# Exploring relationships

**EXPLORATORY DATA ANALYSIS IN PYTHON** 



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#### Height and weight



Q SEARCH

CDC A-Z INDEX V

#### Behavioral Risk Factor Surveillance System







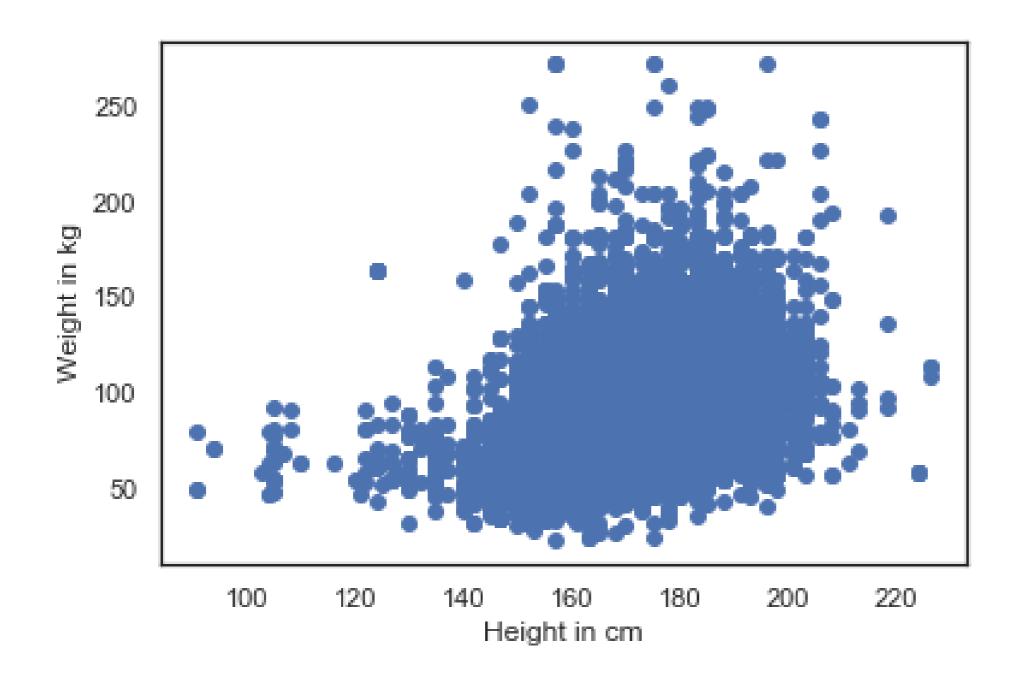


The Behavioral Risk Factor Surveillance System (BRFSS) is the nation's premier system of health-related telephone surveys that collect state data about U.S. residents regarding their health-related risk behaviors, chronic health conditions, and use of preventive services. Established in 1984 with 15 states, BRFSS now collects data in all 50 states as well as the District of Columbia and three U.S. territories. BRFSS completes more than 400,000 adult interviews each year, making it the largest continuously conducted health survey system in the world. See More.



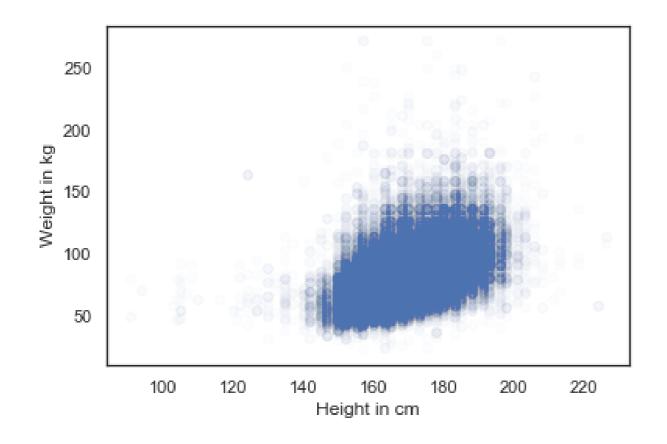
#### Scatter plot

```
brfss = pd.read_hdf('brfss.hdf5', 'brfss')
height = brfss['HTM4']
weight = brfss['WTKG3']
plt.plot(height, weight, 'o')
plt.xlabel('Height in cm')
plt.ylabel('Weight in kg')
plt.show()
```



