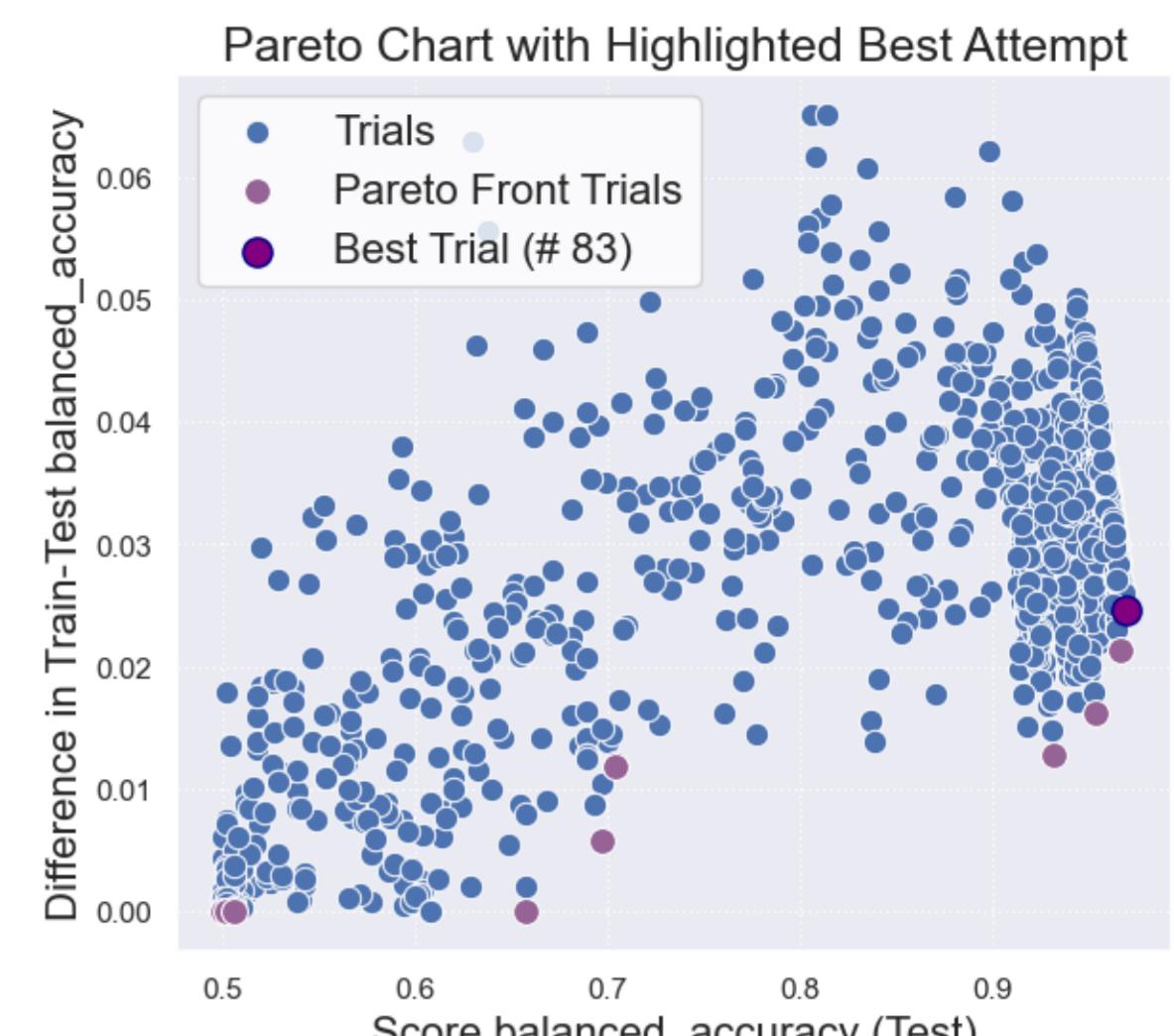
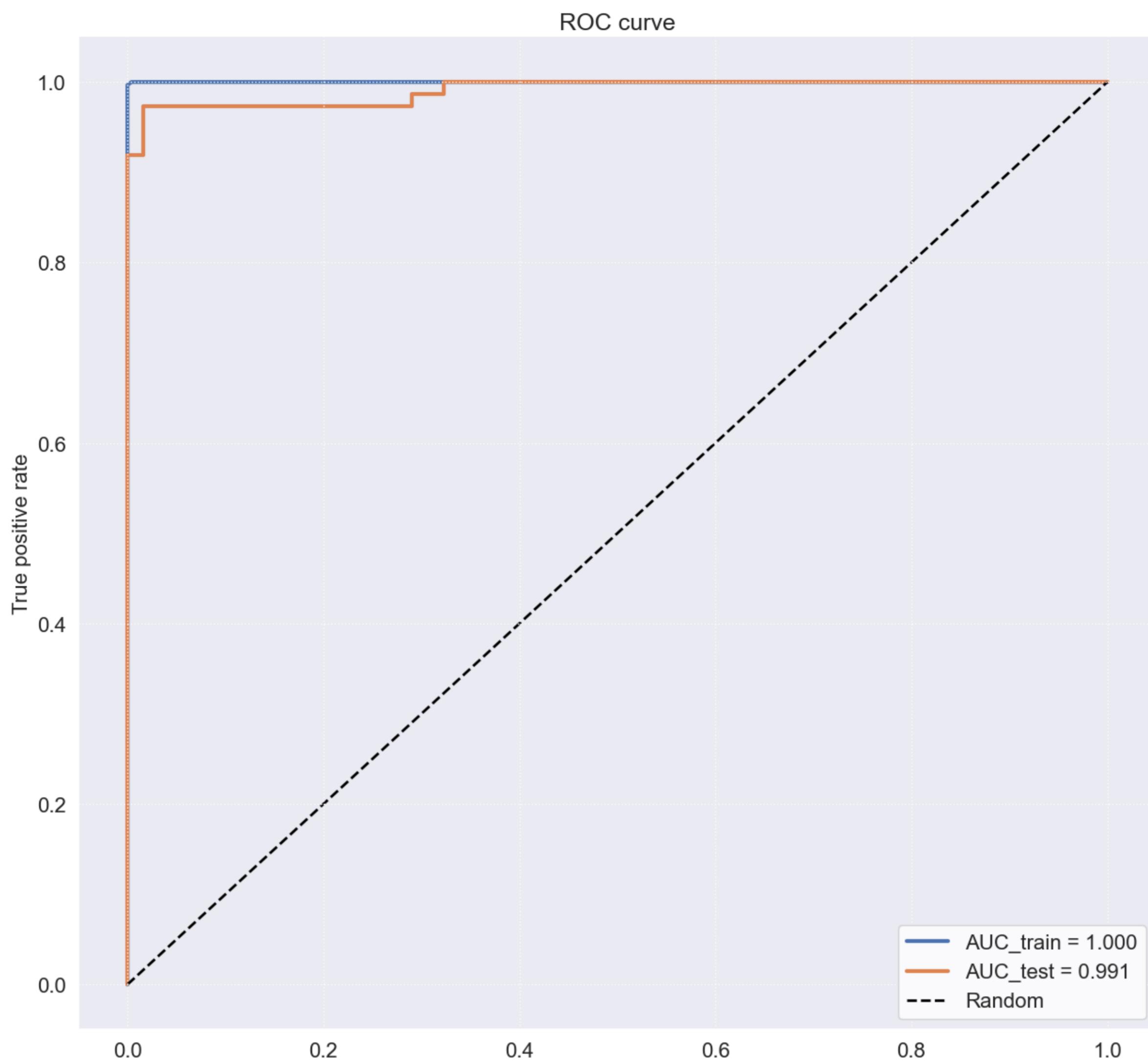
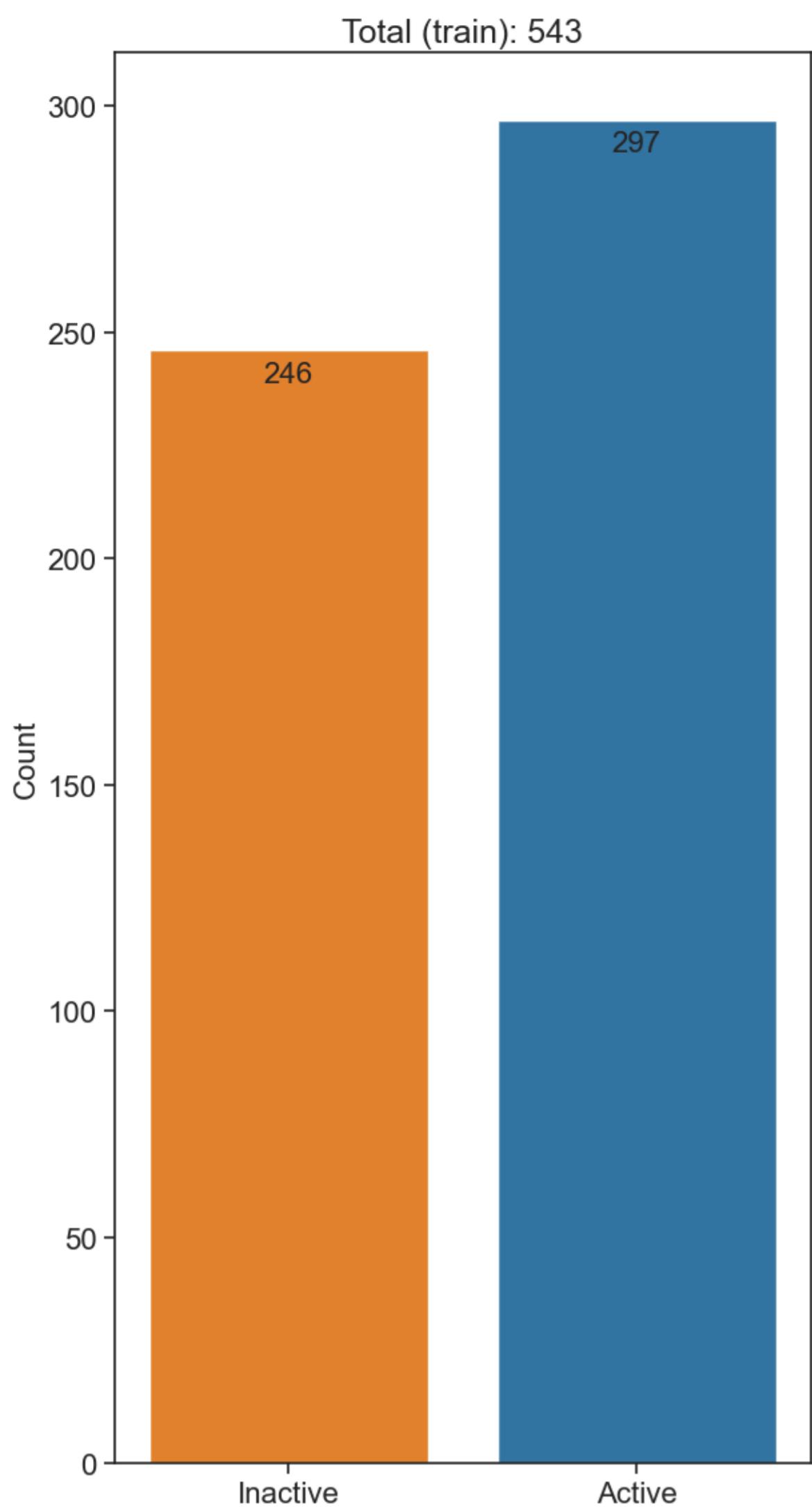
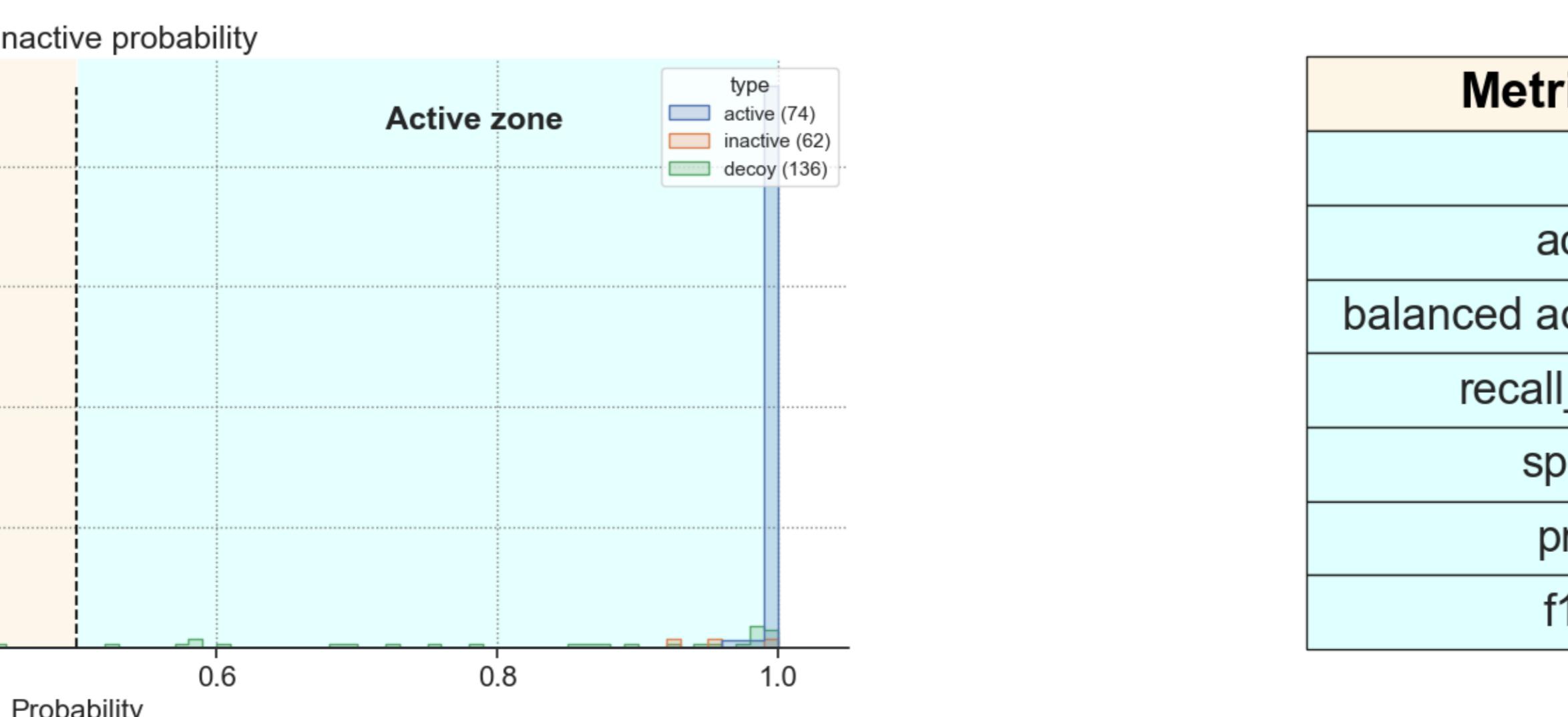
