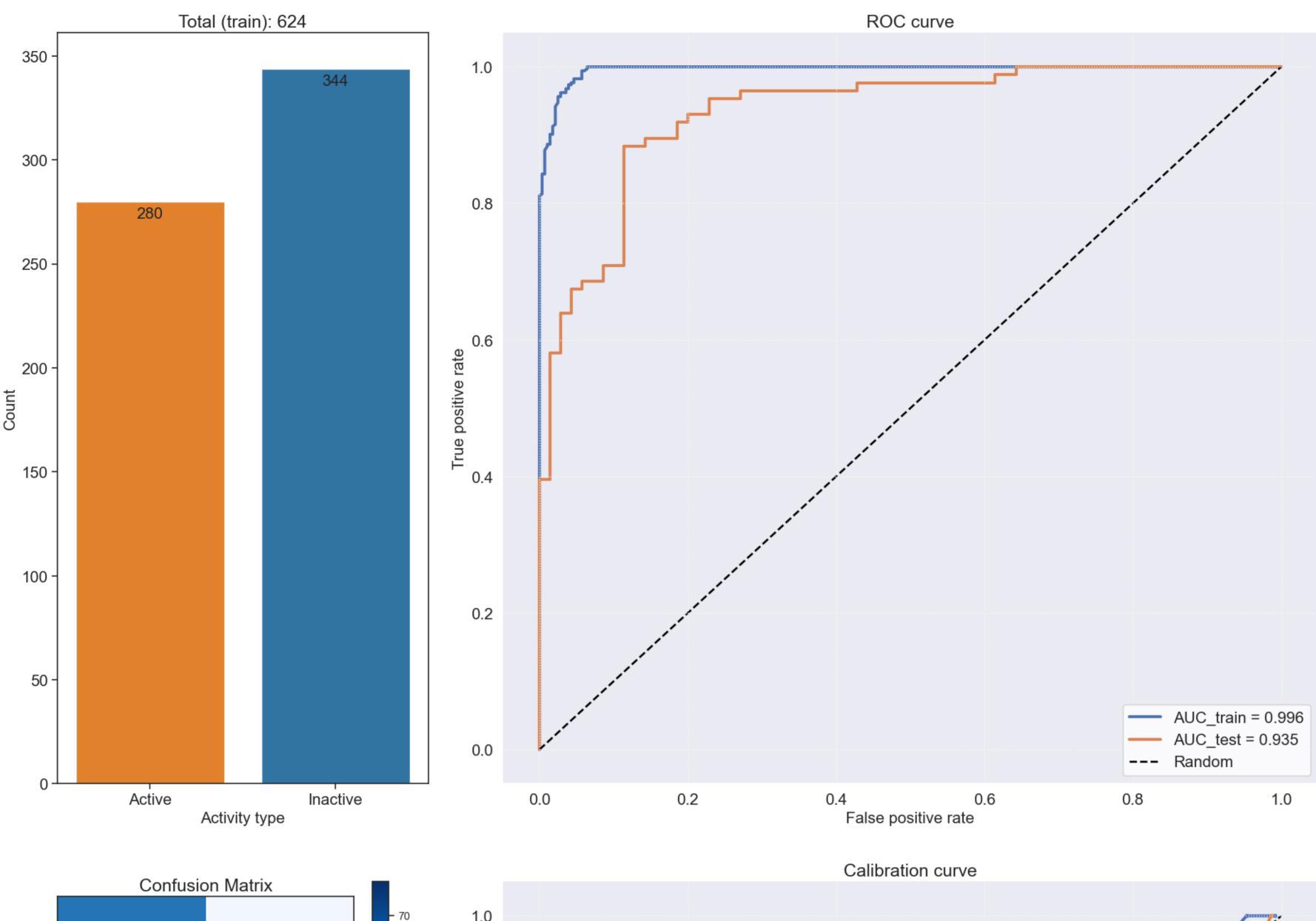
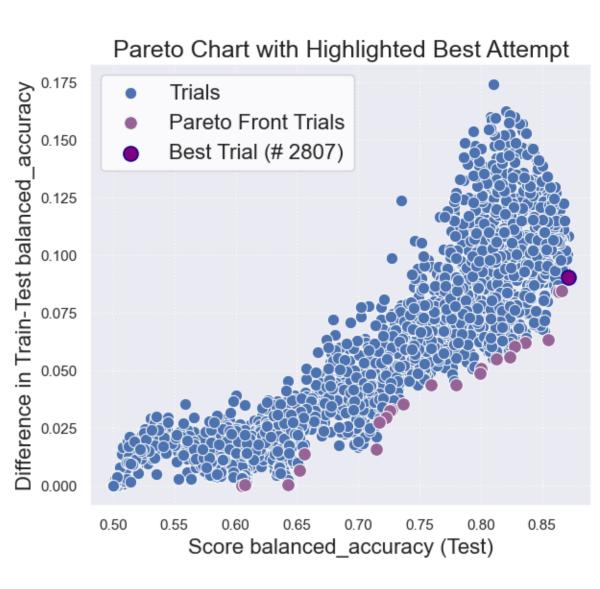
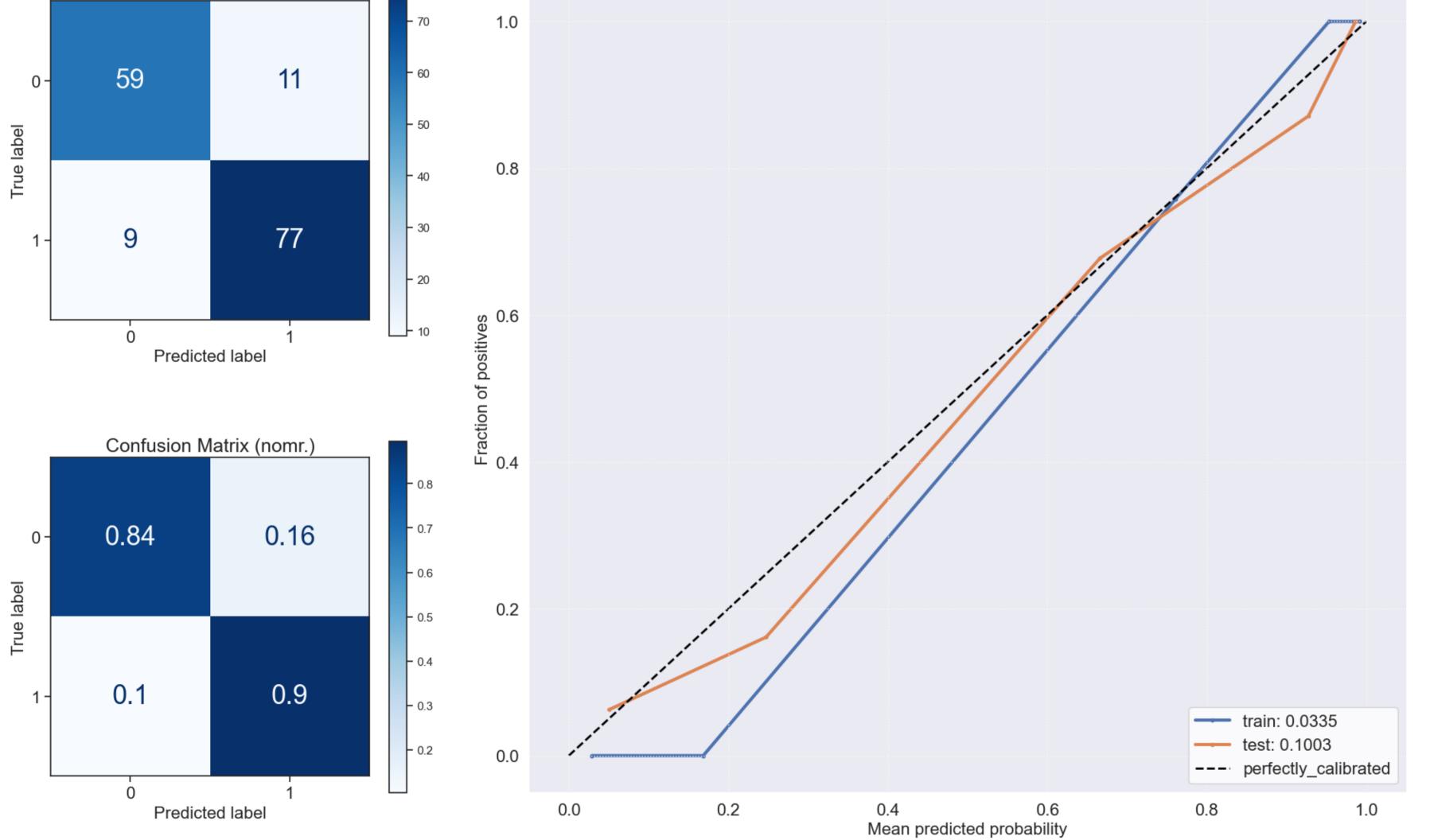
Protein: P48039 - metric: balanced_accuracy





