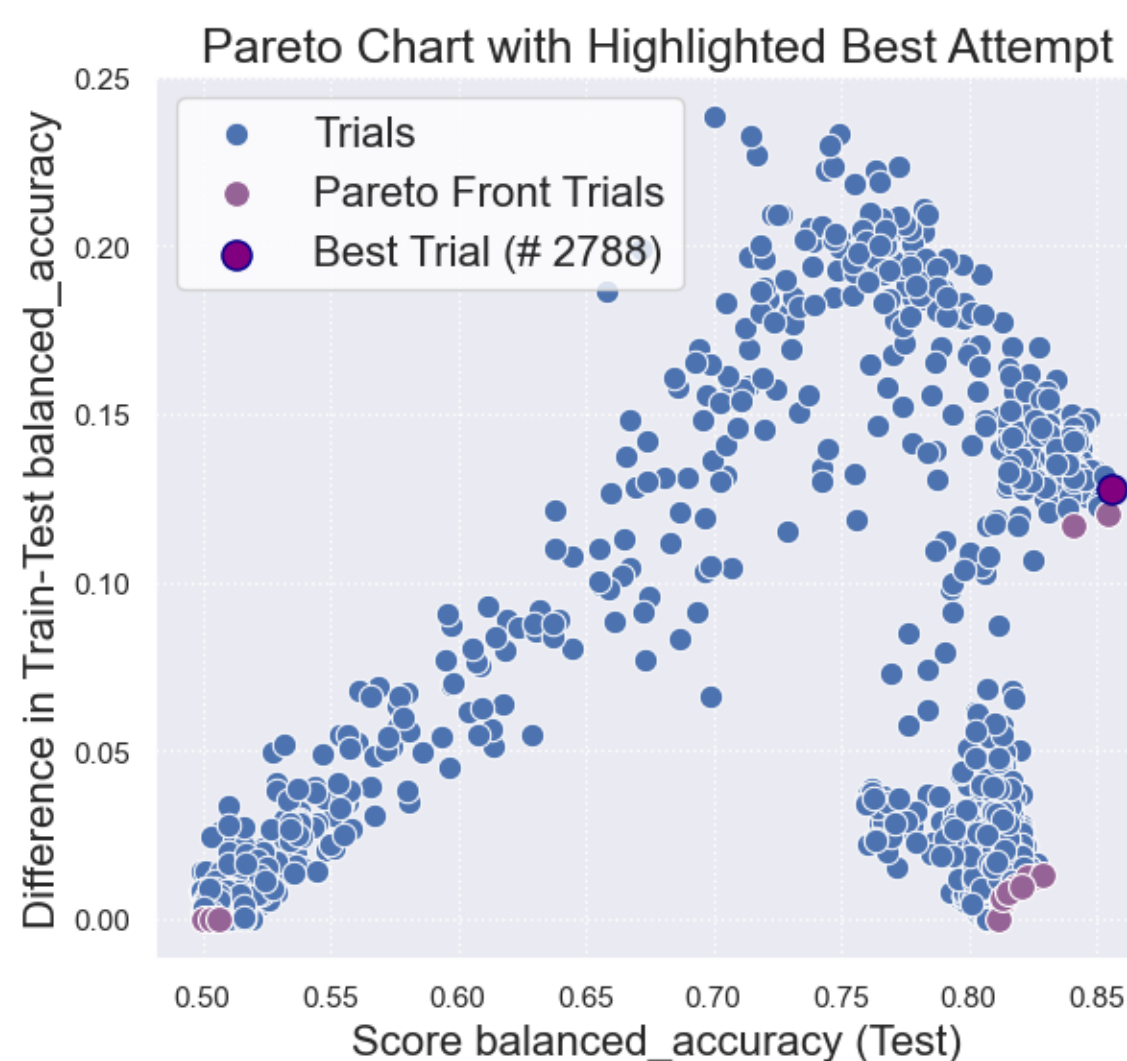
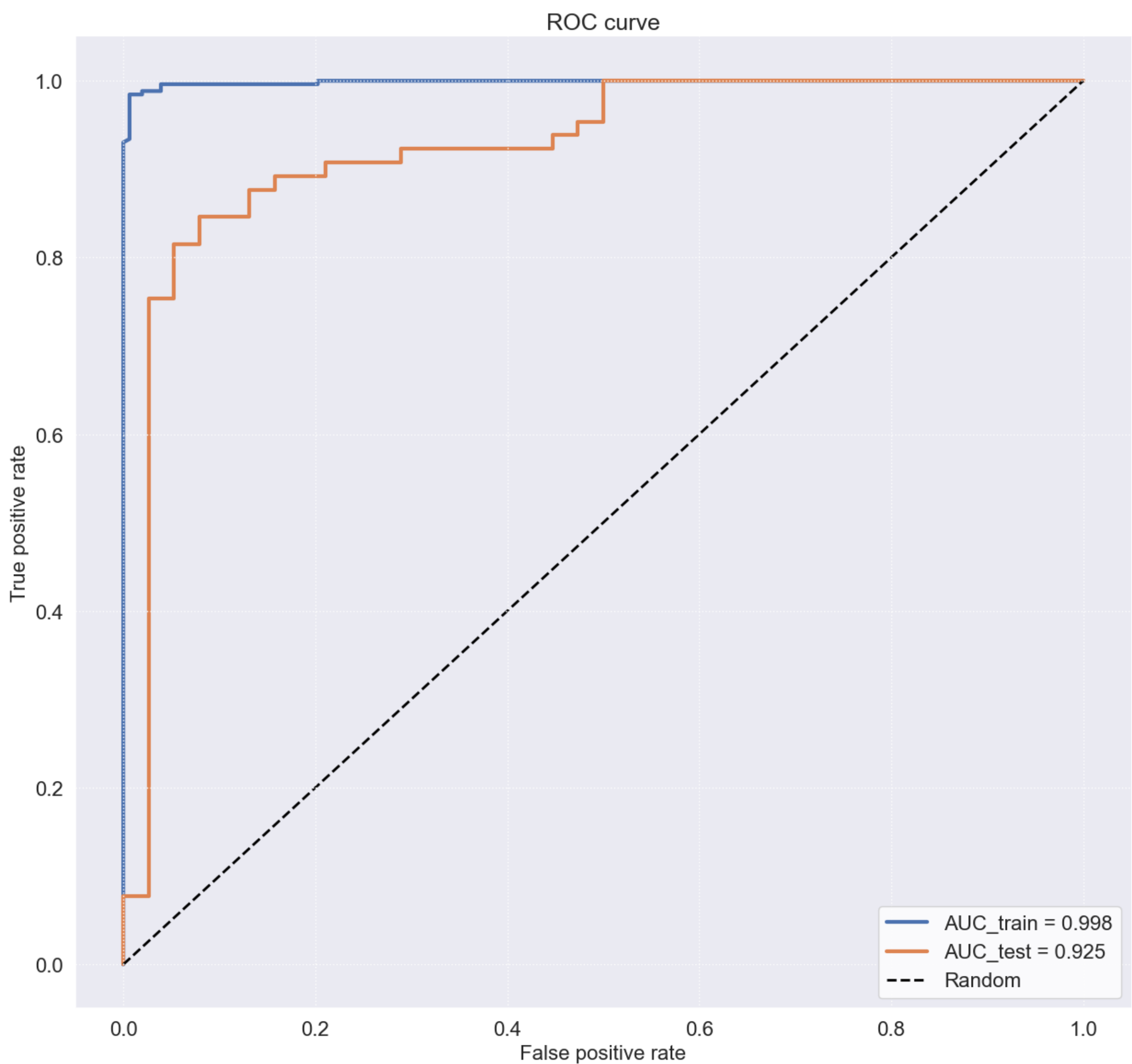
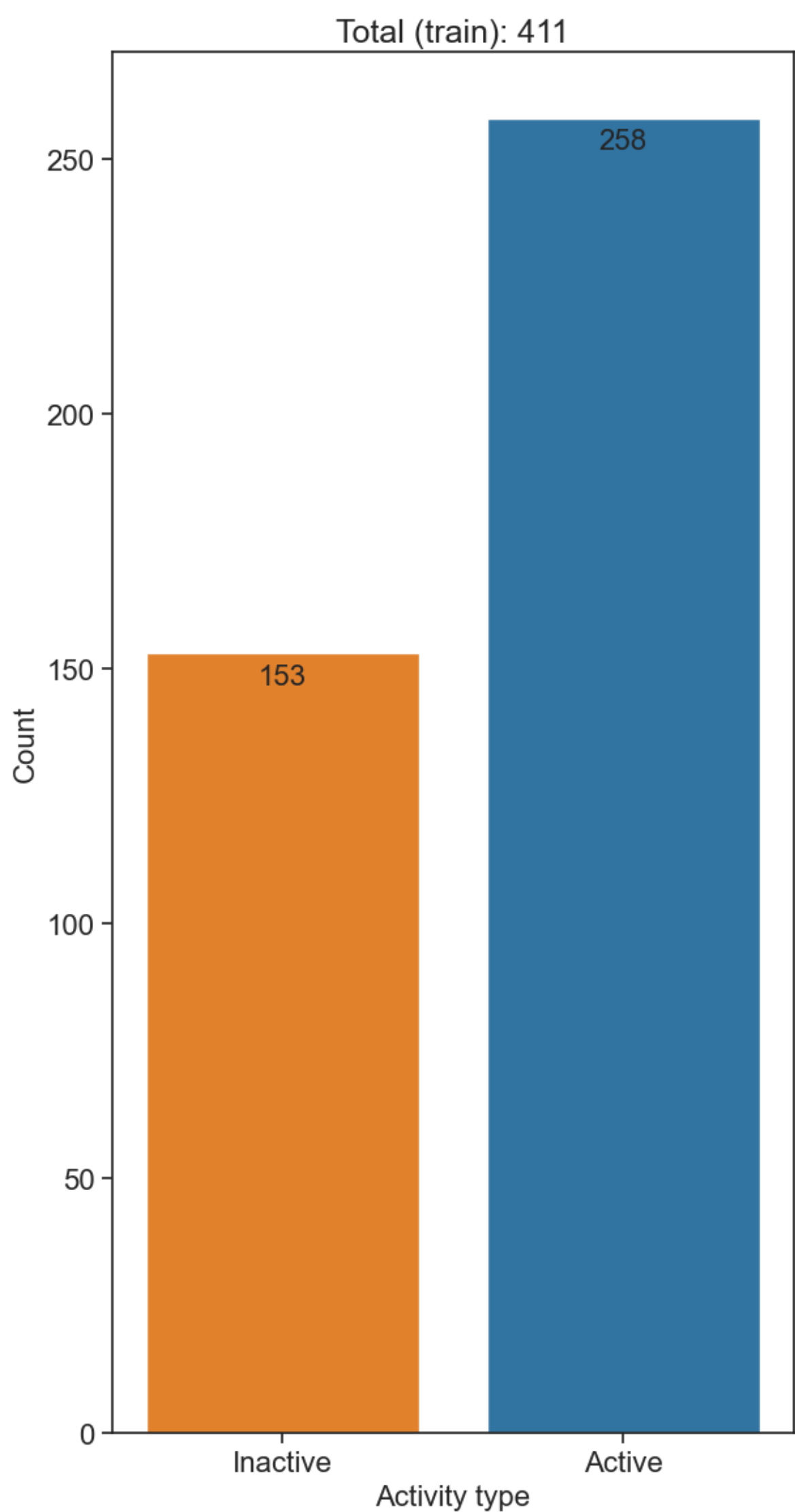
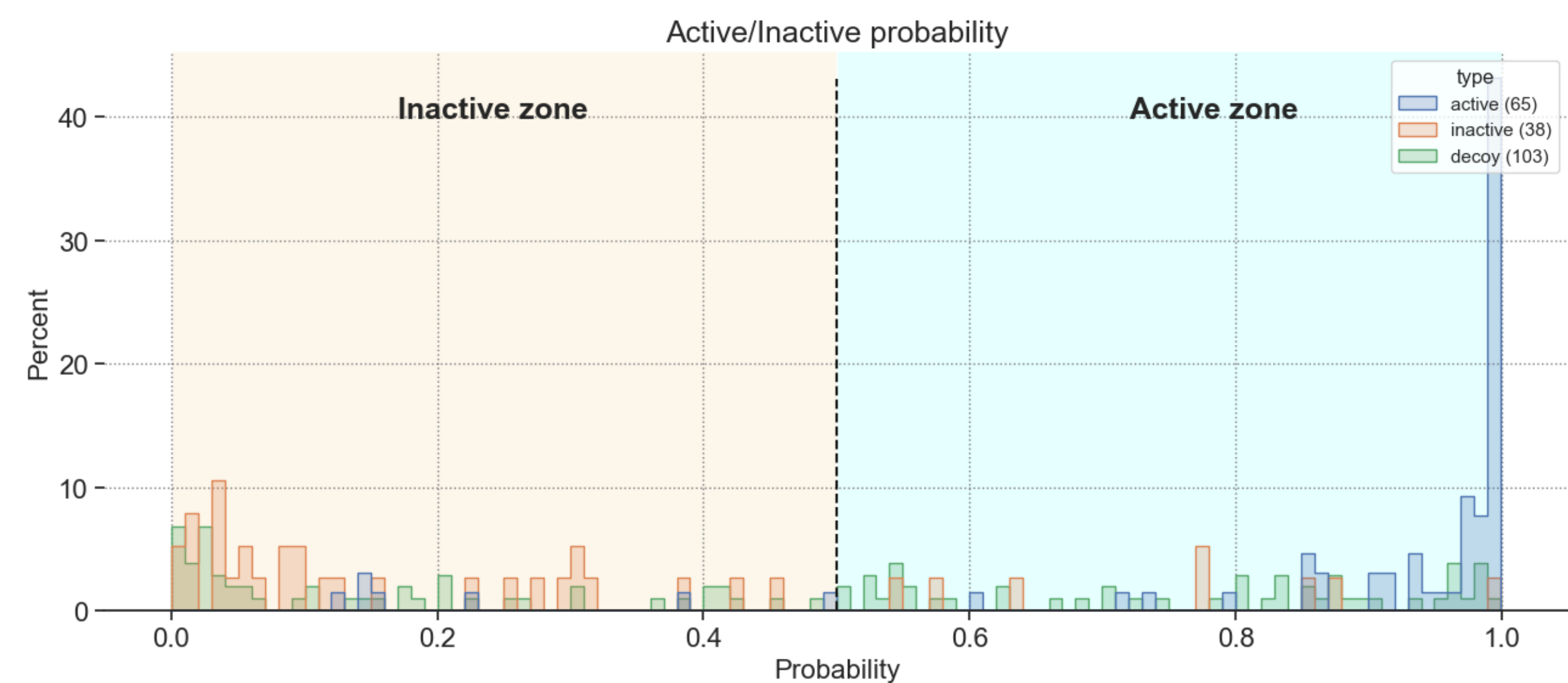
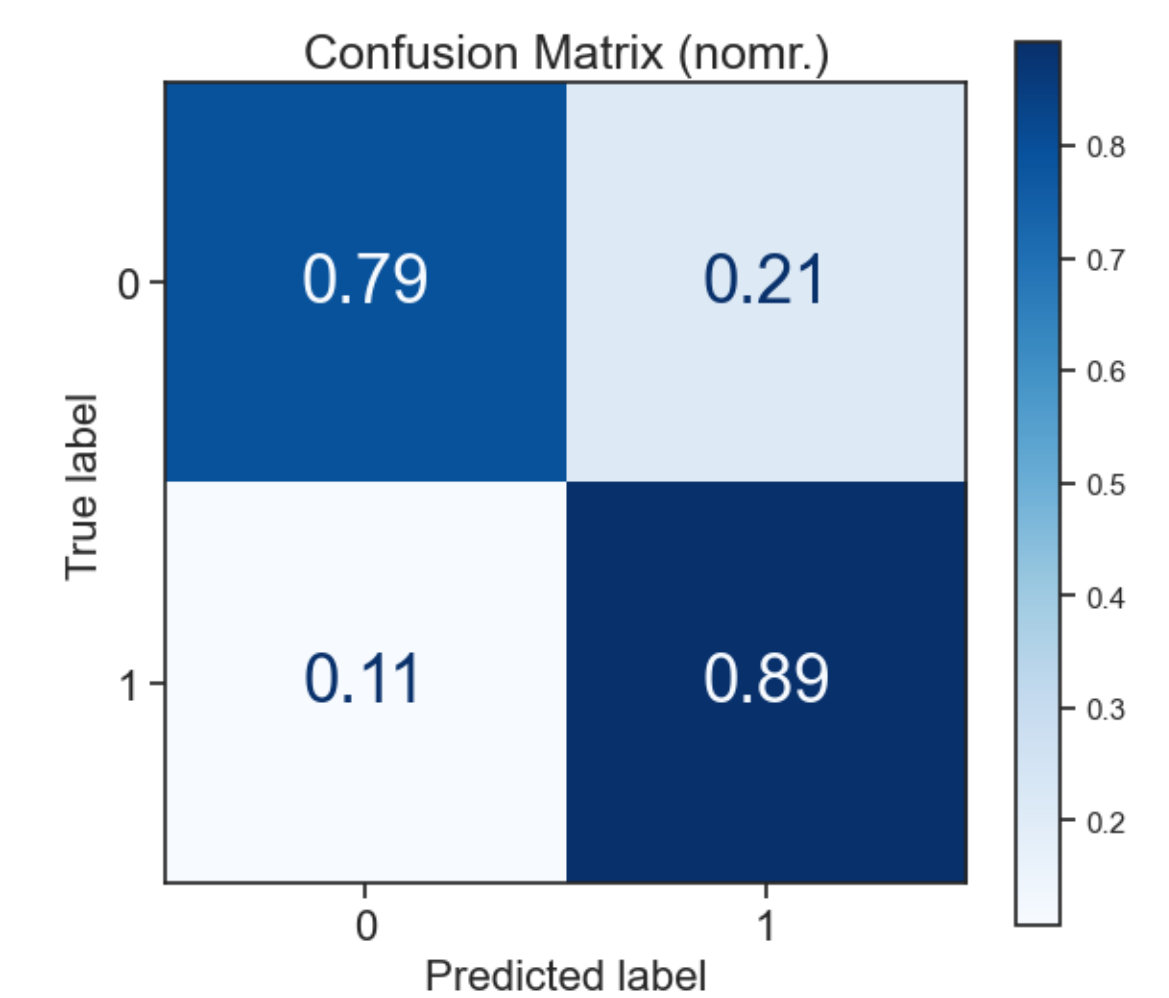
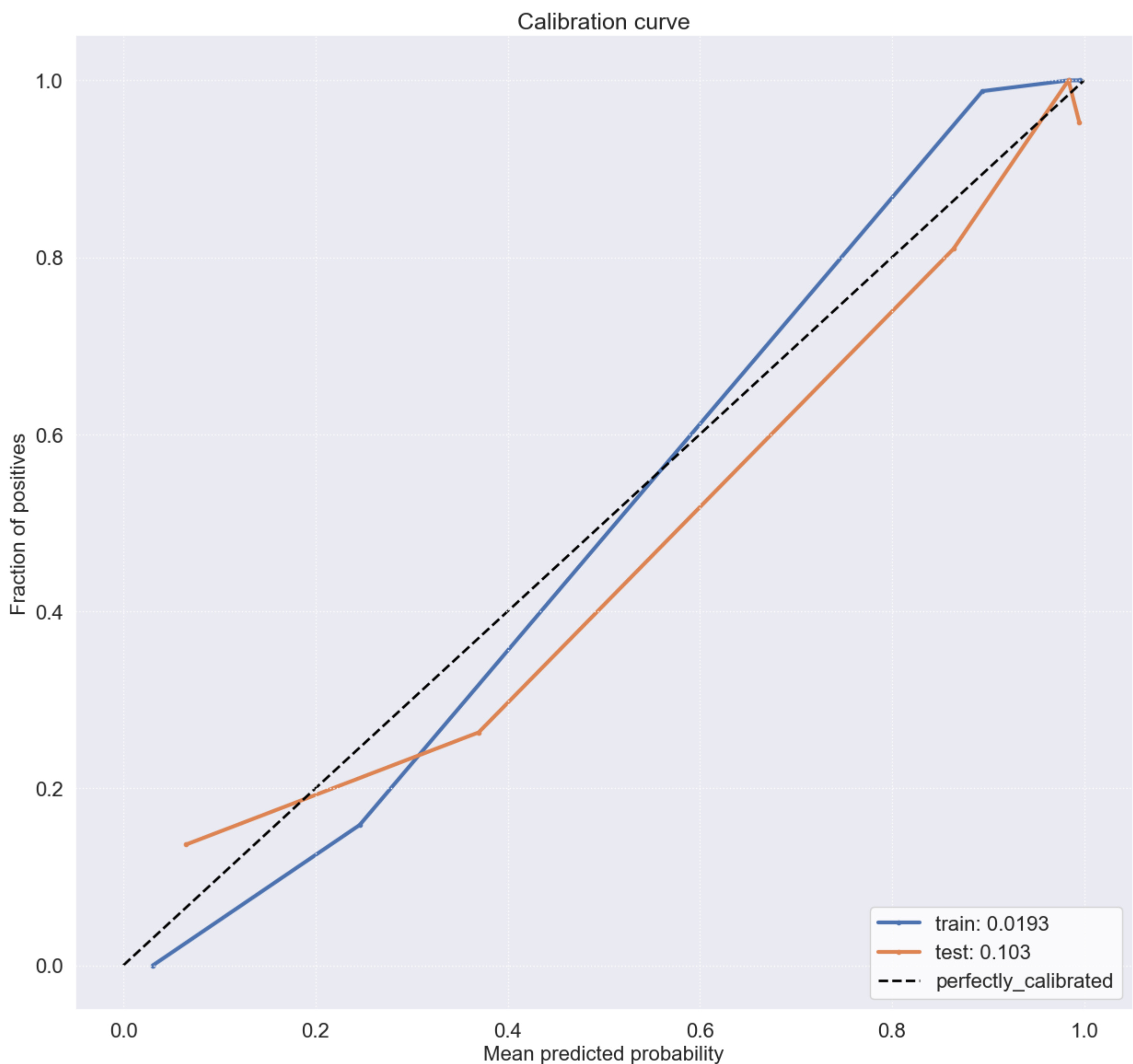
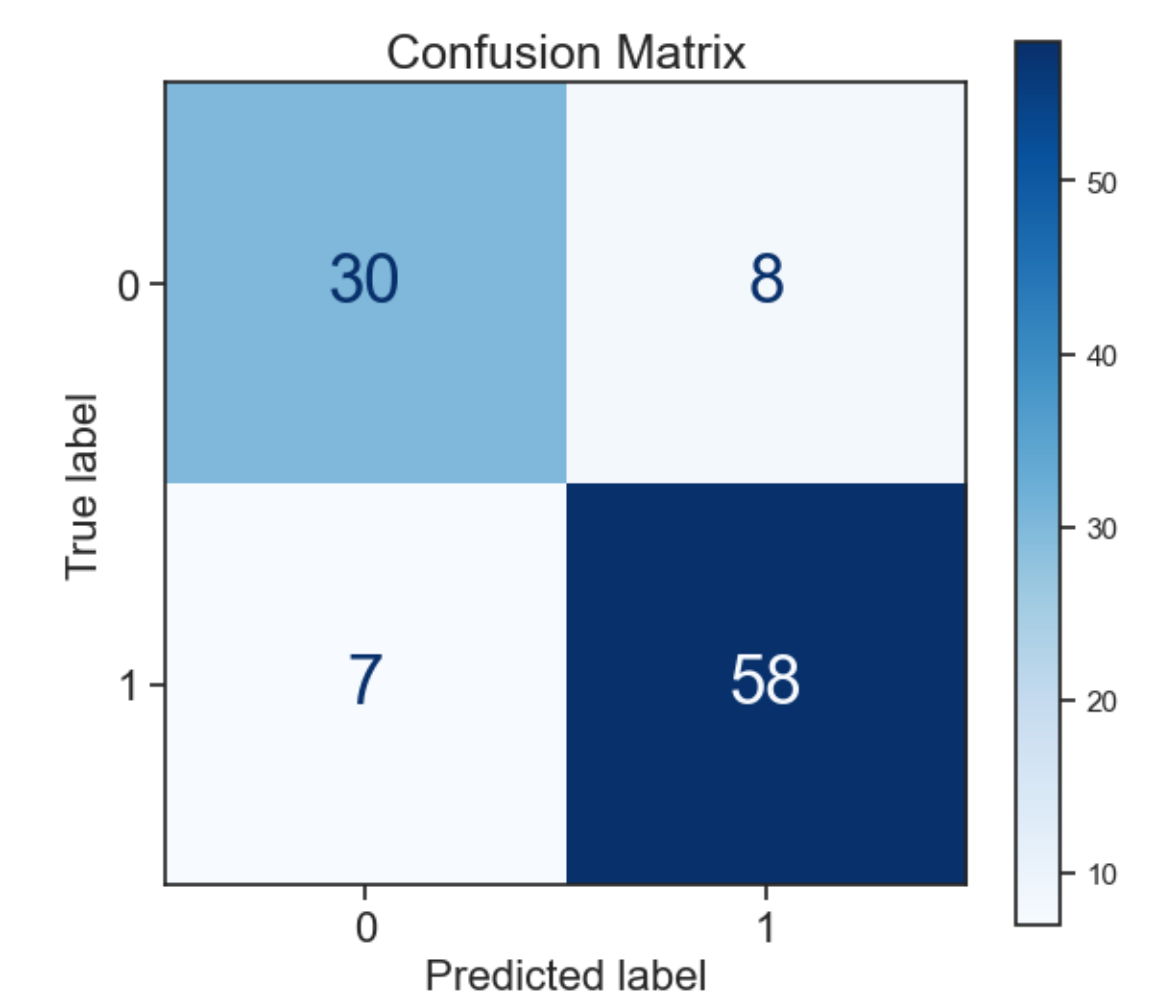


Protein: P06213 - metric: balanced\_accuracy



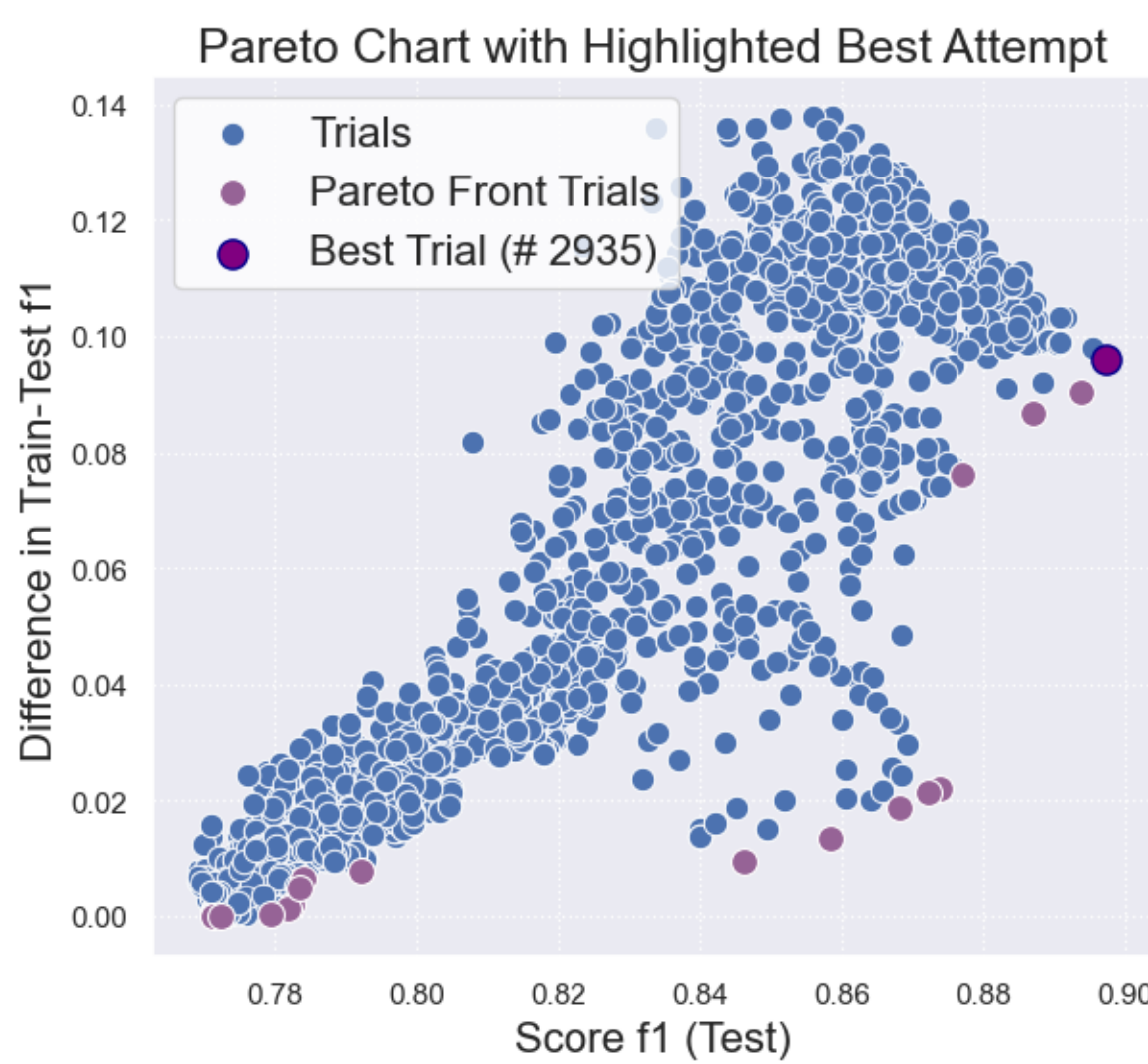
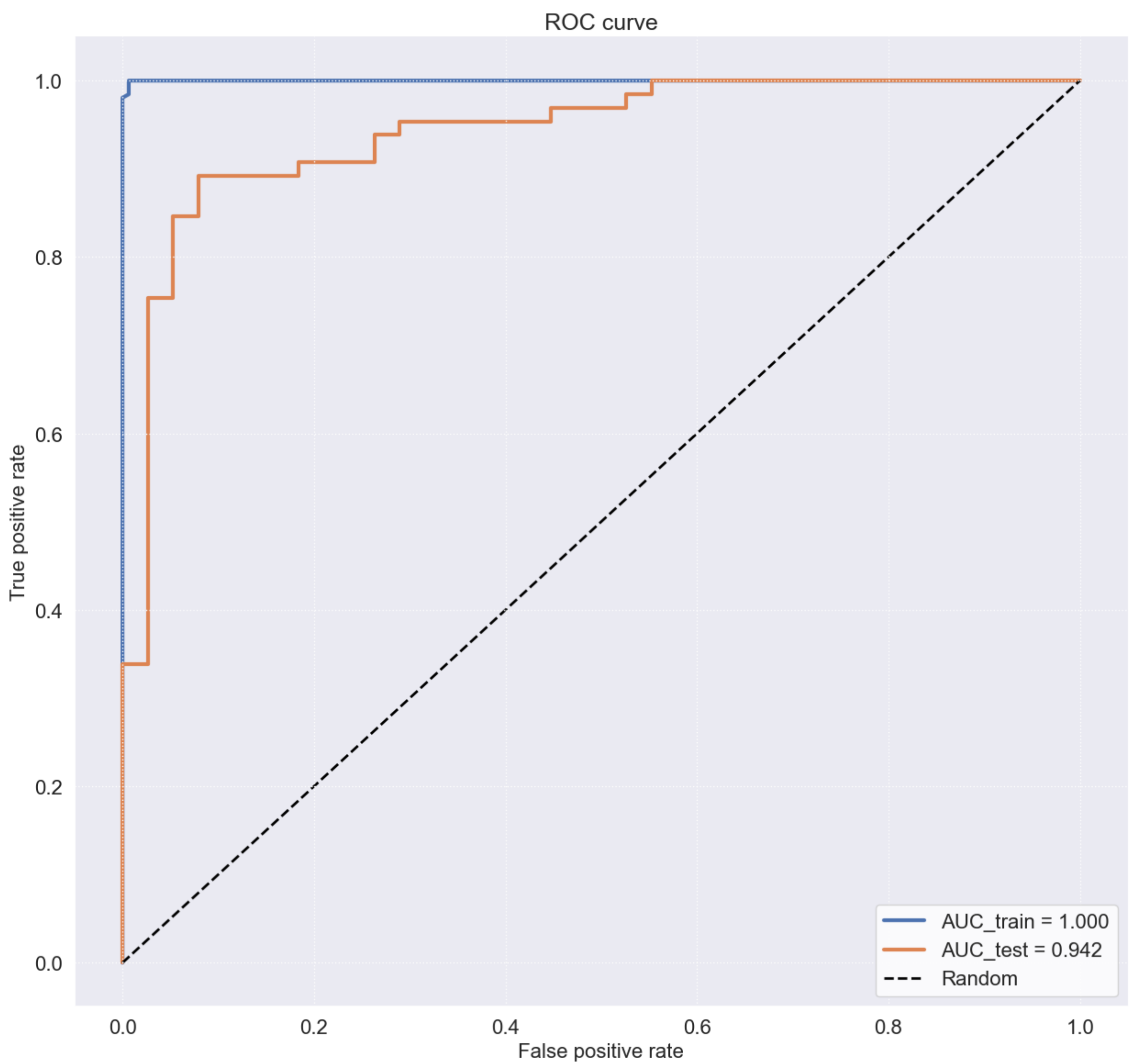
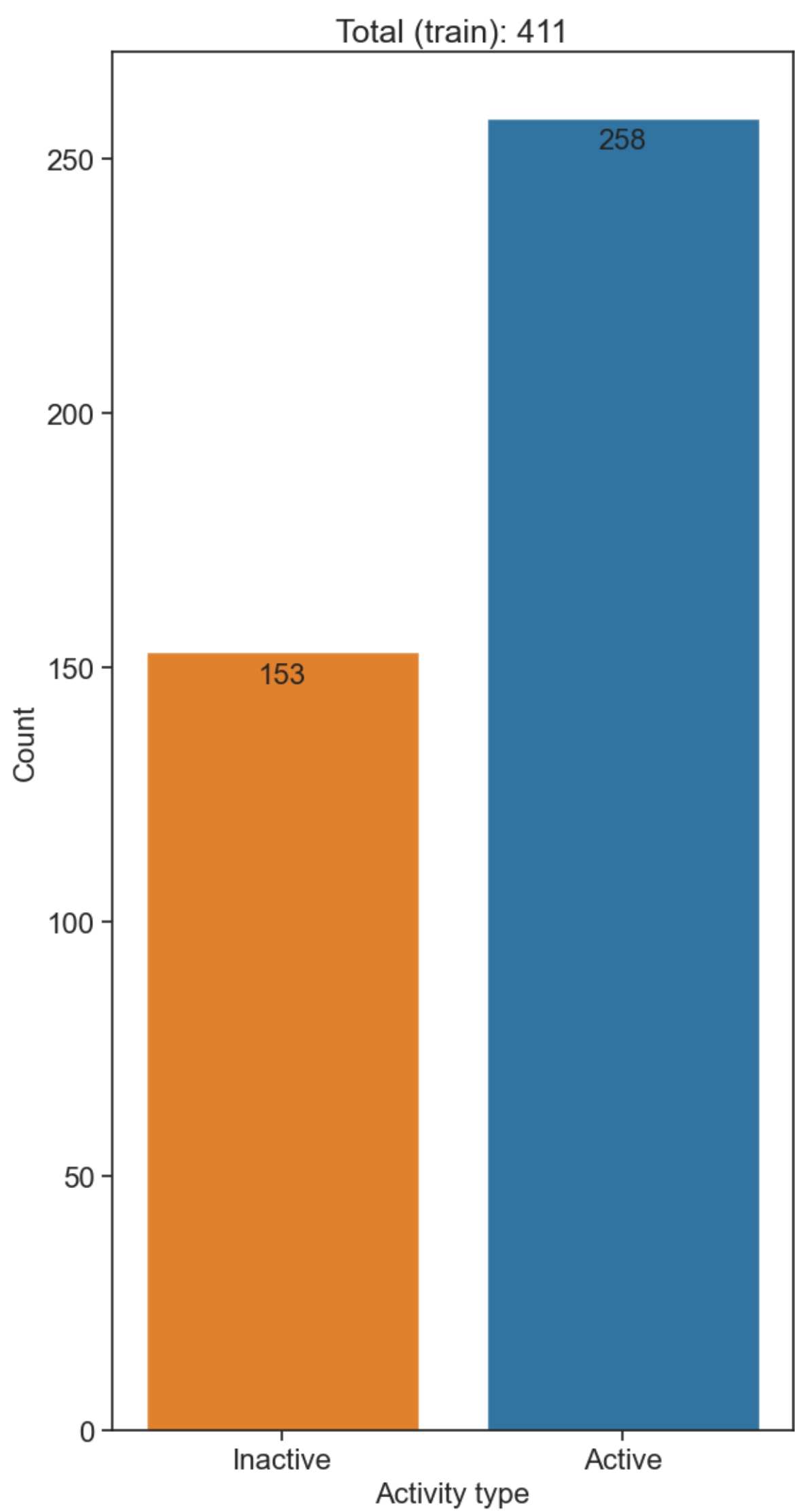
| XGBoost Parameters | Value  |
|--------------------|--------|
