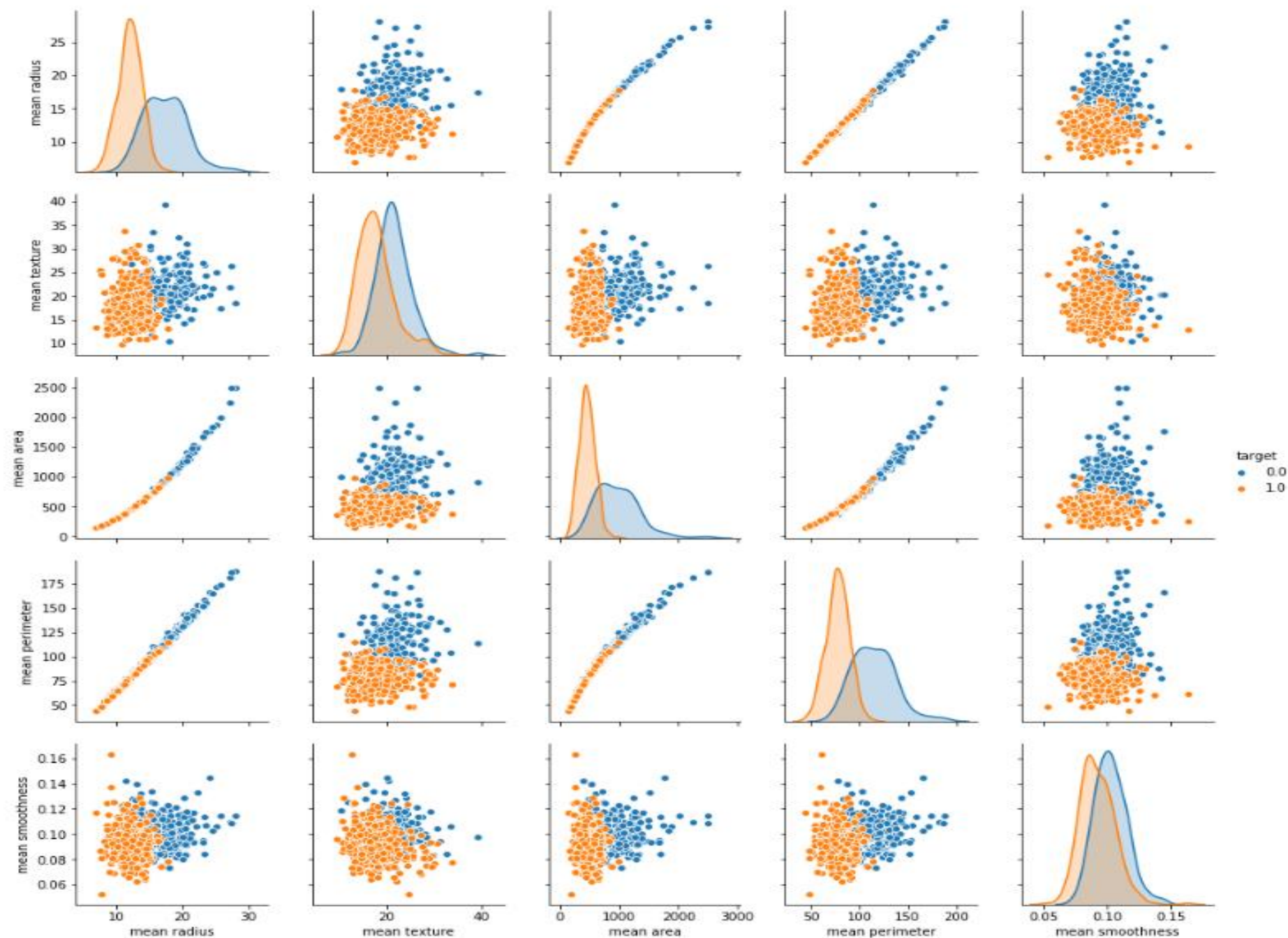
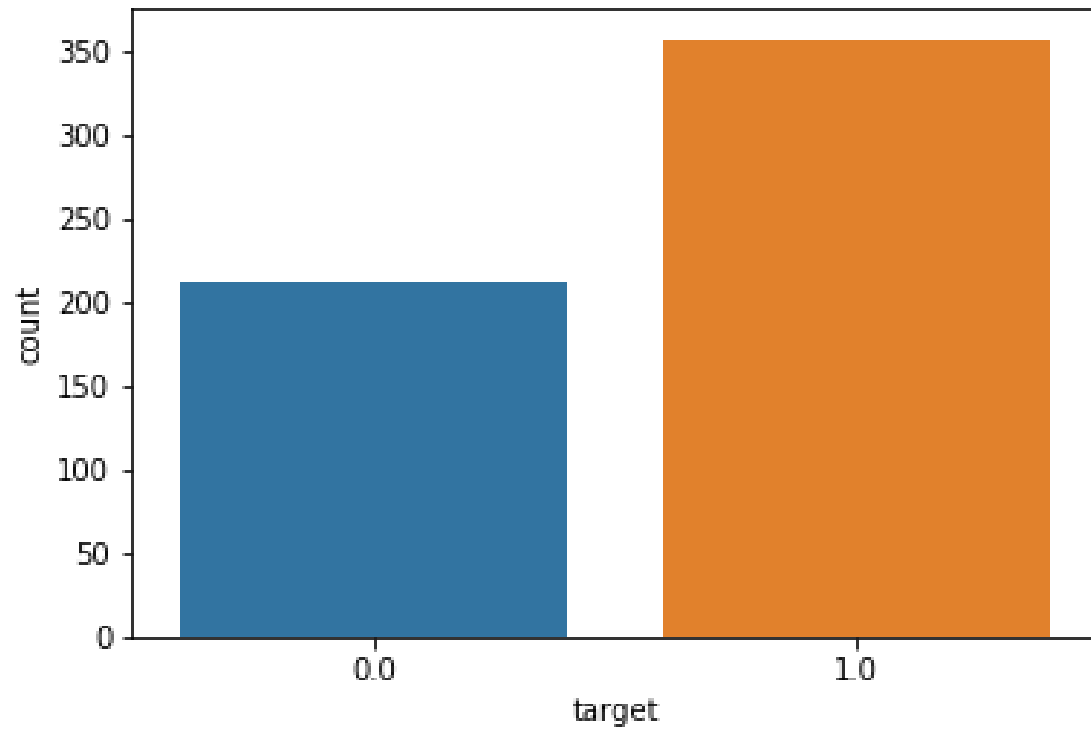


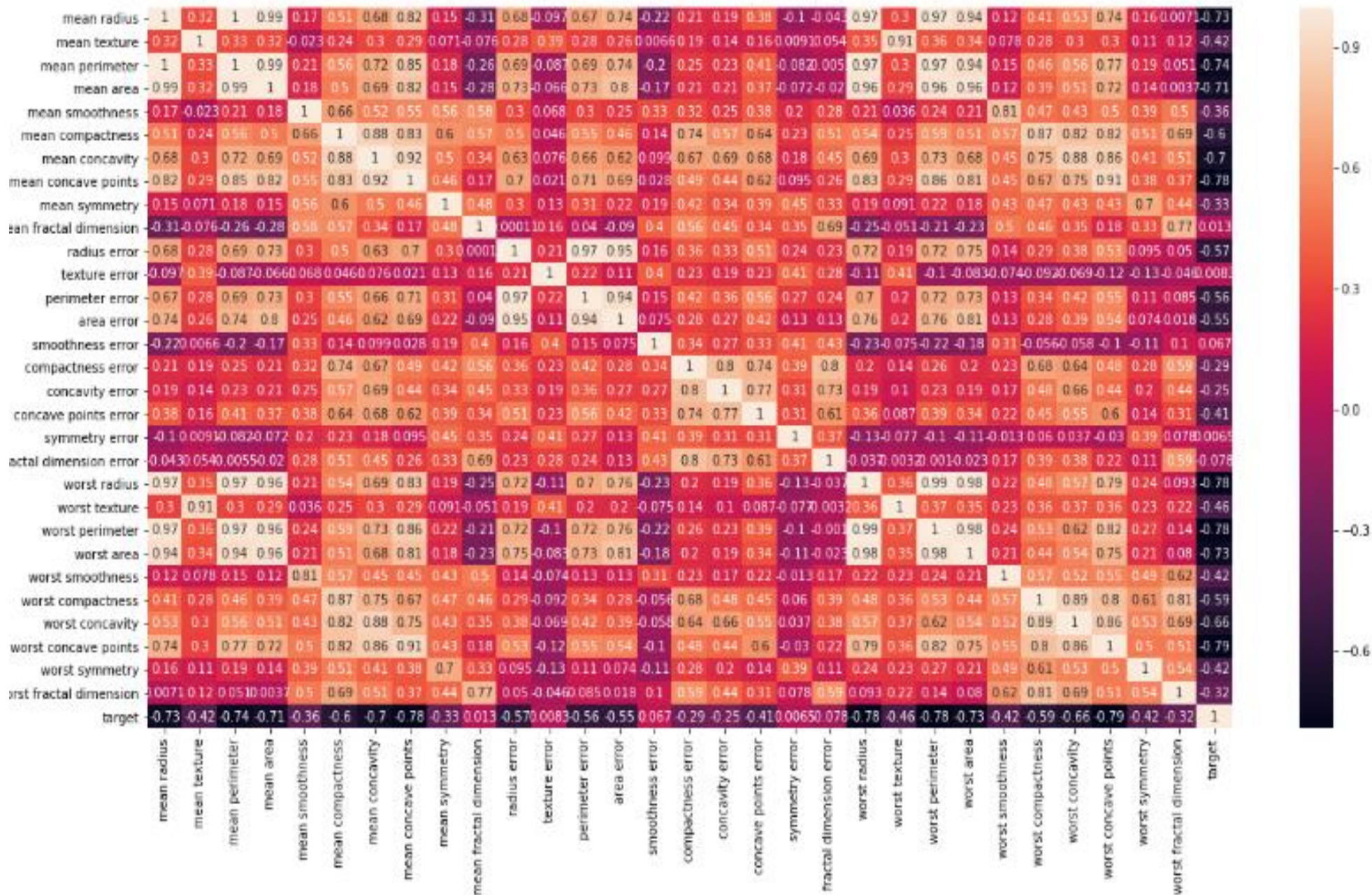
```
: sns.pairplot(df_cancer, hue = 'target', vars = ['mean radius', 'mean texture', 'mean area', 'mean perimeter', 'mean smoothness'])