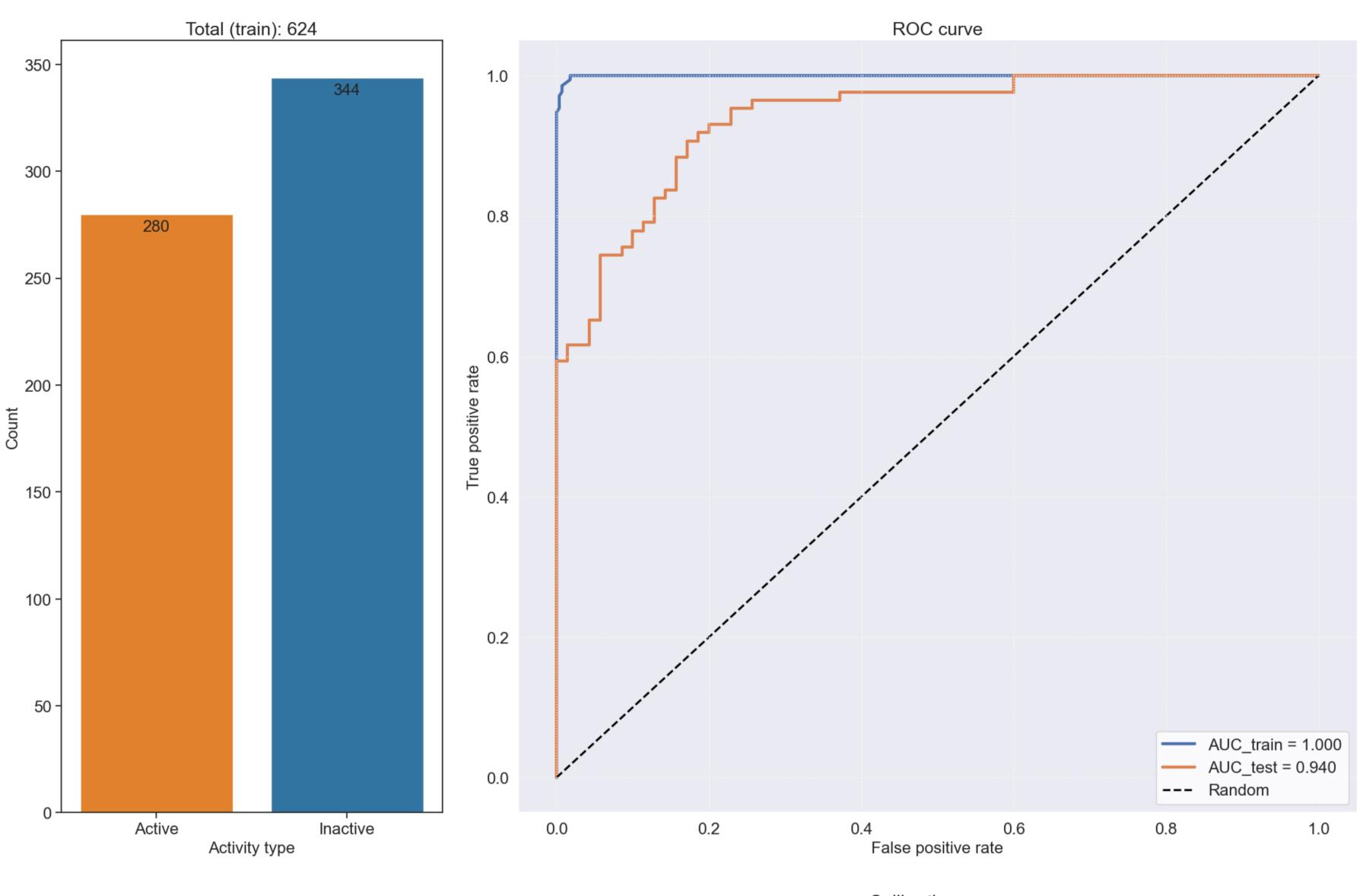
XGBoost Parameters	Value
alpha	0.862
colsample_bytree	0.5079
gamma	0.9621
lambda	2.5963
learning_rate	0.0384
max_depth	9.0
min_child_weight	1.0
n_estimators	968.0
reg_alpha	0.0176
reg_lambda	0.8057
scale_pos_weight	1.7173
subsample	0.8527

