data_checks

January 16, 2023

[1]:
```
from matplotlib.pyplot import subplots
from pandas import read_csv
```

1 EEG Eye State

A worked out solution for this data set is provided.

[2]:
```
DATA_PATH = 'data/eeg_eye_state.csv'

df = read_csv(DATA_PATH)

print(df.dtypes)
print('')

columns = df.columns

print(columns)
print('')

df.head()
```

```
AF3             float64
F7              float64
F3              float64
FC5             float64
T7              float64
P7              float64
O1              float64
O2              float64
P8              float64
T8              float64
FC6             float64
F4              float64
F8              float64
AF4             float64
eyeDetection       bool
dtype: object
```

```
Index(['AF3', 'F7', 'F3', 'FC5', 'T7', 'P7', 'O1', 'O2', 'P8', 'T8', 'FC6',
       'F4', 'F8', 'AF4', 'eyeDetection'],
      dtype='object')
```

[2]:

	AF3	F7	F3	FC5	T7	P7	O1	O2	\
0	4329.23	4009.23	4289.23	4148.21	4350.26	4586.15	4096.92	4641.03	
1	4324.62	4004.62	4293.85	4148.72	4342.05	4586.67	4097.44	4638.97	
2	4327.69	4006.67	4295.38	4156.41	4336.92	4583.59	4096.92	4630.26	
3	4328.72	4011.79	4296.41	4155.90	4343.59	4582.56	4097.44	4630.77	
4	4326.15	4011.79	4292.31	4151.28	4347.69	4586.67	4095.90	4627.69	

	P8	T8	FC6	F4	F8	AF4	eyeDetection
0	4222.05	4238.46	4211.28	4280.51	4635.90	4393.85	False
1	4210.77	4226.67	4207.69	4279.49	4632.82	4384.10	False
2	4207.69	4222.05	4206.67	4282.05	4628.72	4389.23	False
3	4217.44	4235.38	4210.77	4287.69	4632.31	4396.41	False
4	4210.77	4244.10	4212.82	4288.21	4632.82	4398.46	False

2 Breast Cancer

Taken from UCI repository.

[3]:
```
DATA_PATH = 'data/breast_cancer.csv'

df = read_csv(DATA_PATH)

print(df.dtypes)
print('')

columns = df.columns

print(columns)
print('')

print('Rows: ', len(df))
print('')

df.head()
```

```
diagnosis                  int64
radius_mean              float64
texture_mean             float64
perimeter_mean           float64
area_mean                float64
smoothness_mean          float64
compactness_mean         float64
concavity_mean           float64
```

```
concave points_mean        float64
symmetry_mean              float64
fractal_dimension_mean     float64
radius_se                  float64
texture_se                 float64
perimeter_se               float64
area_se                    float64
smoothness_se              float64
compactness_se             float64
concavity_se               float64
concave points_se          float64
symmetry_se                float64
fractal_dimension_se       float64
radius_worst               float64
texture_worst              float64
perimeter_worst            float64
area_worst                 float64
smoothness_worst           float64
compactness_worst          float64
concavity_worst            float64
concave points_worst       float64
symmetry_worst             float64
fractal_dimension_worst    float64
dtype: object

Index(['diagnosis', 'radius_mean', 'texture_mean', 'perimeter_mean',
       'area_mean', 'smoothness_mean', 'compactness_mean', 'concavity_mean',
       'concave points_mean', 'symmetry_mean', 'fractal_dimension_mean',
       'radius_se', 'texture_se', 'perimeter_se', 'area_se', 'smoothness_se',
       'compactness_se', 'concavity_se', 'concave points_se', 'symmetry_se',
       'fractal_dimension_se', 'radius_worst', 'texture_worst',
       'perimeter_worst', 'area_worst', 'smoothness_worst',
       'compactness_worst', 'concavity_worst', 'concave points_worst',
       'symmetry_worst', 'fractal_dimension_worst'],
      dtype='object')

Rows:  569
```

[3]:

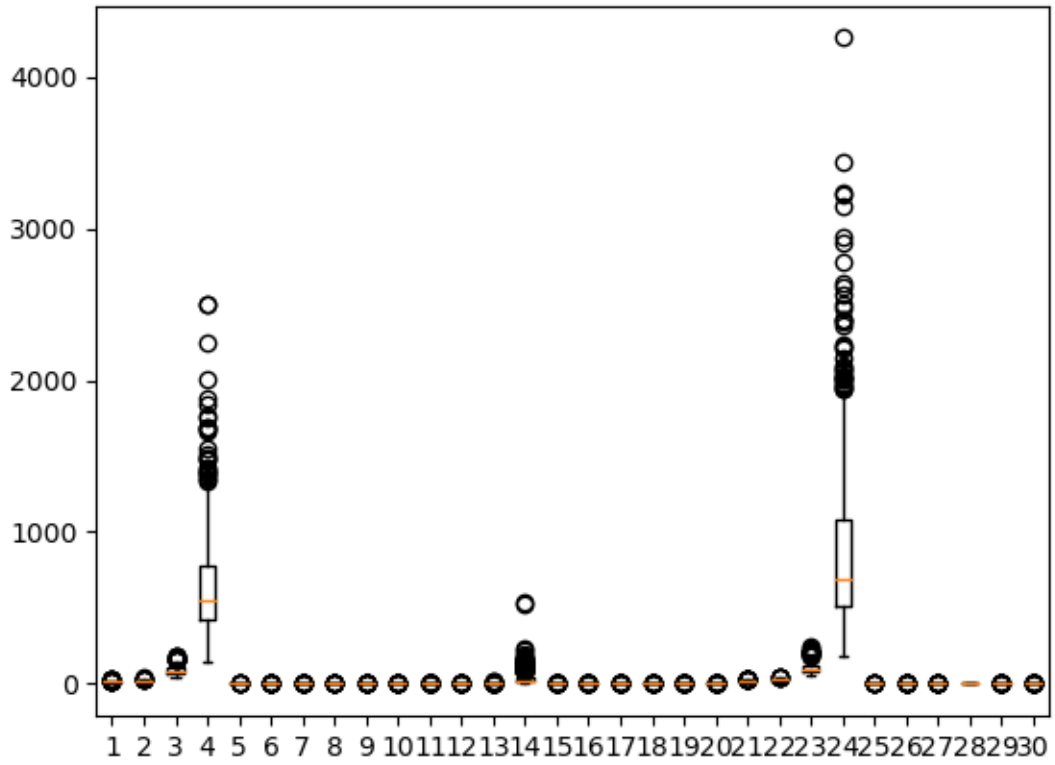
	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	\
0	0	17.99	10.38	122.80	1001.0	
1	0	20.57	17.77	132.90	1326.0	
2	0	19.69	21.25	130.00	1203.0	
3	0	11.42	20.38	77.58	386.1	
4	0	20.29	14.34	135.10	1297.0	

	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

   symmetry_mean  ...  radius_worst  texture_worst  perimeter_worst  \
0         0.2419  ...         25.38          17.33           184.60
1         0.1812  ...         24.99          23.41           158.80
2         0.2069  ...         23.57          25.53           152.50
3         0.2597  ...         14.91          26.50            98.87
4         0.1809  ...         22.54          16.67           152.20

   area_worst  smoothness_worst  compactness_worst  concavity_worst  \
0      2019.0            0.1622             0.6656           0.7119
1      1956.0            0.1238             0.1866           0.2416
2      1709.0            0.1444             0.4245           0.4504
3       567.7            0.2098             0.8663           0.6869
4      1575.0            0.1374             0.2050           0.4000

   concave points_worst  symmetry_worst  fractal_dimension_worst
0                0.2654          0.4601                  0.11890
1                0.1860          0.2750                  0.08902
2                0.2430          0.3613                  0.08758
3                0.2575          0.6638                  0.17300
4                0.1625          0.2364                  0.07678

