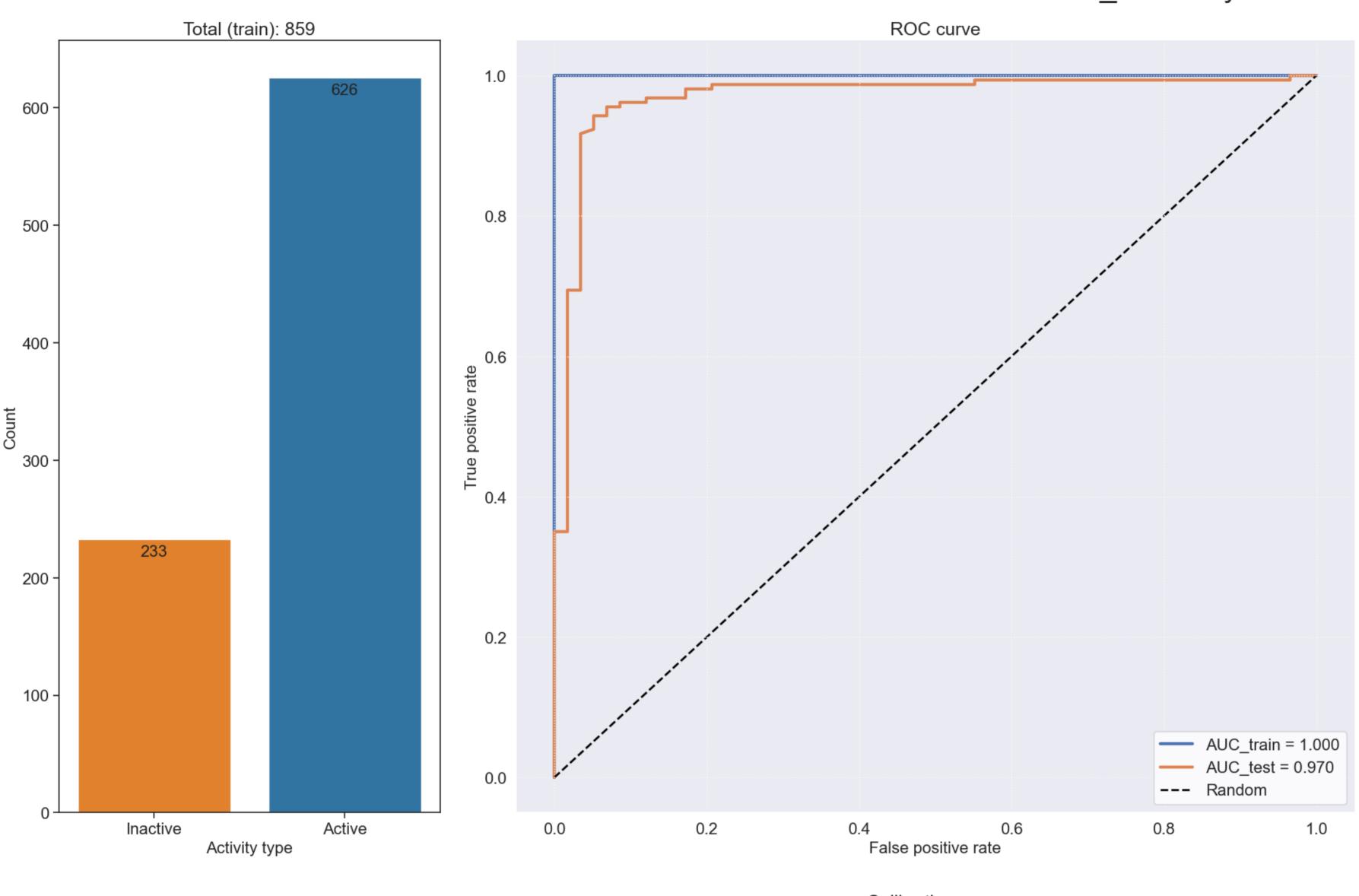
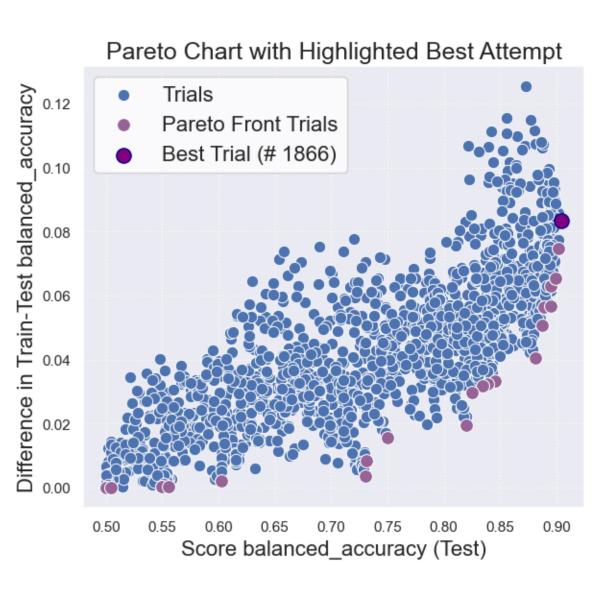
