

# AlkahestResultV4a

July 4, 2023

```
[ ]: import numpy as np
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
from sklearn.impute import SimpleImputer
from sklearn.pipeline import Pipeline
from sklearn.preprocessing import StandardScaler
from sklearn.feature_selection import VarianceThreshold
from stabl.preprocessing import LowInfoFilter
from stabl.stabl import Stabl, save_stabl_results, plot_stabl_path, \
    plot_fdr_graph
from stabl.visualization import boxplot_features
from sklearn.base import clone
from sklearn.linear_model import LogisticRegression, LogisticRegressionCV, \
    LinearRegression
from sklearn.model_selection import LeaveOneOut, RepeatedStratifiedKFold
from sklearn.metrics import roc_auc_score
```

This here provides the code that gets us the result of the best model on the V4a (normalized to V3) only unstim data.

The best model found was a Lasso model, with hyperparameters  $\lambda_1 = 0.25$ . Reminder, the Lasso solves:

$$\operatorname{argmin}_{\beta} \frac{1}{2n} \|y - X\beta\|_2^2 + \lambda_1 \|\beta\|_1$$

We also use the usual preprocessing pipeline, with a variance threshold of 0.001 (i.e. features with lower variance are removed before fitting the model).

## 0.0.1 Step 1: Setup

We import the data and set up the preprocessing pipeline:

```
[ ]: dataV4a = pd.read_csv("./Data/Alkahest-V4a.csv", index_col=0)
dataV4a = dataV4a.iloc[:, [x[-6:] == "Unstim" for x in dataV4a.columns]]
label = pd.read_csv("./Data/AlkahestLabel.csv", index_col=0).iloc[:, 0]

preprocessing = Pipeline(
    steps=[
        ("variance", VarianceThreshold(0.001)),
```

```

        ("l1", LowInfoFilter(0)),
        ("impute", SimpleImputer(strategy="median")),
        ("std", StandardScaler())
    ]
)

X = dataV4a.to_numpy()
y = label.to_numpy()

```

## 0.0.2 Step 2: Cross-Validation

Now we calculate the ROC curve for the cross-validation predictions, using Leave-One-Out cross-validation. We obtain an AUC of 0.62 for the curve.

```

[ ]: from sklearn.metrics import roc_curve, RocCurveDisplay, auc

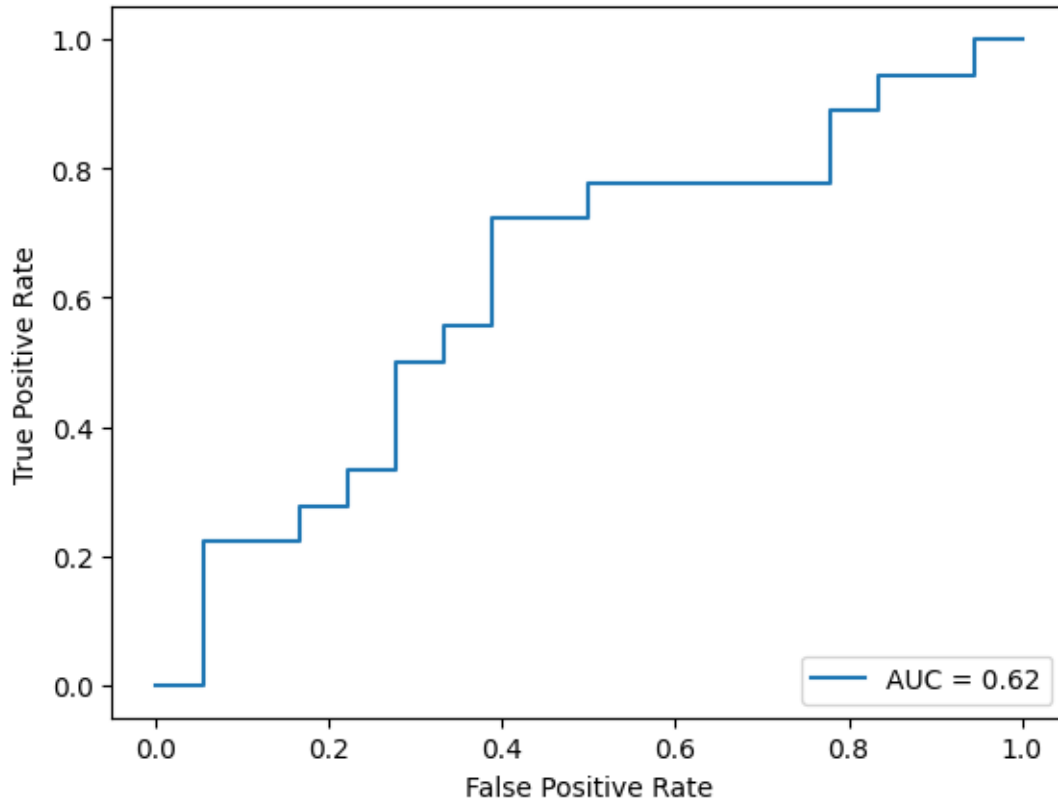
c = 0.25
model = LogisticRegression(penalty="l1", C=c, max_iter=int(1e8), solver="saga",
    ↪class_weight="balanced")
kf = LeaveOneOut()
all_probs=[]
for train, test in kf.split(X, y):
    Xtrain = pd.DataFrame(data=preprocessing.fit_transform(dataV4a.iloc[train,:
    ↪]),
                        index=dataV4a.index[train],
                        columns=preprocessing.get_feature_names_out()
                        ).to_numpy()
    Xtest = pd.DataFrame(data=preprocessing.transform(dataV4a.iloc[test,:]),
                        index=dataV4a.index[test],
                        columns=preprocessing.get_feature_names_out()
                        ).to_numpy()
    all_probs.append(model.fit(Xtrain, y[train]).predict_proba(Xtest)[:,-1])
all_probs = np.array(all_probs).flatten()
fpr, tpr, thresholds = roc_curve(y, all_probs)
roc_auc = auc(fpr, tpr)
disp = RocCurveDisplay(fpr=fpr, tpr=tpr, roc_auc=roc_auc)
disp.plot()

```

```

[ ]: <sklearn.metrics._plot.roc_curve.RocCurveDisplay at 0x1614ed81410>

