# Predicting The Office episode ratings to advise NBC Universal on a reunion episode

### Machine Learning in Python - Group Project 1

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In this report, we explore what makes a great episode of The Office, NBC Universal's widely-regarded situational comedy about the office life of a paper manufacturer.

This is achieved through a machine learning model which predicts the rating of the episode based on a number of episode characteristics - such as whether the episode features a cold open, and how many spoken lines are in it.

We use this model to give a recomendation on episode features which will **maximise IMBD rating** for the episode.

The report is formatted in the style of an interactive workbook in which the client (NBC Universal) can step through cells of code interspersed with explanation of model choices and mathematical background.

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## 1 Introduction

In recent years it has become popular for shows to have reunion episodes aired potentially boosting the market for online streaming of the older episodes. It is important to get these episodes right as failure to do so may taint viewers otherwise good memories of their favourite shows. In this report we aim to use machine learning methods to give informative recommendations on what a reunion episode of the TV show "The US Office" should include to achieve the highest possible IMDb rating. We had an initial data set with a variety of features such as number of lines and main characters. It also had the IMDb rating of each episode. The full feature space for **the\_office.csv** (Source <a href="https://pypi.org/project/schrutepy/">https://pypi.org/project/schrutepy/</a>) contains the following variables:

- season The season that the episode appears in
- episode The number of the episode within the season

- episode\_name The name of the episode
- director The director(s) of the episode
- writer The writer(s) of the episode
- imdb\_rating The IMDb rating of the episode
- total\_votes- The number of ratings of the episode on IMDb
- air\_date The date the episode originally aired on
- n\_lines The number of lines spoken
- n\_directions The number of stage directions
- n\_words The number of words spoken
- n\_speak\_char The number of characters with dialogue
- main\_chars The main characters that feature in the episode

We also used an additional three data sets one containing all of Jim's pranks which are found in many episodes and are a trademark of the show and one contained data for which episodes had cold opens. Cold opens are when there is a short scene before the opening title sequence. Additionally we included data for which episodes were key in the development of Jim and Pam's relationship as this is a central plotline to the show. We also did some feature engineering to create the features 'cs' for calendar season and 'month' for the month of release of the episode.

We fitted the data with a number of regression models and investigated the impact the features had on IMDb ratings. We tried linear regression Lasso and Ridge methods and we found that a Ridge regression model gave us the greatest predictive capabilities with Lasso giving similar results and the linear regression gave little predictive power.

# 1.1 Model and conclusions overview/precis

After comparing a number of models, we found a ridge regression model was the best to fit to the data. It had an  $R^2$  value of 0.4046. This means that the features that we included account for approximately 40% of the variability of the IMDB rating that each episode received. Due to the large number of possible parameters that may or may not be measurable it is difficult to build a perfect predictive model of a TV episode without overfitting. A TV episode contains hundreds of design decisions in each frame, and all the data on these is simply not available.

With that in mind, we make an important first conclusion: the average episode of the office achieved a rating of over 8.25, which is extremely high. In general, NBC already had a successful team and formula for producing good episodes, and this should be replicated as much as possible.

However, our model did indicate that the following varibles correlated with IMDb rating in order of magnatude: 'writer', 'director', 'main\_chars', 'n\_words', 'pam\_and\_jim\_key\_episode', 'cs', 'n\_directions', 'n\_speak\_chars', 'cold\_open', 'jim\_pranks'. From our analysis we make the following recomendations:

#### To include in the episode

- The director Greg Daniels
- The writer Paul Lieberstein
- The characters Michael Scott, Jim, Kelly, Creed and Stanley
- Have the script contain a large number of words which may in fact mean that we have a longer episode
- Make the episode a key episode in Jim and Pam's relationship (perhaps a pregnancy)
- Release the episode in spring
- Have a large number of directions
- Have a large number of speaking characters
- Have the episode start with a cold open

• Include some pranks from Jim

#### To avoid in episode

- Do not have Paul Lieberstein as a director
- Do not have Charlie Grandy as a writer
- Do not have the character Erin feature much or at all in the episode
- Do not release the episode in autumn

Additionally we note that the episode should stick as closely as possible to the formula that has given the US Office consistently high ratings over its 9 seasons. Further we recommend that the chosen directors and writers should be given a degree of free reign to create an episode consistent with their own style and taste. Throughout this report we detail the methods used and decisions we made in order to fit our model to the data and from this model make informed recommendations for a reunion episode for the office.

### 1.2 Data sources

Our primary data source was the office episode ratings dataset the office.csv (Evkaya, 2023).

We also explored the binary variable of whether or not the episode featured a cold-open, where the setting and premise of the episode is introduced in media res at the beginning of the episode, before the credits. This data was web-scraped from Dunderpedia: The Office Encyclopedia (Various authors, 2023a) and included in our dataset after an extensive cleaning process.

Furthermore, we explored two running themes of the episodes themselves: these are how many times Jim plays a prank in an episode, and whether or not the episode is contains key moments for the development of one of the series' key romantic relationships between Jim and Pam. The first of these was also sourced by web-scrapoing Dunderpedia: The Office Encyclopedia and subjecting the data to an extensive cleaning process (Various authors, 2023b), whilst the second was scraped and cleaned from a CinemaBlend article (West, 2020).

In total, this gave us four data sources:

- Initial dataset
- Dunderpedia article on cold opens
- Dunderpedia article on Jim's pranks
- CinemaBlend article on Jim and Pam's relationship.

