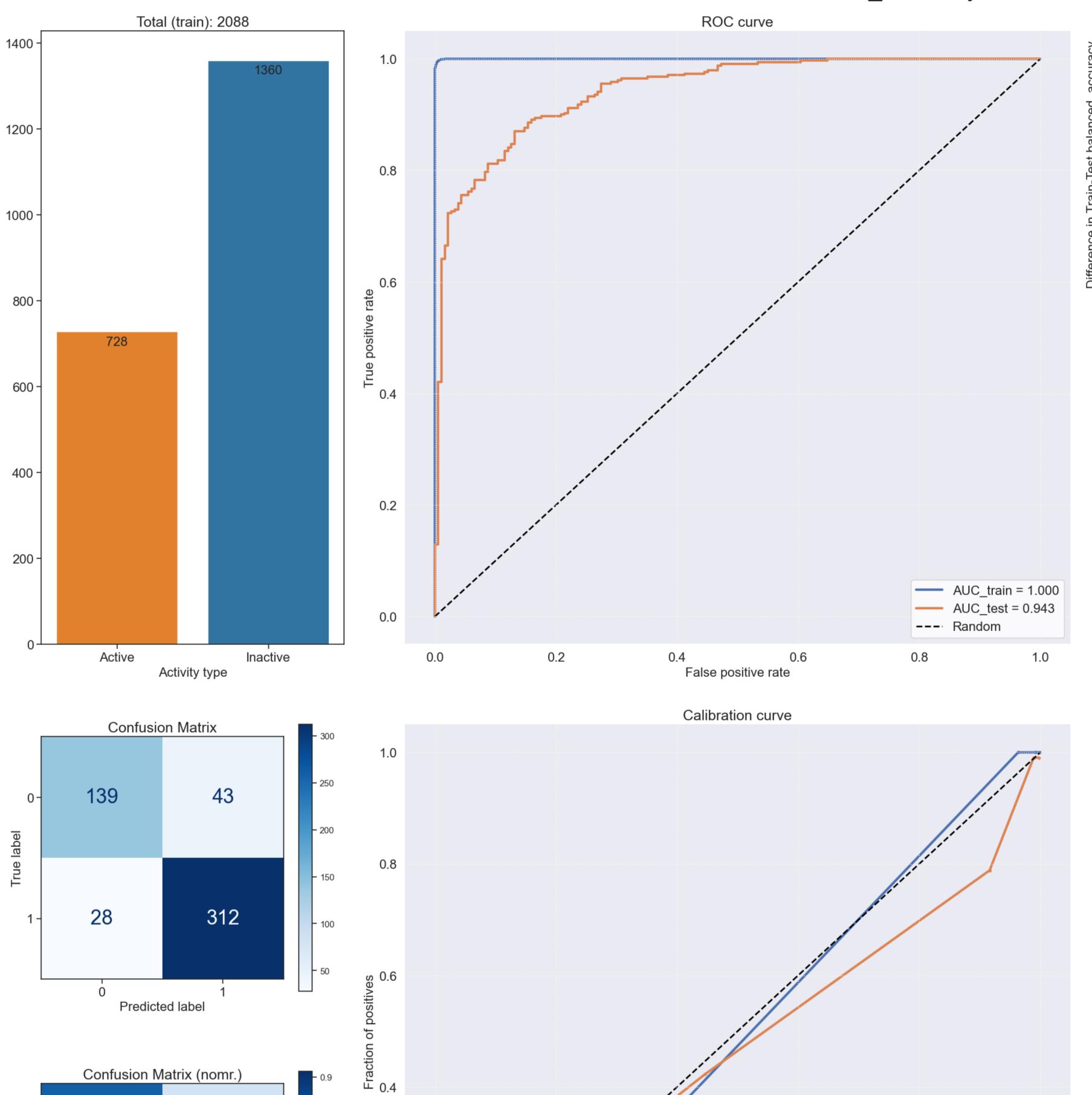
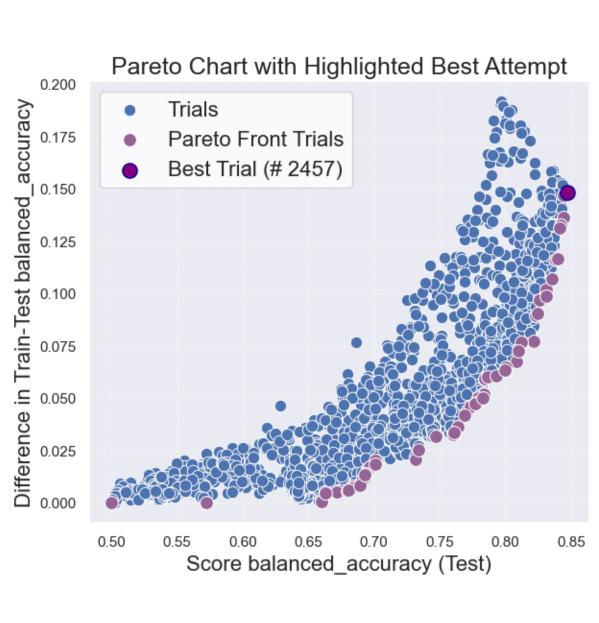
Protein: P27338 - metric: balanced_accuracy





