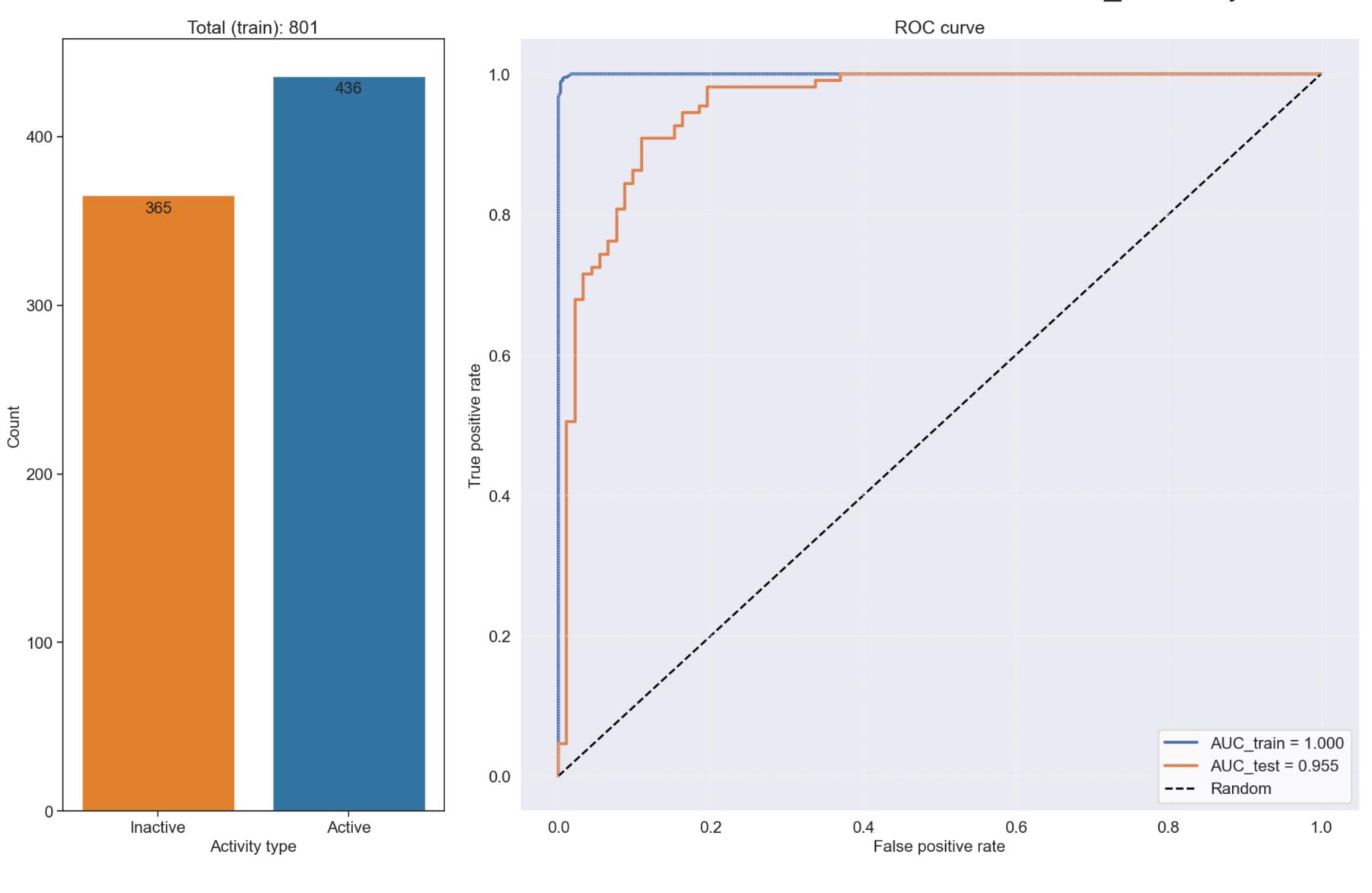
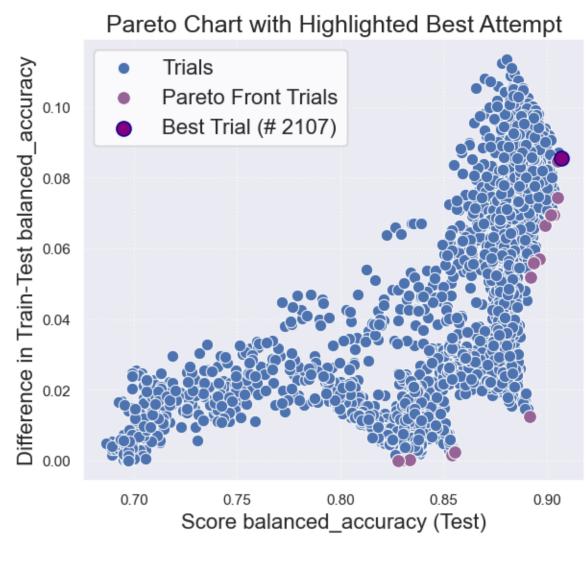