```



We then calculate the p-value of the same predictions, which is about 21.4%.

```
[ ]: from scipy.stats import mannwhitneyu

utest1,upval1 = mannwhitneyu(all_probs[y == 1],all_probs[y == 0],method="exact")
print( "p-value of the U-Test on the CV predictions : ",upval1)
```

p-value of the U-Test on the CV predictions : 0.21424980209385966

The spread of the predictions is in the next figure - 0 is placebo, 1 is Alkahest. The dots are the prediction values. Ideally we want the top line to have all dots at 0, and bottom line all at 1.

```
[ ]: import matplotlib.patches as plt
fig, ax = plt.subplots(1, 1, figsize=(10, 8))
palette = ["#4D4F53", "#C41E3A"]
sns.boxplot(data = [all_probs[y == 0],all_probs[y == 1]],
            ax=ax,
            showfliers=False,
            palette="pastel",
            boxprops=dict(alpha=.2),
            whiskerprops=dict(alpha=.2),
            width=.4,
```

```

        orient="h",
        saturation=1
    )
sns.stripplot(data=[all_probs[y == 0],all_probs[y == 1]],
              ax=ax,
              palette="pastel",
              jitter = False,
              alpha=1,
              size=4,
              marker="D",orient="h"
    )
box_patches = [patch for patch in ax.patches if type(patch) == plt.PathPatch]

num_patches = len(box_patches)
lines_per_boxplot = len(ax.lines) // num_patches

for i, patch in enumerate(box_patches):
    col = patch.get_facecolor()
    patch.set_edgecolor(col)
    patch.set_facecolor(col)

    for line in ax.lines[i * lines_per_boxplot: (i + 1) * lines_per_boxplot]:
        line.set_color(col)
        line.set_mfc(col)
        line.set_mec(col)

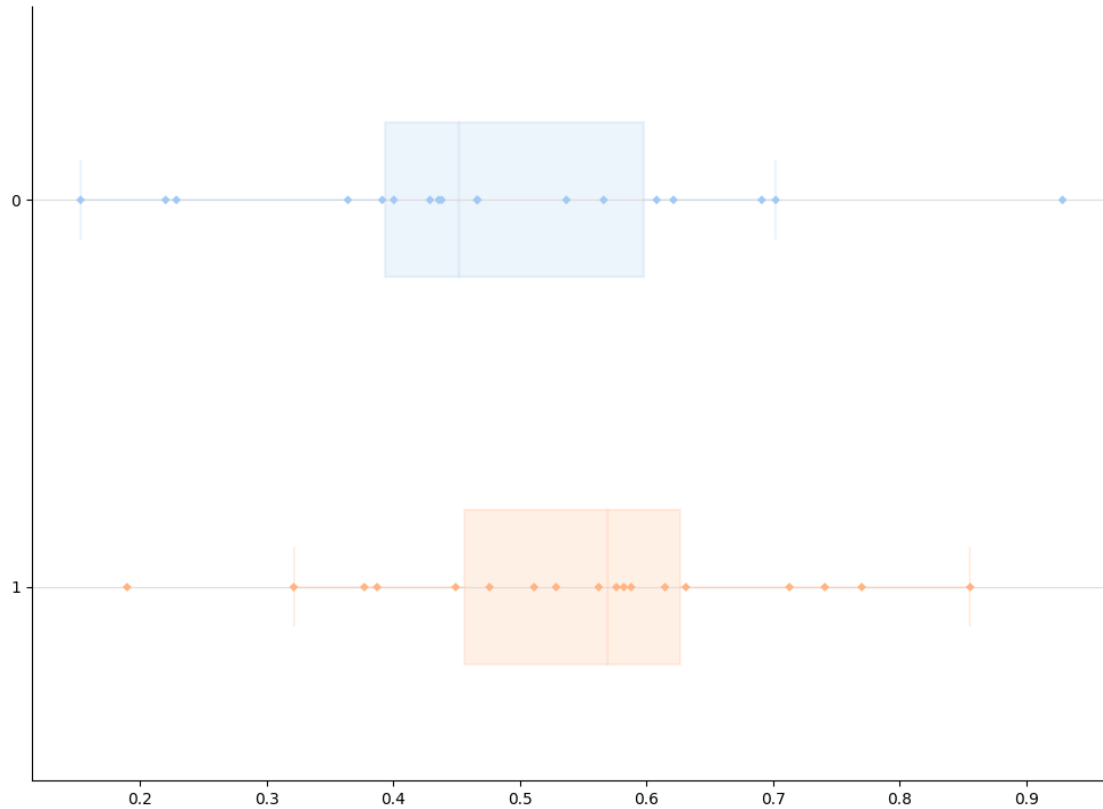
ax.grid(which='major', color='#DDDDDD', linewidth=0.8, axis="y")

ax.spines['top'].set_visible(False)
ax.spines['right'].set_visible(False)

fig.tight_layout()
fig.subplots_adjust(top=0.9)
ax.set_ylabel('')

plt.show()

```



### 0.0.3 Step 3: We get the final model

We fit the model on all of the data.

```
[ ]: Xstd = pd.DataFrame(data=preprocessing.fit_transform(dataV4a),
                        index=dataV4a.index,
                        columns=preprocessing.get_feature_names_out()
                        )
model.fit(Xstd.to_numpy(), y)
```

```
[ ]: LogisticRegression(C=0.25, class_weight='balanced', max_iter=100000000,
                        penalty='l1', solver='saga')
```

Then we look at which features get a non-zero coefficient, to get a similar understanding of which features were selected by the model.

```
[ ]: for i in Xstd.columns[np.where(model.coef_[0] != 0)]:
      print(i)
```

```
CD8Tem_pSTAT6_Unstim
Tregnaive_MAPKAPK2_Unstim
NK-Tcells_pSTAT6_Unstim
gdT_pSTAT6_Unstim
```

```

CD8Tcm_Frequency_Unstim
Th2_STAT5_Unstim
Tregmem_ERK_Unstim
Baso +++_STAT5_Unstim
Th1naive_MAPKAPK2_Unstim

```

```

[ ]: pd.DataFrame(data=all_probs,index=dataV4a.index,columns=["L00 CV predictions"]).
      to_csv("./ResultsFinal/V4a/FinalLASSO_CV_predictions.csv")

```

Boxplots for these features are further down.

#### 0.0.4 Part 2: Stability Selection

With a base estimator a Lasso model, we look at what features it selects:

```

[ ]: ss =Stabl(
      base_estimator=clone(model),
      lambda_name="C",
      lambda_grid=np.linspace(0.01, 3, 50),
      artificial_type=None,
      sample_fraction=0.5,
      replace= True,
      hard_threshold=0.3,
      n_bootstraps=2000,
      random_state=42
    )

ss.fit(Xstd,y)