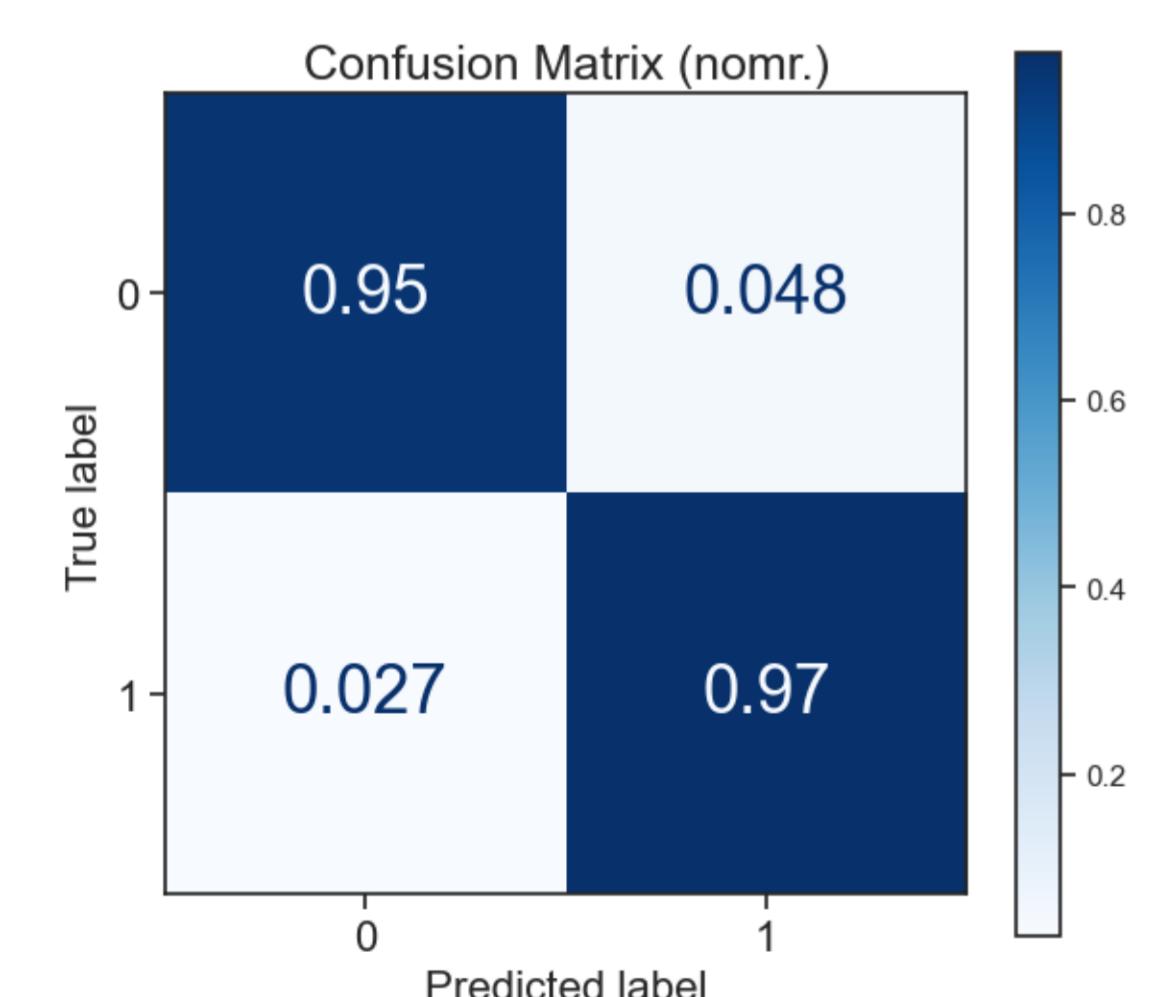
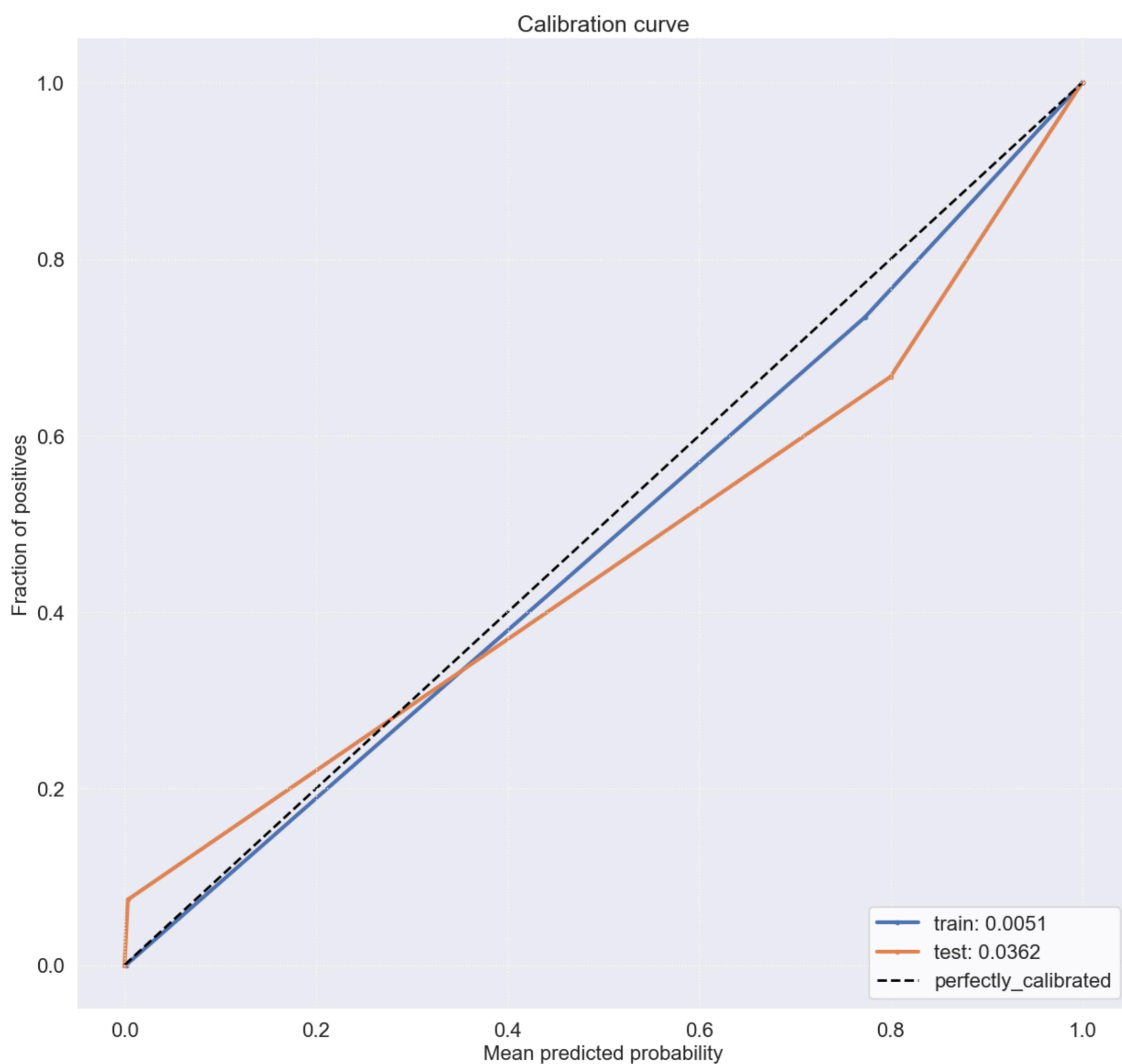
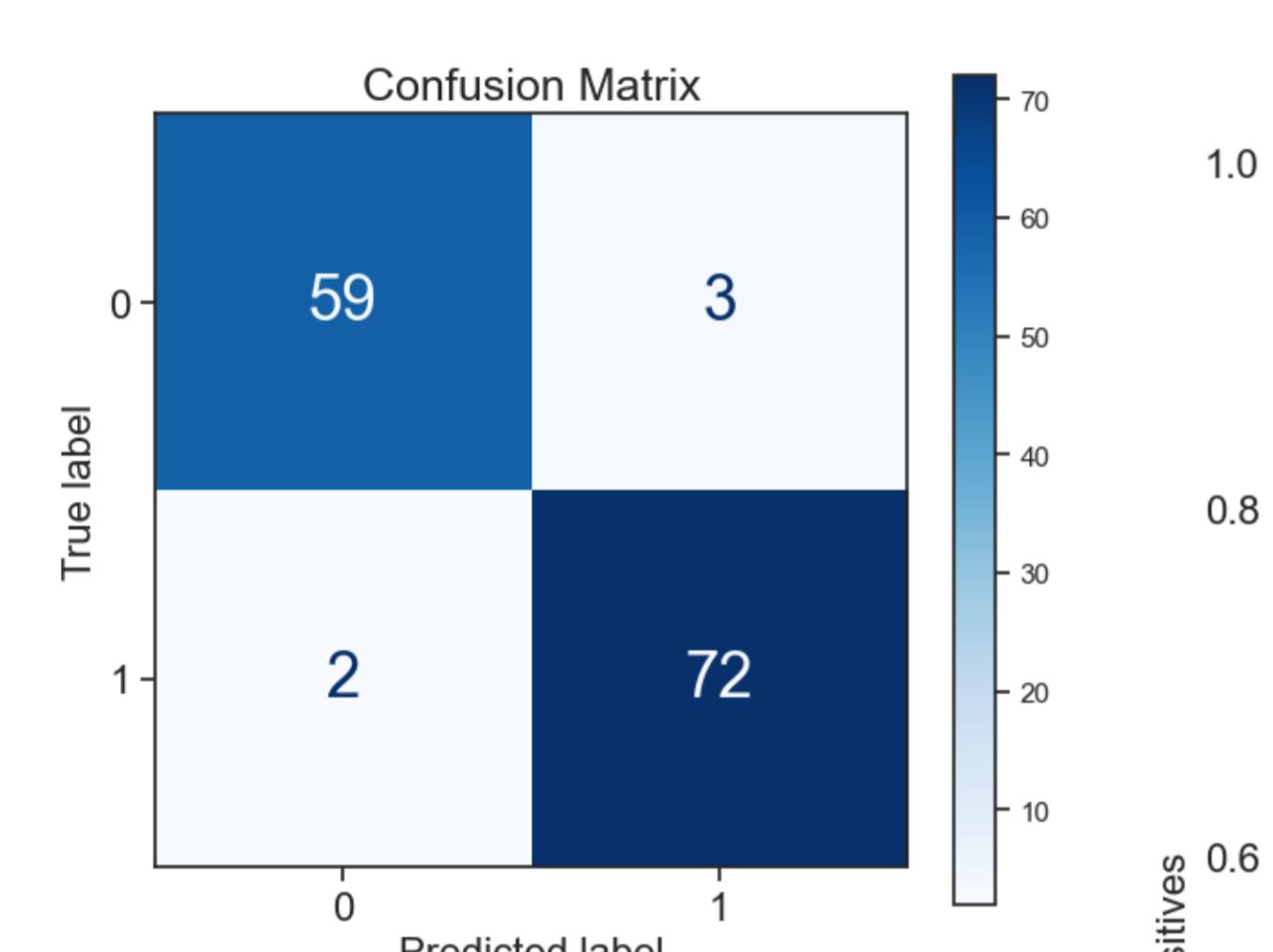


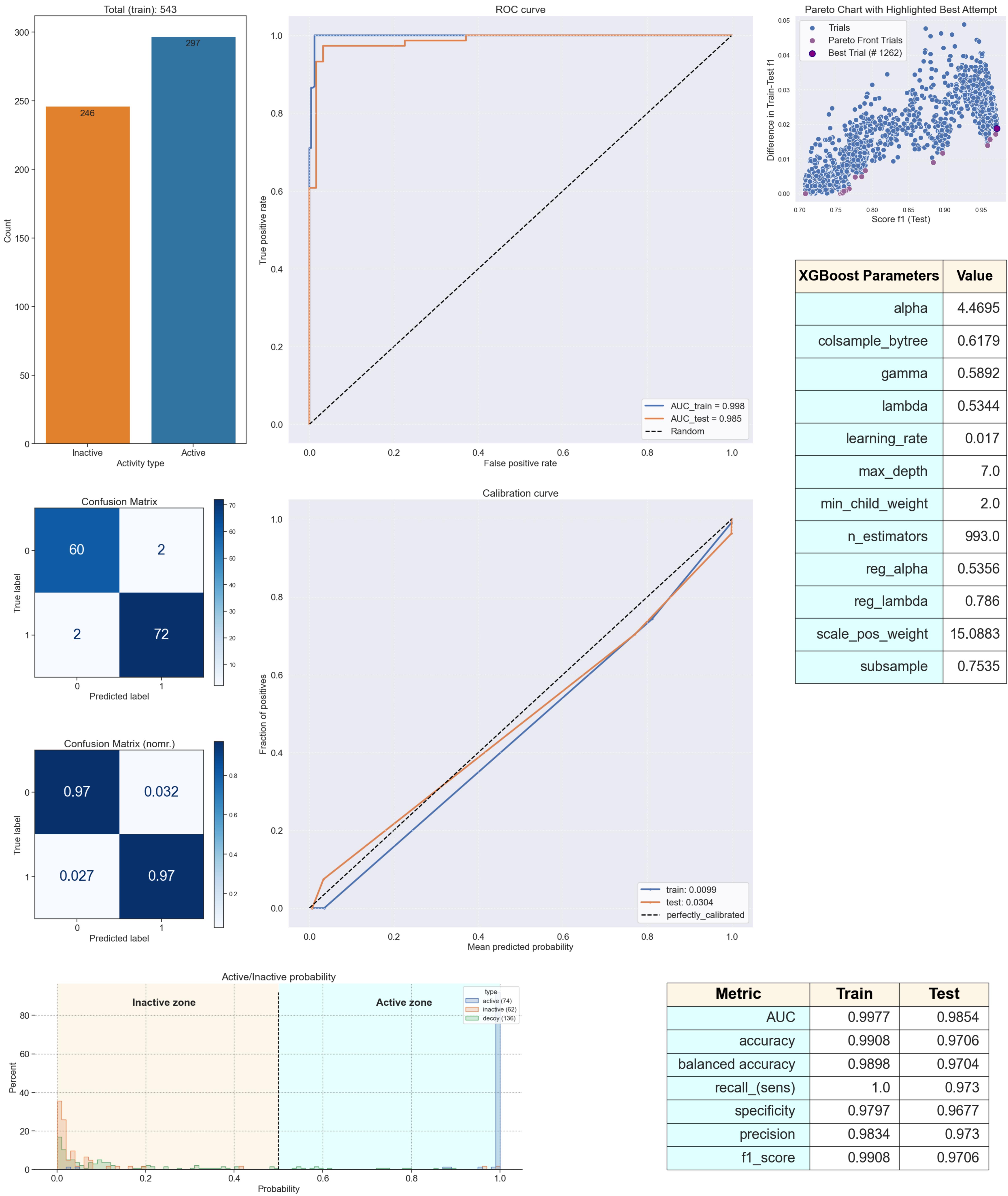
Protein: P01375 - metric: balanced_accuracy



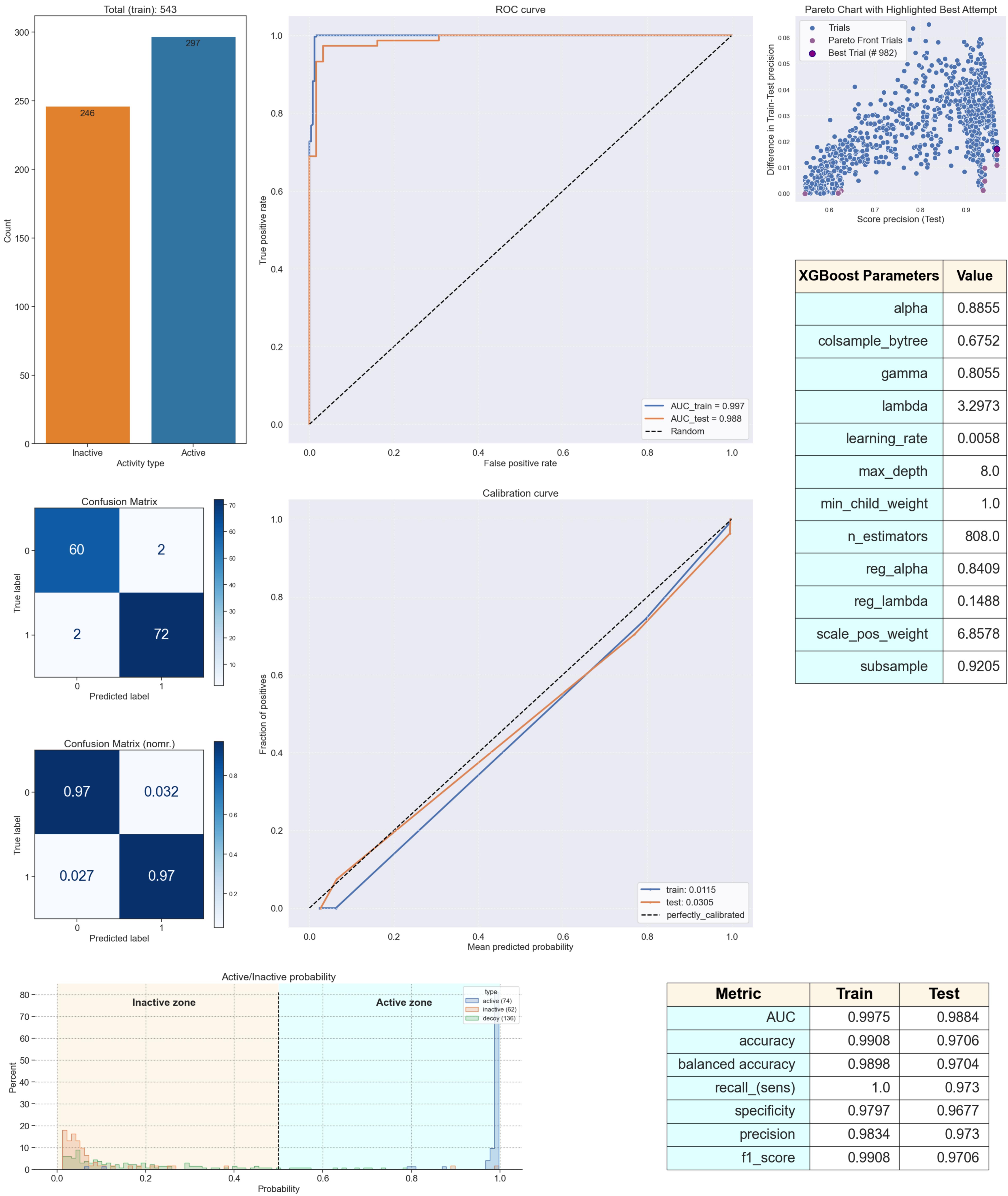
XGBoost Parameters	Value
alpha	1.0303
colsample_bytree	0.8763
gamma	0.1703