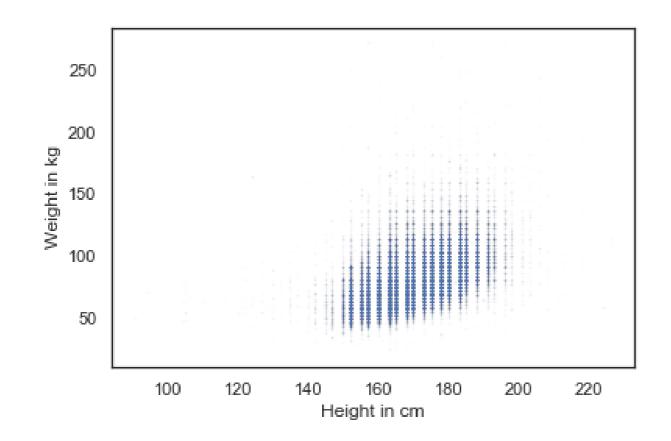
#### **Transparency**

```
plt.plot(height, weight, 'o', alpha=0.02)
plt.show()
```



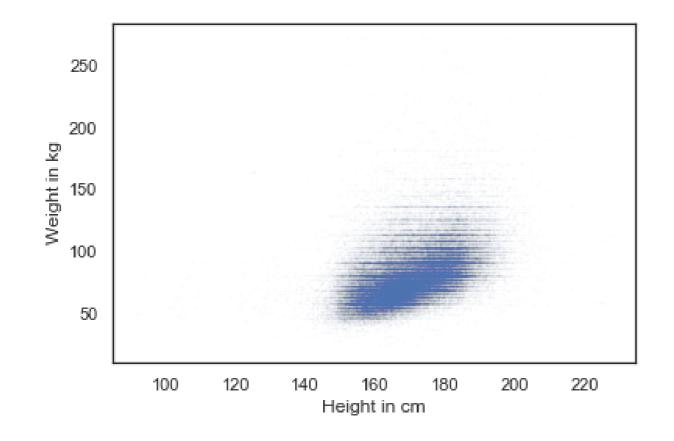
#### Marker size

```
plt.plot(height, weight, 'o', markersize=1, alpha=0.02)
plt.show()
```



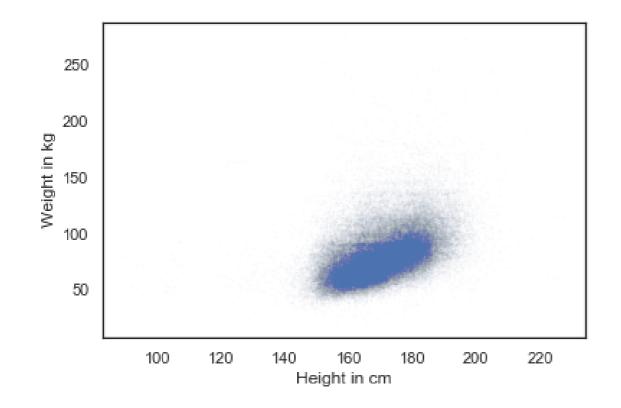
#### **Jittering**

```
height_jitter = height + np.random.normal(0, 2, size=len(brfss))
plt.plot(height_jitter, weight, 'o', markersize=1, alpha=0.02)
plt.show()
```