| alpha              | 1.4574 |
| colsample_bytree   | 0.7944 |
| gamma              | 0.963  |
| lambda             | 0.0386 |
| learning_rate      | 0.0998 |
| max_depth          | 10.0   |
| min_child_weight   | 2.0    |
| n_estimators       | 878.0  |
| reg_alpha          | 0.0532 |
| reg_lambda         | 0.4472 |
| scale_pos_weight   | 1.0672 |
| subsample          | 0.608  |



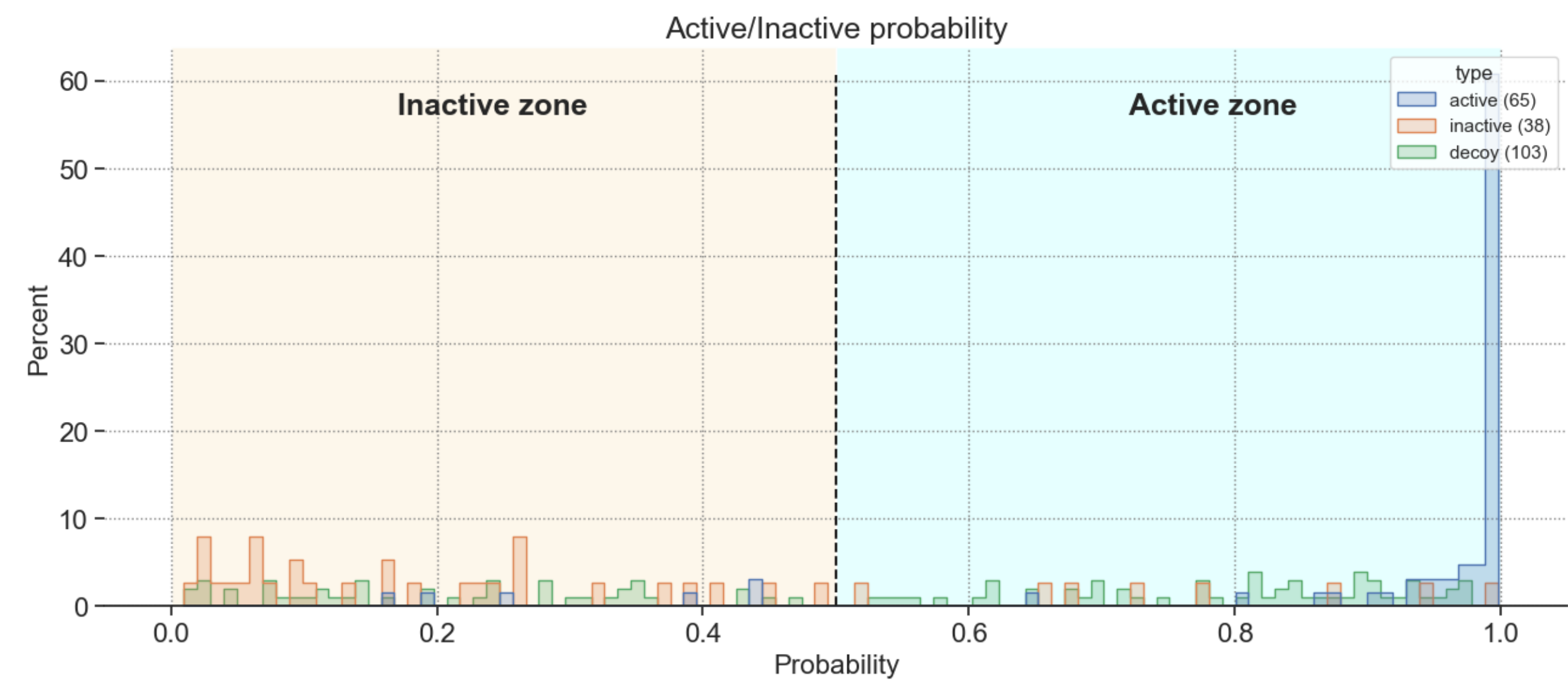
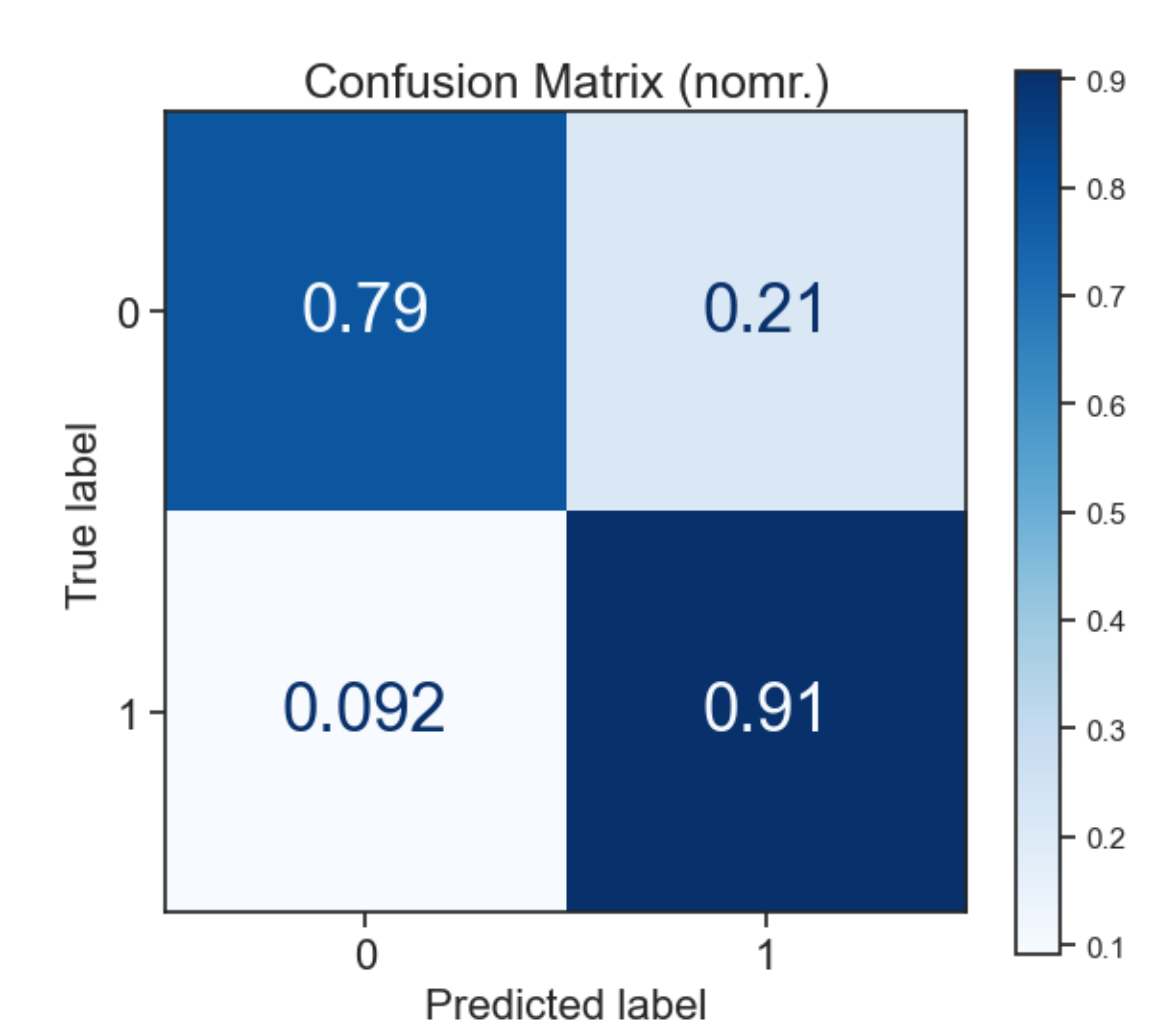
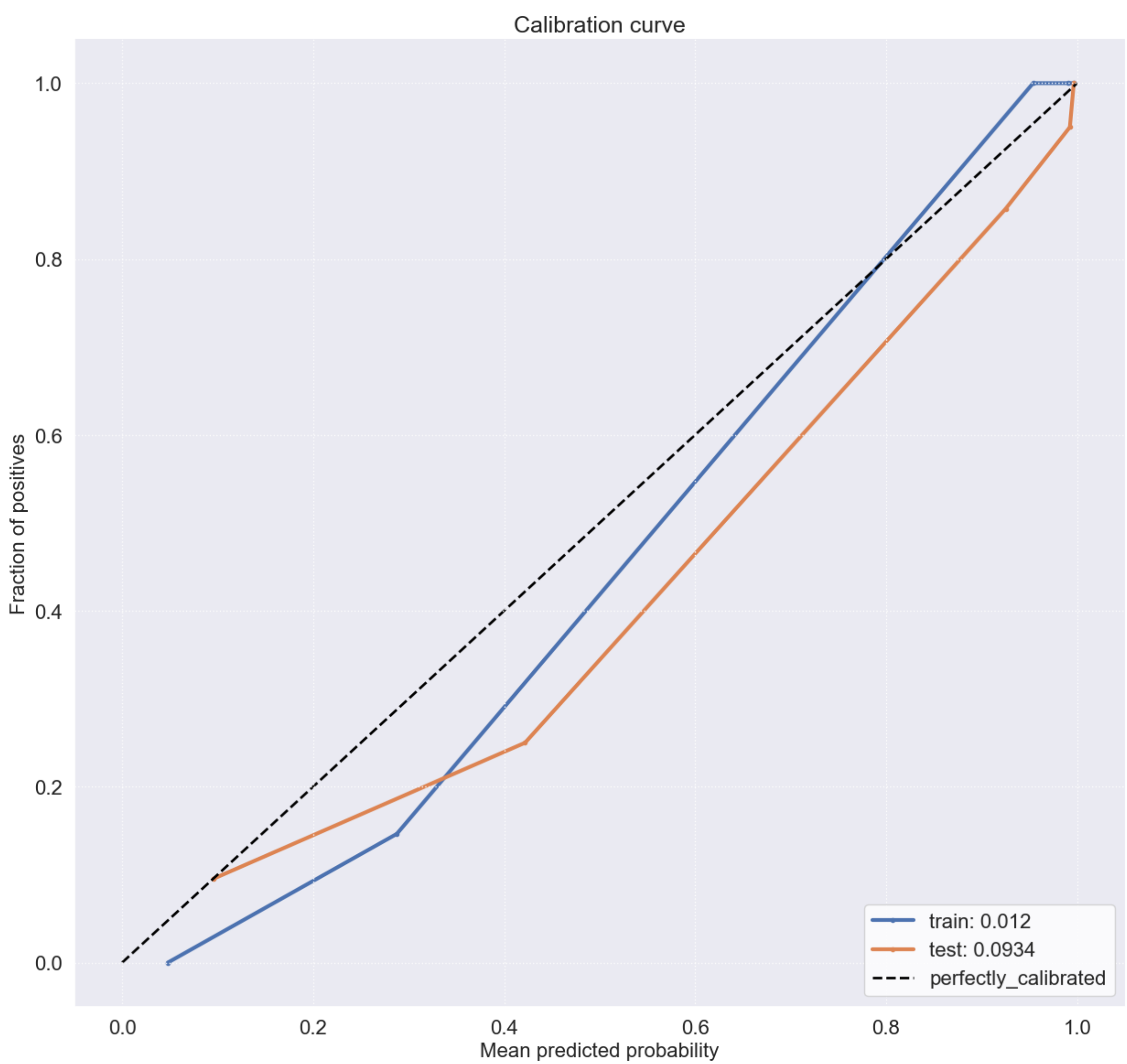