```

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46.97s/it]

```

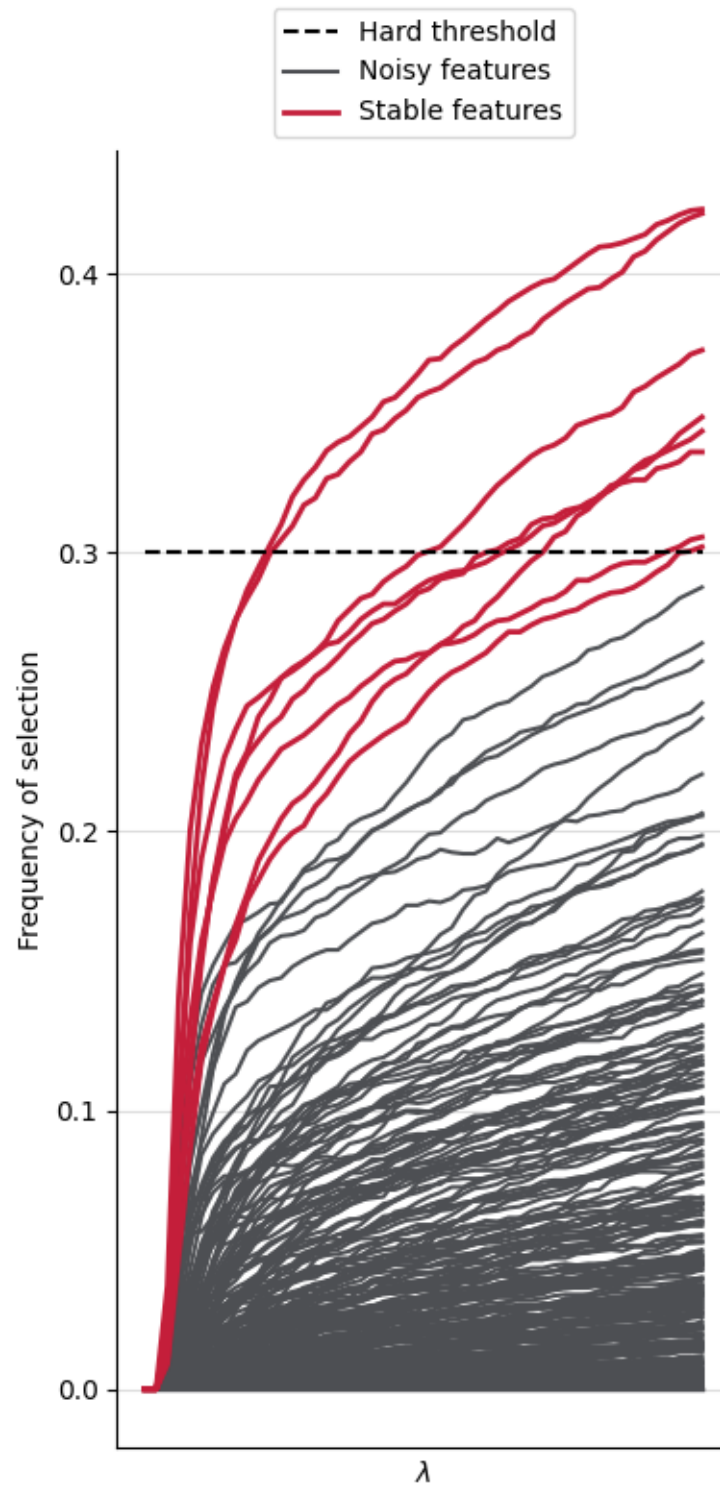
[ ]: Stabl(artificial_type=None,
          base_estimator=LogisticRegression(C=0.25, class_weight='balanced',
                                             max_iter=100000000, penalty='l1',
                                             solver='saga'),
          hard_threshold=0.7,
          lambda_grid=array([0.01          , 0.07102041, 0.13204082, 0.19306122,
0.25408163,
0.31510204, 0.37612245, 0.43714286, 0.49816327, 0.55918367,
0.62020408, 0.68122449, 0.7422449 , 0.80326531, 0.86428571,
0.92530612, 0.98...
1.23040816, 1.29142857, 1.35244898, 1.41346939, 1.4744898 ,
1.5355102 , 1.59653061, 1.65755102, 1.71857143, 1.77959184,
1.84061224, 1.90163265, 1.96265306, 2.02367347, 2.08469388,
2.14571429, 2.20673469, 2.2677551 , 2.32877551, 2.38979592,
2.45081633, 2.51183673, 2.57285714, 2.63387755, 2.69489796,
2.75591837, 2.81693878, 2.87795918, 2.93897959, 3.          ]),
          n_bootstraps=2000, random_state=42, replace=True)

```

The stability path generated:

```
[ ]: plot_stabl_path(ss)
```

```
[ ]: (<Figure size 400x800 with 1 Axes>,  
      <Axes: xlabel='$\\lambda$', ylabel='Frequency of selection'>)
```



The features selected:



```
[ ]: ss.get_feature_names_out()
```

```
[ ]: array(['CD8Tem_pSTAT6_Unstim', 'intMC_Frequency_Unstim',  
          'Tregnaive_MAPKAPK2_Unstim', 'NK-Tcells_pSTAT6_Unstim',  
          'gdT_pSTAT6_Unstim', 'CD8Tcm_Frequency_Unstim',  
          'Th1naive_STAT3_Unstim', 'Baso +++_STAT5_Unstim'], dtype=object)
```

```
[ ]: save_stabl_results(ss, "./ResultsFinal/V4a/", dataV4a, y)
```

```
c:\Users\Max\AppData\Local\Programs\Python\Python311\Lib\site-  
packages\stabl\visualization.py:260: FutureWarning: Passing `palette` without  
assigning `hue` is deprecated.
```

```
sns.stripplot(  

```

```
c:\Users\Max\AppData\Local\Programs\Python\Python311\Lib\site-  
packages\stabl\visualization.py:260: FutureWarning: Passing `palette` without  
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```