lambda	1.0852
learning_rate	0.1909
max_depth	9.0
min_child_weight	1.0
n_estimators	986.0
reg_alpha	0.4337
reg_lambda	0.2819
scale_pos_weight	30.4816
subsample	0.5495



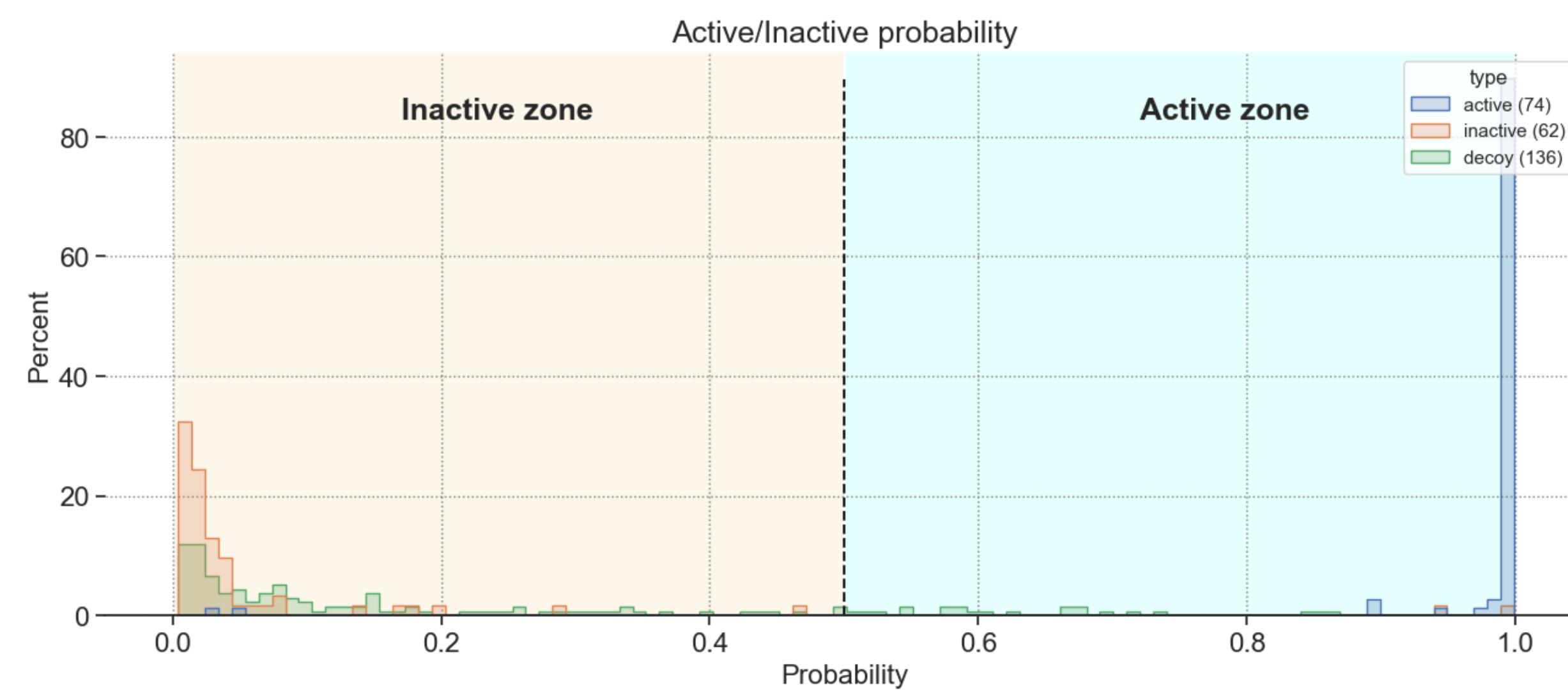
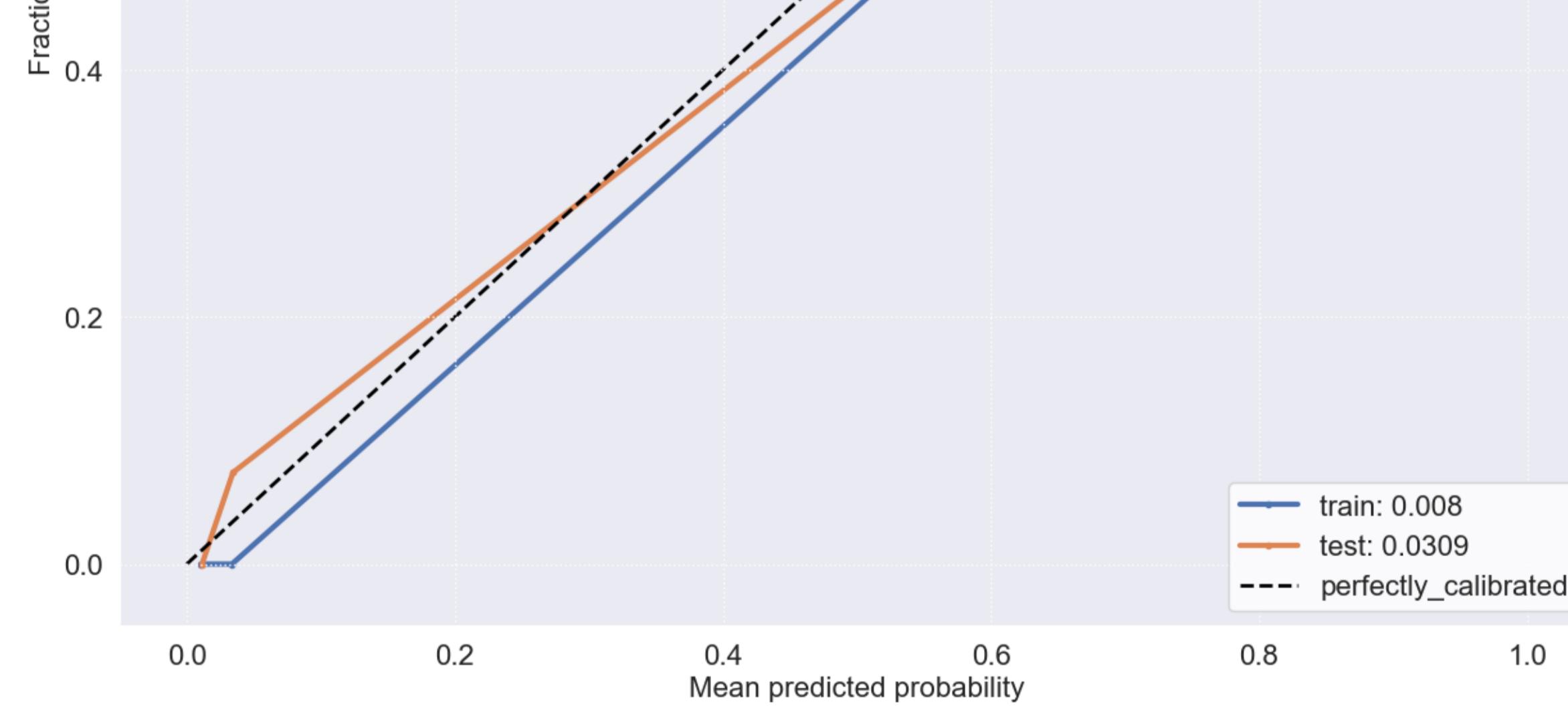
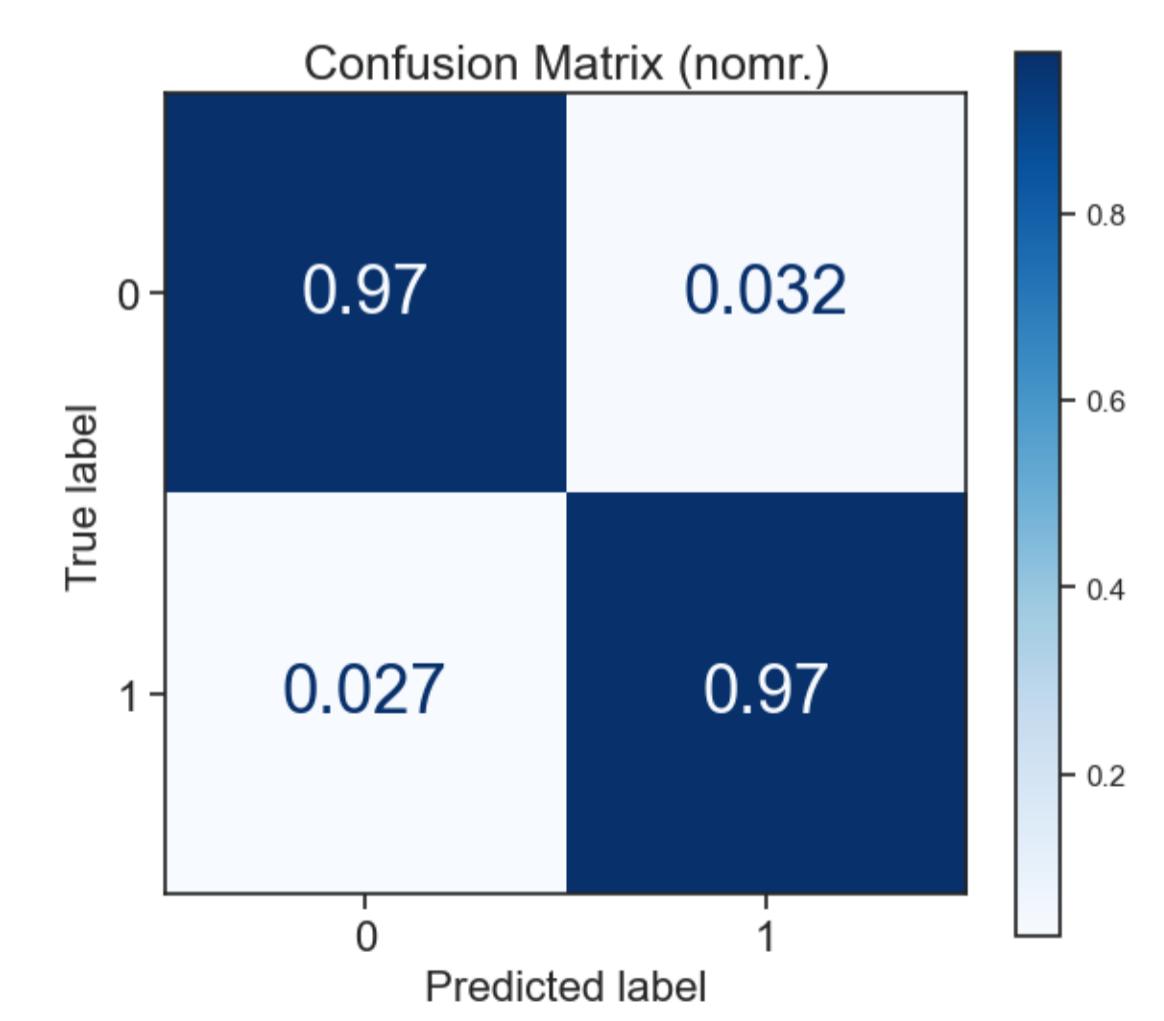
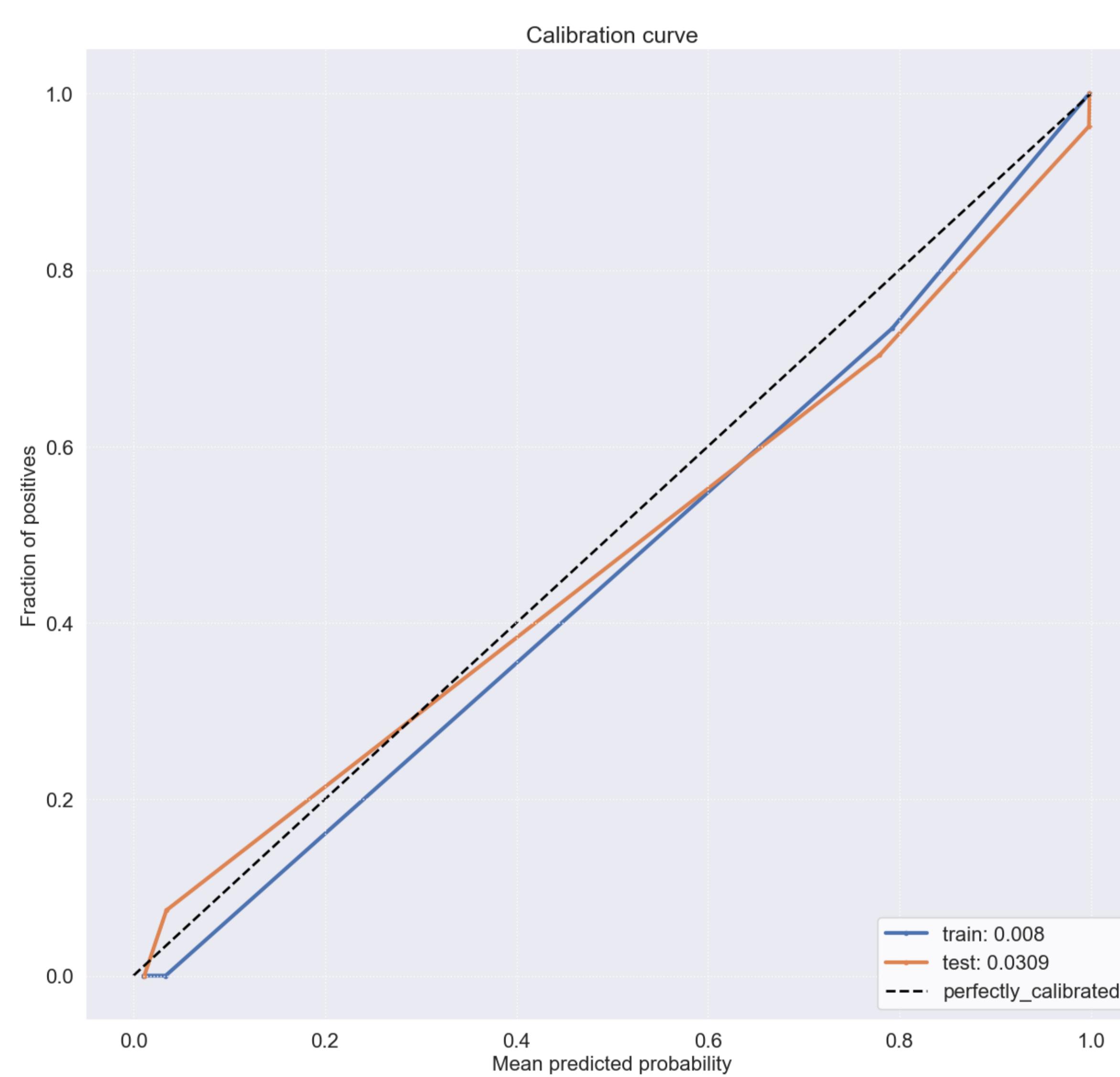