[5 rows x 31 columns]
```

[4]:
```
data = df.to_numpy()

data = data[:, 1:]

print('Data shape: ', data.shape)
print('')


fig, ax = subplots()

ax.boxplot(data);
```

```
Data shape:  (569, 30)
```



3 Cervical Cancer

Also from the UCI repository.

[5]:
```
DATA_PATH = 'data/cervical_cancer.csv'

df = read_csv(DATA_PATH)

print(df.dtypes)
print('')

columns = df.columns

print(columns)
print('')

print('Rows: ', len(df))
print('')

df.head()
```

```
Age                                   int64
Number of sexual partners           float64
First sexual intercourse            float64
Num of pregnancies                  float64
Smokes                              float64
Smokes (years)                      float64
Smokes (packs/year)                 float64
Hormonal Contraceptives             float64
Hormonal Contraceptives (years)     float64
IUD                                 float64
IUD (years)                         float64
STDs                                float64
STDs (number)                       float64
STDs:condylomatosis                 float64
STDs:cervical condylomatosis        float64
STDs:vaginal condylomatosis         float64
STDs:vulvo-perineal condylomatosis  float64
STDs:syphilis                       float64
STDs:pelvic inflammatory disease    float64
STDs:genital herpes                 float64
STDs:molluscum contagiosum          float64
STDs:AIDS                           float64
STDs:HIV                            float64
STDs:Hepatitis B                    float64
STDs:HPV                            float64
STDs: Number of diagnosis             int64
Dx:Cancer                             int64
Dx:CIN                                int64
Dx:HPV                                int64
Dx                                    int64
Hinselmann                            int64
Schiller                              int64
Citology                              int64
Biopsy                                int64
dtype: object

Index(['Age', 'Number of sexual partners', 'First sexual intercourse',
       'Num of pregnancies', 'Smokes', 'Smokes (years)', 'Smokes (packs/year)',
       'Hormonal Contraceptives', 'Hormonal Contraceptives (years)', 'IUD',
       'IUD (years)', 'STDs', 'STDs (number)', 'STDs:condylomatosis',
       'STDs:cervical condylomatosis', 'STDs:vaginal condylomatosis',
       'STDs:vulvo-perineal condylomatosis', 'STDs:syphilis',
       'STDs:pelvic inflammatory disease', 'STDs:genital herpes',
       'STDs:molluscum contagiosum', 'STDs:AIDS', 'STDs:HIV',
       'STDs:Hepatitis B', 'STDs:HPV', 'STDs: Number of diagnosis',
       'Dx:Cancer', 'Dx:CIN', 'Dx:HPV', 'Dx', 'Hinselmann', 'Schiller',
       'Citology', 'Biopsy'],
      dtype='object')
```

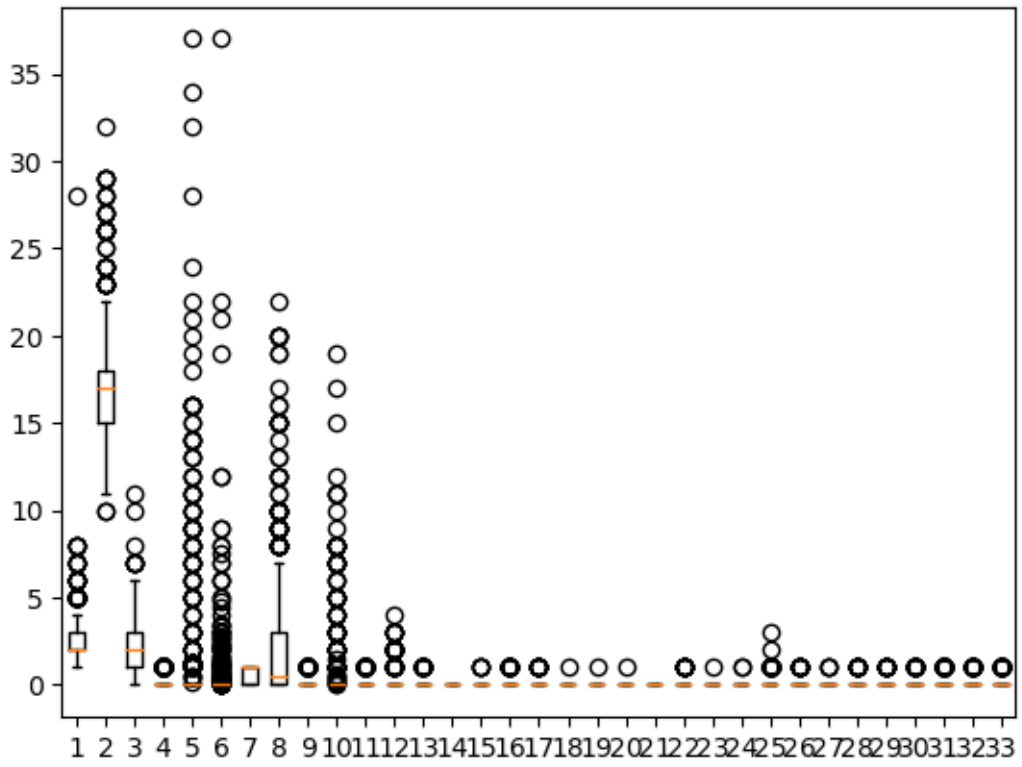
```
Rows:  668
```

[5]:
```
   Age  Number of sexual partners  First sexual intercourse  \
0   18                        4.0                      15.0
1   15                        1.0                      14.0
2   52                        5.0                      16.0
3   46                        3.0                      21.0
4   42                        3.0                      23.0

