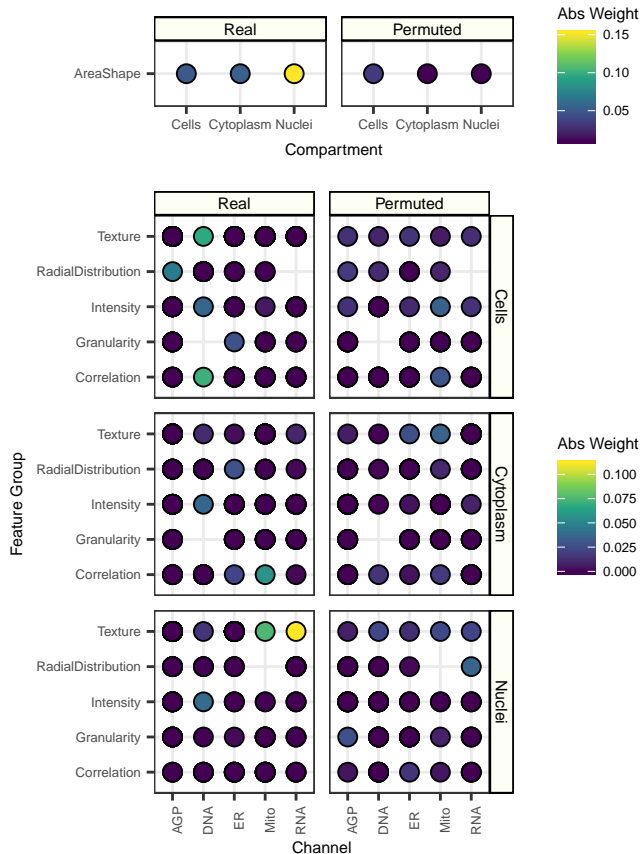
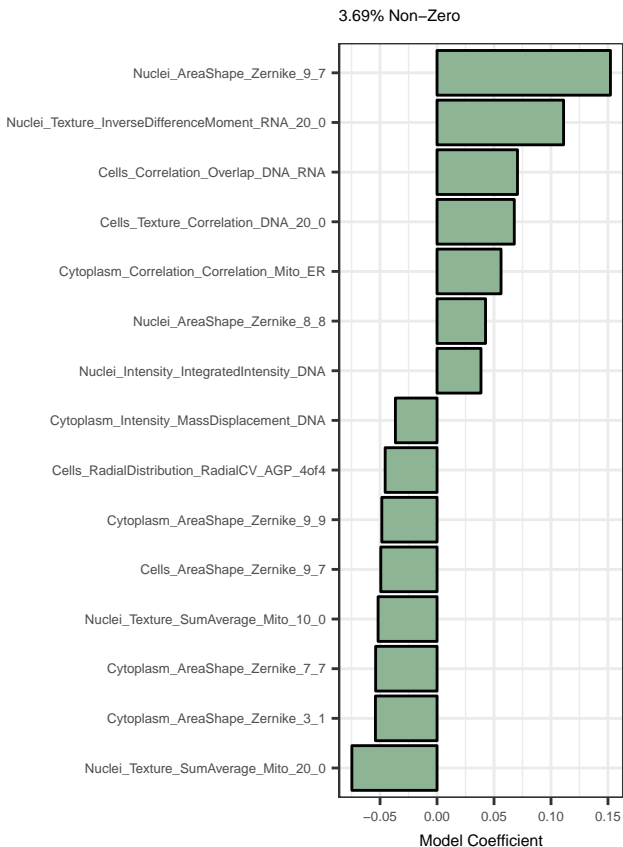
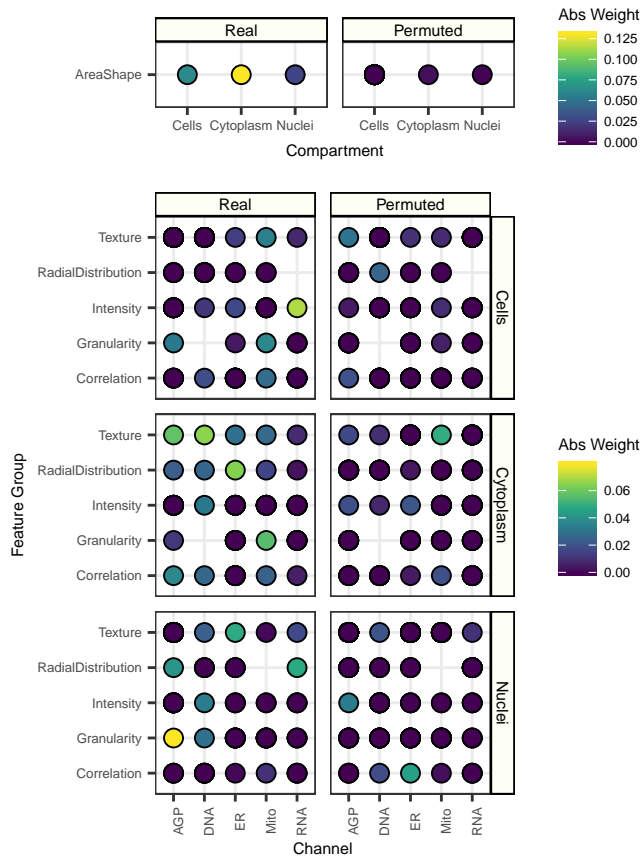
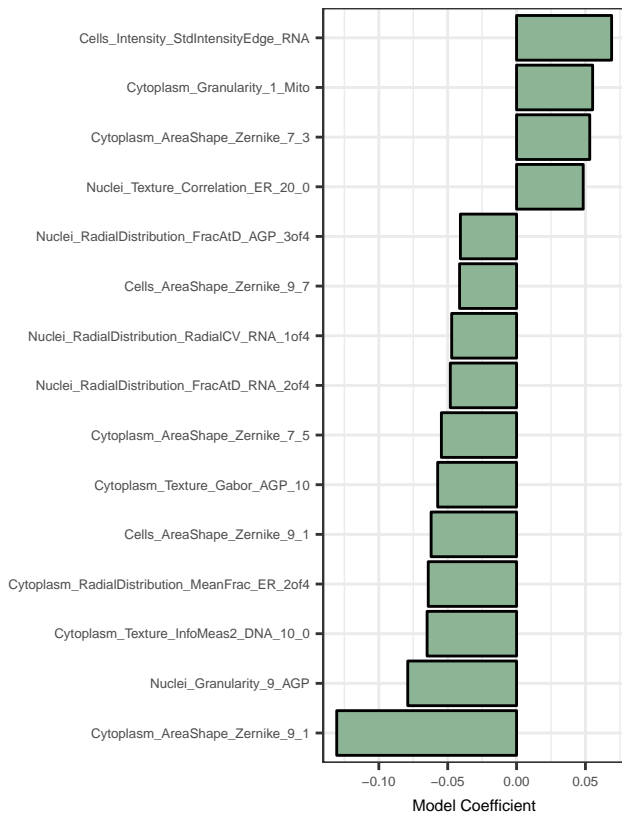