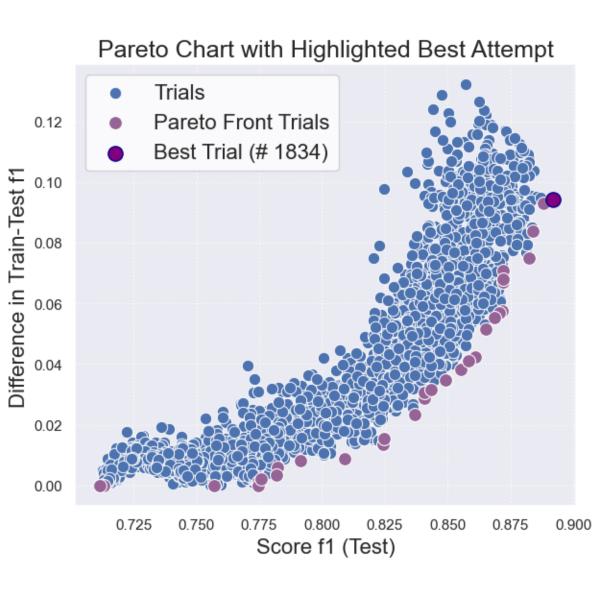


				Active/Ir	nactive prob	oability		
20		Ina	ctive zone				Active zone	type active (86) inactive (70) decoy (156)
Percent								
± 10 5								
0-								
Ŭ	0.0	0	.2	0.4	Probability	0.6	0.8	1.0