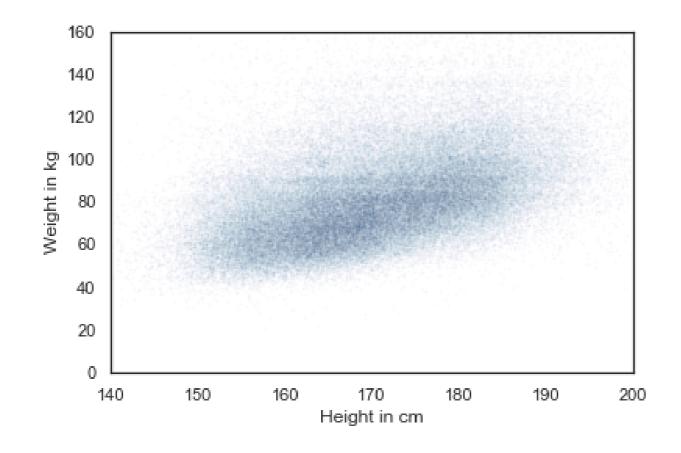
### More jittering

```
height_jitter = height + np.random.normal(0, 2, size=len(brfss))
weight_jitter = weight + np.random.normal(0, 2, size=len(brfss))
plt.plot(height_jitter, weight_jitter, 'o', markersize=1, alpha=0.01
plt.show()
```

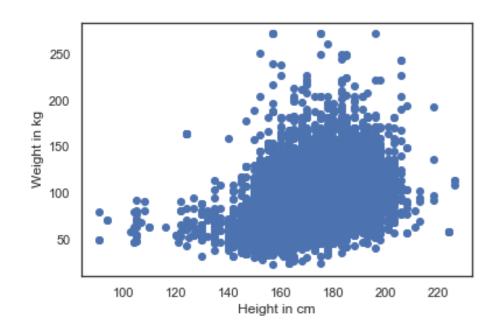


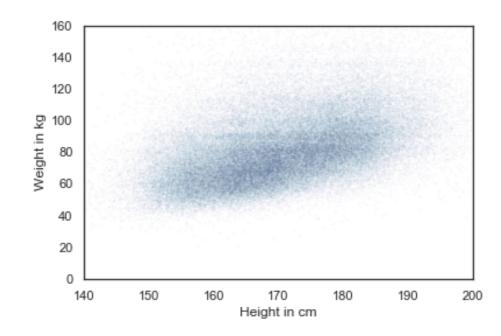
#### Zoom

```
plt.plot(height_jitter, weight_jitter, 'o', markersize=1, alpha=0.02
plt.axis([140, 200, 0, 160])
plt.show()
```



#### Before and after





# Let's explore!

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# Visualizing relationships

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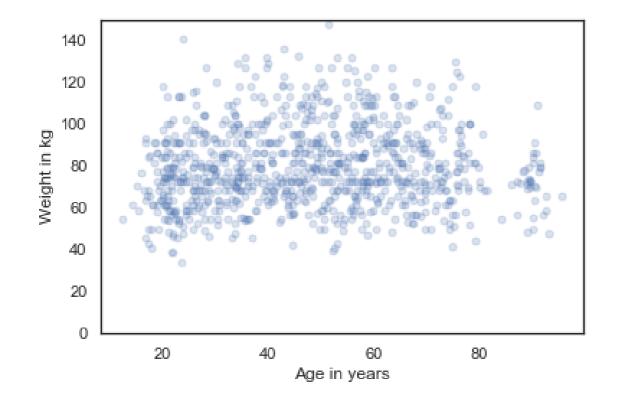


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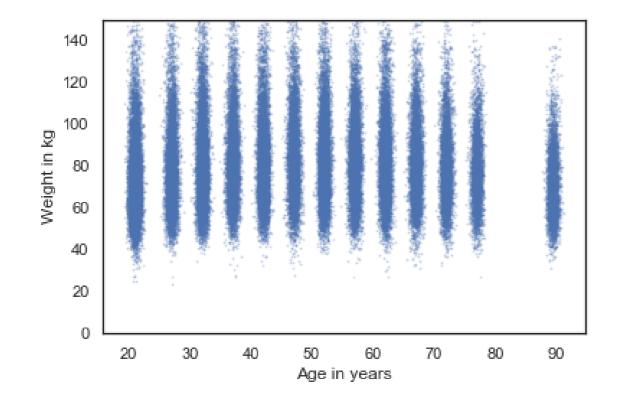
#### Weight and age

```
age = brfss['AGE'] + np.random.normal(0, 2.5, size=len(brfss))
weight = brfss['WTKG3']
plt.plot(age, weight, 'o', markersize=5, alpha=0.2)
plt.show()
```



