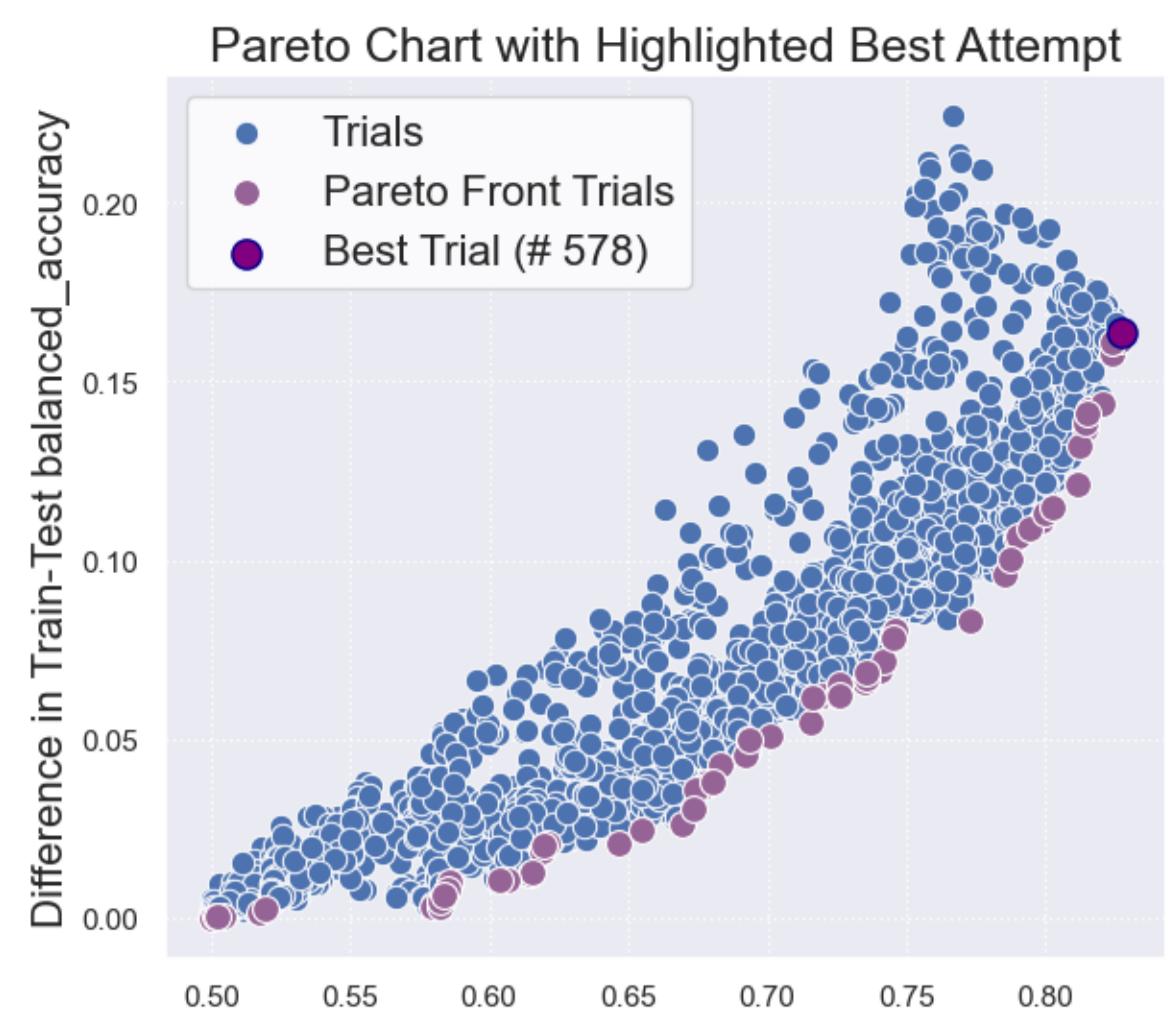
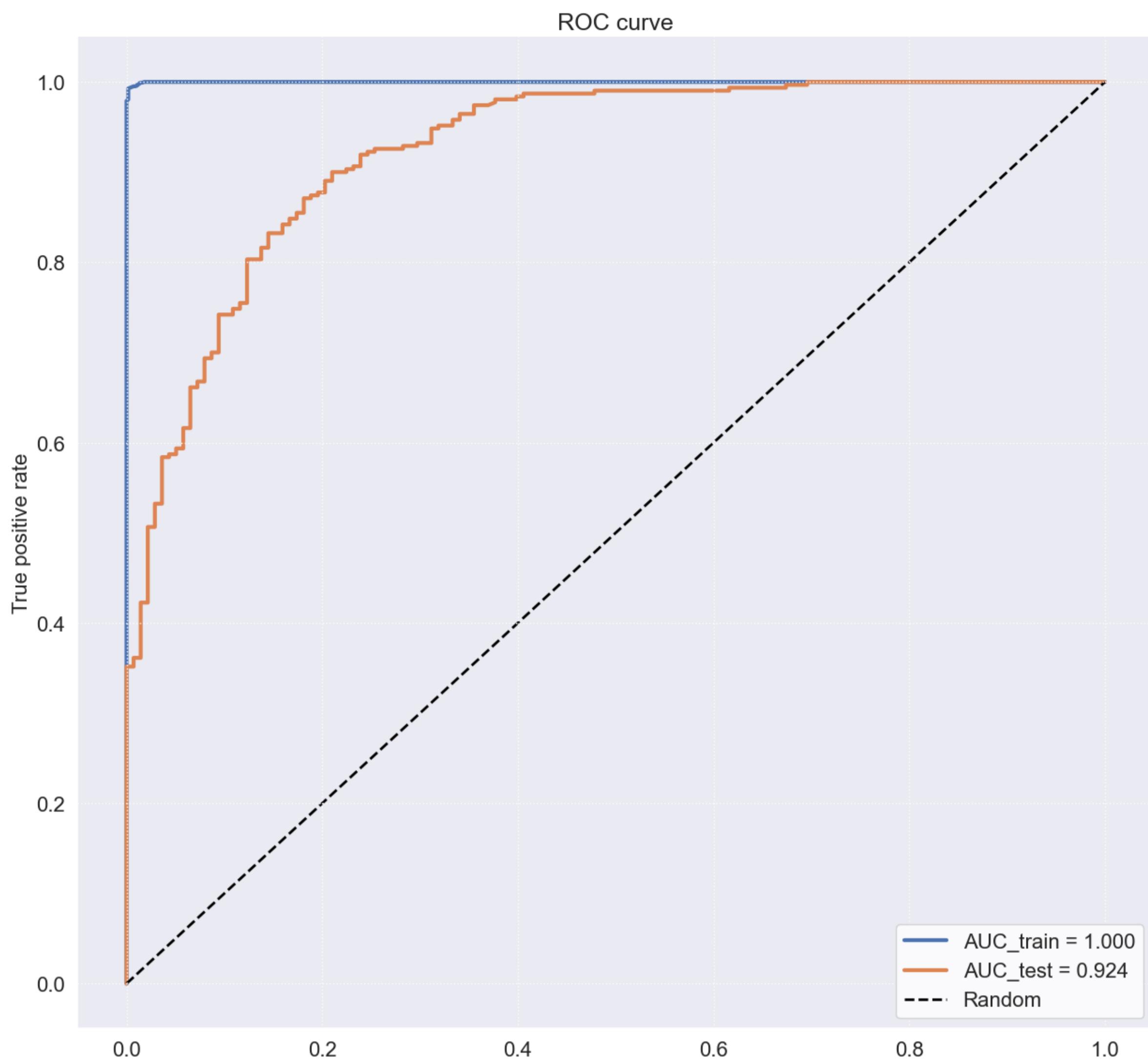
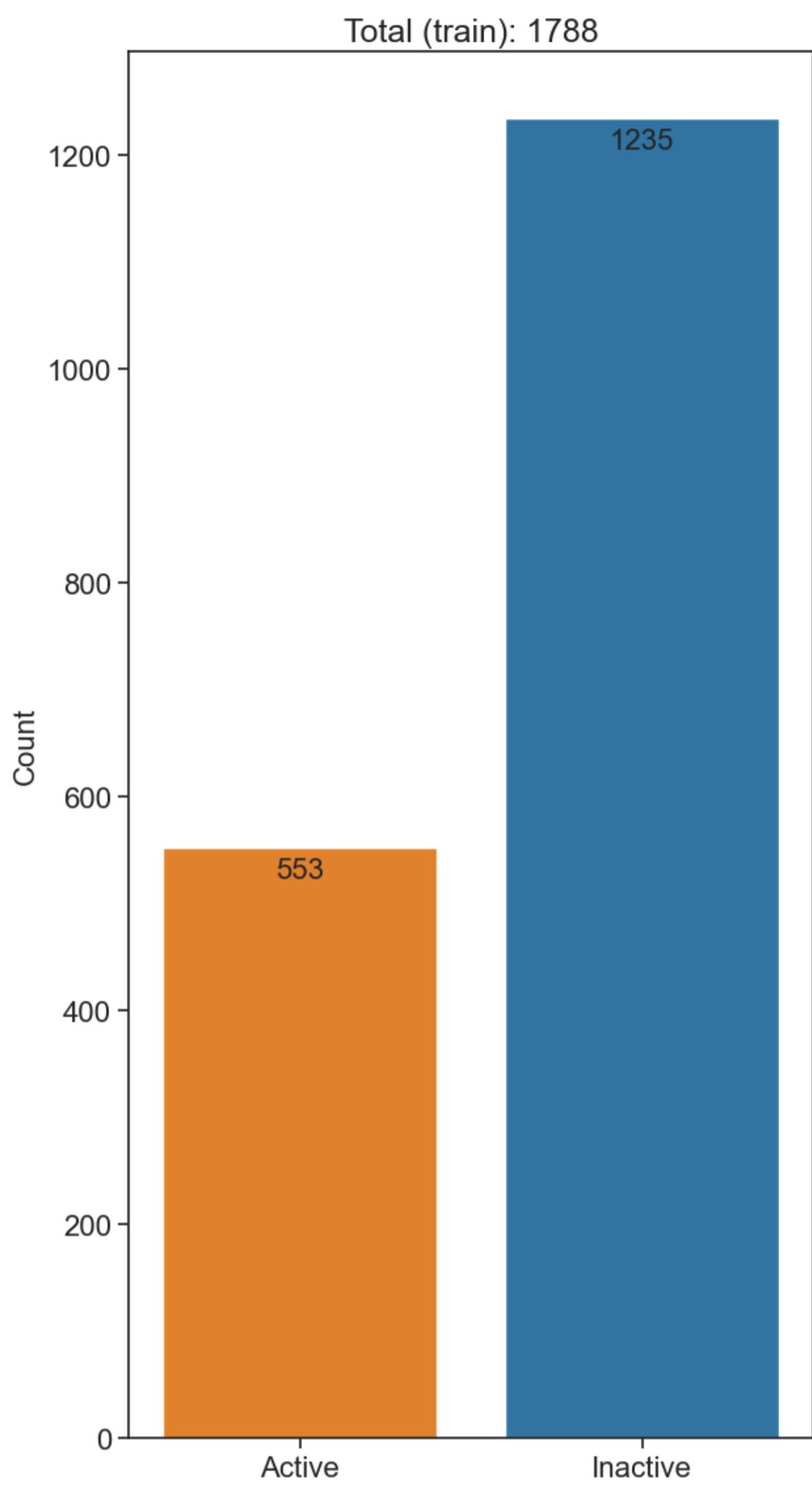
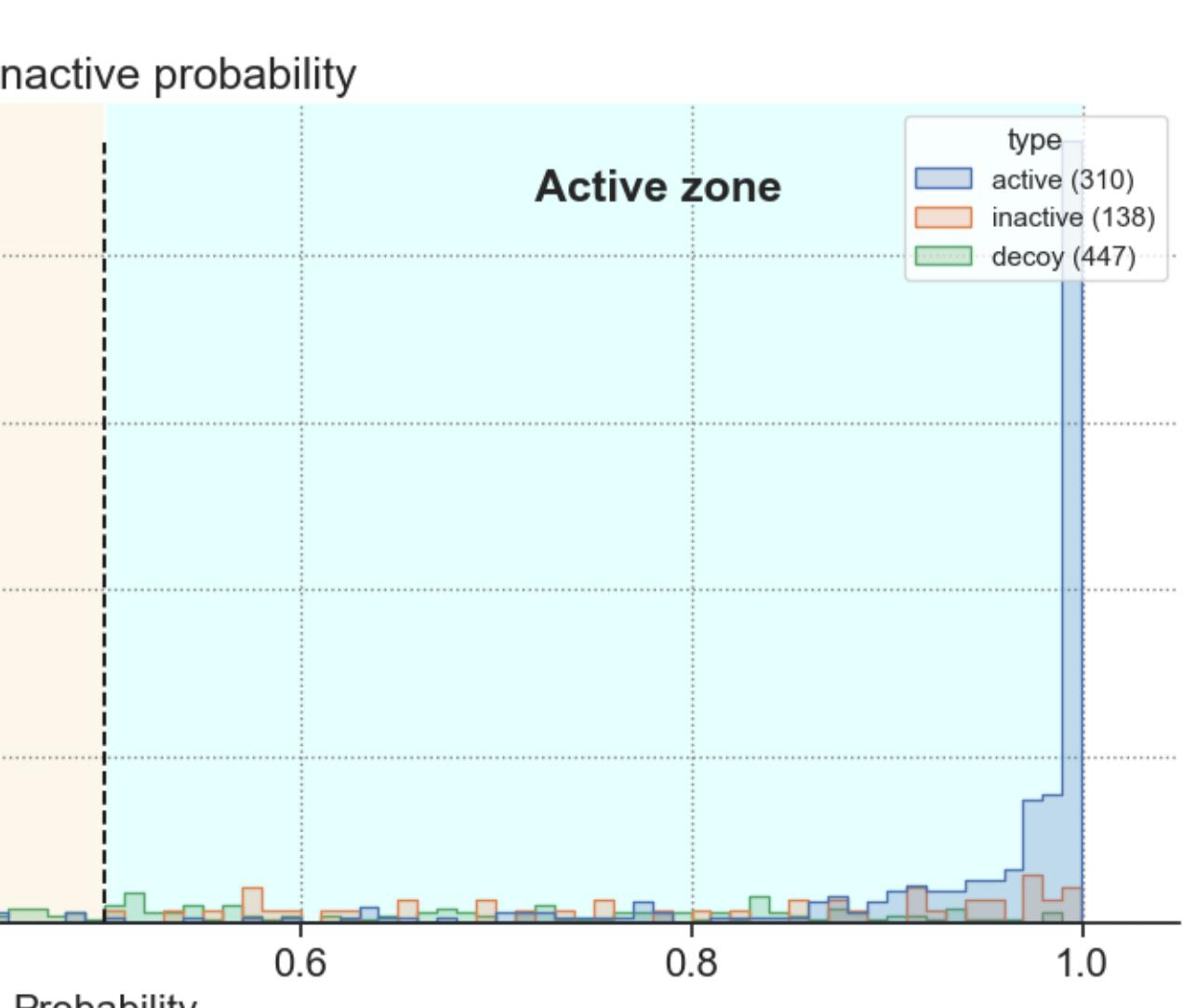
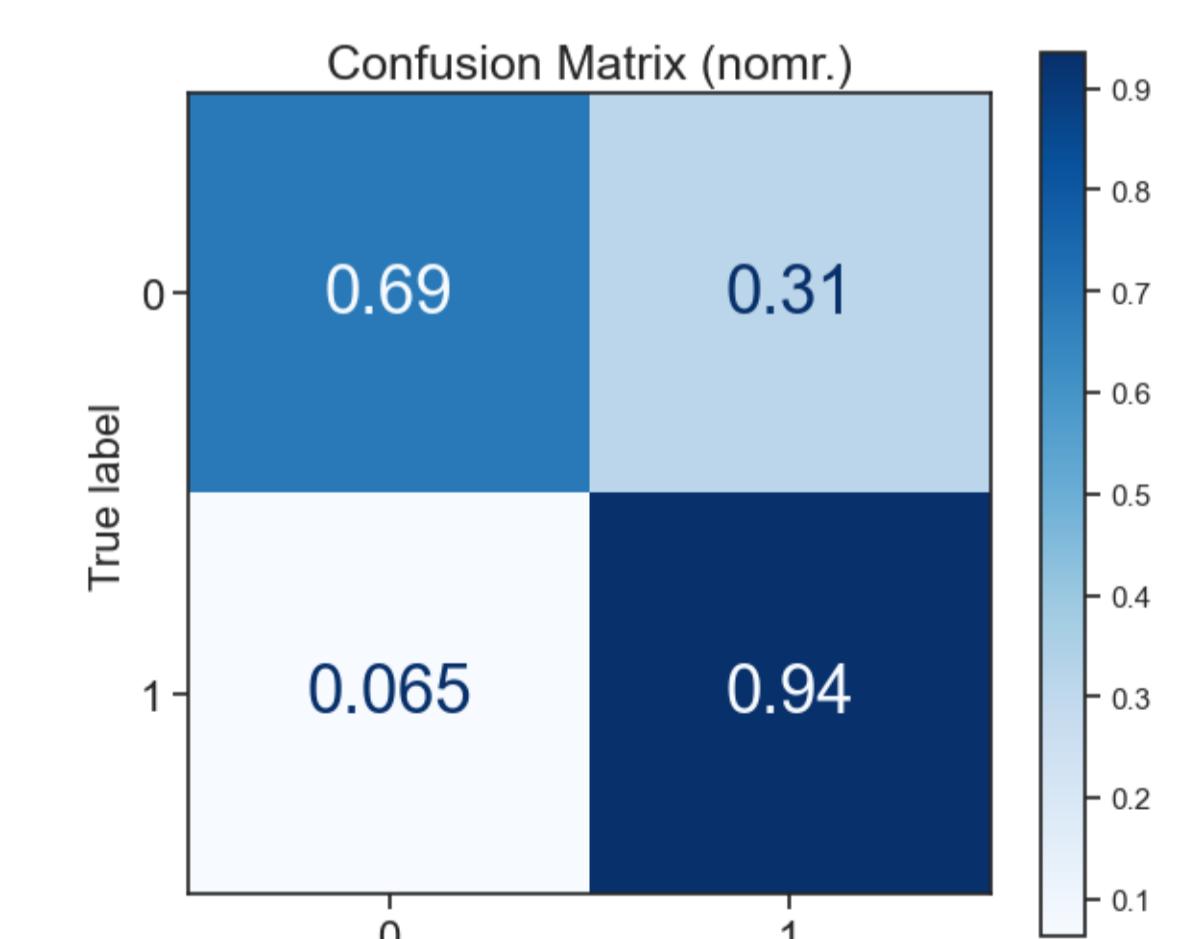
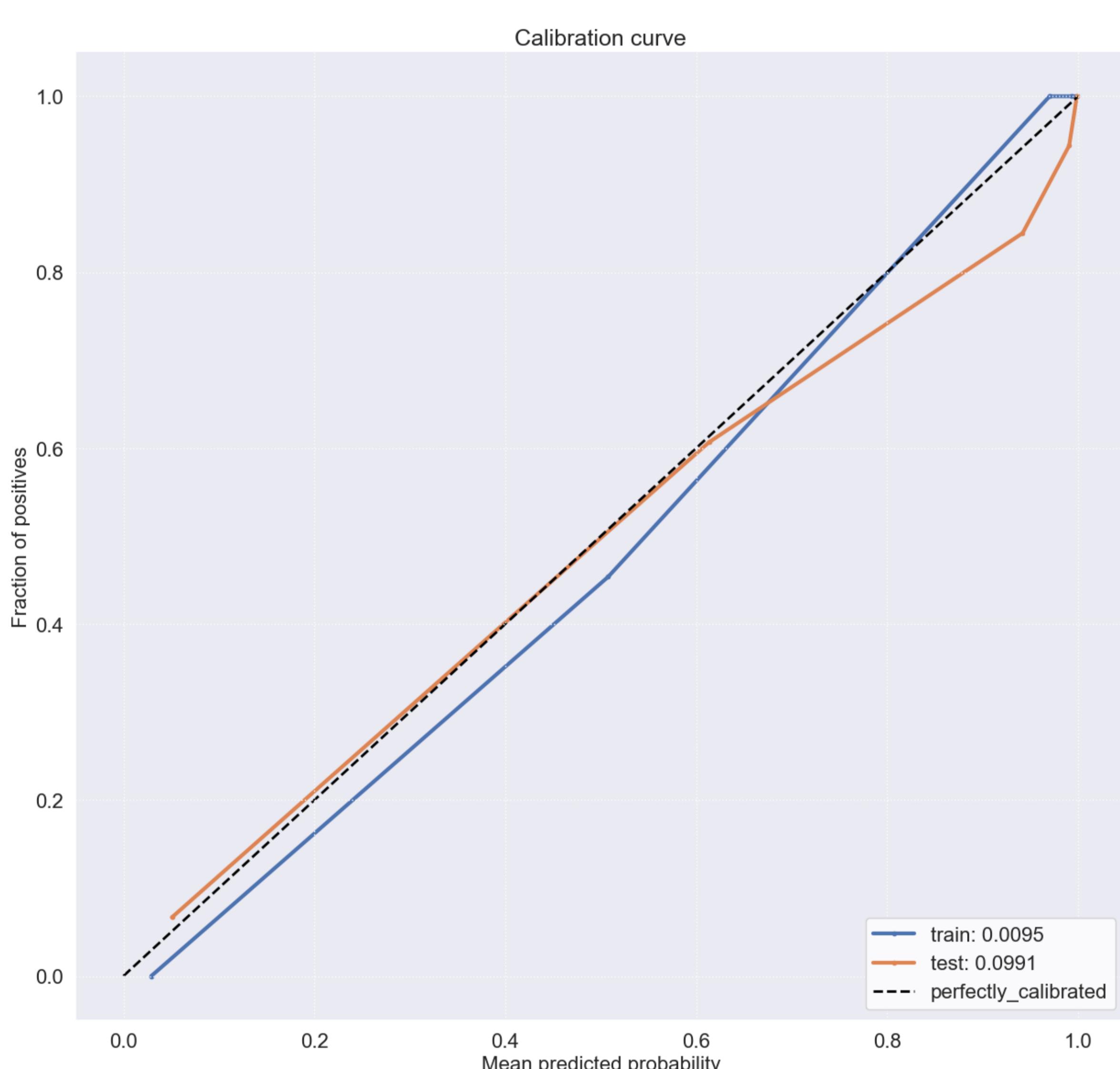
