

# Feature extraction

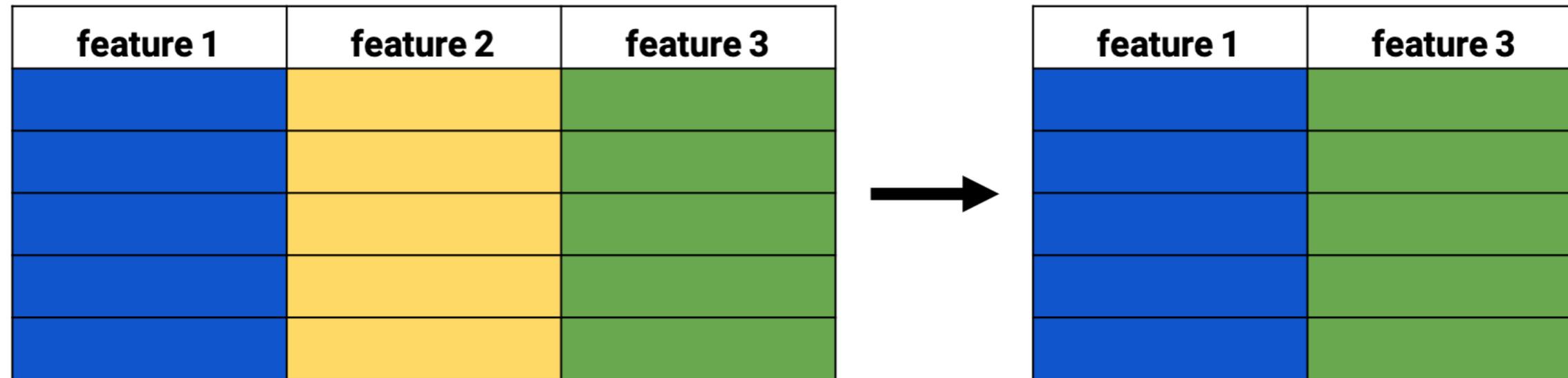
DIMENSIONALITY REDUCTION IN PYTHON



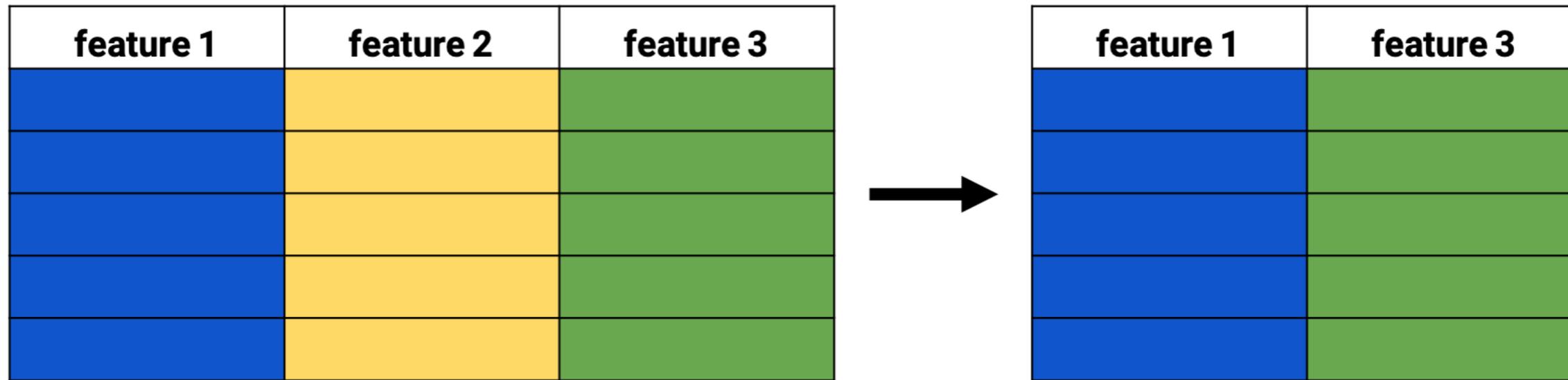
**Jeroen Boeye**

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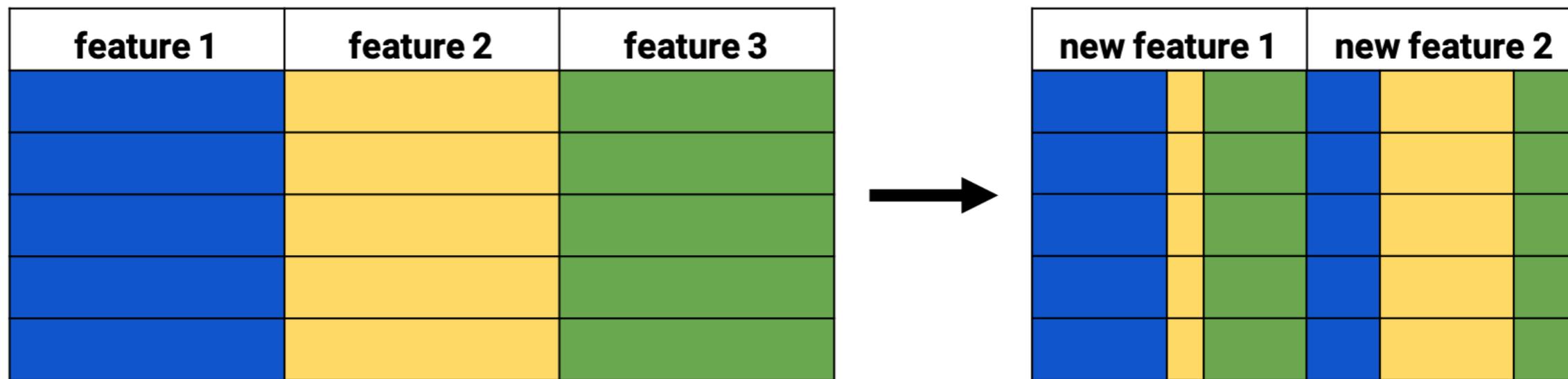
# Feature selection



# Feature selection



# Feature extraction



# Feature generation - BMI

```
df_body['BMI'] = df_body['Weight kg'] / df_body['Height m'] ** 2
```

# Feature generation - BMI

```
df_body['BMI'] = df_body['Weight kg'] / df_body['Height m'] ** 2
```

Weight kg	Height m	BMI
81.5	1.776	25.84
72.6	1.702	25.06
92.9	1.735	30.86

# Feature generation - BMI

```
df_body.drop(['Weight kg', 'Height m'], axis=1)
```

BMI
25.84
25.06
30.86

# Feature generation - averages

left leg mm	right leg mm
882	885
870	869
901	900

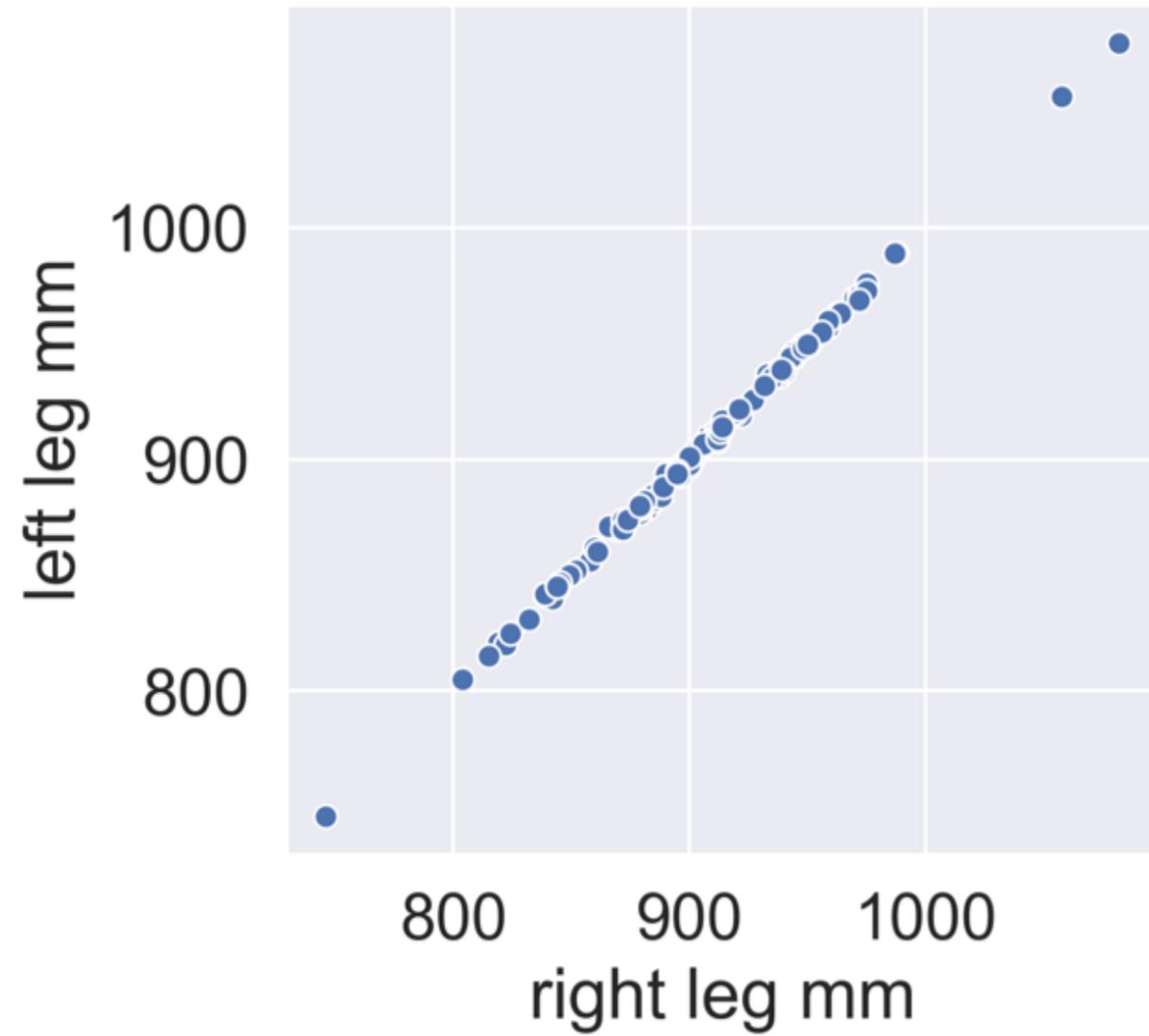
```
leg_df['leg mm'] = leg_df[['right leg mm', 'left leg mm']].mean(axis=1)
```