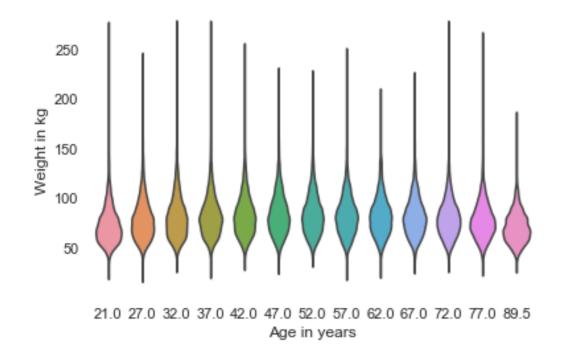
#### More data

```
age = brfss['AGE'] + np.random.normal(0, 0.5, size=len(brfss))
weight = brfss['WTKG3'] + np.random.normal(0, 2, size=len(brfss))
plt.plot(age, weight, 'o', markersize=1, alpha=0.2)
plt.show()
```



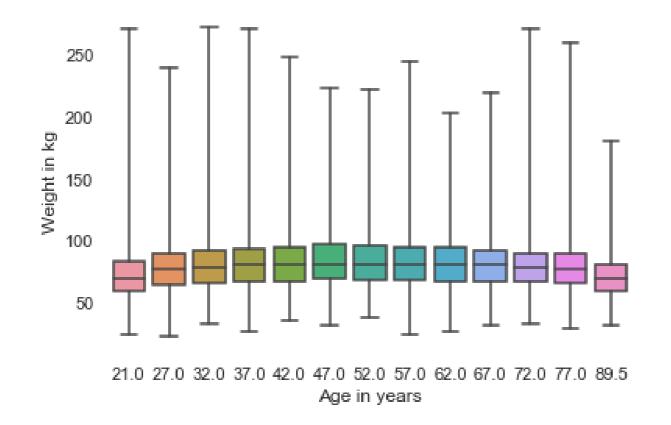
### Violin plot

```
data = brfss.dropna(subset=['AGE', 'WTKG3'])
sns.violinplot(x='AGE', y='WTKG3', data=data, inner=None)
plt.show()
```



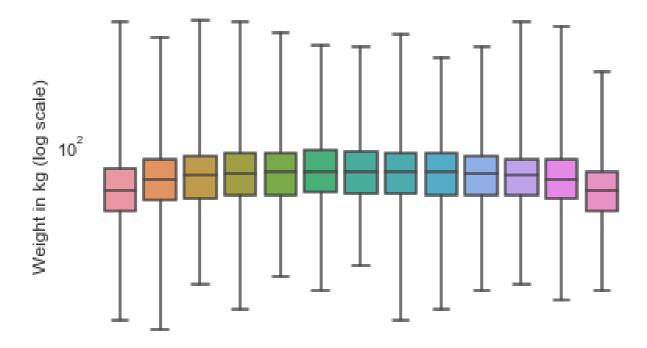
### **Box plot**

```
sns.boxplot(x='AGE', y='WTKG3', data=data, whis=10)
plt.show()
```