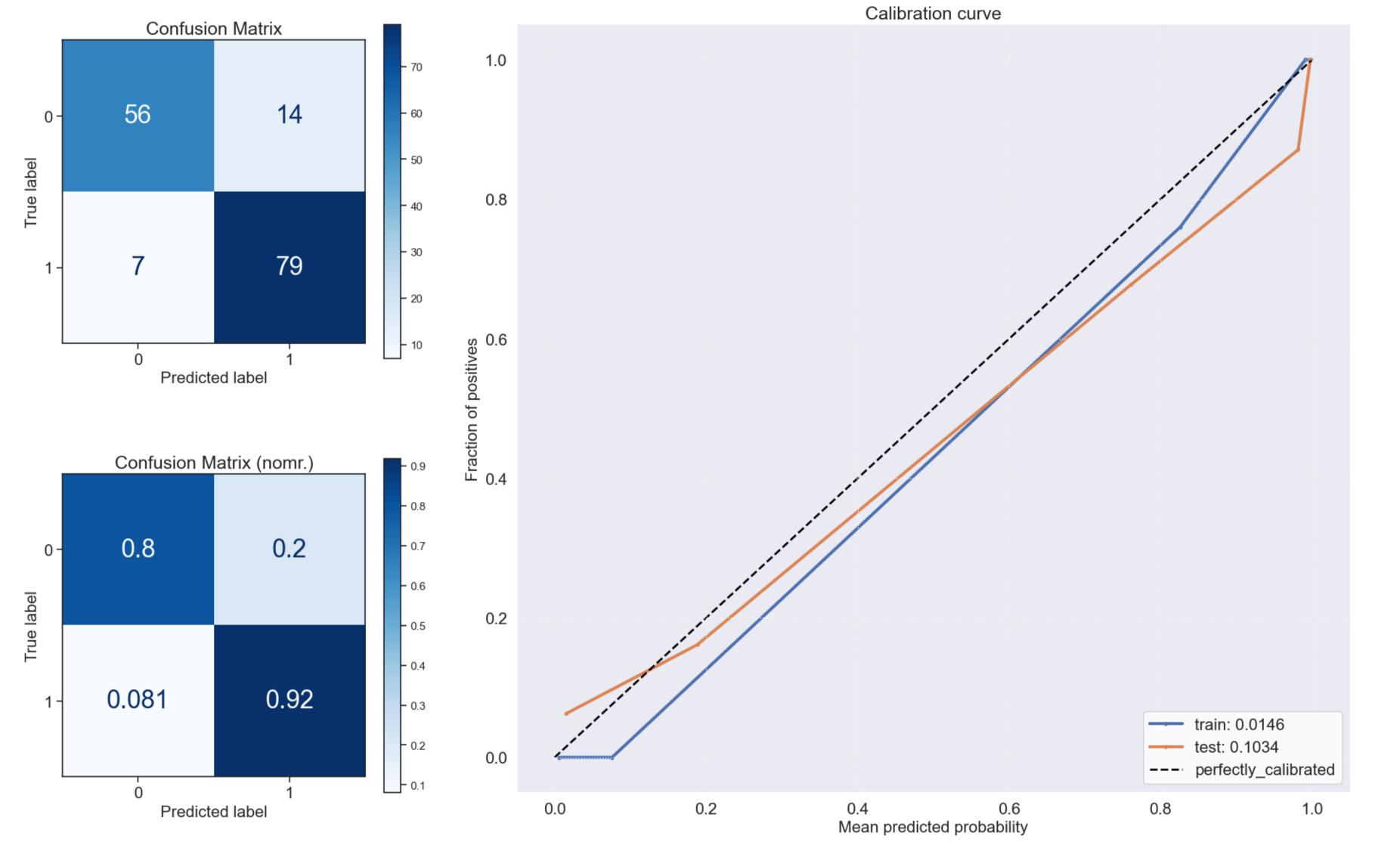
Metric	Train	Test
AUC	0.9961	0.9352
accuracy	0.9663	0.8718
balanced accuracy	0.9625	0.8691
recall_(sens)	1.0	0.8953
specificity	0.925	0.8429
precision	0.9425	0.875
f1_score	0.9662	0.8716

