

pyStatQuest

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Preface

1 PCA (Singular Value Decomposition)

StatQuest 154 singular value decomposition SVD PCA:
python

1.0.1

PCA clustering Sum of square of distances / n-1 (variance) PC
variance scree plot PCA PC1, PC2 scree plot. PCA
Unsupervised dimensionality reduction

- cocktail recipe, linear combination
- singular vector, Eigenvector, loading score

1.0.2

PCA Singular Value Decomposition (SVD) Variances
PCA **linear dimensionality reduction**

- Dimension feature
-
-
-
-

1.0.3 PCA

- Factor Analysis
- Independent Component Analysis (ICA)

1.0.4 1

1.0.4.1 iris data

plotly iris features

```
import plotly.express as px
import pandas as pd

# load iris data
df = px.data.iris()
df.head()
```

/Users/yaochung41/.virtualenvs/statquest/lib/python3.9/site-packages/IPython/core/formatters

In future versions `DataFrame.to_latex` is expected to utilise the base implementation of `S

	sepal_length	sepal_width	petal_length	petal_width	species	species_id
0	5.1	3.5	1.4	0.2	setosa	1
1	4.9	3.0	1.4	0.2	setosa	1
2	4.7	3.2	1.3	0.2	setosa	1
3	4.6	3.1	1.5	0.2	setosa	1
4	5.0	3.6	1.4	0.2	setosa	1

```
# create scatter plot using plotly
features = ["sepal_width", "sepal_length", "petal_width", "petal_length"]

fig = px.scatter_matrix(
    df,
    dimensions=features,
    color="species"
)
fig.update_traces(diagonal_visible=False)
fig.show()
```

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1.0.4.2 sklearn PCA

```
from sklearn.decomposition import PCA

pca = PCA()
components = pca.fit_transform(df[features])
labels = {
    str(i): f"PC {i+1} ({var:.1f}%)"
    for i, var in enumerate(pca.explained_variance_ratio_ * 100)
}

fig = px.scatter_matrix(
    components,
    labels=labels,
    dimensions=range(4),
    color=df["species"]
)
fig.update_traces(diagonal_visible=False)
fig.show()
```

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PCA xy PC1 (92.5%), PC2 (5.3%)

1.0.5 2

1 scale features PCA scale 2 seaborn diamonds PCA

```
# load packages
import seaborn as sns
import matplotlib.pyplot as plt
import pandas as pd
import numpy as np
from sklearn import decomposition

diamond = sns.load_dataset("diamonds")
diamond.head()
```

/Users/yaochung41/.virtualenvs/statquest/lib/python3.9/site-packages/IPython/core/formatters

In future versions `DataFrame.to_latex` is expected to utilise the base implementation of `S

	carat	cut	color	clarity	depth	table	price	x	y	z
0	0.23	Ideal	E	SI2	61.5	55.0	326	3.95	3.98	2.43
1	0.21	Premium	E	SI1	59.8	61.0	326	3.89	3.84	2.31
2	0.23	Good	E	VS1	56.9	65.0	327	4.05	4.07	2.31
3	0.29	Premium	I	VS2	62.4	58.0	334	4.20	4.23	2.63
4	0.31	Good	J	SI2	63.3	58.0	335	4.34	4.35	2.75

features carat, cut, color, clarity, depth, table, price, x, y, z
PCA dummy variables

```
# get dummies and store it in a variable
dummies = pd.get_dummies(diamond[["cut", "clarity"]])

# concat dummies to original dataframe and drop values
merged = pd.concat([diamond, dummies], axis='columns')
merged.drop(['cut', 'clarity'], axis='columns', inplace=True)

#random select rows
merged = merged.sample(n=500)

print(merged.describe())
```

	carat	depth	table	price	x \
count	500.000000	500.000000	500.000000	500.000000	500.000000
mean	0.794720	61.721800	57.508600	3891.870000	5.729640
std	0.462839	1.451799	2.260089	4013.002554	1.120256
min	0.230000	53.300000	52.000000	413.000000	3.900000
25%	0.400000	61.000000	56.000000	951.000000	4.757500
50%	0.700000	61.800000	57.000000	2367.500000	5.680000
75%	1.062500	62.400000	59.000000	5369.250000	6.585000
max	2.540000	67.300000	67.000000	18797.000000	8.800000

	y	z	cut_Ideal	cut_Premium	cut_Very Good \
count	500.000000	500.000000	500.000000	500.000000	500.000000
mean	5.730760	3.535200	0.422000	0.256000	0.188000
std	1.112807	0.685153	0.494373	0.436859	0.391103
min	3.930000	2.380000	0.000000	0.000000	0.000000