### Log scale

```
sns.boxplot(x='AGE', y='WTKG3', data=data, whis=10)
plt.yscale('log')
plt.show()
```



21.0 27.0 32.0 37.0 42.0 47.0 52.0 57.0 62.0 67.0 72.0 77.0 89.5 Age in years

# Let's practice!

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## Correlation

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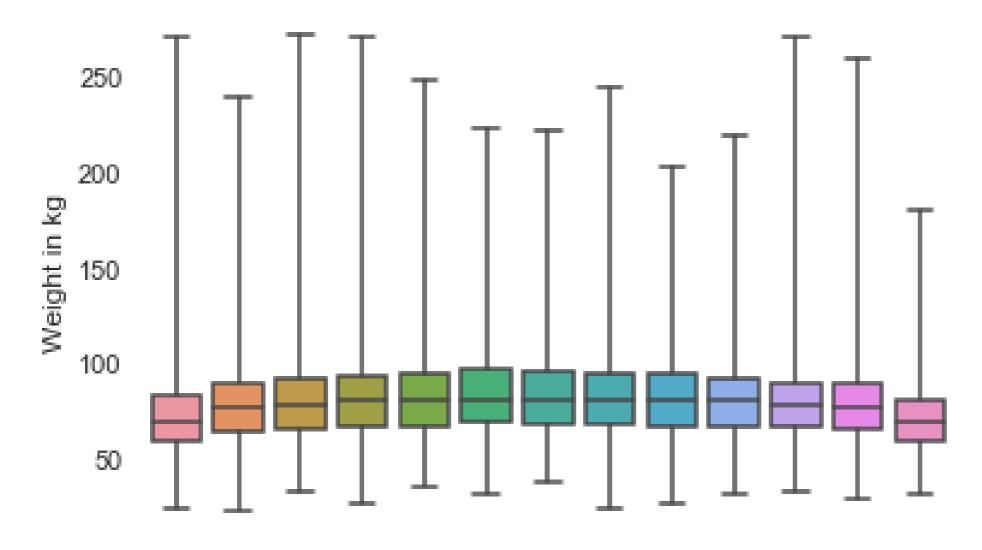
#### **Correlation coefficient**

```
columns = ['HTM4', 'WTKG3', 'AGE']
subset = brfss[columns]
subset.corr()
```

#### **Correlation matrix**

		HTM4	WTKG3	AGE
ı	HTM4	1.000000	0.474203	-0.093684
ı	WTKG3	0.474203	1.000000	0.021641
ı	AGE	-0.093684	0.021641	1.000000

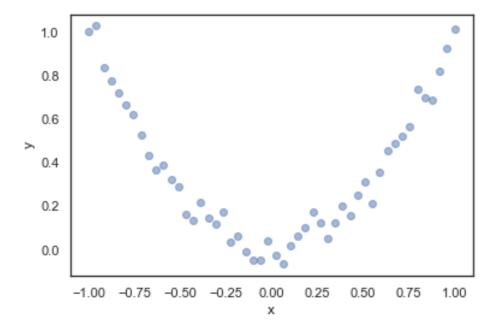
- Height with itself: 1
- Height and weight: 0.47
- Height and age: -0.09
- Weight and age: 0.02



21.0 27.0 32.0 37.0 42.0 47.0 52.0 57.0 62.0 67.0 72.0 77.0 89.5 Age in years

```
xs = np.linspace(-1, 1)
ys = xs**2
ys += normal(0, 0.05, len(xs))
np.corrcoef(xs, ys)
```

```
array([[ 1. , -0.01111647],
[-0.01111647, 1. ]])
```



### You keep using that word

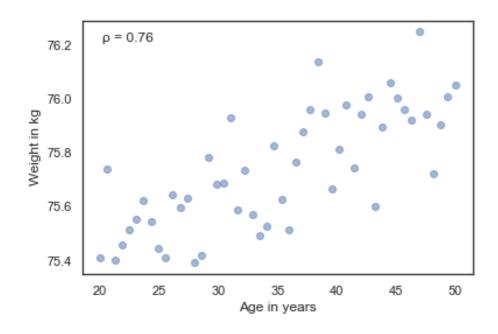
I do not think it means what you think it means