```



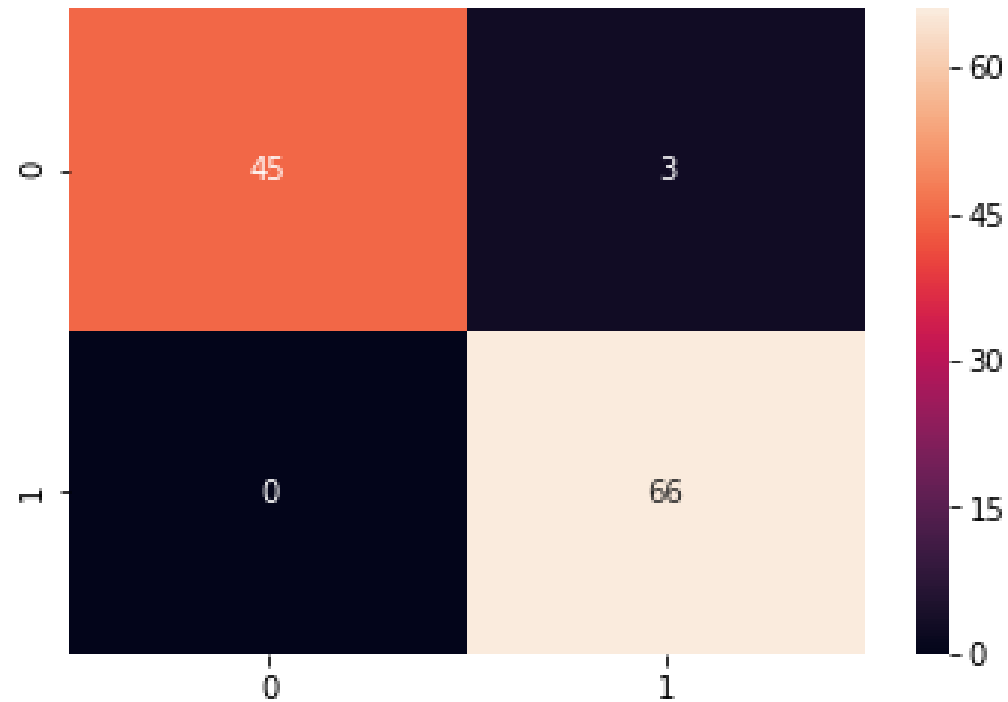
```
: sns.countplot(df_cancer['target'], label = "Count")
```




```
plt.figure(figsize=(20,10))
sns.heatmap(df_cancer.corr(), annot=True)
```



```
sns.heatmap(cm, annot=True)
```



```
: print(classification_report(y_test,grid_predictions))
```

| | precision | recall | f1-score | support |
|-------------|-----------|--------|----------|---------|
| 0.0 | 1.00 | 0.94 | 0.97 | 48 |
| 1.0 | 0.96 | 1.00 | 0.98 | 66 |
| avg / total | 0.97 | 0.97 | 0.97 | 114 |