   Num of pregnancies  Smokes  Smokes (years)  Smokes (packs/year)  \
0                 1.0     0.0             0.0                  0.0
1                 1.0     0.0             0.0                  0.0
2                 4.0     1.0            37.0                 37.0
3                 4.0     0.0             0.0                  0.0
4                 2.0     0.0             0.0                  0.0

   Hormonal Contraceptives  Hormonal Contraceptives (years)  IUD  ...  \
0                      0.0                              0.0  0.0  ...
1                      0.0                              0.0  0.0  ...
2                      1.0                              3.0  0.0  ...
3                      1.0                             15.0  0.0  ...
4                      0.0                              0.0  0.0  ...

   STDs:HPV  STDs: Number of diagnosis  Dx:Cancer  Dx:CIN  Dx:HPV  Dx  \
0       0.0                         0          0       0       0   0
1       0.0                         0          0       0       0   0
2       0.0                         0          1       0       1   0
3       0.0                         0          0       0       0   0
4       0.0                         0          0       0       0   0

   Hinselmann  Schiller  Citology  Biopsy
0           0         0         0       0
1           0         0         0       0
2           0         0         0       0
3           0         0         0       0
4           0         0         0       0

[5 rows x 34 columns]
```

[6]:
```
data = df.to_numpy()

data = data[:, 1:]

print('Data shape: ', data.shape)
print('')
```

```
fig, ax = subplots()

ax.boxplot(data);
```

```
Data shape:  (668, 33)
```



4 Diabetes

[7]:
```
from sklearn import datasets
from pandas import DataFrame

diabetes = datasets.load_diabetes()

data = diabetes.data

df = DataFrame(data)

print(df.dtypes)
print('')
```

```
columns = df.columns

print(columns)
print('')

print('Rows: ', len(df))
print('')

df.head()
```

```
0    float64
1    float64
2    float64
3    float64
4    float64
5    float64
6    float64
7    float64
8    float64
9    float64
dtype: object

RangeIndex(start=0, stop=10, step=1)