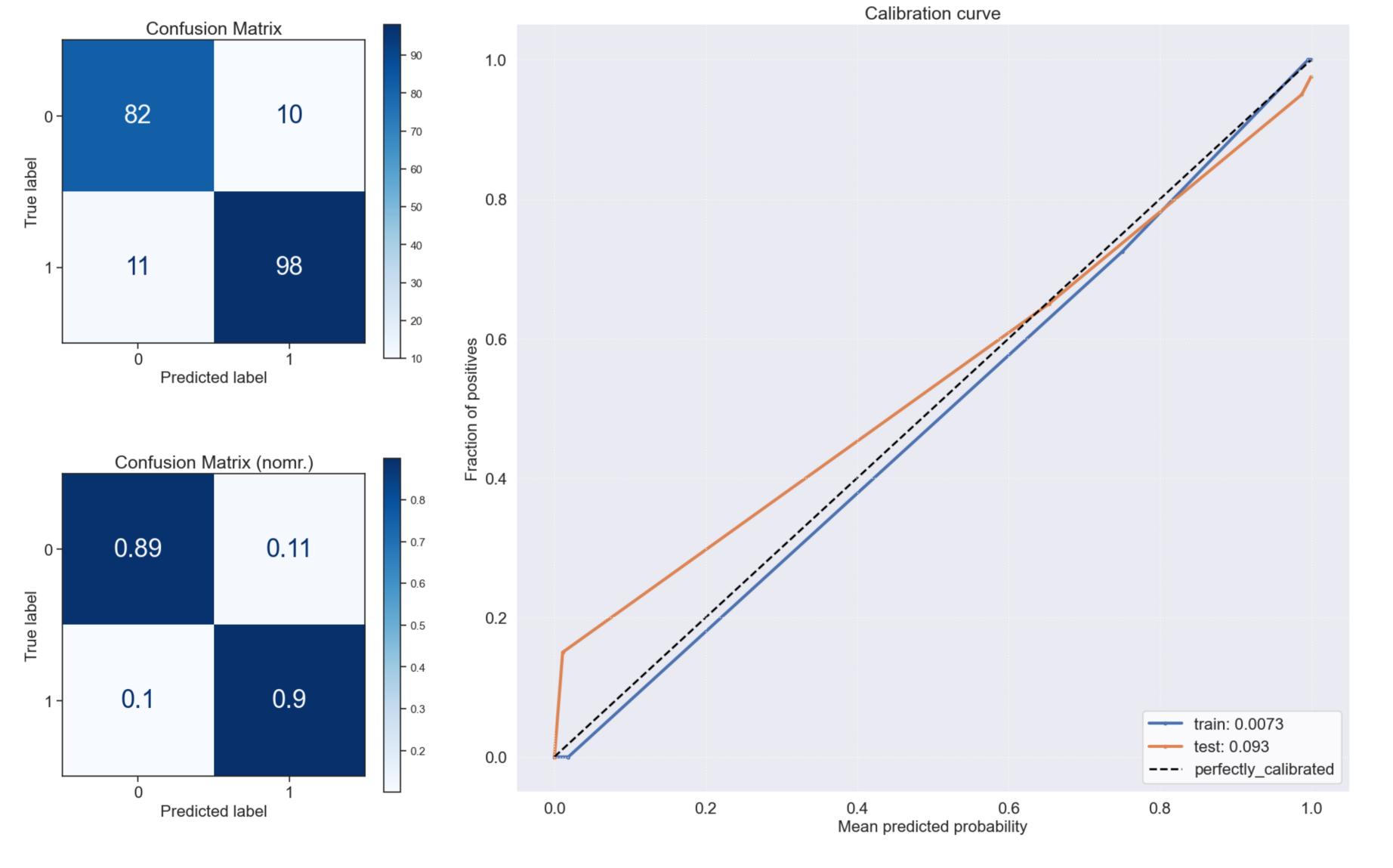
## Protein: P08588 - metric: balanced\_accuracy