25%	4.747500	2.910000	0.000000	0.000000	0.000000
50%	5.695000	3.535000	0.000000	0.000000	0.000000
75%	6.600000	4.060000	1.000000	1.000000	0.000000
max	8.700000	5.360000	1.000000	1.000000	1.000000

	cut_Good	cut_Fair	clarity_IF	clarity_VVS1	clarity_VVS2	\
count	500.000000	500.000000	500.000000	500.000000	500.000000	
mean	0.096000	0.038000	0.032000	0.062000	0.092000	
std	0.294886	0.191388	0.176176	0.241397	0.289315	
min	0.000000	0.000000	0.000000	0.000000	0.000000	
25%	0.000000	0.000000	0.000000	0.000000	0.000000	
50%	0.000000	0.000000	0.000000	0.000000	0.000000	
75%	0.000000	0.000000	0.000000	0.000000	0.000000	
max	1.000000	1.000000	1.000000	1.000000	1.000000	

	clarity_VS1	clarity_VS2	clarity_SI1	clarity_SI2	clarity_I1
count	500.000000	500.000000	500.000000	500.000000	500.000000
mean	0.148000	0.220000	0.244000	0.182000	0.020000
std	0.355456	0.414661	0.429923	0.386231	0.14014
min	0.000000	0.000000	0.000000	0.000000	0.000000
25%	0.000000	0.000000	0.000000	0.000000	0.000000
50%	0.000000	0.000000	0.000000	0.000000	0.000000
75%	0.000000	0.000000	0.000000	0.000000	0.000000
max	1.000000	1.000000	1.000000	1.000000	1.000000

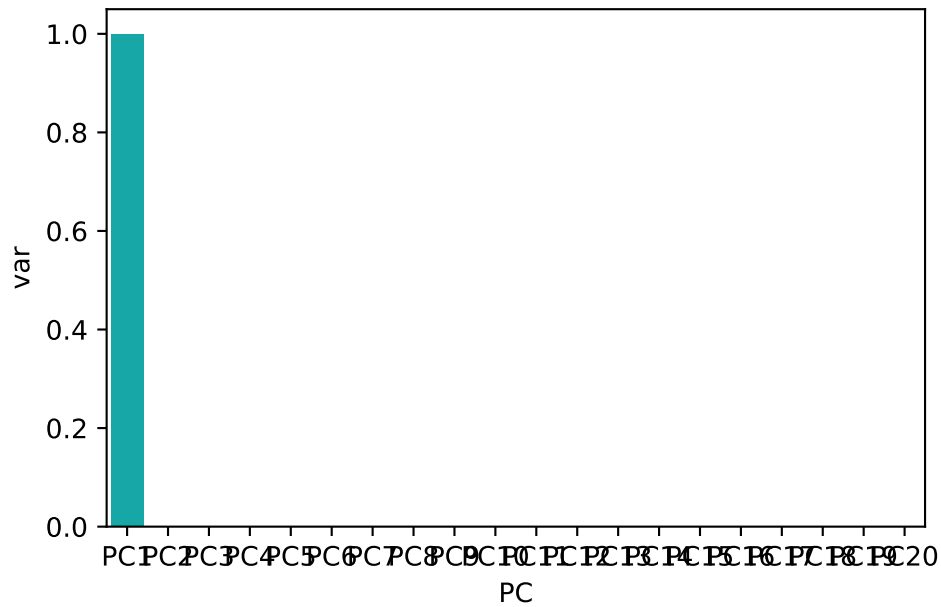
color min, max, std PCA

```
pca = decomposition.PCA()
pc = pca.fit_transform(merged.loc[:, merged.columns!='color'])
pc_df = pd.DataFrame(data=pc)
pc_df.head()
```

```
df = pd.DataFrame({
    'var': pca.explained_variance_ratio_,
    'PC': ["PC" + str(i) for i in list(range(1,21))]
})
```

```
sns.barplot(x = 'PC', y='var', data=df, color="c")
```

```
<AxesSubplot:xlabel='PC', ylabel='var'>
```

