Colin Casazza

Kendall Jordan

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SDSU CS556

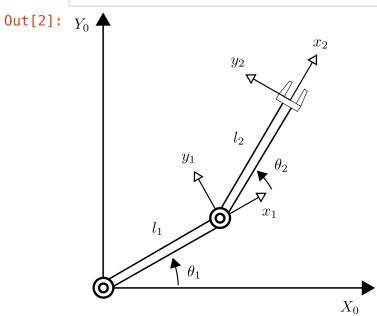
Assignment 3, Problem 2

The purpose of this problem is to determine the inverse kinematics of a manipulator with the method of Decomposition and Approximation

Notebook can be viewed and downloaded from https://github.com/colanzio5/SDSU-CS556-Workspace/tree/master/a3 (https://github.com/colanzio5/SDSU-CS556-Workspace/tree/master/a3)

```
In [1]:
        # packages
        import math
        import random
        import matplotlib
        import tinyik
        import numpy as np
        import sympy as sp
        import pandas as pd
        from sympy.solvers import solve
        from sympy.physics.mechanics import dynamicsymbols
        from sympy.physics.vector import init vprinting
        import plotly.graph objects as go
        from sklearn.cluster import KMeans
        from sklearn.linear model import LinearRegression
        # configure pretty latex like printing
        init vprinting(use latex="mathjax", pretty print=False)
        # number of random input angles
        # to fuzz into the forward kinematics
        # equations for approximation
        NUMBER_TRIALS = 50000
        # link lengths (in centimeters)
        L1 LENGTH = 100
        L2 LENGTH = 100
        # the resolution for our
        # our grid space approximation
        NUMBER ROWS = 50
        NUMBER COLS = 50
```

```
In [2]: # here's an image of the robot arm in question
    from IPython.display import Image
    Image("fig/imgl.png", width=300)
```

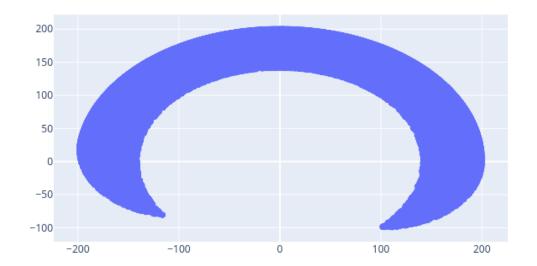


```
In [3]:
        Derive the forward kinematics equations to find the
        corresponding x,y coordnates based on given (\theta 1, \theta 2)
        pair for the 2PR manipulator
        # declare the symbolic variables
        # we'll need to do linear algebra
        theta1, theta2, l1, l2, theta, alpha, a, d = dynamicsymbols(
             "theta1 theta2 l1 l2 theta alpha a d"
        )
        # rotation matrix based off
        # of our dh table
        rotation matrix = sp.Matrix(
                 [sp.cos(theta), -sp.sin(theta) * sp.cos(alpha), sp.sin(theta)
        * sp.sin(alpha)],
                 [sp.sin(theta), sp.cos(theta) * sp.cos(alpha), -sp.cos(theta)
        * sp.sin(alpha)],
                 [0, sp.sin(alpha), sp.cos(alpha)],
        )
        # x,y,z transformation matrix
        transformation matrix = sp.Matrix([a * sp.cos(theta), a * sp.sin(theta))
        a), d])
        # the definition for the last row
        # in a homogenous transformation matrix
        last_row = sp.Matrix([[0, 0, 0, 1]])
        # combine the rotation, transformation and last row
        # to get homogenous transformation matrix
        t = sp.Matrix.vstack(sp.Matrix.hstack(rotation_matrix, transformation
        _matrix), last row)
        # given the homogenous transformation matrix
        # we need to find the transformation from
        # frame 0 to frame 2
        t_02 = t.subs({alpha: 0, a: l1, theta: theta1, d: 0}) * t.subs(
            {alpha: 0, a: 12, theta: theta2, d: 0}
        t 02.simplify()
        display(t 02)
```