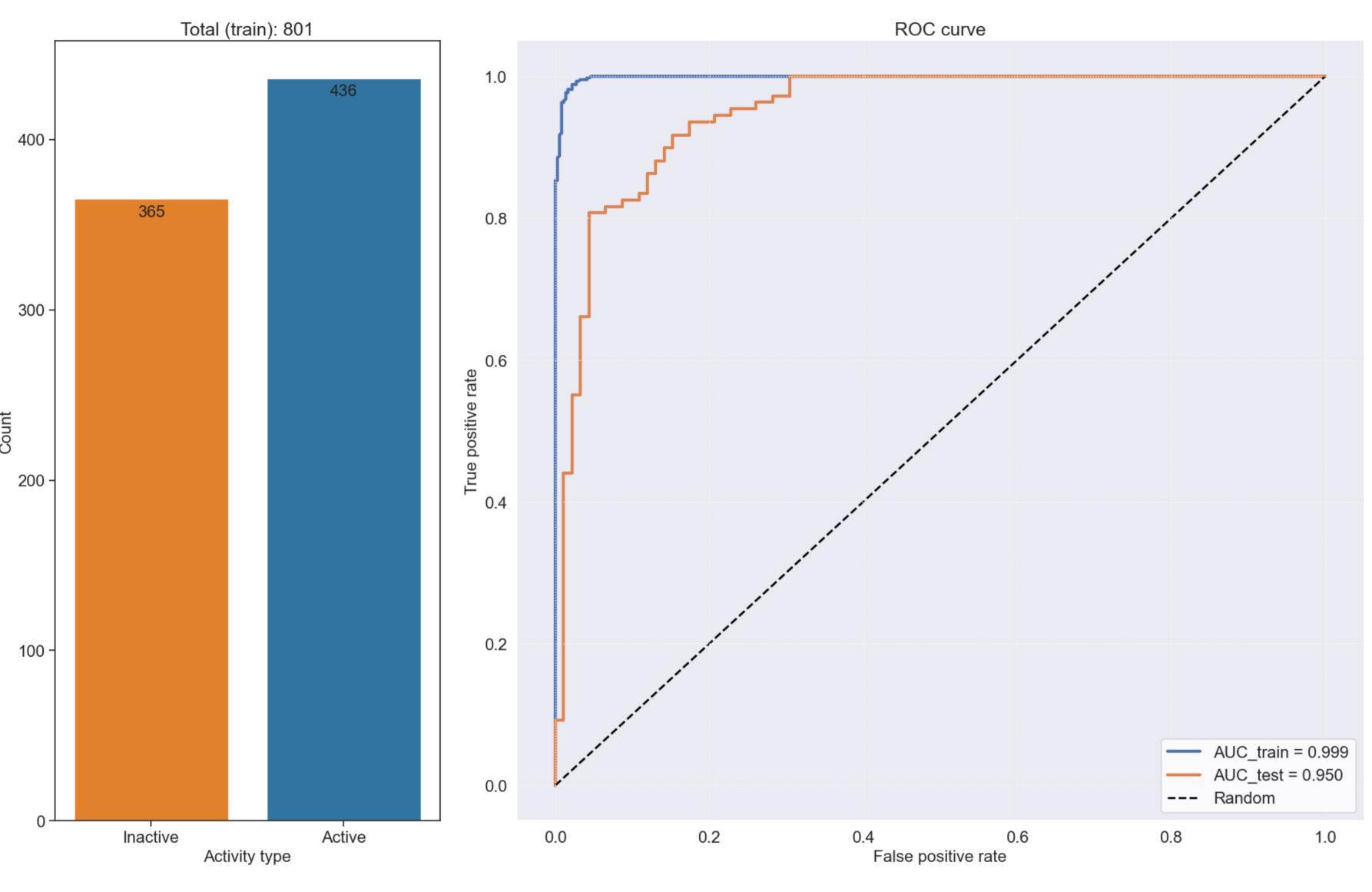
XGBoost Parameters	Value
alpha	2.5798
colsample_bytree	0.9548
gamma	0.1682
lambda	3.1338
learning_rate	0.2508
max_depth	10.0
min_child_weight	1.0
n_estimators	446.0
reg_alpha	0.8781
reg_lambda	0.1112
scale_pos_weight	1.8141
subsample	0.542

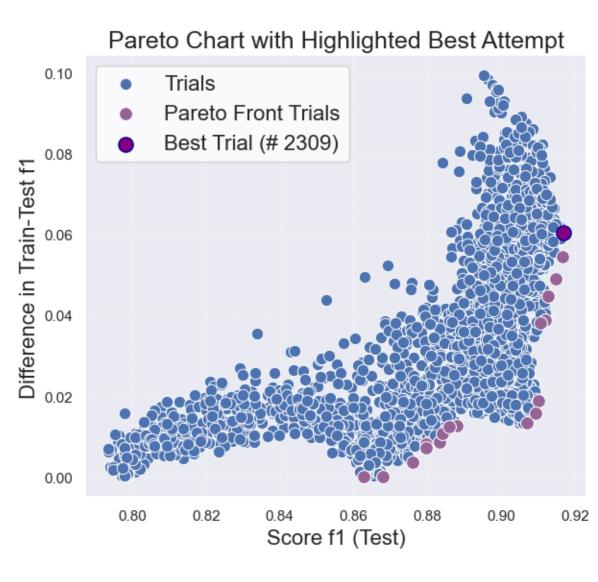


	Active/Inactive probability					
60	Inactive zo	· · · · · · · · · · · · · · · · · · ·		Active zone	type active (109) inactive (92)	
00					decoy (201)	
50						
Percent 0.00						
പ് 30						
20						
10						
0	0.0 0.2	0.4 Probab	0.6 ility	0.8	1.0	

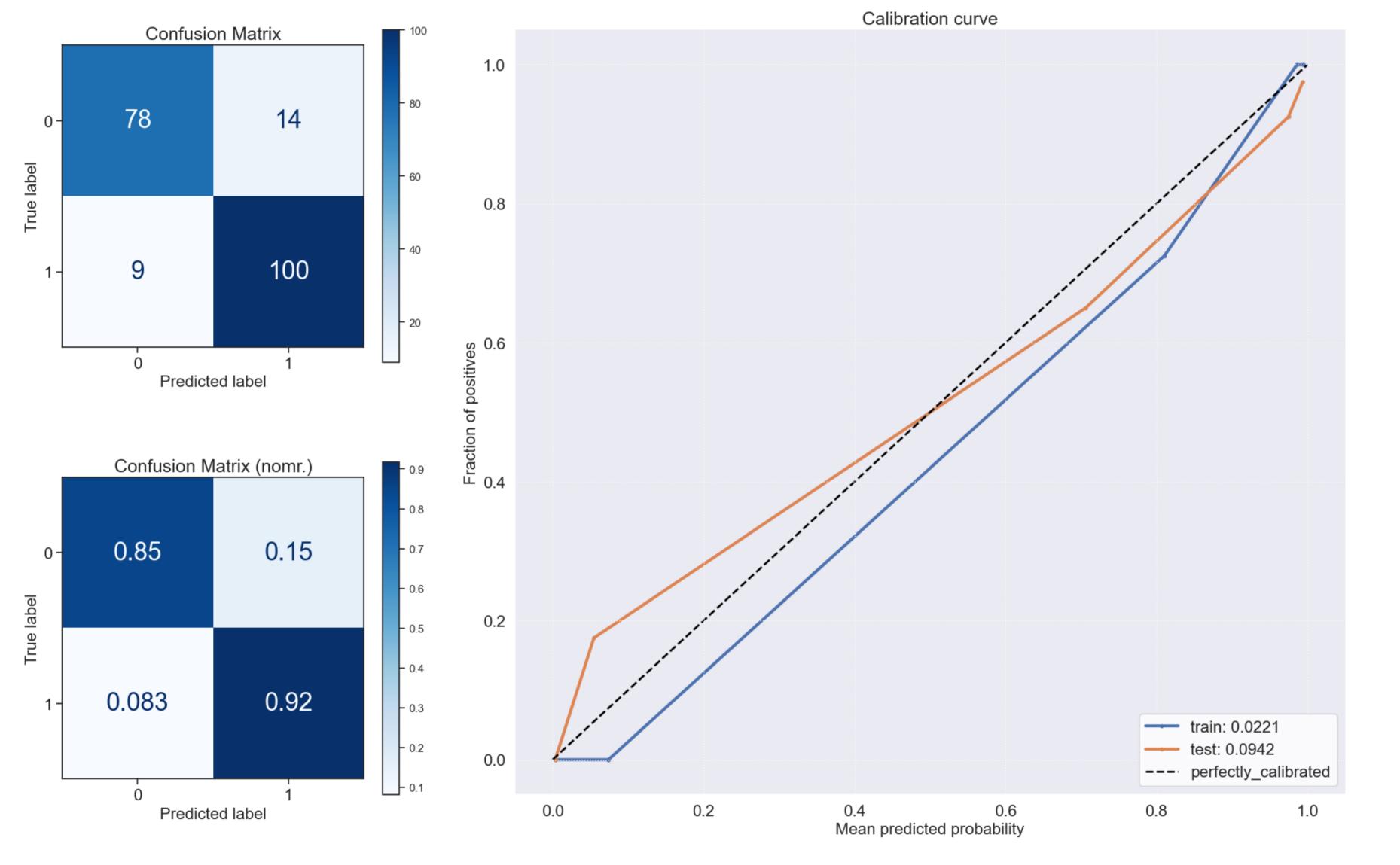
Metric	Train	Test
AUC	0.9999	0.9546
accuracy	0.9913	0.8955
balanced accuracy	0.9904	0.8952
recall_(sens)	1.0	0.8991
specificity	0.9808	0.8913
precision	0.9842	0.9074
f1_score	0.9913	0.8956