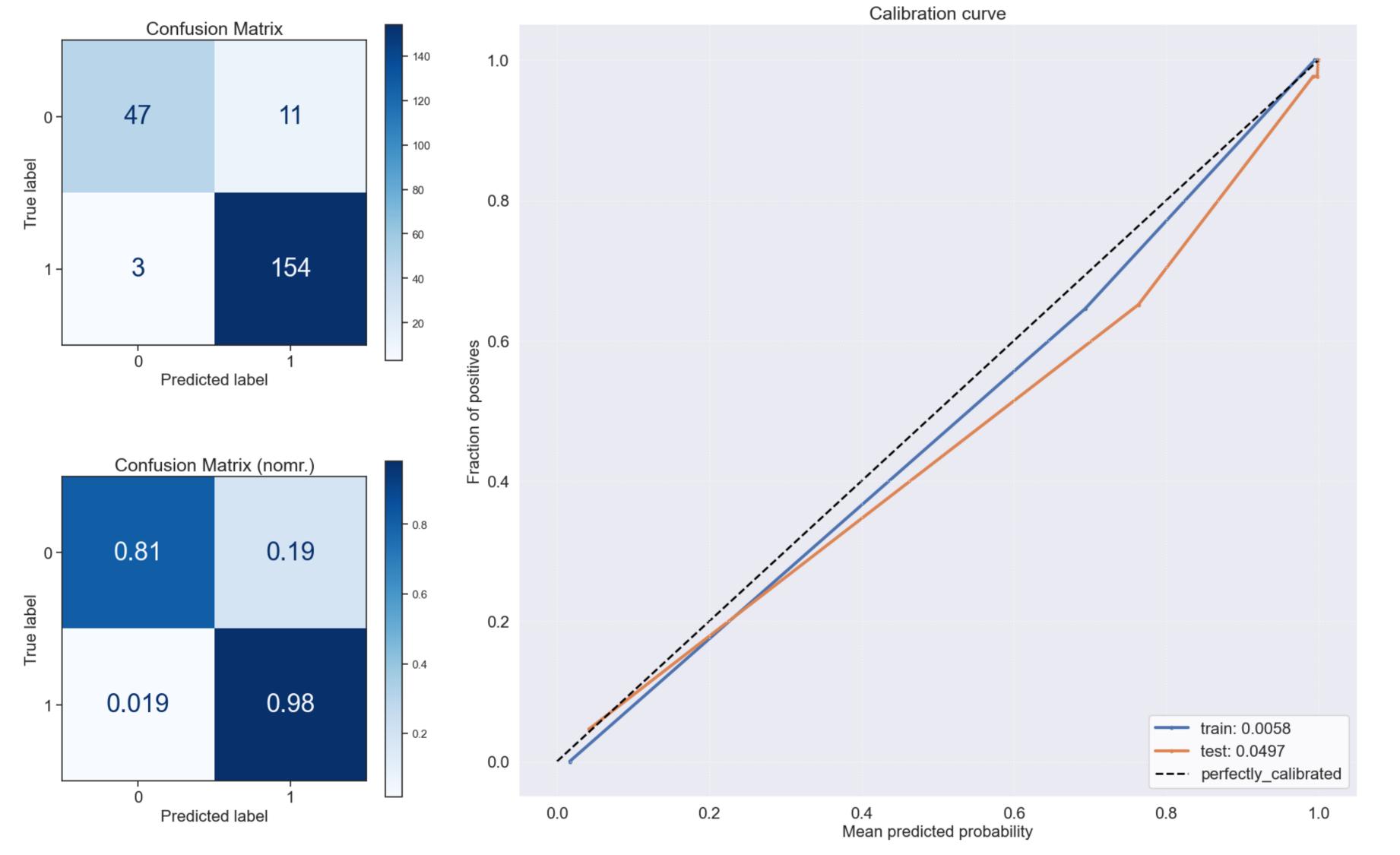
Protein: P05067 - metric: balanced_accuracy





