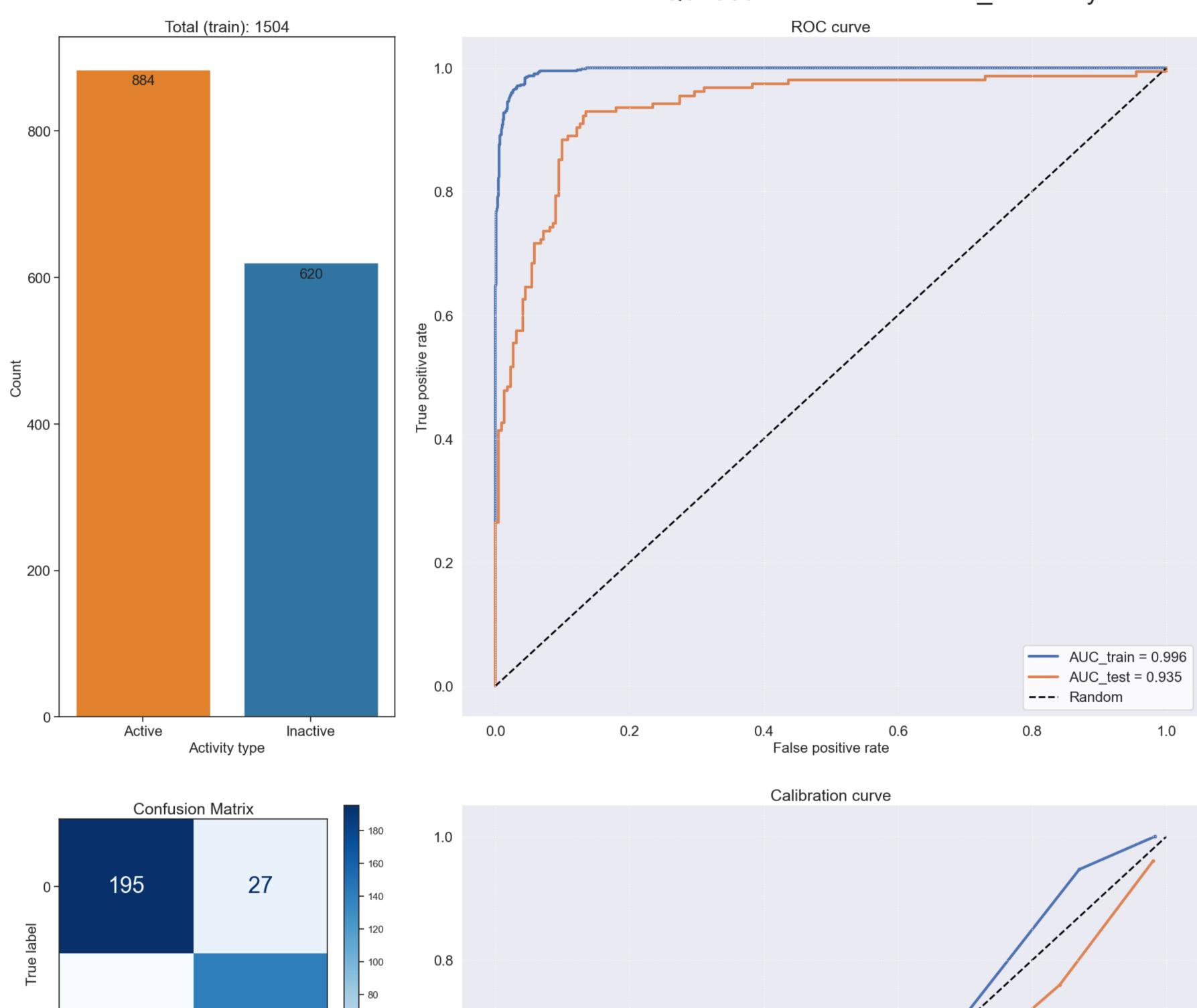
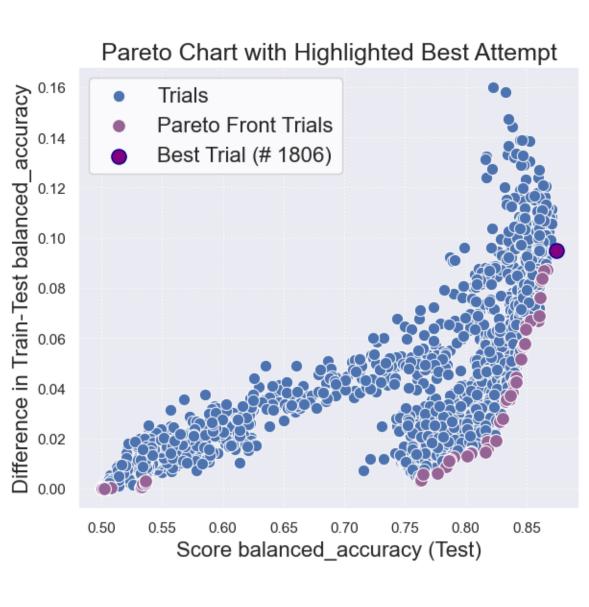
## Protein: Q07869 - metric: balanced\_accuracy





