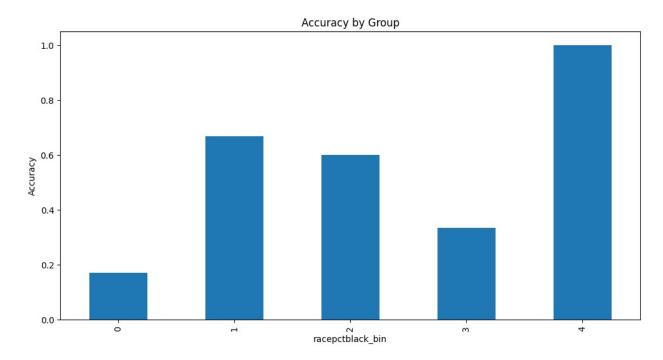
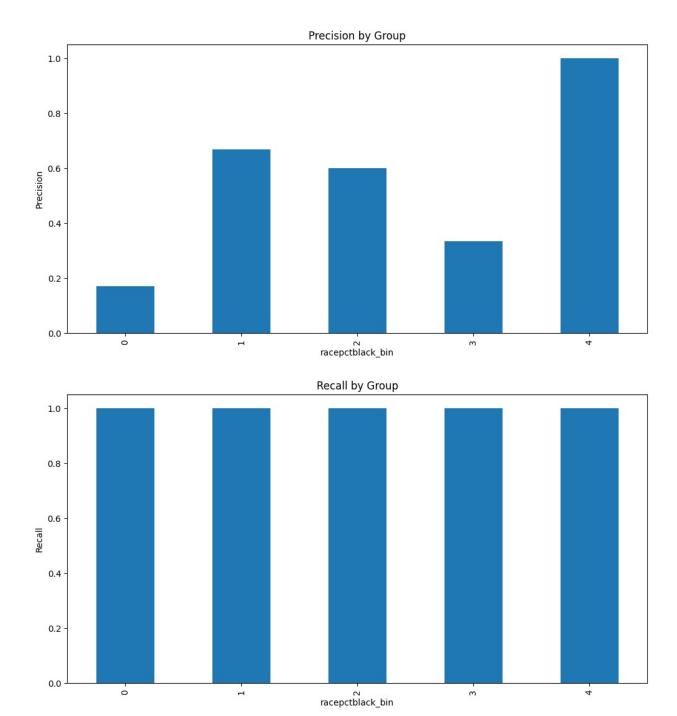
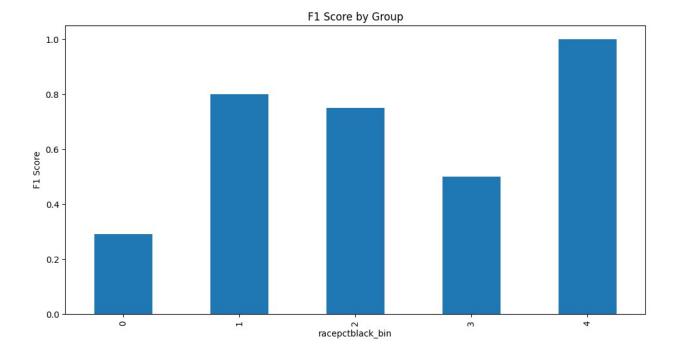
```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.model selection import train test split
from sklearn.ensemble import RandomForestRegressor
from sklearn.metrics import accuracy score, precision score,
recall score, f1 score, confusion matrix, mean squared error,
mean absolute error, r2 score
from fairlearn.metrics import MetricFrame,
demographic parity difference, equalized odds difference,
selection_rate, true_positive_rate, false_positive_rate,
false negative rate
from sklearn.metrics import ConfusionMatrixDisplay
# Load the cleaned dataset
data = pd.read csv('cleaned communities crime data.csv')
# Define the target and features
target = 'ViolentCrimesPerPop'
features = data.drop(columns=[target])
sensitive feature = 'racepctblack'
# Binarize the target variable based on the mean
threshold = data[target].mean()
data['ViolentCrimesPerPop binary'] = (data[target] >
threshold).astype(int)
# Discretize the sensitive feature
data['racepctblack bin'] = pd.cut(data['racepctblack'], bins=5,
labels=False)
# Split the dataset into training and testing sets
X train, X test, y train, y test = train test split(
    features, data['ViolentCrimesPerPop binary'], test size=0.2,
random state=42
# Train a Random Forest model
rf model = RandomForestRegressor(n estimators=100, random state=42)
rf model.fit(X train, y train)
# Predict using the trained model
y pred = rf model.predict(X test)