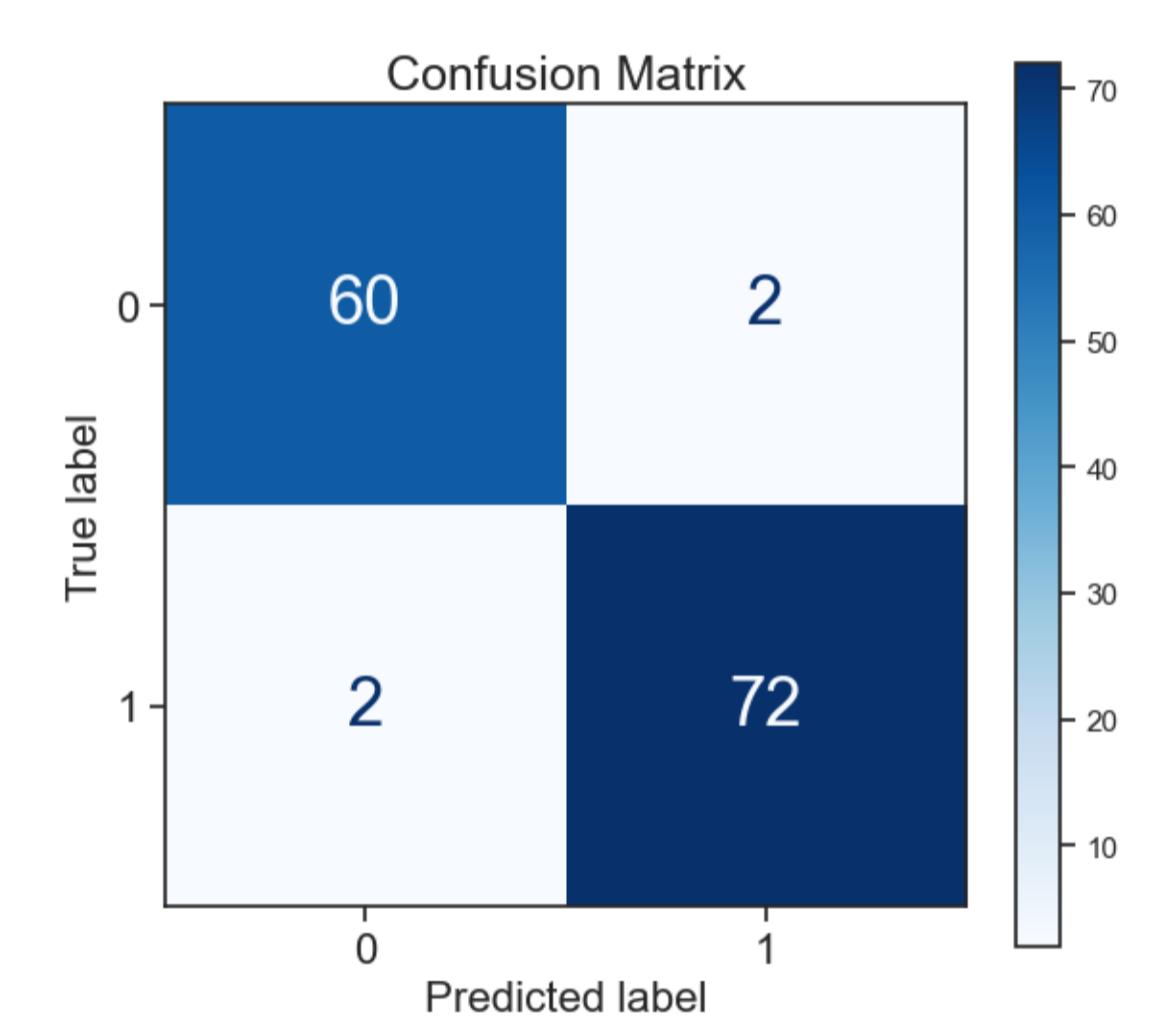
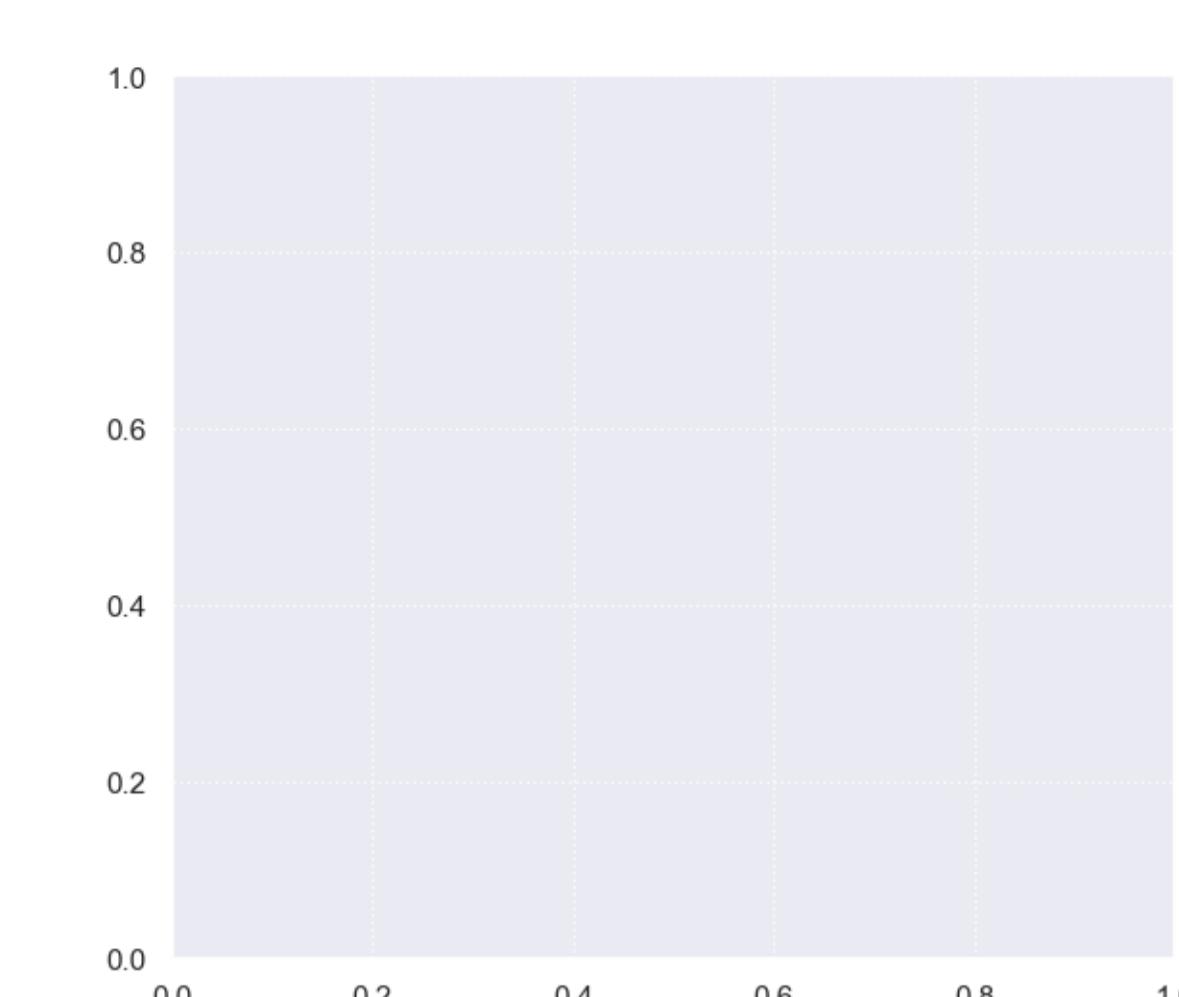
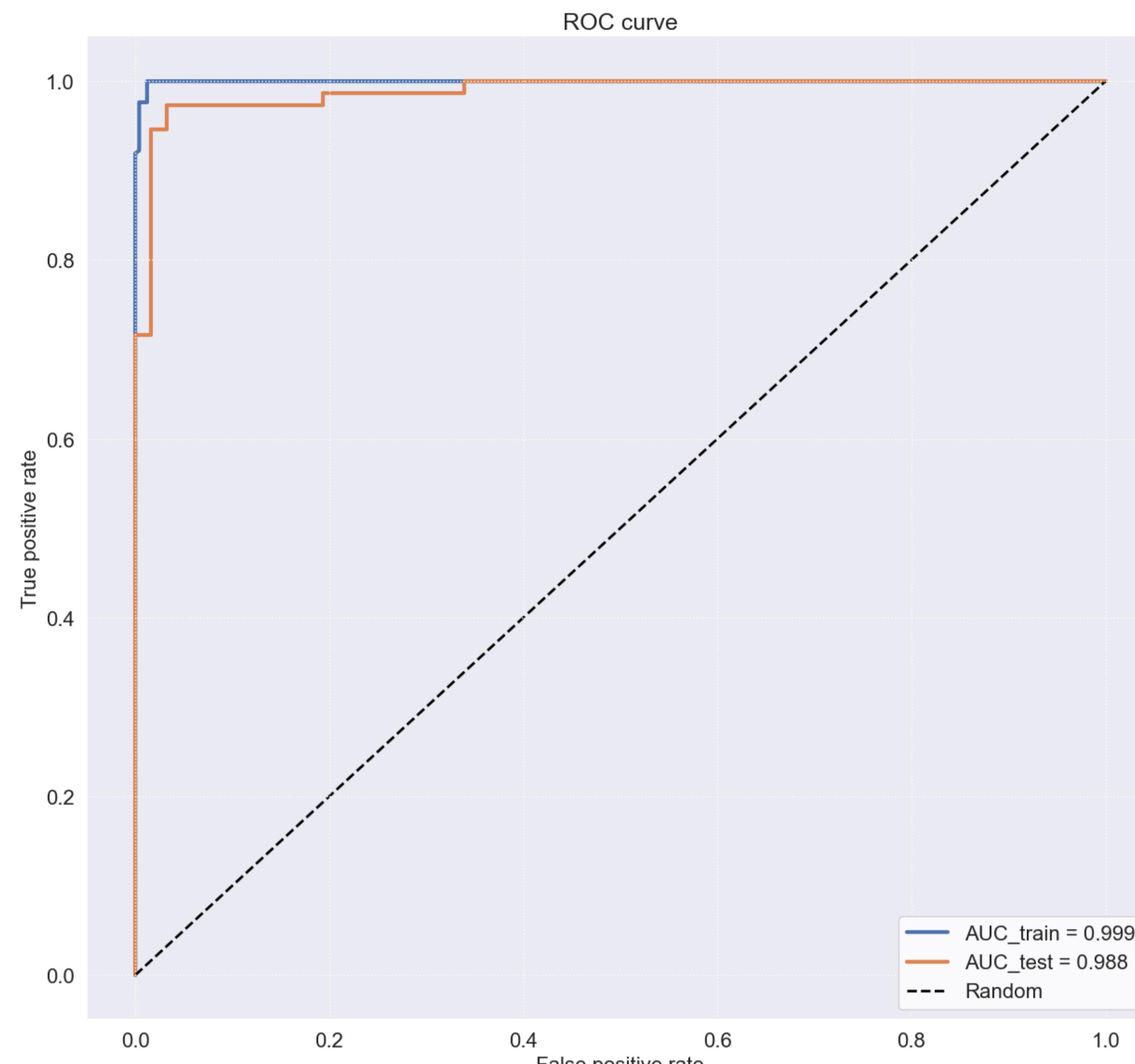
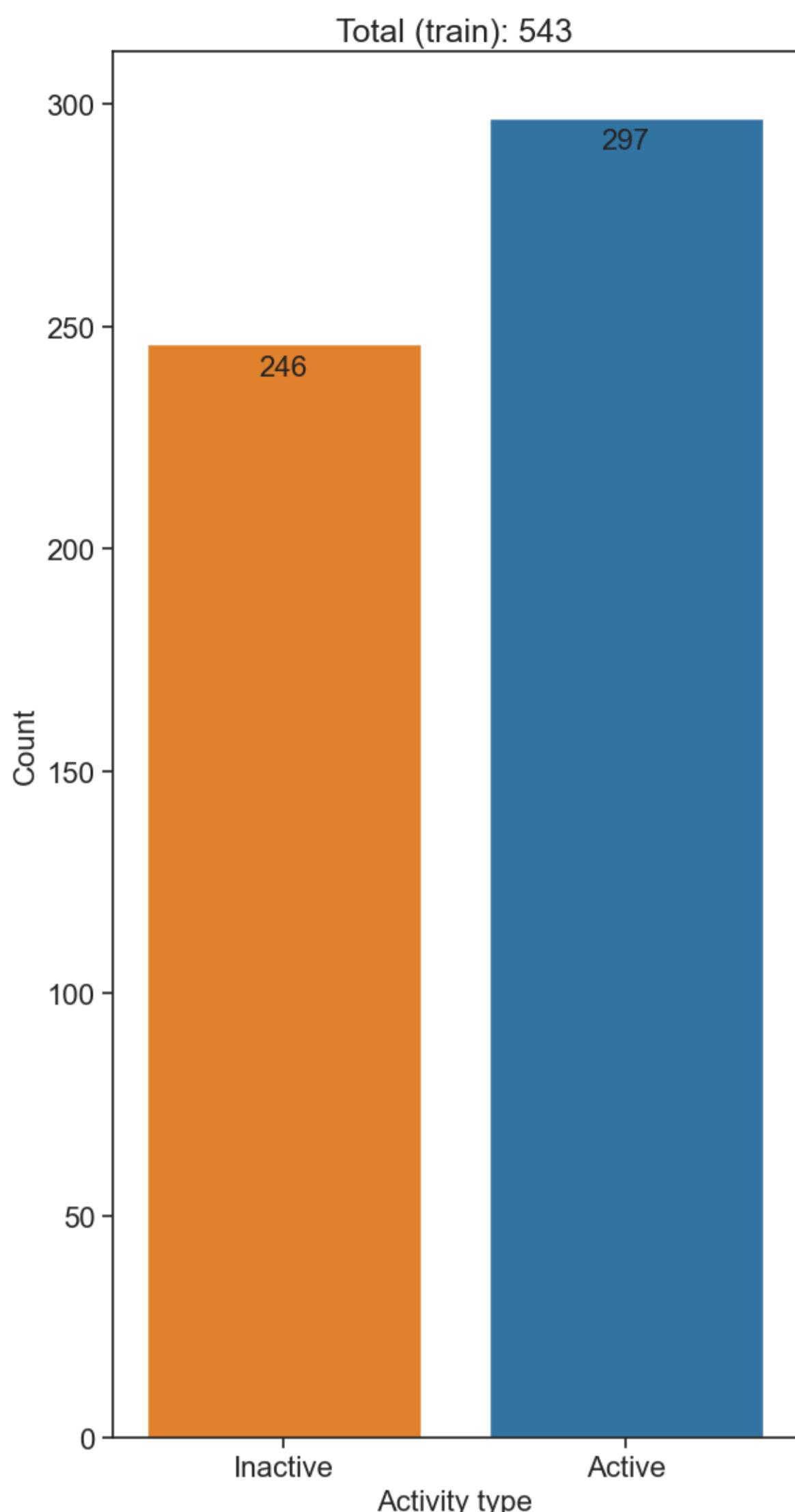
Protein: P01375 - metric: f1



Protein: P01375 - metric: precision



Protein: P01375 - metric: SoftVotingXGBoostEnsemble



Metric	Train	Test
AUC	0.9995	0.9882
accuracy	0.9926	0.9706
balanced accuracy	0.9919	0.9704
recall_(sens)	1.0	0.973
specificity	0.9837	0.9677
precision	0.9867	0.973
f1_score	0.9926	0.9706