Protein: P48039 - metric: f1





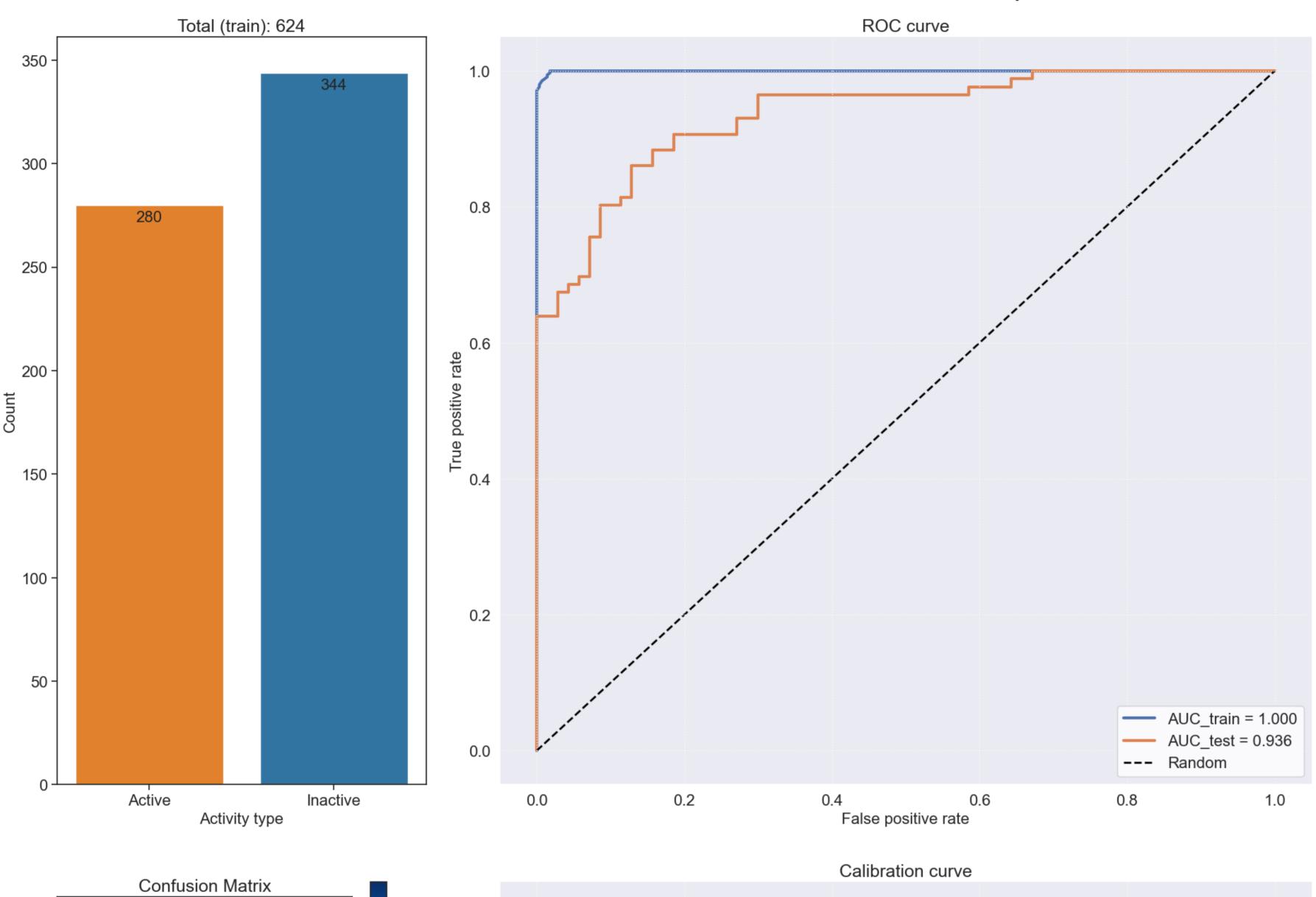
XGBoost Parameters	Value
alpha	0.29
colsample_bytree	0.9376
gamma	0.1099
lambda	3.1744
learning_rate	0.1354
max_depth	9.0
min_child_weight	1.0
n_estimators	414.0
reg_alpha	0.8461
reg_lambda	0.0783
scale_pos_weight	3.6378
subsample	0.8629

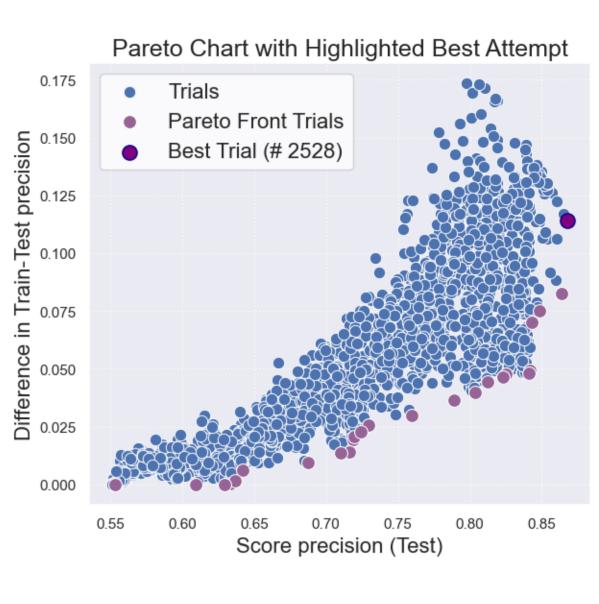


		Active/Inactive	probability			
60	Inactive zone	!		Active zone	active (86) inactive (70)	
50					decoy (156)	
40						
Percent 08						
20						
10						
0 —	0.0 0.2	0.4	0.6	0.8	1.0	
	Probability					

