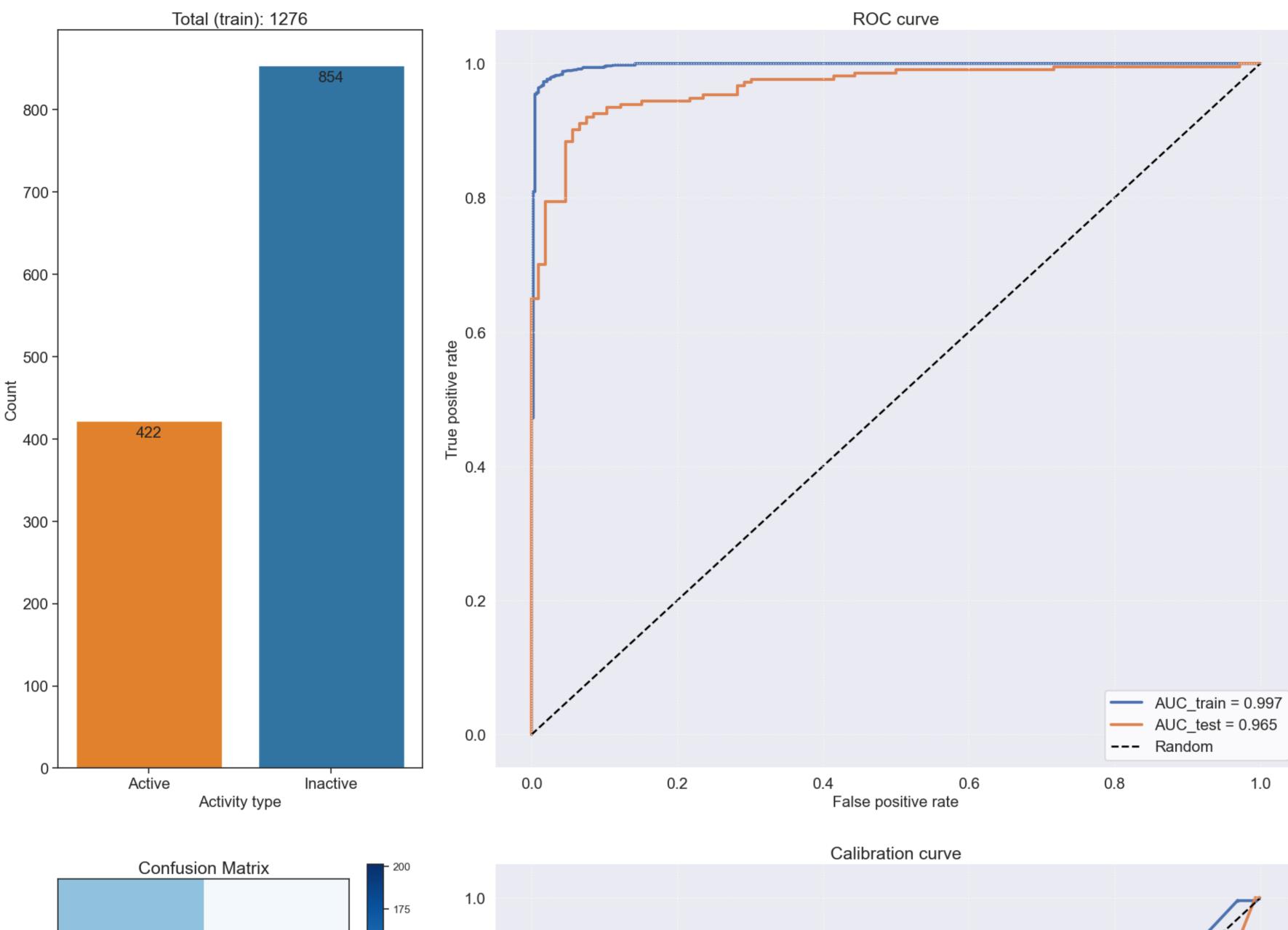
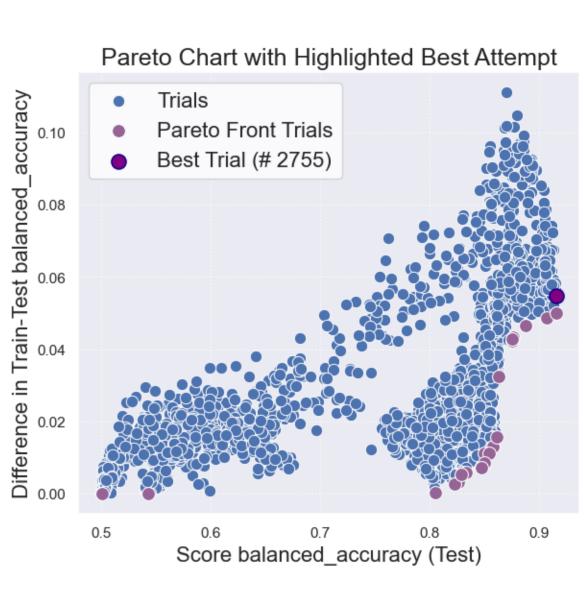