XGBoost Parameters	Value
alpha	0.263
colsample_bytree	0.9724
gamma	0.403
lambda	4.6353
learning_rate	0.3816
max_depth	8.0
min_child_weight	2.0
n_estimators	802.0
reg_alpha	0.4185
reg_lambda	0.2296
scale_pos_weight	2.6048
subsample	0.8684

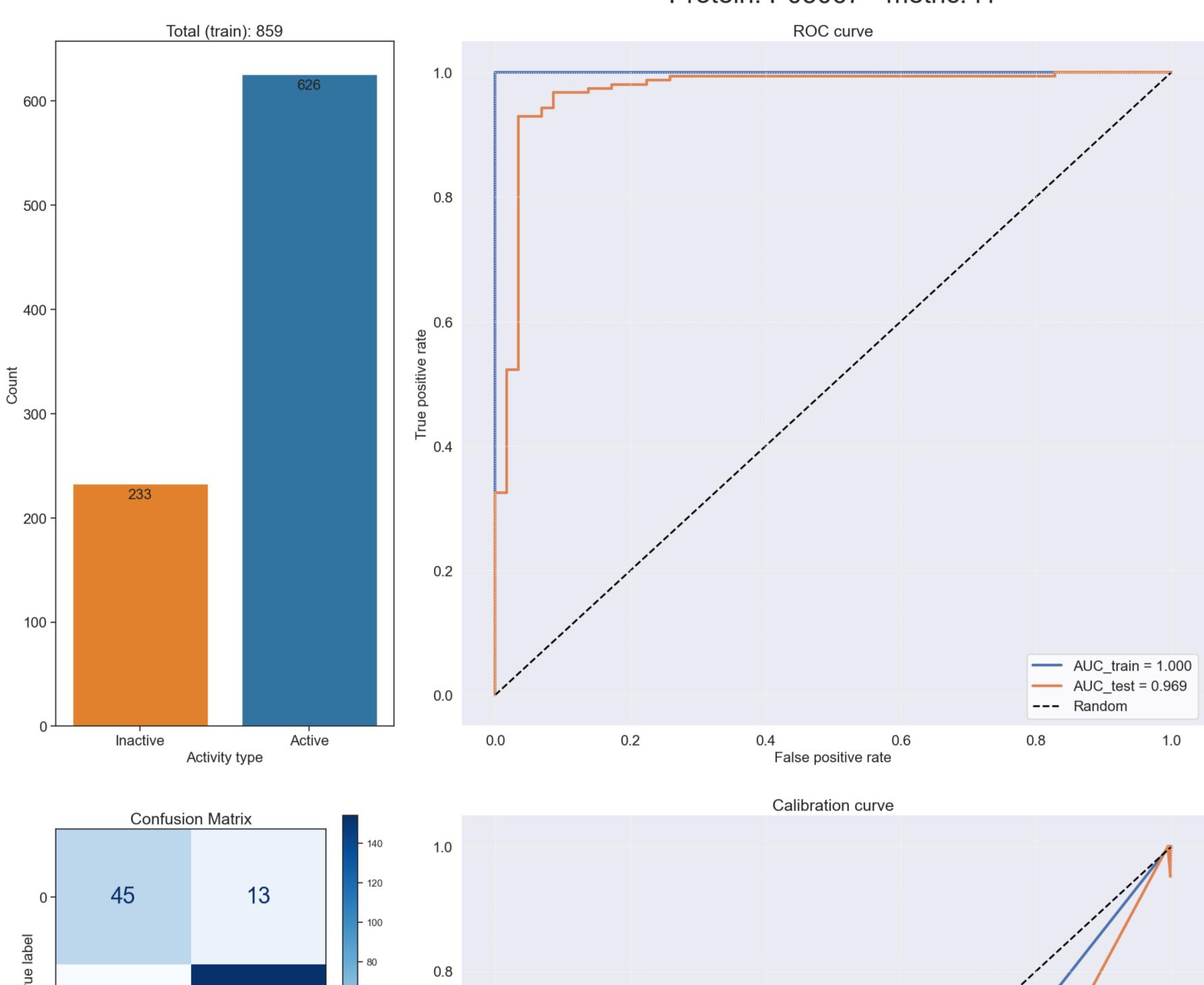


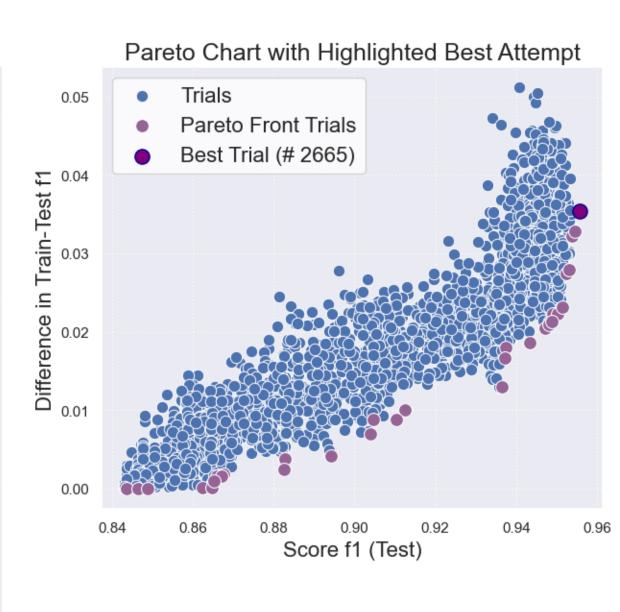
		Inactive zone				inact deco	e (157) ive (58)