# ALL – % High gH2AX Spots

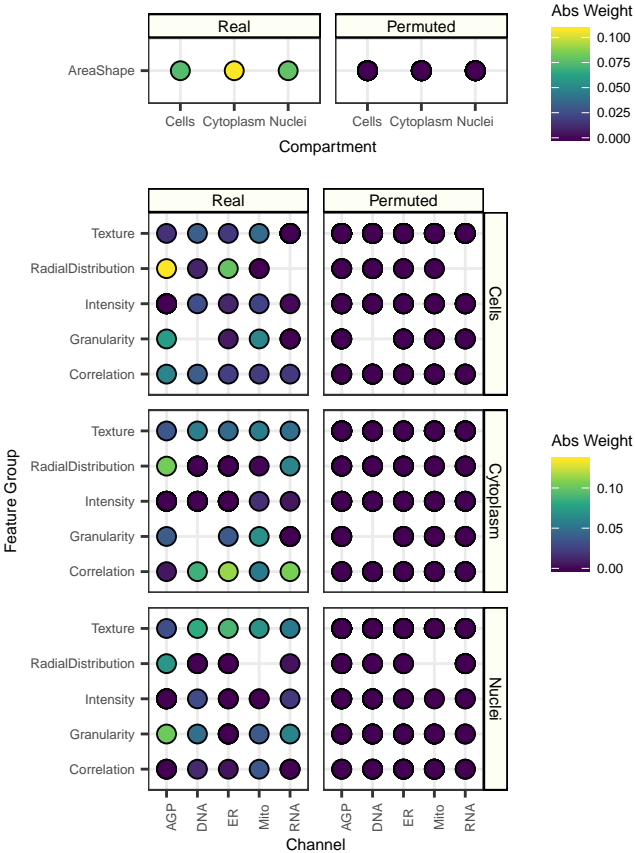
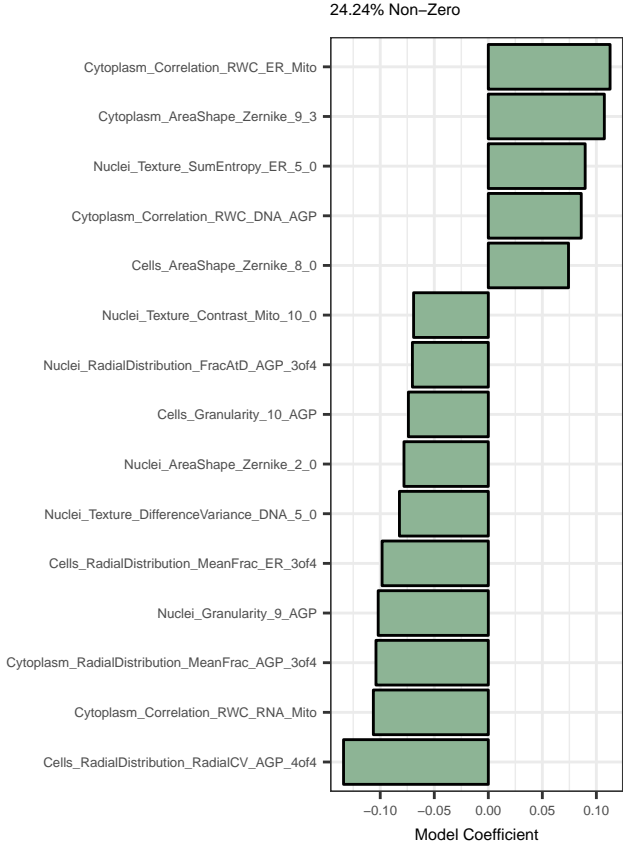