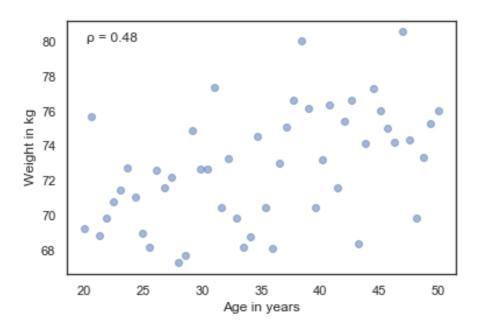


#### Strength of relationship

Hypothetical #1

Hypothetical #2





# Let's practice!

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# Simple regression

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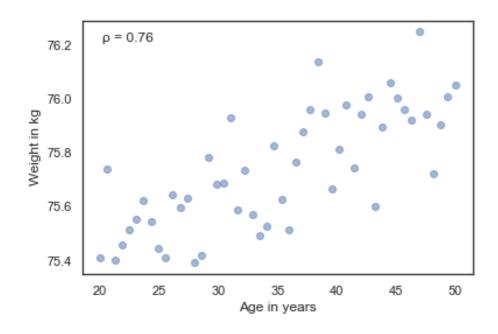
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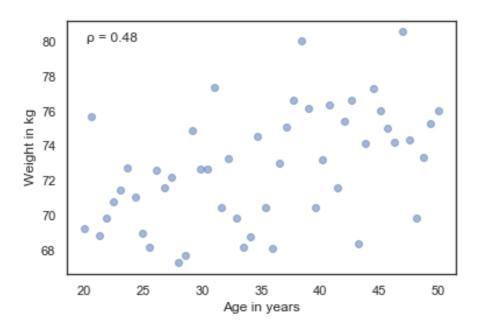


#### Strength of relationship

Hypothetical #1

Hypothetical #2





### Strength of effect

```
from scipy.stats import linregress

# Hypothetical 1
res = linregress(xs, ys)
```

```
LinregressResult(slope=0.018821034903244386,
intercept=75.08049023710964,
rvalue=0.7579660563439402,
pvalue=1.8470158725246148e-10,
stderr=0.002337849260560818)
```



### Strength of effect

```
# Hypothetical 2
res = linregress(xs, ys)
```

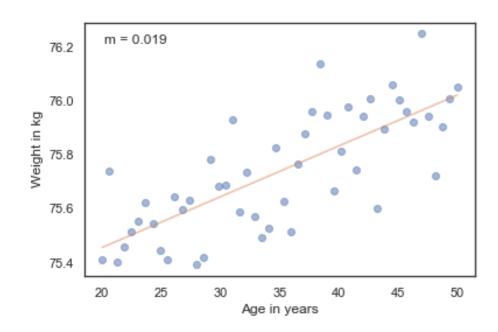
```
LinregressResult(slope=0.17642069806488855,
intercept=66.60980474219305,
rvalue=0.47827769765763173,
pvalue=0.0004430600283776241,
stderr=0.04675698521121631)
```