$$egin{bmatrix} \cos(heta_1 + heta_2) & -\sin(heta_1 + heta_2) & 0 & l_1\cos(heta_1) + l_2\cos(heta_1 + heta_2) \ \sin(heta_1 + heta_2) & \cos(heta_1 + heta_2) & 0 & l_1\sin(heta_1) + l_2\sin(heta_1 + heta_2) \ 0 & 0 & 1 & 0 \ 0 & 0 & 1 \end{bmatrix}$$

```
# give our transformation matrix
# we can derive forward kinematics
# for example
x position = t 02[0, 3]
y position = t 02[1, 3]
fk_x = sp.lambdify((l1, l2, theta1, theta2), x_position, "numpy")
fk y = sp.lambdify((l1, l2, theta1, theta2), y position, "numpy")
def RR forward_kinematics(theta1, theta2):
        inverse kinematics for our 2PR robot
        given joint angles thetal and theta2
        returns the x and y positions of the
        end effector
    return (
        fk x(L1 LENGTH, L2 LENGTH, theta1, theta2),
        fk_y(L1_LENGTH, L2_LENGTH, theta1, theta2),
    )
def RR_inverse_kinematics(x, y):
        forward kinematics for our 2PR robot
        given a desired x any y coordinate
        returns both joint angles for the arm
        https://pythonrobotics.readthedocs.io/en/latest/modules/arm n
avigation.html
    try:
        # calculate both solutions for
        # theta 2
        a = x ** 2 + y ** 2 - L1 LENGTH ** 2 - L2 LENGTH ** 2
        b = 2 * L1 LENGTH * L2 LENGTH
        theta2_positive = math.acos((a / b))
        theta2 negative = -1 * theta2 positive
        # calculate both solutions for theta2
        thetal positive = math.atan2(y, x) - math.atan2(
            L1 LENGTH * math.sin(theta2 positive),
            L1_LENGTH + L2_LENGTH * math.cos(theta2_positive),
        theta1_negative = math.atan2(y, x) - math.atan2(
            L1 LENGTH * math.sin(theta2 negative),
            L1 LENGTH + L2 LENGTH * math.cos(theta2 negative),
        return [(thetal_positive, theta2_positive), (thetal_negative,
theta2 negative)]
    # an error is thrown when this function trys to
    # calculate the inverse kinematics for a position
    # outside the workspace of the 2PR arm
    except ValueError as e:
        return None
```

```
In [5]:
         Generate NUMBER_TRIALS random (\theta 1, \theta 2) pairs,
         and use the forward kinematics equations to find the
         corresponding points in workspace. Keeps a record of
         points in each cell.
         # map the forward kinematics
         # equations onto the input set
         trial_points = [
             {
                 "input_angles": input_angles,
                 "forward_kinematics": RR_forward_kinematics(*input_angles)
             } for input_angles in [
                     np.deg2rad(random.uniform(0, 170)),
                     np.deg2rad(random.uniform(-90, 90))
                 ) for x in range(NUMBER TRIALS)
             ]
         ]
         # plot the fuzzed forward
         # kinematics to show our workspace
         x cord = [ e['forward kinematics'][0] for e in trial points ]
         y_cord = [ e['forward_kinematics'][1] for e in trial_points ]
         go.Figure(
             data=go.Scattergl(
                 x = x \text{ cord}
                 y = y_{cord}
                 mode='markers'
         ).show(renderer="png")
```



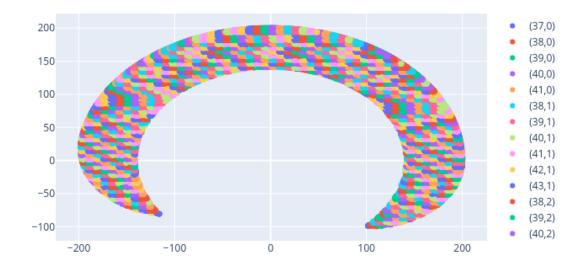
```
In [6]: # helper variables for
        # our workspace frame
        x min = math.floor(min(x cord))
        y min = math.floor(min(y cord))
        x max = math.ceil(max(x cord))
        y max = math.ceil(max(y cord))
        workspace width = x \max - x \min
        workspace_height = y_max - y_min
        # helper variables for our
        # grid representation
        cell width = workspace width / NUMBER COLS
        grid width = math.ceil(workspace width / cell width)
        cell height = workspace height / NUMBER ROWS
        grid height = math.ceil(workspace height / cell height)
        # helper function to map x,y position
        # vector to matching grid coordinates
        def get_grid_coordinates(x, y):
            x = math.floor(((x - x min) / workspace width) * grid width)
            _y = math.floor(((y - y_min) / workspace_height) * grid height)
            return _x, _y
In [7]:
        Divide the joint and workspace in into square cells.
        You will need to decide on the size of the cells in the
        joint space and work space, for example x=y=10 cm.
        # first set up a grid of points
        # so we can determine how to group
        # results from forward kinematics
        grid points = [[[]] for x in range(grid width)] for y in range(grid he
        ight)]
        # for each generated forward kinematics
        # output coordinate, we need to find it's
        # position in the grid and put in the grid
        # array we made previously
        for point in trial points:
            input_angles = (point["input_angles"][0], point["input_angles"][1
        ])
            forward kinematics = (point["forward kinematics"][0], point["forw
```