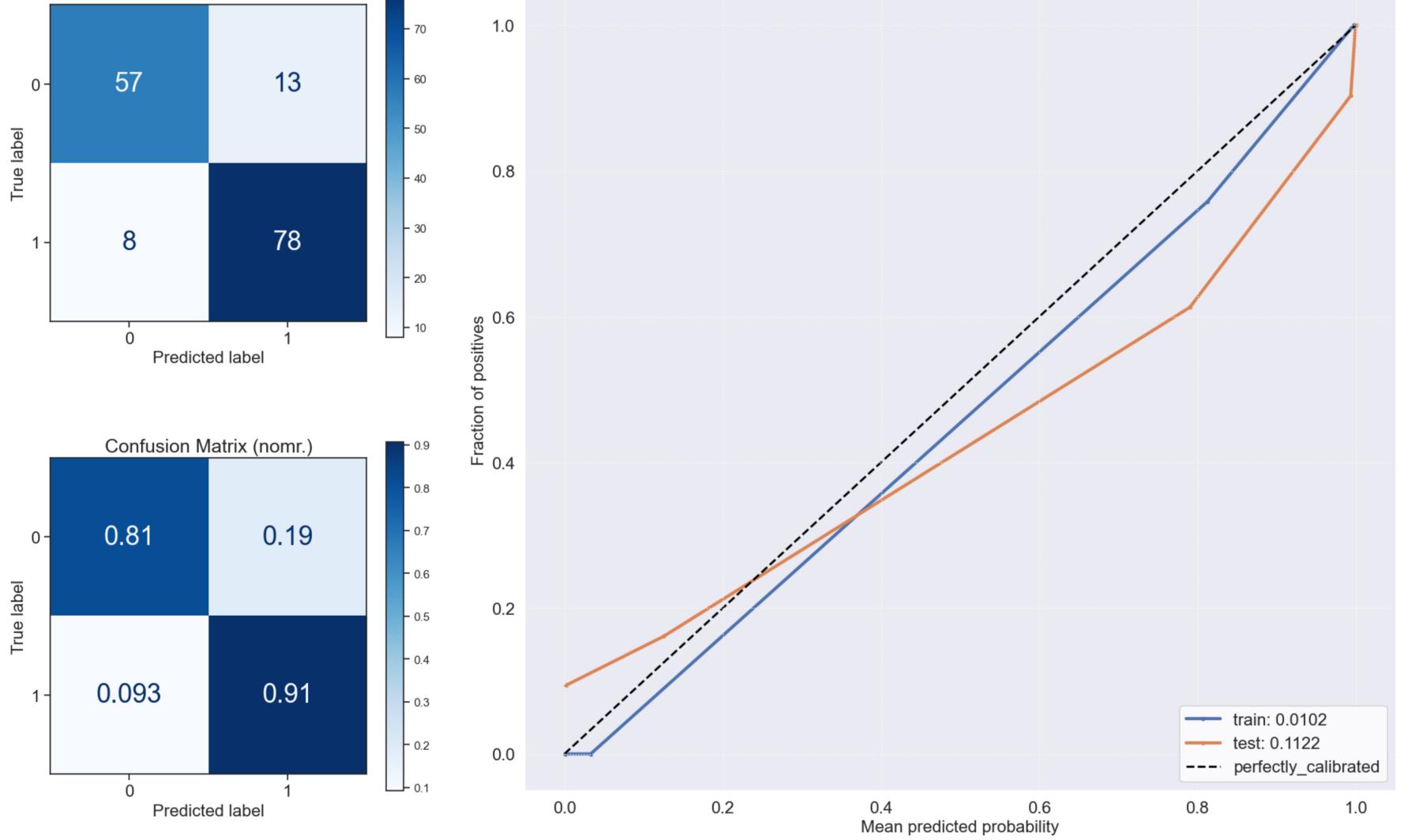
Metric	Train	Test
AUC	0.9996	0.9397
accuracy	0.9856	0.8654
balanced accuracy	0.9839	0.8593
recall_(sens)	1.0	0.9186
specificity	0.9679	0.8
precision	0.9745	0.8495
f1_score	0.9856	0.8645