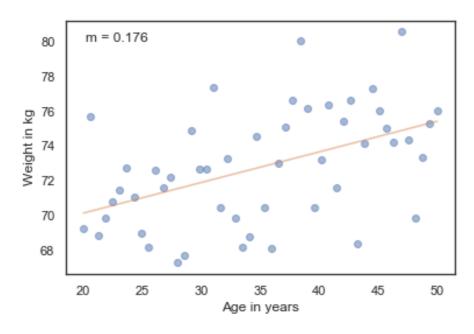


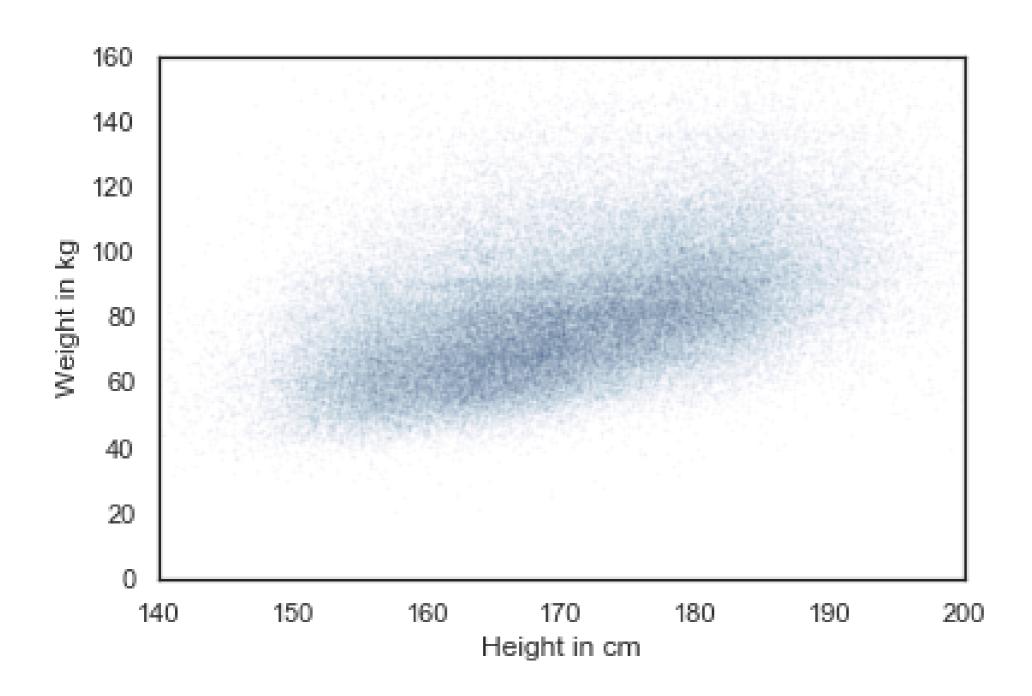
#### Regression lines

```
fx = np.array([xs.min(), xs.max()])
fy = res.intercept + res.slope * fx
plt.plot(fx, fy, '-')
```

```
fx = ...
fy = ...
plt.plot(fx, fy, '-')
```





