In this notebook we work to derive some important information from the paper. We aim to find out

- age distribution
- · distribution of length of time (after first) for check up

And importantly (stat-wise)

We aim to find whether there is a correlation in the age at initial prosthetic implantation and how long it takes for people need a follow up appointment

```
In [1]: # Import the relevant data modules
import numpy as np
import pandas as pd
import matplotlib
import matplotlib.pyplot as pp
import seaborn as sns
from scipy.stats import pearsonr

%matplotlib inline
```

If we first want to visualise the ages in the study we should identify the relevant table (in this case table 2) and work from there

```
In [2]: df = pd.read_csv('Table2extract.csv')
#define variable and attach it to the imported excel comma file from folder
df.head()
#print of only the top of the colloumn (simplified view)
```

Out[2]:

	ID	Age at Prior Fusion (mo)	Age at VEPTR (yr)	Age at Last Follow- up CT (yr)	Length of Follow- up(yr)	Cobb Angle Preoperative	Cobb angle Postoperative	Cobb Angle Last	Thoracic Spine Length (cm) Pre Operative	Thoracic Spine Length (cm) Postoperative	Thoracic Spine Length (cm) Last
0	а	NaN	2.1	7.3	5.2	47.0	29.0	29.0	12.3	15.3	20.7
1	b	NaN	9.9	14.7	4.8	30.0	20.0	32.0	NaN	21.1	26.1
2	С	NaN	1.2	5.2	4.0	49.0	47.0	49.0	107.0	12.2	16.C
3	d	NaN	3.4	8.8	5.4	34.0	49.0	49.0	NaN	NaN	21.2
4	е	NaN	1.0	5.7	4.7	33.0	12.0	16.0	105.0	10.1	13.9

Now we have a visualisation of column location it will help us with the rest, we would like to organise each of the columns into groups (for pie) so we cut out the data into bins of certain value ranges

```
df2 = df.groupby(['Age at Last Follow-up CT (yr)', 'bins2']).size().unstack(fill_value=0
# groupby - groups data!
# size - sums the number of occurences of data
# unstack - pivots the data into a new dataframe where the new bins become collumns
# fillvalue = 0 - makes sure that empty data entry are taken as zero - not always necces
# Display the first DataFrame
print("DataFrame 1:")
print(df1.head())
# Display the second DataFrame
print("\nDataFrame 2:")
print(df2.head())
DataFrame 1:
bins1
                   0-2 2-4 4-6 6-8 8-10
                                             10-12
Age at VEPTR (yr)
0.6
                     1
                          0
                               0
                                    0
                                           0
                                                  0
0.8
                     1
                          0
                               0
                                    0
                                           0
                                                  0
0.9
                     2
                          0
                               0
                                    0
                                           0
                                                  0
1.0
                     2
                                    0
                                           0
                                                  0
                          0
                               0
1.2
                     1
                          0
                               0
                                    0
                                           0
                                                  0
DataFrame 2:
bins2
                               0-1 1-2 2-3 3-4 4-5 5-6
Age at Last Follow-up CT (yr)
                                 2
                                      0
                                            0
                                                      0
                                                           0
1.4
                                                 0
1.9
                                 1
                                       0
                                            0
                                                 0
                                                      0
                                                           0
2.5
                                                      0
                                 0
                                       1
                                            0
                                                 0
                                                           0
3.4
                                 0
                                       1
                                            0
                                                 0
                                                      0
                                                           0
3.5
                                 0
                                       2
                                            0
                                                 0
                                                      0
                                                           0
C:\Users\Luke\AppData\Local\Temp\ipykernel_26896\2923997514.py:11: FutureWarning: The de
fault of observed=False is deprecated and will be changed to True in a future version of
pandas. Pass observed=False to retain current behavior or observed=True to adopt the fut
ure default and silence this warning.
  df1 = df.groupby(['Age at VEPTR (yr)', 'bins1']).size().unstack(fill_value=0)
C:\Users\Luke\AppData\Local\Temp\ipykernel_26896\2923997514.py:12: FutureWarning: The de
fault of observed=False is deprecated and will be changed to True in a future version of
pandas. Pass observed=False to retain current behavior or observed=True to adopt the fut
ure default and silence this warning.
 df2 = df.groupby(['Age at Last Follow-up CT (yr)', 'bins2']).size().unstack(fill_value
```

Now We want to find the magnitude of occurrence of each of the ranges

```
In [4]: # Find the sums
        Total02 = df1['0-2'].sum()
        print(Total02)
        Total24 = df1['2-4'].sum()
        print(Total24)
        Total46 = df1['4-6'].sum()
        print(Total46)
        Total68 = df1['6-8'].sum()
        print(Total68)
        Total810 = df1["8-10"].sum()
        print(Total810)
        Total1012 = df1["10-12"].sum()
        print(Total1012)
        # these functions are simple
        # Variable = dataframe[select data] summing function
        # then print this, again for verification only
```

2

Now we have extracted we re-run to extract a second

```
In [5]: #now the data must be collated again
        Total01 = df2['0-1'].sum()
        print(Total01)
        Total12 = df2['1-2'].sum()
        print(Total12)
        Total23 = df2['2-3'].sum()
        print(Total23)
        Total34 = df2['3-4'].sum()
        print(Total34)
        Total45 = df2['4-5'].sum()
        print(Total45)
        Total56 = df2['5-6'].sum()
        print(Total56)
        # these functions are simple
        # Variable = dataframe[select data] summing function
        # then print this, again for verification only
        7
        5
        6
        4
        6
```

Now, we want to visualise the two extracted dataframes onto piecharts

```
In [6]: # First Pie Chart
        # Crrate a new data frame called df 1 that uses the arrays collated above, indexs each o
        df1 = pd.DataFrame({'Ages': [Total02, Total24, Total46, Total68, Total810, Total1012]},
        # Creates a subplot
        fig, axes = pp.subplots(1, 2, figsize=(14, 7))
        # subplot(rows, columns screen dimensions=14*7)
        # For deviation from default we self-define a color pallete using a seaborn preset
        colors1 = sns.color_palette("Set1")
        # the first pie chart is made
        # pie = var with function pie using df columns and assign to first subplot 'Ages' + cust
        # autopct - creates percentage values for slices
        # explode - highlights slice by extruding
        # shadow - adds 3D effecgt
        # Define titles of the subplot
        axes[0].set_title('Age Distribution at VEPTR insertion')
        # for a cleaner look we remove the label of the y axis - it is unecsessary
        axes[0].set_ylabel('')
        # Legend was overlapping the pies so added this
        legend1 = axes[0].legend(loc='upper left', bbox_to_anchor=(1, 1))
        # adjust legend assigned to subplot 1 position to 'upper left' which is matplot pre-defi
        # Set title on first legend
        legend1.set_title('Age Groups')
        # This code chunk is identical for the second pie chart with some exception, the array d
```

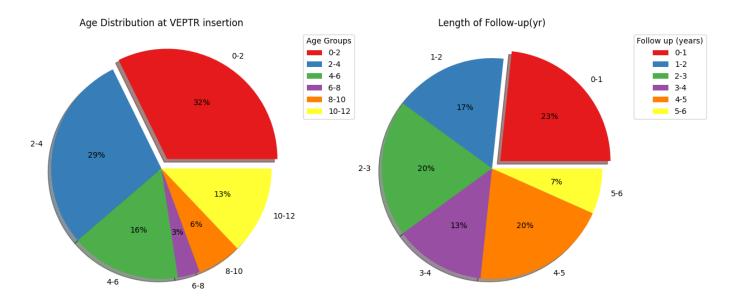
```
# and of course the variable names are different, the strange order is due to the two gr
# they were subplotted

df2 = pd.DataFrame({'Ages': [Total01, Total12, Total23, Total34, Total45, Total56]}, ind
colors2 = sns.color_palette("Set1")
plot2 = df2.plot.pie(y='Ages', autopct='%1.0f%%', ax=axes[1], explode=(0.1, 0, 0, 0, 0, axes[1].set_title('Length of Follow-up(yr)')
axes[1].set_ylabel('')
legend2 = axes[1].legend(loc='upper left', bbox_to_anchor=(1, 1))
legend2.set_title('Follow up (years)')

# Define the title of the two subplots
fig.suptitle('Comparison of Age to follow up length', fontsize=16)

# finally print the stored figures
pp.show()
```

Comparison of Age to follow up length



In [7]: df.head()

Out[7]:

	ID	Age at Prior Fusion (mo)	Age at VEPTR (yr)	Age at Last Follow- up CT (yr)	Length of Follow- up(yr)	Cobb Angle Preoperative	Cobb angle Postoperative	Cobb Angle Last	Thoracic Spine Length (cm) Pre Operative	Thoracic Spine Length (cm) Postoperative	Thoracic Spine Length (cm) Last
0	a	NaN	2.1	7.3	5.2	47.0	29.0	29.0	12.3	15.3	20.7
1	b	NaN	9.9	14.7	4.8	30.0	20.0	32.0	NaN	21.1	26.1
2	С	NaN	1.2	5.2	4.0	49.0	47.0	49.0	107.0	12.2	16.0
3	d	NaN	3.4	8.8	5.4	34.0	49.0	49.0	NaN	NaN	21.2
4	е	NaN	1.0	5.7	4.7	33.0	12.0	16.0	105.0	10.1	13.9

```
In [8]: # Cut out the two ages groups again - we modulated them too much so its easier to re-def
AGEVEP = df.iloc[:,2]
AGEFU = df.iloc[:,3]

# We want to create a new dataframe consisting of the two new arrays we pulled from the
merged_df = pd.merge(AGEFU, AGEVEP, left_index=True, right_index=True)
# we use the merge function and def parameters
```

we want the dataframes to be merged based on their existed indexes so we add it to tru

Due to the specifics of the calculation coming up we need to get rid of Nan values - t
merged_df = merged_df.dropna(subset=['Age at Last Follow-up CT (yr)', 'Age at VEPTR (yr)'

we drop a full print here so we can check rows for NaN
merged_df

Out[8]:		Age at Last Follow-up CT (yr)	Age at VEPTR (yr)
	0	7.3	2.1
	1	14.7	9.9
	2	5.2	1.2
	3	8.8	3.4
	4	5.7	1.0
	5	7.6	3.3
	6	5.6	1.0
	7	7.2	2.5
	8	9.8	5.9
	10	12.9	9.5
	11	9.1	6.5
	12	8.0	4.8
	13	6.4	2.3
	14	5.9	3.8
	15	3.6	1.3
	16	2.5	0.8
	17	12.5	10.2
	18	3.5	1.8
	19	4.2	2.6
	20	6.9	4.4
	21	12.8	10.2
	22	3.5	2.4
	23	3.4	2.0
	24	3.8	2.8
	25	1.9	0.9
	26	4.9	4.4
	27	12.8	12.3
	28	1.4	0.6
	29	1.4	0.9
	30	4.9	4.3

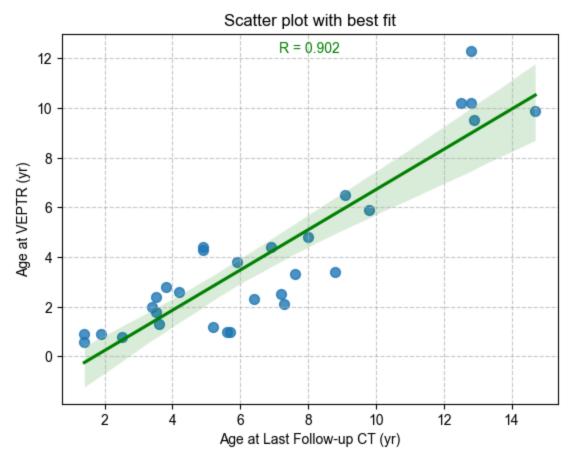
In [9]: # We use scipy module to call def pearson function it computes a corr coef and a two tai Rval,_ = pearsonr(merged_df['Age at Last Follow-up CT (yr)'], merged_df['Age at VEPTR (y # The result is a tuple but the p value isnt relevant for a corr scatter so - unused # we define the params of the calc as the two ages coloumns calling from the merged_df d

```
# Print to verify calc process occur
print(f"Rval: {Rval}")
```

Rval: 0.9016069206365663

Now, the correlation plot (scatter)

```
# We use seaborn here because it can auto-create a line of best fit
In [10]:
         # Create a scatter plot using regression
         ax = sns.regplot(x=AGEFU,y=AGEVEP, data=merged_df, scatter_kws={'s' : 50}, line_kws={'co
         # define X and Y , scatter_kws and line_kws open the settings for the plot and overide t
         # we change size to be 50 , and the line color to be green
         # again changing default settings - change to gridded
         sns.set(style='whitegrid')
         # effect the grid lines created
         ax.grid(True, linestyle='--',alpha=0.7)
         # determine the line type and the transparency
         # Add an annotation using func
         ax.annotate(f'R = \{round(Rval, 3)\}', xy=(0.5, 0.95), xycoords='axes fraction', ha='center'
         # For better veiwing we print the Rval rounded to 3 decs, print Rval at the coords xy=
         # .., define the method of defining coords
         # .., set font size and color of font
         # Set plot title
         ax.set_title('Scatter plot with best fit')
         # print the stored plot
         pp.show()
```



We conclude that there is a **strong and positive correlation** between the two variables, as the age increases the time between appointments also increases