# Feature generation - averages

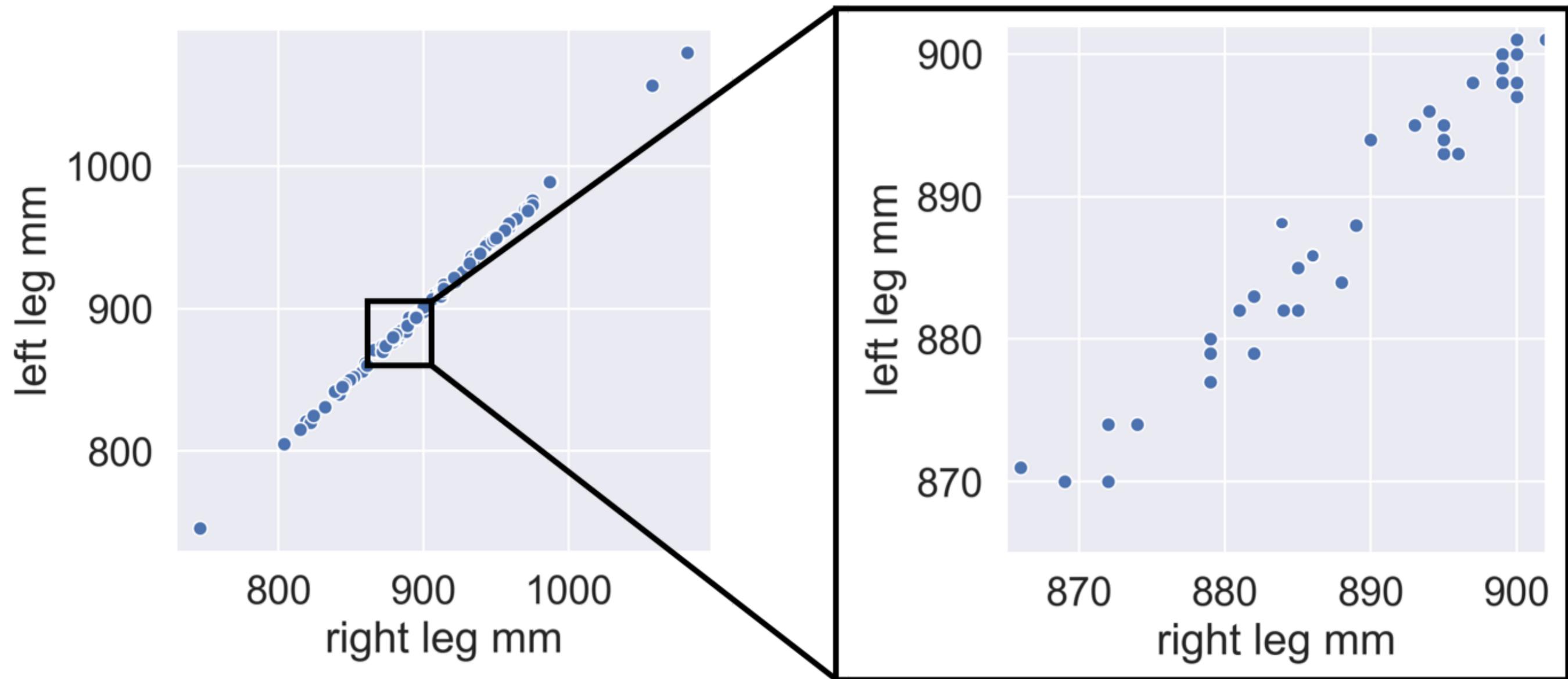
```
leg_df.drop(['right leg mm', 'left leg mm'], axis=1)
```