## 1.3 General Setup

The following cell makes necessary package imports and sets some parameters for plotting.

```
In [1]: # Add any additional libraries or submodules below

# Data Libraries
import numpy as np
import pandas as pd

# Plotting Libraries
import matplotlib.pyplot as plt
import seaborn as sns

# Plotting defaults
```

```
plt.rcParams['figure.figsize'] = (8,5)
plt.rcParams['figure.dpi'] = 80

# sklearn modules that are necessary
import sklearn
import warnings
warnings.filterwarnings('ignore')
```

The next four cells read and join the data from our four sources. We start with the\_office.csv as as the base, and treat other data as additional, cleaning, transforming and appending it.

```
In [2]:
        # Read CSV file data
         data = pd.read csv("the office.csv")
         # Since total votes is a dependent variable we immediately drop it
         data = data.drop('total_votes', axis=1)
         data_explore = data.copy()
         with open('MLP_cold_opens.txt', 'r') as file:
In [3]:
             new_data = file.read().replace('\n', '')
         # Splitting the data up so we can see which episode has a cold open
         x = new_data.split(")")
         hots = []
         for i in range(len(x)):
             if x[i][0:4] == 'None':
                 hots.append(x[i][7:-1])
             elif x[i][0:8] == 'Season 1' and <math>x[i][8:12] == 'None':
                 hots.append(x[i][14:])
             elif x[i][0:6] == 'Season' and <math>x[i][8:12] == 'None':
                 hots.append(x[i][15:-1])
         hots_df = pd.DataFrame (hots, columns = ['episode_name'])
         hots df['cold open'] = np.zeros(12).astype(int)
         # Renaming so it is consistent with our existing data
         hots_df = hots_df.replace(to_replace = 'Weight Loss', value = 'Weight Loss (Parts 1&2)')
         data_ = pd.merge(data, hots_df, on='episode_name', how='left')
         data_['cold_open'] = data_['cold_open'].fillna(1)
         data .loc[data ['cold open'] == 1]
         data = data
In [4]:
         # Reading text file with cold opens data
         with open('MLP jim pranks.txt', 'r') as file:
             pranks str = file.read().replace('\n', '')
         # Data cleaning and pre-processing
         x0 = pranks_str.split('(')
         i = 0
         for i in x0:
             x0[j] = i.split(')')
             j+=1
         j = 0
```

for i in x0:

```
x0[j] = i[0]
             i +=1
         for i in range(len(x0)):
             x0[i] = x0[i][1:-1]
         names = data['episode_name']
         names = names.replace(to replace = 'E-Mail Surveilance', value = 'Email Surveillance')
         names = names.replace(to replace = 'A Benihana Christmas', value = 'A Benihana Christmas')
         names = names.replace(to_replace = 'The Job', value = 'The Job', regex = True)
         names = names.replace(to_replace = 'Launch Party', value = 'Launch Party', regex = True)
         names = names.replace(to replace = 'Goodbye, Toby ', value = 'Goodbye Toby', regex = True)
         names = names.replace(to_replace = 'Niagara', value = 'Niagara', regex = True)
         names = names.replace(to_replace = 'The Cover', value = 'The Cover-Up', regex = True)
         names = names.replace(to_replace = 'Classy Christmas', value = 'Classy Christmas', regex = True)
         names = names.replace(to replace = 'Dwight K. Schrute, (Acting) Manager', value = 'Dwight K. Schrute,')
         names = np.array(names)
         i = 0
         for i in names:
             if i[-11:] == '(Parts 1&2)':
                 names[j] = i[:-11]
             elif i[-8:] == '(Part 1)' or i[-8:] == '(Part 2)':
                 names[i] = i[:-8]
             j +=1
         pranks = []
         for i in names:
             #print(x0.count(i))
             pranks.append(x0.count(i))
         # a += x0.count(i)
         data['jim_pranks'] = pranks
         data
         data_explore = data.copy()
        # Reading text file with Jim and Pam data
In [5]:
         with open('jim_and_pam_key_episodes.txt', 'r') as file:
             text = file.read()
         start = 0
         pam_jim_episodes = []
         while True:
             start = text.find('(', start) + 1
             if start == 0:
                 break
             end = text.find(')', start)
             between_brackets = text[start:end]
             if any(char.isdigit() for char in between brackets):
```

```
season = int(between_brackets[1])
after_E_string = between_brackets[between_brackets.find('E')+1:]

if '/' in after_E_string:
    pam_jim_episodes.append([season, int(after_E_string[:after_E_string.find('/')]])
    pam_jim_episodes.append([season, int(after_E_string[after_E_string.find('/')+1:])])

else:
    pam_jim_episodes.append([season, int(after_E_string]])

start = end + 1

def is_jim_and_pam_key_episode(row, key_episodes):
    season, episode = row['season'], row['episode']
    return int([season, episode] in key_episodes)

data_explore['pam_and_jim_key_episode'] = data_explore.apply(
    lambda row: is_jim_and_pam_key_episode(row, pam_jim_episodes), axis=1
)
```

Since the varibales 'season' and 'episode' are out of our control we drop these from our dataset

```
In [6]: # We drop the variables season and episode
    data = data.drop('season', axis=1)
    data = data.drop('episode', axis=1)

    data_explore = data_explore.drop('season', axis=1)
    data_explore = data_explore.drop('episode', axis=1)
```

# 2. Exploratory Data Analysis and Feature Engineering

# 2.1 Correlation exploration

In the cell below, we show the data frame created from our four sources without any modification, alongside summary statistics. Additionally, we check for NA and duplicated values in the data, finding none - an important step in veryifying the data we collected.

Out[7

There are 0 duplicated values in the data There are 0 Na values in the data

7]:		imdb_rating	n_lines	$n\_directions$	n_words	n_speak_char	cold_open	jim_pranks	pam_and_jim_key_episode
	count	186.000000	186.000000	186.000000	186.000000	186.000000	186.000000	186.000000	186.000000
	mean	8.250538	296.397849	50.150538	3053.510753	20.693548	0.935484	0.521505	0.231183
	std	0.535168	81.998568	23.941797	799.271717	5.092407	0.246333	0.901692	0.422727
	min	6.700000	131.000000	11.000000	1098.000000	12.000000	0.000000	0.000000	0.000000
	25%	7.900000	255.250000	34.000000	2670.250000	17.000000	1.000000	0.000000	0.000000
	50%	8.200000	281.000000	46.000000	2872.500000	20.000000	1.000000	0.000000	0.000000
	75%	8.600000	314.500000	60.000000	3141.000000	23.000000	1.000000	1.000000	0.000000
	max	9.700000	625.000000	166.000000	6076.000000	54.000000	1.000000	8.000000	1.000000

Our summary statistics show us a few important summary points:

- There are 186 episodes in the dataset.
- The mean IMDB rating across the series was 8.25.
- The average number of Jim pranks per episode is 0.5, with a maximum of 8 in one.