y_pred_binary = (y_pred > threshold).astype(int)
# Define a function to calculate different metrics
def compute metrics(y true, y pred):
    metrics = {
        'accuracy': accuracy score(y true, y pred),
```

```
'precision': precision_score(y_true, y_pred, zero_division=0),
        'recall': recall score(y true, y pred, zero division=0),
        'f1': f1_score(y_true, y_pred, zero_division=0)
    return metrics
# Compute metrics for different groups
metrics = MetricFrame(
    metrics=compute metrics,
    y_true=y_test,
    y_pred=y_pred_binary,
    sensitive features=data.loc[X test.index, 'racepctblack bin']
)
# Print the overall metrics
print("Overall Metrics:")
print(metrics.overall)
# Print metrics by sensitive feature groups
print("\nMetrics by Sensitive Feature Groups:")
print(metrics.by group)
# Extract metrics for visualization
metrics by group = metrics.by group.apply(pd.Series)
accuracy = metrics by group['accuracy']
precision = metrics by group['precision']
recall = metrics by group['recall']
f1 = metrics by group['f1']
# Plot Accuracy by Group
accuracy.plot(kind='bar', figsize=(12, 6), title='Accuracy by Group')
plt.ylabel('Accuracy')
plt.show()
# Plot Precision by Group
precision.plot(kind='bar', figsize=(12, 6), title='Precision by
Group')
plt.ylabel('Precision')
plt.show()
# Plot Recall by Group
recall.plot(kind='bar', figsize=(12, 6), title='Recall by Group')
plt.ylabel('Recall')
plt.show()
# Plot F1 Score by Group
f1.plot(kind='bar', figsize=(12, 6), title='F1 Score by Group')
plt.ylabel('F1 Score')
plt.show()
```