50							
ercent 05							
<u> </u>							
20							
0 —	1-2-						
	0.0	0.2	0.4 Prol	0.6 pability	0.8	•	1.0

Active/Inactive probability

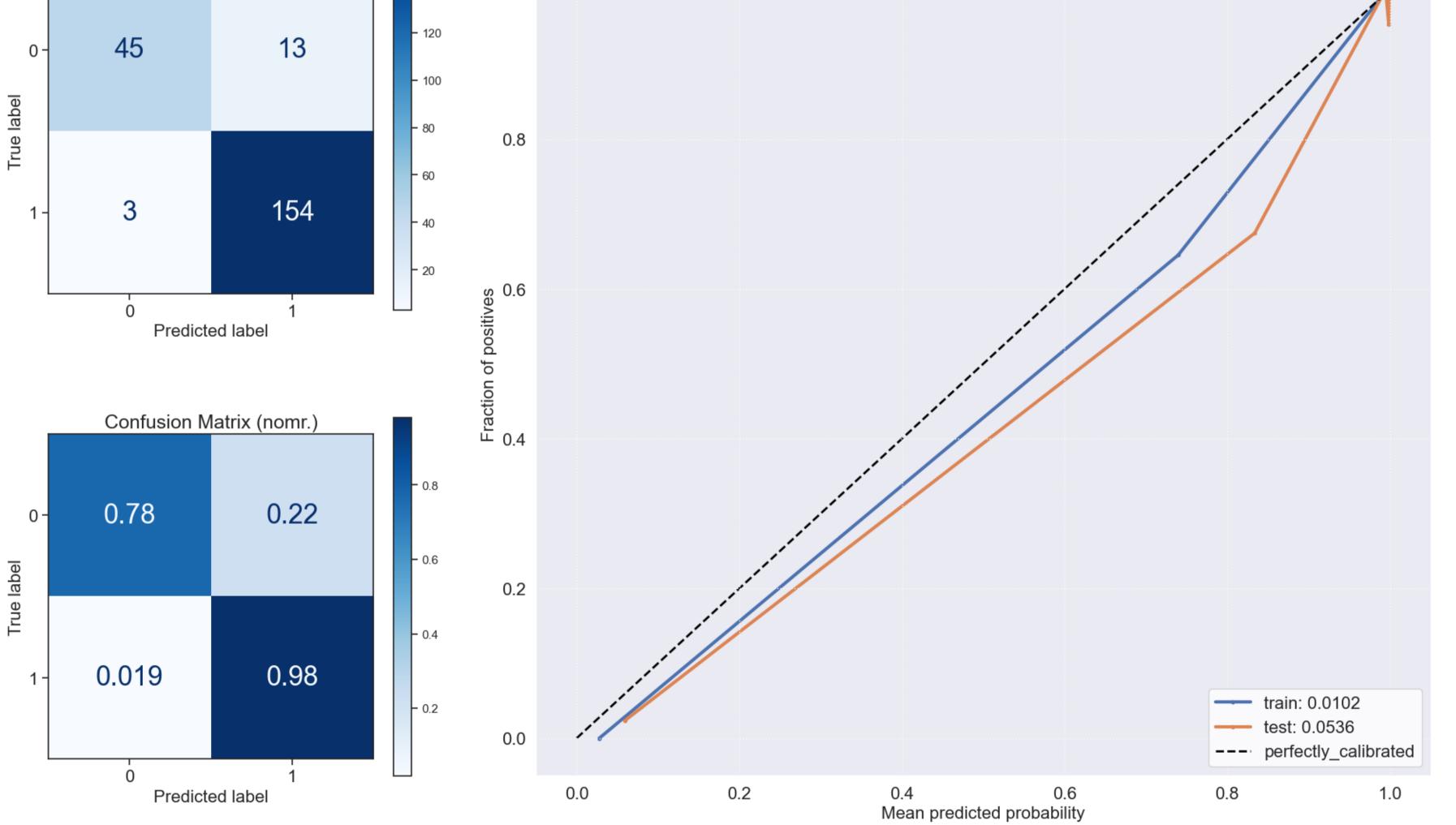
Metric	Train	Test
AUC	1.0	0.9697
accuracy	0.9965	0.9349
balanced accuracy	0.9936	0.8956
recall_(sens)	1.0	0.9809
specificity	0.9871	0.8103
precision	0.9952	0.9333
f1_score	0.9965	0.9333