Rows:  442
```

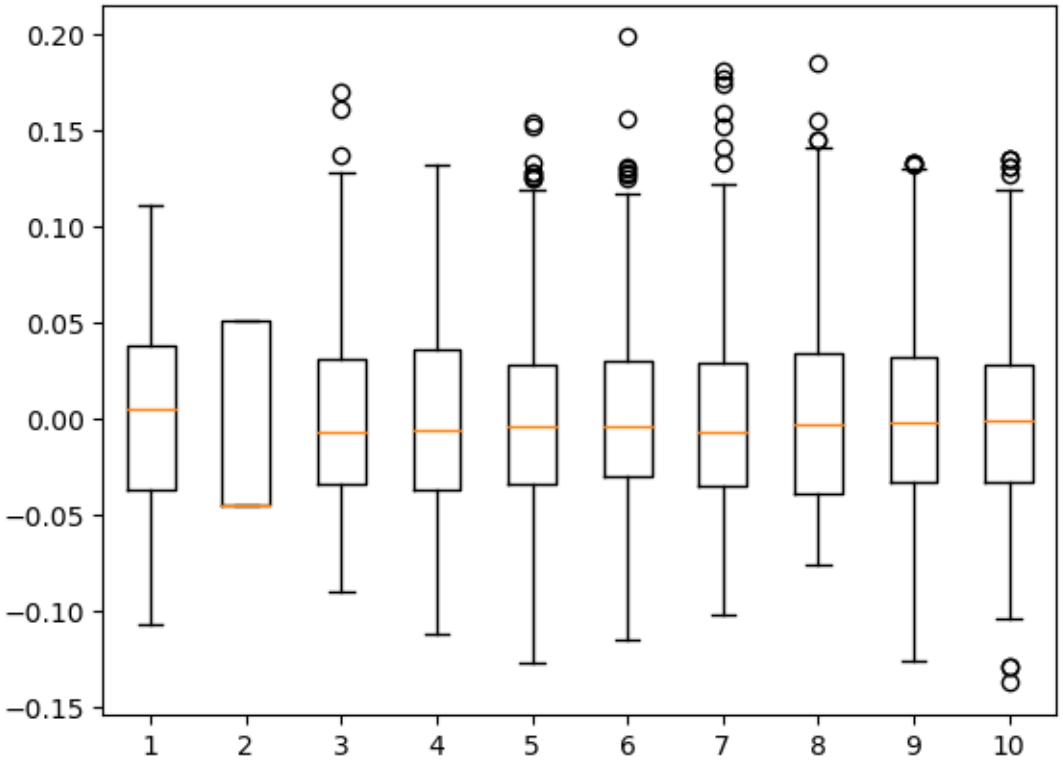
[7]:

	0	1	2	3	4	5	6	7	8	9
0	0.038076	0.050680	0.061696	0.021872	-0.044223	-0.034821	-0.043401	-0.002592	0.019908	-0.017646
1	-0.001882	-0.044642	-0.051474	-0.026328	-0.008449	-0.019163	0.074412	-0.039493	-0.068330	-0.092204
2	0.085299	0.050680	0.044451	-0.005671	-0.045599	-0.034194	-0.032356	-0.002592	0.002864	-0.025930
3	-0.089063	-0.044642	-0.011595	-0.036656	0.012191	0.024991	-0.036038	0.034309	0.022692	-0.009362
4	0.005383	-0.044642	-0.036385	0.021872	0.003935	0.015596	0.008142	-0.002592	-0.031991	-0.046641

[8]:
```
data = df.to_numpy()

data = data[:, :]

print('Data shape: ', data.shape)
print('')
```

```
fig, ax = subplots()

ax.boxplot(data);
```

```
Data shape:  (442, 10)
```



5 Echocardiogram

[9]:
```
DATA_PATH = 'data/echocardio.csv'

df = read_csv(DATA_PATH)

print(df.dtypes)
print('')

columns = df.columns

print(columns)
```

```
print('')

print('Rows: ', len(df))
print('')

df.head()
```

```
attack age               float64
pericardial effusion       int64
fractional shortening    float64
epss                     float64
lvdd                     float64
wall motion score        float64
wall motion index        float64
alive in a year             bool
survival                 float64
alive                      int64
dtype: object

Index(['attack age', 'pericardial effusion', 'fractional shortening', 'epss',
       'lvdd', 'wall motion score', 'wall motion index', 'alive in a year',
       'survival', 'alive'],
      dtype='object')