## Accuracy by Group:

The accuracy of the model varies significantly across different groups based on the racepctblack bin. The group with the highest proportion of black population (racepctblack\_bin = 4) has the highest accuracy, while the group with the lowest proportion (racepctblack\_bin = 0) has the lowest accuracy. This indicates a potential bias in the model, as it performs better for certain racial groups compared to others. Precision by Group:

Similar to accuracy, the precision also varies across groups. The highest precision is observed in the racepctblack\_bin = 4 group, and the lowest in the racepctblack\_bin = 0 group. This suggests that the model is better at identifying true positives in groups with a higher proportion of black population. Recall by Group:

The recall is consistent across all groups, indicating that the model has a uniform ability to identify actual positives regardless of the group. F1 Score by Group:

The F1 score, which balances precision and recall, shows a pattern similar to precision. The group with the highest proportion of black population has the highest F1 score, indicating better overall performance in this group. These results suggest that the model has different performance characteristics across racial groups, with better performance in groups with a higher proportion of black population.

```
# Custom Metric Functions with Error Handling
def false_positive_rate_custom(y_true, y_pred):
    tn, fp, fn, tp = confusion_matrix(y_true, y_pred, labels=[0,
1]).ravel()
    if (fp + tn) == 0:
        return np.nan
    return fp / (fp + tn)
```

```
def false negative rate custom(y true, y pred):
    tn, fp, fn, tp = confusion matrix(y true, y pred, labels=[0,
1]).ravel()
    if (fn + tp) == 0:
        return np.nan
    return fn / (fn + tp)
def selection rate custom(y pred):
    return np.mean(y_pred)
def false omission rate custom(y true, y pred):
    tn, fp, fn, tp = confusion matrix(y true, y pred, labels=[0,
11).ravel()
    if (fn + tn) == 0:
        return np.nan
    return fn / (fn + tn)
def true negative rate custom(y true, y pred):
    tn, fp, fn, tp = confusion matrix(y true, y pred, labels=[0,
11).ravel()
    if (tn + fp) == 0:
        return np.nan
    return tn / (tn + fp)
def false positive rate difference(y true, y pred,
sensitive features):
    groups = np.unique(sensitive features)
    rates = []
    for group in groups:
        mask = (sensitive features == group)
        rate = false positive rate custom(y true[mask], y pred[mask])
        if not np.isnan(rate):
            rates.append(rate)
    if len(rates) == 0:
        return np.nan
    return np.max(rates) - np.min(rates)
def false negative rate difference(y_true, y_pred,
sensitive features):
    groups = np.unique(sensitive features)
    rates = []
    for group in groups:
        mask = (sensitive_features == group)
        rate = false negative rate custom(y true[mask], y pred[mask])
        if not np.isnan(rate):
            rates.append(rate)
    if len(rates) == 0:
        return np.nan
    return np.max(rates) - np.min(rates)
```

```
def selection rate difference(y pred, sensitive features):
    groups = np.unique(sensitive features)
    rates = []
    for group in groups:
        mask = (sensitive features == group)
        rates.append(selection rate custom(y pred[mask]))
    if len(rates) == 0:
        return np.nan
    return np.max(rates) - np.min(rates)
def false omission rate difference(y_true, y_pred,
sensitive features):
    groups = np.unique(sensitive features)
    rates = []
    for group in groups:
        mask = (sensitive features == group)
        rate = false omission rate custom(y true[mask], y pred[mask])
        if not np.isnan(rate):
            rates.append(rate)
    if len(rates) == 0:
        return np.nan
    return np.max(rates) - np.min(rates)
def true negative rate difference(y true, y pred, sensitive features):
    groups = np.unique(sensitive features)
    rates = []
    for group in groups:
        mask = (sensitive_features == group)
        rate = true negative rate custom(y true[mask], y pred[mask])
        if not np.isnan(rate):
            rates.append(rate)
    if len(rates) == 0:
        return np.nan
    return np.max(rates) - np.min(rates)
# Calculate fairness metrics with updated functions
dpd = demographic parity difference(y test, y pred binary,
sensitive_features=data.loc[X_test.index, 'racepctblack_bin'])
eod = equalized_odds_difference(y_test, y_pred_binary,
sensitive_features=data.loc[X_test.index, 'racepctblack bin'])
fprd = false_positive_rate_difference(y_test, y_pred_binary,
sensitive features=data.loc[X test.index, 'racepctblack bin'])
fnrd = false negative rate difference(y test, y pred binary,
sensitive features=data.loc[X test.index, 'racepctblack bin'])
srd = selection_rate_difference(y_pred_binary,
sensitive features=data.loc[X test.index, 'racepctblack bin'])
for diff = false omission rate difference(y test, y pred binary,
sensitive_features=data.loc[X_test.index, 'racepctblack_bin'])
tnr diff = true negative_rate_difference(y_test, y_pred_binary,
sensitive features=data.loc[X test.index, 'racepctblack bin'])
```