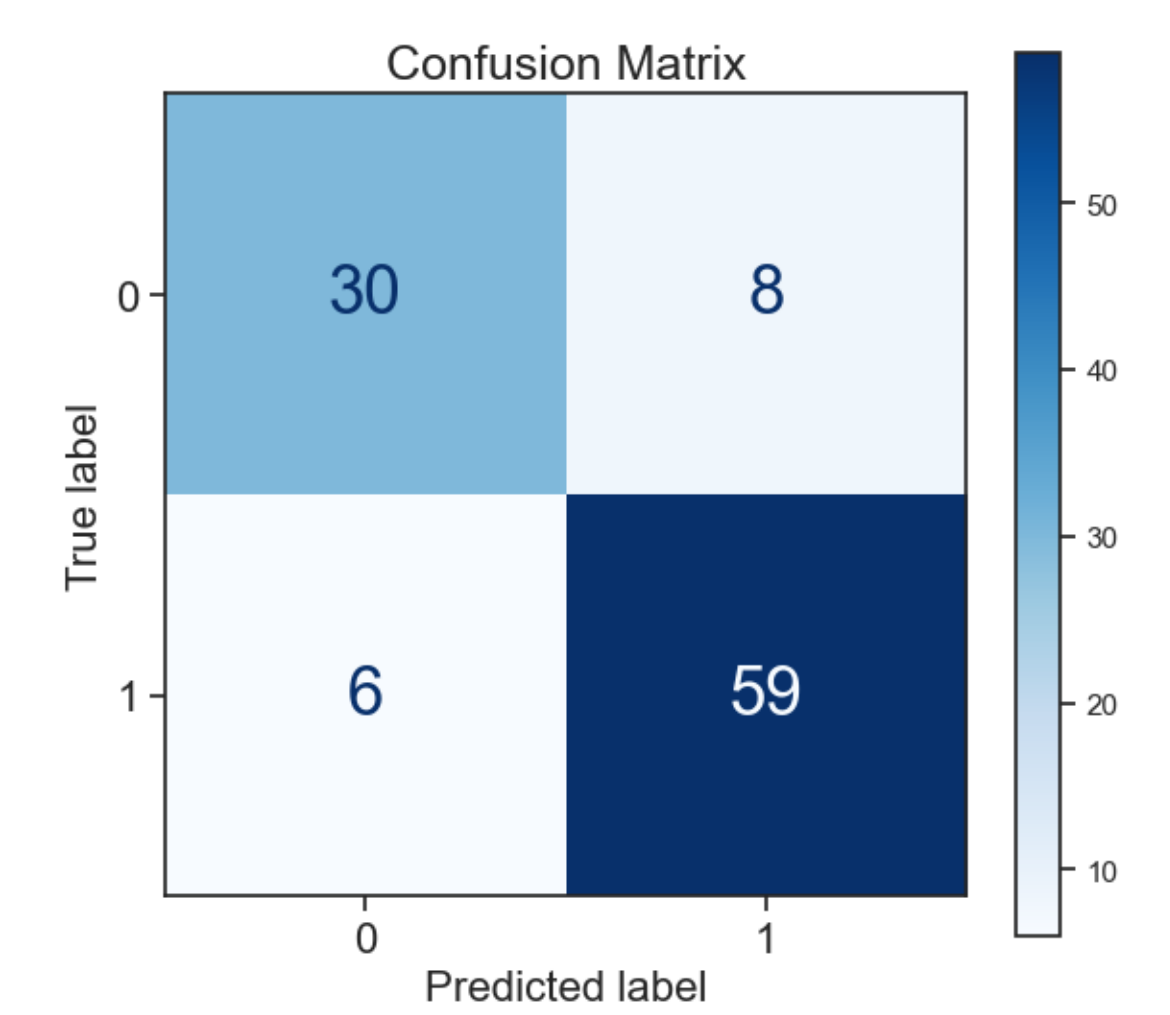
| Metric            | Train  | Test   |
|-------------------|--------|--------|
| AUC               | 0.9985 | 0.9251 |
| accuracy          | 0.9805 | 0.8544 |
| balanced accuracy | 0.9778 | 0.8409 |
| recall_(sens)     | 0.9884 | 0.8923 |
| specificity       | 0.9673 | 0.7895 |
| precision         | 0.9808 | 0.8788 |
| f1_score          | 0.9805 | 0.854  |



Protein: P06213 - metric: f1



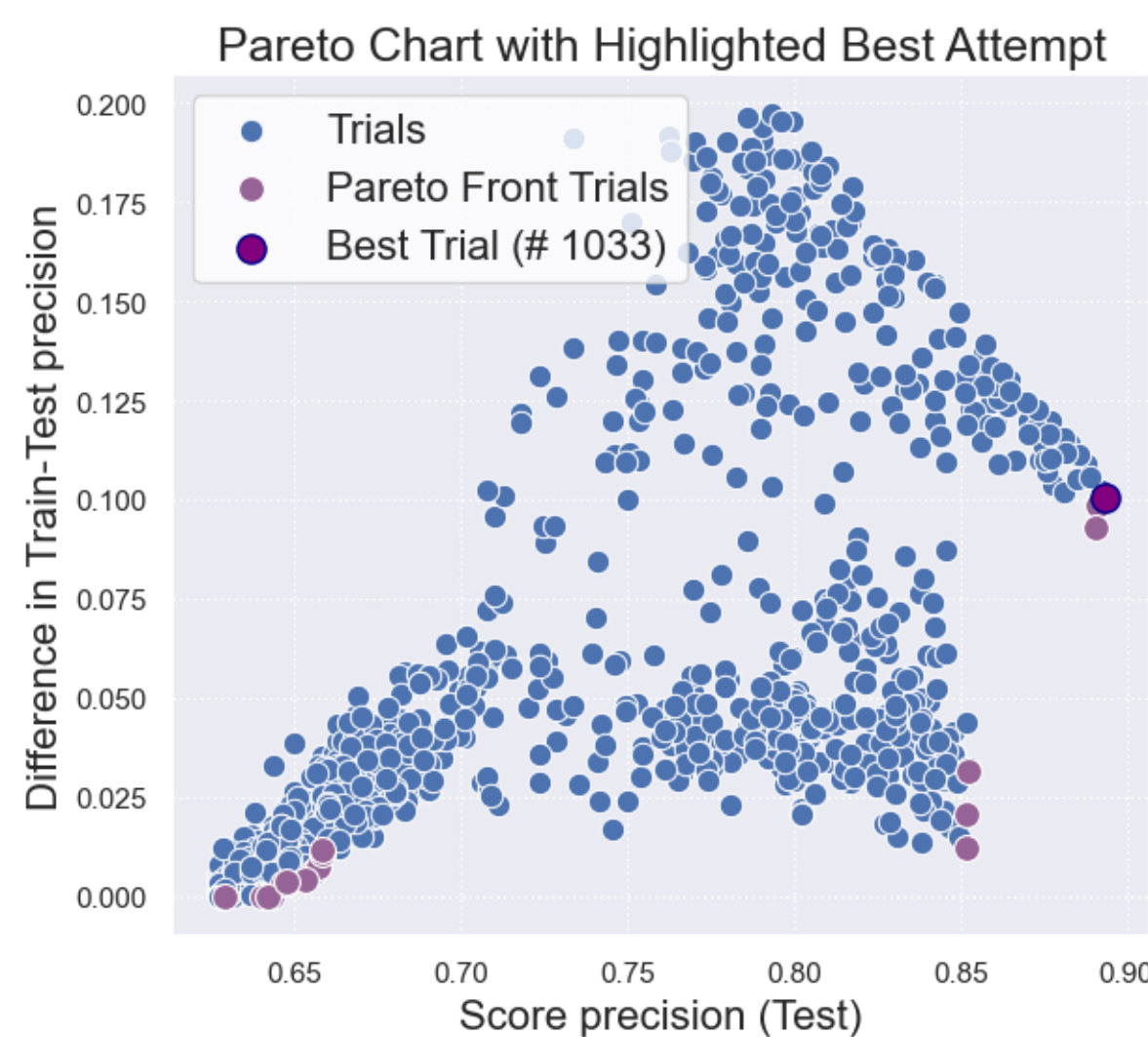
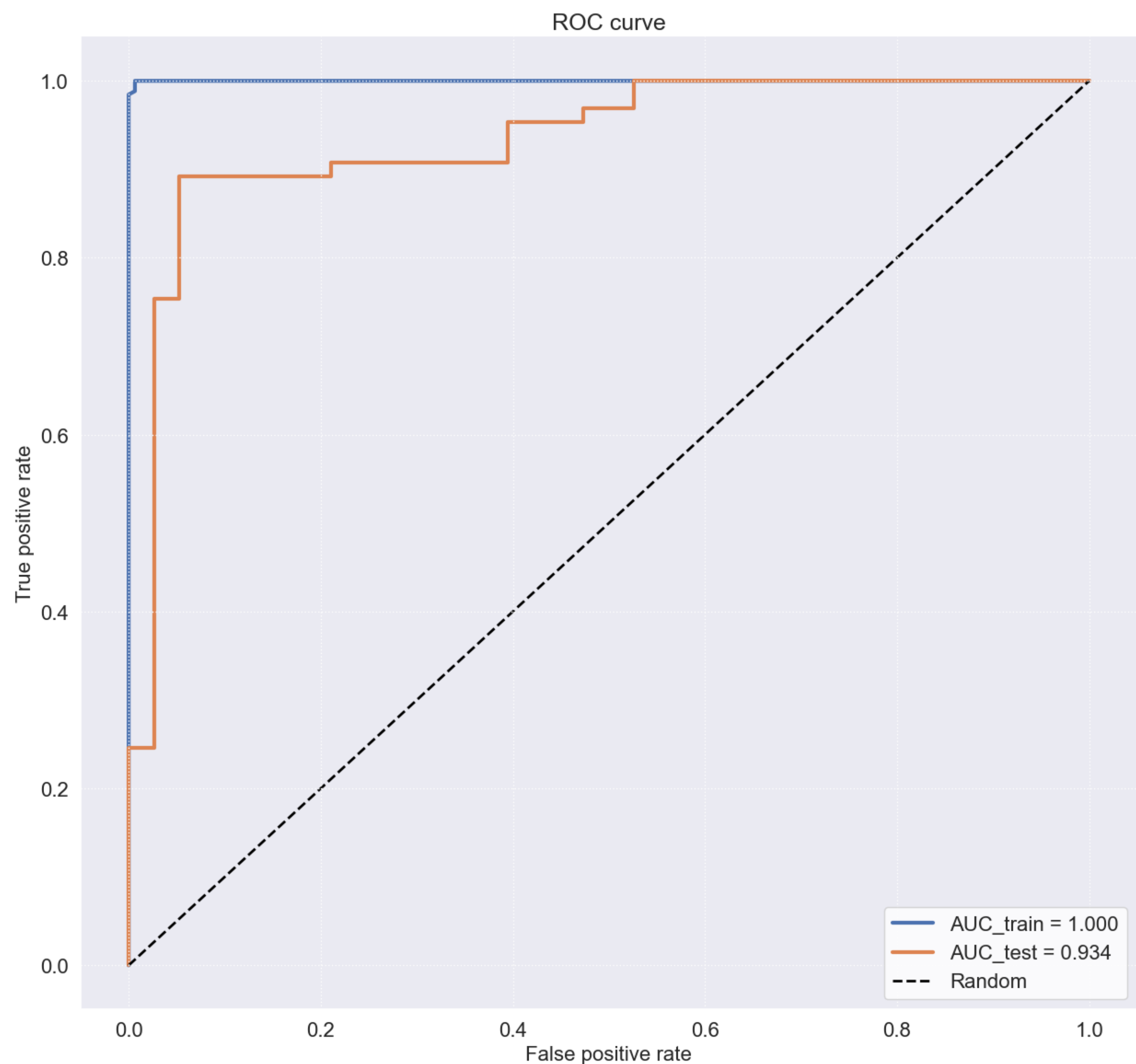