leg mm
883.5
869.5
900.5

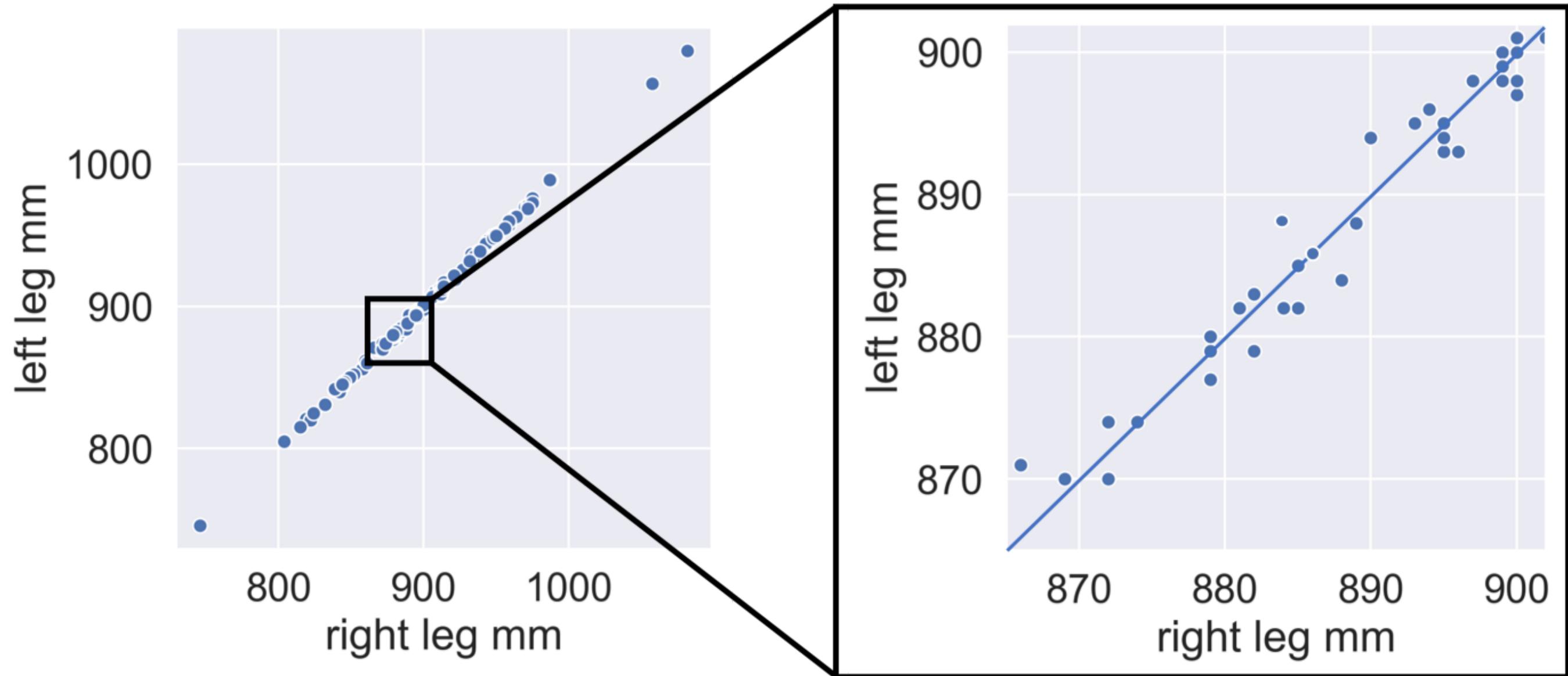
# Cost of taking the average



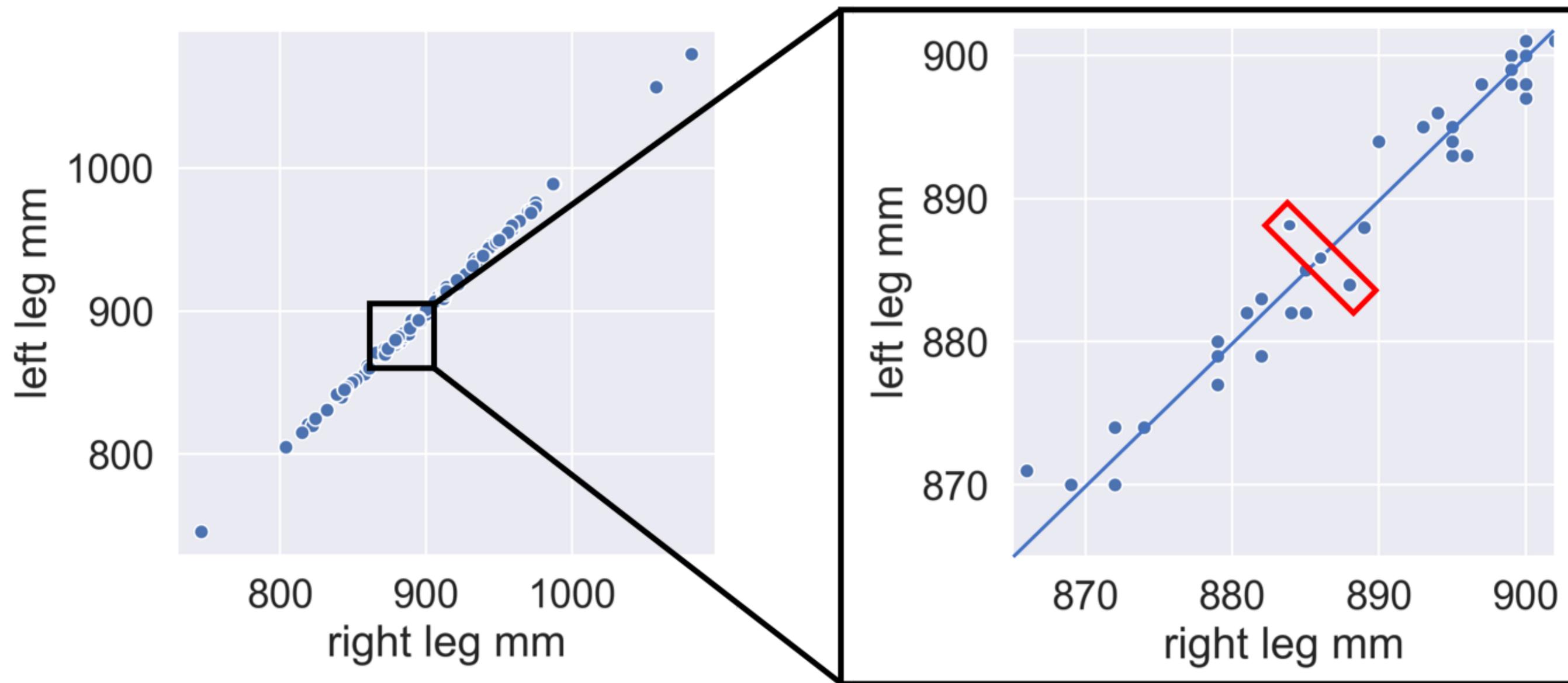
# Cost of taking the average



# Cost of taking the average

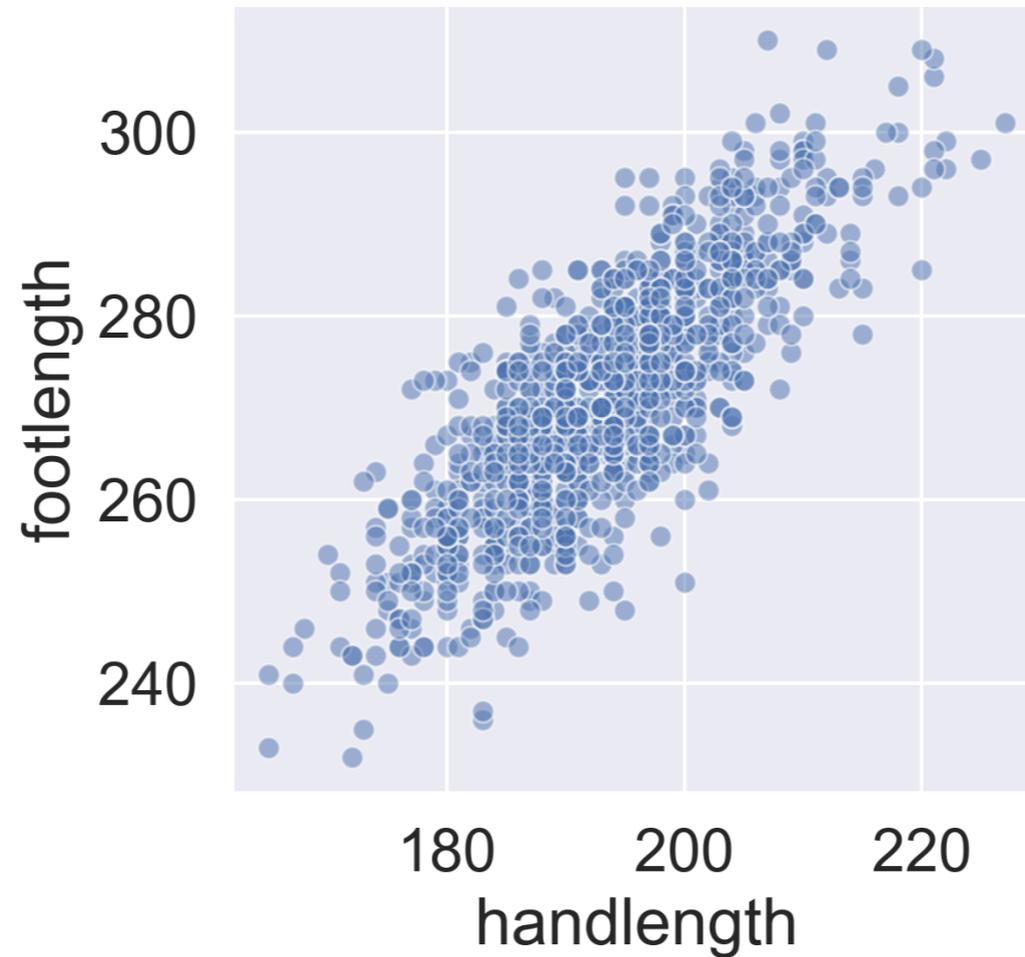


# Cost of taking the average



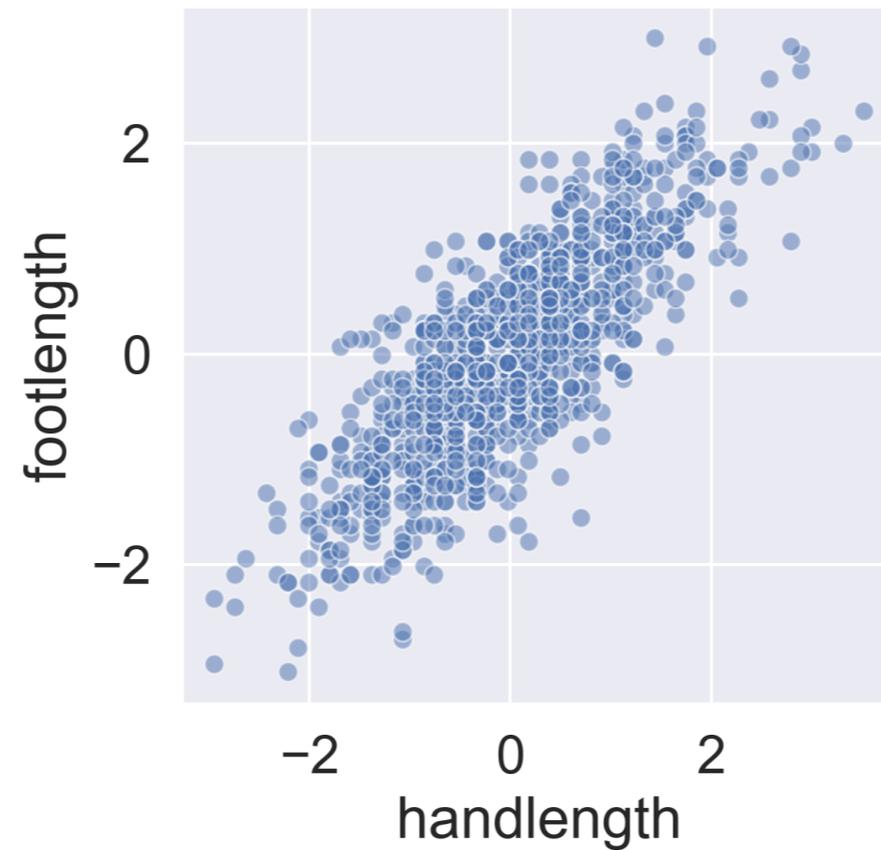
# Intro to PCA

```
sns.scatterplot(data=df, x='handlength', y='footlength')
```



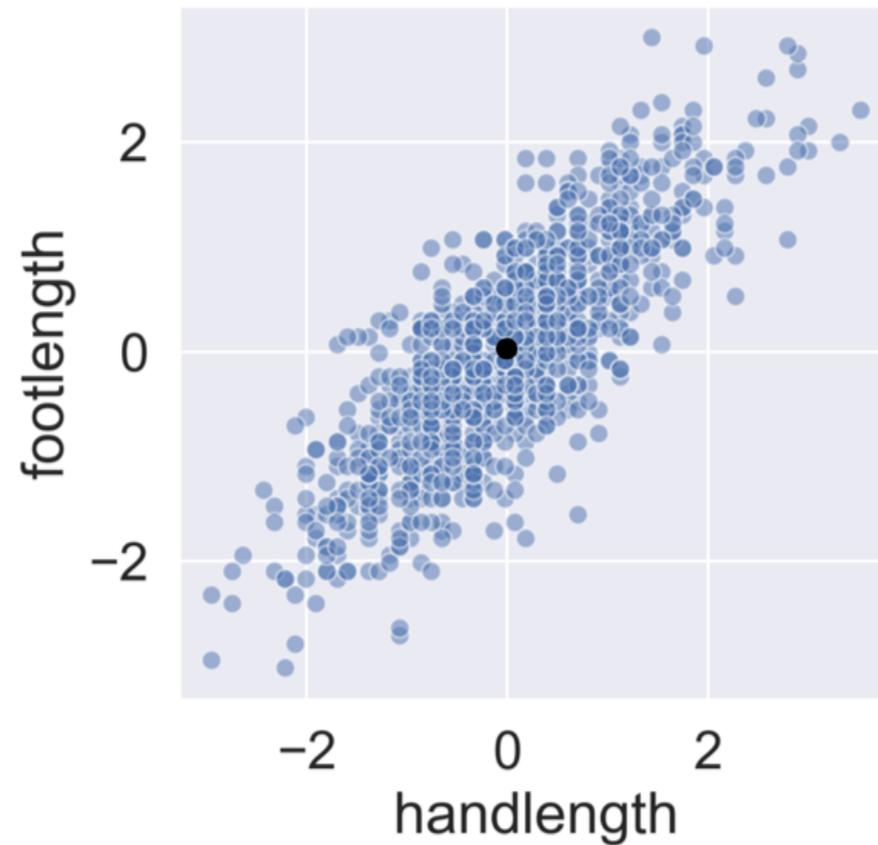
# Intro to PCA

```
scaler = StandardScaler()  
df_std = pd.DataFrame(scaler.fit_transform(df), columns = df.columns)
```



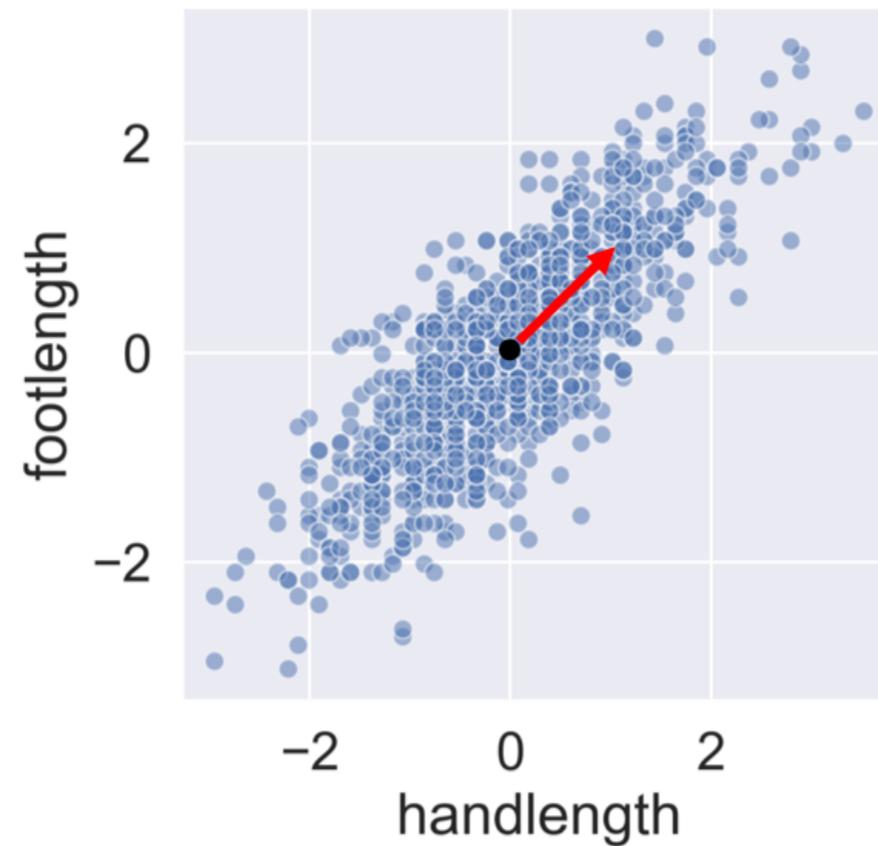
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```



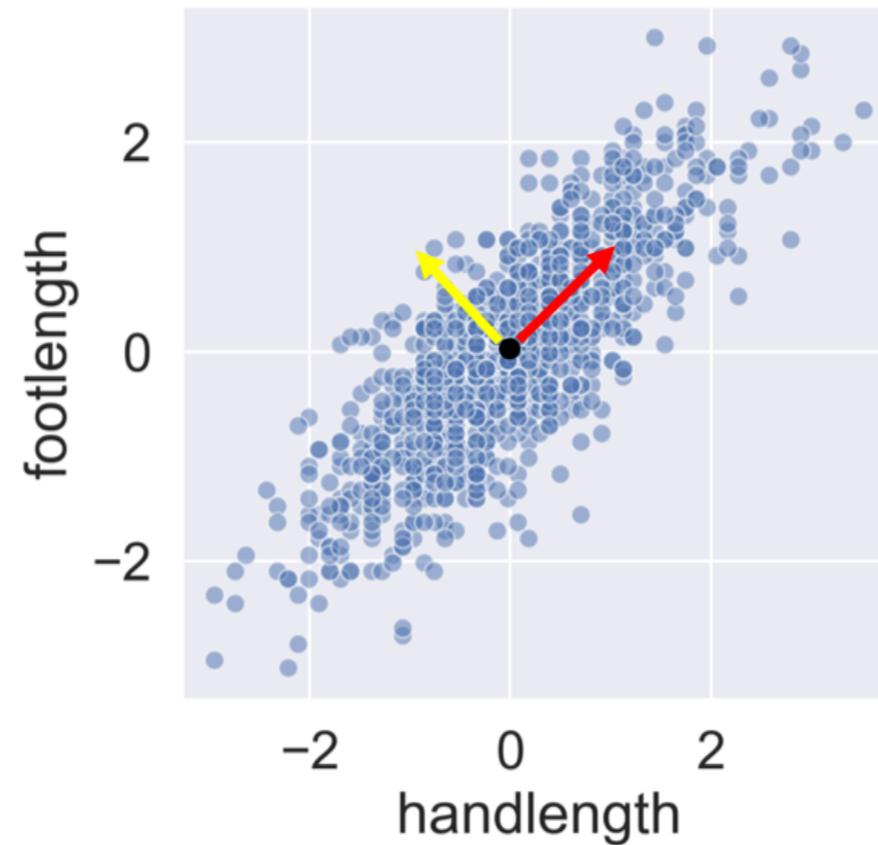
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# Intro to PCA

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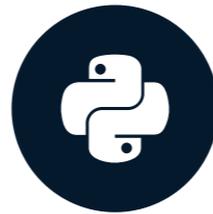