```
sns.stripplot(  

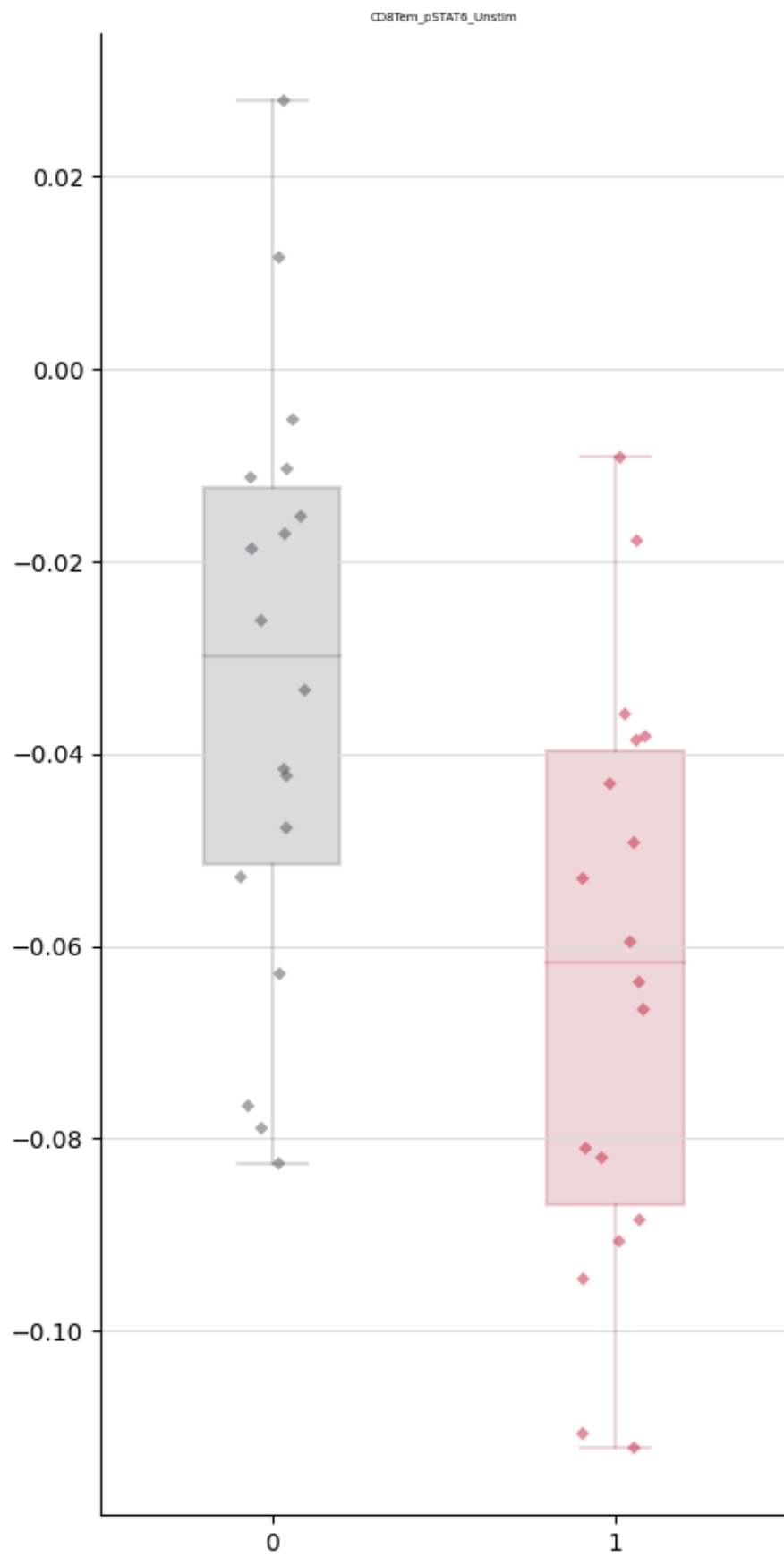
```

Here are the boxplots for these features.

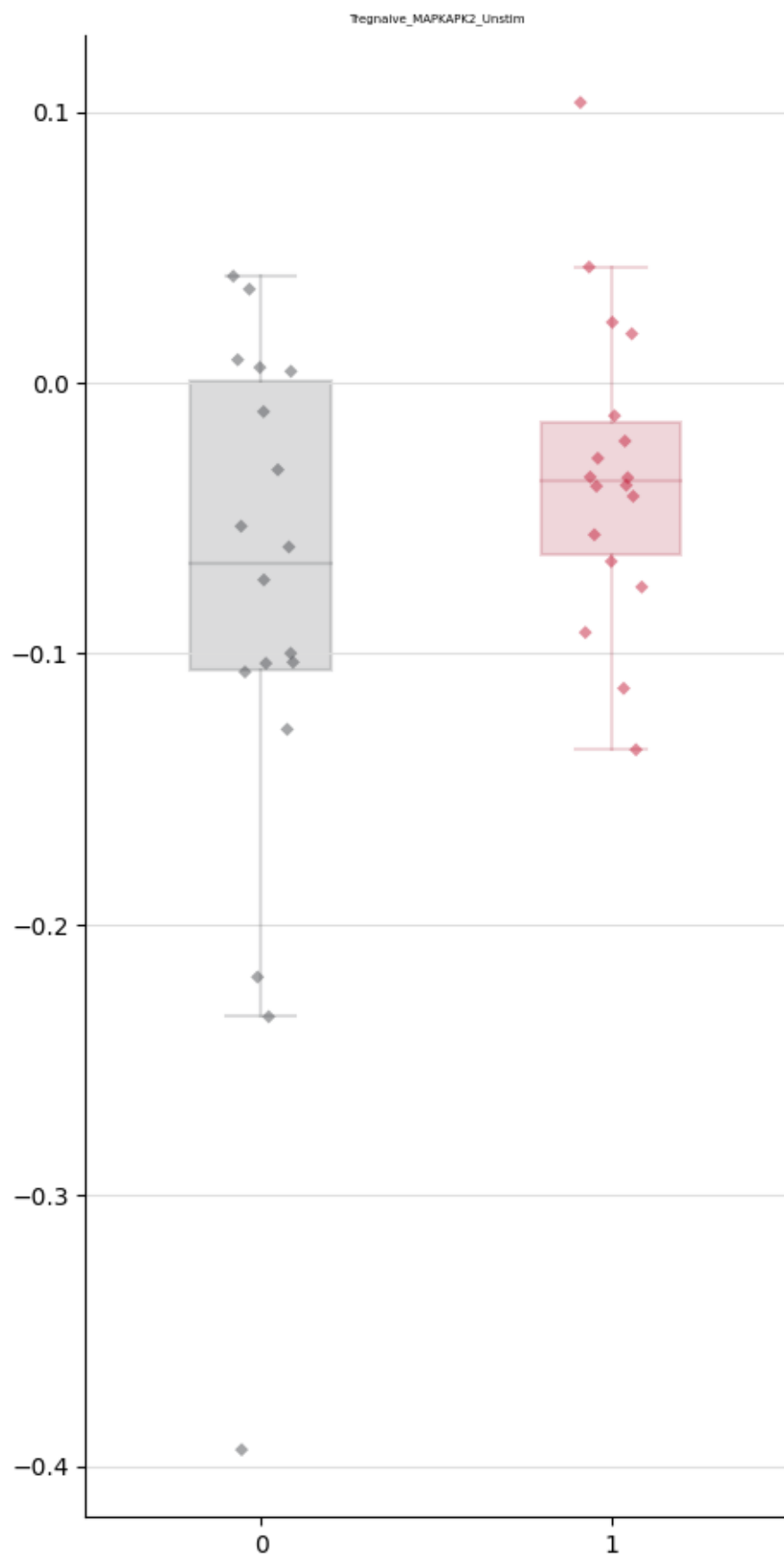
```
[ ]: boxplot_features(Xstd.columns[np.where(model.coef_[0] != 0)], dataV4a, y)
```

