.
- Iterate to calculate each column of the Jacobian matrix:
 - The linear velocity part is obtained by taking the cross product of the joint's z-axis vector and the difference between the end-effector and joint positions.
 - o The angular velocity part is simply the joint's z-axis orientation.

```
# Calculate the Jacobian matrix
end_effector_pos = positions[-1] # Position of the end effector
for i in range(6):
    # Linear velocity component: cross product of z-axis vector with the position difference to the end effector
    jacobian[:3, 1] = np.cross(orientations[i], (end_effector_pos - positions[i]))
    # Angular velocity component: directly take the z-axis direction
    jacobian[3:, i] = orientations[i]
```

Orientation Adjustment:

 Apply an adjustment matrix adjustment to ensure the final end-effector pose aligns with a specific orientation requirement, and apply it to the final rotation matrix part.

Return Final Pose and Jacobian Matrix:

• Extract the 7-dimensional pose (position and orientation) from the final transformation matrix A, and return it along with the computed Jacobian matrix.

```
pose_7d = np.asarray(get_pose_from_matrix(A, 7))
return pose 7d, jacobian
```

1.2 What is the difference between D-H convention and Craig's convention (Modified D-H Conveition)? (2%)

The Denavit-Hartenberg (D-H) convention and Craig's Modified D-H Convention both define transformations between coordinate frames of robotic joints, but they differ in how they assign these frames and the resulting transformation matrices.

1. Frame Assignment:

- **D-H Convention**: Assigns the frame of each joint along the previous joint's z-axis. The transformations focus on aligning each joint's frame using four parameters: θ , d, a, and α .
- Modified D-H Convention: Places the coordinate frame of each link along its own zaxis instead of the previous joint's z-axis. This leads to slightly different parameter definitions and can simplify certain link-to-link transformations, particularly for common robotic configurations.

2. Transformation Matrix:

- In the **D-H Convention**, the transformation from one frame to the next always follows the pattern: rotate around z_{i-1} , translate along z_{i-1} , then translate along x_i , and finally rotate around x_i .
- In the **Modified D-H Convention**, this order changes to: rotate around z_i , translate along z_i , translate along x_i , and finally rotate around x_i . This difference in sequence can make Craig's convention preferable for certain complex manipulations and robot structures.

In summary, while both conventions achieve forward kinematics, Craig's Modified D-H convention provides greater flexibility for frame assignment, which can simplify the math in some robotic configurations.

1.3 Complete the D-H table in your report following D-H convention (10%)

| i | d | α (rad) | a | θ (rad) |
|---|-------|----------------|-------|----------------|
| 1 | d_1 | $-\pi/2$ | 0 | $	heta_1$ |
| 2 | 0 | $\pi/2$ | 0 | $	heta_2$ |
| 3 | d_3 | $-\pi/2$ | a_3 | θ_3 |
| 4 | 0 | $\pi/2$ | a_4 | $	heta_4$ |
| 5 | d_5 | $-\pi/2$ | 0 | $	heta_5$ |
| 6 | 0 | $\pi/2$ | a_6 | θ_6 |
| 7 | d_7 | 0 | 0 | θ_7 |

D-H table example format.

Result

2. About task 2 (10% + 5% bonus)

2.1 Briefly explain how you implement your_ik() function (5%)

- (You can paste the screenshot of your code and explain it)

1. Setting Joint Limits:

The program first defines the joint angle limits for each joint and stores them in the joint_limits array to ensure that angles stay within these bounds during updates.

```
joint_limits = np.asarray([
      [-3*np.pi/2, -np.pi/2], # joint1
      [-2.3562, -1], # joint2
      [-17, 17], # joint3
      [-17, 17], # joint4
      [-17, 17], # joint5
      [-17, 17] # joint6
])
```

2. Initializing Joint Angles:

The initial joint angles of the robot arm are obtained using p.getJointStates, and the first six joint angles are extracted and stored in tmp q as the starting angles.

```
# Get the initial joint angles of the robotic arm
num_q = p.getNumJoints(robot_id)
q_states = p.getJointStates(robot_id, range(0, num_q))
tmp_q = np.asarray([x[0] for x in q_states][2:8]) # Initial angles for the 6 joints
```

3. Retrieving DH Parameters:

The DH parameters of the robot are retrieved using the get_ur5_DH_params() function, which will be used in the forward kinematics calculations.

```
# Retrieve DH parameters
DH_params = get_ur5_DH_params()
```

Setting Step Rate:

The step_rate is set to control the increment in angle updates. This helps adjust the rate of change, preventing overly large updates that may cause instability.

```
# Set step rate parameter to control update increments
step_rate = 0.1
```

Iteratively Updating Joint Angles:

The main iterative loop begins here. During each iteration, the program calculates the current pose of the end-effector and compares it with the target pose to reduce the difference.

- Calculate Current Pose and Jacobian Matrix: Using the your_fk function, the
 program computes the current end-effector pose current_pose and the Jacobian
 matrix J by passing in the DH parameters, current joint angles tmp_q, and base
 position base pos.
- Calculate Position and Rotation Differences: The differences between the current and target positions (delta_pos) and orientation (delta_rot_vec) are computed, which together form delta x.

```
for i in range(max_iters):
    # Use your_fk to obtain the current end-effector pose and Jacobian matrix
    current_pose, J = your_fk(DH_params, tmp_q, base_pos) # Pass DH_params, tmp_q, and base_pos

delta_pos = new_pose[:3] - current_pose[:3]
    delta_rot = R.from_quat(current_pose[3:]).inv() * R.from_quat(new_pose[3:])
    delta_rot_vec = delta_rot.as_rotvec()

delta_x = np.hstack((delta_pos, delta_rot_vec))
```

Check if Target Pose is Achieved:

If the difference in position and orientation (delta_x) is below the threshold stop thresh, the iteration stops, and the current joint angles tmp q are returned.

```
# Check if the change in position and orientation is below the threshold
if np.linalg.norm(delta_x) < stop_thresh:
    return list(tmp_q)</pre>
```

Calculate and Update Joint Angles:

Using the pseudo-inverse of the Jacobian matrix, the joint angle increment delta_q is calculated and applied to the current

```
# Calculate joint angle update and apply it
delta_q = step_rate * (pinv(J).dot(delta_x))
tmp_q += delta_q
```

Limit Joint Angles:

After each update, the joint angles are clamped to stay within the bounds defined in joint_limits to ensure they remain within the robot's allowable range of motion.

```
# Clamp joint angles within their limits
for j in range(len(tmp_q)):
    tmp_q[j] = np.clip(tmp_q[j], joint_limits[j][0], joint_limits[j][1])
```

Return Final Joint Angles:

If the target pose is reached within the maximum number of iterations, the list of final joint angles tmp_q is returned.

```
return list(tmp_q)
```

2.2 What problems do you encounter and how do you deal with them? (5%)

Precision in Pose Control

In the calculation process, it is essential to ensure that the robot's end-effector accurately reaches the target position and orientation, as even a small error can lead to a deviation in the final outcome. To maintain precision, I set a stop_thresh threshold. When the end-effector's deviation falls below this threshold, the iteration stops, preventing endless calculations and ensuring sufficient accuracy in the result.

Iteration Speed and Efficiency

This code uses a loop to iteratively adjust the joint angles, gradually bringing the endeffector closer to the target pose. However, if the iteration speed is too slow, it could impact the program's efficiency. To address this, I introduced the step_rate parameter to control the increment of each update, balancing accuracy and efficiency.

2.3 Bonus! Do you also implement other IK methods instead of pseudo-inverse method? How about the results? (5% bonus)

In addition to using the pseudo-inverse method for intuitive inverse kinematics, I also researched deep learning-based IK solutions. After training, these methods can quickly predict joint configurations to adapt to different target positions and orientations. However, at this stage, I have not yet been able to successfully implement the deep learning-based IK method.

```
Testcase file : ik_test_case_easy.json
- Mean Error : 0.001733
- Error Count : 0 / 300
 Your Score Of Inverse Kinematic : 13.333 / 13.333
- Testcase file : ik_test_case_medium.json
- Mean Error : 0.001371
- Error Count : 0 / 100
- Your Score Of Inverse Kinematic : 13.333 / 13.333

    Testcase file : ik_test_case_hard.json

 Mean Error : 0.001133
 Error Count : 0 / 100
 Your Score Of Inverse Kinematic : 13.333 / 13.333
______
 Your Total Score : 40.000 / 40.000
```

result

3. About task 3 (5%)

This part uses the your ik() function to control the robot and

```
Reward: 1.0 Done: Frue
8/10
Reward: 1.0 Done: True
9/10
Reward: 1.0 Done: True
10/10
Reward: 1.0 Done: True
   ard: 1.0 Done: True
```

In the test results, for each test (from Test: 1/10 to Test: 10/10), the model achieved a Total Reward of 1.0 and showed Done: True, indicating that the model successfully completed and met the objective in all 10 tests.

The final total score is 10.000 / 10.000, representing a perfect performance by the model in this task.

3.1 Compare your results between your ik function and pybullet ik

```
- Testcase file: ik_test_case_easy.json
- Mean Error: 0.001733
- Error Count: 0 / 300
- Your Score Of Inverse Kinematic: 13.333 / 13.333

- Testcase file: ik_test_case_medium.json
- Mean Error: 0.001371
- Error Count: 0 / 100
- Your Score Of Inverse Kinematic: 13.333 / 13.333

- Testcase file: ik_test_case_hard.json
- Mean Error: 0.001133
- Error Count: 0 / 100
- Your Score Of Inverse Kinematic: 13.333 / 13.333

- Tour Score Of Inverse Kinematic: 13.333 / 13.333

- Tour Score Of Inverse Kinematic: 13.333 / 13.333
```

your_ik

```
- Testcase file: ik_test_case_easy.json
- Mean Error: 0.001188
- Error Count: 0 / 300
- Your Score Of Inverse Kinematic: 13.333 / 13.333
- Testcase file: ik_test_case_medium.json
- Mean Error: 0.001459
- Error Count: 0 / 100
- Your Score Of Inverse Kinematic: 13.333 / 13.333
- Testcase file: ik_test_case_hard.json
- Mean Error: 0.001016
- Error Count: 0 / 100
- Your Score Of Inverse Kinematic: 13.333 / 13.333
```

pybullet_ik

When comparing the results of your_ik and pybullet_ik, we can see that both achieved a perfect total score of 40.000, with no errors (Error Count of 0 for each test case). The precision is also very close for both methods, but there are slight differences in the mean error across some test cases. Here's a detailed comparison:

1. ik_test_case_easy.json

o your_ik: Mean Error = 0.001733

- o pybullet_ik: Mean Error = 0.001188
- o Result: pybullet_ik shows a slightly smaller error than your_ik.

2. ik_test_case_medium.json

- o your_ik: Mean Error = 0.001371
- o pybullet_ik: Mean Error = 0.001459
- Result: In this case, your_ik has a slightly smaller error than pybullet_ik.

3. ik_test_case_hard.json

- o your_ik: Mean Error = 0.001133
- o pybullet_ik: Mean Error = 0.001016
- o Result: pybullet_ik shows a slightly smaller error than your_ik.

Summary

- Both methods are very close in precision and achieved a perfect score.
- pybullet_ik has slightly lower mean errors in the easy and hard test cases, while your_ik performs slightly better in the medium difficulty case.