# Let's practice!

DIMENSIONALITY REDUCTION IN PYTHON

# Principal component analysis

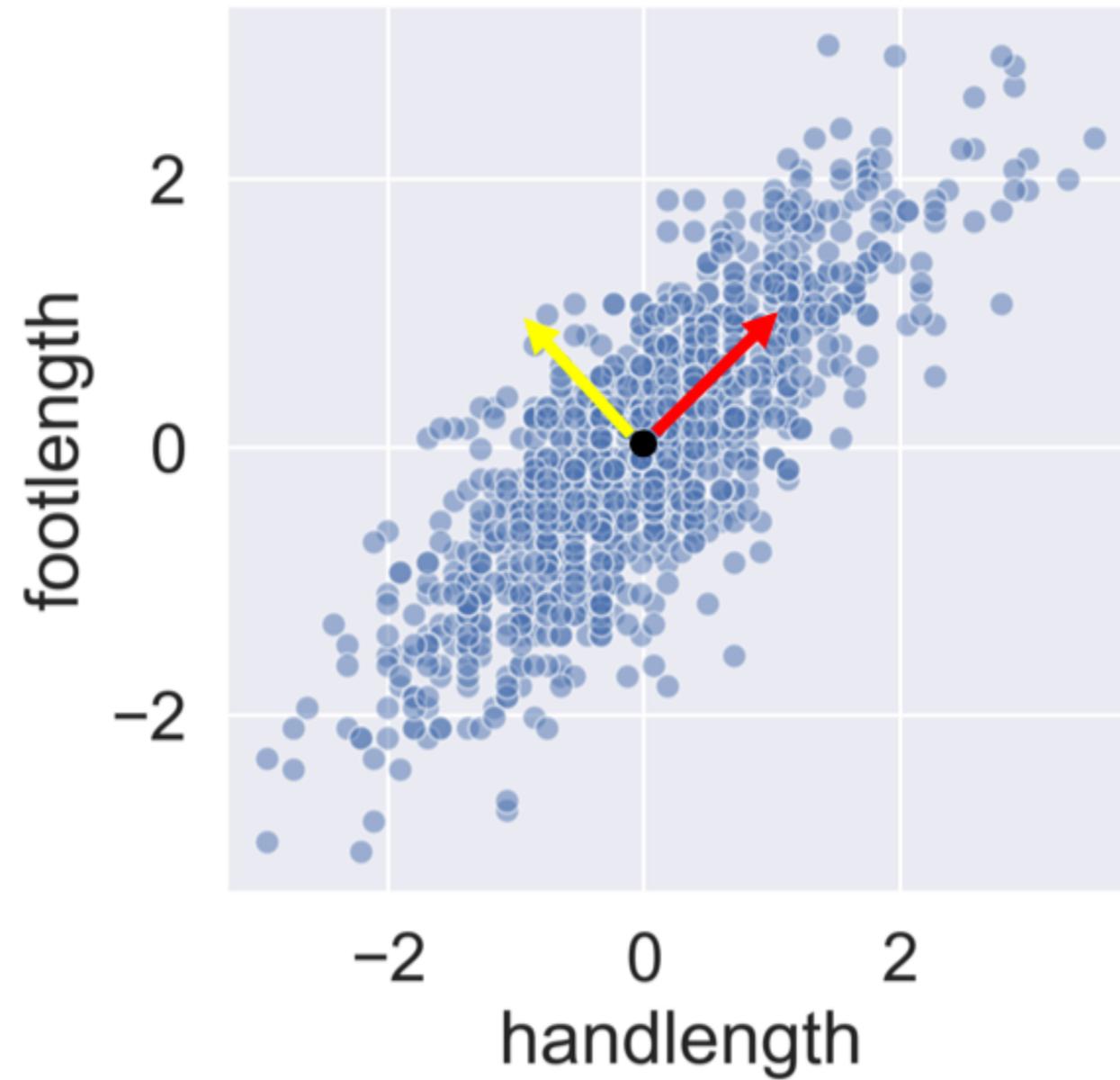
DIMENSIONALITY REDUCTION IN PYTHON



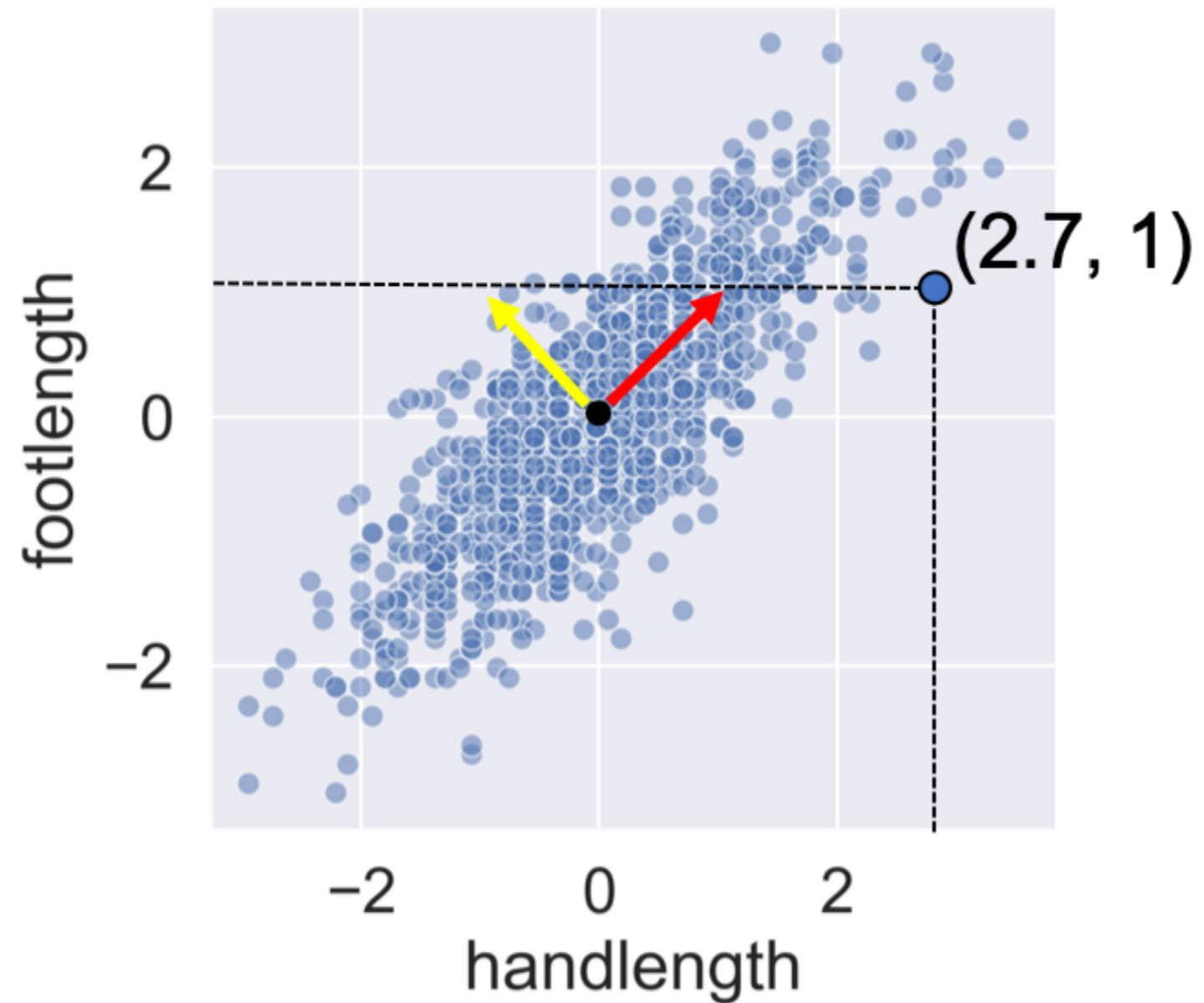
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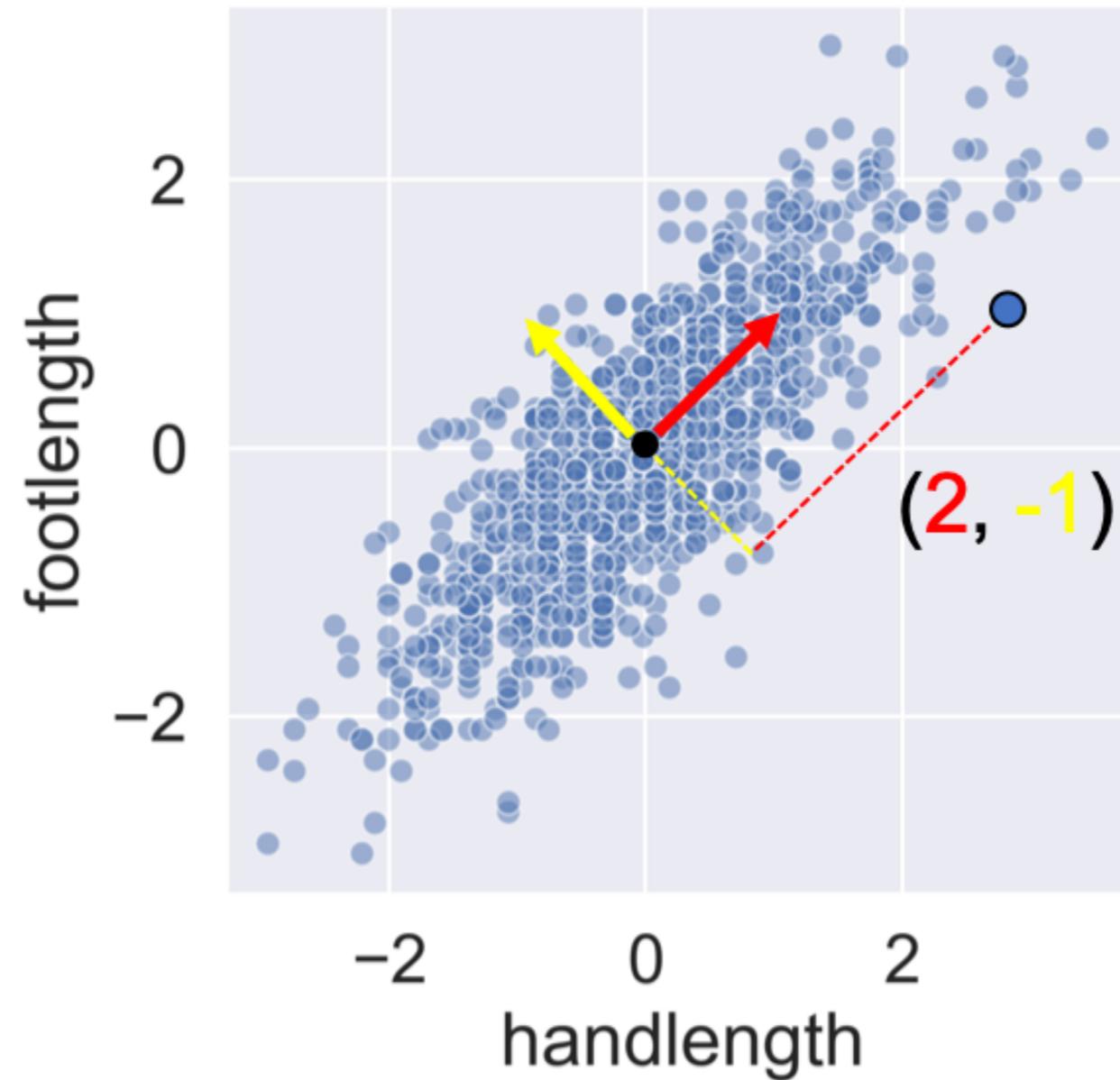
# PCA concept