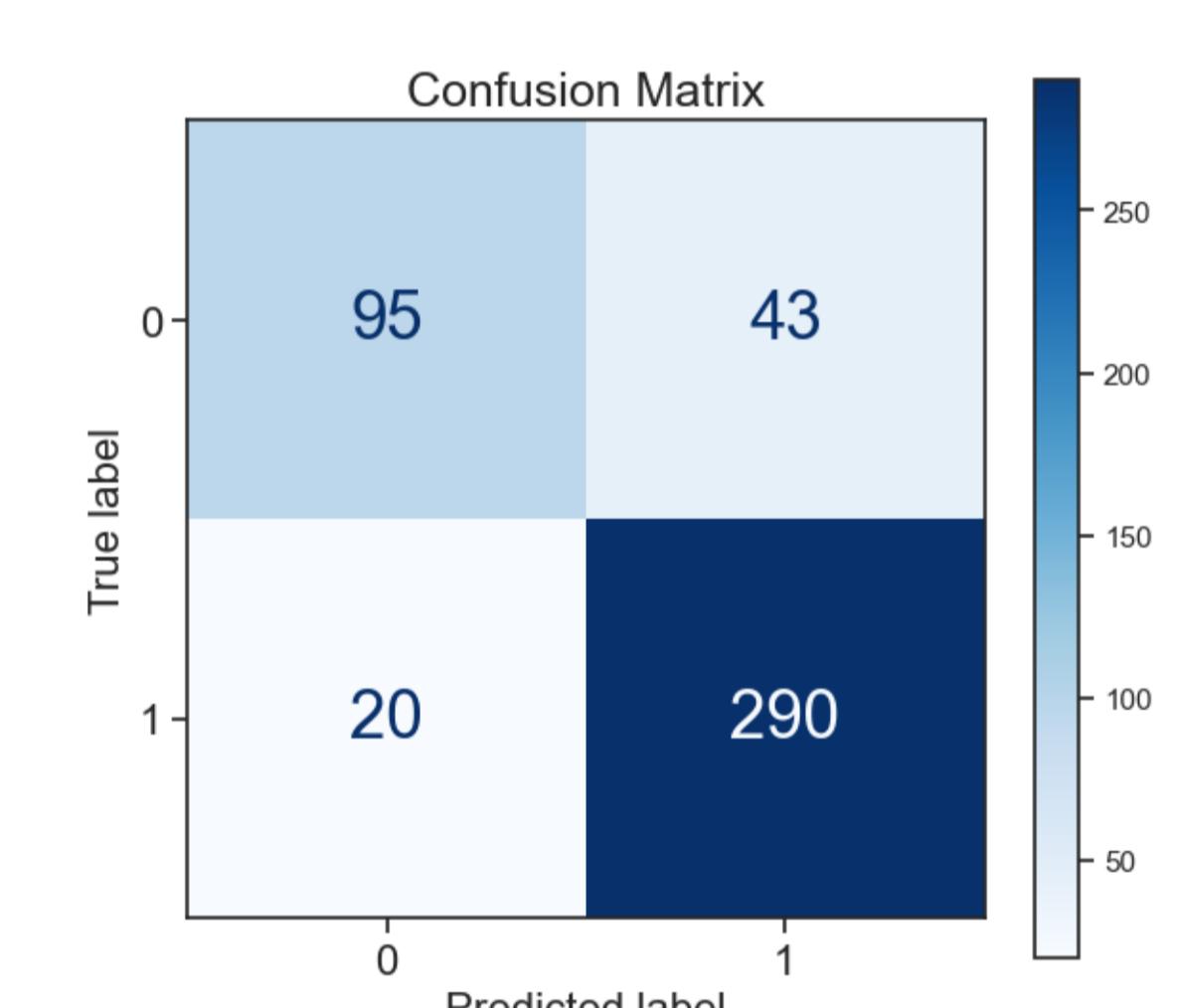


Protein: P28335 - metric: balanced_accuracy

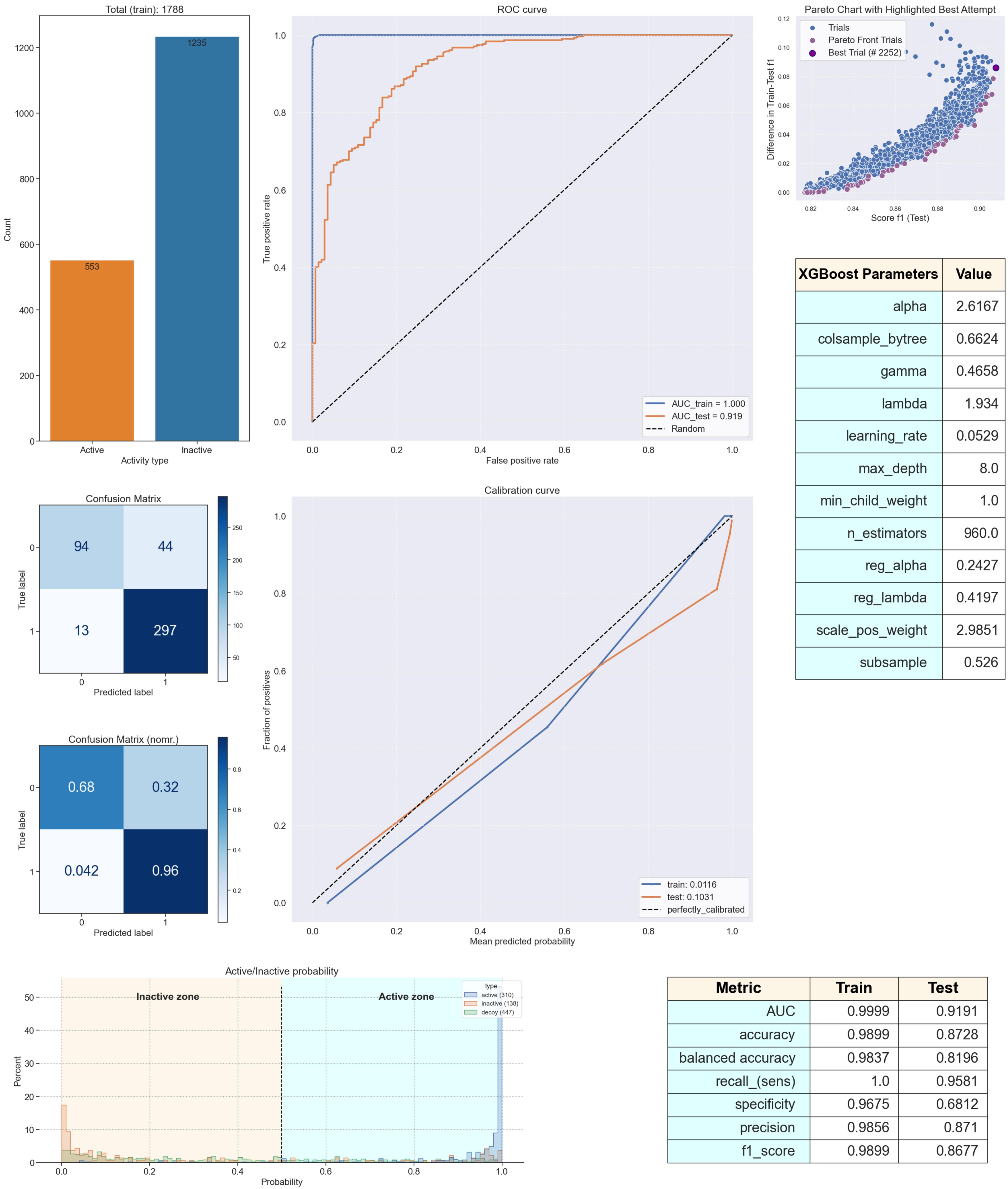


| XGBoost Parameters | Value |
|--------------------|--------|
| alpha | 4.0852 |
| colsample_bytree | 0.6915 |
| gamma | 0.3595 |
| lambda | 2.0192 |