XGBoost Parameters	Value
alpha	3.8932
colsample_bytree	0.8901
gamma	0.0346
lambda	1.7306
learning_rate	0.0416
max_depth	10.0
min_child_weight	3.0
n_estimators	635.0
reg_alpha	0.7349
reg_lambda	0.0227
scale_pos_weight	1.6153
subsample	0.5057

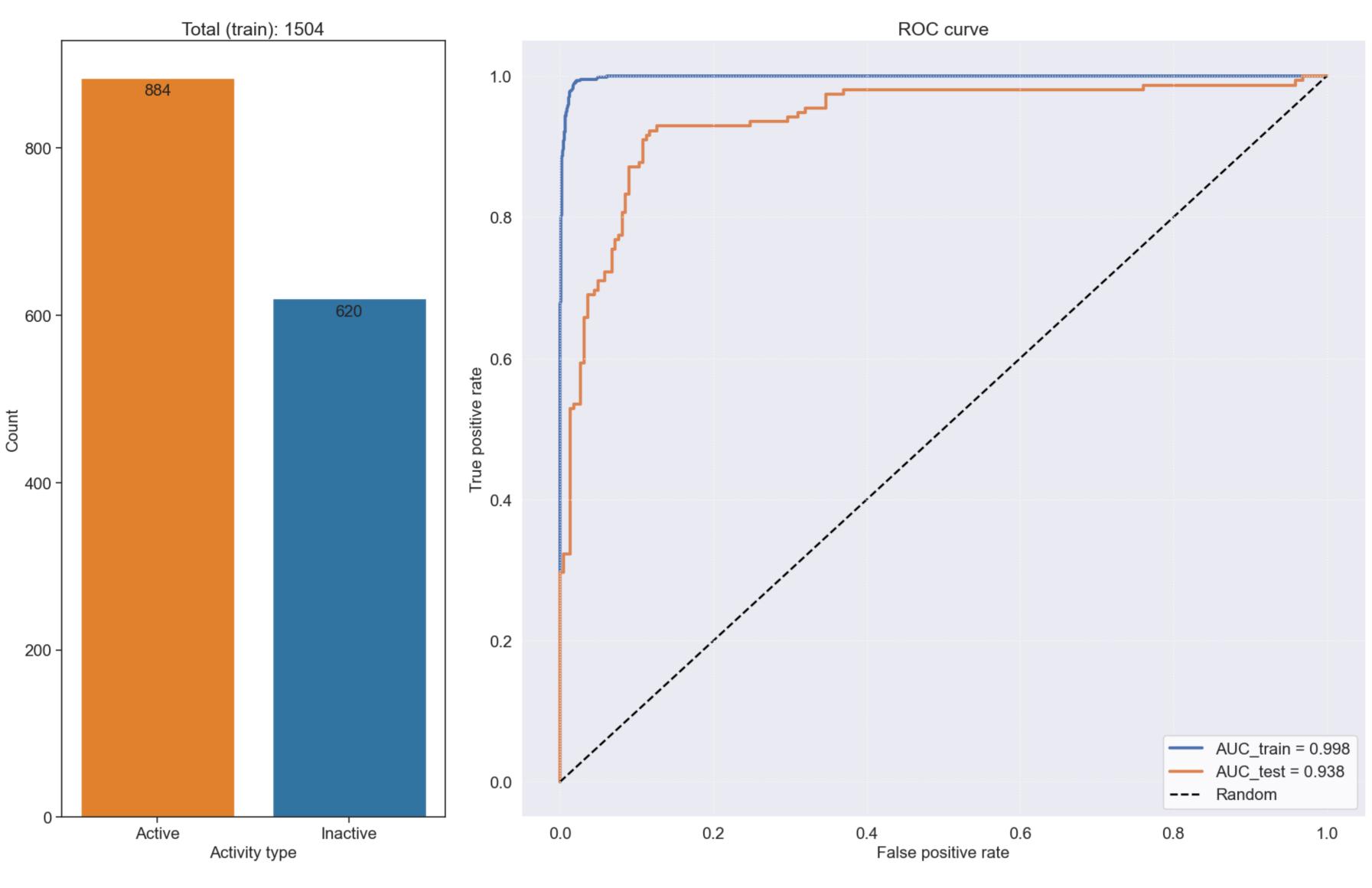
True	15	140	- 100 - 80 - 60 - 40	0.8				
	0 Predicte	1 ed label	4	on of positives				
	Confusion M	latrix (nomr.)	0.9	Laction 4.0				
label O	0.88	0.12	- 0.8 - 0.7 - 0.6 - 0.5	0.4				
True	0.097	0.9	- 0.4 - 0.3 - 0.2	0.0				train: 0.0304 test: 0.0921 perfectly_calibrated
	0 Predicte	1 ed label	□ 0.1		0.0 0.2	0.4 Mean predicted probab	0.6 pility	0.8 1.0

