

Saad+Hamza_Assignment_2.4

April 18, 2023

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[ ]: import pandas as pd
import plotly.express as px
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[ ]: df = pd.read_csv('cancer.csv')
df.head()
```

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[ ]:
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	id	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean
0	842302	M	17.99	10.38	122.80	1001.0
1	842517	M	20.57	17.77	132.90	1326.0
2	84300903	M	19.69	21.25	130.00	1203.0
3	84348301	M	11.42	20.38	77.58	386.1
4	84358402	M	20.29	14.34	135.10	1297.0

	smoothness_mean	compactness_mean	concavity_mean	concave	points_mean
0	0.11840	0.27760	0.3001		0.14710
1	0.08474	0.07864	0.0869		0.07017
2	0.10960	0.15990	0.1974		0.12790
3	0.14250	0.28390	0.2414		0.10520
4	0.10030	0.13280	0.1980		0.10430

...	texture_worst	perimeter_worst	area_worst	smoothness_worst
0	17.33	184.60	2019.0	0.1622
1	23.41	158.80	1956.0	0.1238
2	25.53	152.50	1709.0	0.1444
3	26.50	98.87	567.7	0.2098
4	16.67	152.20	1575.0	0.1374

	compactness_worst	concavity_worst	concave	points_worst	symmetry_worst
0	0.6656	0.7119		0.2654	0.4601
1	0.1866	0.2416		0.1860	0.2750
2	0.4245	0.4504		0.2430	0.3613
3	0.8663	0.6869		0.2575	0.6638
4	0.2050	0.4000		0.1625	0.2364

	fractal_dimension_worst	Unnamed: 32
0	0.11890	NaN
1	0.08902	NaN

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2          0.08758          NaN
3          0.17300          NaN
4          0.07678          NaN

```

[5 rows x 33 columns]

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[ ]: #selecting the columns we want to use with '.loc' method
selected_df = df.loc[:, ['diagnosis', 'radius_mean',
                        'perimeter_mean', 'area_mean',
                        'radius_worst', 'perimeter_worst']]
selected_df

```

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[ ]:      diagnosis  radius_mean  perimeter_mean  area_mean  radius_worst
0           M         17.99         122.80       1001.0        25.380 \
1           M         20.57         132.90       1326.0        24.990
2           M         19.69         130.00       1203.0        23.570
3           M         11.42          77.58        386.1        14.910
4           M         20.29         135.10       1297.0        22.540
..          ...          ...          ...          ...          ...
564          M         21.56         142.00       1479.0        25.450
565          M         20.13         131.20       1261.0        23.690
566          M         16.60         108.30        858.1        18.980
567          M         20.60         140.10       1265.0        25.740
568          B          7.76          47.92        181.0         9.456

```

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      perimeter_worst
0          184.60
1          158.80
2          152.50
3           98.87
4          152.20
..          ...
564         166.10
565         155.00
566         126.70
567         184.60
568          59.16

```

[569 rows x 6 columns]

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[ ]: # with color, we specify the column based on whose values we want to assign
      ↪ colors in our plot
# with dimensions, we choose the columns we want to run the scatter plots
      ↪ against
fig = px.scatter_matrix(selected_df, color='diagnosis',
      ↪ dimensions=['radius_mean', 'perimeter_mean',

```

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↪ 'area_mean', 'radius_worst',  
  
↪ 'perimeter_worst'])  
  
#adjusting height and width for proper viewing  
fig.update_layout(height=800, width=800)  
fig.show()
```

