## HousingRegression

December 28, 2020

[4]: #invite people for the Kaggle party

```
import pandas as pd
     import matplotlib.pyplot as plt
     import seaborn as sns
     import numpy as np
     from scipy.stats import norm
     from sklearn.preprocessing import StandardScaler
     from scipy import stats
     import warnings
     warnings.filterwarnings('ignore')
     %matplotlib inline
[38]: df_train = pd.read_csv('/Users/abhishekvijay/Documents/kaggle/housing/train.
      ⇔csv')
      #imports the file path
[6]: df_train.columns
      #displays the columns in the data
[6]: Index(['Id', 'MSSubClass', 'MSZoning', 'LotFrontage', 'LotArea', 'Street',
             'Alley', 'LotShape', 'LandContour', 'Utilities', 'LotConfig',
             'LandSlope', 'Neighborhood', 'Condition1', 'Condition2', 'BldgType',
             'HouseStyle', 'OverallQual', 'OverallCond', 'YearBuilt', 'YearRemodAdd',
             'RoofStyle', 'RoofMatl', 'Exterior1st', 'Exterior2nd', 'MasVnrType',
             'MasVnrArea', 'ExterQual', 'ExterCond', 'Foundation', 'BsmtQual',
             'BsmtCond', 'BsmtExposure', 'BsmtFinType1', 'BsmtFinSF1',
             'BsmtFinType2', 'BsmtFinSF2', 'BsmtUnfSF', 'TotalBsmtSF', 'Heating',
             'HeatingQC', 'CentralAir', 'Electrical', '1stFlrSF', '2ndFlrSF',
             'LowQualFinSF', 'GrLivArea', 'BsmtFullBath', 'BsmtHalfBath', 'FullBath',
             'HalfBath', 'BedroomAbvGr', 'KitchenAbvGr', 'KitchenQual',
             'TotRmsAbvGrd', 'Functional', 'Fireplaces', 'FireplaceQu', 'GarageType',
             'GarageYrBlt', 'GarageFinish', 'GarageCars', 'GarageArea', 'GarageQual',
             'GarageCond', 'PavedDrive', 'WoodDeckSF', 'OpenPorchSF',
             'EnclosedPorch', '3SsnPorch', 'ScreenPorch', 'PoolArea', 'PoolQC',
             'Fence', 'MiscFeature', 'MiscVal', 'MoSold', 'YrSold', 'SaleType',
             'SaleCondition', 'SalePrice'],
            dtype='object')
```

[7]: df\_train['SalePrice'].describe()

#describes the data with a series of output that will important to understand

the data

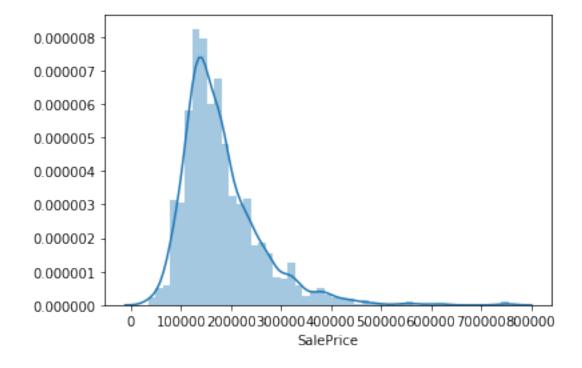
[7]: count 1460.000000 mean 180921.195890 std 79442.502883 min 34900.000000 25% 129975.000000 50% 163000.000000 75% 214000.000000 755000.000000 max

Name: SalePrice, dtype: float64

[92]: #Theres no O values that would potentially ruin the model

[8]: sns.distplot(df\_train['SalePrice'])
#using seaborn to display a distrobution plot

[8]: <matplotlib.axes.\_subplots.AxesSubplot at 0x7f999bde39d0>



[38]: #deviates from the normal distribution
#positive skewness: tail end on the right side is longer
#has a peak

```
[9]: print("skewness: %f" % df_train['SalePrice'].skew())
print("kurtosis: %f" % df_train['SalePrice'].kurt())
#display skewness, still need to figure out %f meaning fully
```