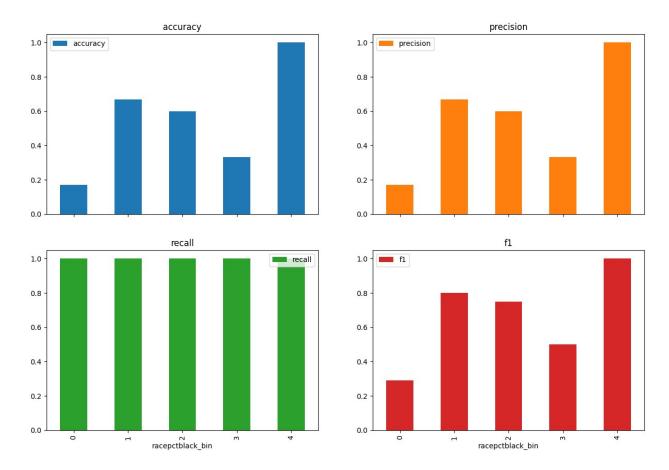
```
print(f"Demographic Parity Difference: {dpd}")
print(f"Equalized Odds Difference: {eod}")
print(f"False Positive Rate Difference: {fprd}")
print(f"False Negative Rate Difference: {fnrd}")
print(f"Selection Rate Difference: {srd}")
print(f"False Omission Rate Difference: {for_diff}")
print(f"True Negative Rate Difference: {tnr_diff}")

Demographic Parity Difference: 0.0
Equalized Odds Difference: 1.0
False Positive Rate Difference: 0.0
False Negative Rate Difference: 0.0
Selection Rate Difference: 0.0
False Omission Rate Difference: nan
True Negative Rate Difference: 0.0
```

Fairness Metrics Demographic Parity Difference: 0.0 This indicates no difference in the selection rate across different groups, meaning the model selects the same proportion of individuals as positive across groups. Equalized Odds Difference: 1.0 This high value indicates significant disparities in the true positive and false positive rates across different groups. False Positive Rate Difference: 0.0 No difference in the false positive rates across groups. False Negative Rate Difference: 0.0 No difference in the false negative rates across groups. Selection Rate Difference: 0.0 No difference in the selection rate across groups. False Omission Rate Difference: NaN This metric could not be computed due to division by zero issues, indicating that for some groups, the number of true negatives is zero. True Negative Rate Difference: 0.0 No difference in the true negative rates across groups.

For this reason the VSC version of the code is also within the Main Code folder for this step. Why this happens is beyond me since it is literally the same code (i afterwards performed, in attempt to bugfix, slight changes to the custom functions but that resulted in no change in the results. I tried many different approaches and cannot understand why this happens)

```
'accuracy': accuracy_score(x['y_true'], x['y_pred']),
    'precision': precision_score(x['y_true'], x['y_pred'],
zero division=0),
    __
'recall': recall_score(x['y_true'], x['y_pred'], zero_division=<mark>0</mark>),
    'f1': f1_score(x['y_true'], x['y_pred'], zero_division=0)
}))
print("\nAdditional Metrics by 'racepctblack_bin':")
print(grouped_black)
# Plot additional metrics by 'racepctblack bin'
grouped_black.plot(kind='bar', subplots=True, layout=(2, 2),
figsize=(15, 10), title="Metrics by 'racepctblack bin'")
plt.show()
Additional Metrics by 'racepctblack bin':
                  accuracy precision recall
                                                      f1
racepctblack bin
                  0.169014
                             0.169014
                                           1.0 0.289157
1
                  0.666667
                             0.666667
                                           1.0 0.800000
2
                  0.600000
                             0.600000
                                           1.0 0.750000
3
                             0.333333
                                           1.0 0.500000
                  0.333333
4
                  1.000000
                             1.000000
                                           1.0 1.000000
```