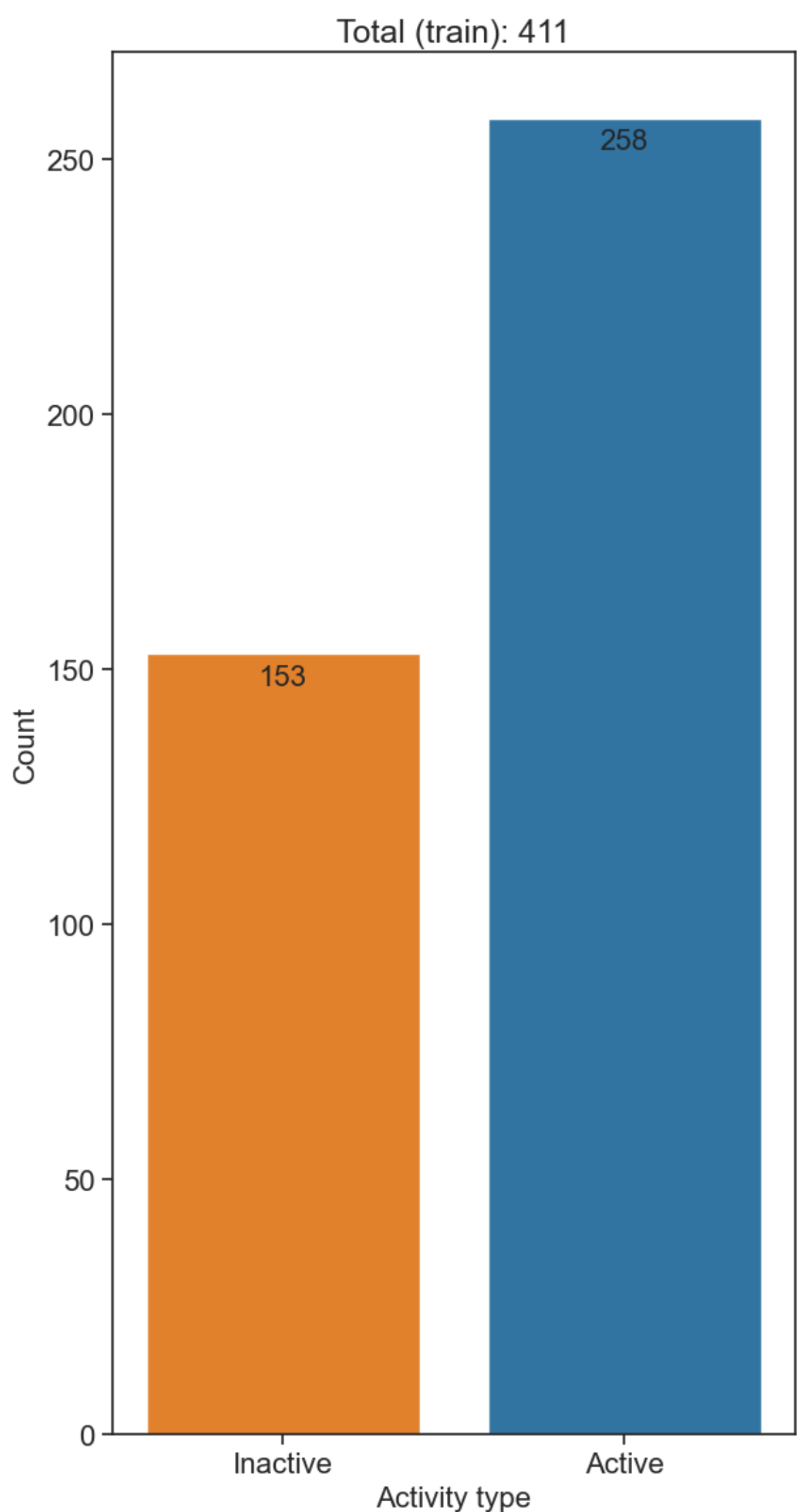
| XGBoost Parameters | Value  |
|--------------------|--------|
| alpha              | 3.533  |
| colsample_bytree   | 0.7193 |
| gamma              | 0.7841 |
| lambda             | 4.5179 |
| learning_rate      | 0.023  |
| max_depth          | 10.0   |
| min_child_weight   | 1.0    |
| n_estimators       | 502.0  |
| reg_alpha          | 0.1448 |
| reg_lambda         | 0.0253 |
| scale_pos_weight   | 2.6219 |
| subsample          | 0.7166 |