# PCA concept



# PCA concept



# Calculating the principal components

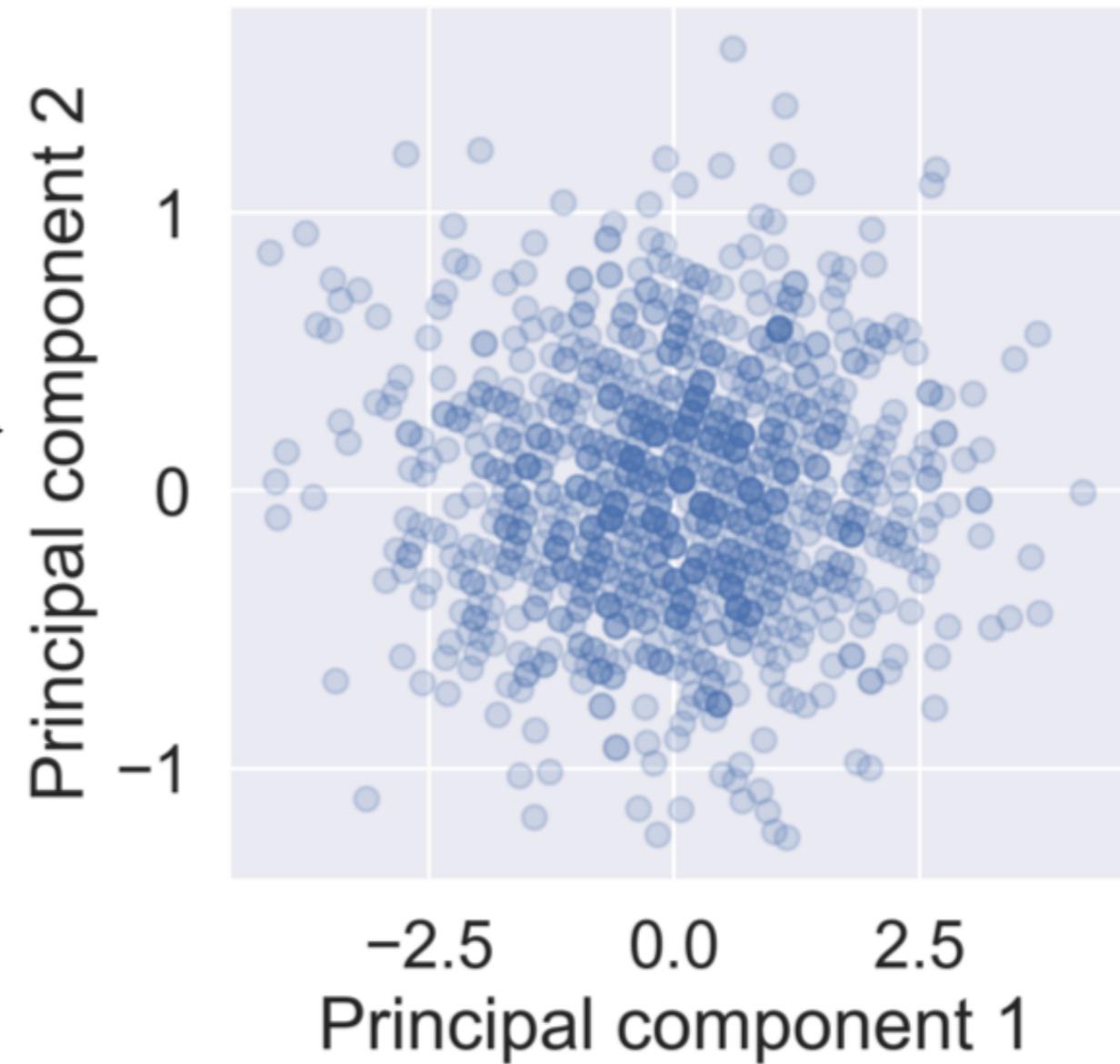
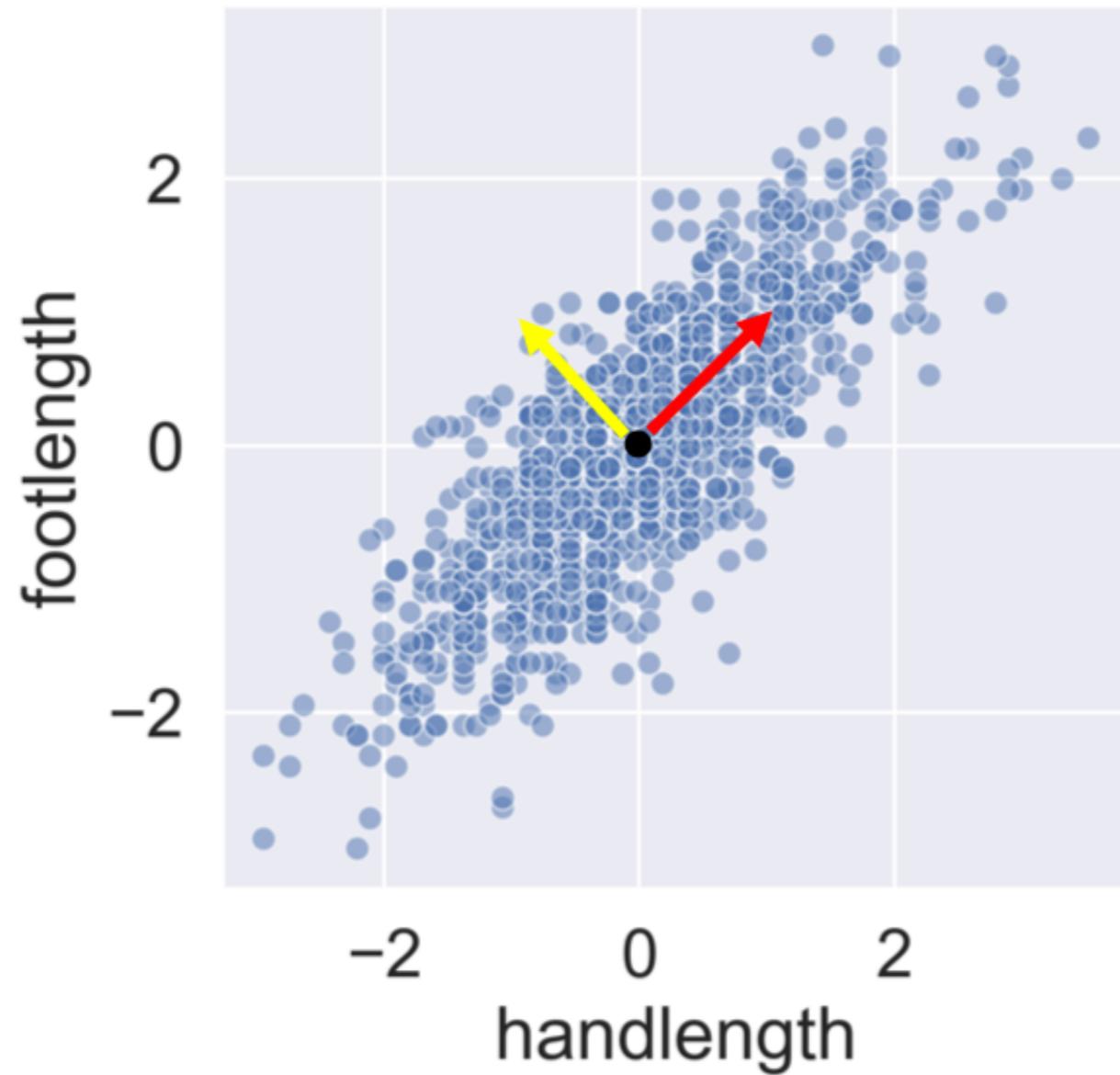
```
from sklearn.preprocessing import StandardScaler

scaler = StandardScaler()
std_df = scaler.fit_transform(df)
from sklearn.decomposition import PCA

pca = PCA()
print(pca.fit_transform(std_df))
```

```
[[[-0.08320426 -0.12242952]
 [ 0.31478004  0.57048158]
 ...
 [-0.5609523  0.13713944]
 [-0.0448304 -0.37898246]]]
```

# PCA removes correlation



# Principal component explained variance ratio

```
from sklearn.decomposition import PCA

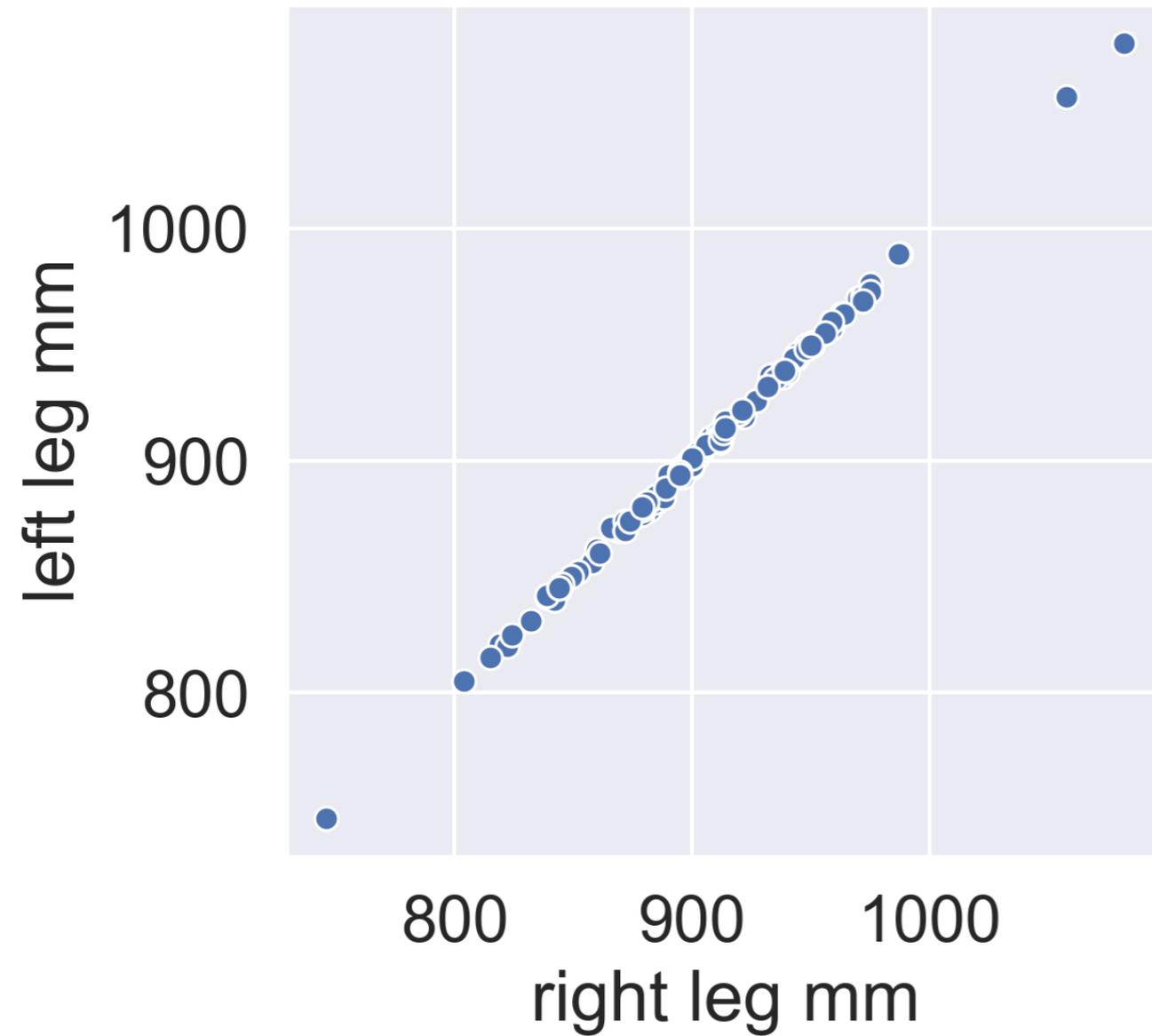
pca = PCA()

pca.fit(std_df)

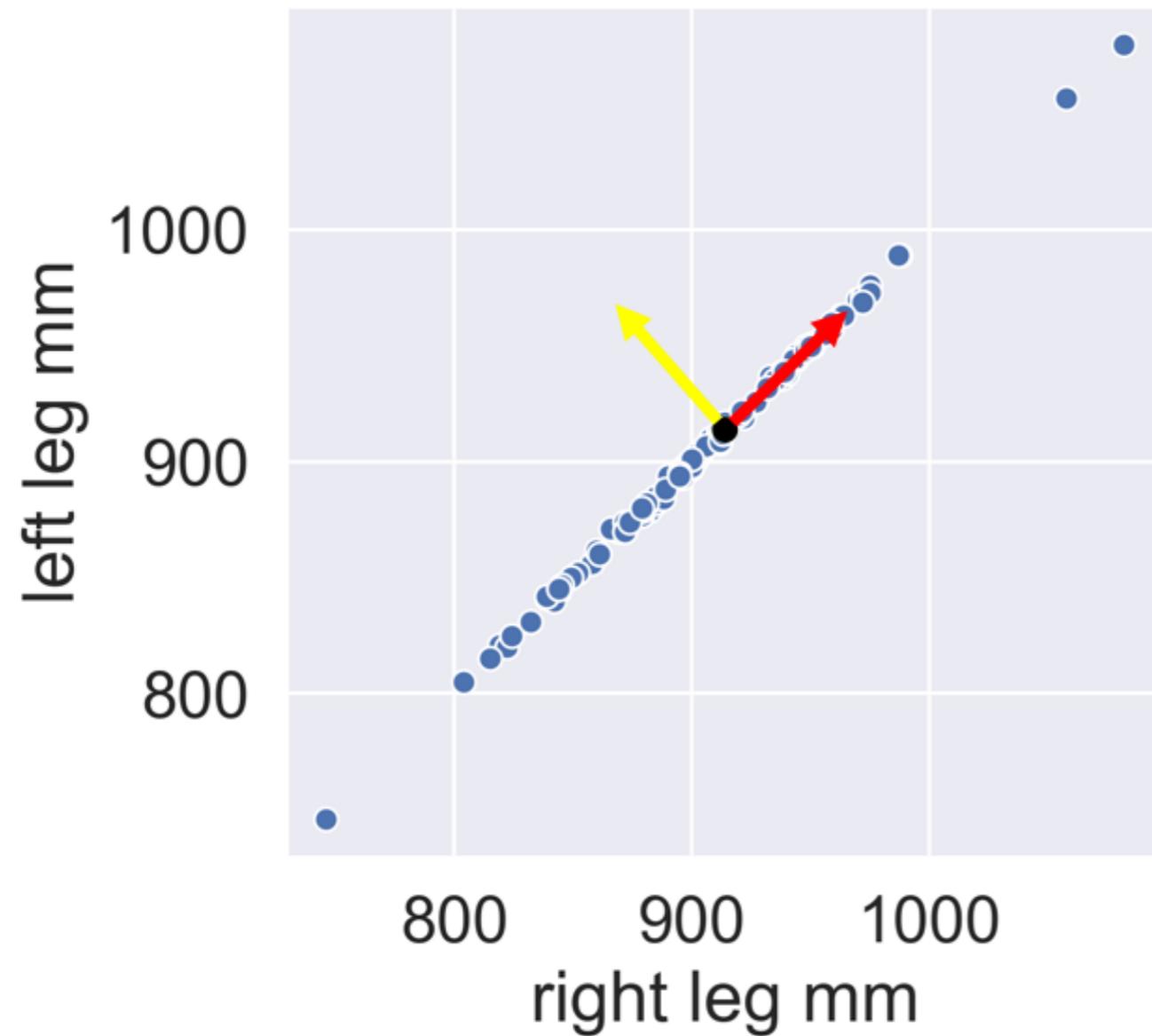
print(pca.explained_variance_ratio_)
```

```
array([0.90, 0.10])
```