XGBoost Parameters	Value
alpha	2.9067
colsample_bytree	0.5045
gamma	0.0659
lambda	3.5222
learning_rate	0.0743
max_depth	9.0
min_child_weight	1.0
n_estimators	992.0
reg_alpha	0.6193
reg_lambda	0.7244
scale_pos_weight	1.2094
subsample	0.7433

		Act	tive/Inactive p	robability		
40		ctive zone			Active zone	type active (340) inactive (182)
20						decoy (520)
Percent						
10						
0	0.0 0	.2 0.	4 Probabilit	0.6 y	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

0.24

0.92

0.76

0.082

0-

True label

Metric	Train	Test
AUC	0.9999	0.9434
accuracy	0.9957	0.864
balanced accuracy	0.9945	0.8407
recall_(sens)	0.9985	0.9176
specificity	0.9904	0.7637
precision	0.9949	0.8789
f1_score	0.9957	0.8625

active/irractive probability					
	Active zone	type active (340)	Metric	Train	Test
		inactive (182) decoy (520)	AUC	0.9999	0.9434
			accuracy	0.9957	0.864
			balanced accuracy	0.9945	0.8407
			recall_(sens)	0.9985	0.9176
			specificity	0.9904	0.7637
			precision	0.9949	0.8789
			f1_score	0.9957	0.8625
0.6	0.8	1.0			

0.4 0.6 Mean predicted probability

--- train: 0.007

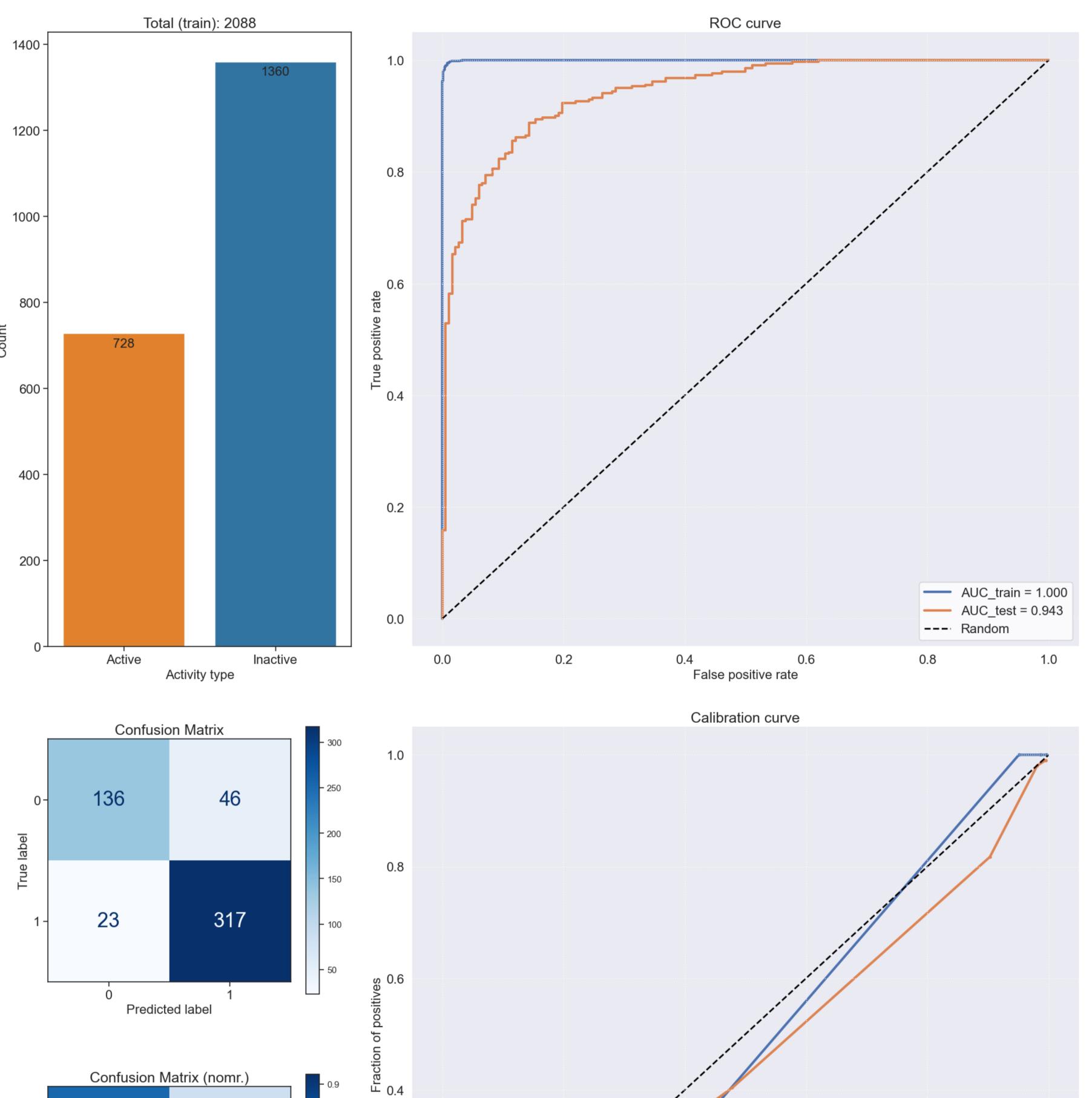
0.8

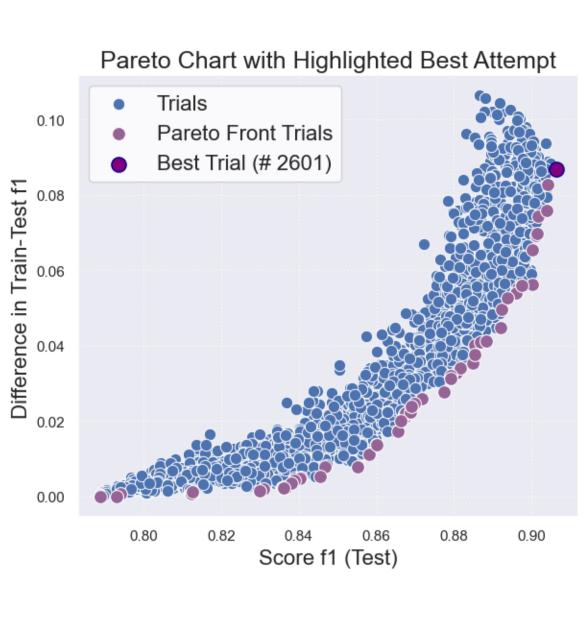
test: 0.0949

--- perfectly_calibrated

1.0

Protein: P27338 - metric: f1





XGBoost Parameters	Value
alpha	1.5326
colsample_bytree	0.6602
gamma	0.3019
lambda	3.1133
learning_rate	0.0318
max_depth	10.0
min_child_weight	1.0
n_estimators	823.0
reg_alpha	0.1337
reg_lambda	0.8143
scale_pos_weight	1.5198
subsample	0.7908

		Activ	e/Inactive probabi	lity		
30	Ina	ctive zone		Active :	zone act	pe ve (340) ctive (182)
25						coy (520)
ant 20						
Percent						
10						
5				┱ <mark>╌</mark> ╃┸┸╌╌┈┈		
0	0.0 0.	2 0.4	0. Probability	6 0	.8	1.0

- 0.8

- 0.6

- 0.5

0.4

0.3

0.2

0.1

0.2

0.0

0.0

0.2

Confusion Matrix (nomr.)