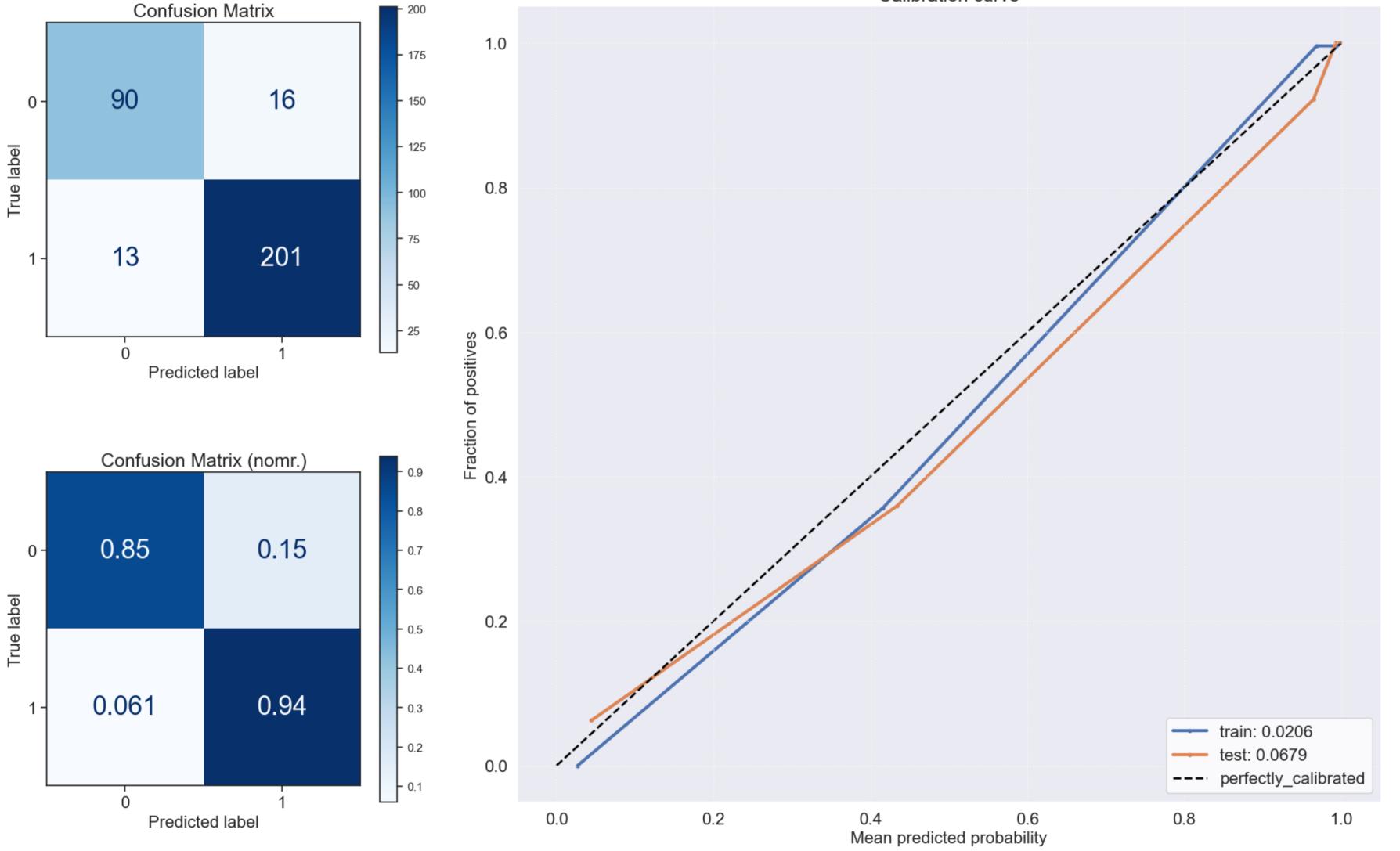
## Protein: O76074 - metric: balanced\_accuracy