skewness: 1.882876 kurtosis: 6.536282

[40]: #skewness is between >1 which means the data is highly skewed

#kurtosis is >3 AKA leptokurtic which means the data is heavy tailed with

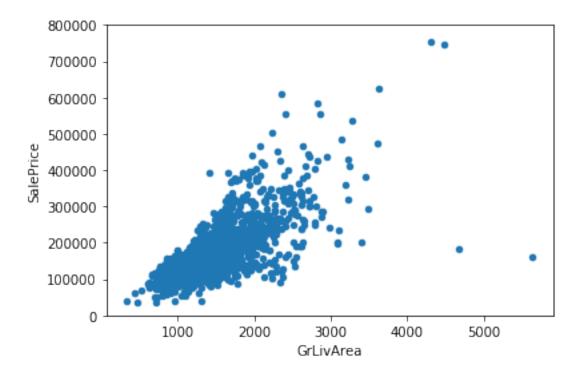
→outliers shown by

#the narrowness of the right tail end

[10]: #lets see the relationship between saleprice and ground living room area

var = 'GrLivArea'
data = pd.concat([df\_train['SalePrice'],df\_train[var]], axis = 1)
#concat used to merge data from saleprice and grlivearea
data.plot.scatter(x=var,y='SalePrice', ylim=(0,800000))
#taking the merged data aka var and displaying it as scatter plot
#by setting the x and y axis variables and setting a limit on the y axis

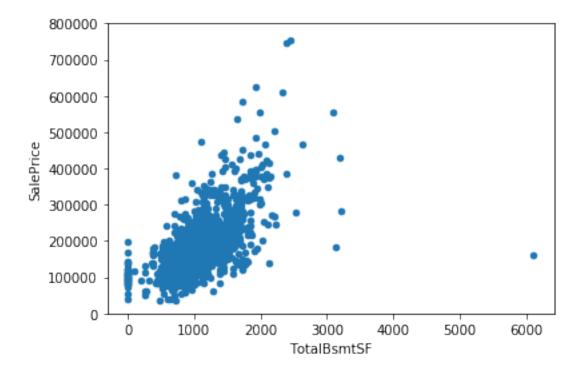
[10]: <matplotlib.axes.\_subplots.AxesSubplot at 0x7f999bfb1d50>



[42]: #there is a linear relationship between saleprice and GrLivArea 
#lets see the relationship between saleprice and total basement square foot $_{\square}$   $_{\longrightarrow}$  TotalBsmntSF

```
[11]: var = 'TotalBsmtSF'
data = pd.concat([df_train['SalePrice'],df_train[var]], axis=1)
data.plot.scatter(x=var,y='SalePrice',ylim=(0,800000))
```

[11]: <matplotlib.axes.\_subplots.AxesSubplot at 0x7f999c955790>



[44]: #strong linear relationship based on the sudden exponential increase #looks like having ~ 500-1500 sq doesnt impact salesprice #lets conduct analysis on the categorical features (not numerical descriptive → traits)

[12]: var = 'OverallQual'

#define OverallQual as the variable being used

data = pd.concat([df\_train['SalePrice'],df\_train[var]],axis = 1)

#concat the data of salesprice and overall quality using pandas

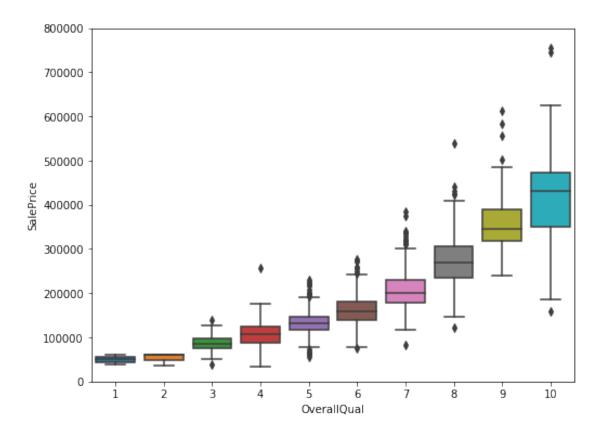
f, ax = plt.subplots(figsize=(8,6))

#plt.sublot is a function that returns a tuple containing a figure and axes

→object(s)

```
#tuple is unpacked into the f and ax variables
fig = sns.boxplot(x=var,y='SalePrice', data=data)
#created the boxplot
fig.axis(ymin=0,ymax=800000)
#setting the axis size
```

#### [12]: (-0.5, 9.5, 0, 800000)

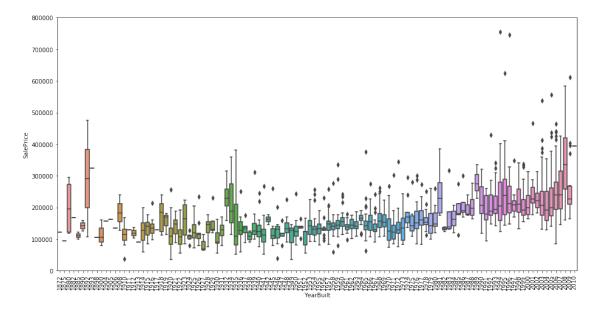


```
[46]: #looks like salesprice is driven up by the overall quality
#lets see how year built looks
```

```
[13]: var = 'YearBuilt'
data = pd.concat([df_train['SalePrice'],df_train[var]], axis = 1)
f, ax = plt.subplots(figsize = (16,8))
fig = sns.boxplot (x=var,y='SalePrice',data=data)
fig.axis(ymin=0,ymax=800000)
plt.xticks(rotation=90)
#use this function to rotate the tick labels 90 degrees to see labels clearly
```