Protein: P05067 - metric: f1





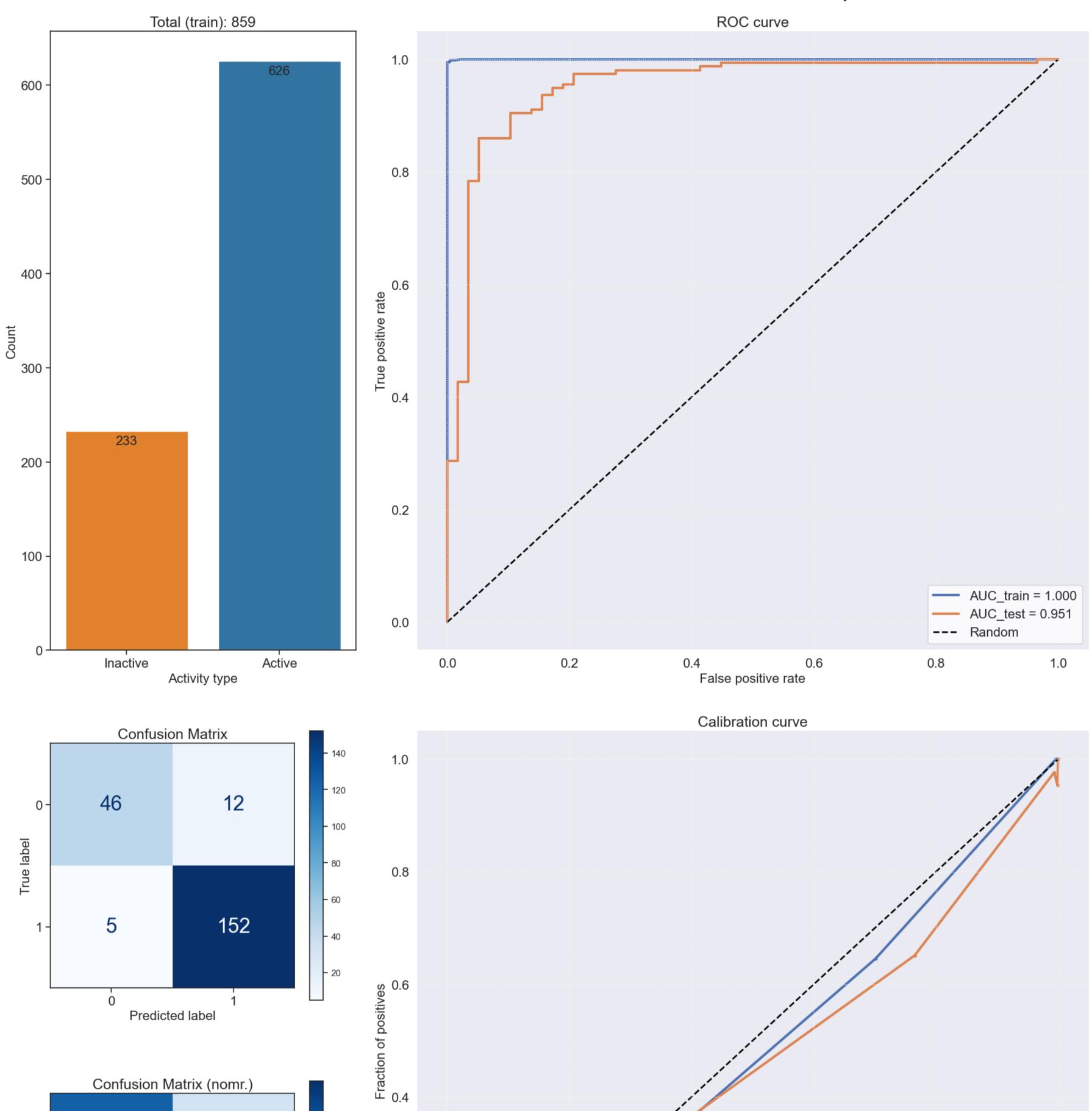
XGBoost Parameters	Value
alpha	2.4641
colsample_bytree	0.938
gamma	0.3353
lambda	3.0646
learning_rate	0.0563
max_depth	8.0
min_child_weight	3.0
n_estimators	973.0
reg_alpha	0.1368
reg_lambda	0.2764
scale_pos_weight	6.636
subsample	0.9612