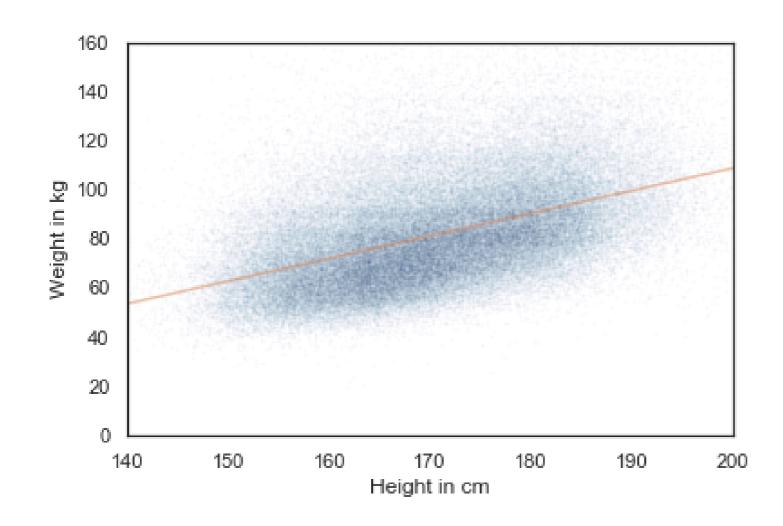


#### Regression line

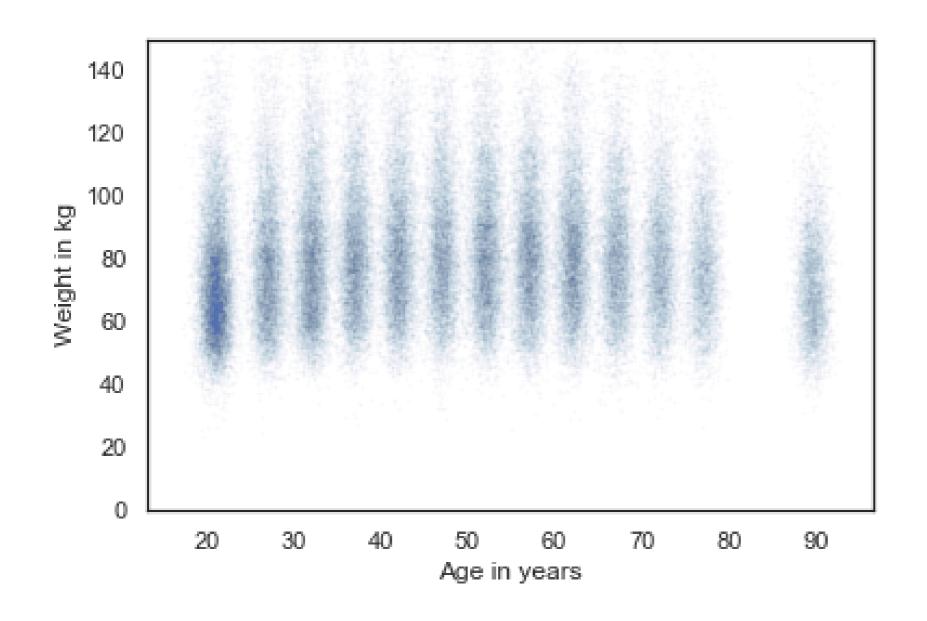
```
subset = brfss.dropna(subset=['WTKG3', 'HTM4'])

xs = subset['HTM4']
ys = subset['WTKG3']
res = linregress(xs, ys)
```

```
fx = np.array([xs.min(), xs.max()])
fy = res.intercept + res.slope * fx
plt.plot(fx, fy, '-')
```



### Linear relationships

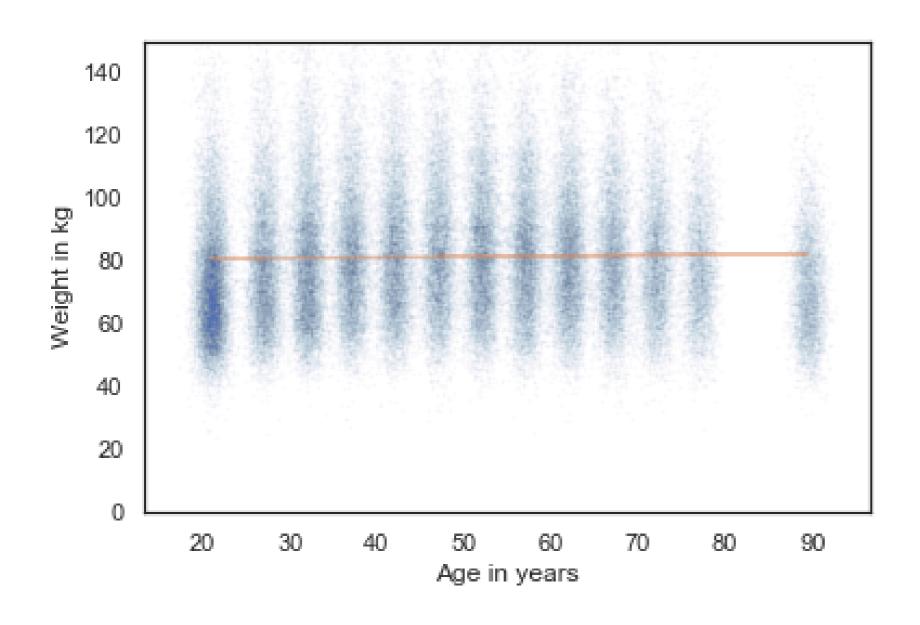


#### Nonlinear relationships

```
subset = brfss.dropna(subset=['WTKG3', 'AGE'])
xs = subset['AGE']
ys = subset['WTKG3']
res = linregress(xs, ys)
```



### Not a good fit



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