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[13]: (array([ 0,
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                                           96, 97,
                                                      98,
                                                            99, 100, 101, 102, 103,
               104, 105, 106, 107, 108, 109, 110, 111]),
```

<a list of 112 Text xticklabel objects>)



### [48]: #Sales price is driven by the year built although its not a strong relationship

### [49]: #Summary

 $\#Ground\ living\ room\ area\ and\ total\ basement\ square\ footage\ seem\ to\ be\ linearly_\sqcup$  $\rightarrow$ related with Sale price.

#Both are positively related, as one variable increases, the other also.

#Overall quality and year built also show a relationship with salesprice, →although it is not as strong with year built

#lets analyze more

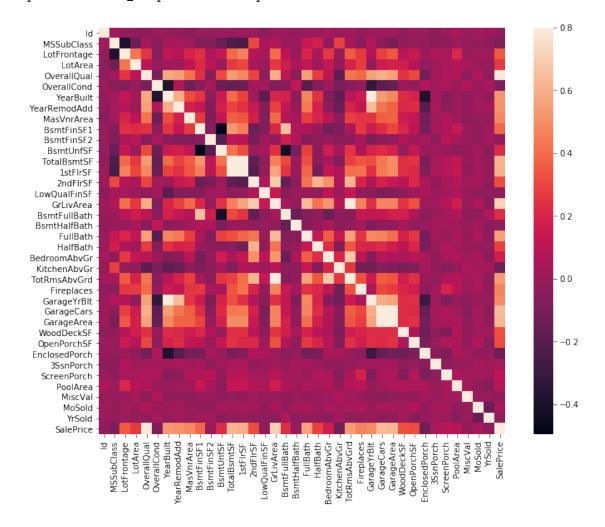
#correlation matrix (heatmap - seaborn)

#salesprice correlation matrix (zoomed heatmap)

#scatter plots between correlated variables

# [14]: #correlation matrix corrmat = df\_train.corr() f, ax = plt.subplots(figsize = (12,9)) sns.heatmap(corrmat, vmax = .8, square=True)

#### [14]: <matplotlib.axes.\_subplots.AxesSubplot at 0x7f999d787390>



```
[51]: #By first looking at this correlation matrix, 2 parts stand out:

# "total bsmt sf" and '1stFlrSF'

#the Garage x variables

#there is a high correlation between these variables which may be a case of → multicollinearity

#if you think about these variables, we can see that multicollinearity is → occuring

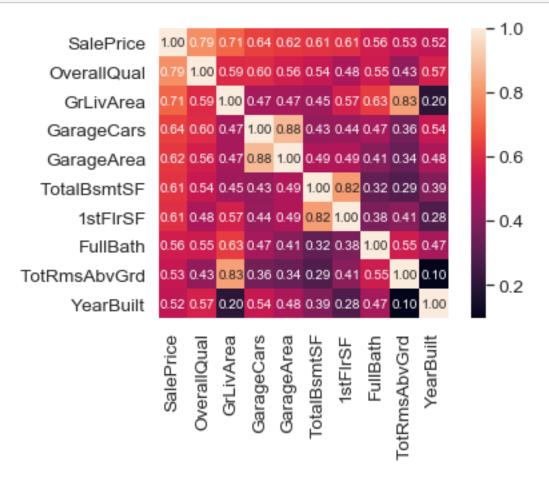
#basement size will usually be the size of the first floor and the bigger the → garage size, the more cars
```

```
#salesprice has a few squares that stand out: OverallQual, GrLivArea and Total

→BsmtSF

#lets take a look at the salesprice correlation matrix (zoomed heatmap style)

→next
```



```
[16]: sns.set()

#apply default seaborn scheme

cols =

□

□ ['SalePrice','OverallQual','GrLivArea','GarageCars','TotalBsmtSF','FullBath','YearBuilt']

#set column names into list

sns.pairplot(df_train[cols],size = 2.5)