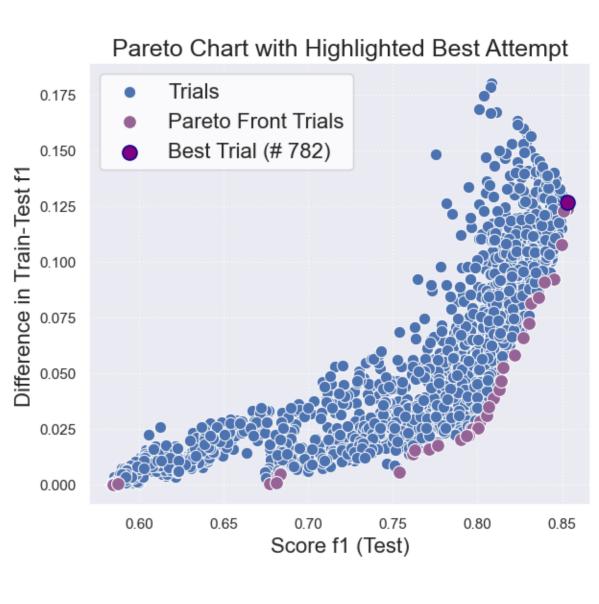
20.0		Inactive zone			Active zo	one active	
17.5						decoy	(377)
15.0							
12.5							
Percent 0.01							
7.5							
5.0							
2.5							
0.0	0.0	0.2	0.4 Proba	0.6 bility	0.8		1.0

