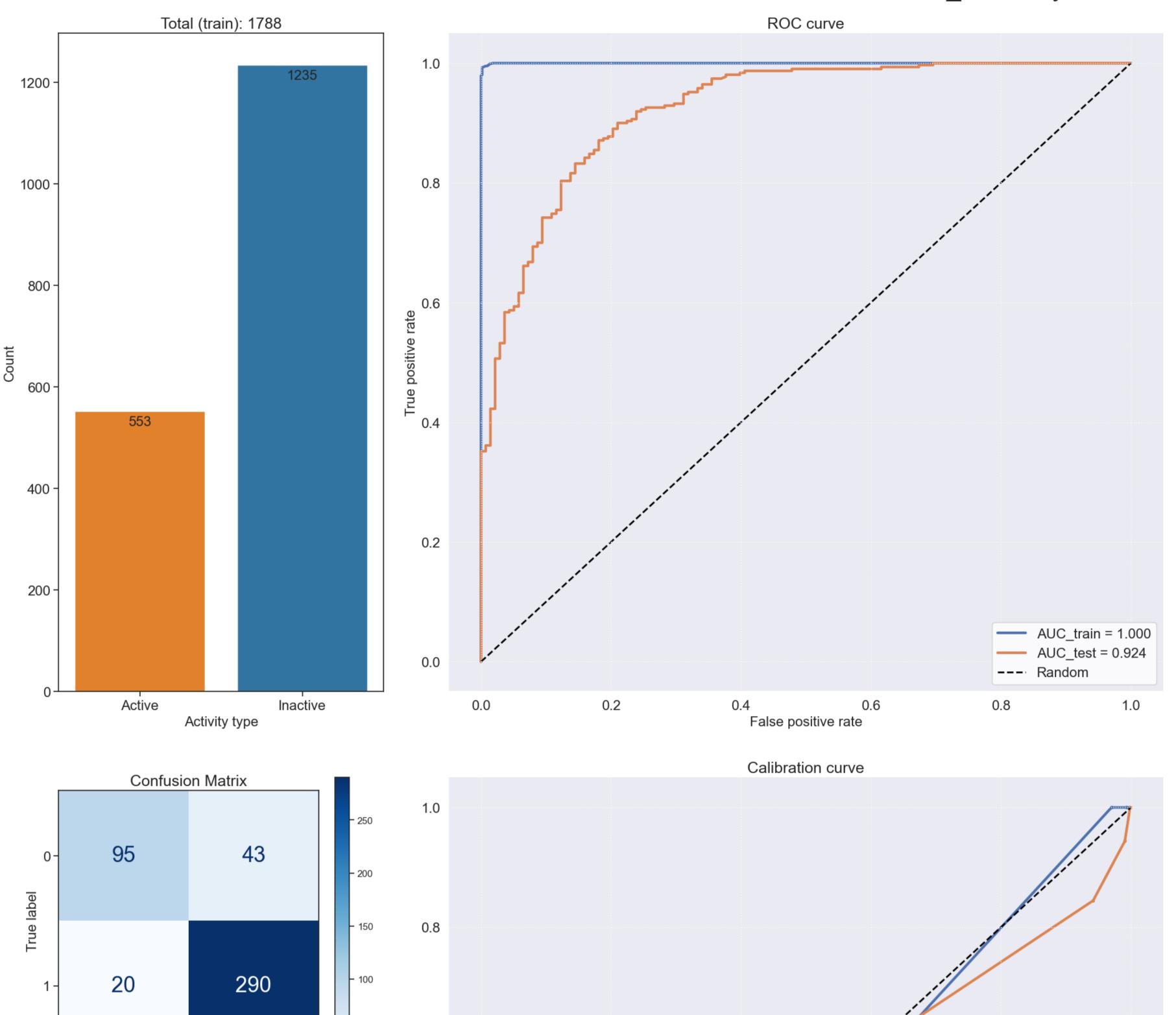
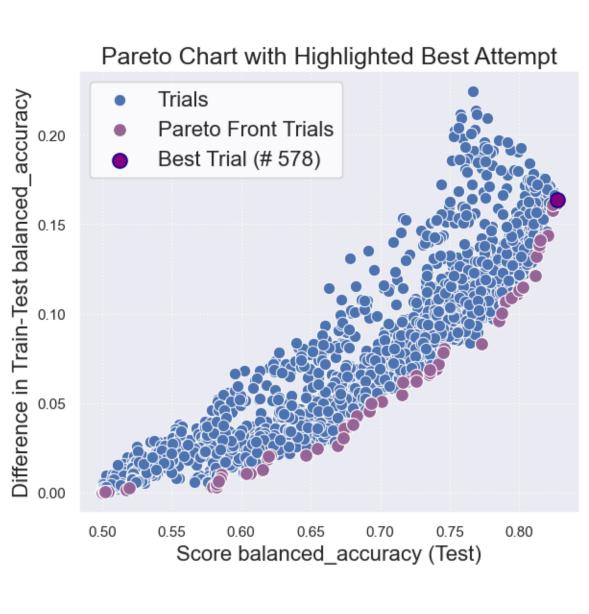
Protein: P28335 - metric: balanced_accuracy





XGBoost Parameters	Value
alpha	4.0852
colsample_bytree	0.6915
gamma	0.3595
lambda	2.0192
learning_rate	0.0923
max_depth	9.0
min_child_weight	1.0
n_estimators	942.0
reg_alpha	0.4933
reg_lambda	0.3234
scale_pos_weight	1.4468
subsample	0.5909

	Active/Inactive probability						
40	Ina	ctive zone			Active z		type active (310) inactive (138) decoy (447)
							decesy (447)
Percent							
10							
0							
	0.0	.2	0.4 Probal	0.6 bility	0.	8	1.0

- 50

- 0.8

- 0.6

- 0.5

0.4

0.3

0.2

Predicted label

Confusion Matrix (nomr.)

Predicted label

0.31

0.94

0.69

0.065

0-

True label

Fraction of positives

0.2

0.0

0.0

0.2

Metric	Train	Test
AUC	0.9999	0.9239
accuracy	0.9938	0.8594
balanced accuracy	0.9901	0.8119
recall_(sens)	1.0	0.9355
specificity	0.9801	0.6884
precision	0.9912	0.8709
f1_score	0.9938	0.8555

--- train: 0.0095

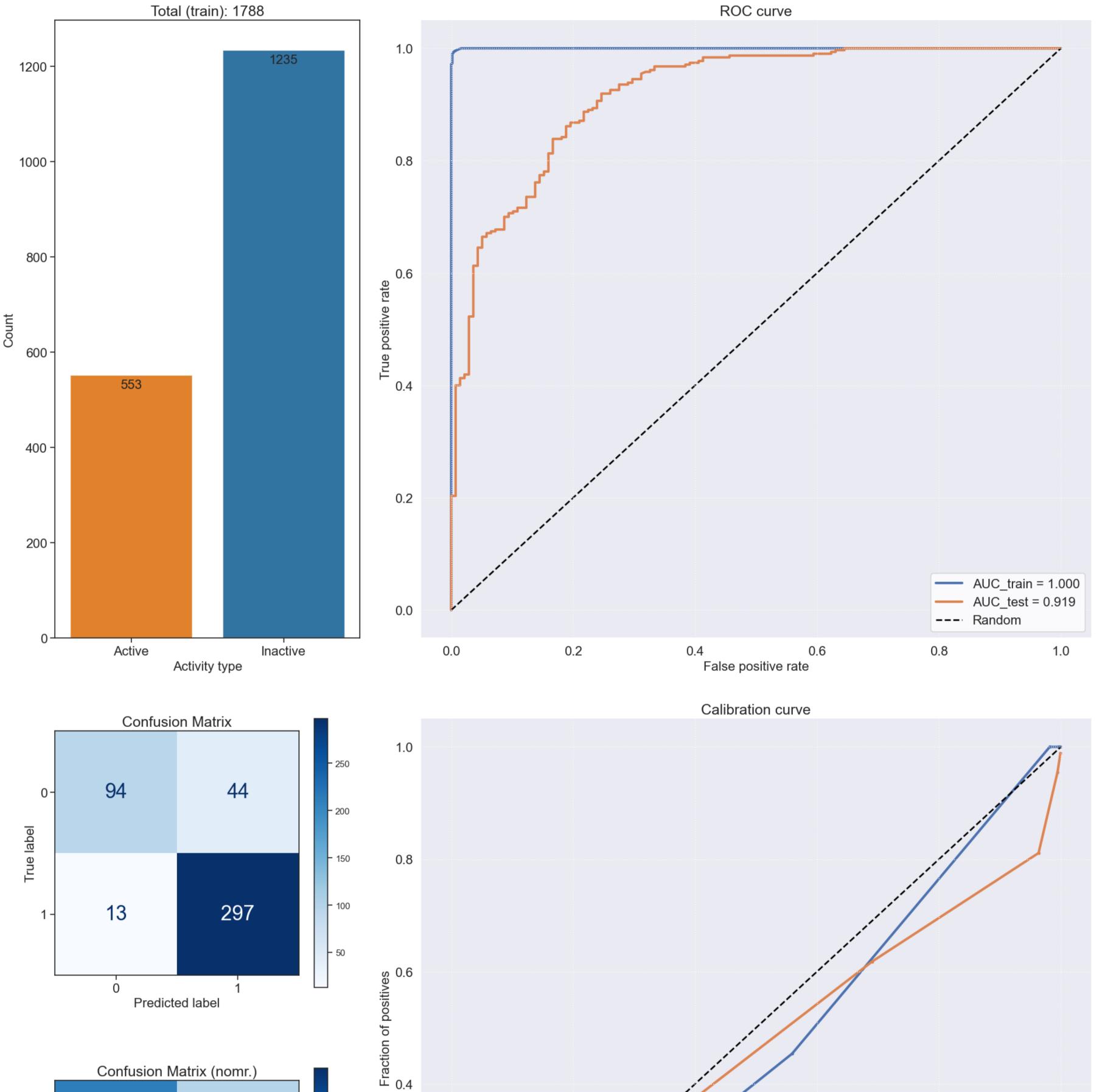
0.8

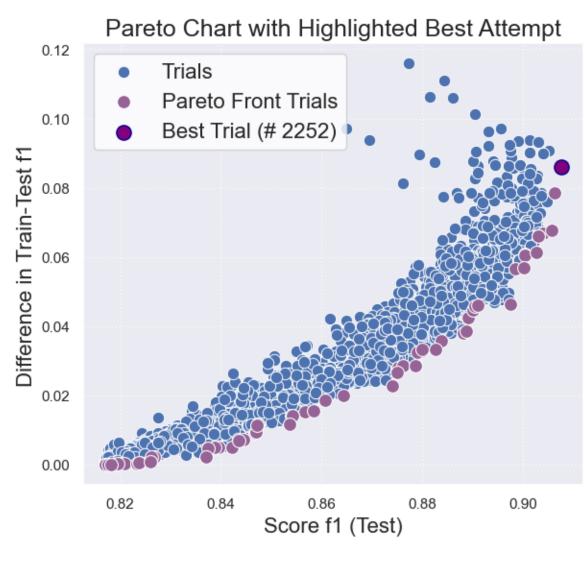
0.4 0.6 Mean predicted probability test: 0.0991

--- perfectly_calibrated

1.0

Protein: P28335 - metric: f1





XGBoost Parameters	Value
alpha	2.6167
colsample_bytree	0.6624
gamma	0.4658
lambda	1.934
learning_rate	0.0529
max_depth	8.0
min_child_weight	1.0
n_estimators	960.0
reg_alpha	0.2427
reg_lambda	0.4197
scale_pos_weight	2.9851
subsample	0.526

	Active/Inactive probability						
50	Ina	ctive zone		Active	zone act	pe ive (310) ctive (138) coy (447)	
Percent 08							
0	0.0 0.	2 0.4	4 (Probability).6 0	.8	1.0	

0.2

0.0

0.0

0.2

0.4

0.2

Confusion Matrix (nomr.)

Predicted label

0.32

0.96

0.68

0.042

0-

True label

Metric	Train	Test
AUC	0.9999	0.9191
accuracy	0.9899	0.8728
balanced accuracy	0.9837	0.8196
recall_(sens)	1.0	0.9581
specificity	0.9675	0.6812
precision	0.9856	0.871
f1_score	0.9899	0.8677

--- train: 0.0116

0.8

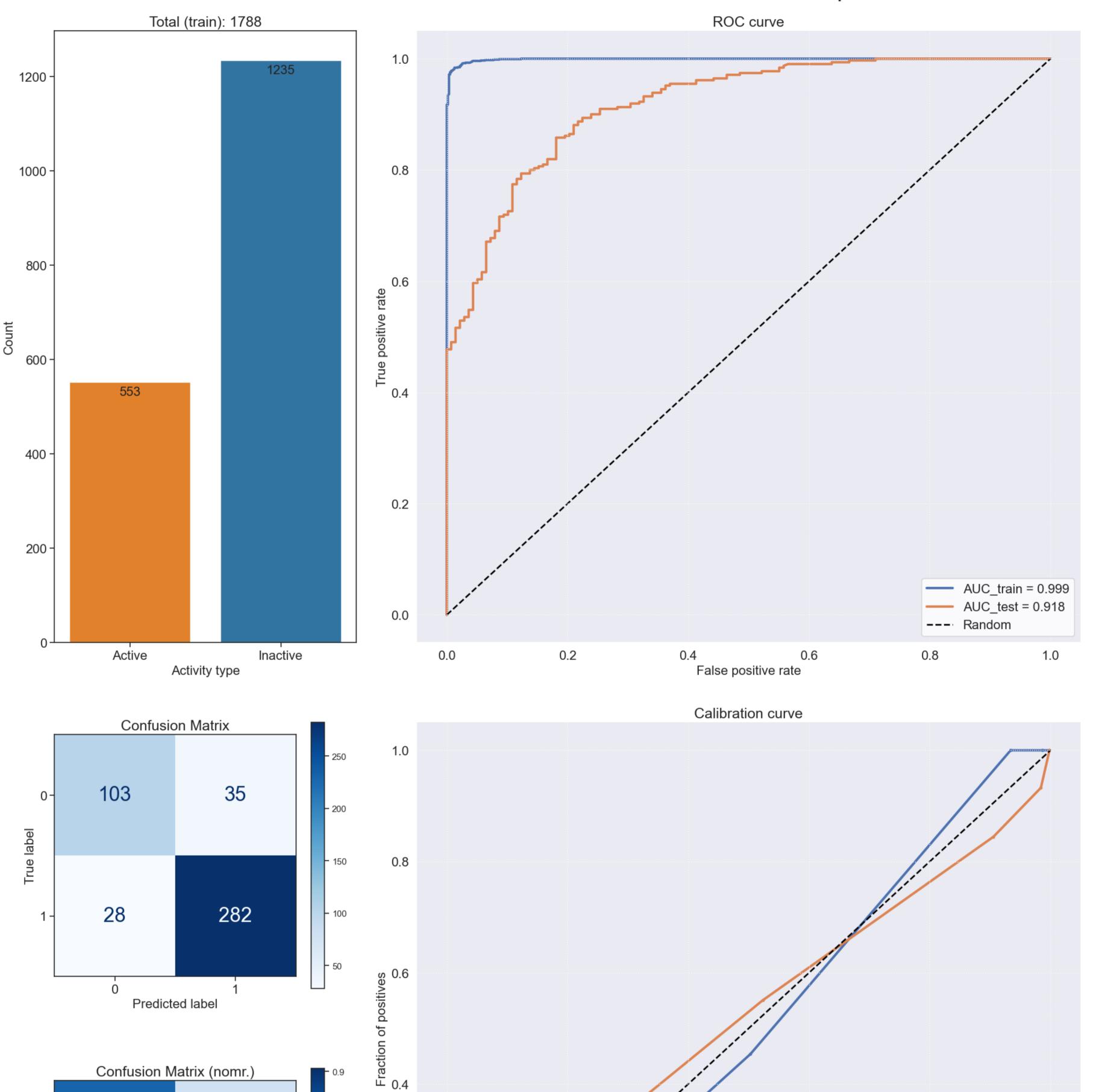
0.4 0.6 Mean predicted probability

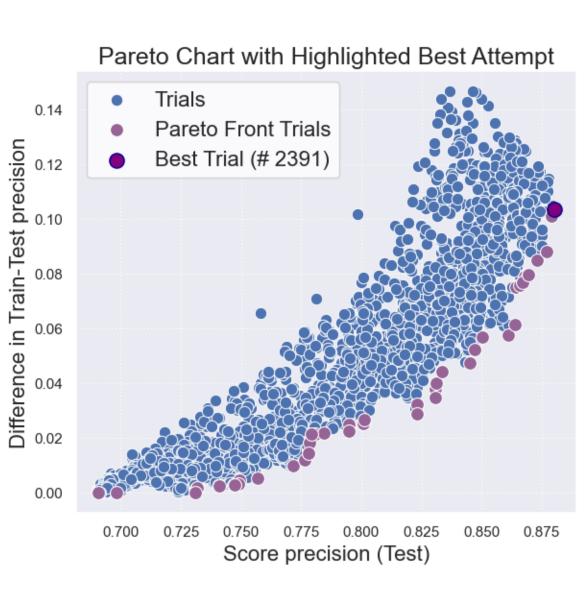
test: 0.1031

--- perfectly_calibrated

1.0

Protein: P28335 - metric: precision





XGBoost Parameters	Value
alpha	2.9929
colsample_bytree	0.9595
gamma	0.1978
lambda	2.4528
learning_rate	0.2287
max_depth	3.0
min_child_weight	3.0
n_estimators	754.0
reg_alpha	0.5149
reg_lambda	0.0868
scale_pos_weight	1.0302
subsample	0.7295

		Ac	tive/Inactiv	e probabil	ity		
40 35 	lna	ctive zone			Active	zone	type active (310) inactive (138) decoy (447)
30							
20							
Dercent 20							
10	_						
5				п			
0.	.0 0.	2 0).4 Proba	0.0 bility	6 0	0.8	1.0

- 0.8

- 0.6

- 0.5

0.4

0.3

0.2

0.2

0.0

0.0

0.2

Confusion Matrix (nomr.)

Predicted label

0.25

0.91

0.75

0.09

0-

True label

Metric	Train	Test
AUC	0.9991	0.918
accuracy	0.9827	0.8594
balanced accuracy	0.9745	0.828
recall_(sens)	0.996	0.9097
specificity	0.953	0.7464
precision	0.9793	0.8896
f1_score	0.9826	0.8583

--- train: 0.02

0.8

test: 0.108

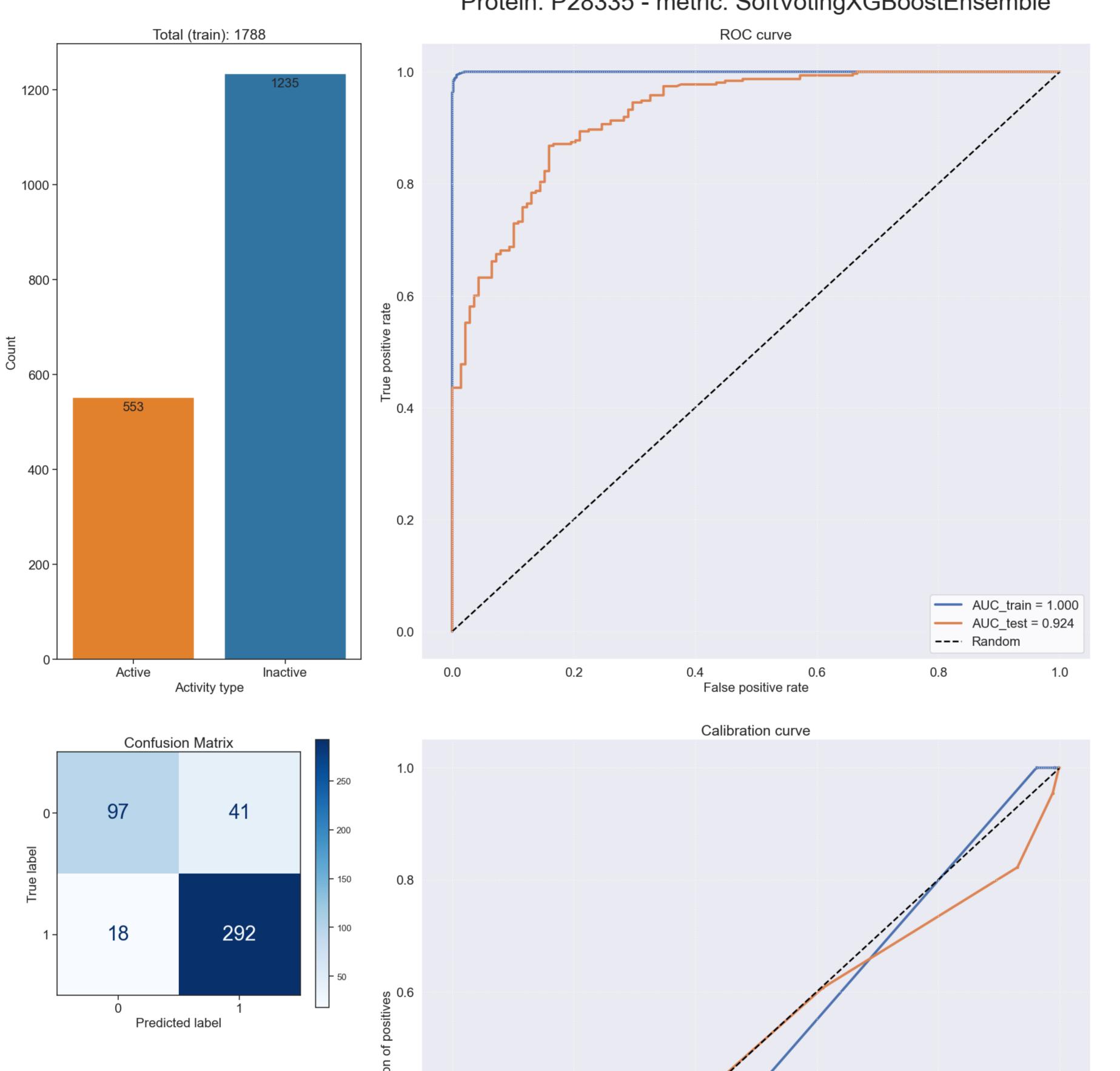
--- perfectly_calibrated

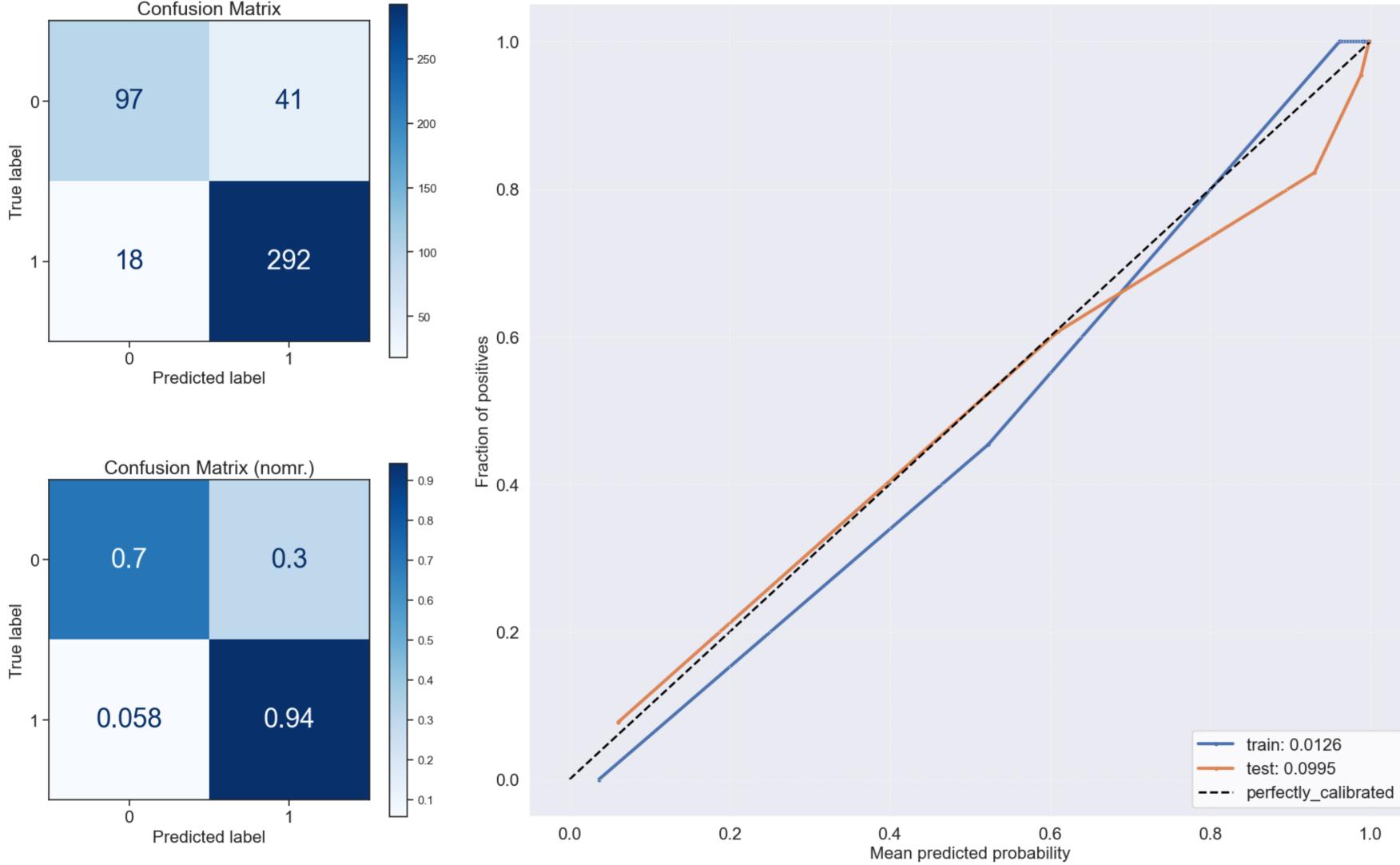
1.0

	Active/Inactive	ve probability						
40	Inactive zone	Act	ive zone	type active (310)	Metr	ic Tr	rain	Test
35				inactive (138) decoy (447)		AUC	0.9991	0.91
30					а	ccuracy	0.9827	0.859
25					balanced a	ccuracy	0.9745	0.82
20					recall	_(sens)	0.996	0.909
15					sp	ecificity	0.953	0.746
10					р	recision	0.9793	0.889
5					f	1_score	0.9826	0.858
0	0.0 0.2 0.4	0.6	0.8	1.0			-	

0.4 0.6 Mean predicted probability

Protein: P28335 - metric: SoftVotingXGBoostEnsemble





	Active/Inactive probability						
40 -		nactive zone			Active zone	type active (310) inactive (138) decoy (447)	
30 -							
- 02 Sercent							
0-							
	0.0	0.2	0.4	0.6	0.8	1.0	
	Probability						

Metric	Train	Test
AUC	0.9998	0.9243
accuracy	0.9916	0.8683
balanced accuracy	0.9864	0.8224
recall_(sens)	1.0	0.9419
specificity	0.9729	0.7029
precision	0.988	0.8769
f1_score	0.9916	0.8647

0.8

0.6

0.4

0.2

0.0

0.0

0.2

0.6