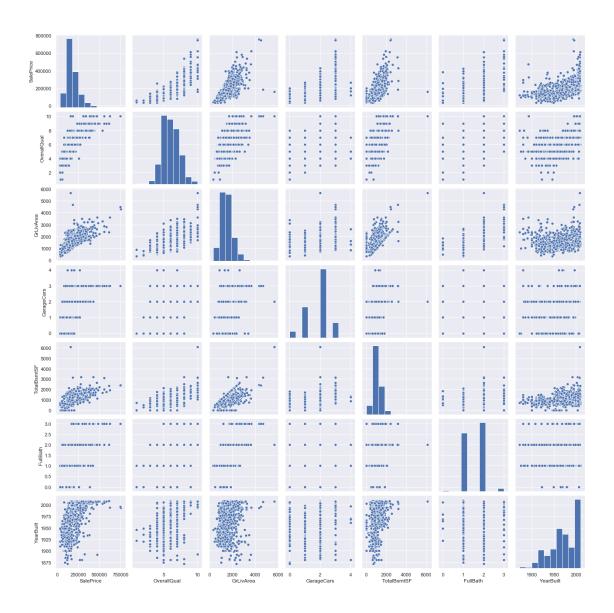
#using seaborn to compare the variables using pairplot

#pairplot plots pairwise relationships in a dataset creating a grid of plots

□ with each variable in x and y axis

plt.show()

#shows the plots
```



- [55]: #looks like Basement size = living room area #prices rise exponentially the newer the house
- [56]: #Important questions to ask about Missing Data
  #How prevalent is the missing data?
  #Is missing data random or does it have a pattern?

#missing data can imply a reduction in sample size which can prevent us from  $\underline{\hspace{0.3cm}}$  -proceeding with the analysis

```
[35]: Total = df_train.isnull().sum().sort_values(ascending = False)

#total number of null values for variable data sorted by most missing missing

percent = (df_train.isnull().sum()/df_train.isnull().count()).

-sort_values(ascending = False)

#setting a percent by dividing sum of missing values

missing_data = pd.concat([Total,percent], axis=1, keys=['Total','Percent'])

missing_data.head(20)
```

```
[35]:
                   Total
                          Percent
     PoolQC
                    1453 0.995205
     MiscFeature
                    1406 0.963014
     Allev
                    1369 0.937671
     Fence
                    1179 0.807534
     FireplaceQu
                     690 0.472603
     LotFrontage
                     259 0.177397
     GarageCond
                      81 0.055479
     GarageType
                      81 0.055479
     GarageYrBlt
                      81 0.055479
     GarageFinish
                      81 0.055479
     GarageQual
                      81 0.055479
     BsmtExposure
                      38 0.026027
     BsmtFinType2
                      38 0.026027
     BsmtFinType1
                      37 0.025342
     BsmtCond
                      37 0.025342
                      37 0.025342
     BsmtQual
     MasVnrArea
                      8 0.005479
                       8 0.005479
     MasVnrType
     Electrical
                       1 0.000685
     Utilities
                       0.000000
```

```
[18]: #the user that is guiding this project feels that variables missing more than

→>15% of data should be deleted

#i will do that in the next cell

#PoolQC, MiscFeature, Alley will be deleted as they are >15% and these arent

→ feature people look for when buying a home

#GarageX variables are missing the same number of variables. Garage cars will

→represent this data

#Same logic for BasementX variables

#For electricalm
```

```
[40]: df_train = df_train.drop((missing_data[missing_data['Total'] > 1]).index,1)

#setting df_train now as a new data frame by dropping the variables where

→missing values are > 1

df_train = df_train.drop(df_train.loc[df_train['Electrical'].isnull()].index)
```

```
#setting df train now as a new data frame by dropping the null variable in the \Box
       \rightarrow electricity columns
[41]: print(len(df train.columns))
      #checking to see if columns were dropped
     63
 []: #lets go on to do analysis on the outliers in our data
      #univariate analysis
      #we need to establish a threshold that defines an observation as an outlier.
      #we will standardize the data - converting data values to have a mean of 0 and \square
       \hookrightarrow standard deviation of 1
[43]: saleprice_scaled = StandardScaler().fit_transform(df_train['SalePrice'][:,np.
       →newaxisl)
      #using the standardscaler for the fit transform function to scale the training_
       \rightarrow data and the scaling parameters
      #the model will learn the mean and variance of the features of the training data
      #https://stackoverflow.com/questions/29241056/
      \rightarrow how-does-numpy-newaxis-work-and-when-to-use-it
      low_range = saleprice_scaled[saleprice_scaled[:,0].argsort()][:10]
      #setting low range of distribution by taking the scaled saleprice
      high_range = saleprice_scaled[saleprice_scaled[:,0].argsort()][-10:]
      #setting low range of distribution by taking the scaled saleprice
      print('outer range (low) of the distribution:')
      print(low_range)
      print('\nouter range (high) of the distribution:')
      print(high range)
     outer range (low) of the distribution:
     [[-1.83820775]
      [-1.83303414]
      [-1.80044422]
      [-1.78282123]
      [-1.77400974]
      [-1.62295562]
      [-1.6166617]
      [-1.58519209]
      [-1.58519209]
      [-1.57269236]]
     outer range (high) of the distribution:
     [[3.82758058]
      [4.0395221]
```

[4.49473628]

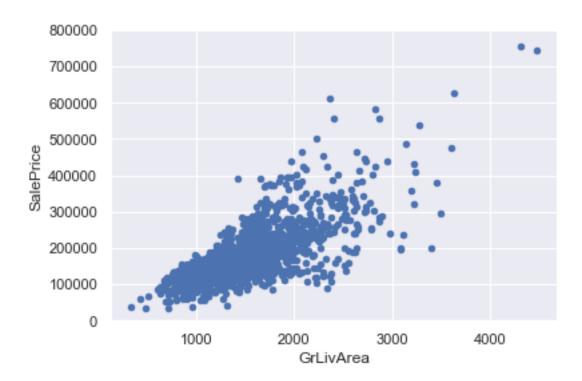
```
[4.728631]
[5.06034585]
[5.42191907]
[5.58987866]
[7.10041987]
[7.22629831]]
```

[4.70872962]