Active/Inactive probability

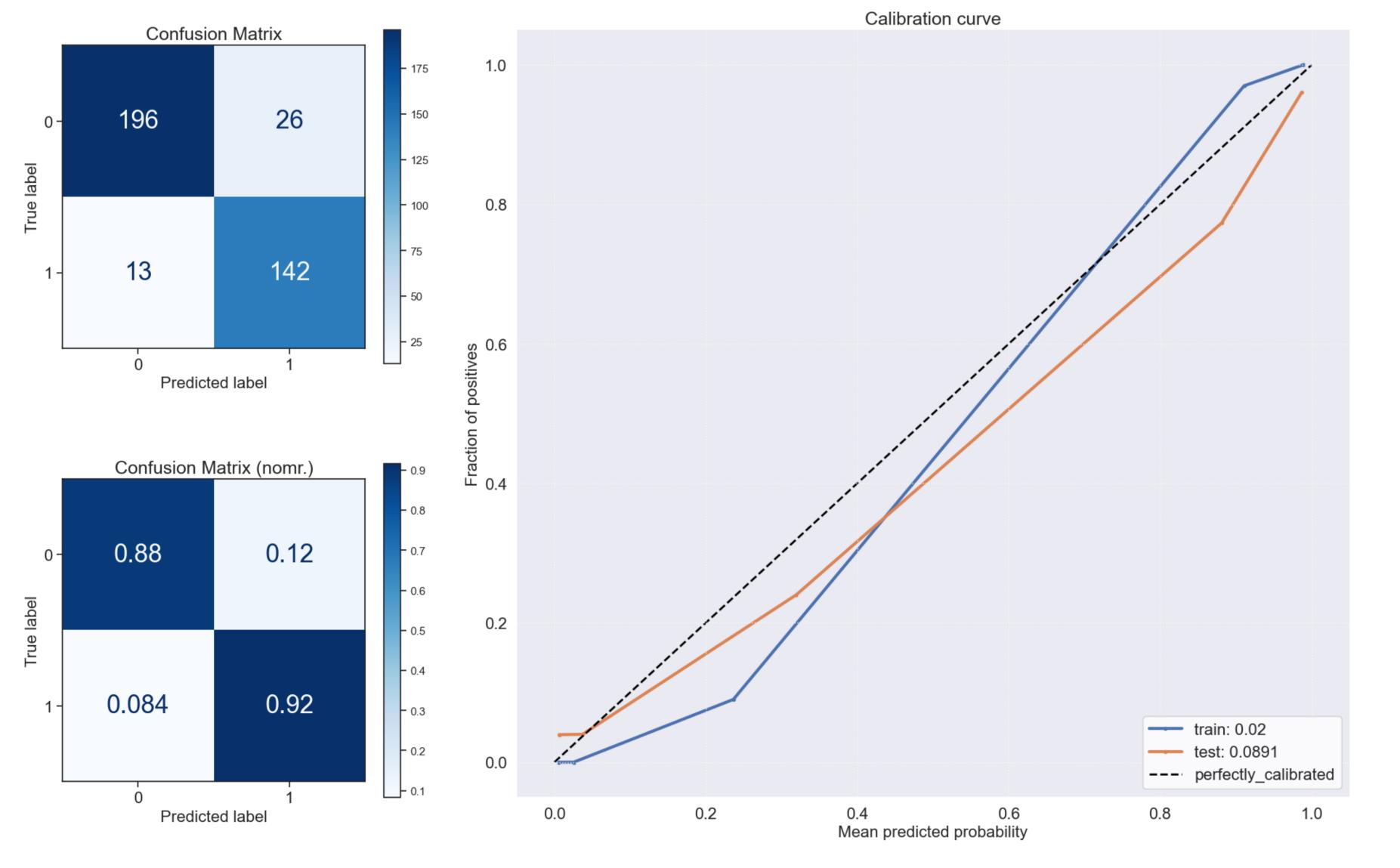
Metric	Train	Test
AUC	0.9961	0.9355
accuracy	0.9661	0.8886
balanced accuracy	0.9683	0.8908
recall_(sens)	0.9806	0.9032
specificity	0.9559	0.8784
precision	0.9397	0.8383
f1_score	0.9662	0.8891

## Protein: Q07869 - metric: f1





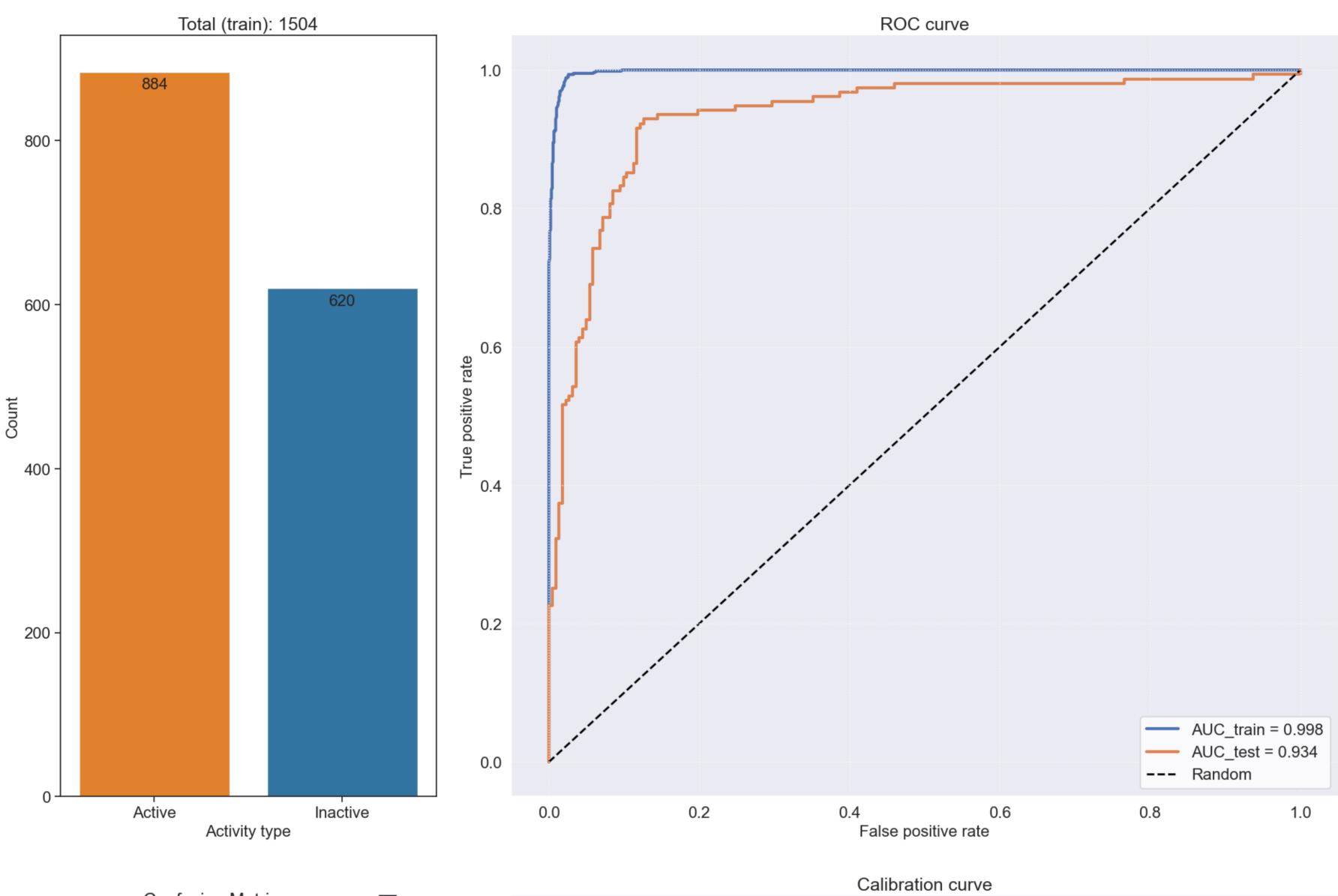
XGBoost Parameters	Value
alpha	2.219
colsample_bytree	0.5377
gamma	0.0037
lambda	3.0693
learning_rate	0.0381
max_depth	9.0
min_child_weight	1.0
n_estimators	604.0
reg_alpha	0.6293
reg_lambda	0.4899
scale_pos_weight	1.9593
subsample	0.649