# PCA for dimensionality reduction



# PCA for dimensionality reduction



```
print(pca.explained_variance_ratio_)
```

```
array([0.9997, 0.0003])
```

# PCA for dimensionality reduction

```
pca = PCA()

pca.fit(ansur_std_df)

print(pca.explained_variance_ratio_)
```

```
array([0.44, 0.18, 0.04, 0.03, 0.02, 0.02, 0.02, 0.01, 0.01, 0.01, 0.01,
       0.01, 0.01, 0.01, 0.01, 0.01, 0.01, 0.01, 0.01, 0.01, 0.01, 0.01,
       0.01, 0.01, 0.01, 0. , 0. , 0. , 0. , 0. , 0. , 0. , 0. ,
       ...,
       0. , 0. , 0. , 0. , 0. , 0. , 0. , 0. , 0. , 0. , 0. ,
       0. , 0. , 0. , 0. , 0. , 0. , 0. , 0. , 0. , 0. , 0. ,
       0. , 0. , 0. , 0. , 0. , 0. ])
```

# PCA for dimensionality reduction

```
pca = PCA()

pca.fit(ansur_std_df)

print(pca.explained_variance_ratio_.cumsum())
```

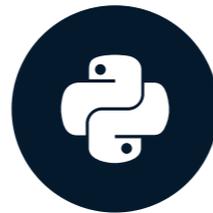
```
array([0.44, 0.62, 0.66, 0.69, 0.72, 0.74, 0.76, 0.77, 0.79, 0.8 , 0.81,
       0.82, 0.83, 0.84, 0.85, 0.86, 0.87, 0.87, 0.88, 0.89, 0.89, 0.9 ,
       0.9 , 0.91, 0.92, 0.92, 0.92, 0.93, 0.93, 0.94, 0.94, 0.94, 0.95,
       ...,
       0.99, 0.99, 0.99, 0.99, 0.99, 1. , 1. , 1. , 1. , 1. , 1. ,
       1. , 1. , 1. , 1. , 1. , 1. , 1. , 1. , 1. , 1. ,
       1. , 1. , 1. , 1. , 1. , 1. ])
```

# Let's practice!

DIMENSIONALITY REDUCTION IN PYTHON

# PCA applications

DIMENSIONALITY REDUCTION IN PYTHON



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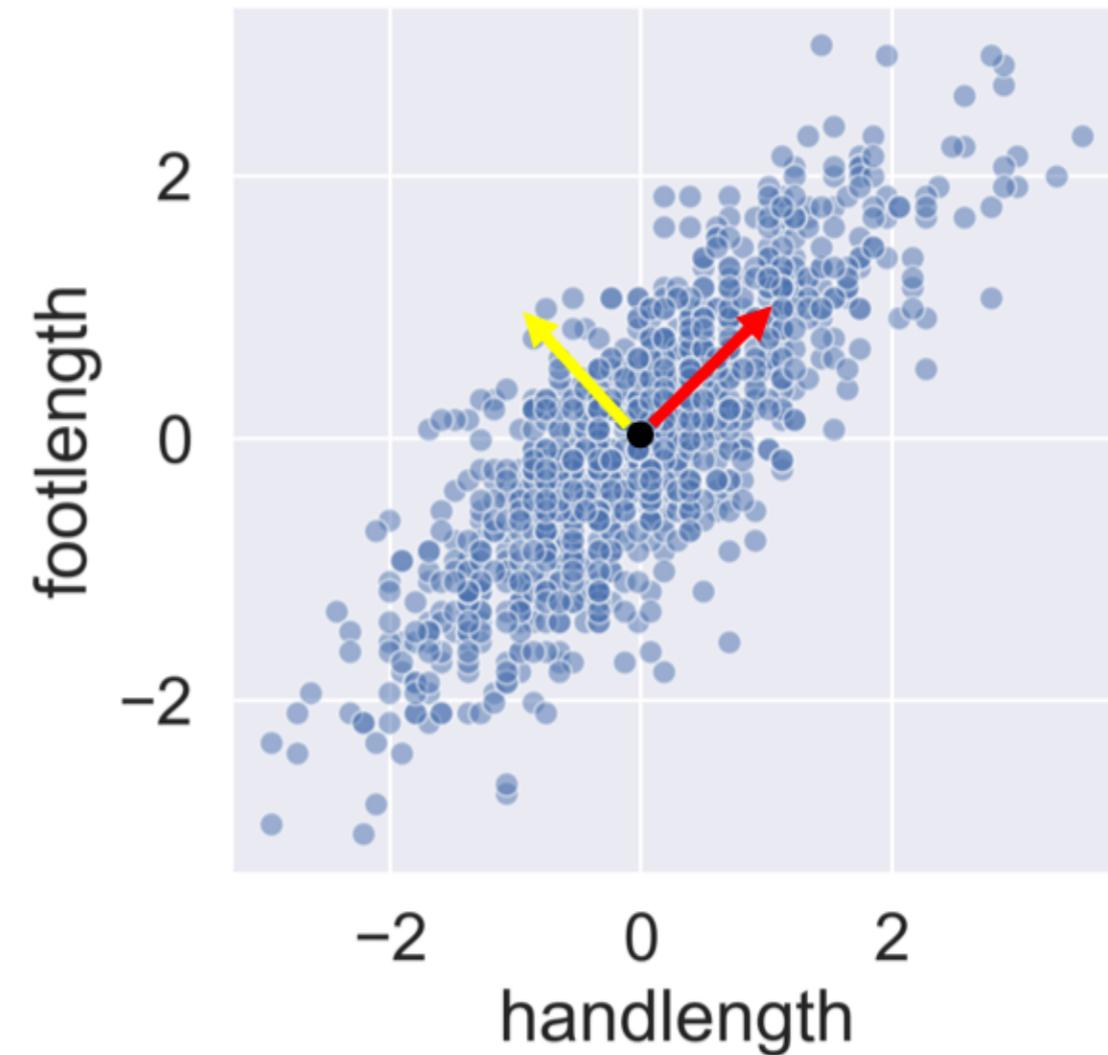
# Understanding the components

```
print(pca.components_)
```

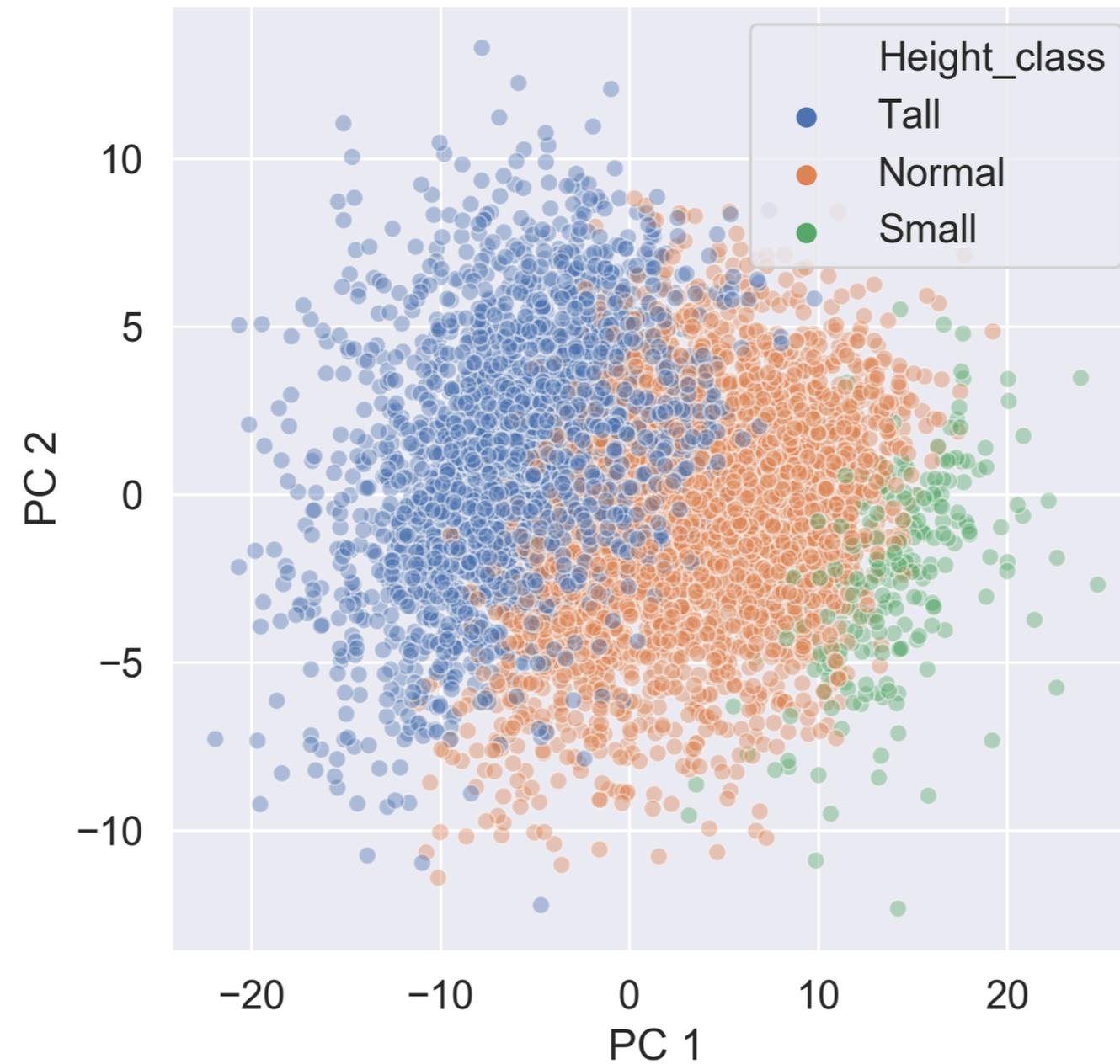
```
array([[ 0.71,  0.71],  
       [-0.71,  0.71]])
```

PC 1 = 0.71 x Hand length + 0.71 x Foot length

PC 2 = -0.71 x Hand length + 0.71 x Foot length



# PCA for data exploration



# PCA in a pipeline

```
from sklearn.preprocessing import StandardScaler
from sklearn.decomposition import PCA
from sklearn.pipeline import Pipeline

pipe = Pipeline([
    ('scaler', StandardScaler()),
    ('reducer', PCA())])
pc = pipe.fit_transform(ansur_df)

print(pc[:, :2])
```

```
array([[ -3.46114925,  1.5785215 ],
       [  0.90860615,  2.02379935],
       ...,
       [10.7569818 , -1.40222755],
       [  7.64802025,  1.07406209]])
```