```
[46]: var = 'GrLivArea'
data = pd.concat([df_train['SalePrice'],df_train[var]],axis=1)
data.plot.scatter(x=var,y='SalePrice', ylim=(0,800000))
```

'c' argument looks like a single numeric RGB or RGBA sequence, which should be avoided as value-mapping will have precedence in case its length matches with 'x' & 'y'. Please use a 2-D array with a single row if you really want to specify the same RGB or RGBA value for all points.

[46]: <matplotlib.axes. subplots.AxesSubplot at 0x7f999bdc7850>



```
[]: #the two at the bottom right do not seem to be following the trend #we will delete those outliers
```

```
[45]: df_train.sort_values(by = 'GrLivArea', ascending = False)[:2]

#sorting df_train by GrLiveArea and keeping the first 2

df_train = df_train.drop(df_train[df_train['Id'] == 1299].index)

df_train = df_train.drop(df_train[df_train['Id'] == 524].index)

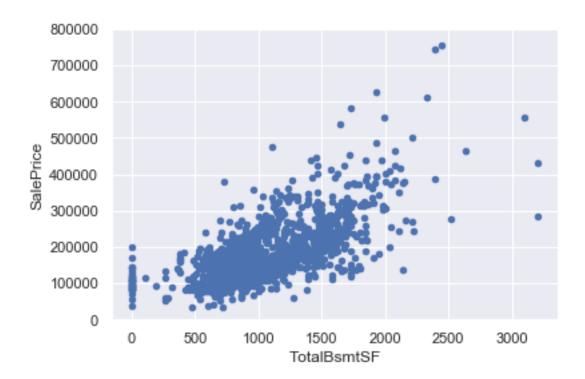
#dropping the rows 1299 and 524
```

[]: #bivariate analysis of salesprice vs totalbsmntsf

```
[47]: var = 'TotalBsmtSF'
data = pd.concat([df_train['SalePrice'],df_train[var]],axis=1)
data.plot.scatter(x=var,y='SalePrice', ylim=(0,800000))
```

'c' argument looks like a single numeric RGB or RGBA sequence, which should be avoided as value-mapping will have precedence in case its length matches with 'x' & 'y'. Please use a 2-D array with a single row if you really want to specify the same RGB or RGBA value for all points.

[47]: <matplotlib.axes.\_subplots.AxesSubplot at 0x7f99a0dff0d0>



```
[]: #looks ok, no need to delete any data

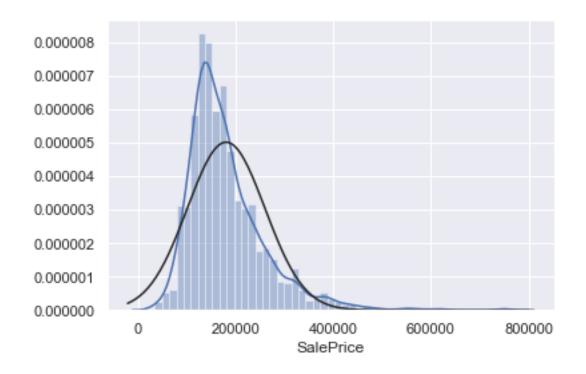
#lets look for normality and homoscedascity

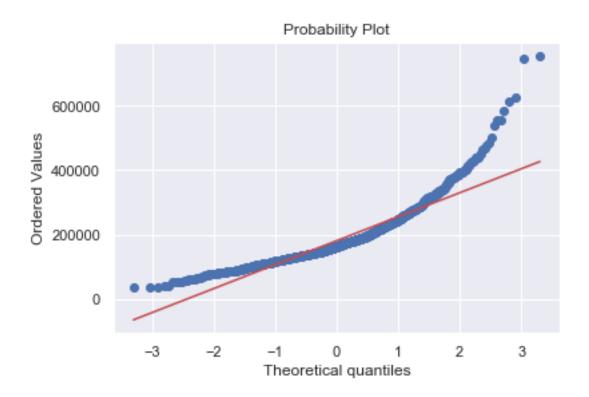
#histogram to see kurtosis and skewness

#normal probability plot - to see if data distribution follows the diagnal that

→represents normal distribution
```

```
[50]: sns.distplot(df_train['SalePrice'],fit=norm)
  #using seaborn to plot histogram by normal distribution
  fig = plt.figure()
  #create figure object
  res = stats.probplot(df_train['SalePrice'],plot = plt)
  #use scipy to create prob plot of saleprice
```



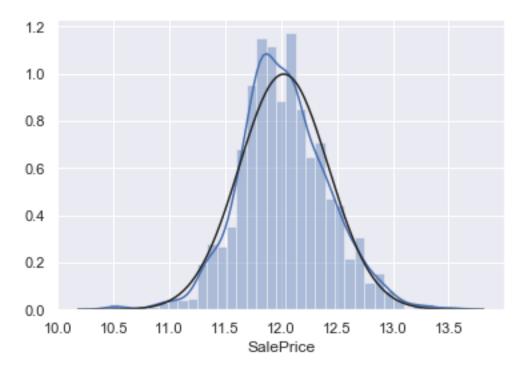


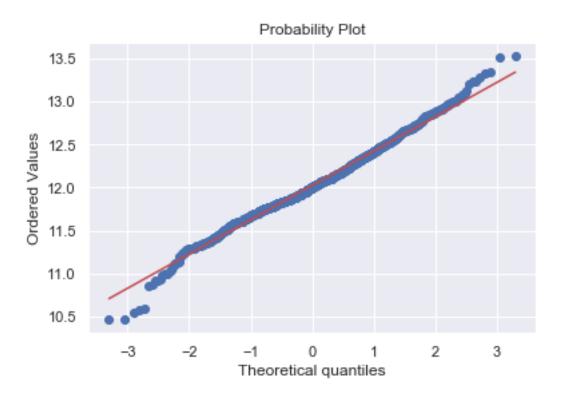
[]: #sale price is not normal distributed
#it shows peakedness, positive skewness does not follow the diagnal line
#lets tranform this data using log transformations

[51]: df\_train['SalePrice'] = np.log(df\_train['SalePrice'])

[52]: #re-running histogram and probability plot

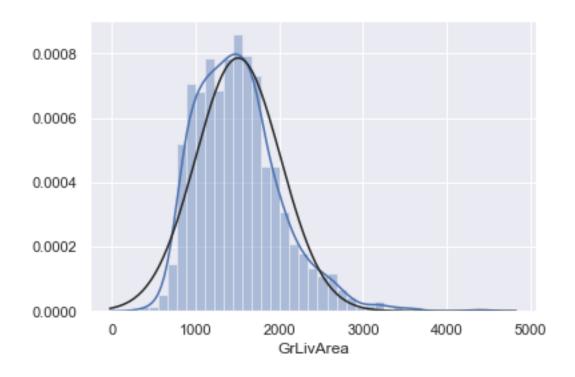
sns.distplot(df\_train['SalePrice'],fit=norm)
#using seaborn to plot histogram by normal distribution
fig = plt.figure()
#create figure object
res = stats.probplot(df\_train['SalePrice'],plot = plt)
#use scipy to create prob plot of saleprice

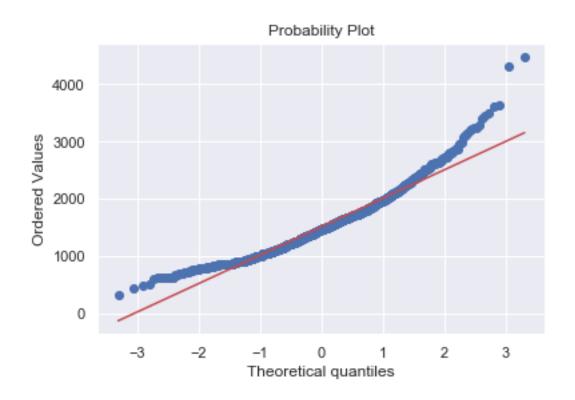




```
[ ]: #looks good, now lets check out GrLivArea

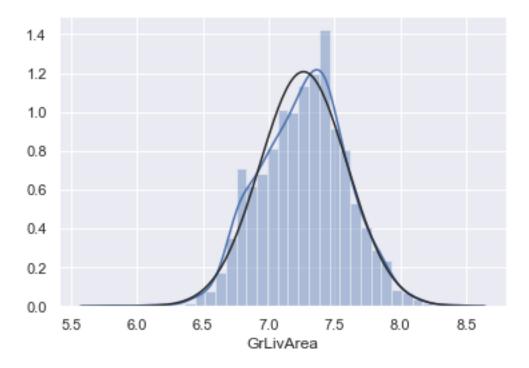
[53]: sns.distplot(df_train['GrLivArea'],fit=norm)
    #using seaborn to plot histogram by normal distribution
    fig = plt.figure()
    #create figure object
    res = stats.probplot(df_train['GrLivArea'],plot = plt)
    #use scipy to create prob plot of saleprice
```

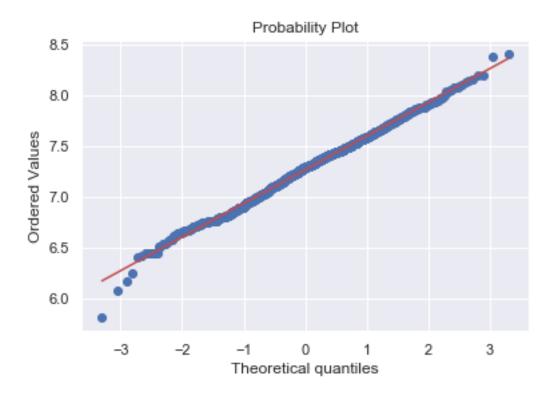




```
[54]: #there is skewness, lets transform this data
df_train['GrLivArea'] = np.log(df_train['GrLivArea'])

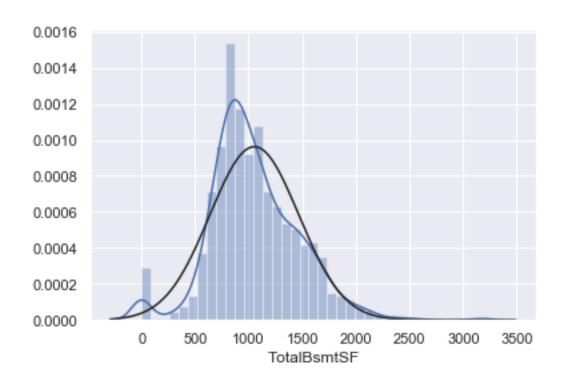
sns.distplot(df_train['GrLivArea'],fit=norm)
#using seaborn to plot histogram by normal distribution
fig = plt.figure()
#create figure object
res = stats.probplot(df_train['GrLivArea'],plot = plt)
#use scipy to create prob plot of saleprice
```

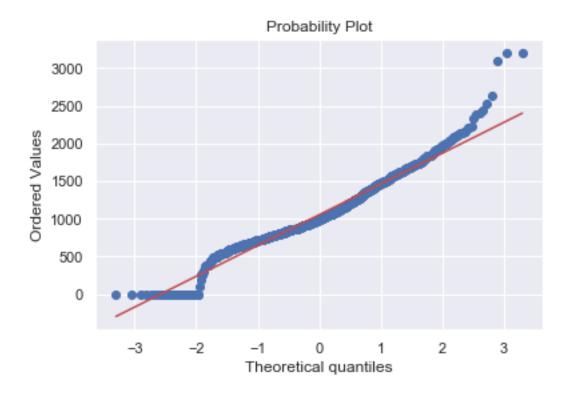




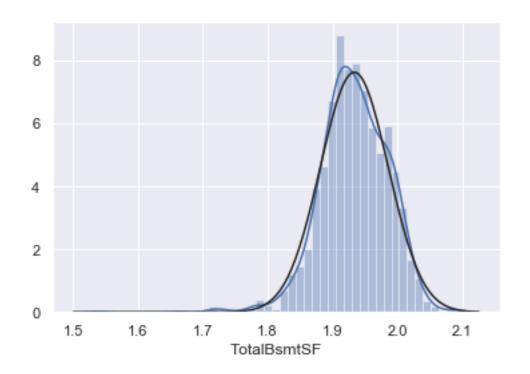
```
[55]: #much better! lets try TotalBsmtSF

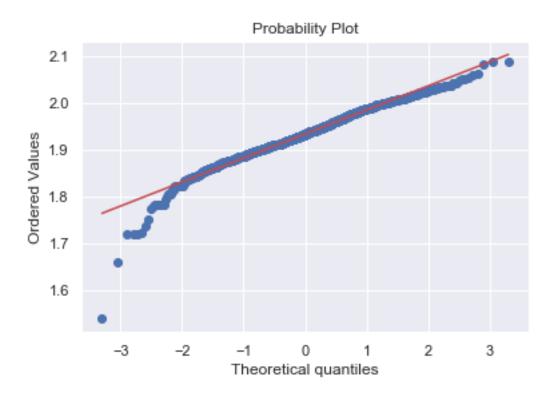
sns.distplot(df_train['TotalBsmtSF'],fit=norm)
    #using seaborn to plot histogram by normal distribution
fig = plt.figure()
    #create figure object
res = stats.probplot(df_train['TotalBsmtSF'],plot = plt)
    #use scipy to create prob plot of salepric
```





```
[]: #there is skewnwess
      #lots of houses with no basements (value = 0)
      #we cannot do log transformation on zero values
      #to conduct a log transformation, we will create a variable that can create the \Box
      →effect of hacing or not having
      #a basement. then we will do a log transformation to all the non-zero \Box
      ⇔observations and then ignore those with
      #value 0
      #a log transform of data set containing zero values can be easily handled by \Box
       →numpy.log1p()
[57]: #create column for new variable (one is enough because it's a binary
      → categorical feature)
      #if area>0 it gets 1, for area==0 it gets 0
      df_train['HasBsmt'] = pd.Series(len(df_train['TotalBsmtSF']), index=df_train.
      ⇒index)
      #create new column "HasBsmt"
      df_train['HasBsmt'] = 0
      #set all values to O
      df_train.loc[df_train['TotalBsmtSF']>0, 'HasBsmt'] = 1
      #qoes thrrough each index of TotalBsmtSF and sets the value for HasBsmt to 1 if_{\sqcup}
       \rightarrow value > 0
[58]: #transform the data
      df_train.loc[df_train['HasBsmt']==1,'TotalBsmtSF'] = np.
       →log(df_train['TotalBsmtSF'])
      #find indexes where there HasBsmt = 1 and conduct a log transformation
[60]: sns.distplot(df_train[df_train['TotalBsmtSF']>0]['TotalBsmtSF'],fit=norm)
      #using seaborn to plot histogram by normal distribution where totalbsmtsf > 0
      fig = plt.figure()
      #create figure object
      res = stats.probplot(df_train[df_train['TotalBsmtSF']>0]['TotalBsmtSF'],plot = __
      #use scipy to create prob plot of saleprice
```

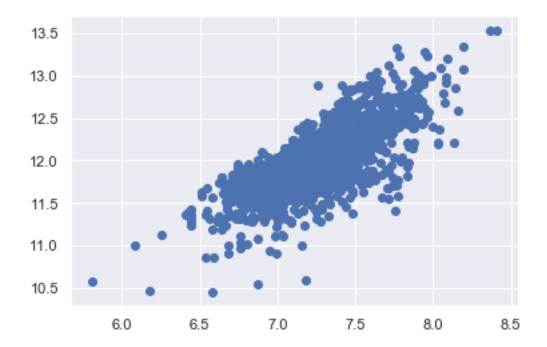




[]: #lets search for homoscedascity

[62]: plt.scatter(df\_train['GrLivArea'],df\_train['SalePrice'])

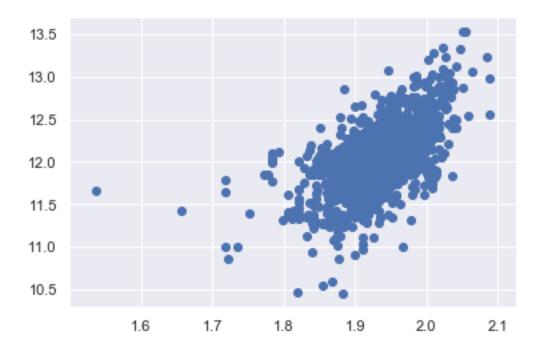
[62]: <matplotlib.collections.PathCollection at 0x7f999bb1b610>



[]: #older versions of this plot had a conic shape - it is normalized #now lets check saleprice with totalbsmtsf

[63]: plt.scatter(df\_train['TotalBsmtSF'],df\_train['SalePrice'])

[63]: <matplotlib.collections.PathCollection at 0x7f99a0689710>



```
[]: #equal level of variance of salesprice across range of totalbsmtsf

#now dummy variables
```

[64]: df\_train = pd.get\_dummies(df\_train)

#### []: #comments

[]: "Understanding the problem" phase is great, "developing sixth sense" is a\_ ⇒practice that is often disregarded. Guessing which columns to drop, in case of correlating features ideally should⊔ ⇒be performed based on some experiments, like their effect on the model accuracy or something of that ilk. It's always better to retain the data than to dismiss it. The threshold of 15% →for dropping appears to me somewhat arbitrary. As a rule of thumb, statisticians dismiss some feature if  $_{\sqcup}$ →it has more than 80 percent missing data and doesn't seem to be significantly important, because at that →point the bias from imputation is very likely to be more problematic than dropping the whole feature. After →that, ideally, it needs a little bit of experimentation to decide which features to impute and which features to  $_{\sqcup}$ →drop. "Best statisticians make the fewest assumptions".

different relationships on higher dimensions.