x_cord, y_cord = get_grid_coordinates(*forward_kinematics)

grid points[y cord][x cord].append(point)

ard kinematics"][1])

```
In [8]: | # some helper objects for the chart
        # we're about to make, dictionary of
        # colors we can randomly select from
        hex colors dic = {}
        rgb colors dic = {}
        hex colors only = []
        for name, hex in matplotlib.colors.cnames.items():
            hex colors only.append(hex)
            hex colors dic[name] = hex
            rgb_colors_dic[name] = matplotlib.colors.to_rgb(hex)
        0.00\,0
        show how we've split the workspace into grid
        sections, each colored, square cluster
        of points share the same grid coordinates
        grid plot = go.Figure()
        for y_index, row in enumerate(grid points):
             for x_index, cell in enumerate(row):
                 # if the cell has no
                 # points, do nothing
                 if len(cell) == 0:
                     continue
                 # plot the x,y pairs
                 # for this cell
                 grid plot.add trace(
                     go.Scattergl(
                         x=[ e['forward_kinematics'][0] for e in cell ],
                         y=[ e['forward_kinematics'][1] for e in cell ],
                         mode="markers",
                         fillcolor=random.choice(hex_colors_only),
                         name = "(" + str(x index) + ", " + str(y index) + ")",
                     )
        # show the plot
        grid plot.show(renderer="png")
```



```
In [9]:
        Using k-mean clustering, divide the points
        in each cell into two clusters, representing
        the two solutions.
        # now lets turn the output grid
        # into a data frame for easier
        # analysis of our approximation
        grid df = pd.DataFrame(e for e in grid points)
        # helper function to find
        # k means clusters for each
        # cell of the dataframe
        # and split coordinates in cell
        # into groups based on k means cluster
        def k means split(cell):
            # if the cell doesn't have at
            # least 2 points, we can't
            # do a kmeans cluster
            if len(cell) < 2:</pre>
                 return None
            # make t1 and t2 lists and
            # pair them for kmeans inputs
            t1 list = [ e['input angles'][0] for e in cell ]
            t2 list = [ e['input angles'][1] for e in cell ]
            t1 t2 pairs = list(zip(t1 list, t2 list))
            # create the kmeans model
            kmeans = KMeans(n clusters=2)
            cluster indexes = kmeans.fit predict(t1 t2 pairs)
            # group the points by cluster
            return [
                 [e for i, e in enumerate(cell) if cluster indexes[i] == 0],
                 [e for i, e in enumerate(cell) if cluster indexes[i] == 1],
             1
        # apply the function to split
        # each cell into two clusters
        cluster grid df = grid df.applymap(k means split)
```

```
In [10]:
         In each cell and each cluster in the cell,
         approximate the inverse kinematics by
         linear relationship of the form
         x=ai1+bi2+ci; x=di1+ei2+fi;
         where i=1,2,\ldots,n are the cell numbers,
         and the coefficients ai to fi are found
         using regression (curve fitting).
         Store the coefficients. This completes
         the decomposition and approximation.
         def lin reg(cell):
             # if there aren't any points
             # in the cell, do nothing
             if cell == None:
                 return None
             # helper function to
             # find the linear regression
             # for one cluster in the cell
             def lin reg helper(cluster):
                 x_list = [ e['forward_kinematics'][0] for e in cluster ]
                 y list = [ e['forward kinematics'][1] for e in cluster ]
                 x y pairs = list(zip(x_list, y_list))
                 t1_list = [ e['input_angles'][0] for e in cluster ]
                 t2 list = [ e['input angles'][1] for e in cluster ]
                 t1 t2 pairs = list(zip(t1 list, t2 list))
                 return LinearRegression().fit(x_y_pairs, t1_t2_pairs)
             # call the helper function for