## Protein: P08588 - metric: f1





XGBoost Parameters	Value
alpha	2.4115
colsample_bytree	0.95
gamma	0.1309
lambda	0.4448
learning_rate	0.0146
max_depth	9.0
min_child_weight	2.0
n_estimators	575.0
reg_alpha	0.3426
reg_lambda	0.0272
scale_pos_weight	2.8993
subsample	0.7619

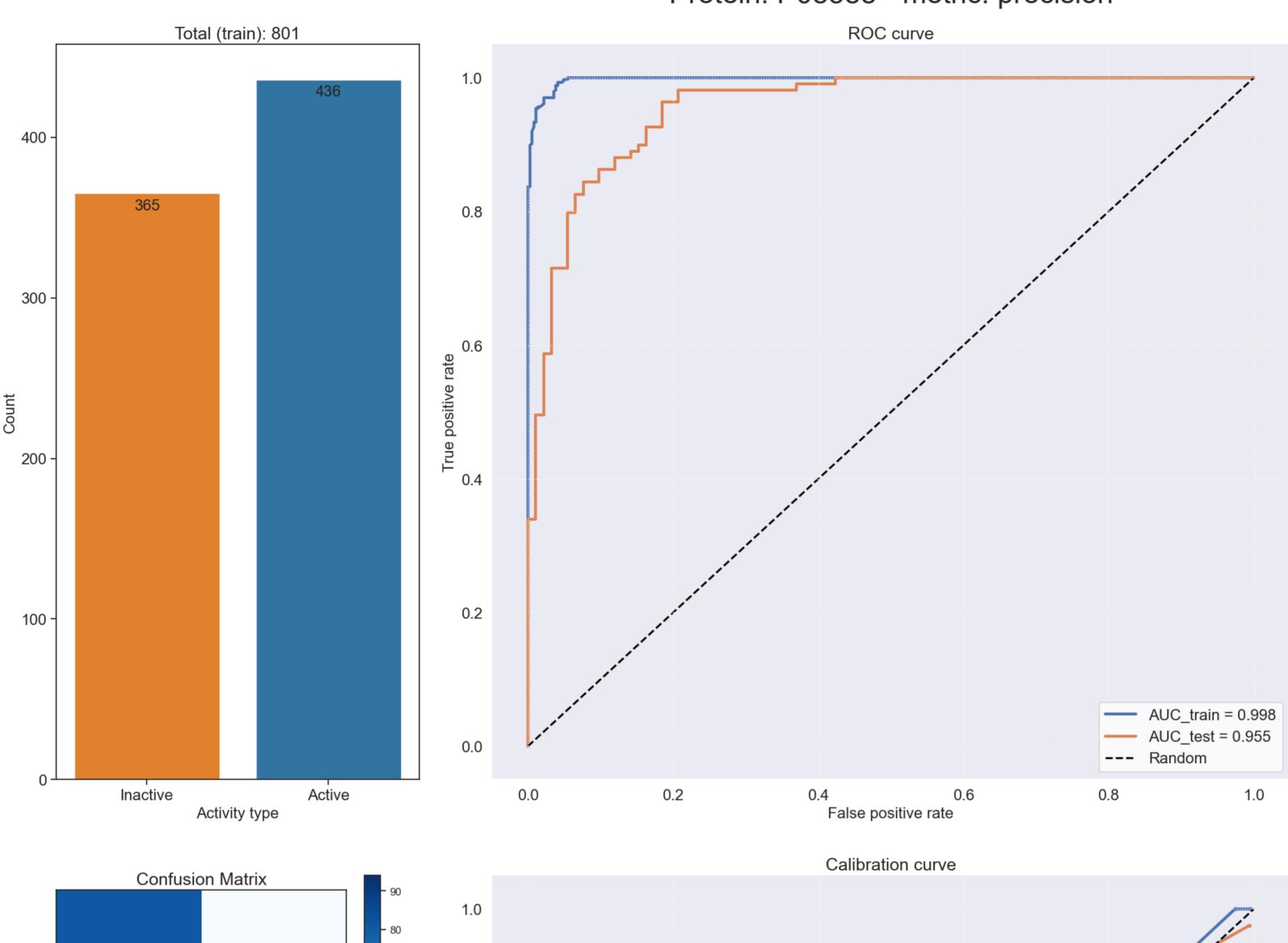


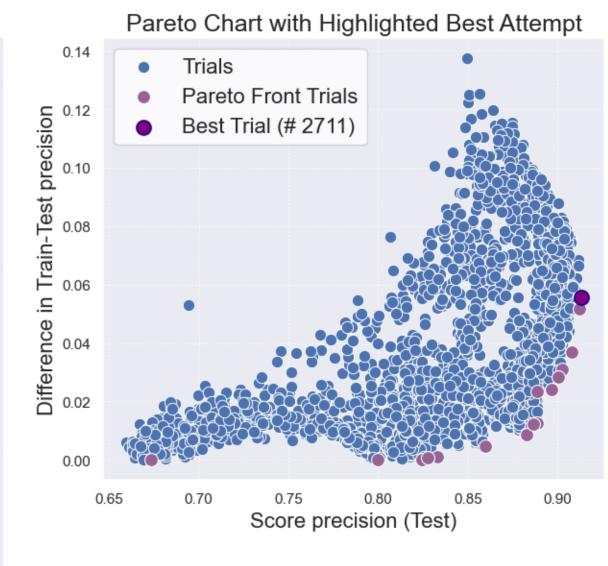
50	Active/mactive probability						
40		nactive zone			Active zone	type active (109) inactive (92) decoy (201)	
30							
Percent 00							
10							
0 —	0.0	0.2	0.4	0.6	0.8	1.0	
			Proba		3.3		

Active/Inactive probability

Metric	Train	Test
AUC	0.9986	0.9498
accuracy	0.9725	0.8856
balanced accuracy	0.9699	0.8826
recall_(sens)	1.0	0.9174
specificity	0.9397	0.8478
precision	0.952	0.8772
f1_score	0.9724	0.8853

## Protein: P08588 - metric: precision





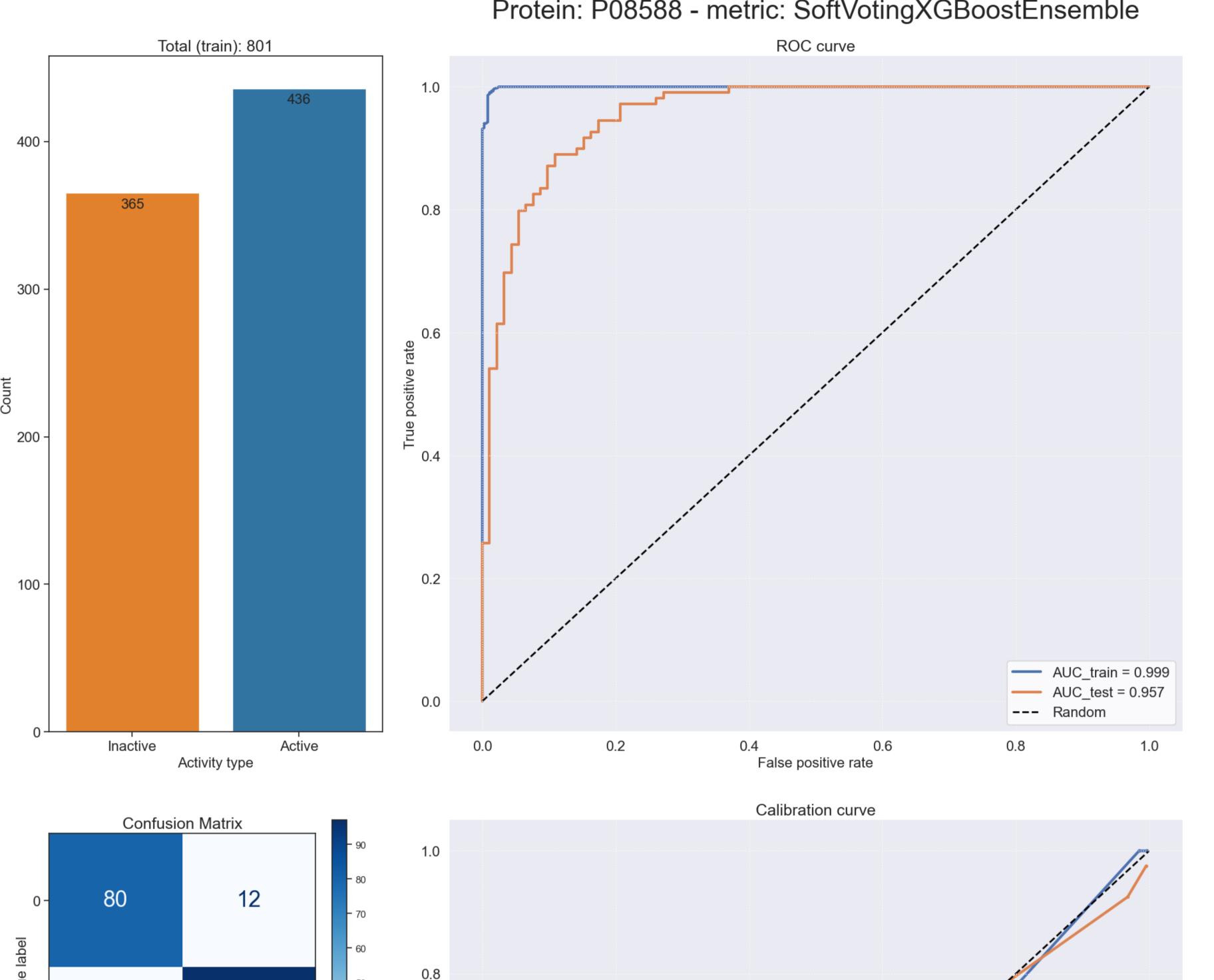
XGBoost Parameters	Value
alpha	2.3133
colsample_bytree	0.6094
gamma	0.511
lambda	4.1654
learning_rate	0.2238
max_depth	9.0
min_child_weight	2.0
n_estimators	379.0
reg_alpha	0.7061
reg_lambda	0.1606
scale_pos_weight	1.025
subsample	0.7457

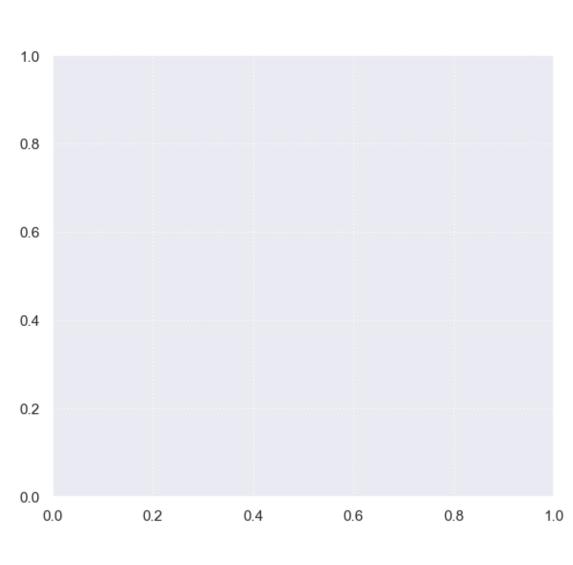
			90	1.0						
abel O	81	11	- 80 - 70 - 60	1.0						
True label	15	94	- 50 - 40 - 30 - 20	0.8						
ı	0 Predicte	1 ed label		Fraction of positives 0 0						
	Confusion M	latrix (nomr.)		ractio						
label O		0.12	- 0.8 - 0.7 - 0.6 - 0.5	0.2						
True label	0.14	0.86	- 0.5 - 0.4 - 0.3 - 0.2	0.0						train: 0.0207 test: 0.0921 perfectly_calibrated
	0 Predicte	1 ed label			0.0	0.2	0.4 Mean predicted prob	0.6 pability	0.8	1.0

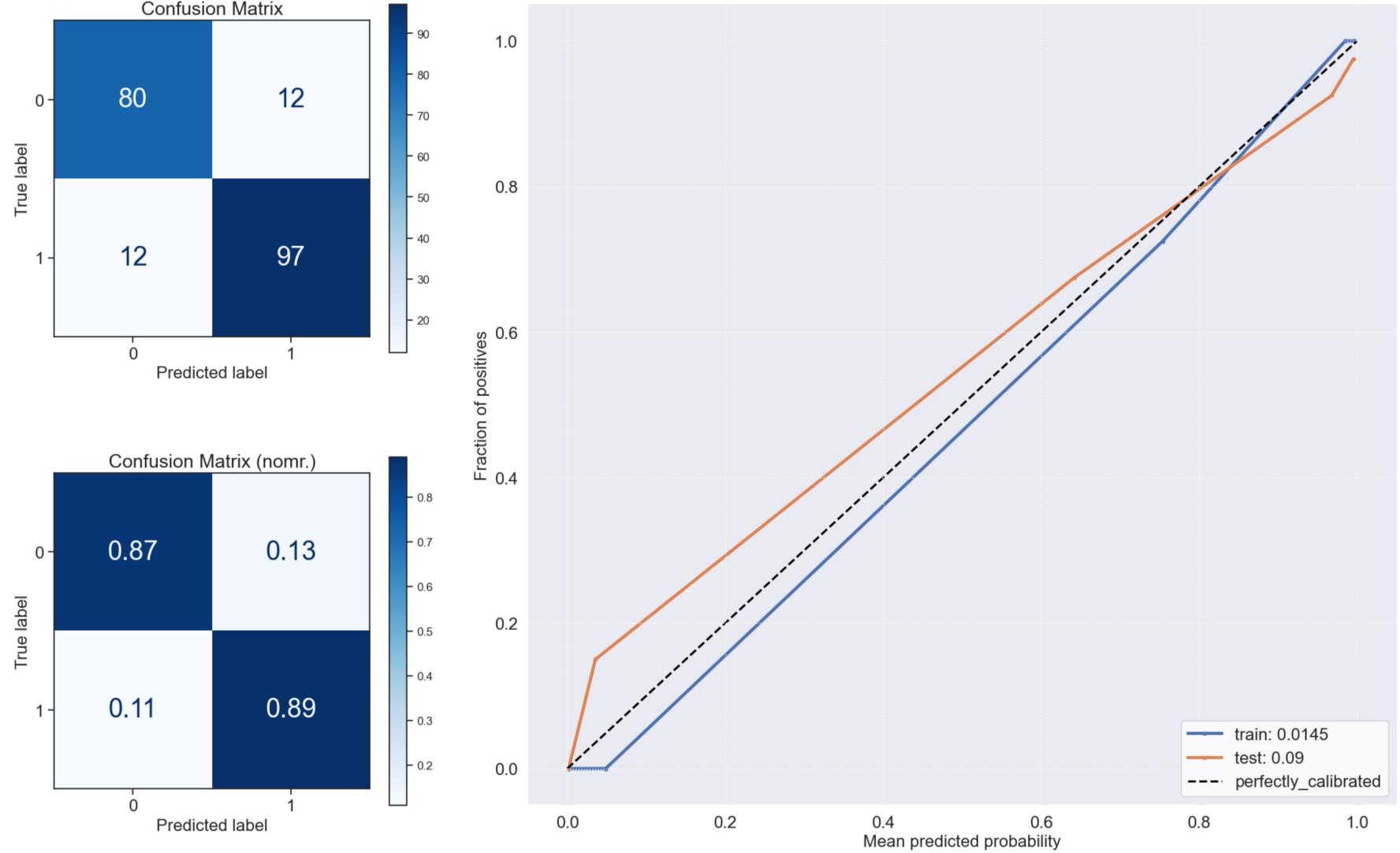
		Active/Ina	ctive probability		
50	Inact	ive zone		Active zone	type active (109) inactive (92) decoy (201)
40 <del>=</del> 30					
Percent 30					
10					
0 —	0.0 0.2	0.4	0.6	0.8	1.0
		Pr	obability		

Metric	Train	Test
AUC	0.9979	0.9554
accuracy	0.9738	0.8706
balanced accuracy	0.9728	0.8714
recall_(sens)	0.9839	0.8624
specificity	0.9616	0.8804
precision	0.9684	0.8952
f1_score	0.9738	0.8708

## Protein: P08588 - metric: SoftVotingXGBoostEnsemble







		Ad	ctive/Inactive	e probability		
50		ictive zone				type active (109) inactive (92) decoy (201)
40						
10						
0	.0 0	.2	D.4 Probal	0.6 pility	0.8	1.0

Metric	Train	Test
AUC	0.9994	0.9565
accuracy	0.9825	0.8806
balanced accuracy	0.9808	0.8797
recall_(sens)	1.0	0.8899
specificity	0.9616	0.8696
precision	0.9689	0.8899
f1_score	0.9825	0.8806