variance

```
from sklearn.preprocessing import StandardScaler

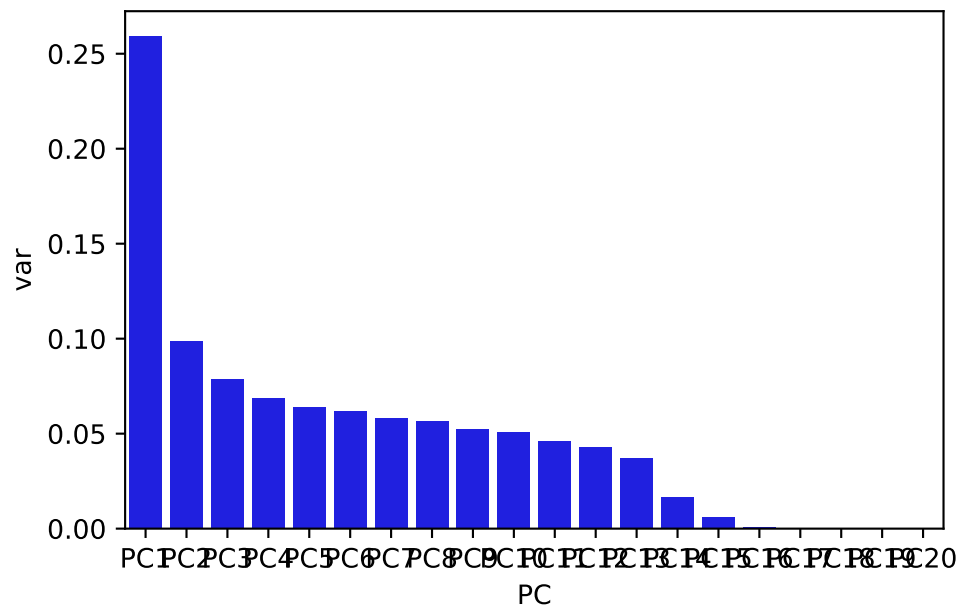
scaler = StandardScaler()
merged_scale = scaler.fit_transform(merged.loc[:, merged.columns!='color'])

pca = decomposition.PCA()
pc_scale = pca.fit_transform(merged_scale)
pc_df_scale = pd.DataFrame(pc_scale, columns = ["PC" + str(i) for i in list(range(1,21))])
pc_df_scale['color'] = merged.color

df_scale = pd.DataFrame({
    'var': pca.explained_variance_ratio_,
    'PC': ["PC" + str(i) for i in list(range(1,21))])
})

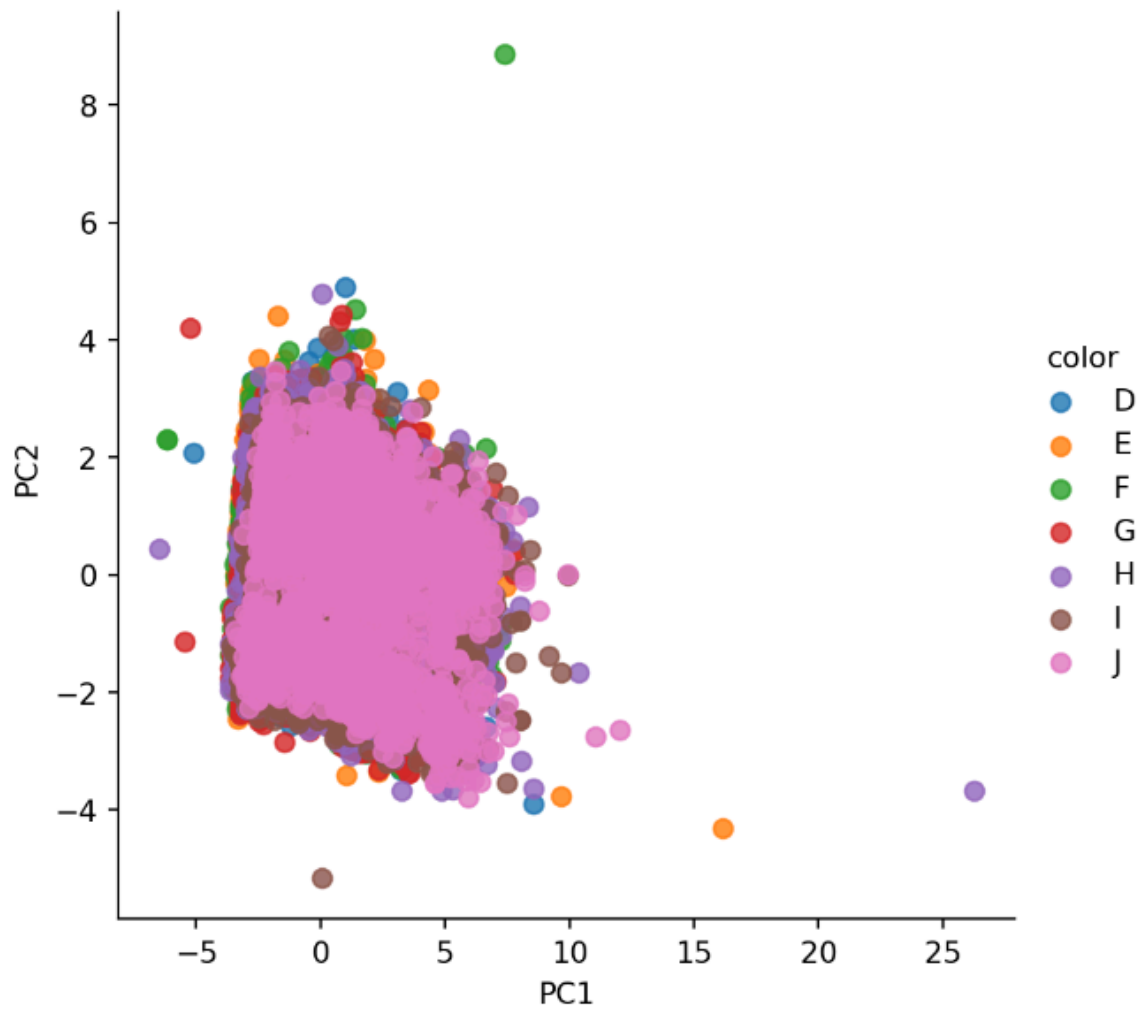
sns.barplot(x = 'PC', y='var', data=df_scale, color="b")
```