             # each cluster in the cell
             return [ lin reg helper(cluster) for cluster in cell]
         lin_reg_cluster_grid = cluster_grid_df.applymap(lin_reg)
```

```
In [11]:
         Given a set of 100 position (x,y) points
         find the corresponding two solution (\theta 1, \theta 2) pairs
         for each position.
         trial errors = []
         predicted solutions = []
         inputs = []
         while len(trial errors) < 1000:
             # generate some random input positions
             # NOTE: these aren't guaranteed to be within
             # the workspace of the robot.
             px, py = random.uniform(x min, x max), random.uniform(y min, y max)
         X)
             # get the grid coordinates
             # for the regression lookup
             x, y = get grid coordinates(px, py)
             ik = RR inverse kinematics(px, py)
             # don't do anything if there's nothing in the cell
             # or if the inverse kinematics returns None
             if lin reg cluster grid[x][y] is None or ik is None:
                  continue
             # save the inputs and actual
             # inverse kinematics
             inputs.append((px,py))
             # get the inverse kinematic solutions
             # based of the cluster linear regressions
             c1, c2 = lin_reg_cluster_grid[x][y]
             # using the two linear regression objects
             # for each cell, pridict the two inverse
             # kinematic solutions for the given coordinates
             solution 1 = c1.predict([[px, py]])[-1]
             solution_2 = c2.predict([[px, py]])[-1]
             # given the predictetd solutions, see where
             # the arm would go if we used these input angles
             solution1 fk = RR forward kinematics(*solution 1)
             solution2 fk = RR forward kinematics(*solution 2)
             # helper function for finding the
             # distance between two numbers when
             # we don't know which is larger
             def distance(a, b):
                  if (a < 0) and (b < 0) or (a > 0) and (b > 0):
                      return abs(abs(a) - abs(b))
                 if (a < 0) and (b > 0) or (a > 0) and (b < 0):
                      return abs(abs(a) + abs(b))
             # determine the total error in our approximation
```

```
# by finding the distance between the actual point and
    # the two points predicted by linear approximation
    solution_1_error = abs(math.hypot(px - solution1_fk[0], py - solu
tion1 fk[1]))
    solution 2 error = abs(math.hypot(px - solution2 fk[0], py - solution2 fk[0])
tion2_fk[1]))
    # depending on which solution
    # is more accurate, add the most
    # accurate soluton's error to the
    # array of trial errors
    if(solution_1_error <= solution_2_error):</pre>
        predicted solutions.append(solution1 fk)
        trial_errors.append(solution_1_error)
    else:
        predicted solutions.append(solution2 fk)
        trial_errors.append(solution_2_error)
# take the average of all of the
# total errors for each trial
average_error = sum(trial_errors) / len(trial_errors)
print("average error: ", average_error, " cm")
```

average error: 0.12355501508197836 cm

```
In [12]:
         here's a chart showing the inputs to
         out test above on the x y plane
         each datapoint's size and color
         coresponds to the amount of error encountered
         during our approximation step, where
         smaller, darker datapoints are more accurate
         and larger lighter datapoints have more error
         Note: sizes and colors of data points
         in the scatter plot are not to scale
         in terms of the approximation error.
         Only for a relative scale.
         x_{inputs} = list(map(lambda x: x[0], inputs))
         y_{inputs} = list(map(lambda x: x[1], inputs))
         normalized_error = [ (x+5) for x in trial_errors]
         error chart = go.Figure(
             data=go.Scattergl(
                 x = x_inputs,
                 y = y_inputs,
                 mode='markers',
                 marker=dict(
                      size=normalized_error,
                      color=trial errors,
                      colorscale='Viridis'
              )
         error_chart.show(renderer="png")
```