In the cell below, we create a correlation heatmap between the current columns of a dataset, to give some understanding of the data. We are looking for variables with a high correlation with each other (as they may not both add a lot of value to the model) and the correlation of variables with IMDB ratings.

```
In [8]: # Plotting heatmap
sns.set(rc={'figure.figsize': (10, 6)})
sns.heatmap(data_explore.corr(), annot = True, fmt = '.2f', linewidths = 2)
plt.title("Correlation Heatmap for current numerical variables")
plt.tight_layout()
plt.show()
```



We see that the number of lines and number of words features are highly correlated so we will chose to drop 'n\_lines' as including both would be redundant.

```
In [9]: data = data.drop('n_lines', axis=1)
    data_explore = data_explore.drop('n_lines', axis=1)
```

# 2.2 Feature engineering and justification

This section contains the feature engineering for the model, including columns added and dropped and the justifications as to why.

We will first drop the episode name and episode number features because they are not relevant to the model.

Out[12]:	director	writer	imdb_rating	n_directions	n_words	n_speak_char	main_chars	cold_open	jim_pranks	pam_and_jim_key_episode	calendar_season	mc
	<b>0</b> Ken Kwapis	Ricky Gervais;Stephen Merchant;Greg Daniels	7.6	27	2757	15	Angela;Dwight;Jim;Kevin;Michael;Oscar;Pam;Phyl	0.0	3	1	winter	
	1 Ken Kwapis	B.J. Novak	8.3	20	2808	12	Angela; Dwight; Jim; Kelly; Kevin; Michael; Oscar; Pa	0.0	0	0	winter	
	2 Ken Whittingham	Paul Lieberstein	7.9	21	2769	13	$\label{thm:meredith:meredith:Michael;Oscar} Angela; Dwight; Jim; Kevin; Meredith; Michael; Oscar$	0.0	2	0	spring	
	Bryan Gordon	Michael Schur	8.1	24	2939	14	Angela; Dwight; Jim; Kevin; Meredith; Michael; Oscar	0.0	1	0	spring	
	4 Greg Daniels	Greg Daniels	8.4	49	2437	18	Angela; Darryl; Dwight; Jim; Kevin; Michael; Oscar; P	0.0	0	0	spring	

Our current data contains columns such as main\_chars which have a number of non-numerical and non-continuous variables. We need to split these into categorical variables. For example, if an episode had Michael in it, we'd want to be able to quickly access a column called mc\_michael and if it was = 1, we'd know he was in the episode.

The following code defines and deploys a one hot encoder that assigns a binary attribute per category for the categorical variables ("director, "writer", "calendar\_season" and "main\_chars"):

```
In [13]: def OneHot_encoder(df, cat_variable):
    oneHot_df = df.copy()
    variable_initials = ''.join([word[0] for word in cat_variable.split("_")])

unique_chars = set(';'.join(oneHot_df[cat_variable]).split(';'))

# Create binary attributes for each unique character
for char in unique_chars:
    oneHot_df[variable_initials+"_"+char] = oneHot_df[cat_variable].apply(lambda x: 1 if char in x else 0)

# Drop the original main_chars column
oneHot_df = oneHot_df.drop(columns=[cat_variable])
return oneHot df
```

```
cat_variables = ['director', 'writer', 'main_chars', 'calendar_season']

for cat_variable in cat_variables:
    data_explore = OneHot_encoder(data_explore, cat_variable)

data_explore.head(5)
```

Out[13]:	imdb_rating	n_directions	n_words	n_speak_char	cold_open	jim_pranks	pam_and_jim_key_episode	month	d_Rainn Wilson	d_Joss Whedon	mc_Eri	n mc_Pam	mc_Andy	mc_Stanley	mc_Oscar	mc_Toby	c
O	7.6	27	2757	15	0.0	3	1	3	0	0		) 1	0	1	1	0	_
1	8.3	20	2808	12	0.0	0	0	3	0	0		) 1	0	1	1	1	
2	2. 7.9	21	2769	13	0.0	2	0	4	0	0		) 1	0	1	1	0	
3	8.1	24	2939	14	0.0	1	0	4	0	0		) 1	0	1	1	1	
4	8.4	49	2437	18	0.0	0	0	4	0	0		) 1	0	1	1	0	

5 rows × 129 columns

By inspection we realised that there are some repeated directors where in some data entries the name of the director is spelled wrong. The specific cases were:

- Charles McDougall (correct name) and Charles McDougal
- Claire Scanlon (correct name) and Claire Scanlong
- Greg Daniels (correct name) and Greg Daneils
- Ken Whittingham (correct name) and Ken Wittingham
- Paul Lieberstein (correct name) and Paul Lieerstein

The following code removes columns with the wrong name, where we will put a 1 in the correct name column if the incorrect one had a 1 and the correct one didn't.

```
In [14]:
    correct_names = ['d_Charles McDougal','d_Claire Scanlon','d_Greg Daniels', 'd_Ken Whittingham', 'd_Paul Lieberstein']
    wrong_names = ['d_Charles McDougal','d_Claire Scanlong','d_Greg Daniels', 'd_Ken Wittingham', 'd_Paul Lieberstein']

for i, correct_name in enumerate(correct_names):
    data_explore.loc[data_explore[wrong_names[i]] == 1, correct_name] = 1
    data_explore = data_explore.drop(wrong_names[i], axis=1)

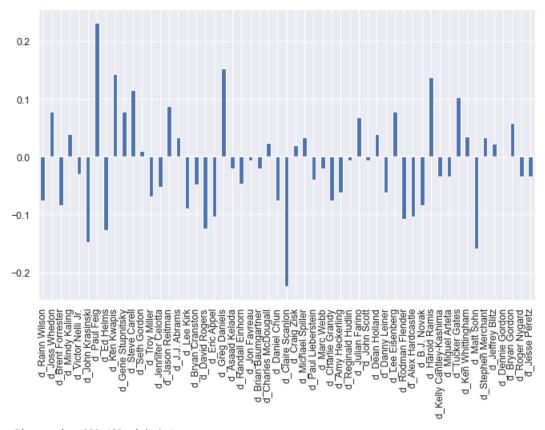
data_explore.head(5)
```

Out[14]:	imdb_rating	n_directions	n_words	n_speak_char	cold_open	jim_pranks	pam_and_jim_key_episode	month	d_Rainn Wilson	d_Joss Whedon	mc_Erin	mc_Pam	mc_Andy	mc_Stanley	mc_Oscar	mc_Toby	c
-	0 7.6	5 27	2757	15	0.0	3	1	3	0	0	0	1	0	1	1	0	_
	1 8.3	3 20	2808	12	0.0	0	0	3	0	0	0	1	0	1	1	1	
	2 7.9	21	2769	13	0.0	2	0	4	0	0	0	1	0	1	1	0	
	3 8.	1 24	2939	14	0.0	1	0	4	0	0	0	1	0	1	1	1	

	imdb_rating	n_directions	n_words	n_speak_char	cold_open	jim_pranks	pam_and_jim_key_episode	month	d_Rainn Wilson	d_Joss Whedon	mc_Erin	mc_Pam	mc_Andy	mc_Stanley	mc_Oscar	mc_Toby	c
4	8.4	49	2437	18	0.0	0	0	4	0	0	0	1	0	1	1	0	

5 rows × 124 columns

Having defined our data, we now need to explore the predictive power of each piece of data in creating a good episode with high imdb ratings. The following plot gives the correlation between each variable individually and the imdb rating.



<Figure size 800x480 with 0 Axes>

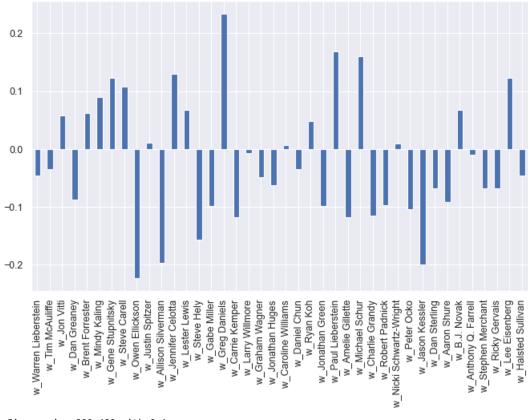
# 2.3 Exploration of potential dimensionality reduction

To reduce the dimenionsionality of our data and avoid overfitting, the following code cells look at directors, writers and main characters in the dataset with a greater than 0.1 absolute correlation value with IMDB rating.

In our model fine tuning, this was important to prevent overfitting and to reduce dimensionality, but is only presented here as EDA.

```
corr_mat_d_siginificant = corr_mat_d[abs(corr_mat_d) > 0.1]
In [16]:
          corr mat d siginificant
Out[16]: d_John Krasinski
                              -0.148062
         d_Paul Feig
                              0.232619
         d_Ed Helms
                              -0.127075
         d_Ken Kwapis
                              0.143246
         d_Steve Carell
                              0.115819
         d_David Rogers
                              -0.124627
         d_Eric Appel
                              -0.103387
         d_Greg Daniels
                              0.153229
         d_Claire Scanlon
                              -0.224743
         d_Rodman Flender
                              -0.107541
```

```
d_Alex Hardcastle
                             -0.103387
         d_Harold Ramis
                              0.138732
         d_Tucker Gates
                              0.104012
         d_Matt Sohn
                             -0.159096
         Name: imdb_rating, dtype: float64
          # Exploring correlation between IMDb and directors
In [17]:
          w_list = [columns[i] for i in range(len(columns)) if columns[i][:2] == "w_"]
          corr_mat_w = data_explore[w_list + ["imdb_rating"]].corr()["imdb_rating"]
          corr_mat_w = corr_mat_w[:-1]
          print()
          corr_mat_w.plot(kind='bar')
          plt.xticks(rotation ='vertical')
          plt.show()
          plt.tight_layout()
```



<Figure size 800x480 with 0 Axes>

```
Out[18]: w_Gene Stupnitsky
                               0.123633
         w_Steve Carell
                               0.107331
         w_Owen Ellickson
                               -0.222360
         w_Allison Silverman -0.196041
         w_Jennifer Celotta
                               0.129994
         w_Steve Hely
                               -0.156059
         w_Greg Daniels
                               0.233452
         w_Carrie Kemper
                               -0.118199
         w_Paul Lieberstein
                               0.168538
         w_Amelie Gillette
                              -0.117308
         w_Michael Schur
                               0.160538
         w_Charlie Grandy
                              -0.114410
         w_Peter Ocko
                              -0.103387
         w_Jason Kessler
                              -0.199813
         w Lee Eisenberg
                               0.123633
         Name: imdb_rating, dtype: float64
```

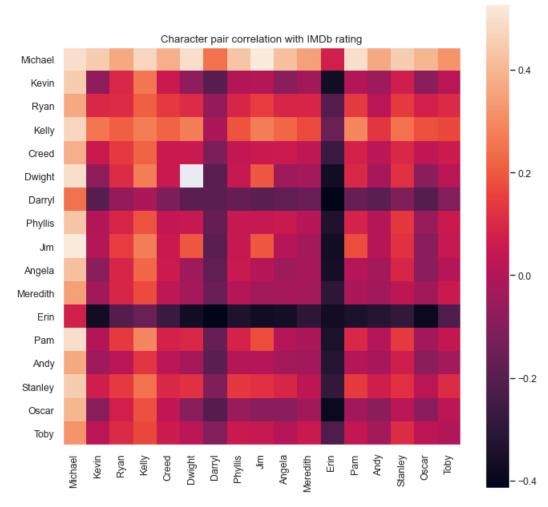
# 2.4 Paired-characters feature exploration

In this section, we explore the effect of character pairings. For example, Jim and Dwight had a number of running jokes and gags which may have affected IMDB rating of episodes that they both featured in.

This cell creates lists of main character names, and then adds main character pairings to a data exploration frame. The purpose of this analysis is to determine whether some characters are better on screen with others.

```
In [19]: | # Exploring the correlation of character pairs agains IMDb
          mc_list = [columns[i] for i in range(len(columns)) if columns[i][:3] == "mc_"]
          n = len(mc_list)
          pairnames = []
          data_explore_2 = data_explore.copy()
          for i in range(n-1):
              for j in range(i+1,n):
                  pairname = "pair_" + mc_list[i][3:] + "_"+ mc_list[j][3:]
                  pairnames.append(pairname)
                  data_explore_2[pairname] = ((data_explore_2[mc_list[i]] + data_explore_2[mc_list[j]])//2)
In [20]:
          corr_mat_p = data_explore_2.corr()["imdb_rating"]
          corr array p = np.zeros((n,n))
          for i in range(n-1):
              for j in range(i+1,n):
                  pairname = "pair_" + mc_list[i][3:] + "_"+ mc_list[j][3:]
                  corr_array_p[i][j] = corr_mat_p[pairname]
                  corr_array_p[j][i] = corr_mat_p[pairname]
              corr_array_p[i][i] = corr_mat_p[mc_list[i]]
          fig, axes = plt.subplots(figsize = (10,10), ncols = 1)
In [21]:
          mc_names = [mc_name[3:] for mc_name in mc_list]
          plt.grid(b=False)
          plt.imshow(corr array p)
          plt.colorbar()
```

```
plt.xticks(range(n), mc_names, rotation ='vertical')
plt.yticks(range(n), mc_names)
plt.title('Character pair correlation with IMDb rating')
```



We note the positive impact of Michael and Kelly with all other characters and the negative impact of Erin and Darryl. Dwight's correlation with himself is so high because he is in every episode: this is an important observation as we will need to drop Dwight from the final model, his presence in an epsiode cannot be compared with episodes he isn't in so we shouldn't include it in the model.

# 3. Model Fitting and Tuning

In this section, we fit our model to the data and fine-tune it based on the EDA conducted in section 2. We tested a number of models and eventually settled on on a ridge regression model, preliminary results for our other models are noted below.

The following cell imports necessary SciKitLearn packages and sets up get\_coefs functions.

```
In [22]: from sklearn.linear model import LinearRegression
          from sklearn.pipeline import Pipeline
          from sklearn.model_selection import train_test_split
          from sklearn.preprocessing import StandardScaler
          from sklearn.metrics import mean squared error, r2 score
          from sklearn.linear_model import Ridge
          from sklearn.linear model import Lasso
          def get coefs(m):
              """Returns the model coefficients from a Scikit-learn model object as an array,
              includes the intercept if available.
              # If pipeline, use the last step as the model
              if (isinstance(m, sklearn.pipeline.Pipeline)):
                  m = m.steps[-1][1]
              if m.intercept is None:
                  return m.coef_
              return m.coef
          def model_fit(m, X, y, plot = False):
              """Returns the mean squared error, root mean squared error and R^2 value of a fitted model based
              on provided X and y values.
              Args:
                  m: sklearn model object
                  X: model matrix to use for prediction
                  y: outcome vector to use to calculating rmse and residuals
                  plot: boolean value, should fit plots be shown
              y hat = m.predict(X)
              MSE = mean squared error(y, y hat)
              RMSE = np.sqrt(mean_squared_error(y, y_hat))
              Rsqr = r2 \ score(y, y \ hat)
              Metrics = (round(MSE, 4), round(RMSE, 4), round(Rsqr, 4))
              res = pd.DataFrame(
                  data = {'y': y, 'y_hat': y_hat, 'resid': y - y_hat}
              if plot:
                  plt.figure(figsize=(11, 5.5))
                  plt.subplot(121)
                  sns.lineplot(x='y', y='y_hat', color="grey", data = pd.DataFrame(data={'y': [min(y),max(y)], 'y_hat': [min(y),max(y)]}))
                  sns.scatterplot(x='y', y='y_hat', data=res).set_title("Actual vs Fitted plot")
```

```
plt.subplot(122)
sns.scatterplot(x='y_hat', y='resid', data=res).set_title("Fitted vs Residual plot")
plt.hlines(y=0, xmin=np.min(y), xmax=np.max(y), linestyles='dashed', alpha=0.3, colors="black")

plt.subplots_adjust(left=0.0)

plt.suptitle(r"Model (MSE, RMSE, $R^2$) = " + str(Metrics), fontsize=14)
plt.show()

return MSE, RMSE, Rsqr

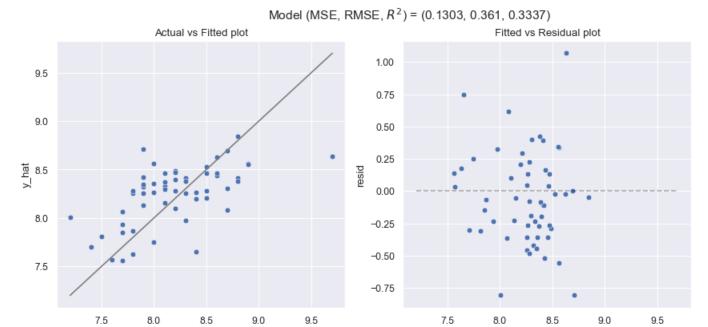
# Having done all the data exploration and feature engineering, we now set our starting dataframe to the
# engineered dataframe.
data = data_explore

X_train, X_test, y_train, y_test = train_test_split(
    data.drop('imdb_rating', axis=1), data['imdb_rating'],
    test_size=0.3, random_state=42
}
```

### Selcection of the Ridge model

From testing out Ridge, Lasso and Linear regression models we found that Ridge gave us the best performance so we opted to choose it for fine tuning.

```
from sklearn.pipeline import make pipeline
In [23]:
          from sklearn.model selection import GridSearchCV, KFold
          # Applying a gridsearch procedure to find the optimal alpha value
          alphas = np.linspace(0, 200, num=200)
          gs ridg3 = GridSearchCV(
              make pipeline(
                  StandardScaler(),
                  Ridge()
              param grid = {'ridge alpha': alphas},
              cv = KFold(5, shuffle = True, random state = 42),
              scoring = "neg root mean squared error"
          gs ridg3.fit(X train, y train)
Out[23]: GridSearchCV(cv=KFold(n_splits=5, random_state=42, shuffle=True),
                      estimator=Pipeline(steps=[('standardscaler', StandardScaler()),
                                                ('ridge', Ridge())]),
                                                                        1.00502513, 2.01005025, 3.01507538,
                      param_grid={'ridge__alpha': array([ 0.
                  4.0201005, 5.02512563, 6.03015075, 7.03517588,
                  8.04020101, 9.04522613, 10.05025126, 11.05527638,
                 12.06030151, 13.06532663, 14.07035176, 15.0753768...
                172.86432161, 173.86934673, 174.87437186, 175.87939698,
                176.88442211, 177.88944724, 178.89447236, 179.89949749,
                180.90452261, 181.90954774, 182.91457286, 183.91959799,
                184.92462312, 185.92964824, 186.93467337, 187.93969849,
                188.94472362, 189.94974874, 190.95477387, 191.95979899,
                192.96482412, 193.96984925, 194.97487437, 195.9798995,
```



# Fine Tuning the model

In this section we will drop features from our model in order to avoid overfitting. Fist of all, we will investigate how many times each director appears in the data, and we will fit a linear regression (against IMDb rating) for each and measure R^2 both for the train data and test data. We will also carry out the same process for writers and main characters.

y\_hat

```
In [26]: directors = X_train.filter(regex='^d_').columns.values
    writers = X_train.filter(regex='^w_').columns.values
    main_characters = X_train.filter(regex='^mc_').columns.values
```

У

```
rest_of_features = [var for var in X_train.columns.values
               if var not in np.concatenate((directors, writers, main characters), axis=0)]
def get_features_rsqr(features, label):
   features_frequencies = data[features].sum(axis=0)
   rsqr_train_all = []
   rsqr_test_all = []
   # Looping through each feature and fit a simple linear regression
   for col in features:
       X = np.c_[X_train[col]]
       m = LinearRegression(fit_intercept=True).fit(X, y_train)
       mse, rmse, rsqr_train = model_fit(m, X, y_train)
       mse, rmse, rsqr_test = model_fit(m, np.c_[X_test[col]], y_test)
       rsqr_train_all.append(rsqr_train)
       rsqr_test_all.append(rsqr_test)
   # Print out table
   df = pd.DataFrame({label: features, 'Frequency': features_frequencies,
                       'R^2 train': rsqr_train_all, 'R^2 test': rsqr_test_all})
   df = df.sort_values(by=['R^2 train'], ascending=False)
   return df.reset index(drop=True)
get features rsqr(directors, 'Directors').head(10)
```

Out[26]:		Directors	Frequency	R^2 train	R^2 test
	0	d_Claire Scanlon	2	0.066846	-0.044416
	1	d_Paul Feig	14	0.038677	0.079539
2	2	d_John Krasinski	3	0.030056	-0.041402
:	3	d_Harold Ramis	4	0.022449	-0.018241
4	4	d_Ed Helms	2	0.022007	-0.037281
!	5	d_Greg Daniels	15	0.019921	0.009524
(	6	d_Steve Carell	3	0.015557	-0.020942
;	7	d_Eric Appel	1	0.014403	-0.033435
:	8	d_Alex Hardcastle	1	0.014403	-0.033435
9	9	d_B.J. Novak	5	0.013737	-0.053040
In [27]:	ge	t_features_rsq	r(writers,	'Writers	').head(10)
Out[27]:		Writers	Frequency	R^2 train	R^2 test

4 0.066846 -0.089808

1 0.052296 -0.038073

**0** w\_Owen Ellickson

w\_Jason Kessler

Out[29]:

	Writers	Frequency	R^2 train	R^2 test
2	w_Paul Lieberstein	16	0.047219	-0.101873
3	w_Steve Hely	3	0.033270	-0.042130
4	w_Greg Daniels	12	0.028189	0.110828
5	w_Lee Eisenberg	15	0.024714	-0.067359
6	w_Gene Stupnitsky	15	0.024714	-0.067359
7	w_Jon Vitti	2	0.020706	-0.165538
8	w_Jennifer Celotta	11	0.018910	-0.023061
9	w_Carrie Kemper	4	0.018858	-0.044368

We can see that both directors and writers correlate very poorly with IMDb rating, since in both cases all the  $R^2$  values are very small, not exceeding 0.06. We also notice that many of them take part in very few episodes, indicating that they are most likely not relevant features. Therefore, we have decided to drop all the directors and writers that appear in 5 episodes or less.

```
In [28]: data_tuning = data.copy()

# Getting df with frequencies of each director and writer.
frequencies = data_tuning[list(writers) + list(directors)].apply(pd.Series.value_counts)
# Getting the ones we will drop
to_drop = frequencies.columns[frequencies.loc[1] <= 5]
# Dropping them.
data_tuning = data_tuning.drop(to_drop, axis=1)</pre>
In [29]: main_chars_stats = get_features_rsqr(main_characters, 'Main characters')
```

In [29]: main\_chars\_stats = get\_features\_rsqr(main\_characters, 'Main characters')
main\_chars\_stats

	Main characters	Frequency	R^2 train	R^2 test
0	mc_Michael	137	0.244101	0.233410
1	mc_Erin	98	0.140186	0.099177
2	mc_Kelly	142	0.072594	0.047324
3	mc_Jim	185	0.052296	-0.038073
4	mc_Darryl	105	0.045861	-0.028430
5	mc_Stanley	165	0.008692	0.004926
6	mc_Ryan	139	0.007197	0.002627
7	mc_Creed	131	0.006404	-0.043715
8	mc_Oscar	164	0.005232	-0.015092
9	mc_Kevin	180	0.005082	-0.021952
10	mc_Pam	182	0.005070	-0.001006
11	mc_Meredith	131	0.002727	-0.037586

	Main characters	Frequency	R^2 train	R^2 test
12	mc_Angela	171	0.002535	-0.031828
13	mc_Phyllis	165	0.001949	-0.026248
14	mc_Toby	108	0.000766	-0.041249
15	mc_Andy	143	0.000098	-0.027077
16	mc_Dwight	186	0.000000	-0.028662

In main characters on the other hand we see that Dwight appears in all the episodes, which thus leads to an  $R^2$  of 0. Therefore, we will drop Dwight, and in our recommendations we will say that he should be included. Additionally, we see that in this case we two  $R^2$  scores in train that are significant, 0.24 for Michael and 0.14 for Erin, but the other onces are once again very small. In order to decrease overfitting, we have decided to only keep the characters with the 10 biggest  $R^2$ .