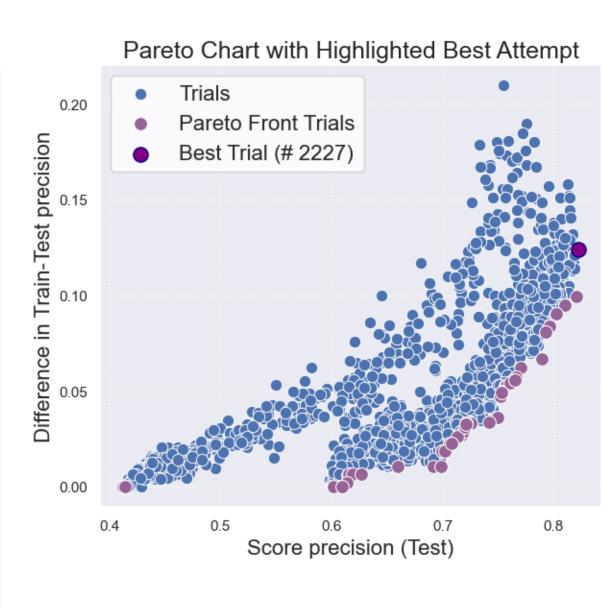


			Active/Inactive pr	obability		
25	Ina	active zone			Active zone	type active (155) inactive (222) decoy (377)
Percent 55						
10 <del></del>						
5 <del></del>						
	0.0	).2	0.4 Probability	0.6	0.8	1.0