<AxesSubplot:xlabel='PC', ylabel='var'>



standard scaler

```
sns.lmplot(
    x="PC1",
    y="PC2",
    data=pc_df_scale,
    hue="color",
    fit_reg = False,
    legend=True,
    scatter_kws={"s": 40}
)
```

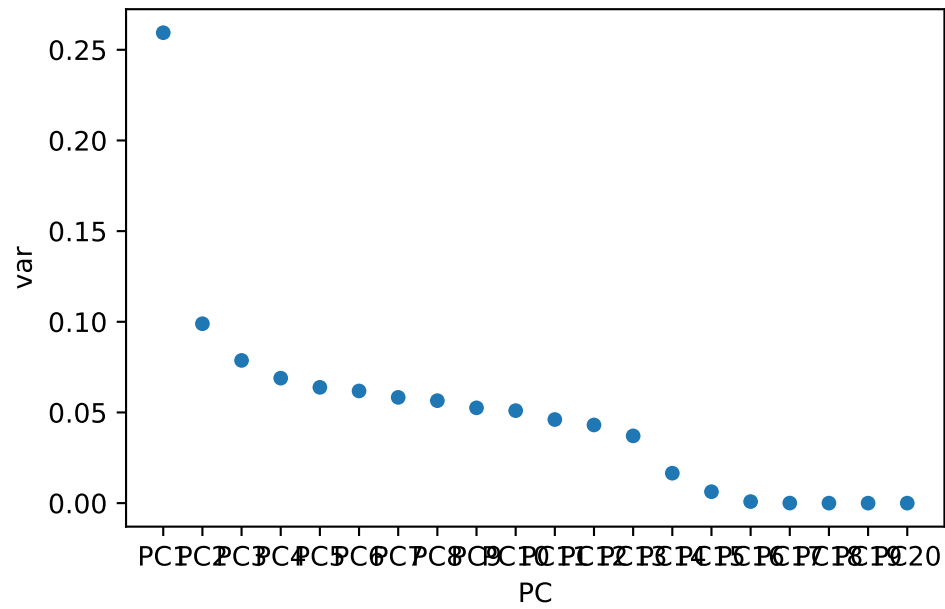


PCA clustering scree plot variance

scree plot using `sns.scatterplot`

```
pc_value = np.arange(pca.n_components_) + 1
sns.scatterplot(
    x='PC',
    y='var',
    data=df_scale
)
```

<AxesSubplot:xlabel='PC', ylabel='var'>



1.0.6

- PCA
- `normalization, standardization)` PCA
- (1) xy PC1, PC2 (2) Scree plot
- PCs Variance (Ex: 70%)

1.0.7

[PCA](#)
[Plotly PCA](#)
[PCA](#)
[Python and R tips](#)

2 Logistic Regression

StatQuest Logistic Regression

-
- True/ False
- (Walid Test)
- fit the line with Maximum-likelihood

python

3 Introduction

This is a book created from markdown and executable code.

See Knuth (1984) for additional discussion of literate programming.

4 Summary

In summary, this book has no content whatsoever.

References

Knuth, Donald E. 1984. “Literate Programming.” *Comput. J.* 27 (2): 97–111. <https://doi.org/10.1093/comjnl/27.2.97>.