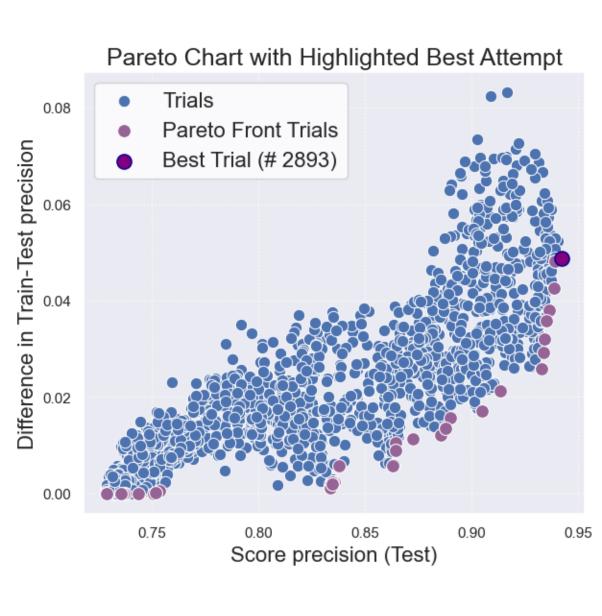


		Act	ive/Inactiv	e probabi	lity			
80	Inactiv	ve zone			Active	: _	type active (157) inactive (58)	
70							decoy (215)	
60								
Percent 0.								
30								
20	П							
10								
0 —	0.0 0.2	0.	4 Proba	0. ability	6 ().8	1.0	
	1 Tobability							

Metric	Train	Test
AUC	1.0	0.9691
accuracy	0.986	0.9256
balanced accuracy	0.9742	0.8784
recall_(sens)	1.0	0.9809
specificity	0.9485	0.7759
precision	0.9812	0.9222
f1_score	0.9859	0.9232

Protein: P05067 - metric: precision





XGBoost Parameters	Value
alpha	4.9234
colsample_bytree	0.8605
gamma	0.5457
lambda	2.9011
learning_rate	0.8991
max_depth	3.0
min_child_weight	2.0
n_estimators	110.0
reg_alpha	0.3692
reg_lambda	0.251
scale_pos_weight	2.6572
subsample	0.7904

	Active/Inactive probability						
70	Ina	ctive zone		Active	inact	e (157) ive (58) y (215)	
50							
Percent 5 6							
30							
10							
U	0.0	.2 0	.4 (Probability	0.6 0	.8	1.0	

Confusion Matrix (nomr.)

Predicted label

0.21

0.97

- 0.6

- 0.4

0.2

0.2

0.0

0.0

0.2

0.79

0.032

True label

Metric	Train	Test
AUC	1.0	0.9511
accuracy	0.9907	0.9209
balanced accuracy	0.9828	0.8806
recall_(sens)	1.0	0.9682
specificity	0.9657	0.7931
precision	0.9874	0.9268
f1_score	0.9906	0.9193

train: 0.0072

test: 0.0695

0.8

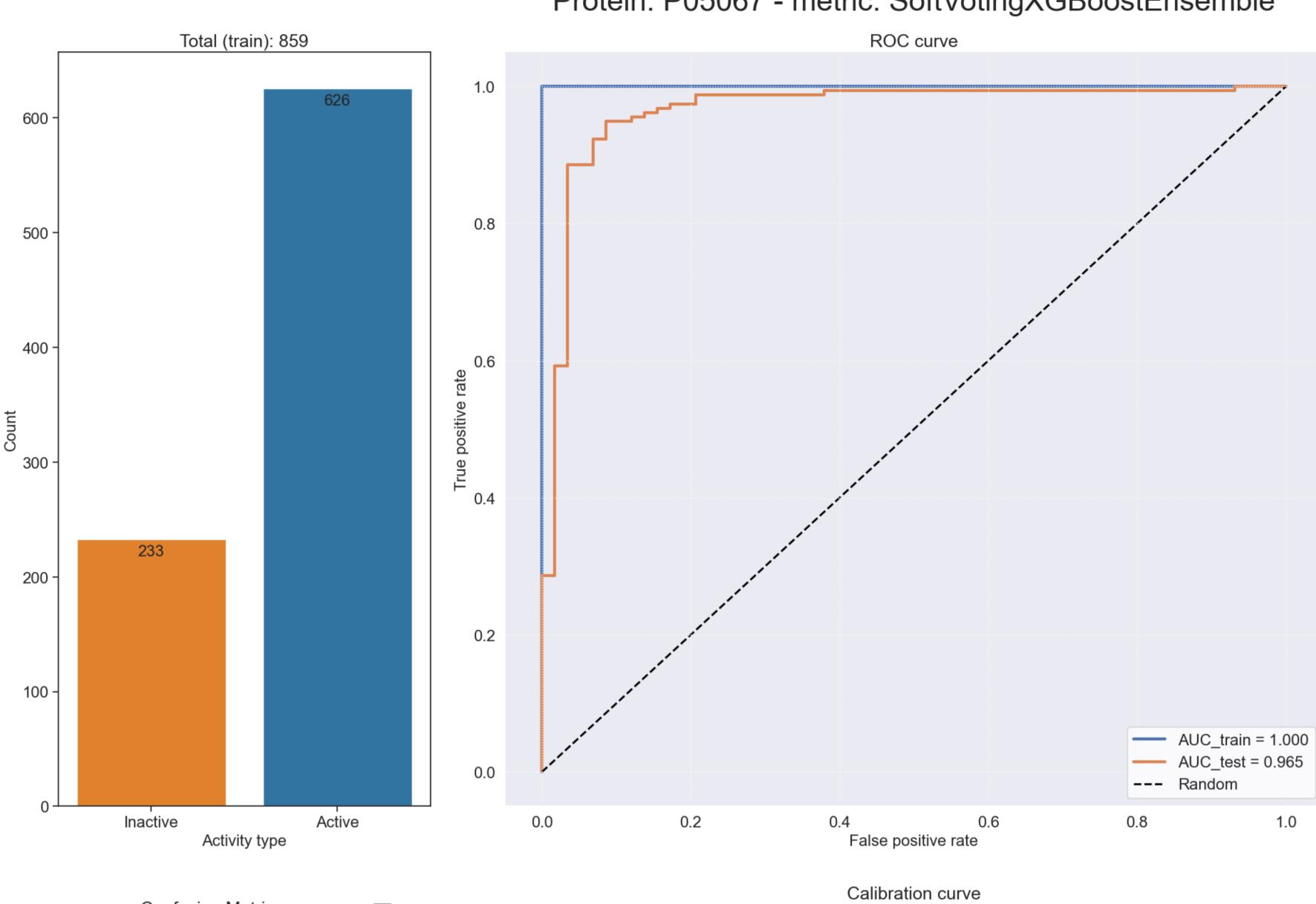
--- perfectly_calibrated

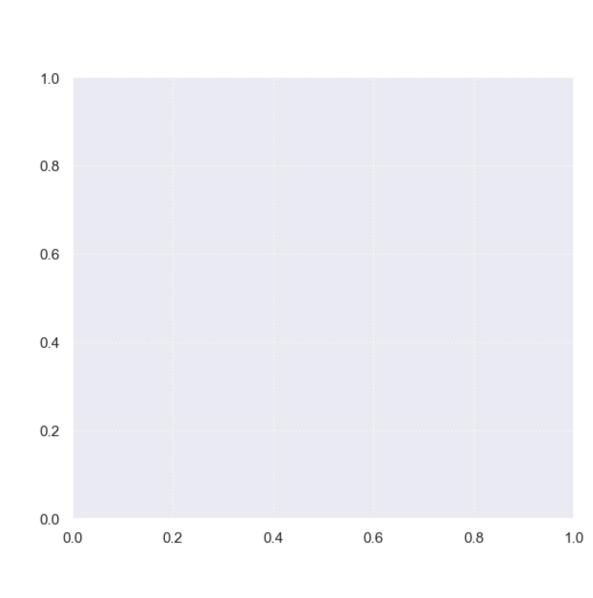
1.0

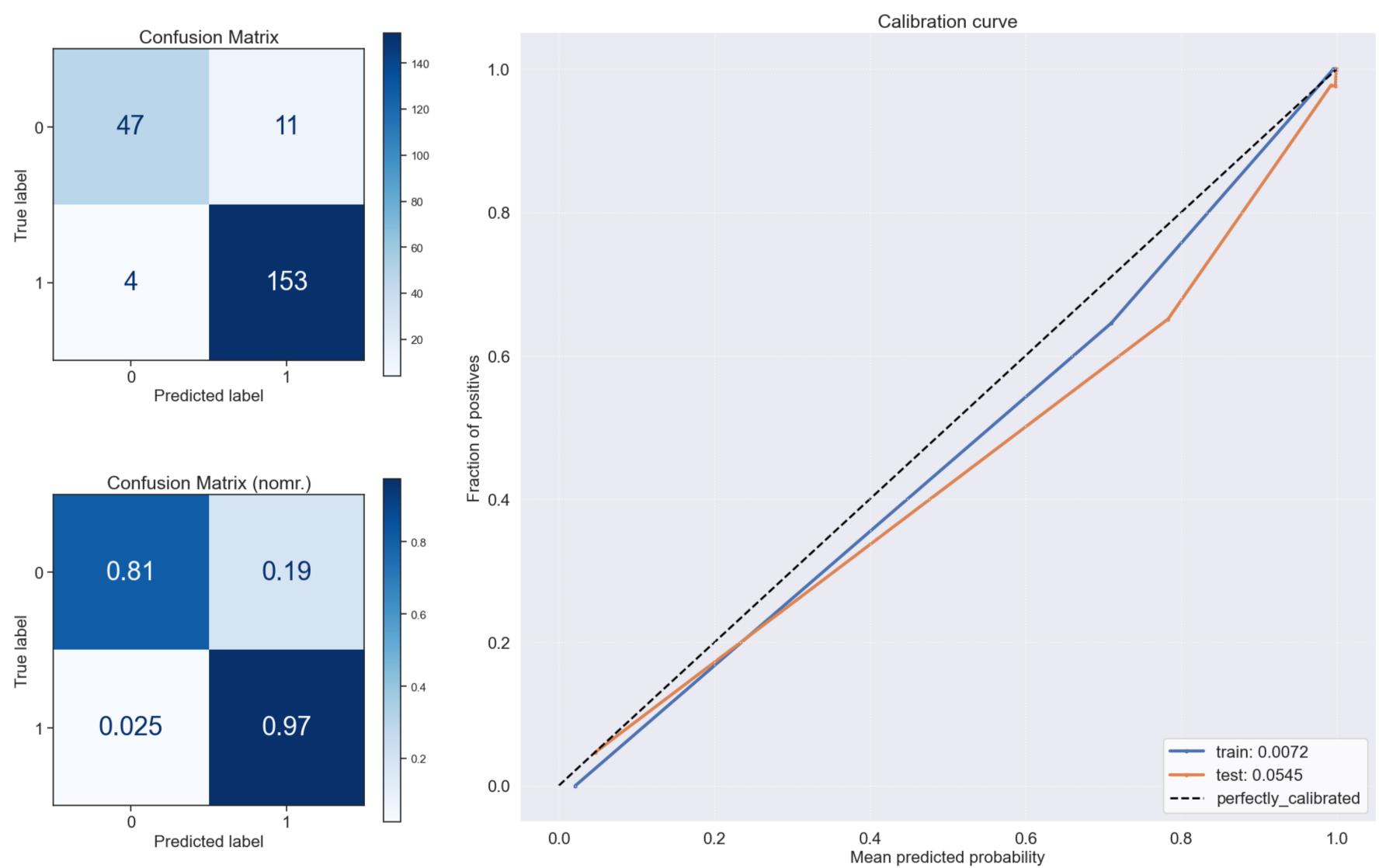
	Ac	tive/Inactive probabi	lity				
70	Inactive zone		Active 2	zone	type active (157)	Metric	Train
60				i i	decoy (215)	AUC	1.0
50						accuracy	0.9907
						balanced accuracy	0.9828
Percent 30						recall_(sens)	1.0
						specificity	0.9657
20						precision	0.9874
10						f1_score	0.9906
0.	0 0.2 0	0.4			1.0		

0.4 0.6 Mean predicted probability

Protein: P05067 - metric: SoftVotingXGBoostEnsemble







	Ina					zone	inactive (58) decoy (215)
50							
ercent - 04							
<u> </u>							
20							
10							
0	0.0 0	.2 0	.4 Probab	0.6	0.	.8	1.0

Active/Inactive probability

Metric	Train	Test
AUC	1.0	0.9651
accuracy	0.9895	0.9302
balanced accuracy	0.9807	0.8924
recall_(sens)	1.0	0.9745
specificity	0.9614	0.8103
precision	0.9858	0.9329
f1_score	0.9895	0.9288