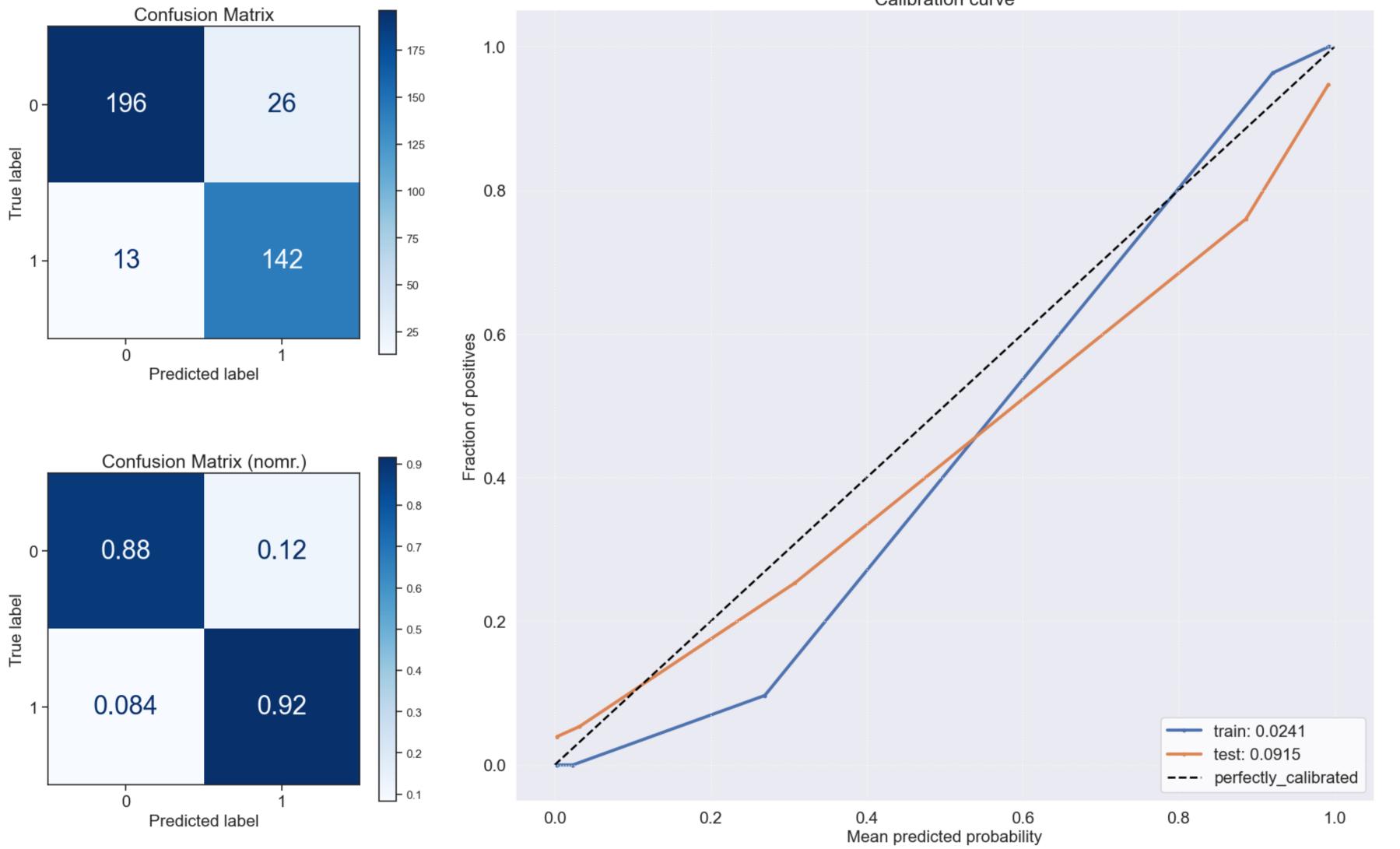
Metric	Train	Test
AUC	0.9984	0.9381
accuracy	0.9807	0.8966
balanced accuracy	0.9829	0.8995
recall_(sens)	0.9952	0.9161
specificity	0.9706	0.8829
precision	0.9596	0.8452
f1_score	0.9808	0.8971

## Protein: Q07869 - metric: precision





XGBoost Parameters	Value
alpha	4.6742
colsample_bytree	0.8802
gamma	0.6787
lambda	4.4581
learning_rate	0.1114
max_depth	8.0
min_child_weight	4.0
n_estimators	493.0
reg_alpha	0.1061
reg_lambda	0.1717
scale_pos_weight	2.2971
subsample	0.7011