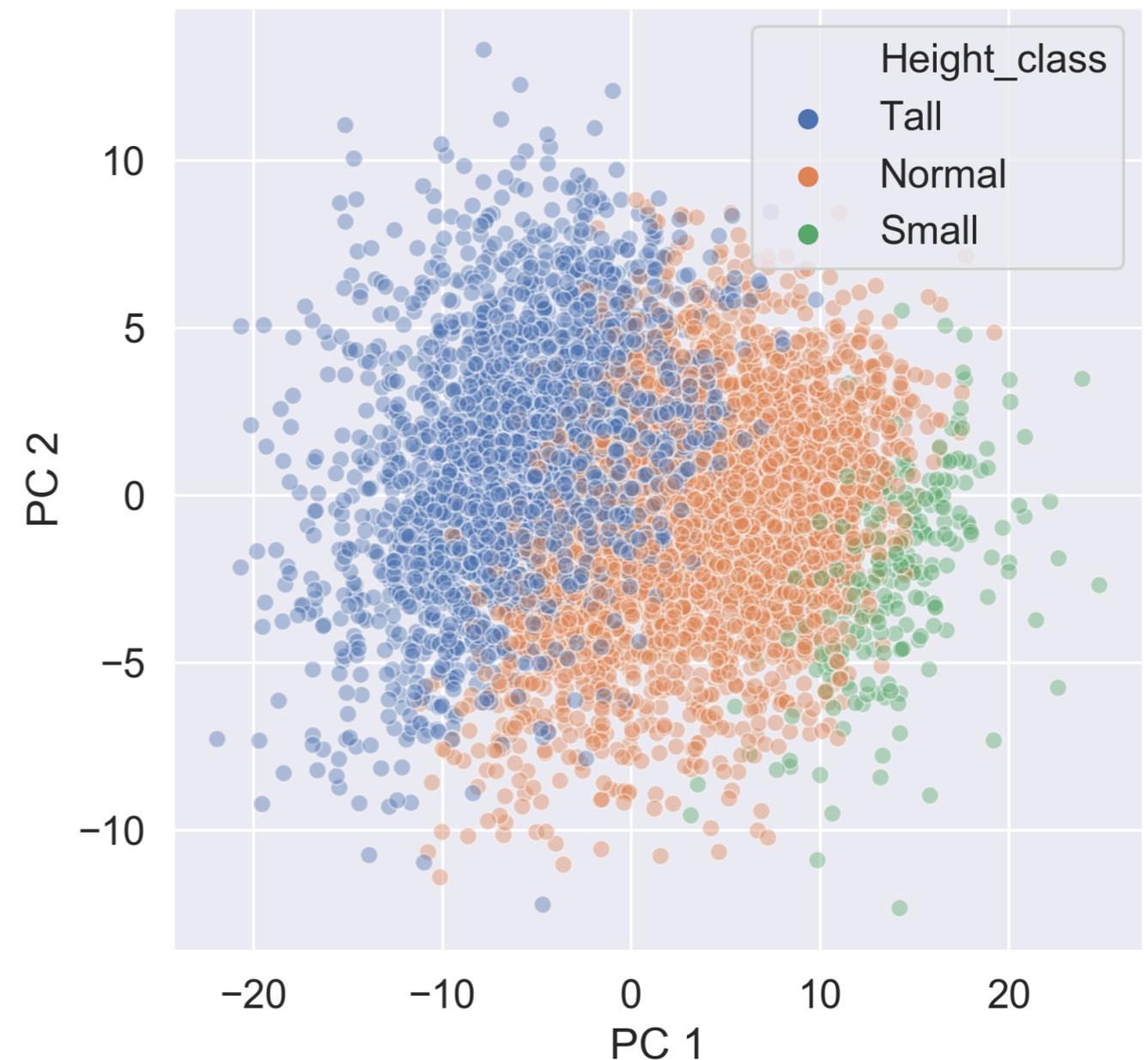
# Checking the effect of categorical features

```
print(ansur_categories.head())
```

```
   Branch      Component  Gender  BMI_class  Height_class
0  Combat Arms  Regular Army  Male  Overweight  Tall
1  Combat Support  Regular Army  Male  Overweight  Normal
2  Combat Support  Regular Army  Male  Overweight  Normal
3  Combat Service Support  Regular Army  Male  Overweight  Normal
4  Combat Service Support  Regular Army  Male  Overweight  Tall
```

# Checking the effect of categorical features

```
ansur_categories['PC 1'] = pc[:,0]
ansur_categories['PC 2'] = pc[:,1]
sns.scatterplot(data=ansur_categories,
                x='PC 1', y='PC 2',
                hue='Height_class', alpha=0.4)
```



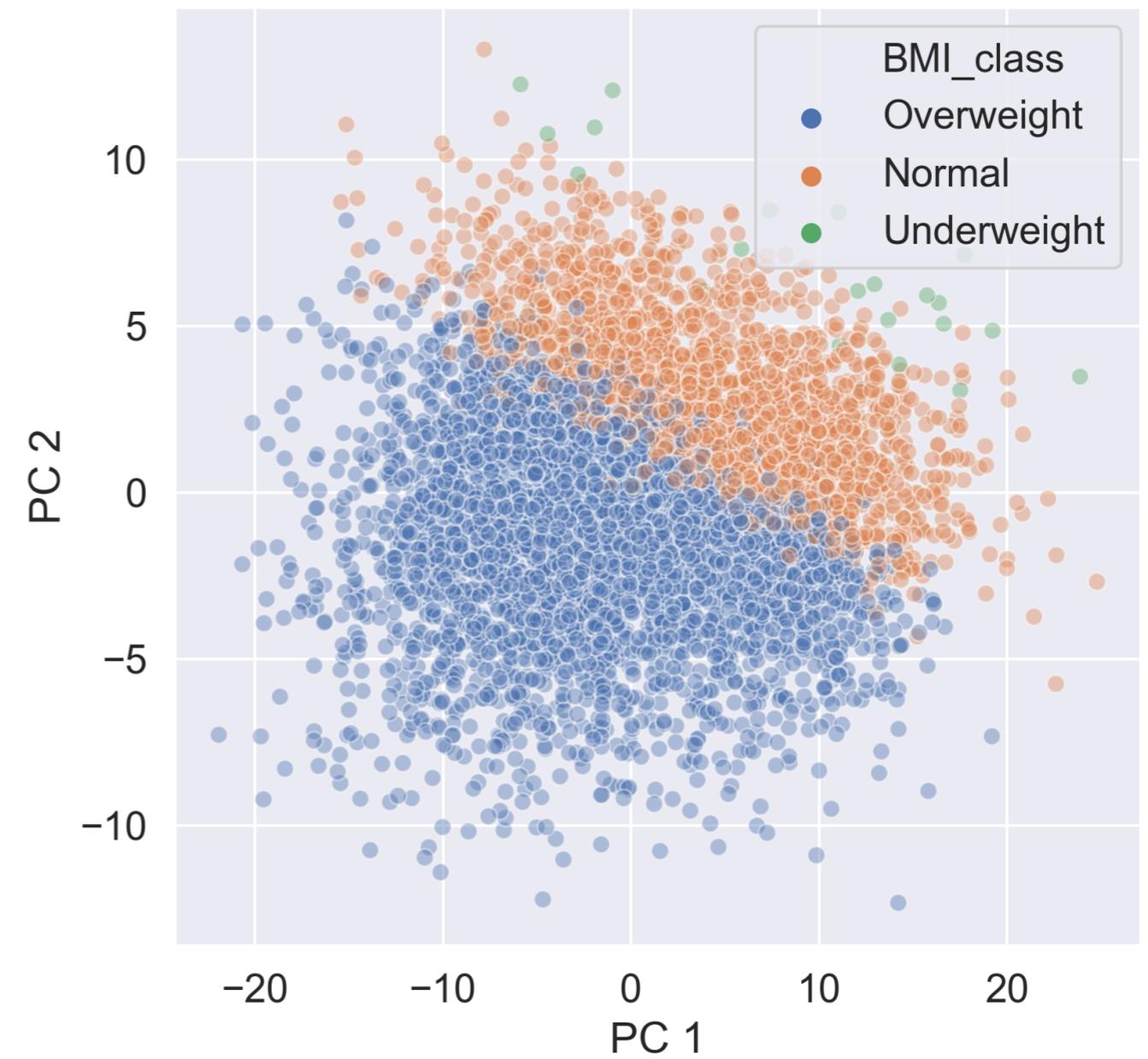
# Checking the effect of categorical features

```
sns.scatterplot(data=ansur_categories,  
               x='PC 1', y='PC 2',  
               hue='Gender', alpha=0.4)
```



# Checking the effect of categorical features

```
sns.scatterplot(data=ansur_categories,  
               x='PC 1', y='PC 2',  
               hue='BMI_class', alpha=0.4)
```



# PCA in a model pipeline

```
pipe = Pipeline([
    ('scaler', StandardScaler()),
    ('reducer', PCA(n_components=3)),
    ('classifier', RandomForestClassifier())])
print(pipe['reducer'])
```

```
PCA(n_components=3)
```

# PCA in a model pipeline

```
pipe.fit(X_train, y_train)  
pipe['reducer'].explained_variance_ratio_
```

```
array([0.56, 0.13, 0.05])
```

```
pipe['reducer'].explained_variance_ratio_.sum()
```

```
0.74
```

```
print(pipe.score(X_test, y_test))
```

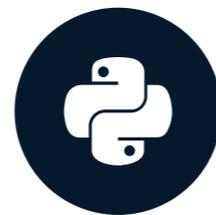
```
0.986
```

# Let's practice!

DIMENSIONALITY REDUCTION IN PYTHON

# Principal Component selection

DIMENSIONALITY REDUCTION IN PYTHON



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Head of Machine Learning, Faktion

# Setting an explained variance threshold

```
pipe = Pipeline([
    ('scaler', StandardScaler()),
    ('reducer', PCA(n_components=0.9))])

# Fit the pipe to the data
pipe.fit(poke_df)

print(len(pipe['reducer'].components_))
```

5

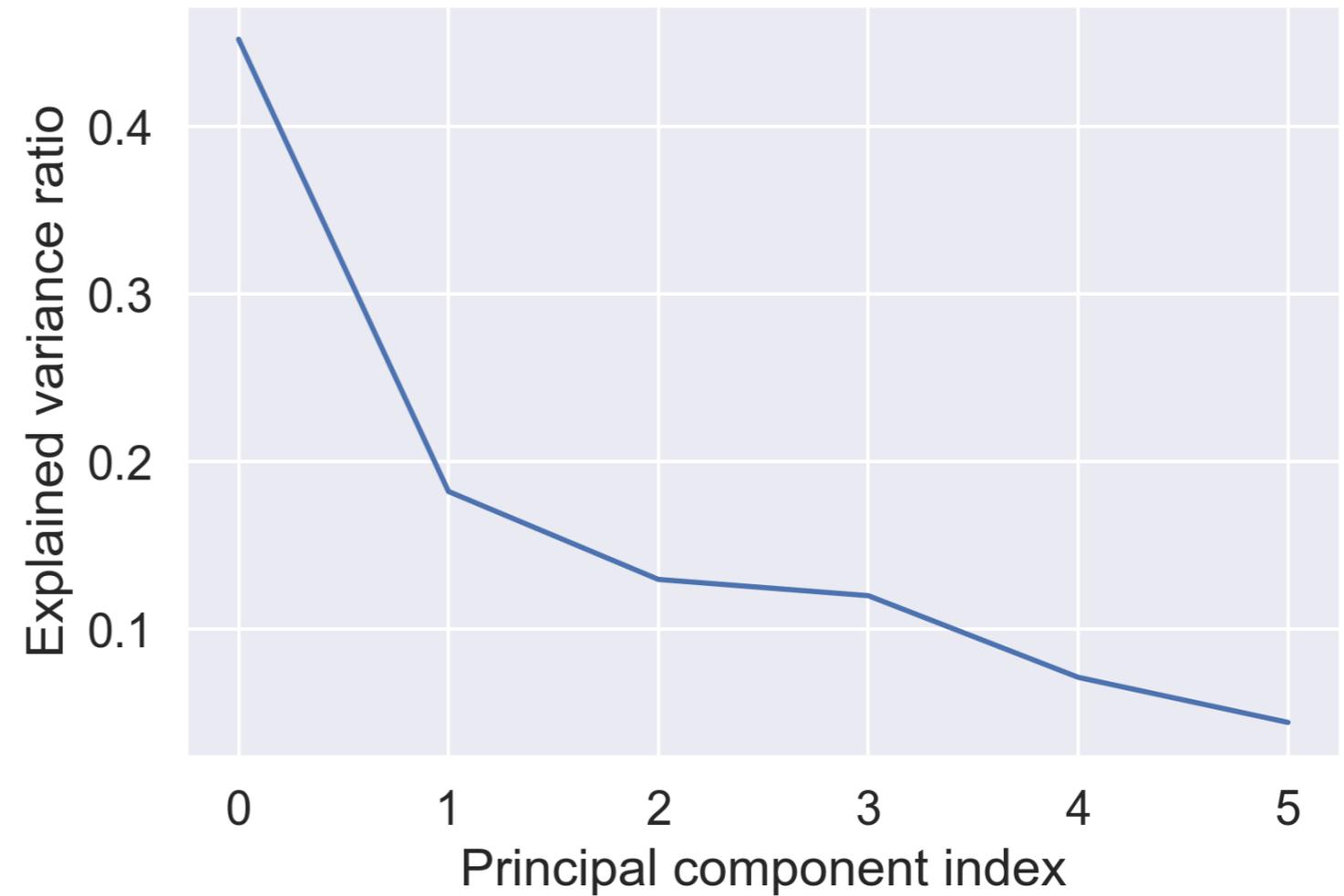
# An optimal number of components

```
pipe.fit(poke_df)

var = pipe['reducer'].explained_variance_ratio_

plt.plot(var)

plt.xlabel('Principal component index')
plt.ylabel('Explained variance ratio')
plt.show()
```