```
c:\Users\Max\AppData\Local\Programs\Python\Python311\Lib\site-  
packages\stabl\visualization.py:260: FutureWarning: Passing `palette` without  
assigning `hue` is deprecated.
```

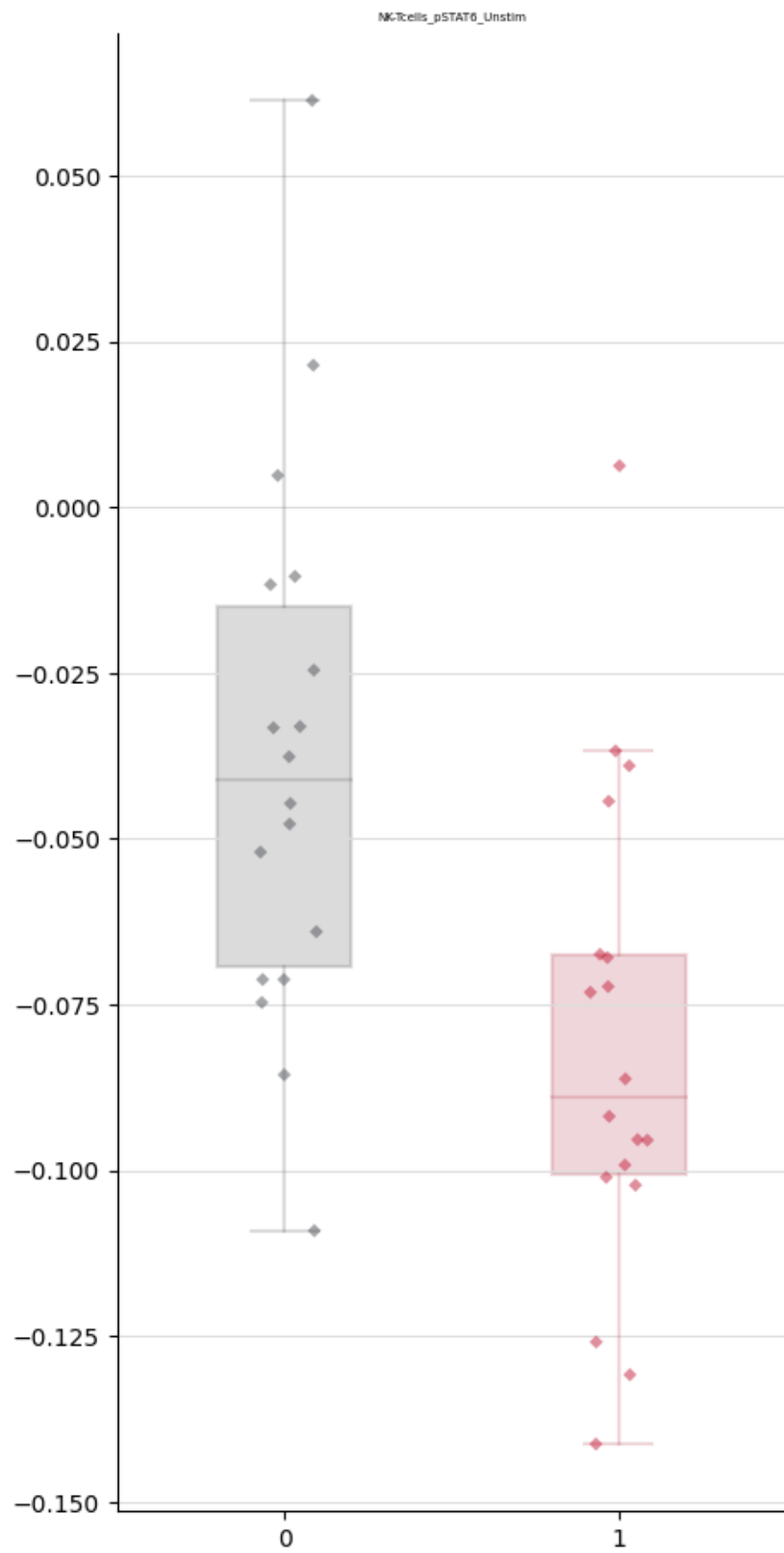






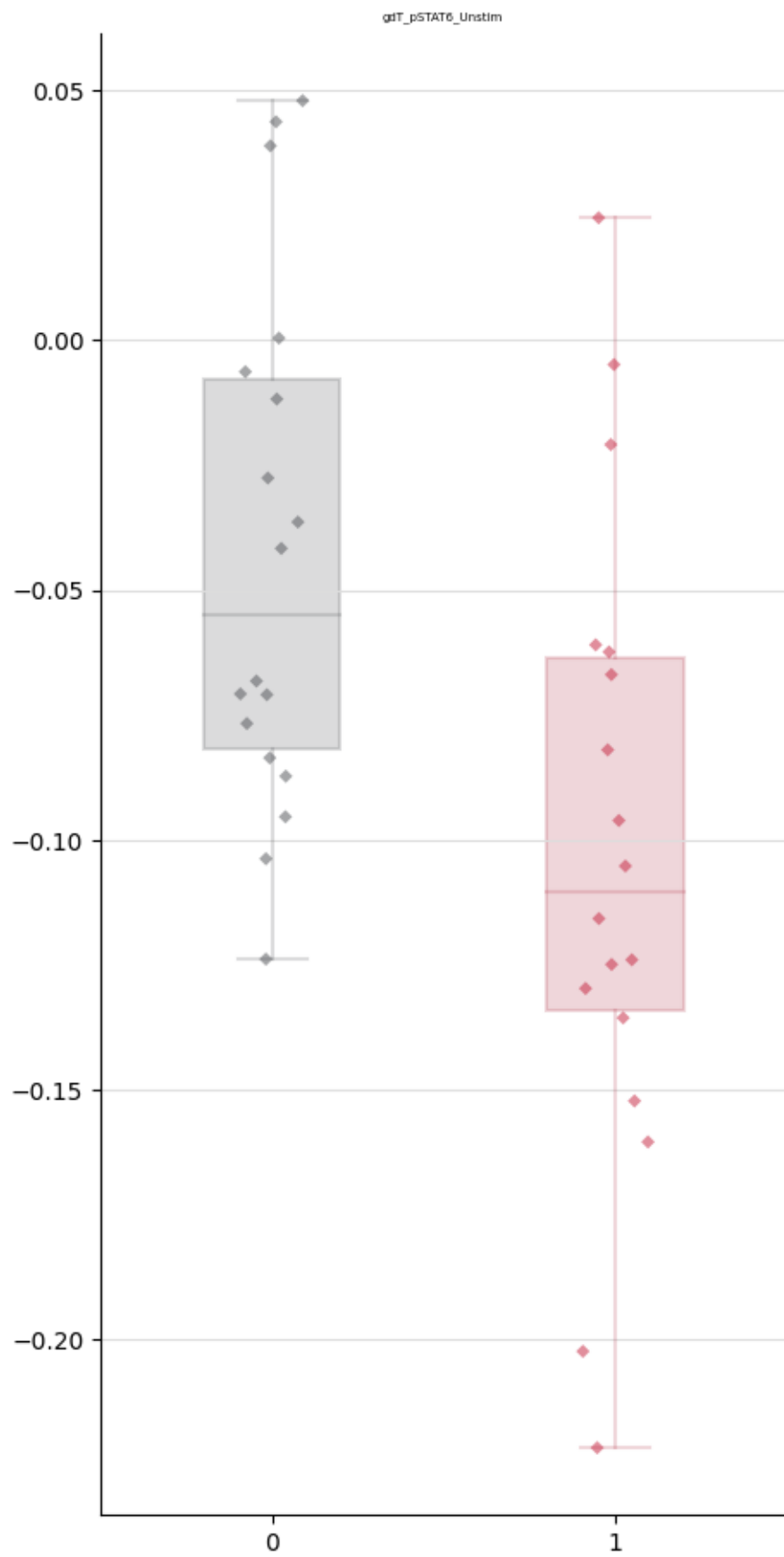