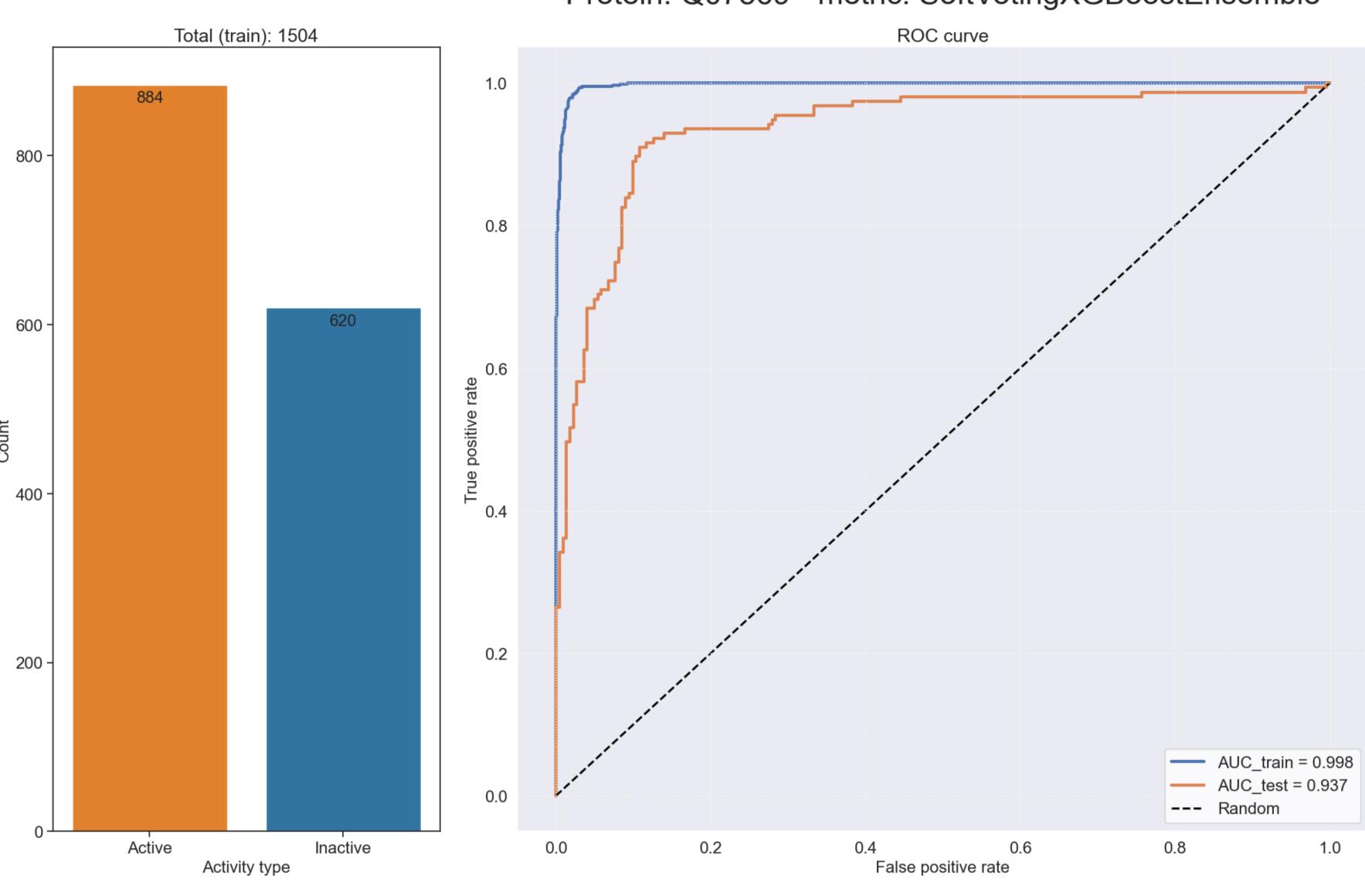


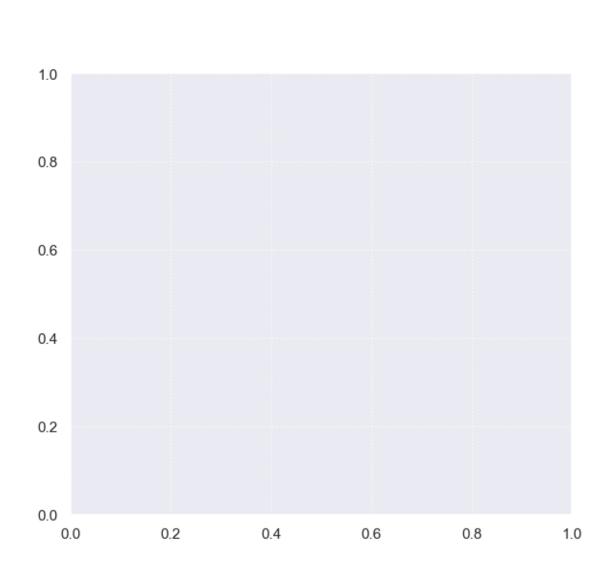
35		ictive zone				zone ac	ype ctive (155) active (222) ecoy (377)
30							
±25							
20 15							
10							
5							
0							L
,	0.0 0	.2 (	).4 Proba	0.6 ability	0.	8	1.0