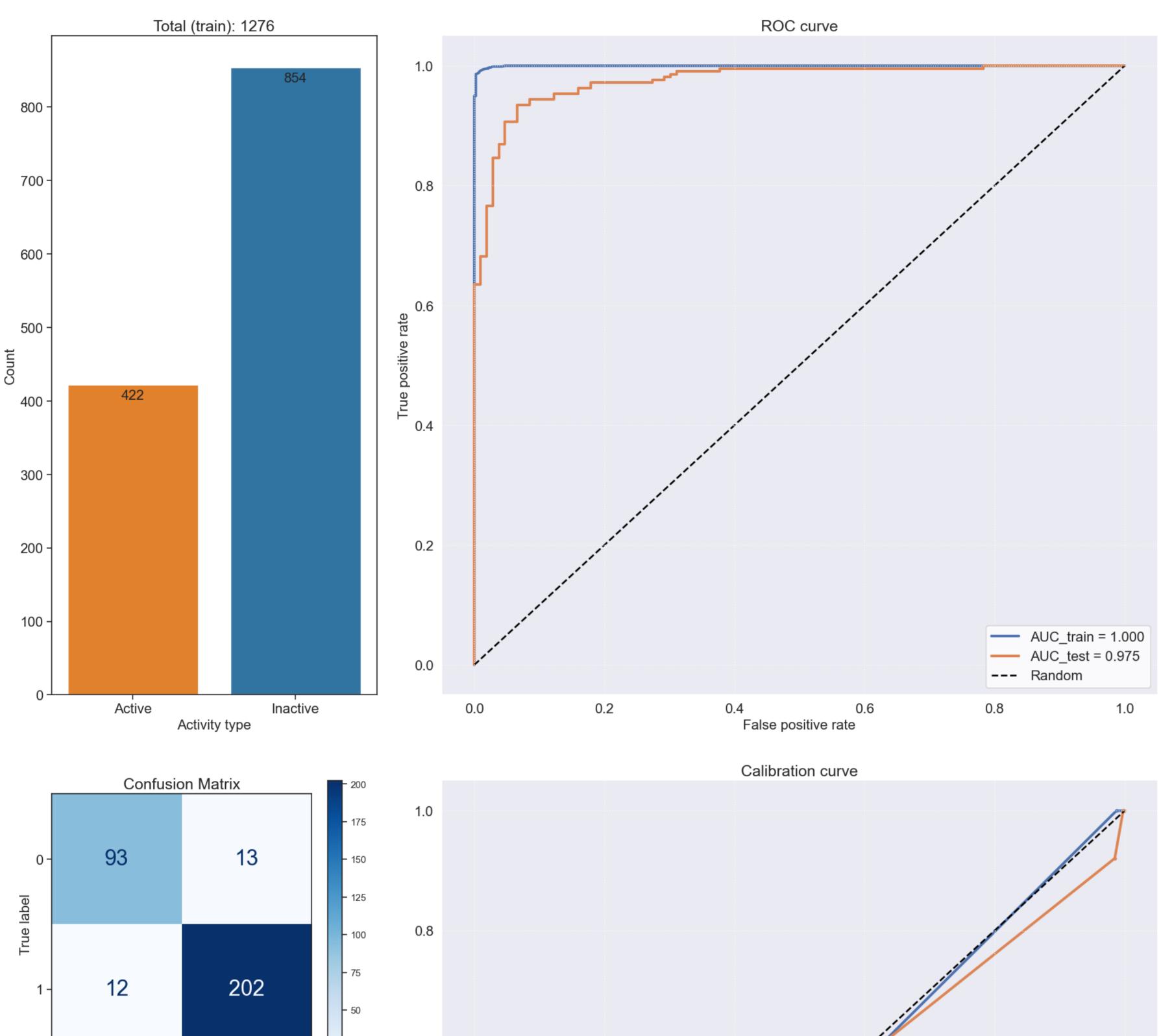
Value
2.0617
0.5892
0.9741
0.1609
0.0729
5.0
3.0
523.0
0.1237
0.3237
1.4201
0.5071