Protein: P48039 - metric: precision





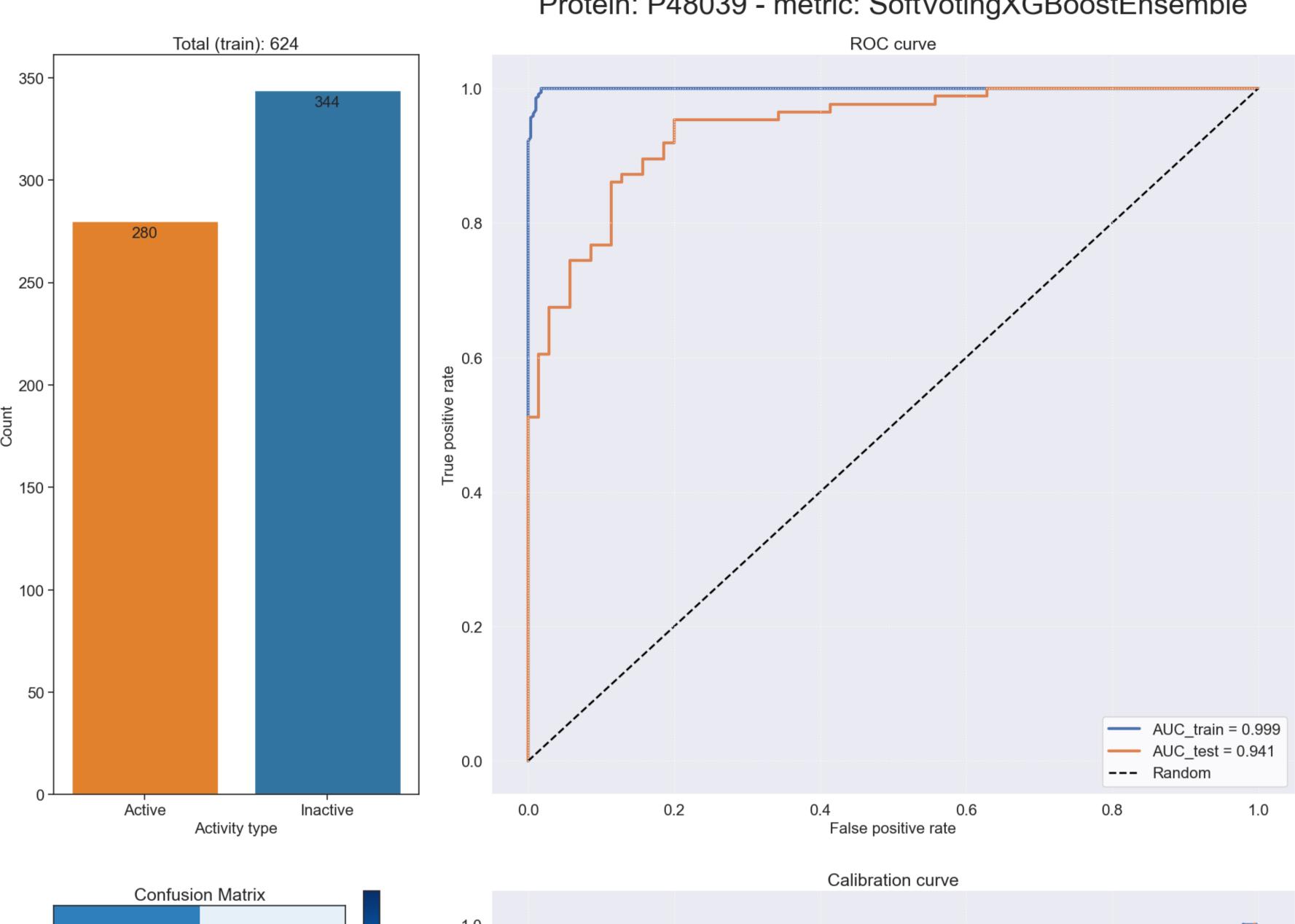
XGBoost Parameters	Value
alpha	3.8401
colsample_bytree	0.8441
gamma	0.438
lambda	1.9722
learning_rate	0.3497
max_depth	7.0
min_child_weight	1.0
n_estimators	488.0
reg_alpha	0.1234
reg_lambda	0.4879
scale_pos_weight	3.9827
subsample	0.5224

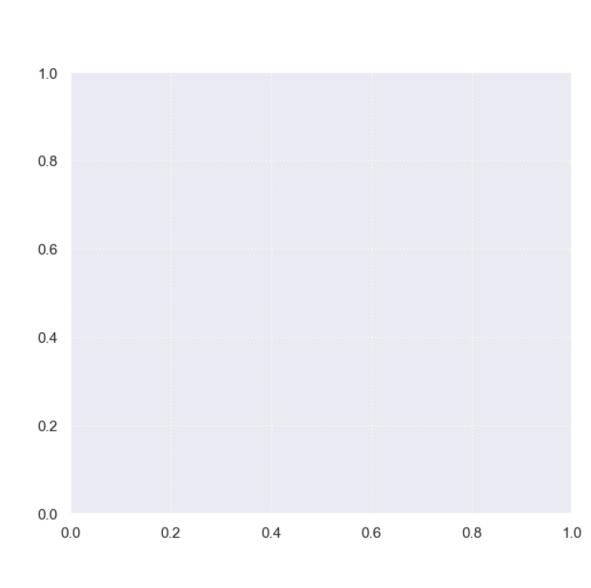


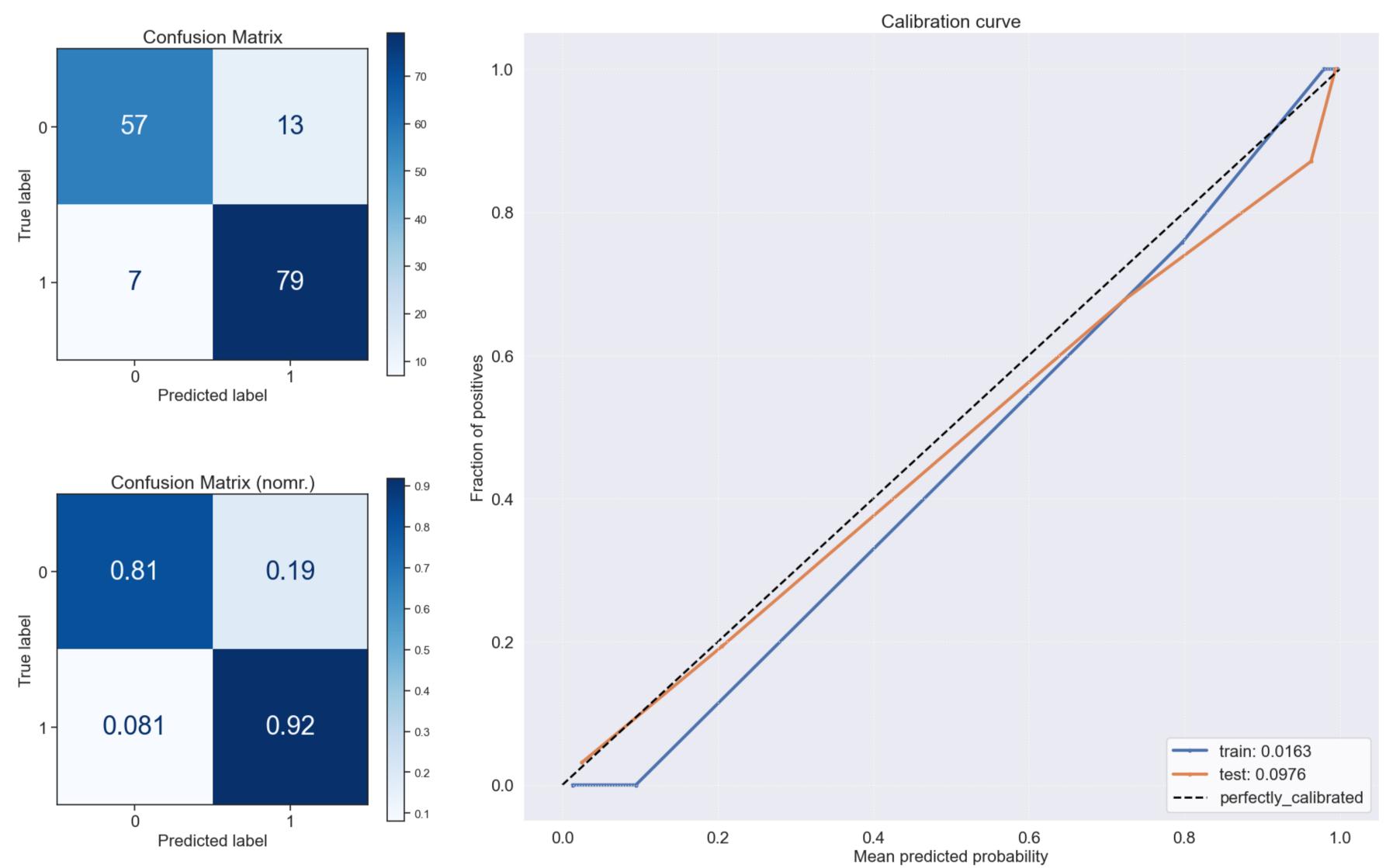
	Active/Inactive probability				
80	Inactive zone		Active zone	type active (86) inactive (70)	
				decoy (156)	
60					
Percent 6					
പ് 40					
20					
0			1		
	0.0 0.2	0.4 Probability	0.6	1.0	

Metric	Train	Test
AUC	0.9997	0.9355
accuracy	0.9888	0.8654
balanced accuracy	0.9875	0.8606
recall_(sens)	1.0	0.907
specificity	0.975	0.8143
precision	0.9801	0.8571
f1_score	0.9888	0.8648

Protein: P48039 - metric: SoftVotingXGBoostEnsemble







			Active/Ina	ctive probabili	ity		
40		Inactive zone			Active 2	zone	type active (86) inactive (70) decoy (156)
30							
Percent 05							
10							
0							
Ü	0.0	0.2	0.4 P	0.6 robability	0.	.8	1.0

Metric	Train	Test
AUC	0.9994	0.941
accuracy	0.984	0.8718
balanced accuracy	0.9821	0.8664
recall_(sens)	1.0	0.9186
specificity	0.9643	0.8143
precision	0.9718	0.8587
f1_score	0.9839	0.8711