# An optimal number of components

```
pipe.fit(poke_df)

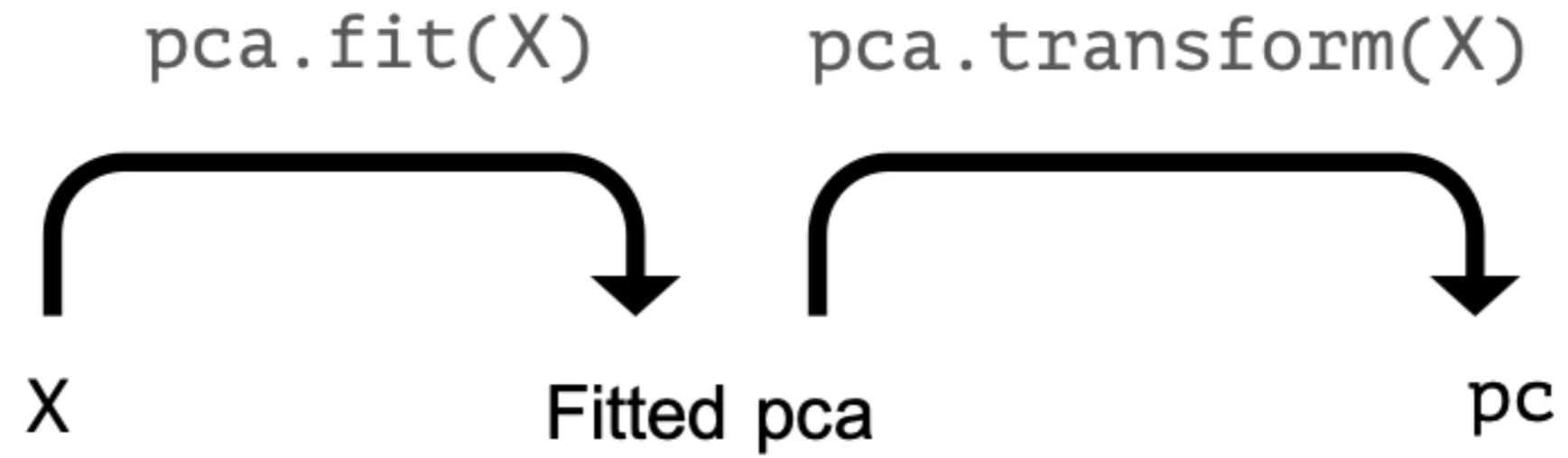
var = pipe['reducer'].explained_variance_ratio_

plt.plot(var)

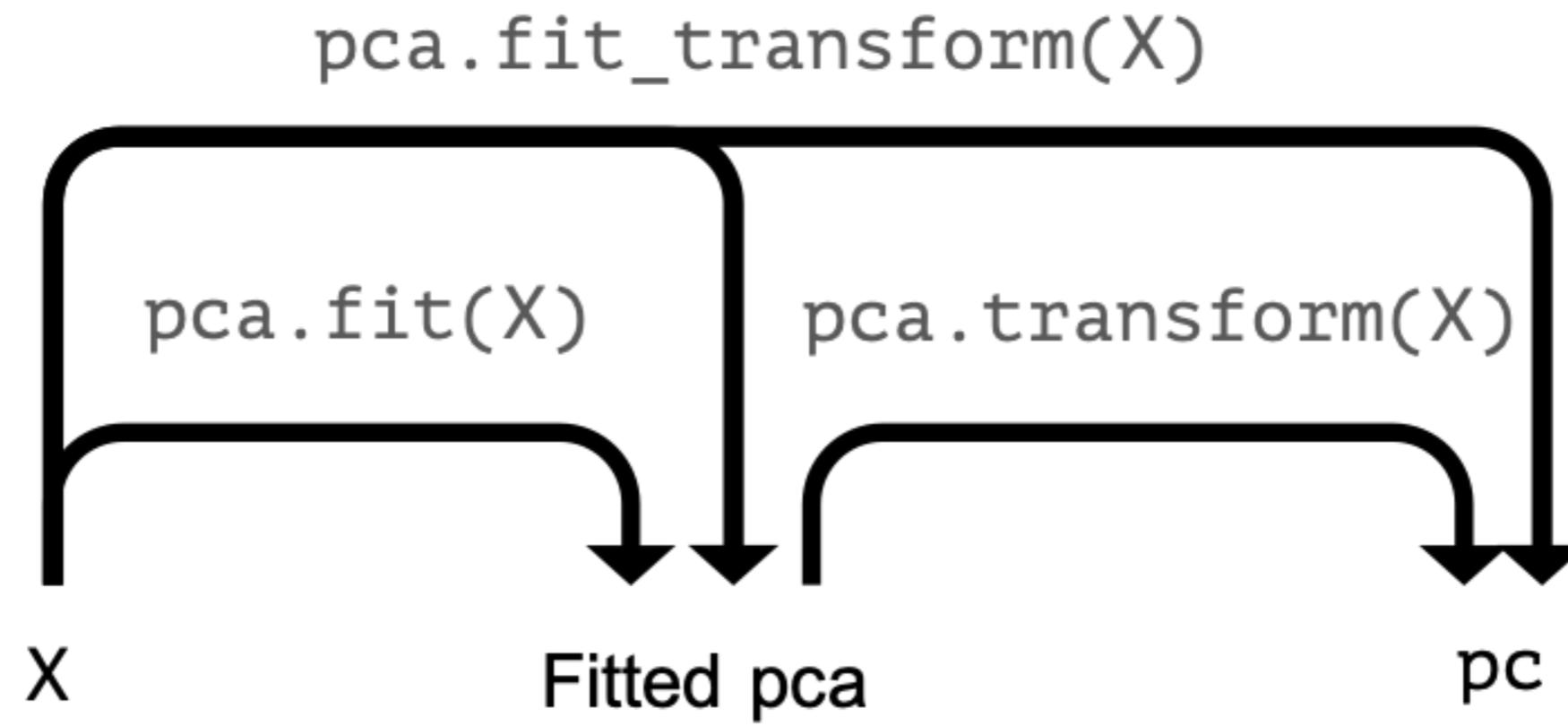
plt.xlabel('Principal component index')
plt.ylabel('Explained variance ratio')
plt.show()
```



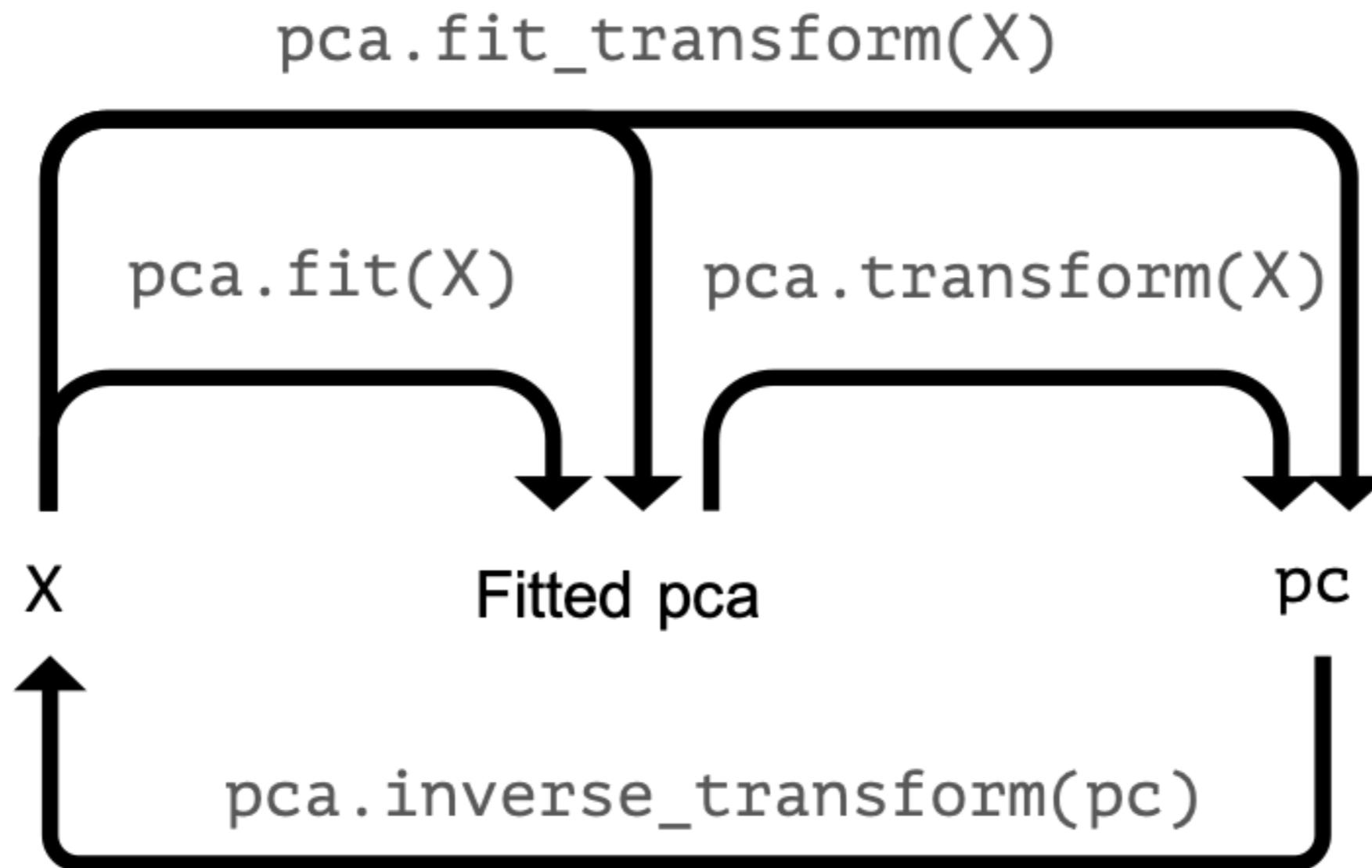
# PCA operations



# PCA operations



# PCA operations



# Compressing images



# Compressing images

```
print(X_test.shape)
```

```
(15, 2914)
```

62 x 47 pixels = 2914 grayscale values

```
print(X_train.shape)
```

```
(1333, 2914)
```

# Compressing images

```
pipe = Pipeline([
    ('scaler', StandardScaler()),
    ('reducer', PCA(n_components=290))]
)
pipe.fit(X_train)
pc = pipe.fit_transform(X_test)

print(pc.shape)
```

```
(15, 290)
```

# Rebuilding images

```
pc = pipe.transform(X_test)
```

```
print(pc.shape)
```

```
(15, 290)
```

```
X_rebuilt = pipe.inverse_transform(pc)
```

```
print(X_rebuilt.shape)
```

```
(15, 2914)
```

```
img_plotter(X_rebuilt)
```



# Rebuilding images

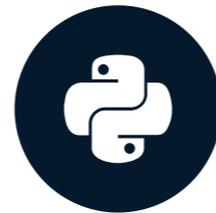


# Let's practice!

DIMENSIONALITY REDUCTION IN PYTHON

# Congratulations!

DIMENSIONALITY REDUCTION IN PYTHON



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# What you've learned

- Why dimensionality reduction is important & when to use it
- Feature selection vs. extraction
- High dimensional data exploration with t-SNE & PCA
- Use models to find important features
- Remove unimportant ones

# Thank you!

DIMENSIONALITY REDUCTION IN PYTHON