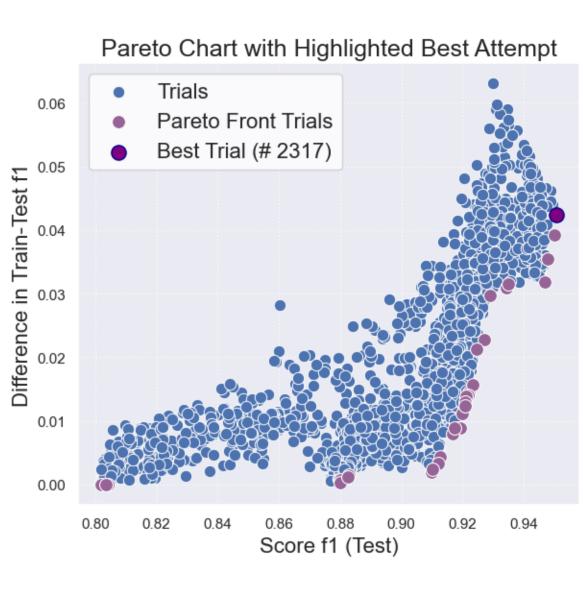


	Active/Inactive probability						
30	In	active zone		i	Active zone	type active (214) inactive (106) decoy (320)	
Percent 08							
10							
O	0.0	0.2	0.4 Probability	0.6	0.8	1.0	

Metric	Train	Test
AUC	0.9967	0.9654
accuracy	0.9765	0.9094
balanced accuracy	0.9716	0.8942
recall_(sens)	0.9859	0.9393
specificity	0.9573	0.8491
precision	0.9791	0.9263
f1_score	0.9764	0.909

## Protein: O76074 - metric: f1





XGBoost Parameters	Value
alpha	4.3534
colsample_bytree	0.7359
gamma	0.0244
lambda	3.4388
learning_rate	0.0878
max_depth	10.0
min_child_weight	2.0
n_estimators	451.0
reg_alpha	0.1454
reg_lambda	0.1742
scale_pos_weight	1.3989
subsample	0.587

		Ac	tive/Inactive pro	bability		
70		ctive zone			Active zone	active (214) inactive (106) decoy (320)
50						
Percent						
20						
10						
	0.0	.2 0	.4 Probability	0.6	0.8	1.0

- 25

- 0.6

- 0.5

0.4

0.3

0.2

- 0.1

Predicted label

Confusion Matrix (nomr.)

Predicted label

0.12

0.94

0.88

0.056

True label

Fraction of positives

0.2

0.0

0.0

0.2

Metric	Train	Test
AUC	0.9997	0.9752
accuracy	0.9898	0.9219
balanced accuracy	0.9858	0.9106
recall_(sens)	0.9977	0.9439
specificity	0.9739	0.8774
precision	0.9873	0.9395
f1_score	0.9898	0.9218

train: 0.0081

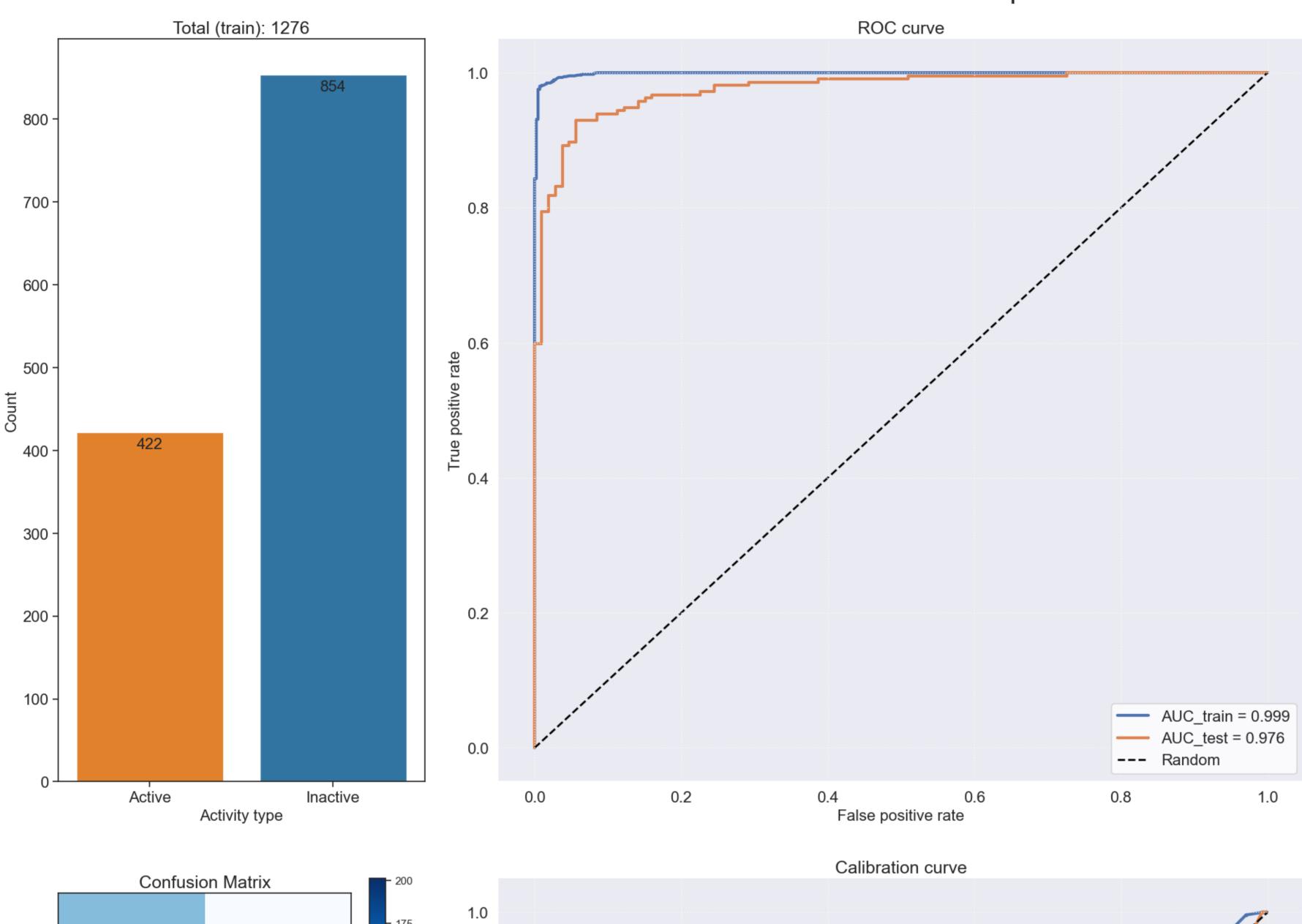
- test: 0.0567

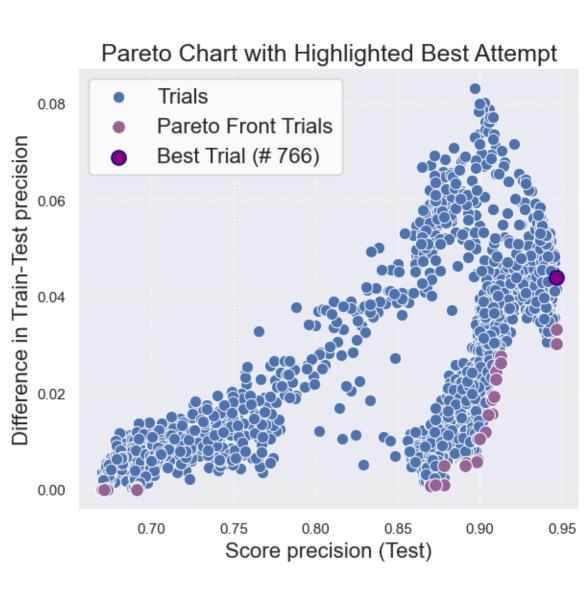
0.8

0.4 0.6 Mean predicted probability --- perfectly\_calibrated

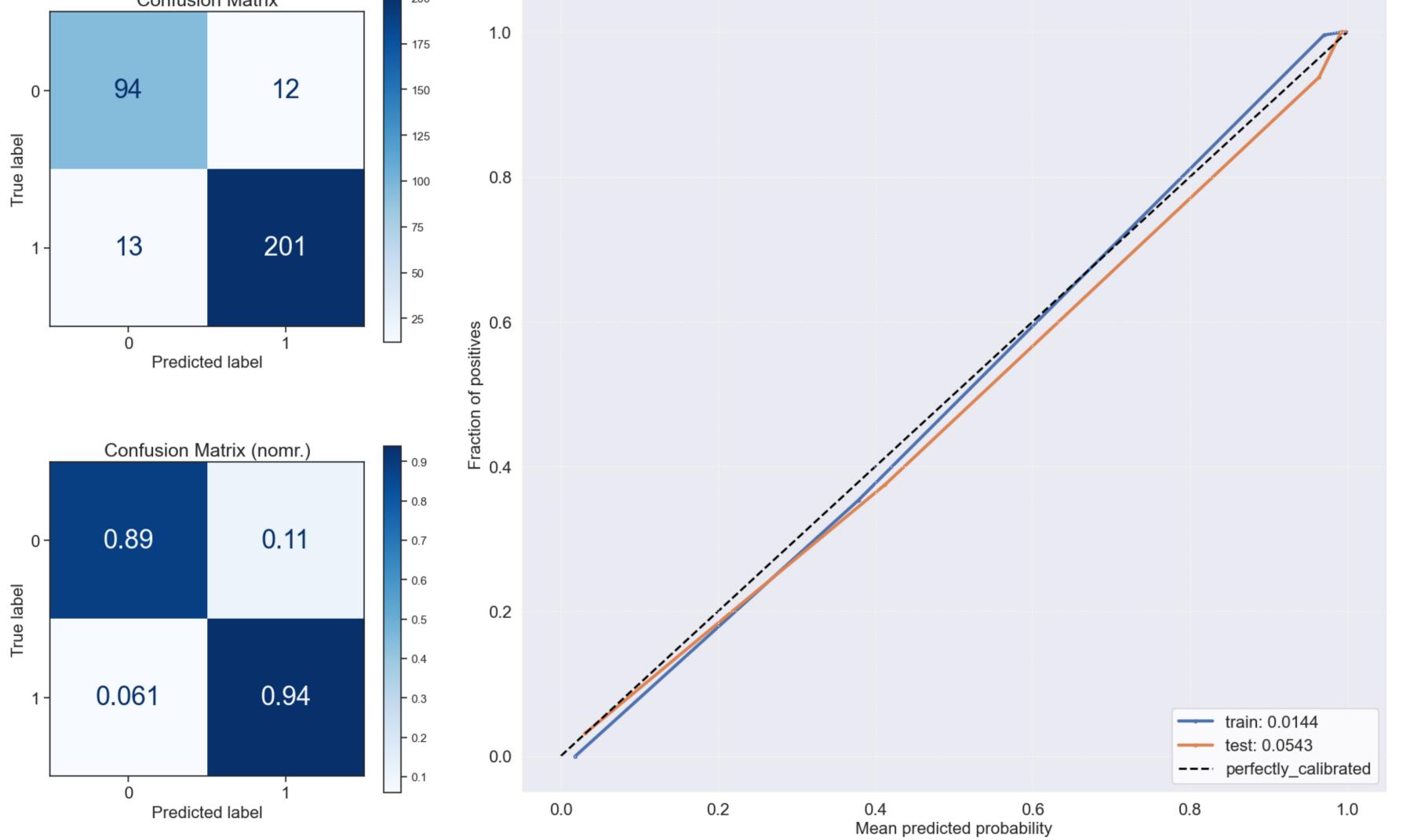
1.0

## Protein: O76074 - metric: precision





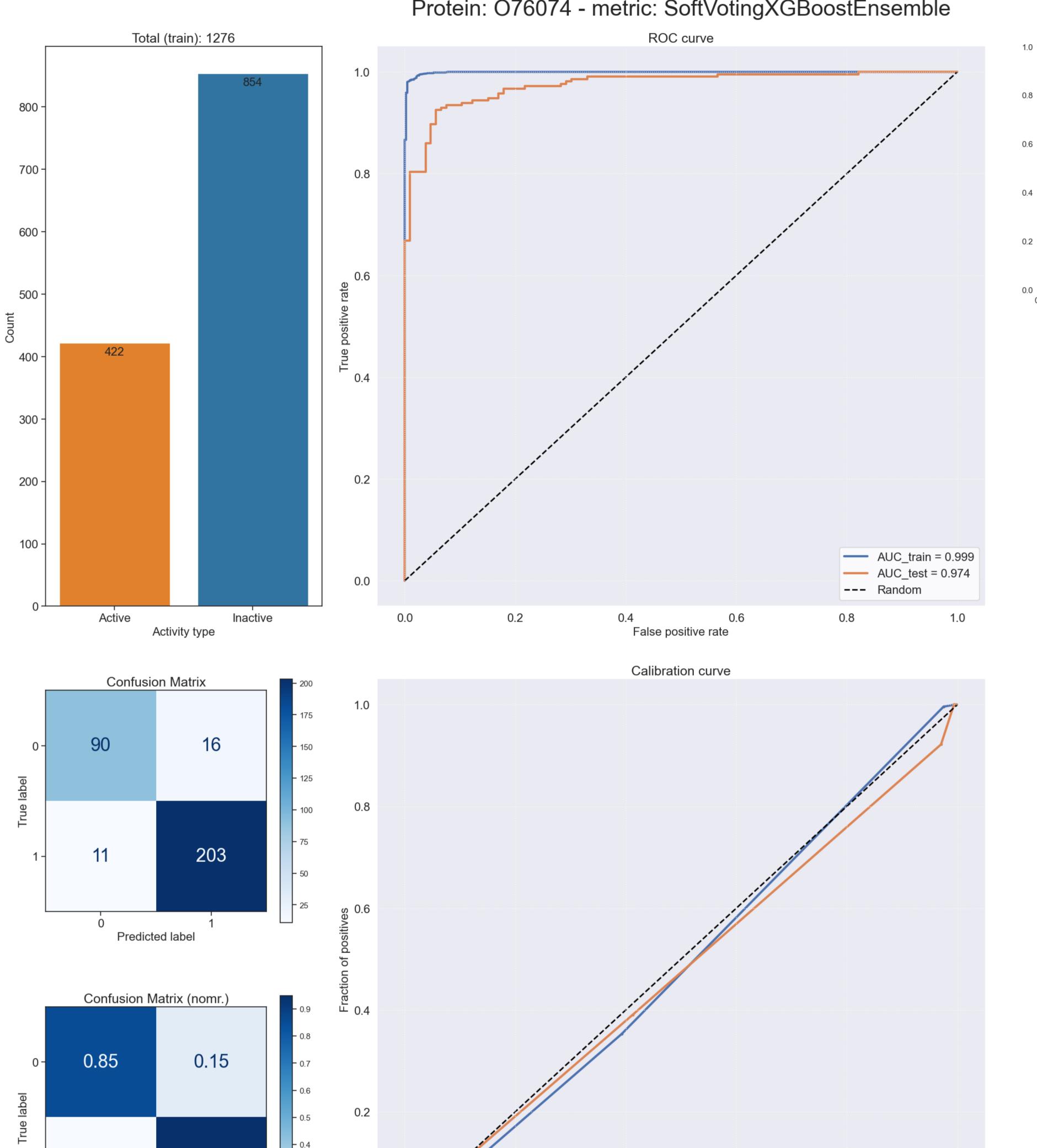
XGBoost Parameters	Value
alpha	0.8118
colsample_bytree	0.5977
gamma	0.1306
lambda	4.0969
learning_rate	0.1234
max_depth	10.0
min_child_weight	2.0
n_estimators	166.0
reg_alpha	0.5474
reg_lambda	0.5239
scale_pos_weight	1.0925
subsample	0.6609



	Active/Inactive probability						
50	lna	active zone			Active 2	inactiv	(214) ve (106)
40						decoy	(320)
Percent 08							
20							
10							
0 —				<u></u>			
	0.0	0.2	0.4 Proba	0.6 bility	0	.8	1.0

Metric	Train	Test
AUC	0.9988	0.976
accuracy	0.9835	0.9219
balanced accuracy	0.9829	0.913
recall_(sens)	0.9848	0.9393
specificity	0.981	0.8868
precision	0.9906	0.9437
f1_score	0.9836	0.922

## Protein: O76074 - metric: SoftVotingXGBoostEnsemble



Metric	Train	Test
AUC	0.9992	0.9744
accuracy	0.9867	0.9156
balanced accuracy	0.9847	0.8988
recall_(sens)	0.9906	0.9486
specificity	0.9787	0.8491
precision	0.9895	0.9269
f1_score	0.9867	0.9151

--- train: 0.0132

--- test: 0.0581

0.8

0.4 0.6 Mean predicted probability

--- perfectly\_calibrated

1.0

0.2

0.6

0.0

		Active/Inacti	ve probability		
00	Inactive	zone		Active zone	type active (214) inactive (106) decoy (320)
40					
Percent 0.					
20	<u></u>				
10					
0	0.2	0.4 Prob	0.6 pability	0.8	1.0

0.2

0.0

0.0

0.2

- 0.5

0.4

0.3

0.2

0.1

0.95

0.051

Predicted label