| learning_rate | 0.0923 |
| max_depth | 9.0 |
| min_child_weight | 1.0 |
| n_estimators | 942.0 |
| reg_alpha | 0.4933 |
| reg_lambda | 0.3234 |
| scale_pos_weight | 1.4468 |
| subsample | 0.5909 |

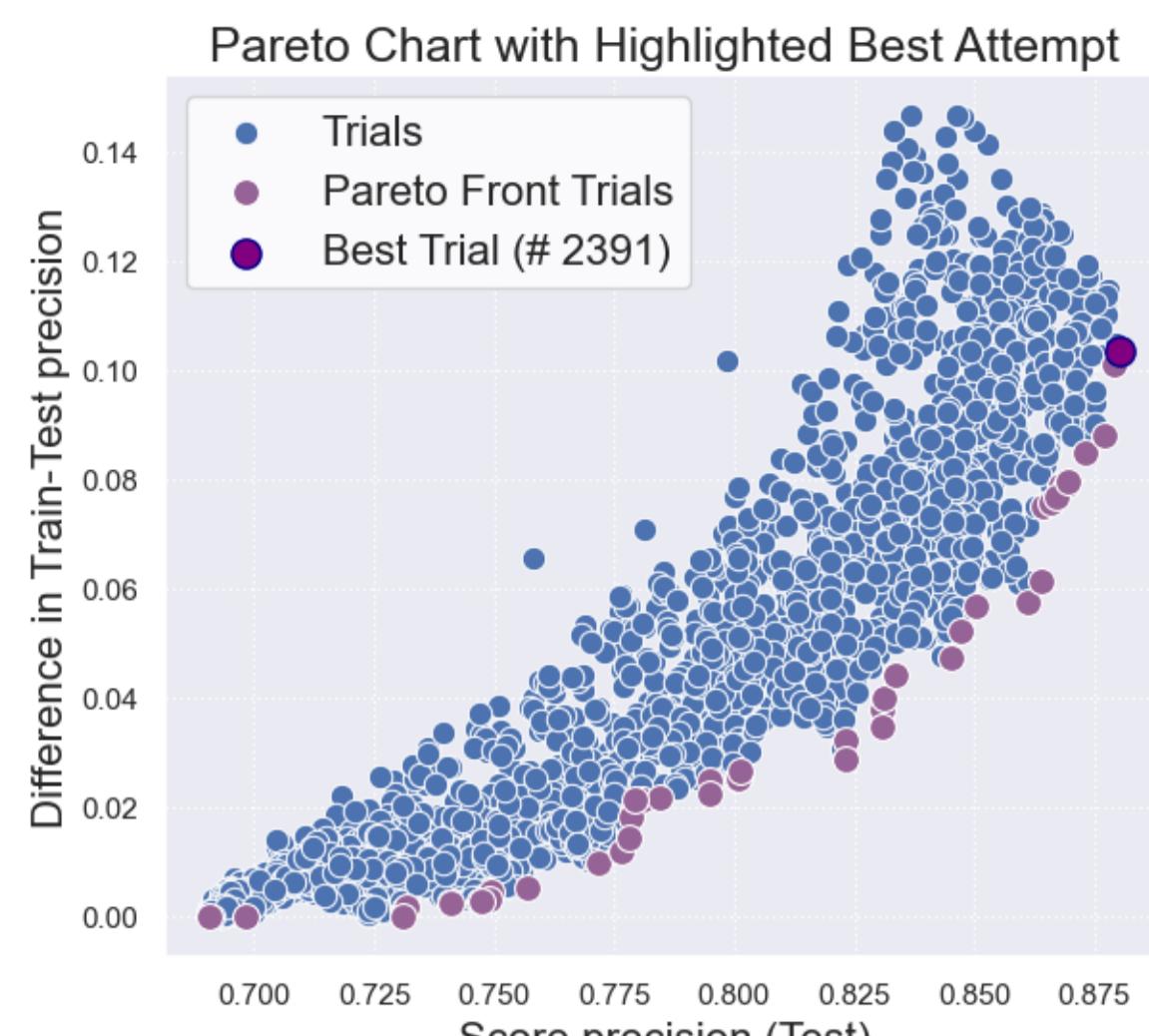
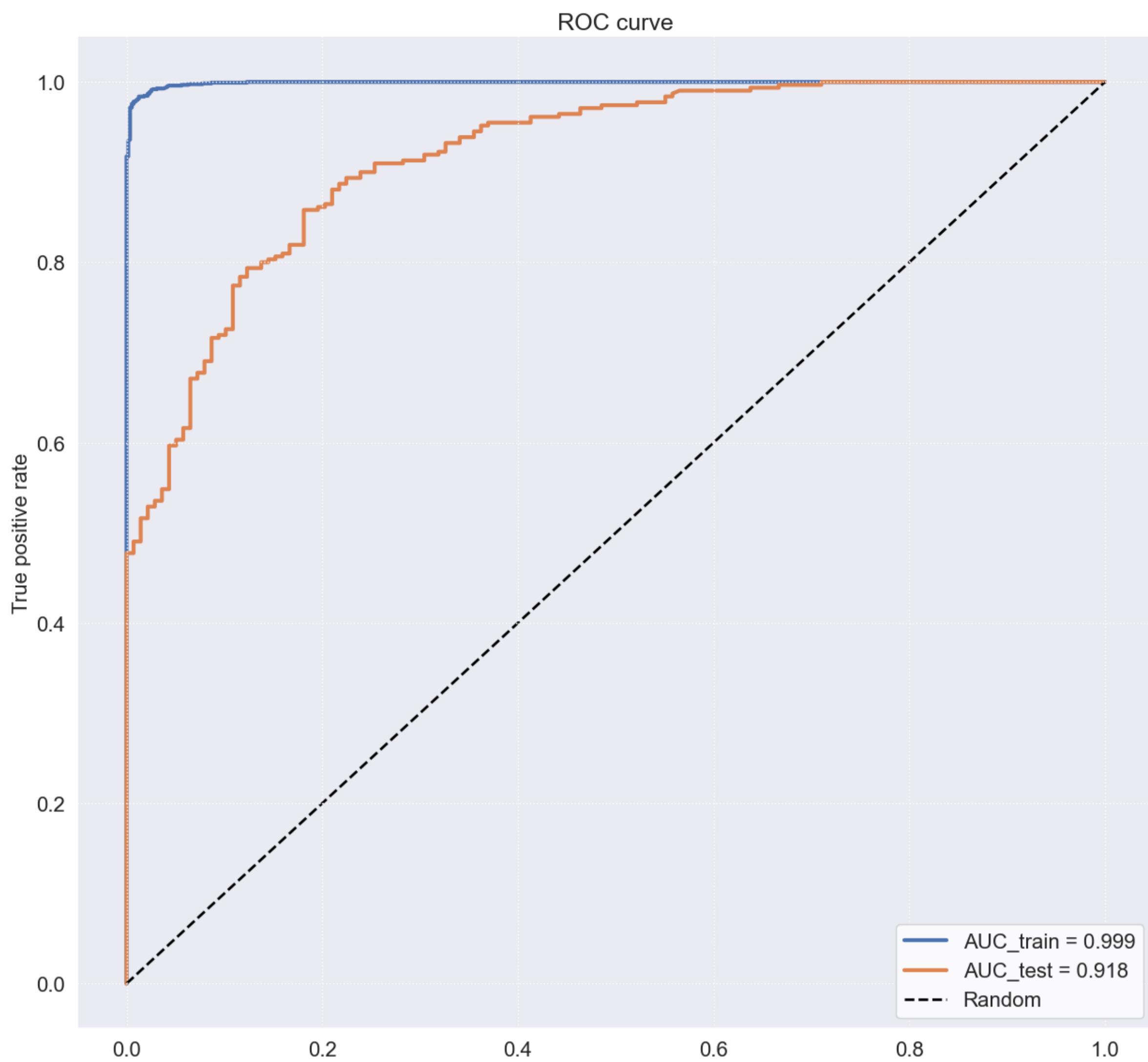
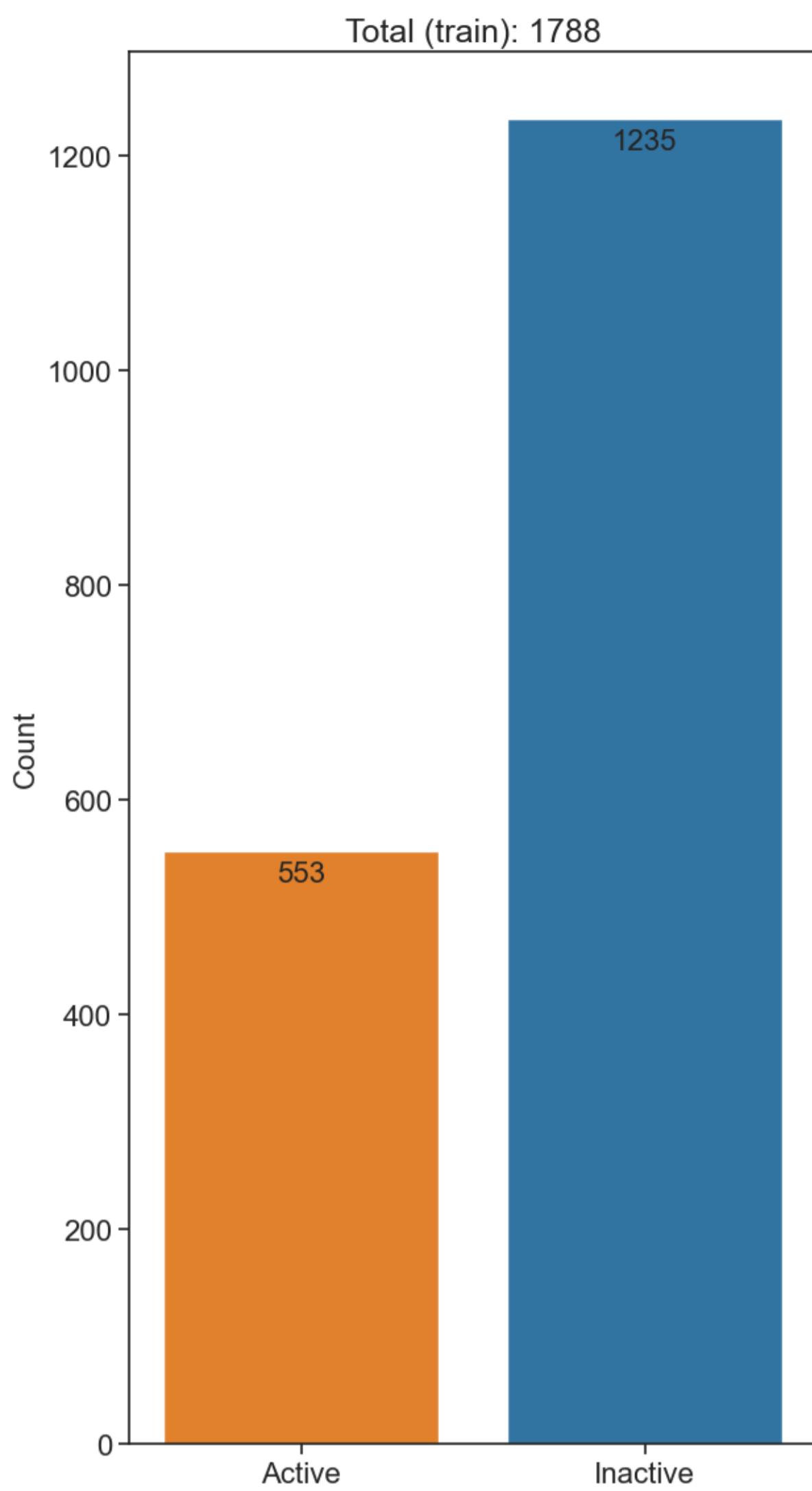


| Metric | Train | Test |
|-------------------|--------|--------|
| AUC | 0.9999 | 0.9239 |
| accuracy | 0.9938 | 0.8594 |
| balanced accuracy | 0.9901 | 0.8119 |
| recall_(sens) | 1.0 | 0.9355 |
| specificity | 0.9801 | 0.6884 |
| precision | 0.9912 | 0.8709 |
| f1_score | 0.9938 | 0.8555 |

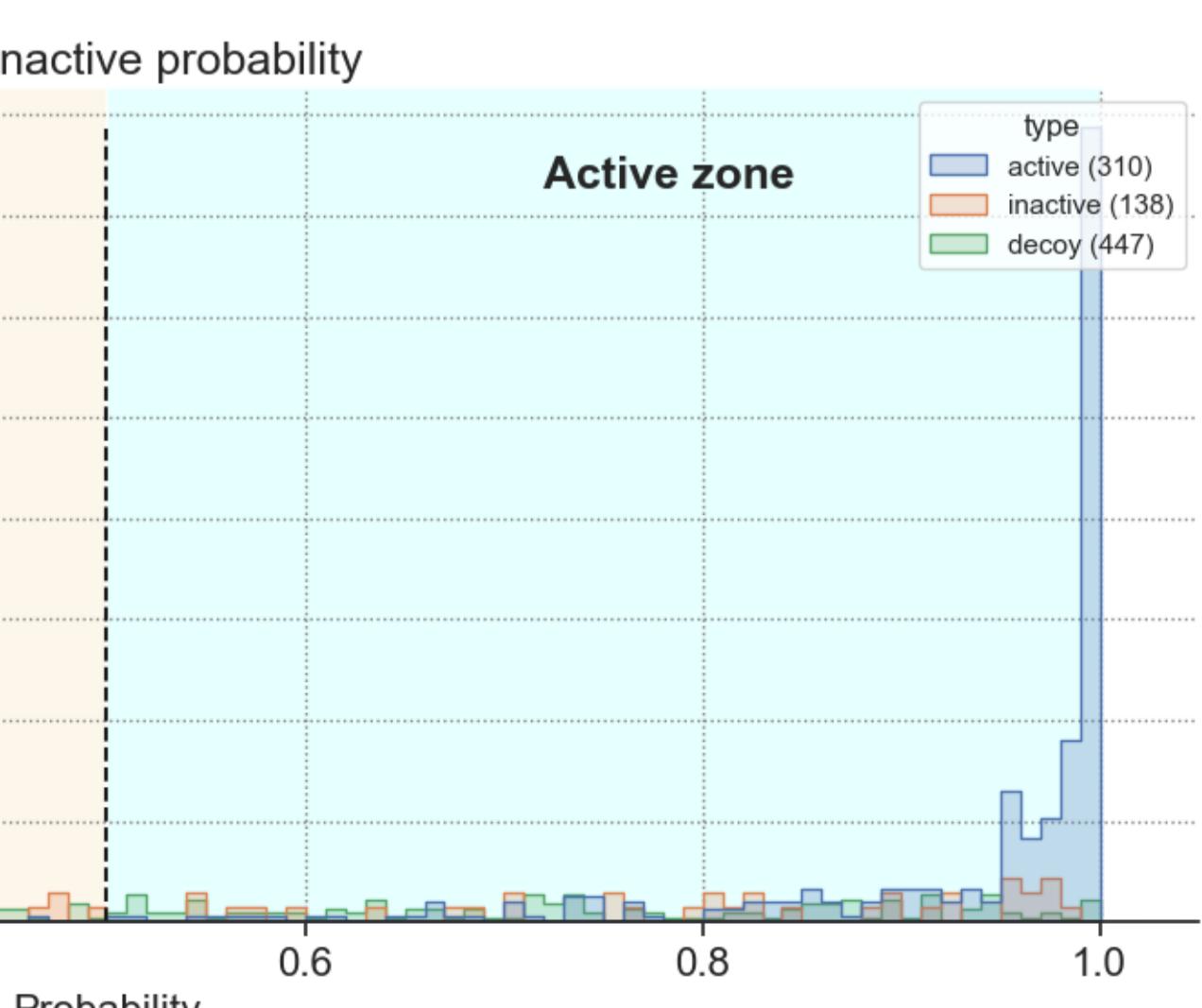
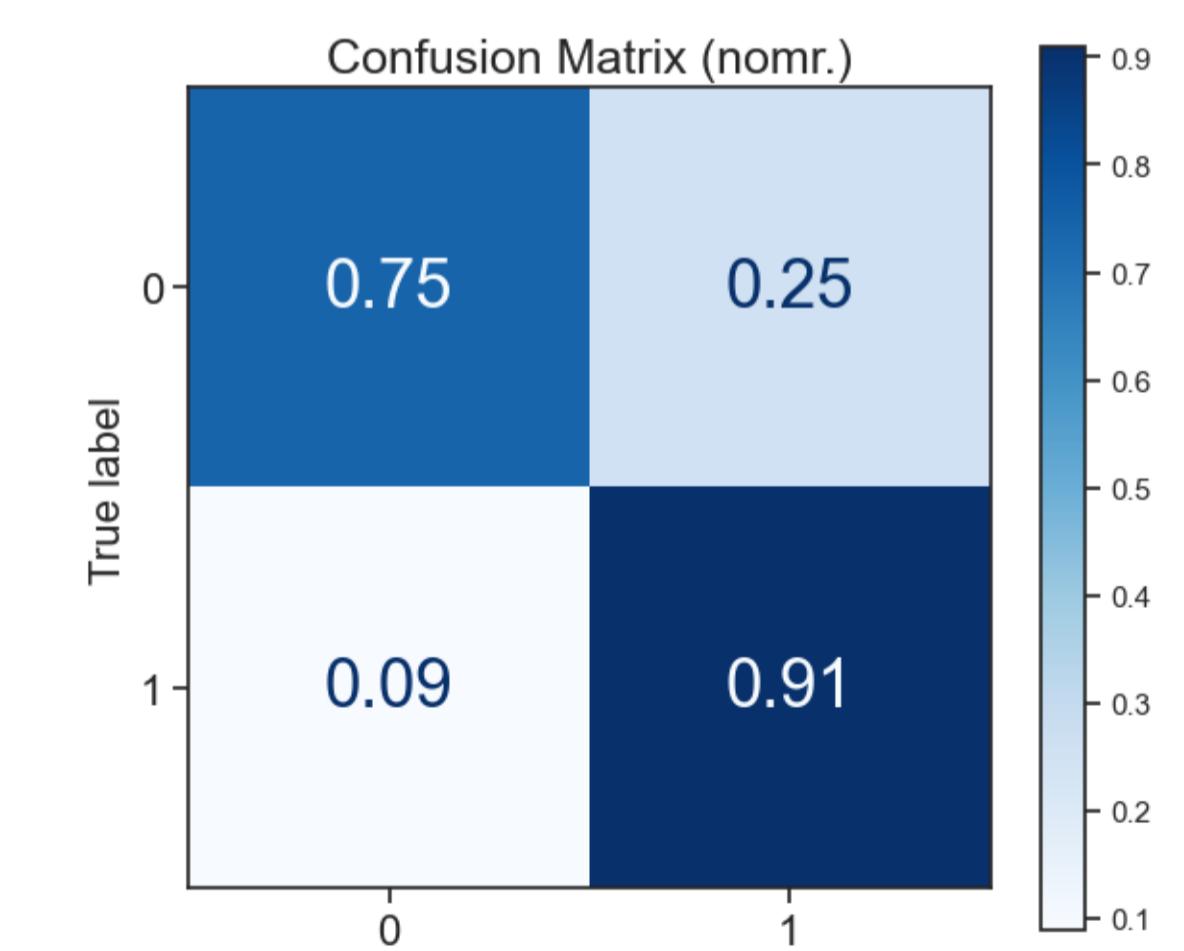
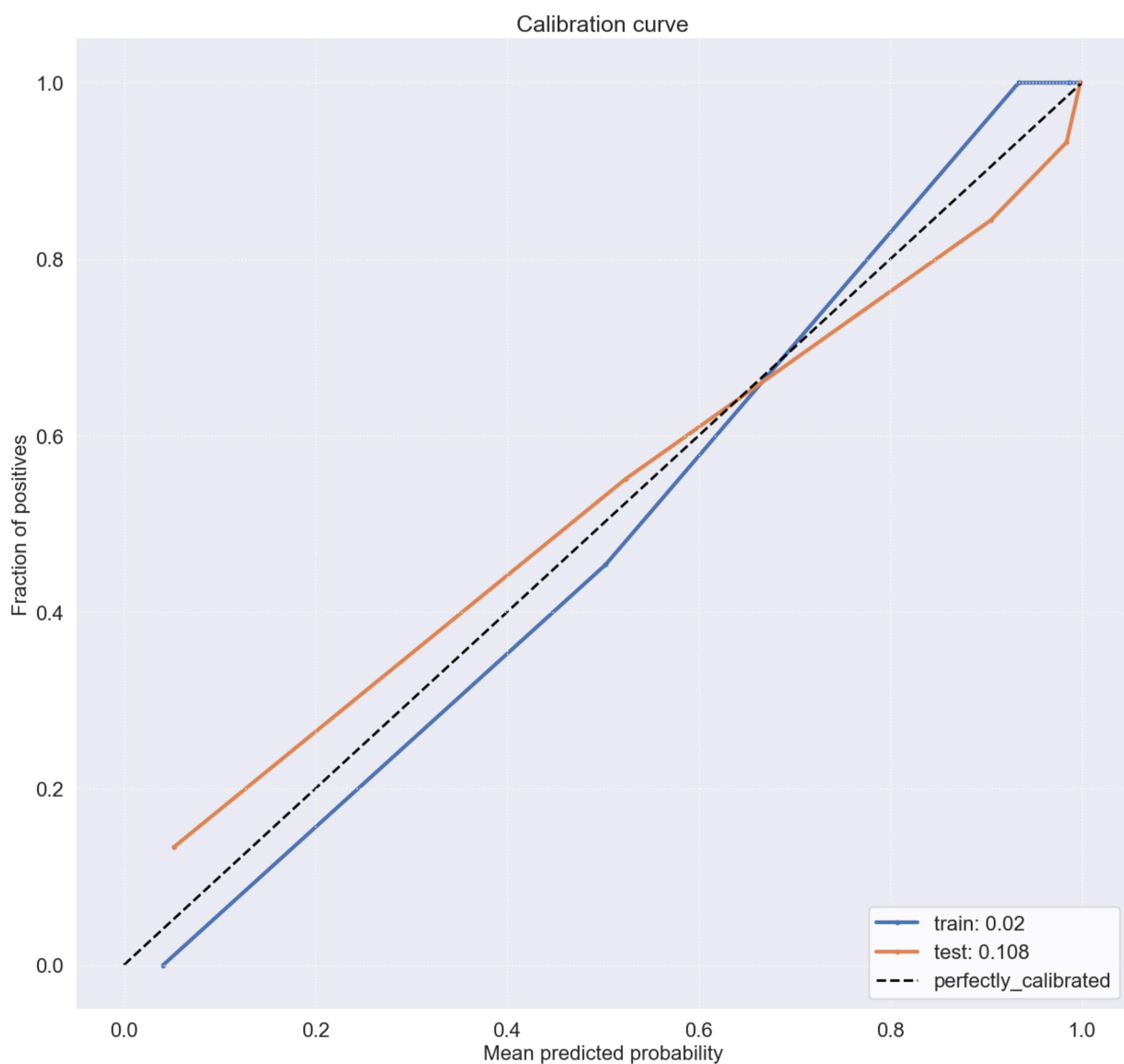
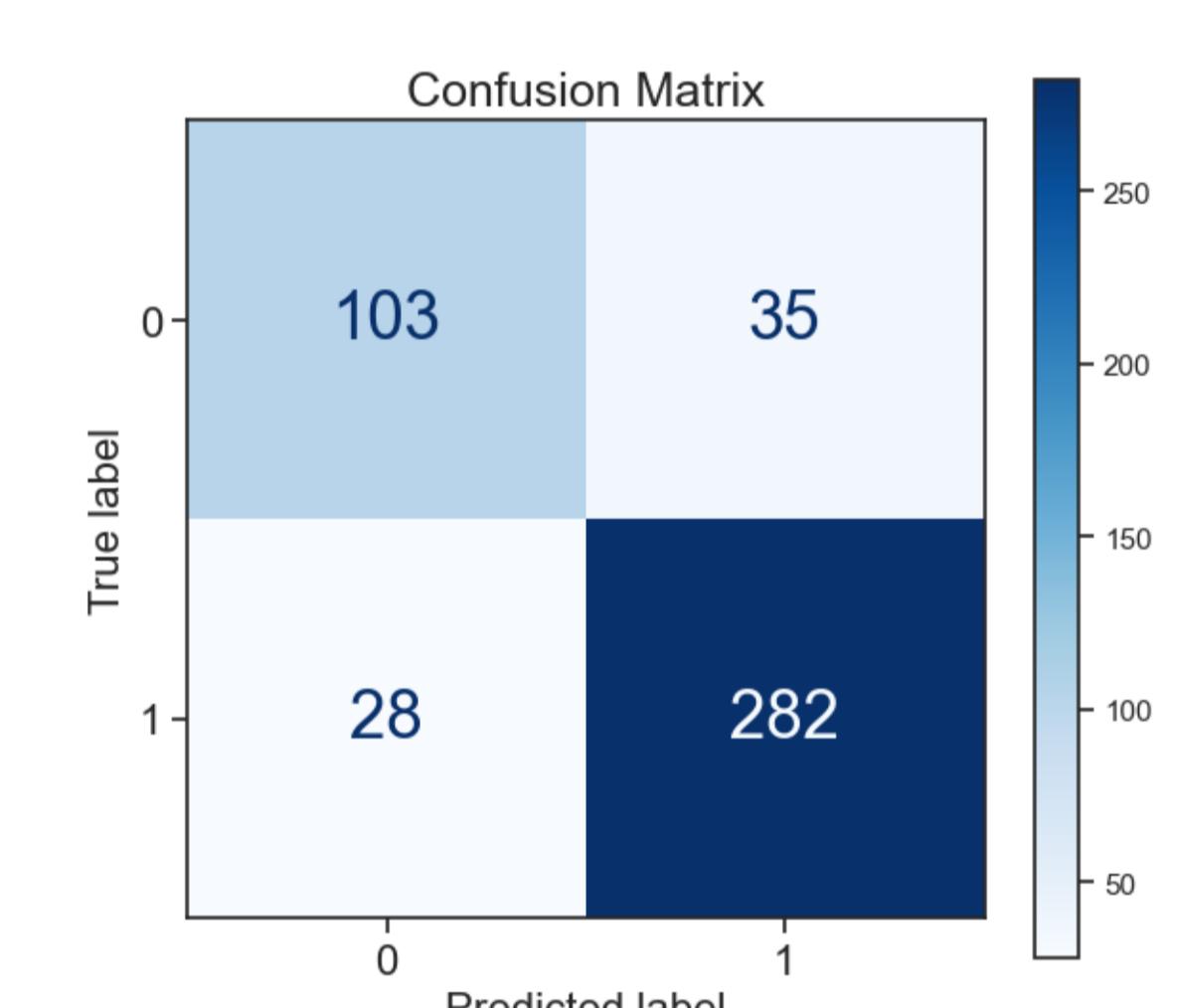
Protein: P28335 - metric: f1



Protein: P28335 - metric: precision



| XGBoost Parameters | Value |
|--------------------|--------|
| alpha | 2.9929 |
| colsample_bytree | 0.9595 |
| gamma | 0.1978 |
| lambda | 2.4528 |
| learning_rate | 0.2287 |
| max_depth | 3.0 |
| min_child_weight | 3.0 |
| n_estimators | 754.0 |
| reg_alpha | 0.5149 |
| reg_lambda | 0.0868 |
| scale_pos_weight | 1.0302 |
| subsample | 0.7295 |



| Metric | Train | Test |
|-------------------|--------|--------|
| AUC | 0.9991 | 0.918 |
| accuracy | 0.9827 | 0.8594 |
| balanced accuracy | 0.9745 | 0.828 |
| recall_(sens) | 0.996 | 0.9097 |
| specificity | 0.953 | 0.7464 |
| precision | 0.9793 | 0.8896 |
| f1_score | 0.9826 | 0.8583 |

Protein: P28335 - metric: SoftVotingXGBoostEnsemble