Rows:  61
```

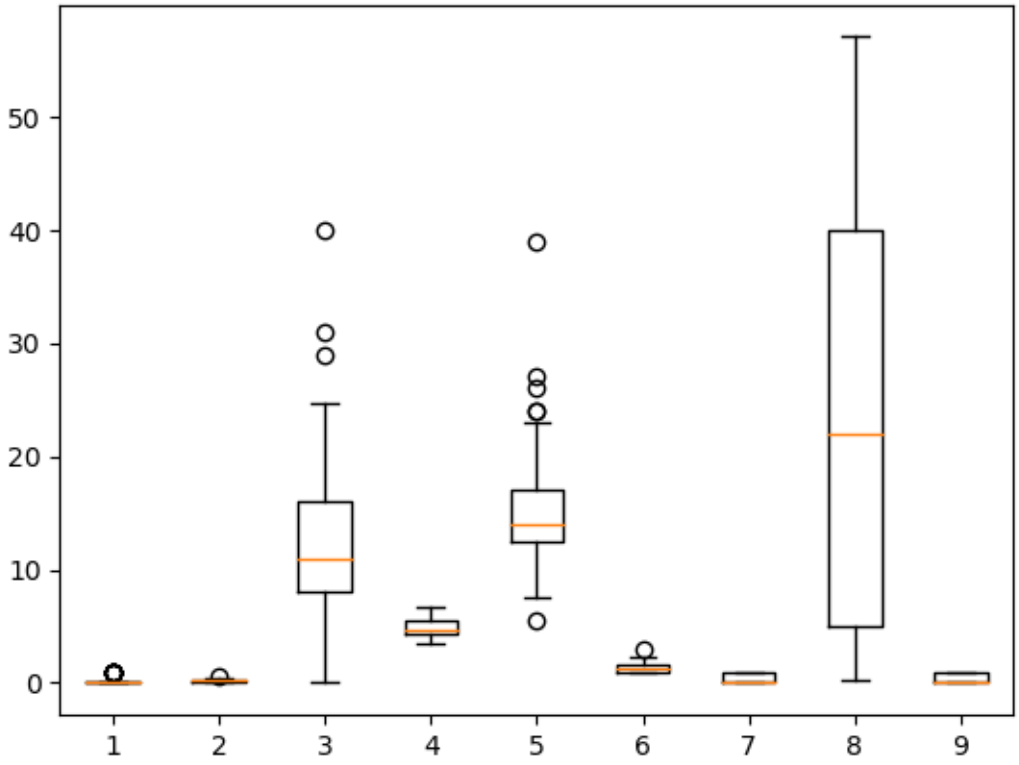
[9]:

	attack age	pericardial effusion	fractional shortening	epss	lvdd	wall motion score	wall motion index	alive in a year	survival	alive
0	71.0	0	0.260	9.000	4.600	14.0	1.00	False	11.0	0
1	72.0	0	0.380	6.000	4.100	14.0	1.70	False	19.0	0
2	55.0	0	0.260	4.000	3.420	14.0	1.00	False	16.0	0
3	60.0	0	0.253	12.062	4.603	16.0	1.45	False	57.0	0
4	57.0	0	0.160	22.000	5.750	18.0	2.25	False	19.0	1

[10]:
```
data = df.to_numpy()

data = data[:, 1:]

print('Data shape: ', data.shape)
print('')
```

```
fig, ax = subplots()

ax.boxplot(data);
```

```
Data shape:  (61, 9)
```



6 Gene Expression

From Gene Expression Omnibus

There is an accompanying paper called A Beginner's Guide to Analysis of RNA Sequencing Data

[11]:
```
data_file = "data/GSE116583_transplant.am.htseq.all.rpkm.txt"

df = read_csv(data_file, delimiter='\s+')

column_names = df.columns

print('Number of columns: ', len(column_names))
print('Number of rows: ', len(df))
```

```
print('')

