Jigsaw_Analysis

August 22, 2019

```
[2]: %load_ext autoreload
%autoreload 2
[3]: import time
import tensorflow as tf
import pandas as pd
import gensim

import p8_util
import p8_util_config
import p9_util
```

Using TensorFlow backend.

1 1. Data loading

'sad', 'likes', 'disagree', 'sexual_explicit',

```
'identity_annotator_count', 'toxicity_annotator_count'], dtype='object')
```

2 2. Data pre-processing

2.1 2.1. Check NAN rate into columns

[236]:	ser_train_isna = df_train.isna().sum()	
	ser_train_isna		
[236]:	id	0	
	target	0	
	comment_text	0	
	severe_toxicity	0	
	obscene	0	
	identity_attack	0	
	insult	0	
	threat	0	
	asian	1399744	
	atheist	1399744	
	bisexual	1399744	
	black	1399744	
	buddhist	1399744	
	christian	1399744	
	female	1399744	
	heterosexual	1399744	
	hindu	1399744	
	homosexual_gay_or_lesbian	1399744	
	<pre>intellectual_or_learning_disability</pre>	1399744	
	jewish	1399744	
	latino	1399744	
	male	1399744	
	muslim	1399744	
	other_disability	1399744	
	other_gender	1399744	
	other_race_or_ethnicity	1399744	
	other_religion	1399744	
	other_sexual_orientation	1399744	
	physical_disability	1399744	
	<pre>psychiatric_or_mental_illness</pre>	1399744	
	transgender	1399744	
	white	1399744	
	created_date	0	
	<pre>publication_id</pre>	0	
	parent_id	778646	
	article_id	0	
	rating	0	

```
funny
                                                 0
                                                 0
wow
sad
                                                 0
likes
                                                 0
disagree
                                                 0
sexual_explicit
                                                 0
identity_annotator_count
                                                 0
toxicity_annotator_count
                                                 0
dtype: int64
```

2.1.1 Data description

2.2 2.2. Filter features without Nan values

Number of valid columns= 20

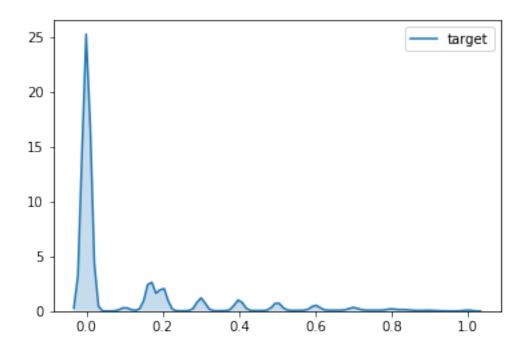
2.2.1 Columns are filtered from dataframe

Columns with Nan values are droped: this leads to having 20 features.

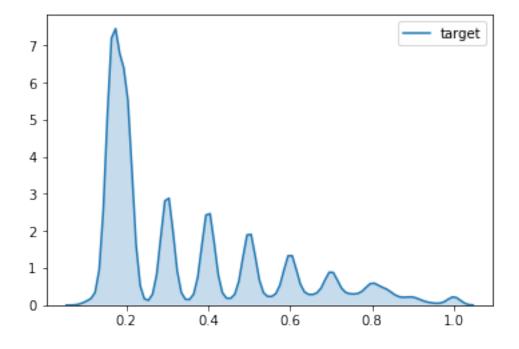
```
[96]: df_train_filtered = df_train[list_colname]
    df_train_filtered.shape, df_train.shape

[96]: ((1804874, 20), (1804874, 45))
```

2.3 2.3. Target (toxicity) distribution

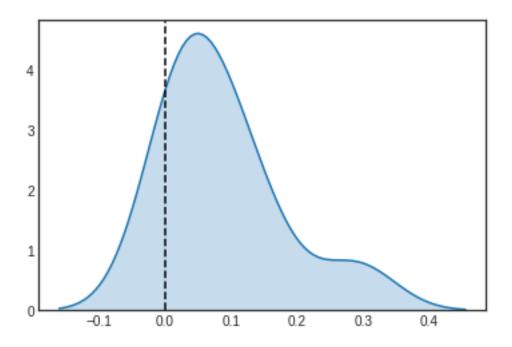


[189]: _=sns.kdeplot(ser_text_10pcent, shade=True)

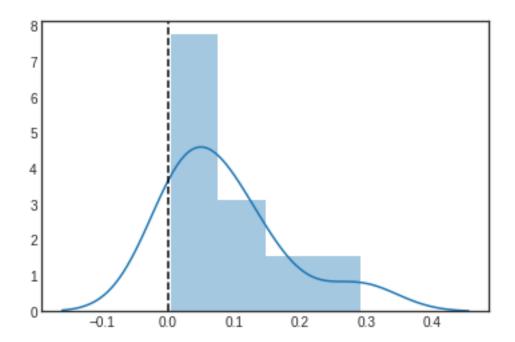


[226]: import p9_util

```
total = df_train_filtered.shape[0]
     dict_percent = dict()
     verbose = True
     for val in range(1,10,1):
         threshold = val/10
         percent = p9_util.print_col_stat(df_train_filtered, col,threshold,__
       →verbose=verbose)
         if verbose:
             print()
         dict_percent[val] = percent
     Number of texts where toxicity > 0.1 : 524054
     Percentage of texts where toxicity > 0.1 : 0.290355
     Number of texts where toxicity > 0.2 : 270194
     Percentage of texts where toxicity > 0.2 : 0.149702
     Number of texts where toxicity > 0.3 : 206991
     Percentage of texts where toxicity > 0.3 : 0.114684
     Number of texts where toxicity > 0.4 : 151463
     Percentage of texts where toxicity > 0.4 : 0.083919
     Number of texts where toxicity > 0.5 : 106438
     Percentage of texts where toxicity > 0.5 : 0.058973
     Number of texts where toxicity > 0.6 : 72235
     Percentage of texts where toxicity > 0.6 : 0.040022
     Number of texts where toxicity > 0.7 : 45451
     Percentage of texts where toxicity > 0.7 : 0.025182
     Number of texts where toxicity > 0.8 : 23802
     Percentage of texts where toxicity > 0.8 : 0.013188
     Number of texts where toxicity > 0.9 : 7517
     Percentage of texts where toxicity > 0.9 : 0.004165
[227]: pd.Series(dict_percent).min()
[227]: 0.004164833667059307
[228]: import pandas as pd
     _=sns.kdeplot(pd.Series(dict_percent), shade=True)
     plt.axvline(0, color="k", linestyle="--");
```



[229]: sns.distplot(pd.Series(dict_percent), kde=True);
plt.axvline(0, color="k", linestyle="--");



2.4 2.4. Extract a test dataset from train file

- Dataset with excluded Nan values results in df_train_filtered dataframe.
- Features are studied over a fraction of the train dataset (1%) resulting in df_train_sample.
- A test dataset is extracted from df_train_sample using indexes excluded from df_train_filtered. Result is df_test_filtered dataframe.

```
[237]: df_train_sample = df_train_filtered.sample(frac=0.01,random_state=0)
      print(df_train_sample.shape)
     (18049, 20)
     Get test dataset from indexes into df_train_filtered that do not belongs to train dataset
[238]: df_test_filtered = df_train_filtered.query("index!="+str(list(df_train_sample.
       →index)))
     Check consistency after building test dataset and train dataset operations from df_train_filter
     It is expected to have equality: Size of train dataset + size of test dataset = size of sample
[239]: df_train_filtered.shape[0],df_test_filtered.shape[0] + df_train_sample.shape[0]
[239]: (1804874, 1804874)
[240]: print(df_train_sample.shape)
      print(df_test_filtered.shape)
     (18049, 20)
     (1786825, 20)
        • df_train_filtered holds data with excluded Nan features values.
        • 1% of df_train_filtered is used for analysis.
        • 30% from df_test_filtered is used leading to df_test_sample.
[241]: # 1% of df_train_filtered is used for analysis.
      frac_train = df_train_filtered.shape[0]//100
      print("\nNumber rows for analysis train dataset= {}".format(frac_train))
      # 30% issued from 1% of df_train filtered is used for test part analysis.
```

Number rows for analysis train dataset= 18048 Number rows for analysis test part of dataset= 6016

frac_test = frac_train//3

```
[242]: df_test_sample = df_test_filtered.sample(n=frac_test,random_state=0)
    print(df_test_sample.shape)
(6016, 20)
```

print("Number rows for analysis test part of dataset= {}".format(frac test))

3 3. Toxicity contributors analysis

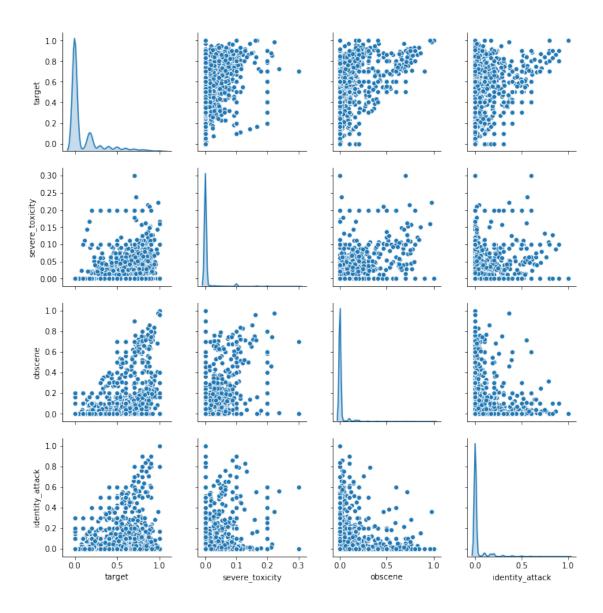
• df_train_part_1 and df_train_part_2 contain toxicity contributors for analysis.

/home/bangui/.local/lib/python3.6/site-packages/matplotlib/__init__.py:886: MatplotlibDeprecationWarning:

examples.directory is deprecated; in the future, examples will be found relative to the 'datapath' directory.

"found relative to the 'datapath' directory.".format(key))

[13]: <seaborn.axisgrid.PairGrid at 0x7fe4bf356b70>

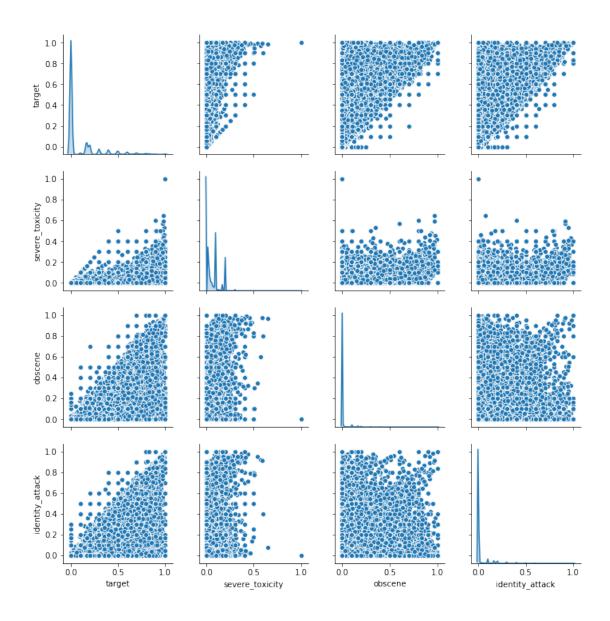


```
[46]: import seaborn as sns sns.pairplot(df_train_part_1, diag_kind="kde")
```

examples.directory is deprecated; in the future, examples will be found relative to the 'datapath' directory.

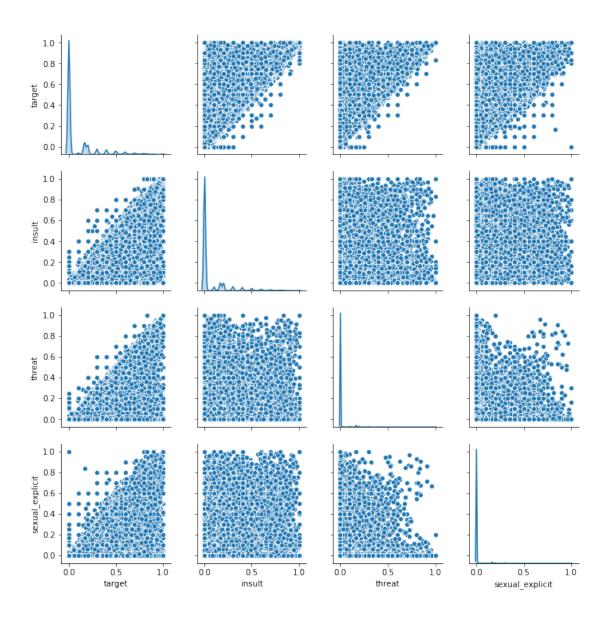
"found relative to the 'datapath' directory.".format(key))

[46]: <seaborn.axisgrid.PairGrid at 0x7f399aada860>



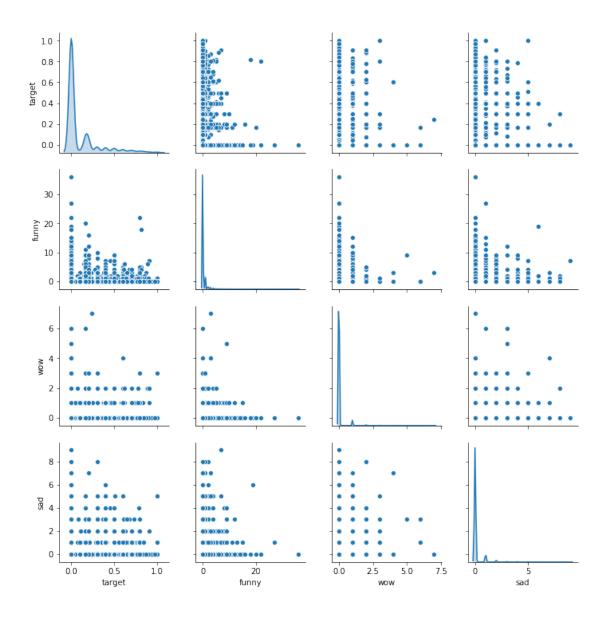
```
[47]: import seaborn as sns sns.pairplot(df_train_part_2, diag_kind="kde")
```

[47]: <seaborn.axisgrid.PairGrid at 0x7f3957043048>



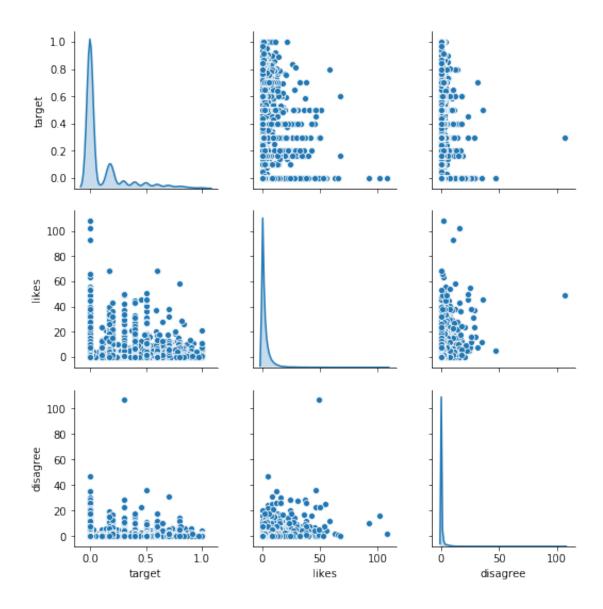
```
[256]: import seaborn as sns sns.pairplot(df_train_part_3, diag_kind="kde")
```

[256]: <seaborn.axisgrid.PairGrid at 0x7f8411b52d30>



```
[257]: import seaborn as sns sns.pairplot(df_train_part_4, diag_kind="kde")
```

[257]: <seaborn.axisgrid.PairGrid at 0x7f8404e6ccc0>



3.1 3.1. Feature engineering: new_feature

Combinaison of features that contribute to toxicity detection and forming new_feature shows that rectangle-triangle is replaced with a shape in which linear correlation between target and new_feature is more obvious.

All features does not contribute with the same weight to the toxicity. new_feature is then formed from weights issued from correlations matrix.

3.1.1 Features to toxicity contribution are filtered

• Result lies on dataframe df_train_sample

```
[103]: list_feature_excluded =
      list_feature_excluded +=['rating','funny','wow','sad','likes','disagree']
     list feature excluded
[103]: ['comment_text',
      'created_date',
      'publication_id',
      'article_id',
      'identity_annotator_count',
      'toxicity_annotator_count',
      'rating',
      'funny',
      'wow',
      'sad',
      'likes',
      'disagree']
[104]: list_feature = df_train_sample.columns
     list_feature = list_feature[1:-1]
     list_feature = [feature for feature in list_feature if feature not in_
      →list_feature_excluded ]
     list_feature
[104]: ['target',
      'severe_toxicity',
      'obscene',
      'identity_attack',
      'insult',
      'threat',
      'sexual explicit']
[271]: df = df_train_sample[list_feature]
     res = df.query('target>0.3').sample(2).head()
     print(df_train_sample['comment_text'].loc[res.index[0]])
     print()
     print(df_train_sample['comment_text'].loc[res.index[1]])
```

So why not worry about the scandal of heterosexual priests? Why not act against the "straight" sexually active priests and hierarchs with families or girlfriends on the side? Why not act against the "straight" priests and hierarchs who abuse or hide abusers? There's been a helluva lot more scandal from that, despite some people's efforts to portray it as a gay issue. The claim of protecting the Church by keeping gay men out of the seminaries is, bluntly, hypocrisy.

Pathetic. Is that the best the Globe can do?
A drop of (so far) less than 1% is somehow relevant to judge Trump?

How many 1% drops did we have during the Obama years?

When the market goes down big time, AND IT WILL, it will have little to do with Trump, but with China, Fed policies of the past decade, and a war somewhere that probably is mostly due to Obama's weak leadership in foreign affairs.

By spewing this kind of nonsense, the media enfeeble themselves and take themselves out of the conversation of serious people discussing serious issues.

```
[271]:
                target severe_toxicity
                                          obscene identity_attack
                                                                      insult \
      357023 0.400000
                                    0.1 0.100000
                                                               0.2 0.300000
      555712 0.833333
                                    0.0 0.166667
                                                               0.0 0.833333
              threat sexual_explicit
      357023
                0.0
      555712
                0.0
                                  0.0
[267]:
```

The sore losers will be out heavy on this one. The truth drives them batty. Say... come on now king kong... er i mean waveloser. Lets hear your brilliant analysis. Wootwoot!!!!

I only got the email notification which said something about Smug. I believe in full disclosure - just so all our presuppositions are on the table. Yes I am a white, Christian, conservative male - just the type you most likely spent most of your years persecuting. You use a clearly isolated incident with the EPD, and ignore the hundreds of thousands of good things they do. Pretty pathetic if you ask me, and the taxpayers are paying you for this as well. SAD

```
[272]: len("I only got the email notification which said something about Smug. I_{\sqcup} \rightarrow believe in full disclosure - just so all our presuppositions are on the \sqcup \rightarrow table. Yes I am a white, Christian, conservative male - just the type you \sqcup \rightarrow most likely spent most of your years persecuting. You use a clearly isolated \sqcup \rightarrow incident with the EPD, and ignore the hundreds of thousands of good things \sqcup \rightarrow they do. Pretty pathetic if you ask me, and the taxpayers are paying you for \sqcup \rightarrow this as well. SAD")
```

[272]: 461

```
[105]: df_train_sample = df_train_sample[list_feature]
```

3.1.2 Feature engineering: building new_feature

- Features such as:
 - threat, insult, sexual_explicit, identity_attack, obscene, severe_toxicity are combined all-together forming to a new feature.

- Those features do contribute to toxicity text detection.
- Those features draw a rectangle-triangle compared with target column.
- A linear correlation relation is suspected between those features and target.

In a first step, all weights are fixed to 1. It is supposed that all those features have the same weight considering target value.

Weights are handled into ser_weight_unit Series.

```
[106]: import numpy as np
      import pandas as pd
      df_train_sample.shape
      arr_unit = np.array([1. for value in list_feature])
      ser_weight_unit =pd.Series( arr_unit, index=list_feature)
      print(ser_weight_unit)
                         1.0
     target
     severe_toxicity
                         1.0
                         1.0
     obscene
     identity_attack
                         1.0
     insult
                         1.0
     threat
                         1.0
     sexual_explicit
                         1.0
     dtype: float64
```

• df_train_sample is added with a new column named new_feature. This new feature is the combination of all other columns with weight values to 1.

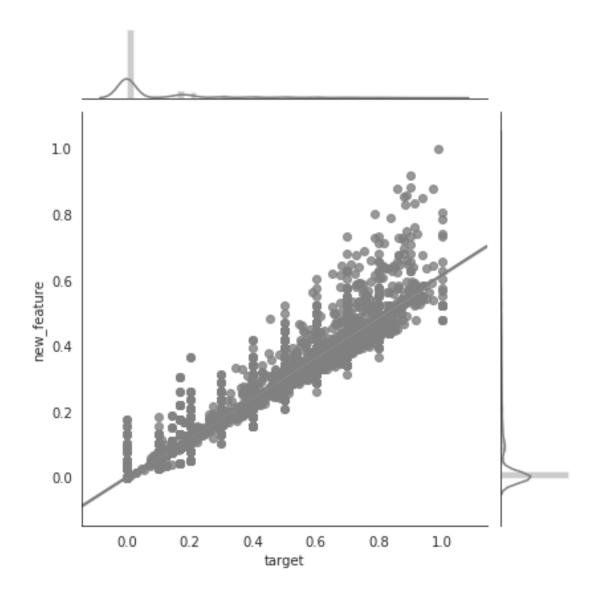
```
[112]: import p9_util
      df_train_sample = p9_util.df_weight_newFeature(df_train_sample,__

¬ser_weight_unit, list_feature, 'new_feature')
      print()
      print(df_train_sample.columns)
      print(df_train_sample.shape)
     target 1.0
     severe_toxicity 1.0
     obscene 1.0
     identity_attack 1.0
     insult 1.0
     threat 1.0
     sexual_explicit 1.0
     Index(['target', 'severe_toxicity', 'obscene', 'identity_attack', 'insult',
            'threat', 'sexual explicit', 'new feature'],
           dtype='object')
     (18049, 8)
```

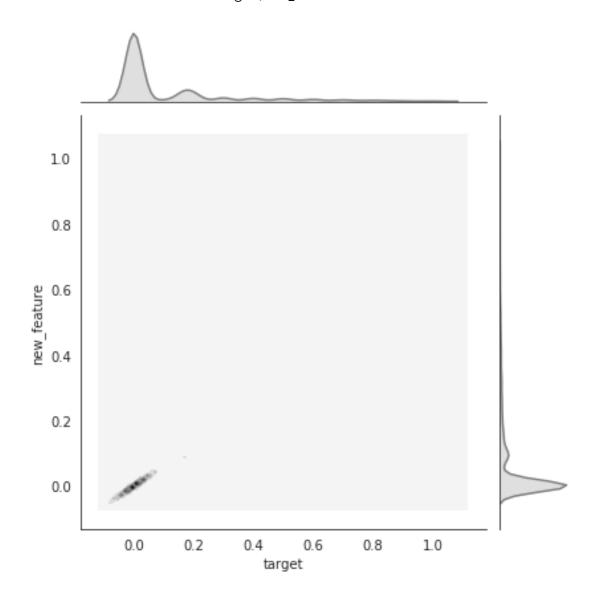
Pearson coefficient between (target,new_feature) = 0.9772645985025511

/home/bangui/.local/lib/python3.6/site-packages/scipy/stats/stats.py:1713: FutureWarning: Using a non-tuple sequence for multidimensional indexing is deprecated; use `arr[tuple(seq)]` instead of `arr[seq]`. In the future this will be interpreted as an array index, `arr[np.array(seq)]`, which will result either in an error or a different result.

return np.add.reduce(sorted[indexer] * weights, axis=axis) / sumval



Displaying density distributions give additional interpretation. Features are correlated strongly in interval [0.0, 0.2]

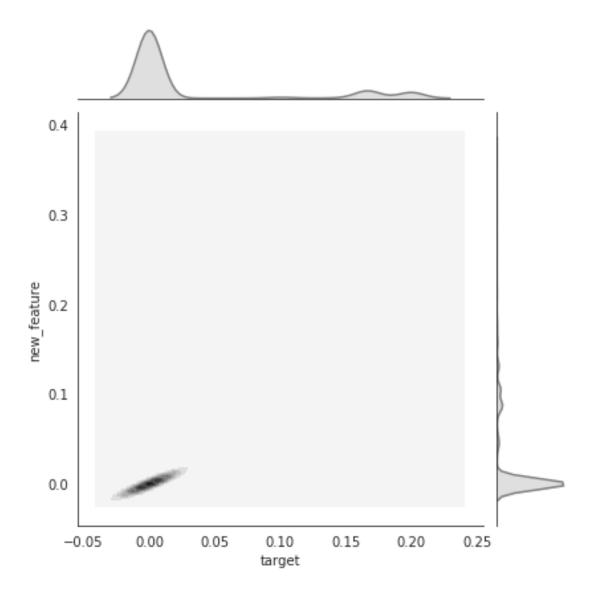


Then let's plot target values >0.2 and target values <= 0.2 This is achieved applying query to df_train_sample dataframe.

```
[29]: select="target <= 0.2"
    df_train_sample_select_0 = df_train_sample.query(select, inplace=False)
    select="target > 0.2"
    df_train_sample_select_1 = df_train_sample.query(select, inplace=False)

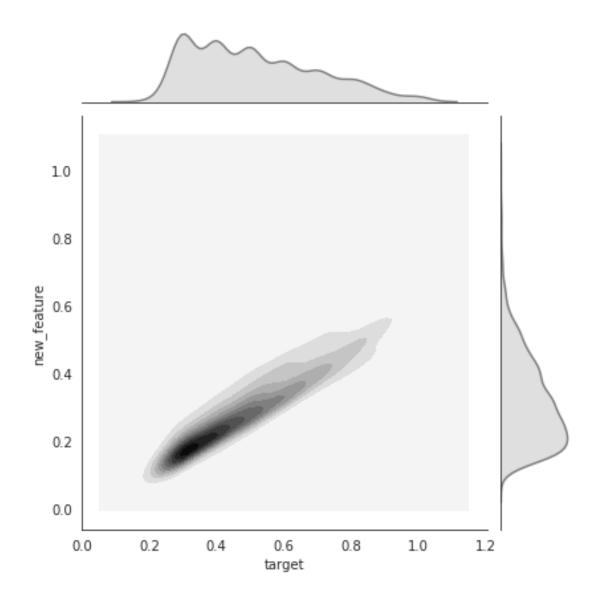
[30]: import scipy
    import p3_util
```

Pearson coefficient between (target,new_feature) = 0.929778393032554



```
[31]: import scipy import p3_util
```

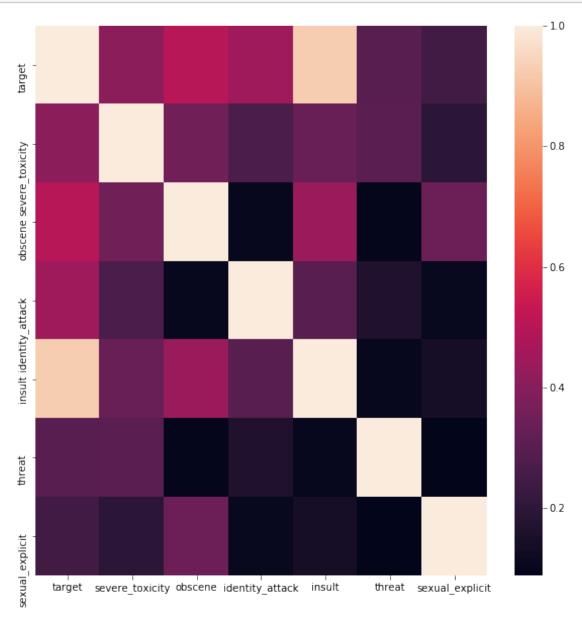
Pearson coefficient between (target,new_feature) = 0.9035771413019859



3.2 Weighting features with matrix correlations

```
[32]: df_train_sample.max()
[32]: target
                        1.0
     severe_toxicity
                        0.3
                        1.0
     obscene
     identity_attack
                        1.0
                        1.0
     insult
                        1.0
     threat
                        1.0
     sexual_explicit
    new_feature
                        1.0
    dtype: float64
[33]: if 'new_feature' in df_train_sample :
         del(df_train_sample['new_feature'])
[34]: from sklearn.preprocessing import StandardScaler
     std scaler = StandardScaler()
     print(std_scaler.fit(df_train_sample))
     X_std = std_scaler.transform(df_train_sample)
    StandardScaler(copy=True, with_mean=True, with_std=True)
[35]: df_train_sample.columns
[35]: Index(['target', 'severe_toxicity', 'obscene', 'identity_attack', 'insult',
            'threat', 'sexual_explicit'],
           dtype='object')
[36]: df_train_sample_std = pd.DataFrame(X_std, columns=df_train_sample.columns)
[37]: df_train_sample_std.head(3)
[37]:
          target severe_toxicity obscene identity_attack
                                                                 insult
                                                                           threat \
                        -0.209535 -0.215539
     0 -0.527767
                                                   -0.288058 -0.466598 -0.190872
     1 -0.527767
                        -0.209535 -0.215539
                                                   -0.288058 -0.466598 -0.190872
     2 -0.527767
                       -0.209535 -0.215539
                                                   -0.288058 -0.466598 -0.190872
        sexual_explicit
     0
              -0.136735
              -0.136735
     1
              -0.136735
     2
[38]: import pandas as pd
     import seaborn as sns
     import matplotlib.pyplot as plt
     X_std_corr = df_train_sample_std.corr()
```

```
f, ax = plt.subplots(figsize=(10, 10))
_z = sns.heatmap(X_std_corr, annot=False)
```



[45]: X_std_corr[:]	<pre>X_std_corr[:]</pre>						
[45]:	target	severe_toxicity	obscene	identity_attack	\		
target	1.000000	0.405802	0.500494	0.448081			
severe_toxicity	0.405802	1.000000	0.352787	0.267649			
obscene	0.500494	0.352787	1.000000	0.104657			
identity_attack	0.448081	0.267649	0.104657	1.000000			
insult	0.925399	0.333349	0.438020	0.295346			
threat	0.296270	0.303908	0.096228	0.164630			

```
insult
                                   threat
                                           sexual_explicit
     target
                      0.925399 0.296270
                                                  0.244588
     severe_toxicity 0.333349 0.303908
                                                  0.193290
                                0.096228
                                                  0.338451
     obscene
                      0.438020
     identity_attack 0.295346 0.164630
                                                  0.107064
     insult
                      1.000000 0.104562
                                                  0.139459
     threat
                      0.104562 1.000000
                                                  0.087756
     sexual_explicit 0.139459 0.087756
                                                   1.000000
    Weights assignement for building new_feature Weights are issued from correlation matrix.
[49]: list_feature
[49]: ['target',
      'severe_toxicity',
      'obscene',
      'identity_attack',
      'insult',
      'threat',
      'sexual_explicit']
[50]: X_std_corr['target']
[50]: target
                        1.000000
     severe_toxicity
                        0.405802
                        0.500494
     obscene
     identity_attack
                        0.448081
     insult
                        0.925399
                        0.296270
     threat
     sexual explicit
                        0.244588
     Name: target, dtype: float64
[51]: df_train_filtered_sample = df_weight_newFeature(df_train_sample,__

¬X_std_corr['target'], list_feature, 'new_feature')
    target 1.0
    severe_toxicity 0.40580207453035044
    obscene 0.5004938914679019
    identity attack 0.44808065477165393
    insult 0.9253985594798301
    threat 0.29627000932665676
    sexual_explicit 0.24458800527543303
[52]: df_train_filtered_sample['new_feature'].min(),__

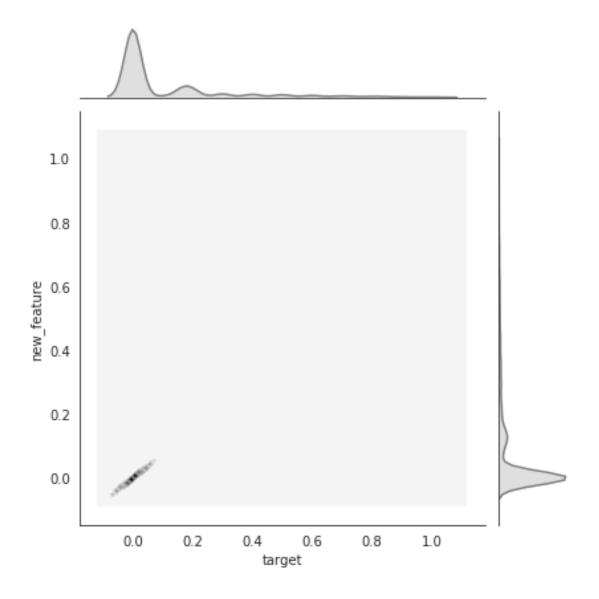
→df_train_filtered_sample['new_feature'].max()
[52]: (0.0, 1.0)
```

0.193290 0.338451

0.107064

sexual_explicit 0.244588

Pearson coefficient between (target,new_feature) = 0.9897087258200451



Using correlation matrix for weighting features when building new_feature improve global correlation between new_feature and target (0.989 vs 0.977).

Display Pearson coefficient between all features against target

```
[83]: df_train_filtered_sample.columns
[83]: Index(['severe_toxicity', 'obscene', 'identity_attack', 'insult', 'threat',
            'sexual explicit'],
           dtype='object')
[55]: import scipy
     import p3_util_plot
     var1 = 'target'
     for var2 in df_train_filtered_sample.columns :
         if var2 != 'target' :
             #p3 util_plot.df_sns_joint_plot(df_train_filtered_sample, var1, var2,__
      →parameter_kind='reg', parameter_color='grey')
             pearson = scipy.stats.

-pearsonr(df_train_filtered_sample[var1],df_train_filtered_sample[var2])
             print("\nPearson coefficient between ({},{}) = {}".format(var1,var2,__
      →pearson[0]))
         else :
             pass
```

```
Pearson coefficient between (target,severe_toxicity) = 0.4058020745302237

Pearson coefficient between (target,obscene) = 0.5004938914680612

Pearson coefficient between (target,identity_attack) = 0.448080654771592

Pearson coefficient between (target,insult) = 0.9253985594798431

Pearson coefficient between (target,threat) = 0.29627000932655323

Pearson coefficient between (target,sexual_explicit) = 0.2445880052754865

Pearson coefficient between (target,new_feature) = 0.9897087258200451
```

Correlation between insult and target

```
[56]: import scipy import p3_util_plot
```

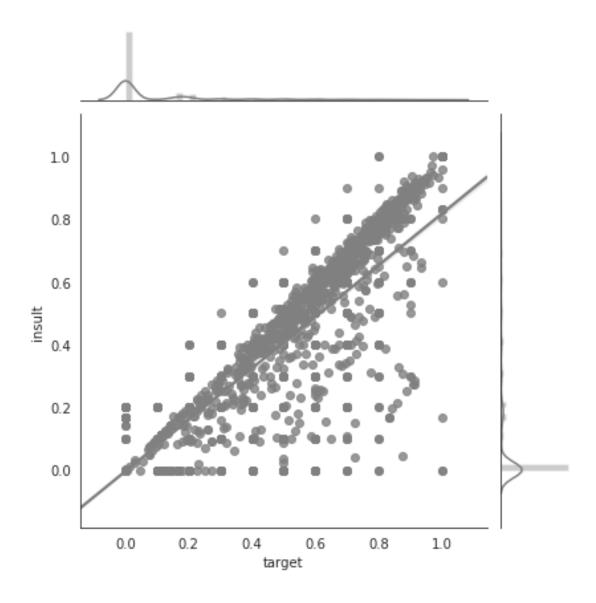
```
var1 = 'target'
var2 = 'insult'
p3_util_plot.df_sns_joint_plot(df_train_filtered_sample, var1, var2,
parameter_kind='reg', parameter_color='grey')

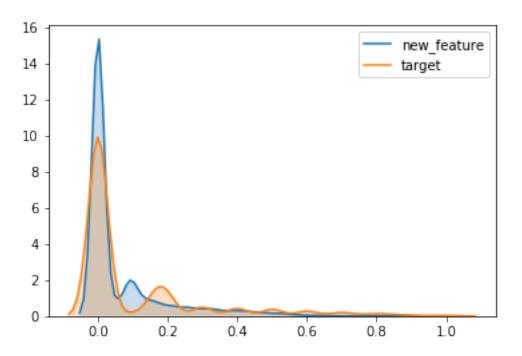
pearson = scipy.stats.
pearsonr(df_train_filtered_sample[var1],df_train_filtered_sample[var2])
print("\nPearson coefficient between ({{}},{{}}) = {{}}".format(var1,var2,
pearson[0]))
```

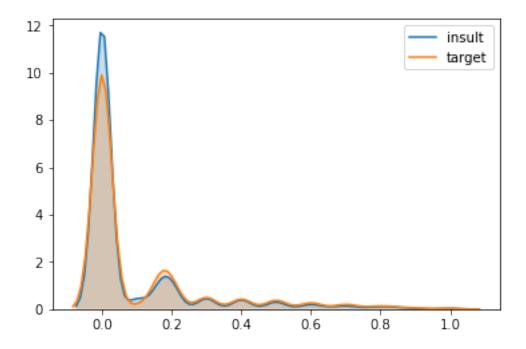
/home/bangui/.local/lib/python3.6/site-packages/scipy/stats/stats.py:1713: FutureWarning: Using a non-tuple sequence for multidimensional indexing is deprecated; use `arr[tuple(seq)]` instead of `arr[seq]`. In the future this will be interpreted as an array index, `arr[np.array(seq)]`, which will result either in an error or a different result.

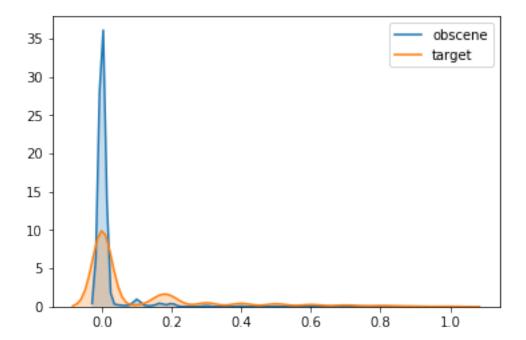
return np.add.reduce(sorted[indexer] * weights, axis=axis) / sumval

Pearson coefficient between (target, insult) = 0.9253985594798431









3.2.1 Weights issued from a linear regression

Target is extracted from dataframe Dataframe keeps features used only for linear regression.

Data is standardized

```
[60]: from sklearn.preprocessing import StandardScaler
if False :
    std_scaler = StandardScaler()
    print(std_scaler.fit(df_train_sample))

    X_std = std_scaler.transform(df_train_sample)
else :
    X_std = df_train_sample.copy()
```

Linear regression model is built

```
[61]: from sklearn.linear_model import LinearRegression

model_regression = LinearRegression().fit(X_std, df_train_label)
```

Displaying correlation coefficient Coefficient issued from linear regression and those issued from matrix correlation are both displayed.

Some of them have closed values, such as threat and identity_attack, others strongly differ.

```
[62]: for feature, lr_coef, corr_coef in zip(df_train_sample.columns, □ → model_regression.coef_, X_std_corr['target'][1:]):

print("Feature: {} : linear regression coeff= {} / Correlation matrix □ → coeff= {}".format(feature, lr_coef, corr_coef))
```

```
Feature: severe_toxicity : linear regression coeff= -0.05362561443422439 / Correlation matrix coeff= 0.40580207453035044

Feature: obscene : linear regression coeff= 0.2619629875316454 / Correlation matrix coeff= 0.5004938914679019

Feature: identity_attack : linear regression coeff= 0.40888237843007297 / Correlation matrix coeff= 0.44808065477165393

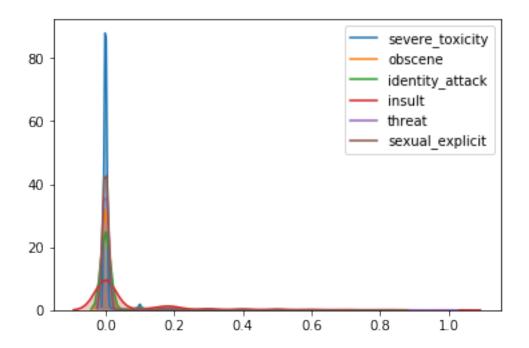
Feature: insult : linear regression coeff= 0.9155839990091565 / Correlation matrix coeff= 0.9253985594798301

Feature: threat : linear regression coeff= 0.6734955460989804 / Correlation matrix coeff= 0.29627000932665676

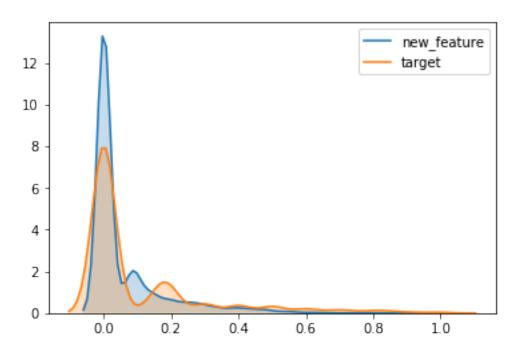
Feature: sexual_explicit : linear regression coeff= 0.31847938964928624 / Correlation matrix coeff= 0.24458800527543303
```

```
[70]: if 'target' in df_test_sample.columns :
          df test target = df test sample.pop('target')
      df_test_sample.columns
 [70]: Index(['id', 'comment_text', 'severe_toxicity', 'obscene', 'identity_attack',
             'insult', 'threat', 'created_date', 'publication_id', 'article_id',
             'rating', 'funny', 'wow', 'sad', 'likes', 'disagree', 'sexual explicit',
             'identity_annotator_count', 'toxicity_annotator_count'],
            dtype='object')
 [71]: list_feature
      if 'target' in list feature :
          list_feature.remove('target')
      print(list_feature)
     ['severe_toxicity', 'obscene', 'identity_attack', 'insult', 'threat',
     'sexual_explicit']
 [72]: df_test_sample = df_test_sample[list_feature]
  : X std = std scaler.transform(df test sample)
 [76]: model_regression.score(X_std, df_test_target)
 [76]: -48.750536382222116
     3.2.2 Distribution of target from test dataset
[122]: df_test_sample.columns
[122]: Index(['id', 'target', 'comment_text', 'severe_toxicity', 'obscene',
             'identity_attack', 'insult', 'threat', 'created_date', 'publication_id',
             'article_id', 'rating', 'funny', 'wow', 'sad', 'likes', 'disagree',
             'sexual_explicit', 'identity_annotator_count',
             'toxicity_annotator_count'],
            dtype='object')
[123]: import numpy as np
      import pandas as pd
      df_train_sample.shape
      arr_unit = np.array([1. for value in list_feature])
      ser_weight_unit =pd.Series( arr_unit, index=list_feature)
      print(ser_weight_unit)
     target
                         1.0
                         1.0
     severe_toxicity
     obscene
                         1.0
     identity_attack
                         1.0
     insult
                         1.0
```

```
threat
                         1.0
     sexual_explicit
                        1.0
     dtype: float64
[124]: import p9_util
      df_test_sample = p9_util.df_weight_newFeature(df_test_sample, ser_weight_unit,__
      →list_feature, 'new_feature')
      print()
      print(df_test_sample.columns)
      print(df_test_sample.shape)
     target 1.0
     severe_toxicity 1.0
     obscene 1.0
     identity_attack 1.0
     insult 1.0
     threat 1.0
     sexual_explicit 1.0
     Index(['id', 'target', 'comment_text', 'severe_toxicity', 'obscene',
            'identity_attack', 'insult', 'threat', 'created_date', 'publication_id',
            'article_id', 'rating', 'funny', 'wow', 'sad', 'likes', 'disagree',
            'sexual_explicit', 'identity_annotator_count',
            'toxicity_annotator_count', 'new_feature'],
           dtype='object')
     (6016, 21)
 [74]: import seaborn as sns
      for col in df_test_sample.columns:
          sns.kdeplot(df_test_sample[col], shade=True)
```

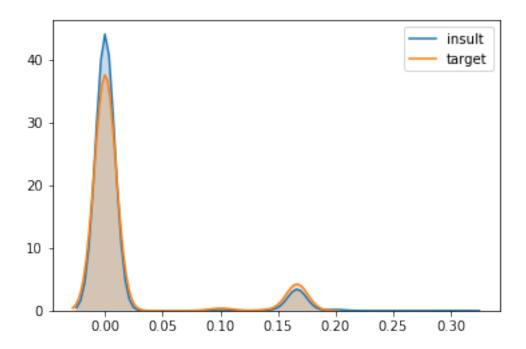


This display shows a long trail above 0.2 target column is added to df_test_sample. This will allow to extract a subset of df_test_sample filtered with target >0.2



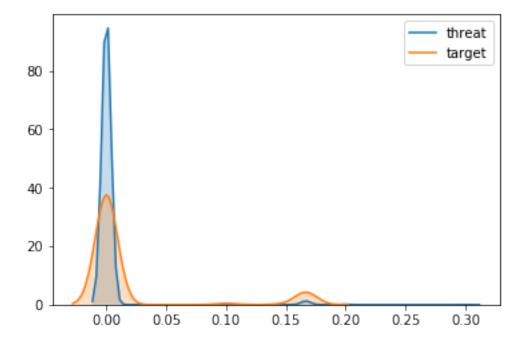
```
[79]: import seaborn as sns

for col in ['insult','target']:
    z_ = sns.kdeplot(df_test_sample_filtered[col], shade=True)
```



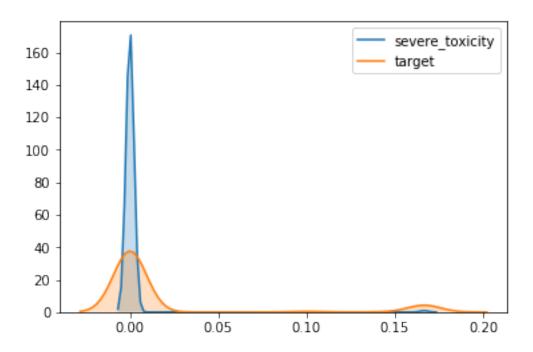
```
[430]: import seaborn as sns

for col in ['threat','target']:
    z_ = sns.kdeplot(df_test_sample_filtered[col], shade=True)
```



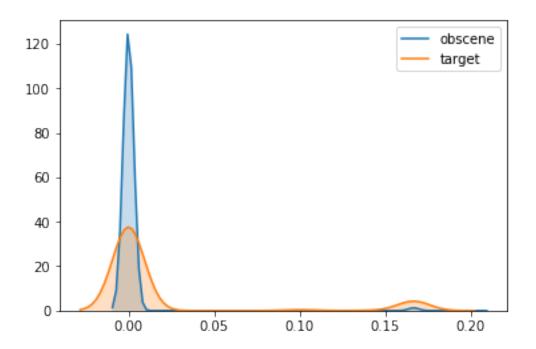
```
[431]: import seaborn as sns

for col in ['severe_toxicity','target']:
    z_ = sns.kdeplot(df_test_sample_filtered[col], shade=True)
```



```
[432]: import seaborn as sns

for col in ['obscene', 'target']:
    z_ = sns.kdeplot(df_test_sample_filtered[col], shade=True)
```

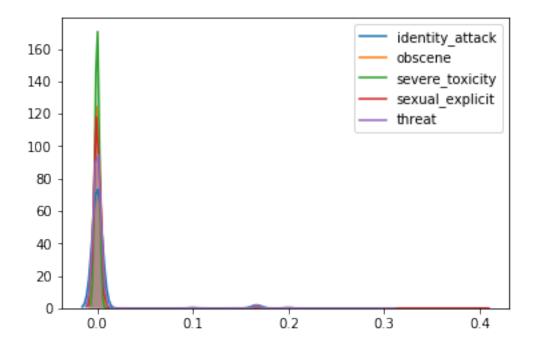


```
[433]: import seaborn as sns

for col in ['identity_attack','obscene',

□ 'severe_toxicity','sexual_explicit','threat']:

z_ = sns.kdeplot(df_test_sample_filtered[col], shade=True)
```



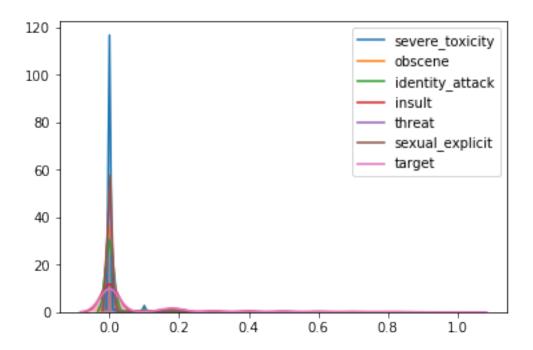
```
[436]: from scipy.stats import kstest col='insult' kstest(df_test_sample_filtered[col], 'norm')
```

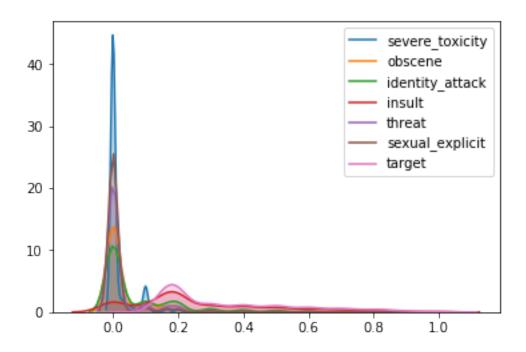
[436]: KstestResult(statistic=0.5, pvalue=0.0)

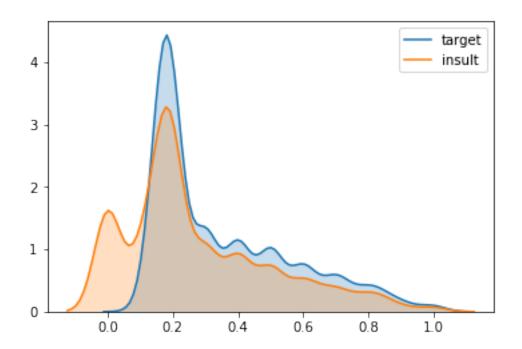
3.2.3 Distribution of target from train dataset

```
[439]: import seaborn as sns

for col in df_train_sample.columns:
    sns.kdeplot(df_train_sample[col], shade=True)
```







We can conclude that insult contributes the most to a toxic comment

```
[121]: df_test_sample.shape, df_train_sample.shape
[121]: ((6016, 20), (18049, 8))
[]:
```