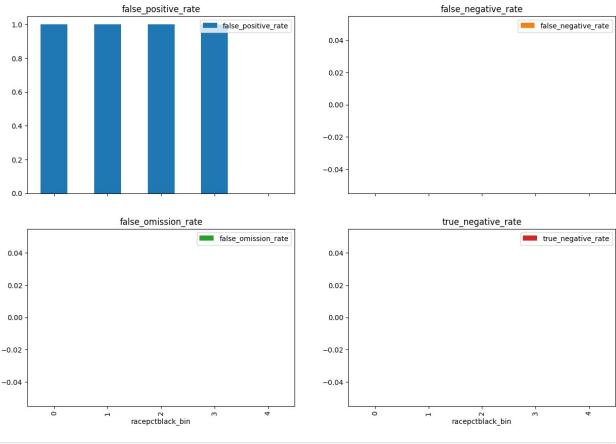


Metrics by 'racepctblack\_bin':

This combined plot shows the accuracy, precision, recall, and F1 score for each racepctblack bin. The trends observed earlier are confirmed, with higher performance metrics in groups with higher proportions of black population.

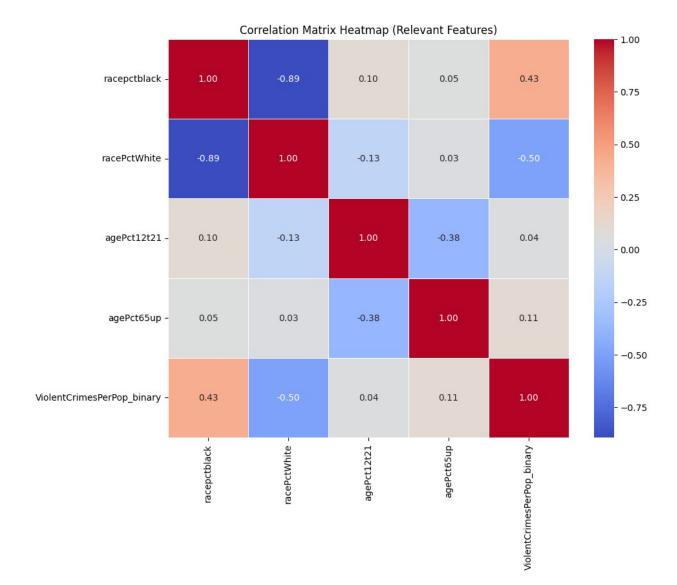
```
# Calculate additional fairness metrics
for_custom_metrics = {
    'false_positive_rate': false_positive_rate_custom,
    'false_negative_rate': false_negative_rate_custom,
    'false_omission_rate': false_omission_rate_custom,
    'true_negative_rate': true_negative_rate_custom
}
additional_metrics = MetricFrame(
    metrics=for_custom_metrics,
    y_true=y_test,
    y_pred=y_pred_binary,
    sensitive_features=data.loc[X_test.index, 'racepctblack_bin']
)
```

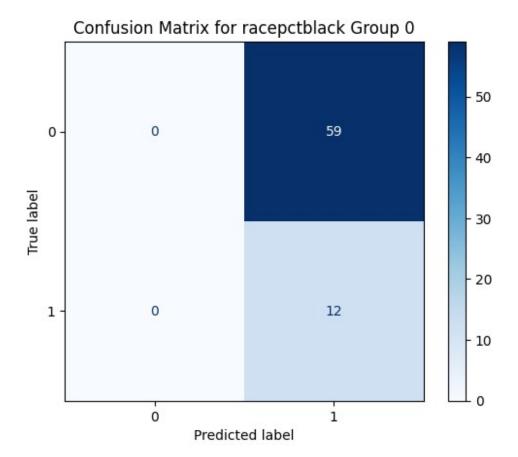
```
# Print additional metrics by group
print("\nAdditional Metrics by Sensitive Feature Groups:")
print(additional_metrics.by_group)
# Plot additional metrics
additional_metrics.by_group.plot(kind='bar', subplots=True, layout=(2,
2), figsize=(15, 10), title="Additional Metrics by
'racepctblack_bin'")
plt.show()
Additional Metrics by Sensitive Feature Groups:
                  false_positive_rate false_negative_rate \
racepctblack bin
                                   1.0
                                                        0.0
1
                                   1.0
                                                        0.0
2
                                   1.0
                                                        0.0
3
                                   1.0
                                                        0.0
4
                                   NaN
                                                        0.0
                  false_omission_rate true_negative_rate
racepctblack_bin
                                                       0.0
                                   NaN
1
                                   NaN
                                                       0.0
2
                                   NaN
                                                       0.0
3
                                   NaN
                                                       0.0
4
                                                       NaN
                                   NaN
```