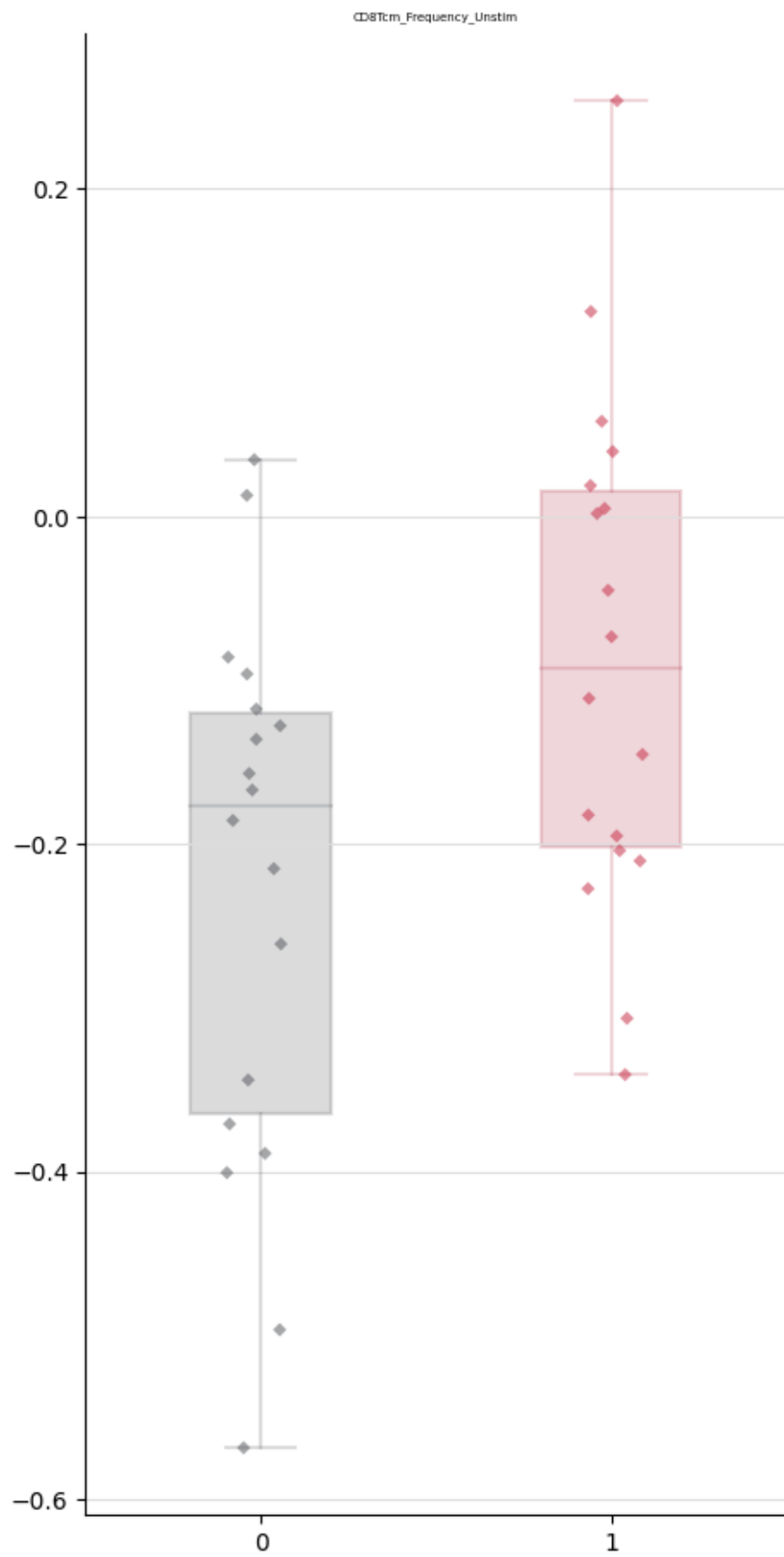




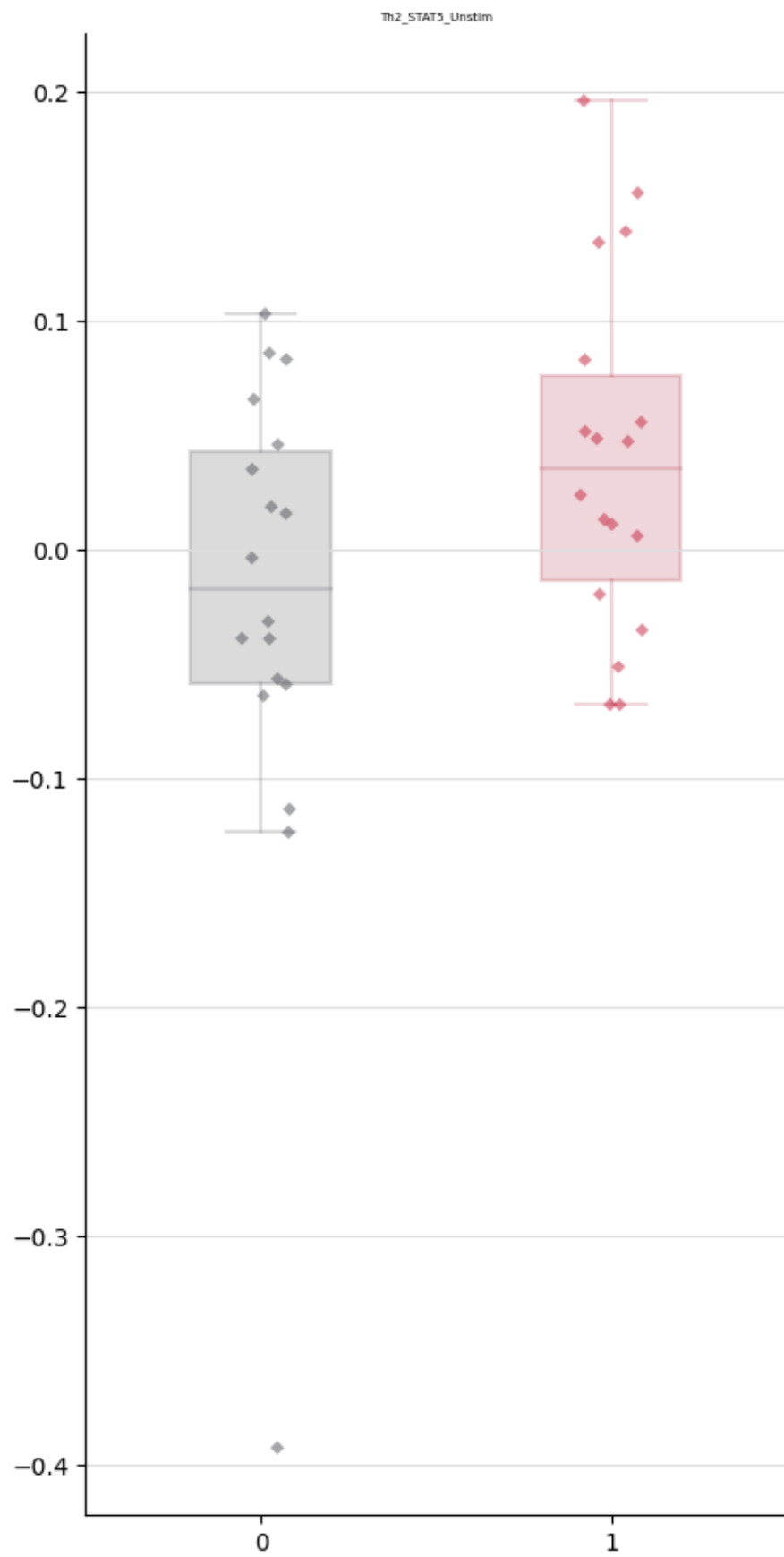