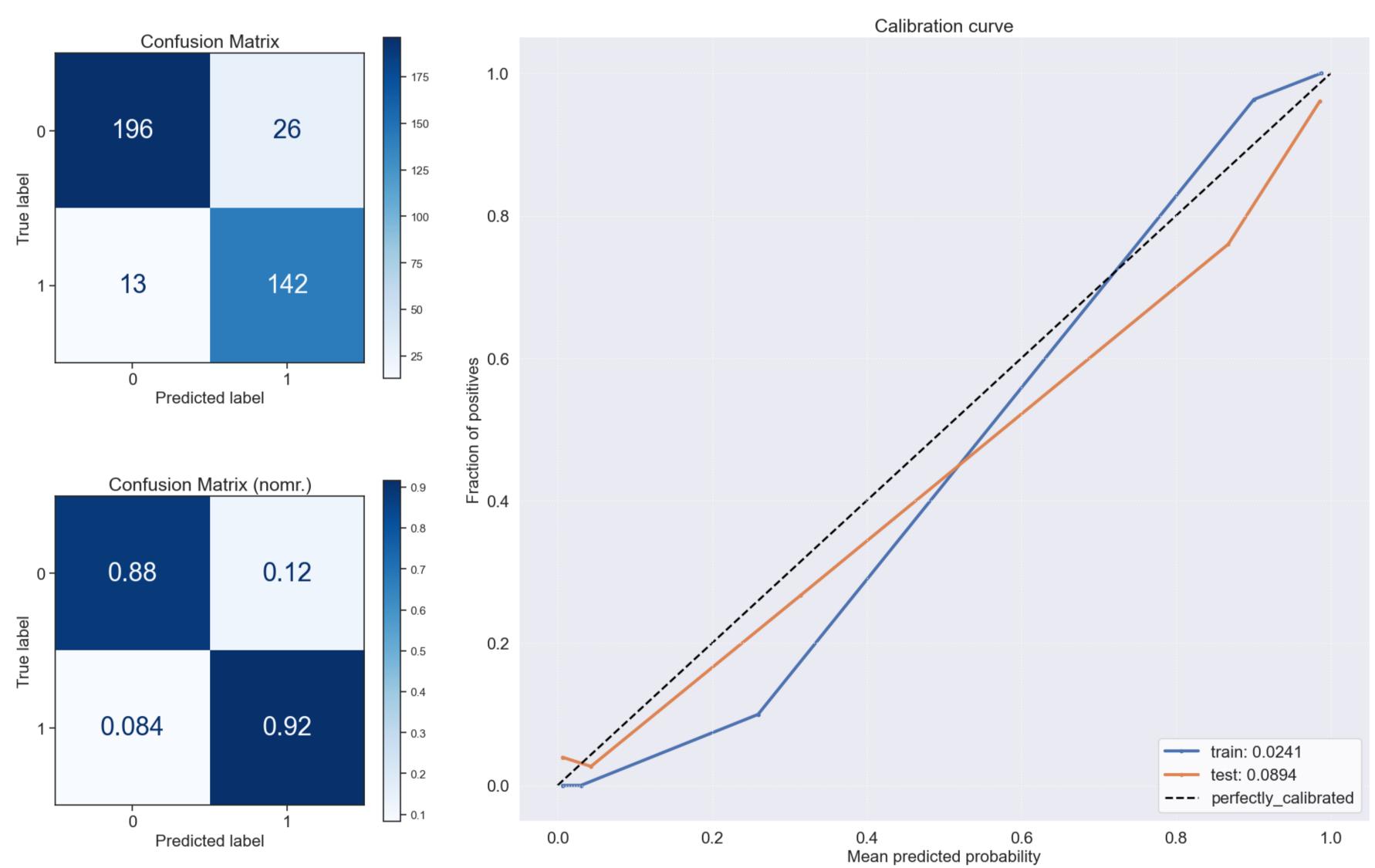
Active/Inactive probability

Metric	Train	Test
AUC	0.9978	0.9341
accuracy	0.9734	0.8966
balanced accuracy	0.9767	0.8995
recall_(sens)	0.9952	0.9161
specificity	0.9581	0.8829
precision	0.9434	0.8452
f1_score	0.9735	0.8971

## Protein: Q07869 - metric: SoftVotingXGBoostEnsemble







		Ac	tive/Inactive	probability	у		
25	Ina	active zone			Active z	one	type active (155) inactive (222) decoy (377)
20							
Percent							
5							
0 —							
	0.0	0.2	.4 Probab	0.6 oility	0.8	გ გ	1.0

Metric	Train	Test
AUC	0.9977	0.9365
accuracy	0.9774	0.8966
balanced accuracy	0.98	0.8995
recall_(sens)	0.9952	0.9161
specificity	0.9649	0.8829
precision	0.9522	0.8452
f1_score	0.9775	0.8971