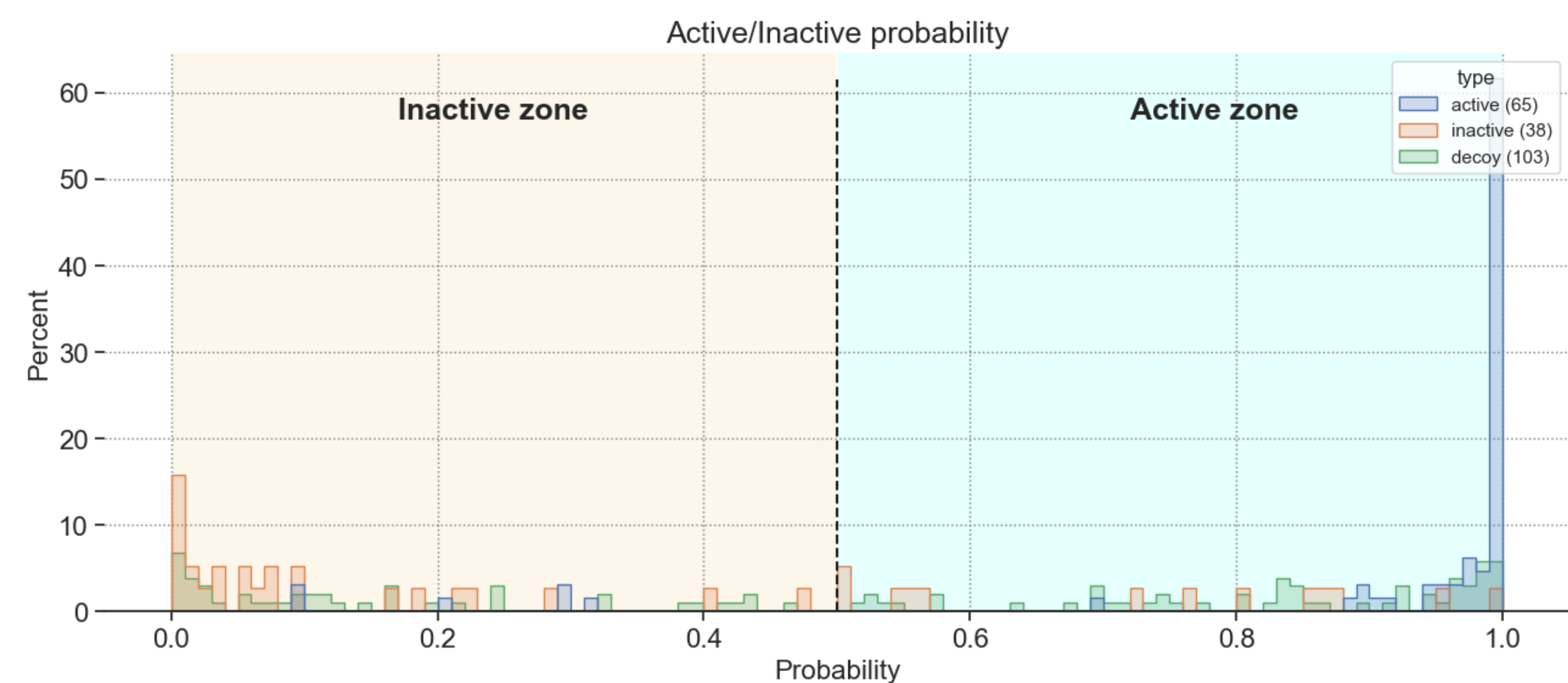
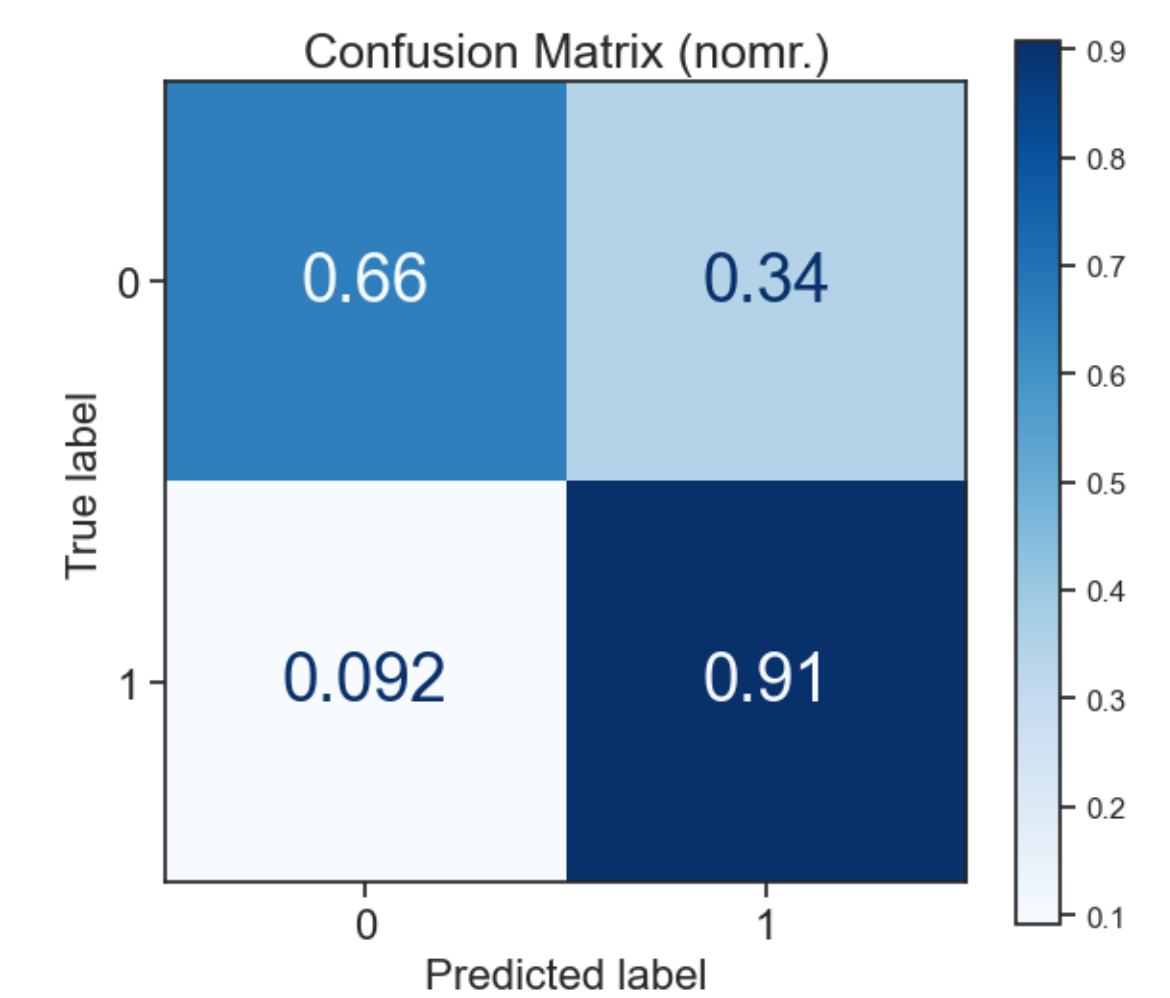
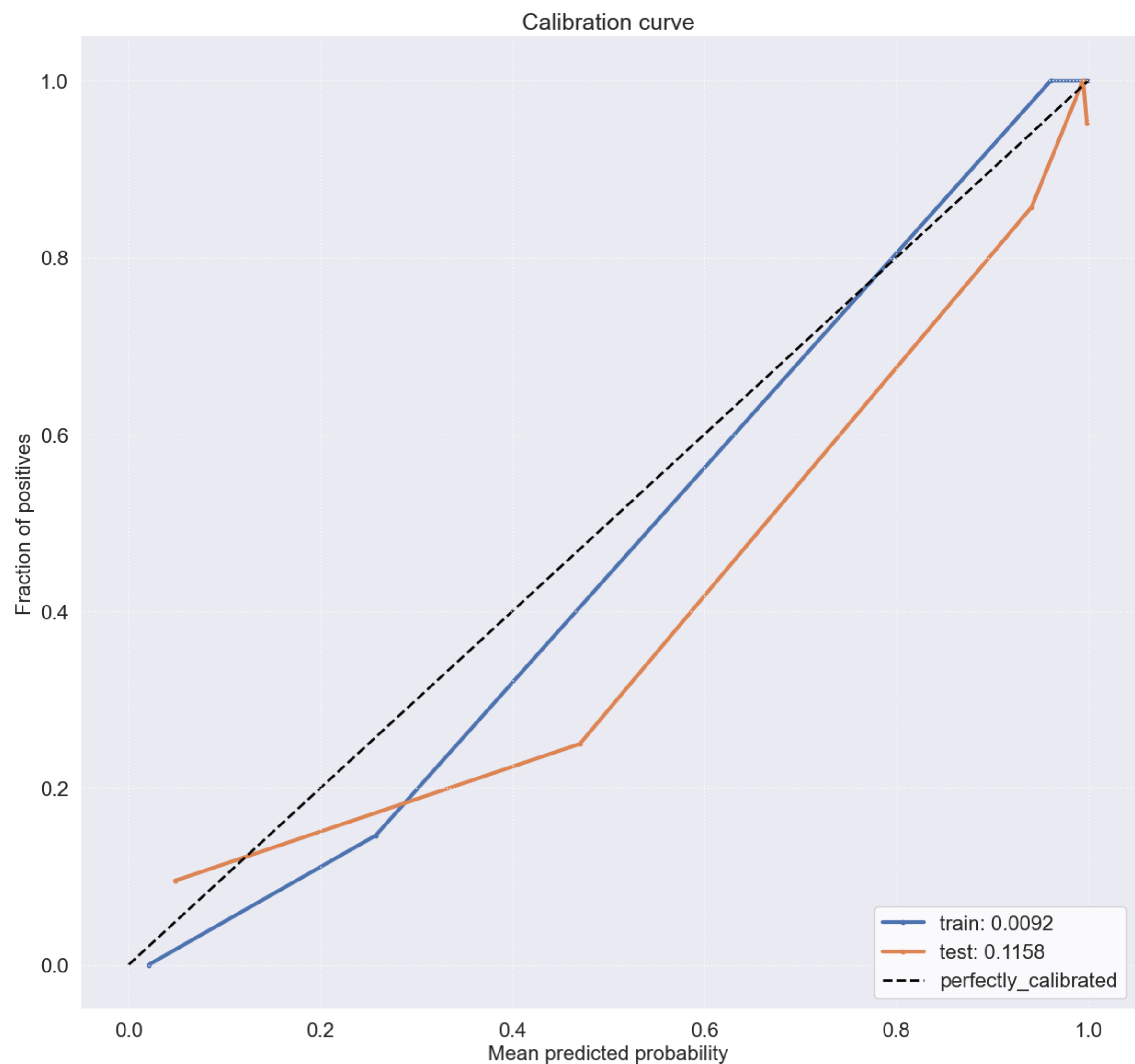
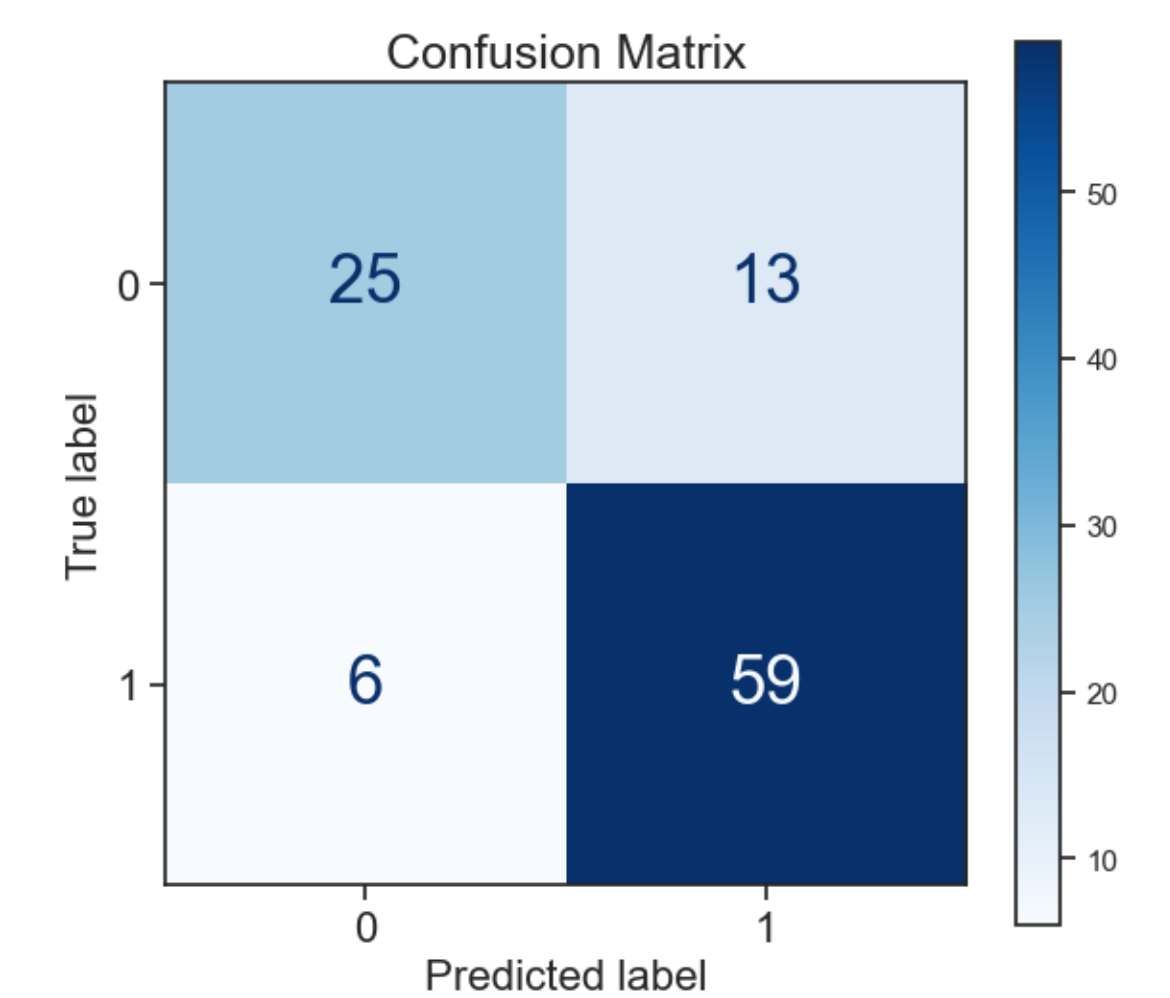
| Metric            | Train  | Test   |
|-------------------|--------|--------|
| AUC               | 0.9999 | 0.9417 |
| accuracy          | 0.9927 | 0.8641 |
| balanced accuracy | 0.9902 | 0.8486 |
| recall_(sens)     | 1.0    | 0.9077 |
| specificity       | 0.9804 | 0.7895 |
| precision         | 0.9885 | 0.8806 |
| f1_score          | 0.9927 | 0.8633 |