df.head()
```

```
Number of columns:  13
Number of rows:  43430
```

[11]:

	Symbol	N01_AM_Naive_01	N02_AM_Naive_02	N03_AM_Naive_03	\
0	ENSMUSG00000000001	66.567260	72.604388	67.217287	
1	ENSMUSG00000000003	0.000000	0.000000	0.000000	
2	ENSMUSG00000000028	2.407605	2.163268	2.490079	
3	ENSMUSG00000000031	0.000000	0.000000	0.000000	
4	ENSMUSG00000000037	0.009852	0.005464	0.017368	

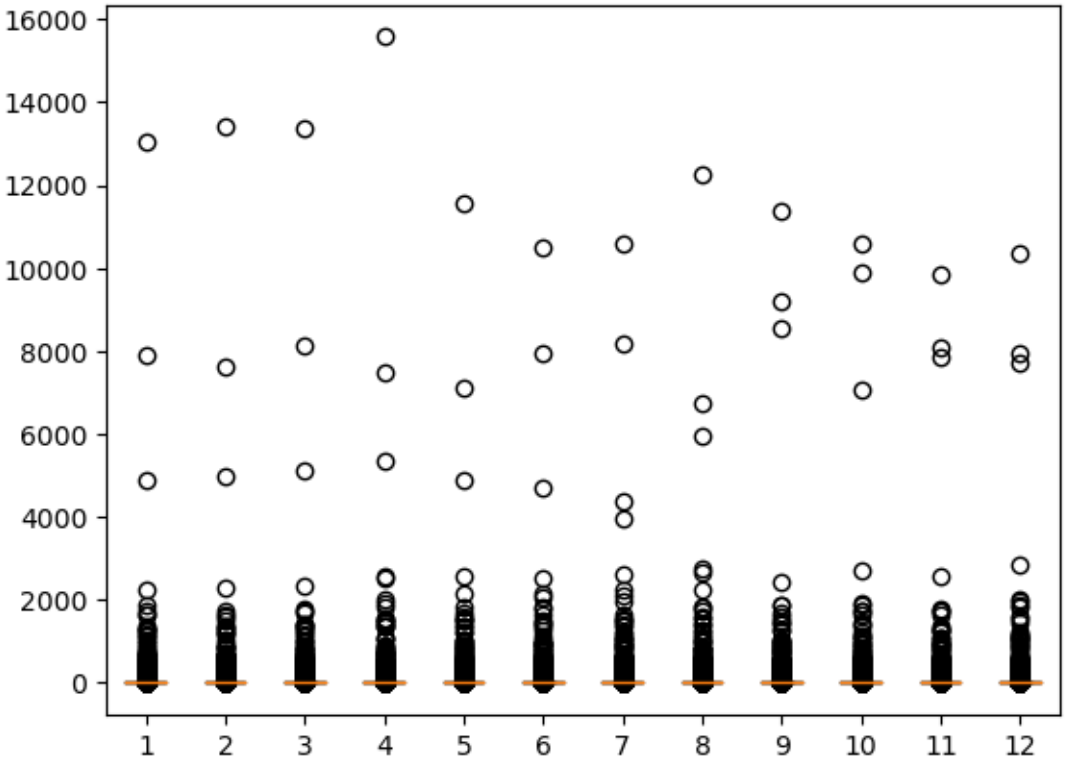
	N04_AM_Naive_04	R01_AM_Allo_02H_01	R02_AM_Allo_02H_02	\
0	70.263406	70.839475	64.616903	
1	0.000000	0.000000	0.000000	
2	1.593701	1.441733	1.002870	
3	0.000000	0.000000	0.000000	
4	0.014177	0.012641	0.000000	

	R03_AM_Allo_02H_03	R04_AM_Allo_02H_04	R05_AM_Allo_24H_01	\
0	69.642272	72.673890	68.579085	
1	0.000000	0.000000	0.000000	
2	1.700592	1.666637	11.861413	
3	0.000000	0.000000	0.000000	
4	0.010085	0.011964	0.028890	

	R06_AM_Allo_24H_02	R07_AM_Allo_24H_03	R08_AM_Allo_24H_04
0	69.754822	68.286563	68.031853
1	0.000000	0.000000	0.000000
2	13.541675	11.626514	9.789112
3	0.000000	0.000000	0.000000
4	0.015992	0.018193	0.018984

[12]:
```
data = df.to_numpy()

data = data[:, 1:]

print('Data shape: ', data.shape)
print('')


fig, ax = subplots()

ax.boxplot(data);
```

```
Data shape:  (43430, 12)
```



7 Imaging or Other Data

If you see something that looks interesting, please contact us with a public link to database or a suggestion for discussion of suitability. Do not use confidential data.

Good success!