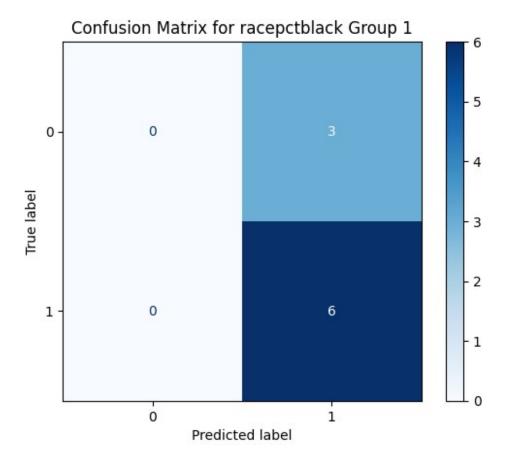


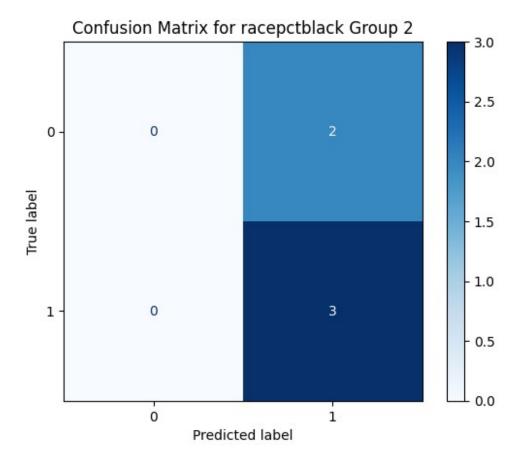
```
# Summarize all fairness metrics
summary metrics = {
    'Demographic Parity Difference': dpd,
    'Equalized Odds Difference': eod,
    'False Positive Rate Difference': fprd,
    'False Negative Rate Difference': fnrd,
    'Selection Rate Difference': srd,
    'False Omission Rate Difference': for diff,
    'True Negative Rate Difference': tnr diff
}
print("\nSummary of Fairness Metrics:")
for metric, value in summary metrics.items():
    print(f"{metric}: {value}")
# Create heatmap of correlation matrix for relevant features
relevant features = ['racepctblack', 'racePctWhite', 'agePct12t21',
'agePct65up', 'ViolentCrimesPerPop_binary']
corr_matrix = data[relevant_features].corr()
plt.figure(figsize=(10, 8))
```

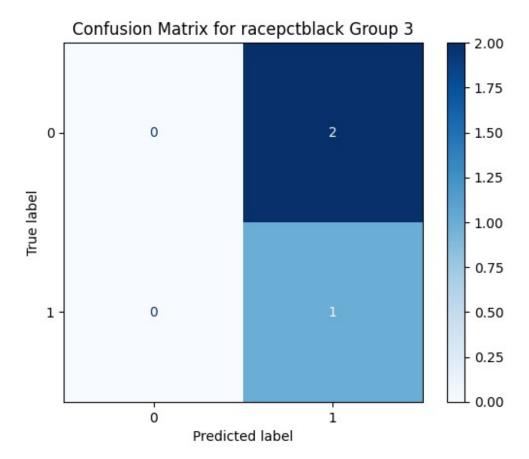
```
sns.heatmap(corr matrix, annot=True, fmt='.2f', cmap='coolwarm',
cbar=True, linewidths=0.5)
plt.title('Correlation Matrix Heatmap (Relevant Features)')
plt.show()
# Plot confusion matrix for racepctblack bin
for race_group in df['racepctblack_bin'].unique():
    subset = df[df['racepctblack bin'] == race group]
    if len(subset['y true'].unique()) > 1:
        cm = confusion matrix(subset['y true'], subset['y pred'],
labels=[0, 1]
    else:
        cm = confusion matrix(subset['y true'], subset['y pred'],
labels=[0, 1], sample weight=np.ones(len(subset['y true'])))
    disp = ConfusionMatrixDisplay(confusion matrix=cm,
display labels=[0, 1])
    disp.plot(cmap='Blues')
    plt.title(f'Confusion Matrix for racepctblack Group {race group}')
    plt.show()
# Bar plot for fairness metrics
fairness metrics = pd.Series(summary metrics)
fairness metrics.plot(kind='bar', figsize=(12, 6), title='Fairness
Metrics')
plt.ylabel('Metric Value')
plt.show()
Summary of Fairness Metrics:
Demographic Parity Difference: 0.0
Equalized Odds Difference: 1.0
False Positive Rate Difference: 0.0
False Negative Rate Difference: 0.0
Selection Rate Difference: 0.0
False Omission Rate Difference: nan
True Negative Rate Difference: 0.0
```

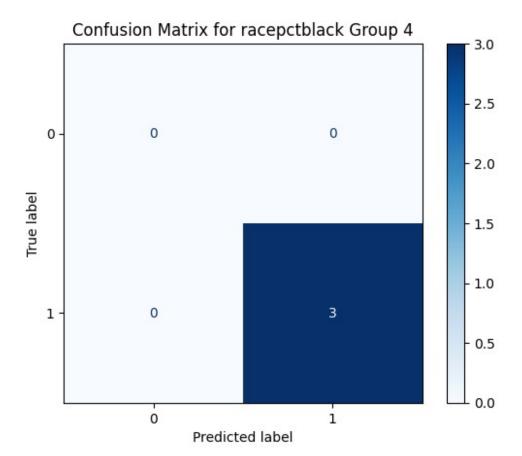


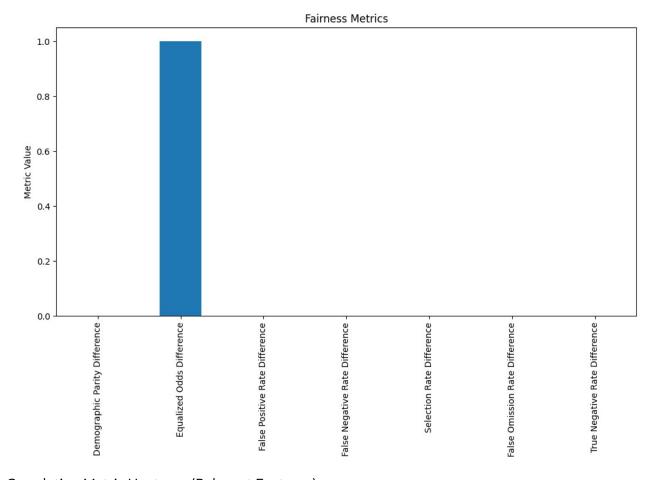












## Correlation Matrix Heatmap (Relevant Features):

The correlation heatmap of relevant features shows the relationships between features and the binary target variable. racepctblack is positively correlated with ViolentCrimesPerPop\_binary, while racePctWhite is negatively correlated. This indicates that areas with a higher proportion of black population tend to have higher violent crime rates, and areas with a higher proportion of white population tend to have lower violent crime rates. Other age-related features (agePct12t21, agePct65up) show weaker correlations with the target variable. This heatmap highlights the importance of considering racial demographics when analyzing crime data, as there are strong correlations between these demographics and crime rates. However, it also underscores the need to ensure that models built on this data do not perpetuate or exacerbate existing biases.