Protein: P06213 - metric: precision



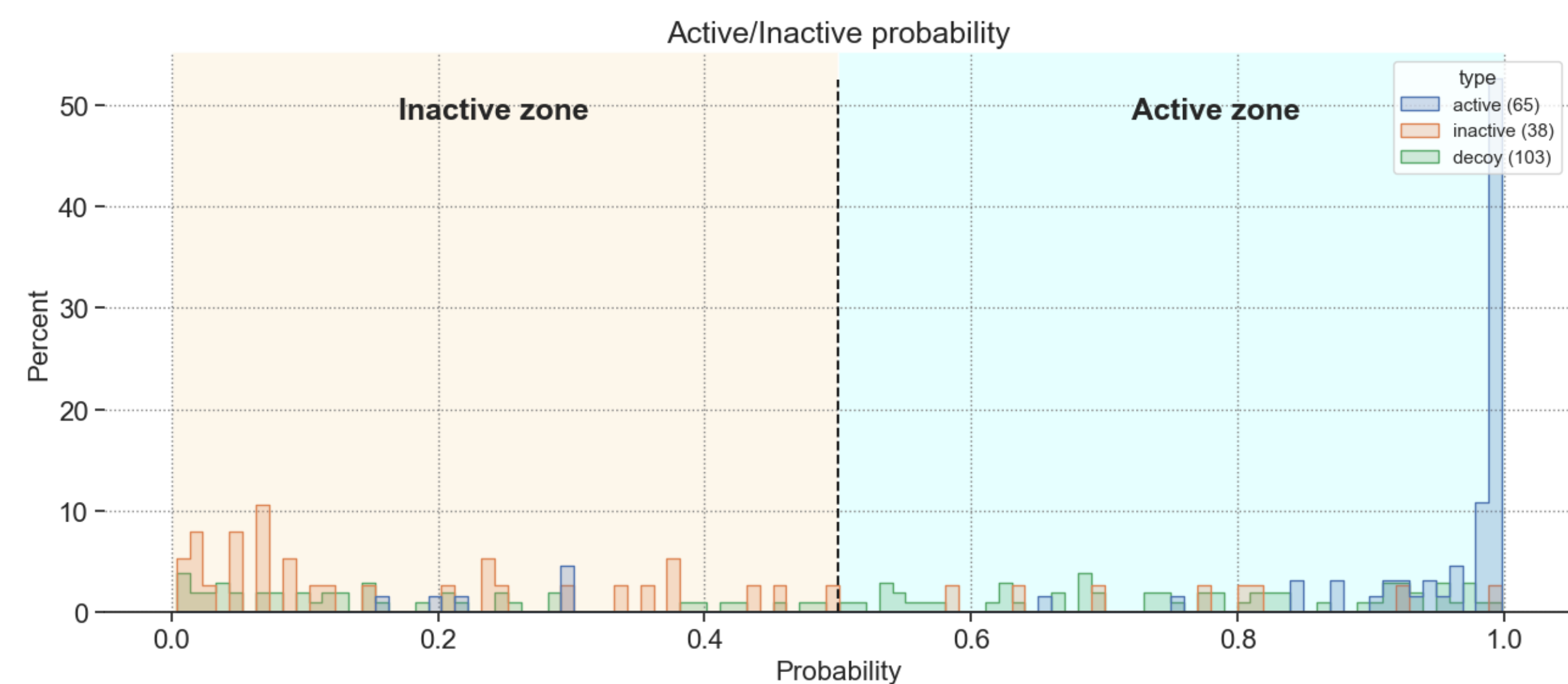
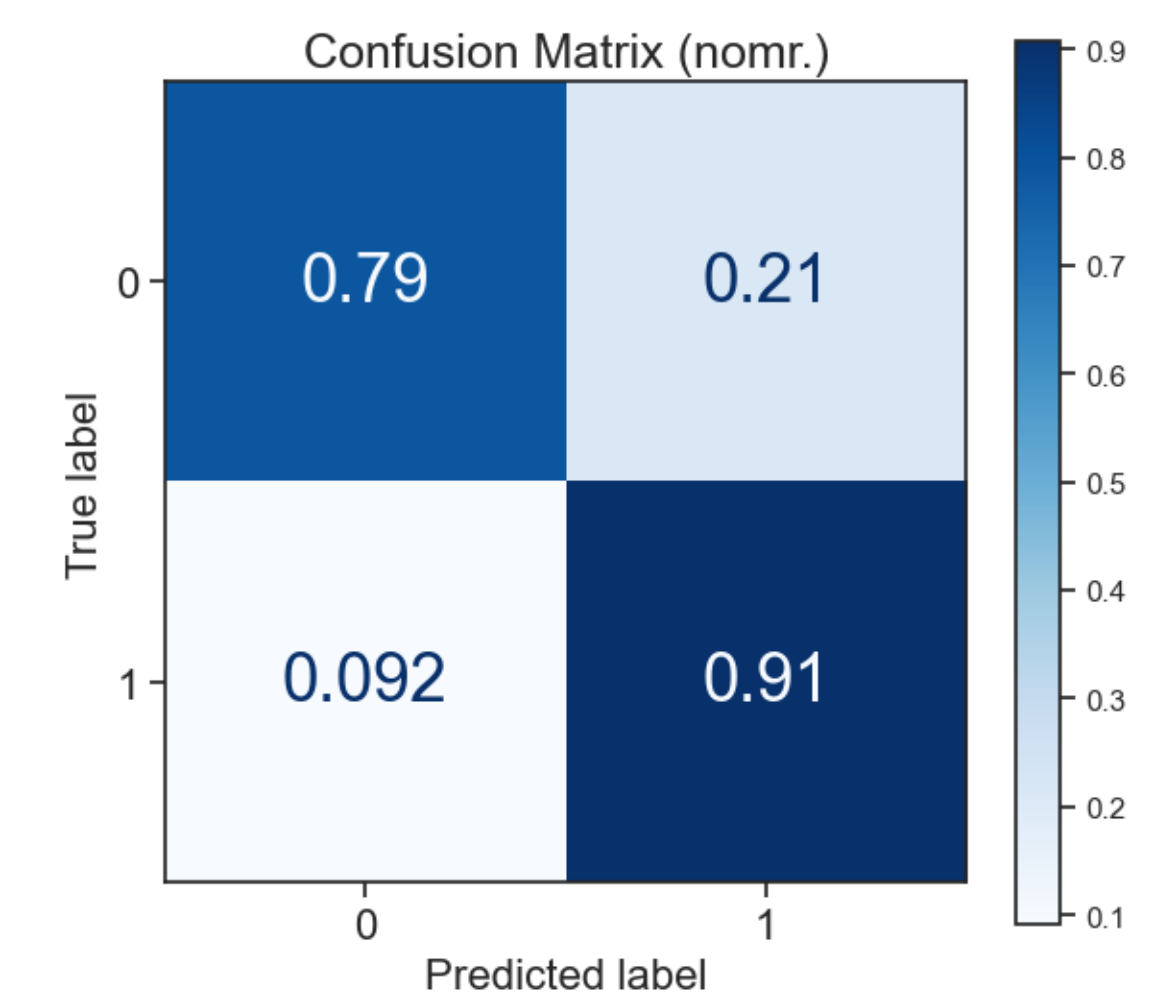
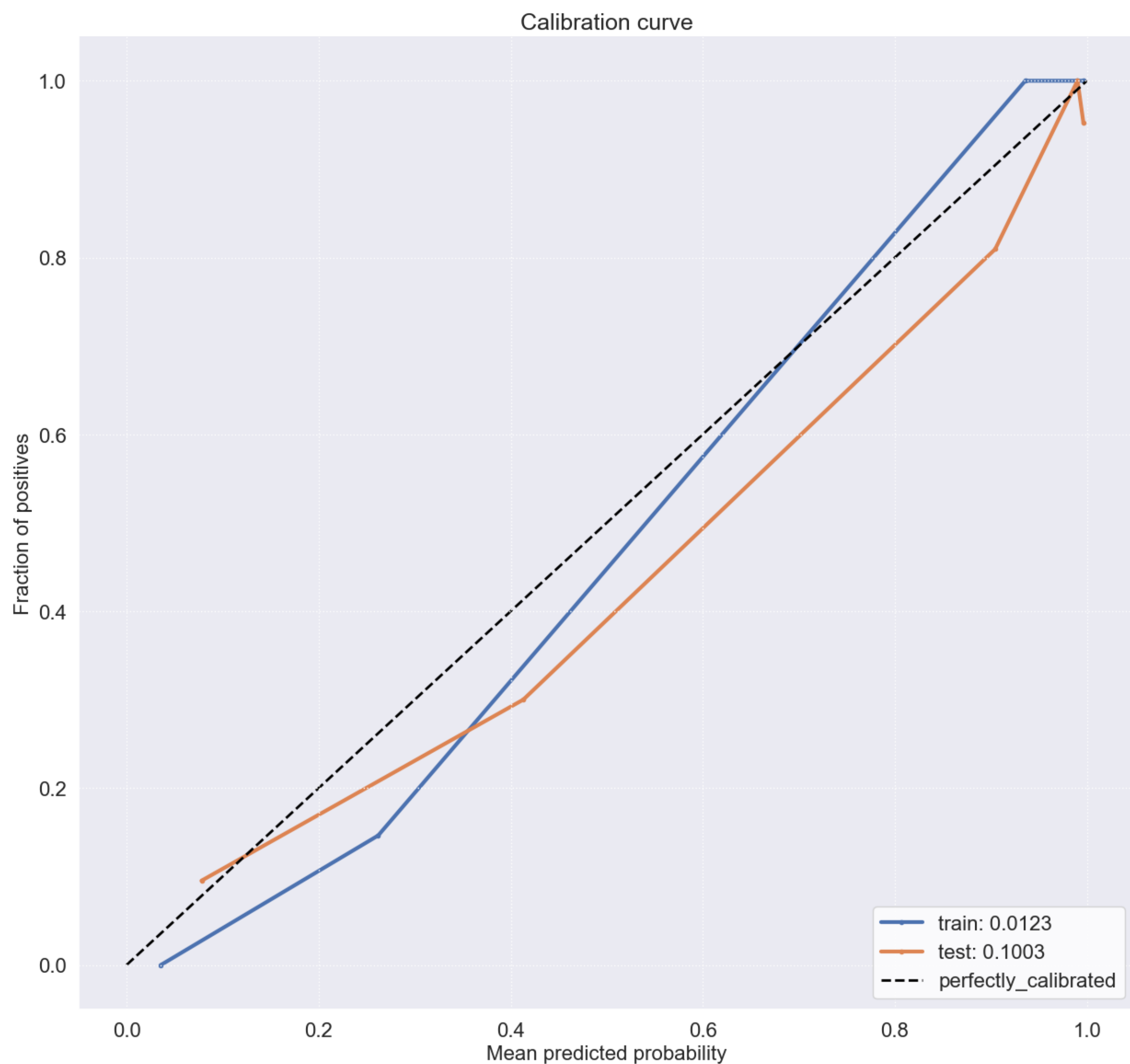
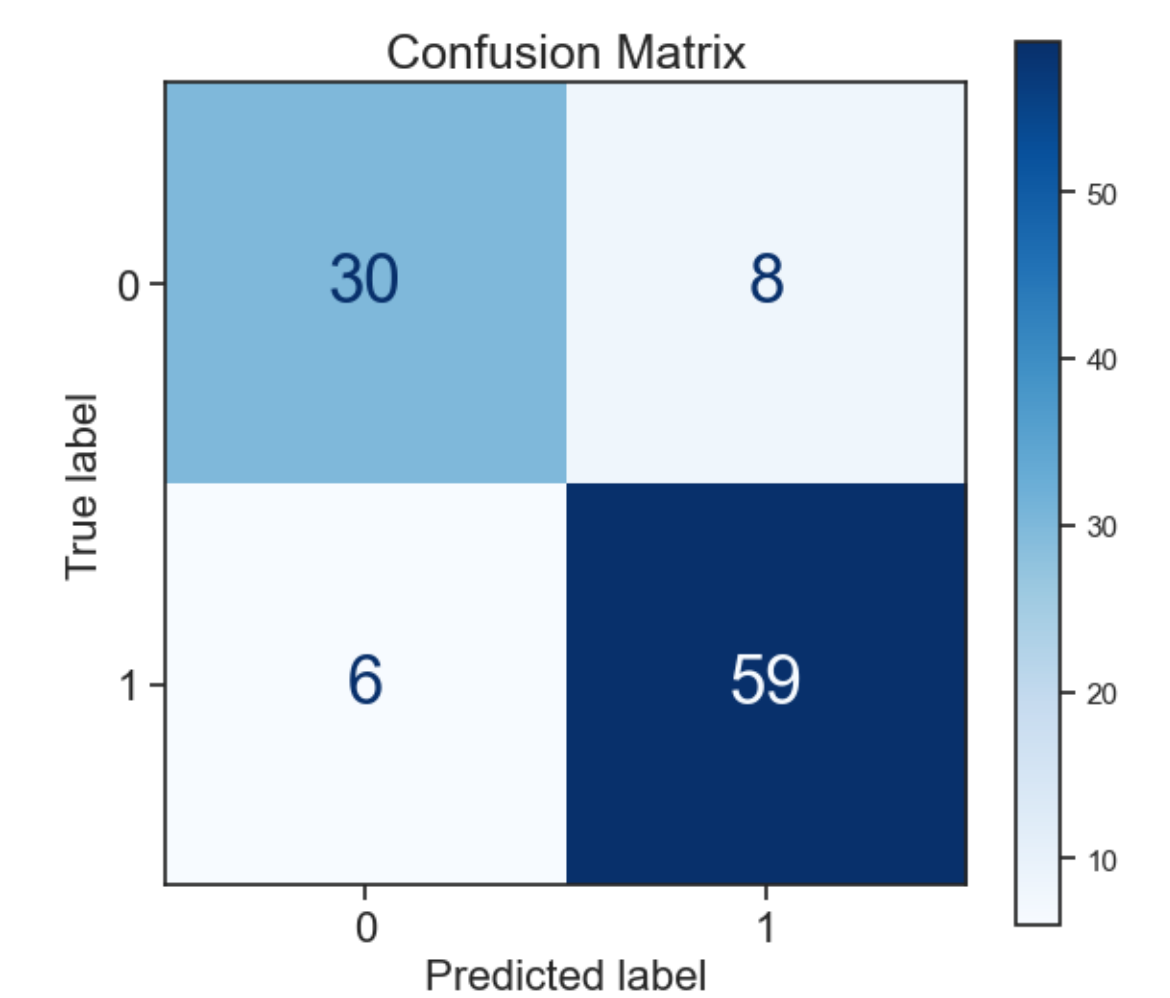
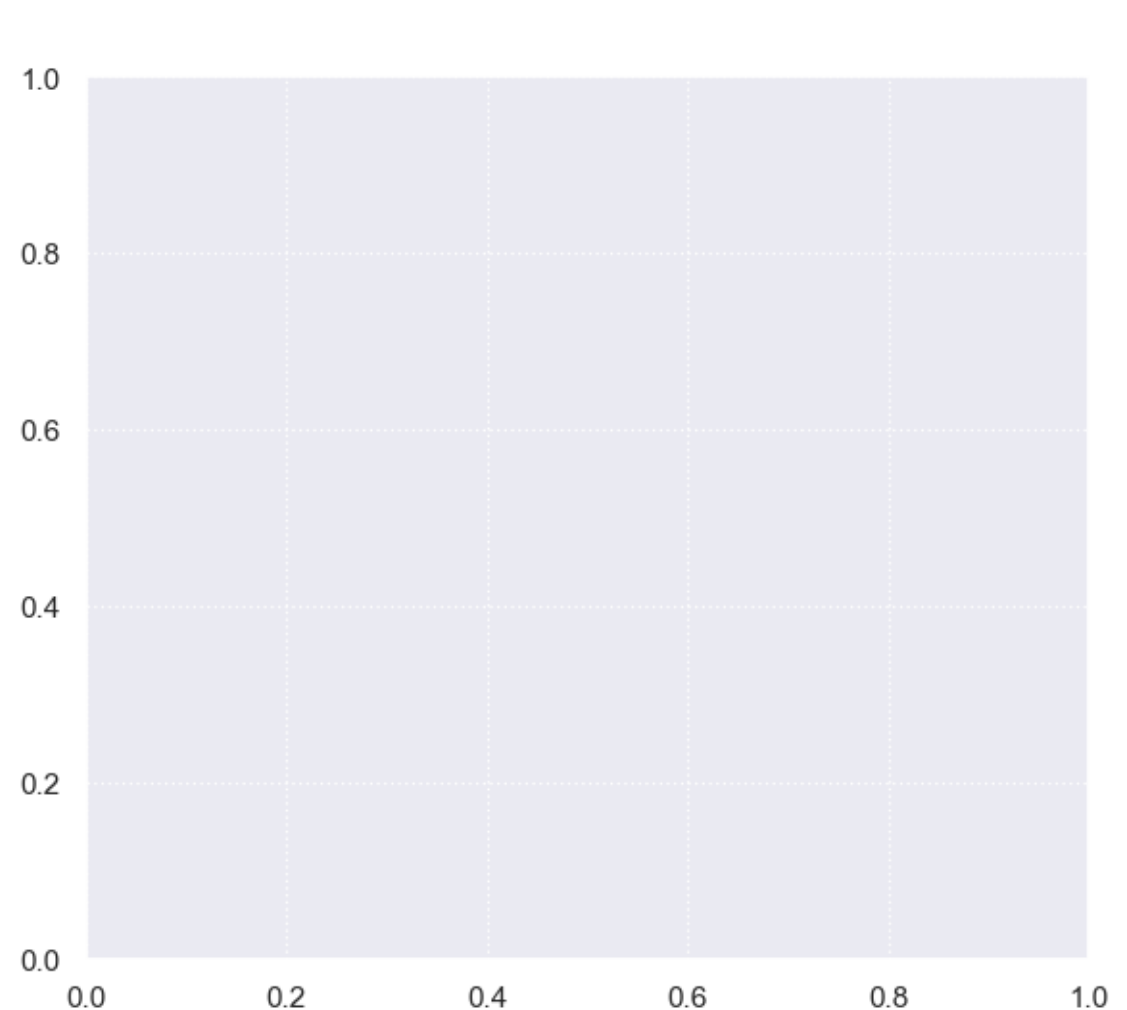
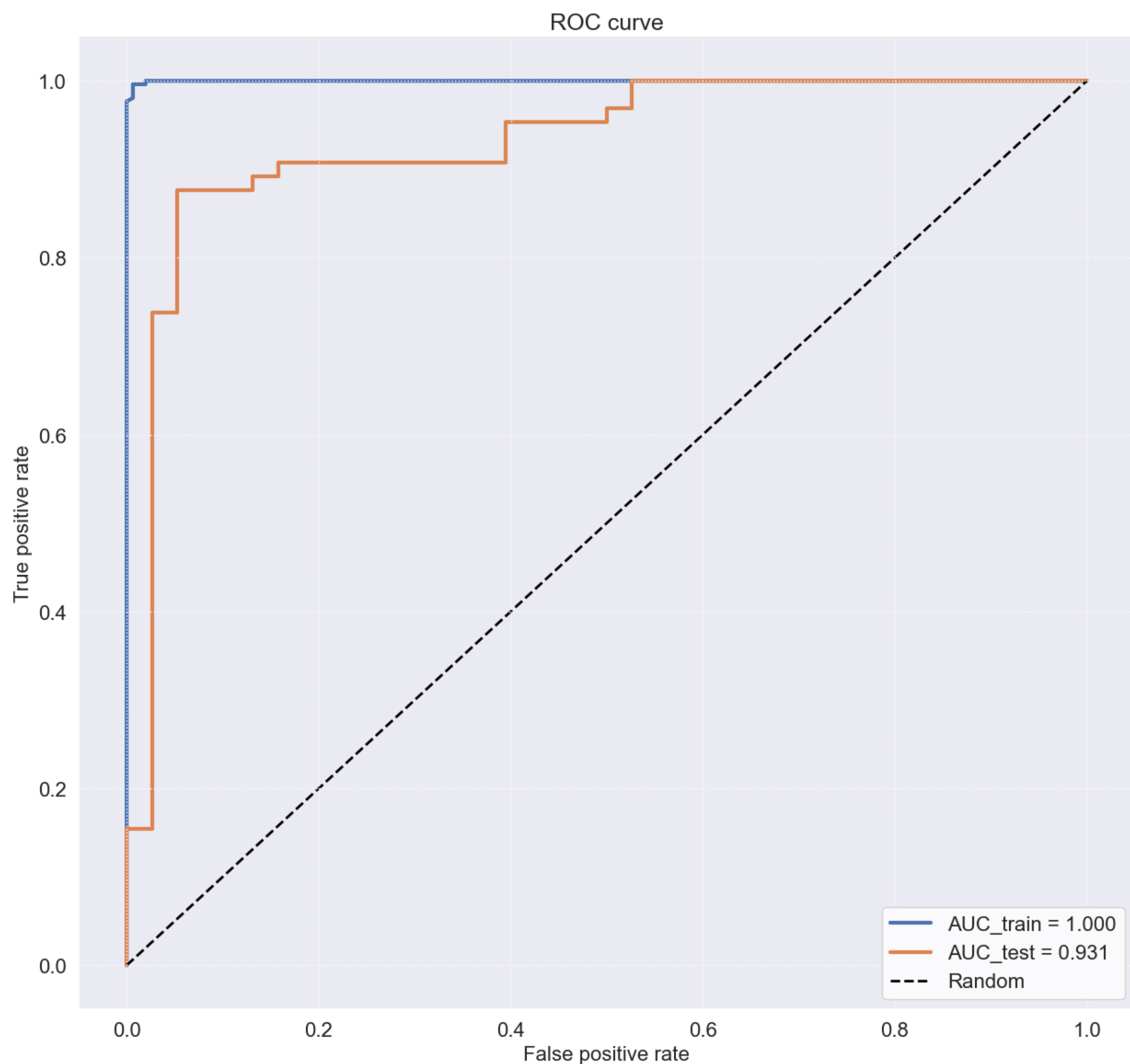
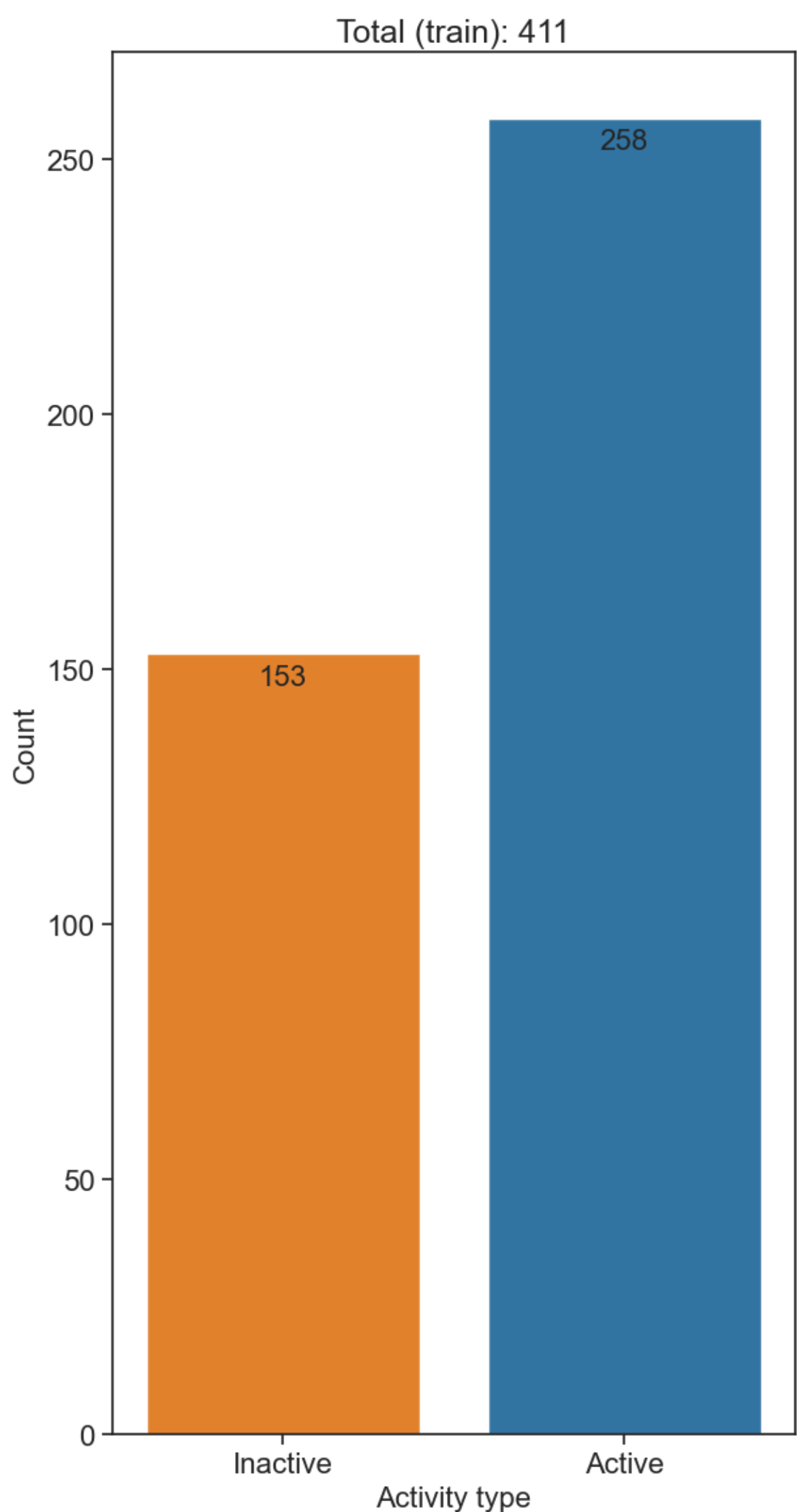
| XGBoost Parameters | Value  |
|--------------------|--------|
| alpha              | 2.728  |
| colsample_bytree   | 0.7487 |
| gamma              | 0.5311 |
| lambda             | 3.6587 |
| learning_rate      | 0.1589 |
| max_depth          | 5.0    |
| min_child_weight   | 1.0    |
| n_estimators       | 975.0  |
| reg_alpha          | 0.3405 |
| reg_lambda         | 0.8213 |
| scale_pos_weight   | 2.2065 |
| subsample          | 0.6879 |



| Metric            | Train  | Test   |
|-------------------|--------|--------|
| AUC               | 0.9999 | 0.9344 |
| accuracy          | 0.9927 | 0.8155 |
| balanced accuracy | 0.9902 | 0.7828 |
| recall_(sens)     | 1.0    | 0.9077 |
| specificity       | 0.9804 | 0.6579 |
| precision         | 0.9885 | 0.8194 |
| f1_score          | 0.9927 | 0.8109 |



Protein: P06213 - metric: SoftVotingXGBoostEnsemble



| Metric            | Train  | Test   |
|-------------------|--------|--------|
| AUC               | 0.9998 | 0.9308 |
| accuracy          | 0.9903 | 0.8641 |
| balanced accuracy | 0.9869 | 0.8486 |
| recall_(sens)     | 1.0    | 0.9077 |
| specificity       | 0.9739 | 0.7895 |
| precision         | 0.9847 | 0.8806 |
| f1_score          | 0.9902 | 0.8633 |