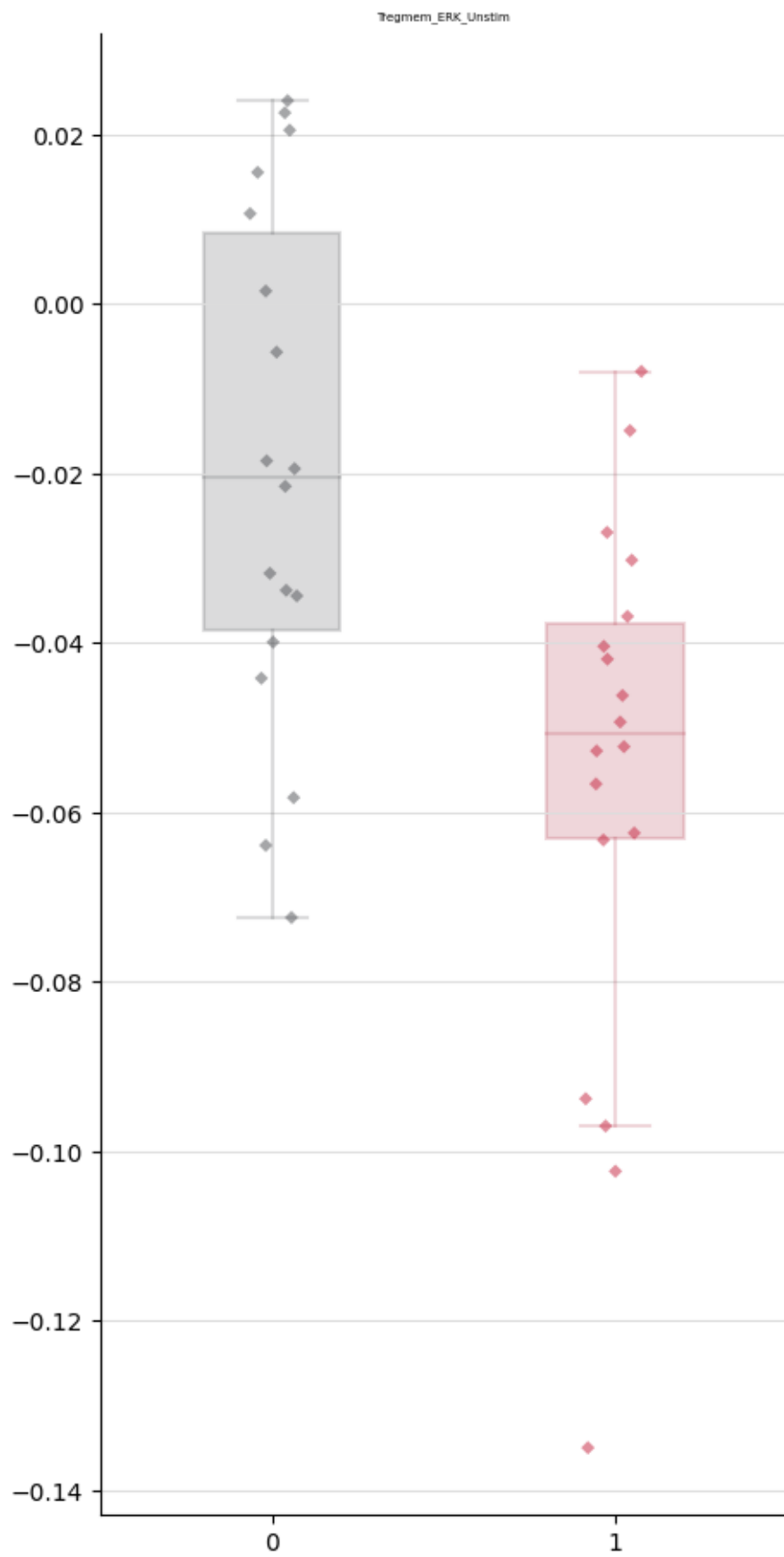






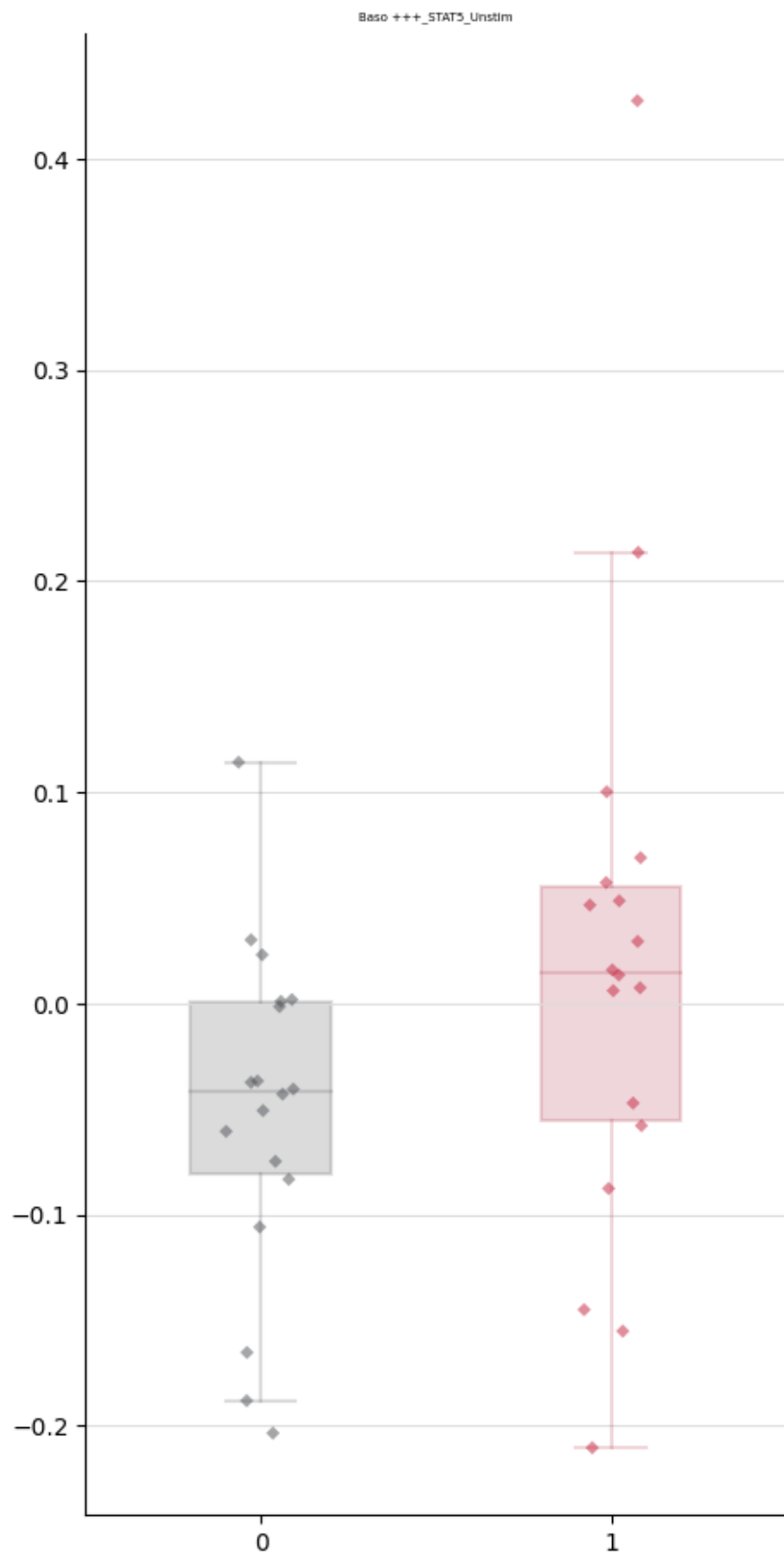




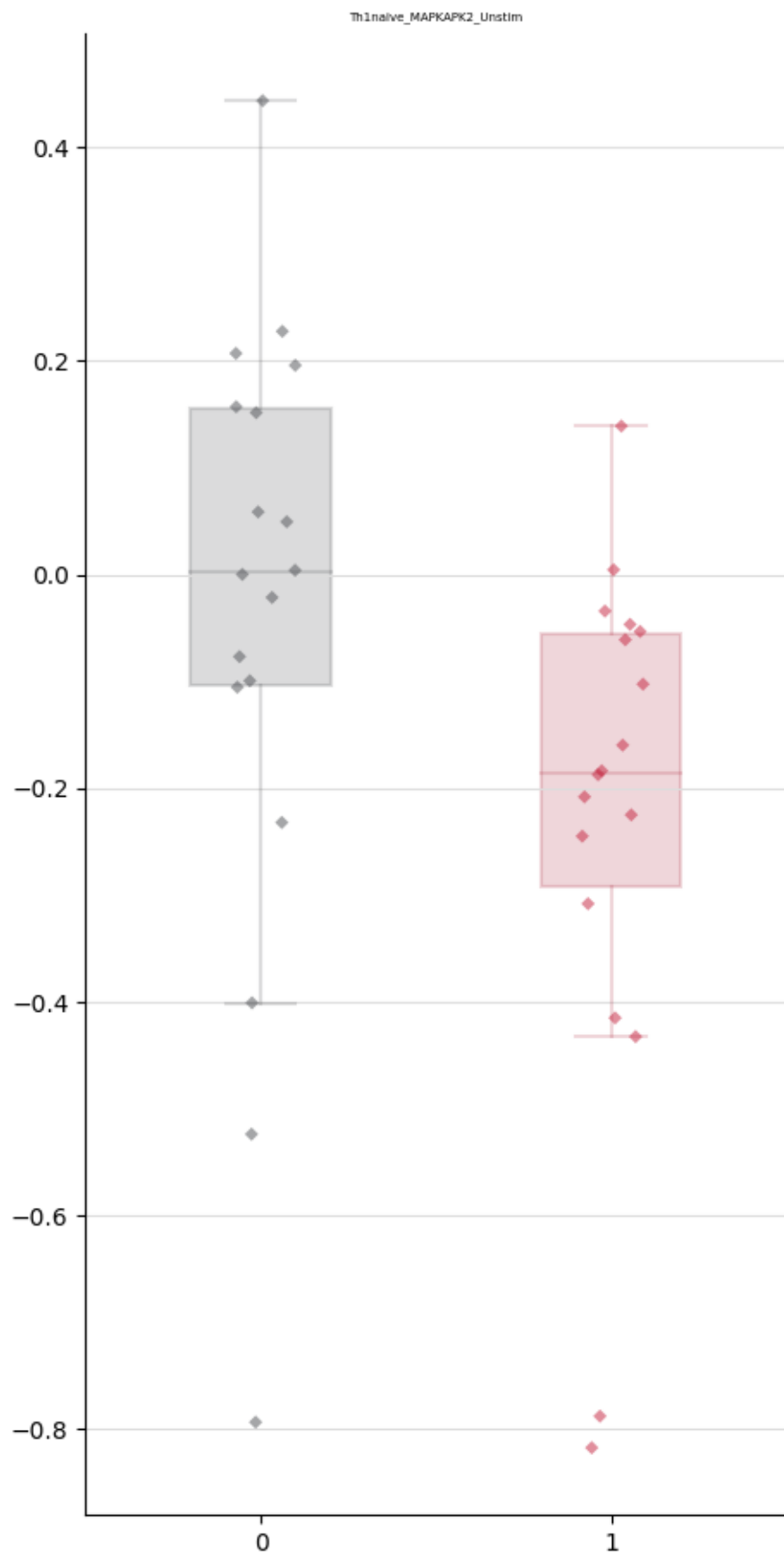












[ ]: