## PN balance

January 10, 2022

```
import os
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import time
from imblearn.combine import SMOTETomek
from imblearn.under_sampling import NearMiss
from imblearn.over_sampling import RandomOverSampler
from collections import Counter
```

[2]: %run /data/emo/notebooks/source/pipeline/dataset\_loader.ipynb

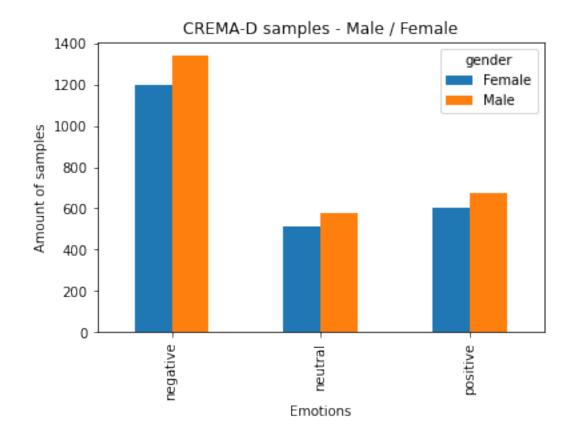
## 1 CremaBinair

```
[3]: # Histo Pos - Neg - Neu
df_male = MaleSplitCremaBinair.load_dataset()
df_female = FemaleSplitCremaBinair.load_dataset()

df = pd.concat([df_male, df_female])

df.groupby(['emotion', 'gender']).size().unstack(level=1).plot(kind='bar')
plt.title('CREMA-D samples - Male / Female')
plt.ylabel('Amount of samples')
plt.xlabel('Emotions')
plt.show()

print(df_male.groupby('emotion').size())
print("")
print(df_female.groupby('emotion').size())
```



emotion
negative 1341
neutral 575
positive 671
dtype: int64
emotion

negative 1199 neutral 512 positive 600 dtype: int64

```
# Oversampeling RandomSampler Male

# https://github.com/ufoym/imbalanced-dataset-sampler

# https://www.youtube.com/watch?v=OJedgzdipCO&ab_channel=KrishNaik

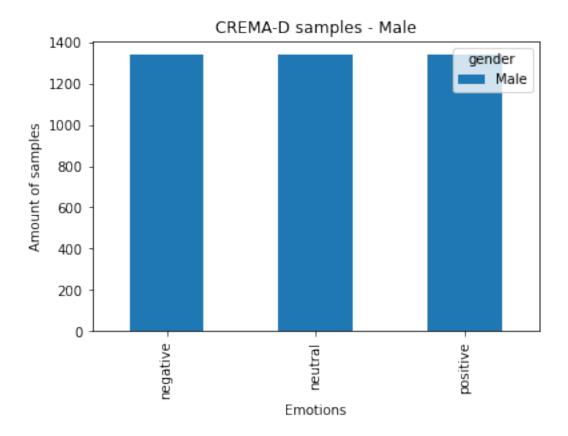
# https://github.com/krishnaikO6/Handle-Imbalanced-Dataset/blob/master/

→ Handling%20Imbalanced%20Data-%20Over%20Sampling.ipynb

# https://imbalanced-learn.org/stable/over_sampling.html
```

```
columns = df_male.columns.tolist()
columns = [c for c in columns if c not in ["emotion"]]
X = df_male[columns]
Y = df_male['emotion']
# Oversampeling
male_grouped_pos = df_male[df_male['emotion'] == 'positive']
male_grouped_neu = df_male[df_male['emotion'] == 'neutral']
male_grouped_neg = df_male[df_male['emotion'] == 'negative']
print(f"Pos:{male_grouped_pos.shape}, Neu:{male_grouped_neu.shape}, Neg:_u
 →{male_grouped_neg.shape}")
print(f"Original X:{X.shape}, Y:{Y.shape}")
ros = RandomOverSampler()
X_train_res, y_train_res = ros.fit_resample(X, Y)
y_train_pos = y_train_res[y_train_res == 'positive']
y_train_neu = y_train_res[y_train_res == 'neutral']
y_train_neg = y_train_res[y_train_res == 'negative']
print(f"Pos:{y_train_pos.shape}, Neu:{y_train_neu.shape}, Neg: {y_train_neg.
 →shape}")
print(f"Resampled X:{X_train_res.shape}, Y:{y_train_res.shape}")
print(X_train_res)
print(y_train_res)
X_train_res['emotion'] = y_train_res
Pos: (671, 4), Neu: (575, 4), Neg: (1341, 4)
Original X: (2587, 3), Y: (2587,)
Pos: (1341,), Neu: (1341,), Neg: (1341,)
Resampled X:(4023, 3), Y:(4023,)
     gender subset
                                                             file path
0
      Male
              None /data/emo/notebooks/source/datasets/crema/1023...
1
       Male
              None /data/emo/notebooks/source/datasets/crema/1001...
2
      Male
              None /data/emo/notebooks/source/datasets/crema/1040...
3
      Male
              None /data/emo/notebooks/source/datasets/crema/1034...
4
      Male
              None /data/emo/notebooks/source/datasets/crema/1035...
              None /data/emo/notebooks/source/datasets/crema/1086...
4018
      Male
      Male
              None /data/emo/notebooks/source/datasets/crema/1038...
4019
              None /data/emo/notebooks/source/datasets/crema/1051...
4020
      Male
```

```
4021
           Male
                  None /data/emo/notebooks/source/datasets/crema/1031...
    4022
           Male
                  None
                        /data/emo/notebooks/source/datasets/crema/1069...
    [4023 rows x 3 columns]
            negative
    0
             neutral
    1
    2
            negative
    3
             neutral
            negative
              •••
    4018
            positive
    4019
            positive
    4020
            positive
    4021
            positive
    4022
            positive
    Name: emotion, Length: 4023, dtype: object
[5]: # Histo Male
     df = X_train_res
     df.groupby(['emotion', 'gender']).size().unstack(level=1).plot(kind='bar')
     plt.title('CREMA-D samples - Male')
     plt.ylabel('Amount of samples')
     plt.xlabel('Emotions')
     plt.show()
     print(df.groupby('emotion').size())
```



```
emotion
negative 1341
neutral 1341
positive 1341
dtype: int64
```

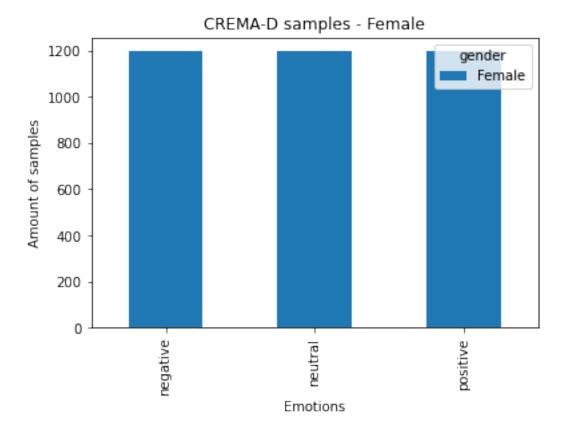
```
ros = RandomOverSampler()
X_train_res, y_train_res = ros.fit_resample(X, Y)
y_train_pos = y_train_res[y_train_res == 'positive']
y_train_neu = y_train_res[y_train_res == 'neutral']
y_train_neg = y_train_res[y_train_res == 'negative']
print(f"Pos:{y_train_pos.shape}, Neu:{y_train_neu.shape}, Neg: {y_train_neg.
 ⇒shape}")
print(f"Resampled X:{X_train_res.shape}, Y:{y_train_res.shape}")
print(X_train_res)
print(y_train_res)
X_train_res['emotion'] = y_train_res
Pos: (600, 4), Neu: (512, 4), Neg: (1199, 4)
Original X:(2311, 3), Y:(2311,)
Pos: (1199,), Neu: (1199,), Neg: (1199,)
Resampled X: (3597, 3), Y: (3597,)
      gender subset
                                                              file_path
0
     Female
               None /data/emo/notebooks/source/datasets/crema/1037...
1
     Female None /data/emo/notebooks/source/datasets/crema/1058...
2
              None /data/emo/notebooks/source/datasets/crema/1054...
      Female
3
     Female
               None /data/emo/notebooks/source/datasets/crema/1003...
               None /data/emo/notebooks/source/datasets/crema/1007...
4
     Female
3592 Female
              None /data/emo/notebooks/source/datasets/crema/1082...
3593 Female
               None /data/emo/notebooks/source/datasets/crema/1013...
3594 Female
               None /data/emo/notebooks/source/datasets/crema/1061...
3595 Female
               None /data/emo/notebooks/source/datasets/crema/1013...
3596 Female
               None /data/emo/notebooks/source/datasets/crema/1009...
[3597 rows x 3 columns]
0
        negative
1
        negative
2
        neutral
3
        negative
4
        negative
3592
        positive
3593
        positive
3594
        positive
3595
        positive
3596
        positive
```

Name: emotion, Length: 3597, dtype: object

```
[7]: # Histo Female
df = X_train_res

df.groupby(['emotion', 'gender']).size().unstack(level=1).plot(kind='bar')
plt.title('CREMA-D samples - Female')
plt.ylabel('Amount of samples')
plt.xlabel('Emotions')
plt.show()

print(df.groupby('emotion').size())
```



```
emotion
negative 1199
neutral 1199
positive 1199
dtype: int64
```

```
[8]: # Oversampeling SMOTETomek Male
```

```
# columns = df_male.columns.tolist()
# columns = [c for c in columns if c not in ["emotion"]]
\# X = df_male[columns]
\# Y = df_male['emotion']
# print(X)
# male_grouped_pos = df_male[df_male['emotion'] == 'positive']
# male grouped neu = df male[df male['emotion'] == 'neutral']
# male_grouped_neg = df_male[df_male['emotion'] == 'negative']
# print(f"Pos:{male_grouped_pos.shape}, Neu:{male_grouped_neu.shape}, Neq:_u
→ {male_grouped_neg.shape}")
# print(f"Original X:{X.shape}, Y:{Y.shape}")
# smk = SMOTETomek()
\# X_{res,y_{res=smk.fit_{resample}(X,Y)}
# y_train_pos = y_train_res[y_train_res == 'positive']
# y_train_neu = y_train_res[y_train_res == 'neutral']
# y train neg = y train res[y train res == 'negative']
# X_res.shape, y_res.shape
# print(f"Pos:{y_train_pos.shape}, Neu:{y_train_neu.shape}, Neg: {y_train_neg.
⇒shape}")
# print(f"Resampled X:{X train res.shape}, Y:{y train res.shape}")
# print(X_train_res)
# print(y_train_res)
# X_train_res['emotion'] = y_train_res
```

## ${\bf 2}\quad {\bf Quaternair Combined PN}$

```
[10]: df = QuaternairCombinedPN.load_dataset()

df.groupby(['emotion', 'gender']).size().unstack(level=1).plot(kind='bar')
    plt.title('QuaternairCombinedPN samples')
    plt.ylabel('Amount of samples')
    plt.xlabel('Emotions')
    plt.show()

print(df.groupby(['emotion', 'gender']).size())
```

```
Traceback (most recent call last)
/tmp/ipykernel_11595/2278170828.py in <module>
----> 1 df = QuaternairCombinedPN.load_dataset()
      3 df.groupby(['emotion', 'gender']).size().unstack(level=1).
→plot(kind='bar')
      4 plt.title('QuaternairCombinedPN samples')
      5 plt.ylabel('Amount of samples')
/tmp/ipykernel_11595/1532011642.py in load_dataset(cls)
                        value['emotion'] = 'neutral'
     21
---> 22
                    components = np.array([value, "Female", None, os.path.
 →join(path, file)])
     23
                    components.append(component)
     24
NameError: name 'path' is not defined
```

```
[]: # Oversampeling RandomSampler
     columns = df.columns.tolist()
     columns = [c for c in columns if c not in ["emotion"]]
     X = df[columns]
     Y = df['emotion']
     #Oversampeling
     gender_grouped_pos = df[df['emotion'] == 'positive']
     gender_grouped_neu = df[df['emotion'] == 'neutral']
     gender_grouped_neg = df[df['emotion'] == 'negative']
     print(f"Pos:{gender_grouped_pos.shape}, Neu:{gender_grouped_neu.shape}, Neg:__
     →{gender_grouped_neg.shape}")
     print(f"Original X:{X.shape}, Y:{Y.shape}")
     ros = RandomOverSampler()
     X_train_res, y_train_res = ros.fit_resample(X, Y)
     y_train_pos = y_train_res[y_train_res == 'positive']
     y_train_neu = y_train_res[y_train_res == 'neutral']
     y_train_neg = y_train_res[y_train_res == 'negative']
     print(f"Pos:{y_train_pos.shape}, Neu:{y_train_neu.shape}, Neg: {y_train_neg.
     ⇒shape}")
```

```
print(f"Resampled X:{X_train_res.shape}, Y:{y_train_res.shape}")

print(X_train_res)
print(y_train_res)

X_train_res['emotion'] = y_train_res

[]: df = X_train_res

df.groupby(['emotion', 'gender']).size().unstack(level=1).plot(kind='bar')
plt.title('QuaternairCombinedPN samples')
plt.ylabel('Amount of samples')
plt.xlabel('Emotions')
plt.show()

print(df.groupby(['emotion', 'gender']).size())
```