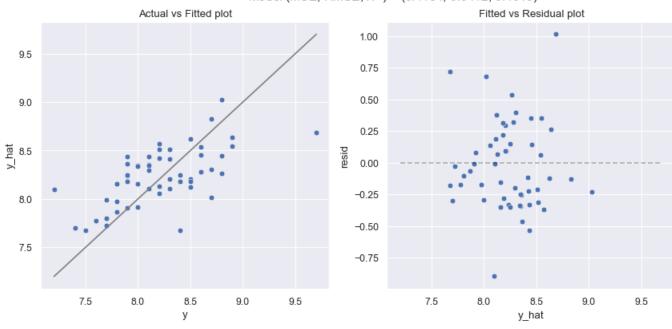
```
In [30]: # Dropping Dwigth.
    data_tuning = data_tuning.drop('mc_Dwight', axis=1)

# Getiting the the characters not in the top 10 and dropping them.
    bottom_characters = main_chars_stats["Main characters"].values[11:-1]
    data_tuning = data_tuning.drop(bottom_characters, axis=1)
```

We now fit the model again with the new set of features.

```
90.45226131, 90.95477387, 91.45728643, 91.95979899,
                 92.46231156, 92.96482412, 93.46733668, 93.96984925,
                 94.47236181, 94.97487437, 95.47738693, 95.9798995,
                 96.48241206, 96.98492462, 97.48743719, 97.98994975,
                 98.49246231, 98.99497487, 99.49748744, 100.
                                                                     ])},
                      scoring='neg_root_mean_squared_error')
In [32]:
          print(gs.best_params_)
         {'ridge__alpha': 74.87437185929647}
In [33]:
          final_model = Pipeline([
              ('scaler', StandardScaler()),
              ('model', Ridge(alpha=74.87))
          1)
          # Fitting the model
          final_model.fit(X_train, y_train)
          mse, rmse, rsqr = model_fit(final_model, X_test, y_test, plot=True)
```

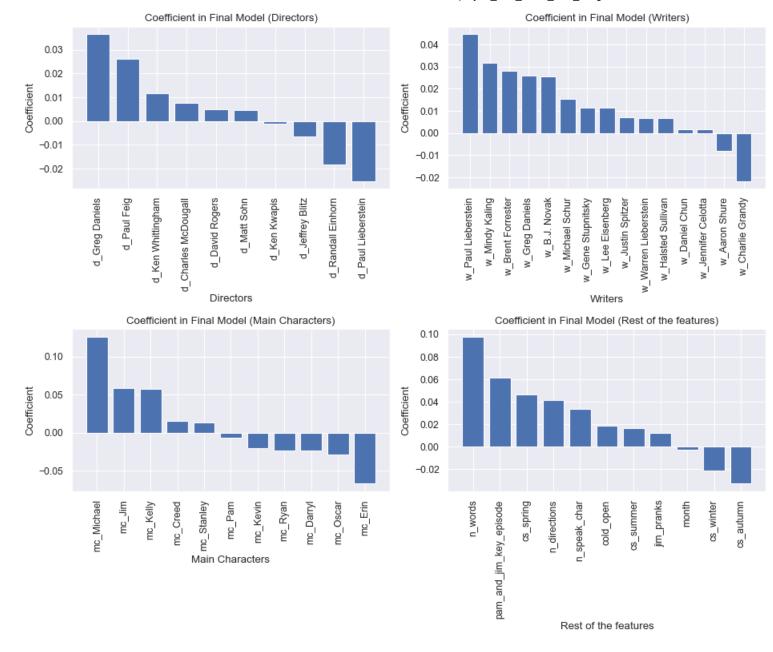
### Model (MSE, RMSE, $R^2$ ) = (0.1164, 0.3412, 0.4046)



## 4. Discussion and Conclusions

```
In [34]: # Getting the new reduced number of directors, writers and main characters
directors = X_train.filter(regex='^d_').columns.values
writers = X_train.filter(regex='^w_').columns.values
```

```
main_characters = X_train.filter(regex='^mc_').columns.values
def plot_features_coef_magnitude_for_given_model(m, features, label, ax):
    # Getting the coefficients from the model and selecting subset
    # corresponding to features.
    coefs_series = pd.Series(get_coefs(m), index=X_train.columns)
    coefs_features = coefs_series.loc[features]
    # Sorting coefficients
    sorted coef = coefs features.sort values(ascending=False)
    ax.bar(sorted_coef.index, sorted_coef.values)
    ax.set_xticklabels(sorted_coef.index, rotation=90)
    ax.set xlabel(label)
    ax.set_ylabel('Coefficient')
    ax.set_title('Coefficient in Final Model ({})'.format(label))
fig, axs = plt.subplots(2, 2, figsize=(12, 10))
plot_features_coef_magnitude_for_given_model(final_model, directors, "Directors", axs[0, 0])
plot_features_coef_magnitude_for_given_model(final_model, writers, "Writers", axs[0, 1])
plot_features_coef_magnitude_for_given_model(final_model, main_characters, "Main Characters", axs[1, 0])
plot_features_coef_magnitude_for_given_model(final_model, rest_of_features, "Rest_of_the features", axs[1, 1])
plt.tight layout()
nlt.show()
```

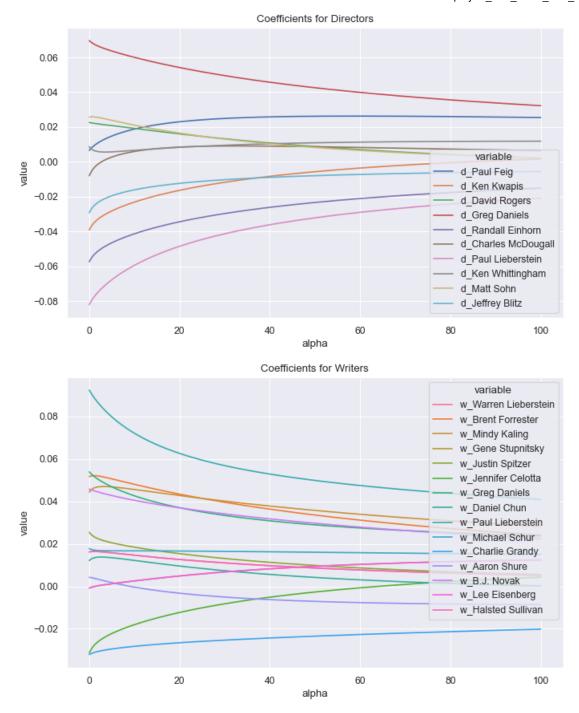


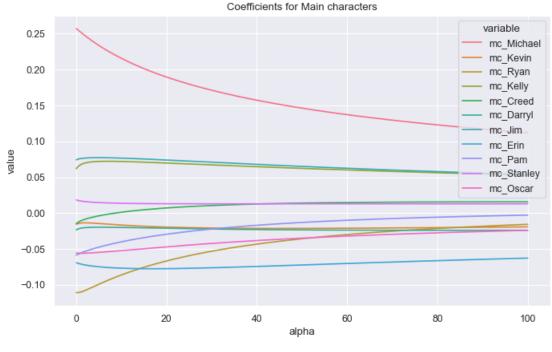
## 4.1 Model Performance

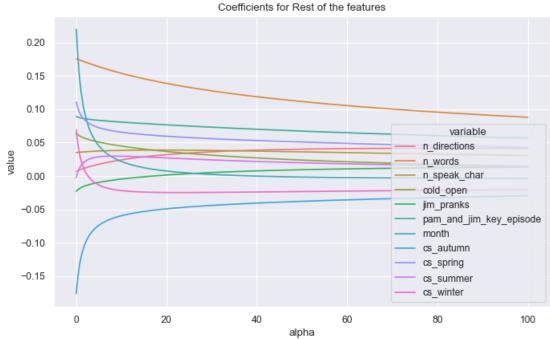
After fine tuning the model and selecting certain features we achived an  $R^2$  value of 0.4046 which suggests that we our model accounts for about 40% of the variability in IMDb rating given our features. This means that a large degree of the variability is outwith out predictive capabilities so it is important to not overestate the importence of our results. Other factors such as actors performance and plot details may affect the IMDb rating more than the features we had in our data set. That being said this  $R^2$  value is high enough for us to make meaningful suggestions to NBC Universal as to how the reunion episode should be made. The **RMSE** value for our final model was 0.3412 so on avergae our predicted IMDb ratings were this far from the true IMDb ratings that the episodes recieved. This indicated that our model predicts with a resonable level of accuracy given IMDb rating are on a scale from 0-10. Additionally we removed features such as 'total votes' and 'season' which would be outwith the control of producers meaning that the predictions we make are relevent for the production of a future episode.

Below we plot the value of the coefficients the model gives depending on various selctions of  $\alpha$ . In ection 3 we used a gridsearch method to find the optimal  $\alpha$  value for our model. The  $\alpha$  parameter is intended to punish models with large amounts of predictive features. These plots are included to shed some light on how this affects the model.

```
def coefs_vs_alpha(features, model_type, title):
    alphas = np.logspace(-2, 2, num=200) # from 10^-2 to 10^2
    # Arrays to store coefficients and rmses
    betas = []
    rmses = []
    for a in alphas:
        m = make pipeline(
            StandardScaler().
            model type(alpha=a)
        ).fit(X train, y train)
        # Getting the coefficients for the specified features, and
        # getting rmses.
        coefs series = pd.Series(get coefs(m), index=X train.columns)
        variables_coefs = coefs_series.loc[features]
        betas.append(variables_coefs)
        rmses.append(model fit(m, X test, y test))
    # Createing DataFrame of coefficients and RMSEs
    res = pd.DataFrame(
        data=betas,
        columns=features,
    ).assign(
        alpha = alphas,
        rmse = rmses
    ).melt(
        id vars = ('alpha', 'rmse')
    sns.lineplot(x='alpha', y='value', hue='variable', data=res).set title("Coefficients for "+title)
    plt.show()
coefs vs alpha(directors, Ridge, "Directors")
coefs vs alpha(writers, Ridge, "Writers")
coefs_vs_alpha(main_characters, Ridge, "Main characters")
coefs_vs_alpha(rest_of_features, Ridge, "Rest of the features")
```







## 4.2 Recomendations

From the plots above we can see that certain directors and writers gave significant improvements in the predicted IMDb rating of the episodes. Additionally certain characters and other features such as 'n\_words' and 'pam\_and\_jim\_key\_episode' had positive correlations with IMDb rating. From these results we make the following recomendations:

**Directors: Greg Daniels or Paul Feig** 

Writers: Paul Liberstein, Mindy Kaling, Greg Daniels, or B. J. Novak

Minimal list of Main Characters: Dwight, Michael, Jim, Creed, Stanley, and Pam.

Characters to consider not including: Erin.

As per other Pam and Jim key episodes, the plot should atleast partially follow their relationship, maybe showing how this has progressed since the finale episode. That said, following our choice of directors and writers, these should be given creative freedom as our model shows that this has had a positive influence on the success of historic episodes. We do not provide advice on the number of lines and directions to be included, as we trust these will be determined by the choice of director and writer. However we refer them to our analysis that suggests a high number of lines and directions has a possitive impact on the rating. We leave it up to the writers and directors to decide on how to implement this advice into their visons for the reunion episode. Similarly our analysis suggests that the rating responds positively to the episode starting with a cold open, and containing Jim pranks, which is something we suggest the writers consider including.

Our analysis suggests that releasing the episode during spring, or failing this summer, may have a positive impact of the rating, whilst releasing in autumn may best be avoided.

## 5. References

In this section, you should present a list of external sources (except the course materials) that you used during the project, if any

Additional data sources can be cited here, in addition to related python documentations, any other webpage sources that you benefited from

- Evkaya, Ozan. 2023. The Office CSV Data File, MATH11205: Machine Learning in Python, The University of Edinburgh.
- Various authors. 2023a. List of Cold Opens, Dunderpedia: The Office Encylopedia. Accessed on 08/03/2023. URL: https://theoffice.fandom.com/wiki/List\_of\_Cold\_Opens
- Various authors. 2023b. List of Jim's Pranks, Dunderpedia: The Office Encylopedia. Accessed on 06/03/2023. URL: https://theoffice.fandom.com/wiki/List\_of\_Jim%27s\_pranks
- West, Kelly. 2020. The Office: Jim Halpert And Pam Beesly's Relationship Told In 45 Episodes, CinemaBlend. Accessed on 05/03/2023. URL: https://www.cinemablend.com/television/2487769/the-office-jim-halpert-and-pam-beeslys-relationship-episode-playlist