Predicted label

0.25

0.93

0.75

0.068

0-

True label

Metric	Train	Test
AUC	0.9998	0.9433
accuracy	0.9909	0.8678
balanced accuracy	0.9876	0.8398
recall_(sens)	0.9985	0.9324
specificity	0.9766	0.7473
precision	0.9876	0.8733
f1_score	0.9909	0.8655

Metric	IIaiii	1621
AUC	0.9998	0.9433
accuracy	0.9909	0.8678
balanced accuracy	0.9876	0.8398
recall_(sens)	0.9985	0.9324
specificity	0.9766	0.7473
precision	0.9876	0.8733
f1 score	0.9909	0.8655

--- train: 0.0126

0.8

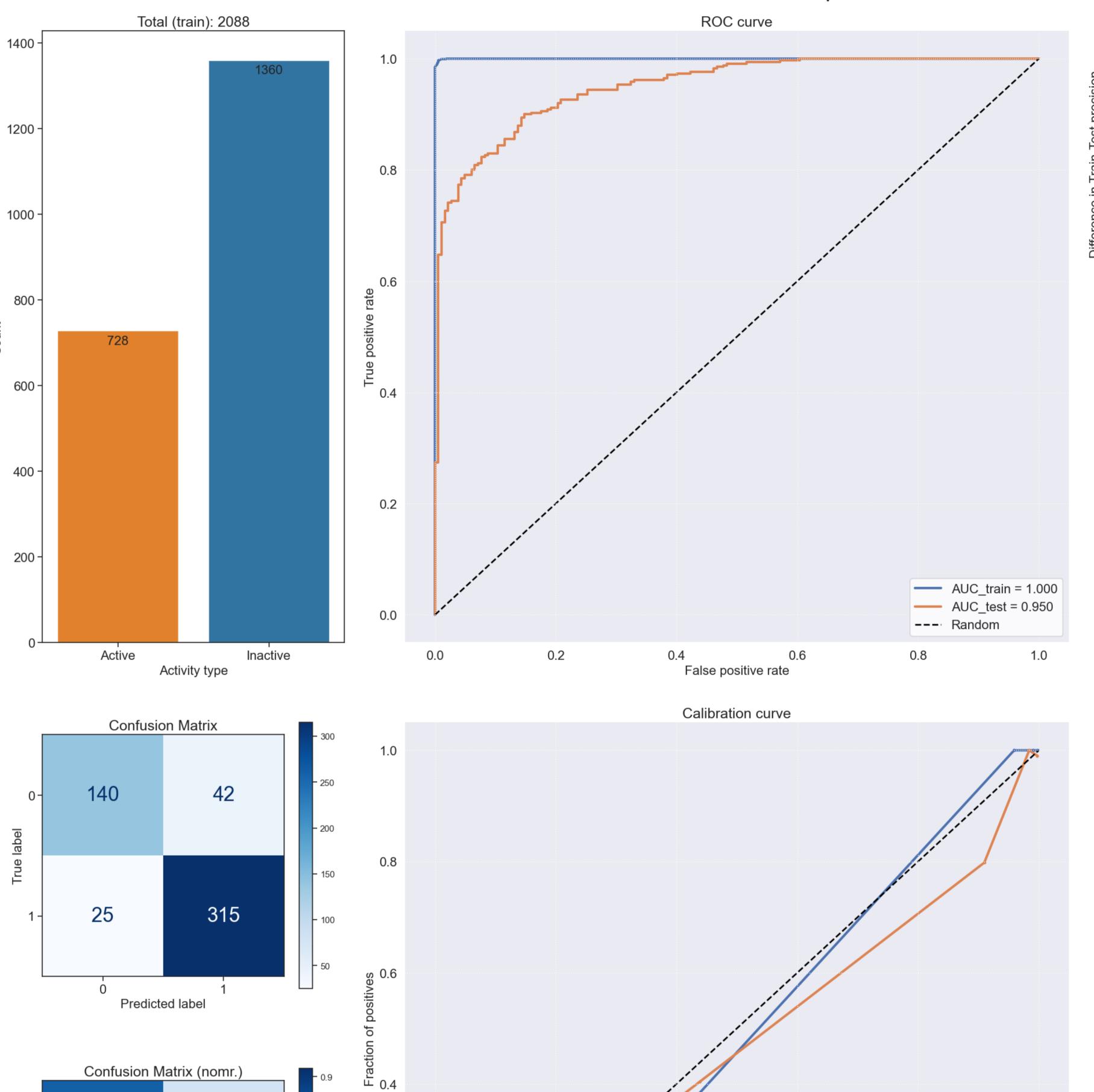
0.4 0.6 Mean predicted probability

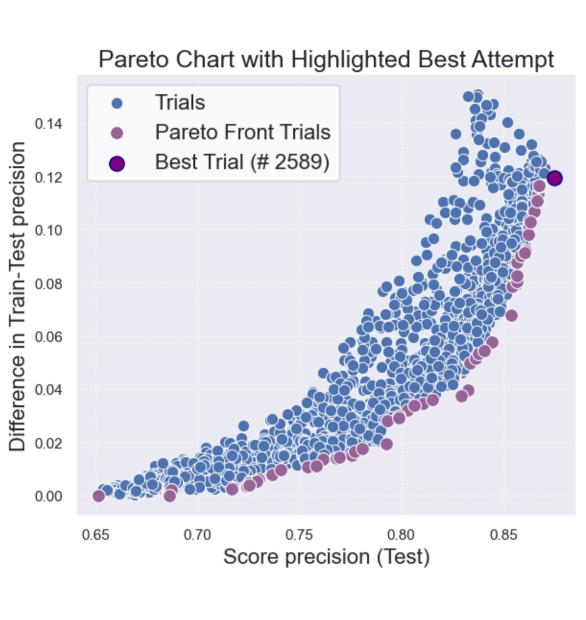
test: 0.0932

--- perfectly_calibrated

1.0

Protein: P27338 - metric: precision





XGBoost Parameters	Value
alpha	4.5009
colsample_bytree	0.7063
gamma	0.1131
lambda	0.0241
learning_rate	0.0639
max_depth	8.0
min_child_weight	1.0
n_estimators	675.0
reg_alpha	0.1111
reg_lambda	0.6137
scale_pos_weight	1.4019
subsample	0.944

		Ac	tive/Inactiv	e probabi	lity			
35	lna	ctive zone			Active :	zone	type active (34 inactive (
30							decoy (52	
25								
Percent 								
15								
10								
5								
U	0.0	.2 0	0.4 Proba	0. ability	6 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

0.23

0.93

0.77

0.074

0-

True label

Metric	Train	Test
AUC	0.9999	0.9502
accuracy	0.9943	0.8716
balanced accuracy	0.9921	0.8479
recall_(sens)	0.9993	0.9265
specificity	0.9849	0.7692
precision	0.992	0.8824
f1_score	0.9942	0.8701

--- train: 0.0088

0.8

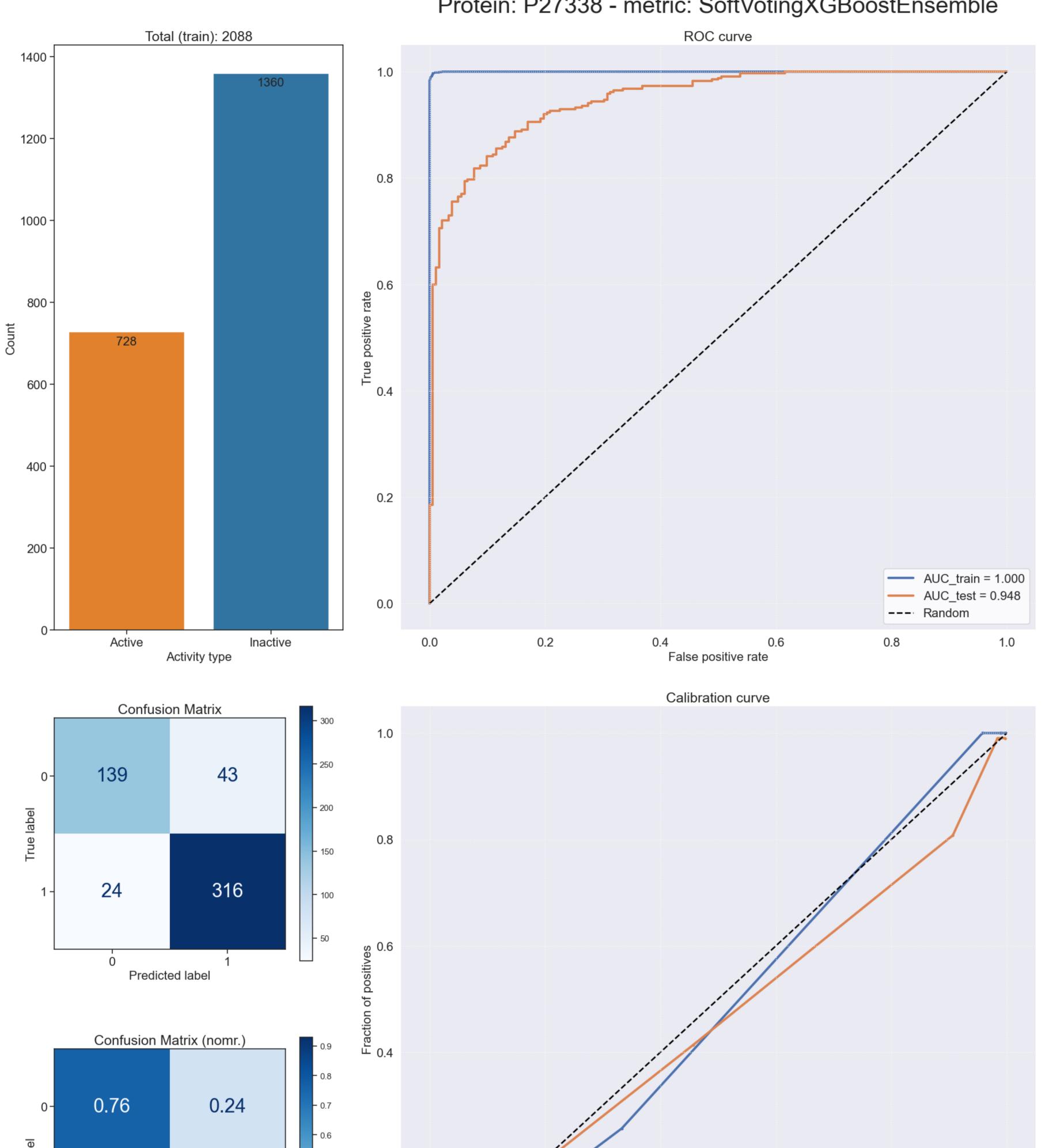
0.4 0.6 Mean predicted probability

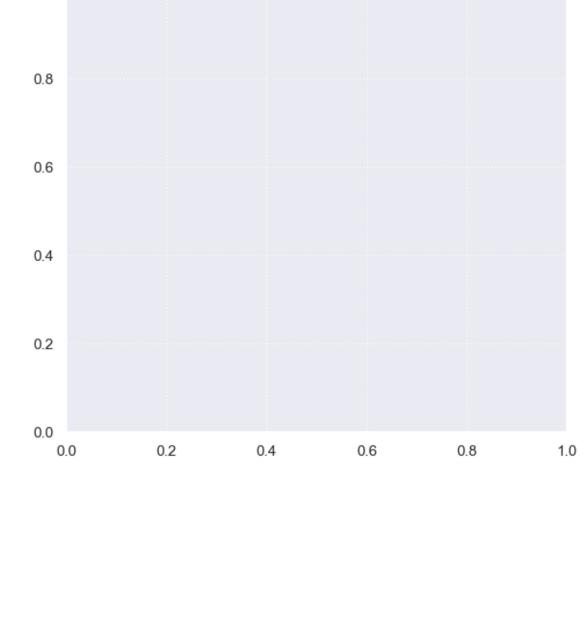
test: 0.0902

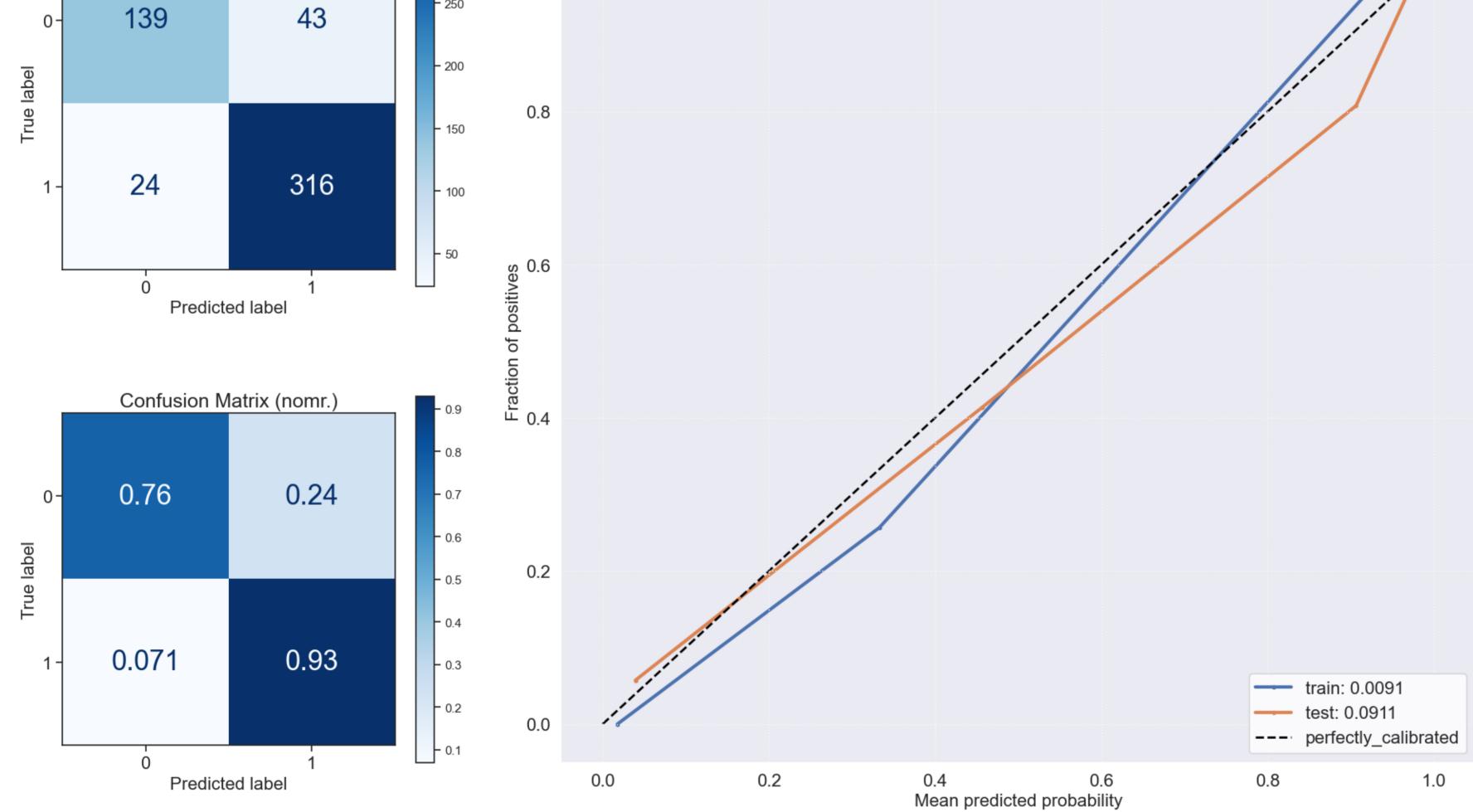
--- perfectly_calibrated

1.0

Protein: P27338 - metric: SoftVotingXGBoostEnsemble







		Acti	ve/Inactive probat	oility		
35	lna	ctive zone		Active	zone type active inactive decoy	e (182)
30						
20						
- 20						
10						
5						
0		2 0.4		0.6	8 1	0
	0.0	0.	Probability	0.0		

Metric	Train	Test
AUC	0.9999	0.9475
accuracy	0.9933	0.8716
balanced accuracy	0.991	0.8466
recall_(sens)	0.9985	0.9294
specificity	0.9835	0.7637
precision	0.9912	0.8802
f1_score	0.9933	0.8699