# ALL – Polynuclear

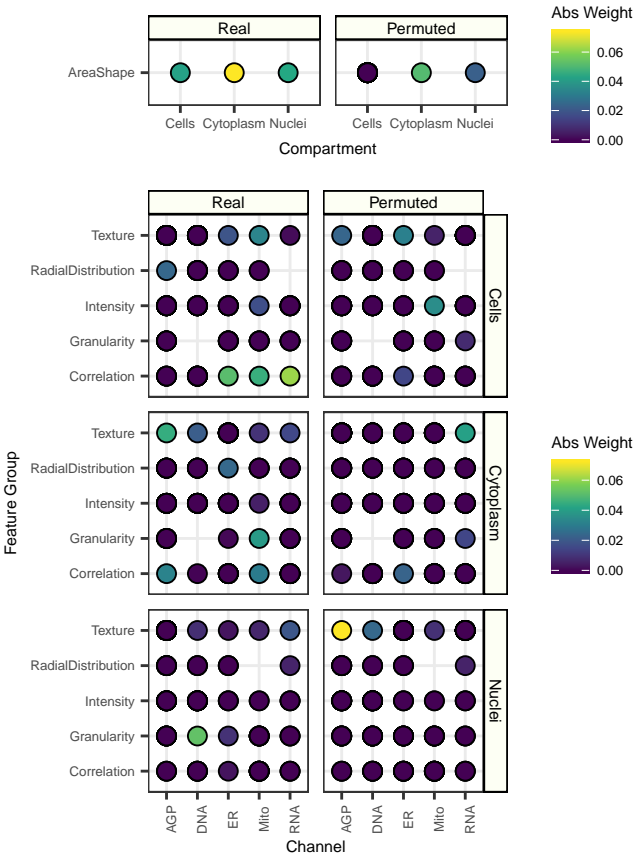
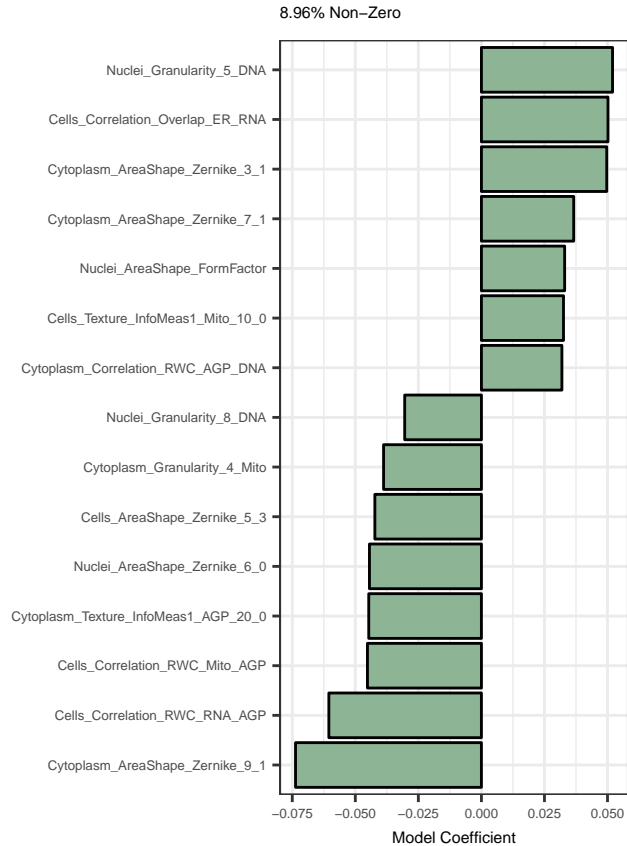
14.54% Non-Zero



# ALL – Polyploid

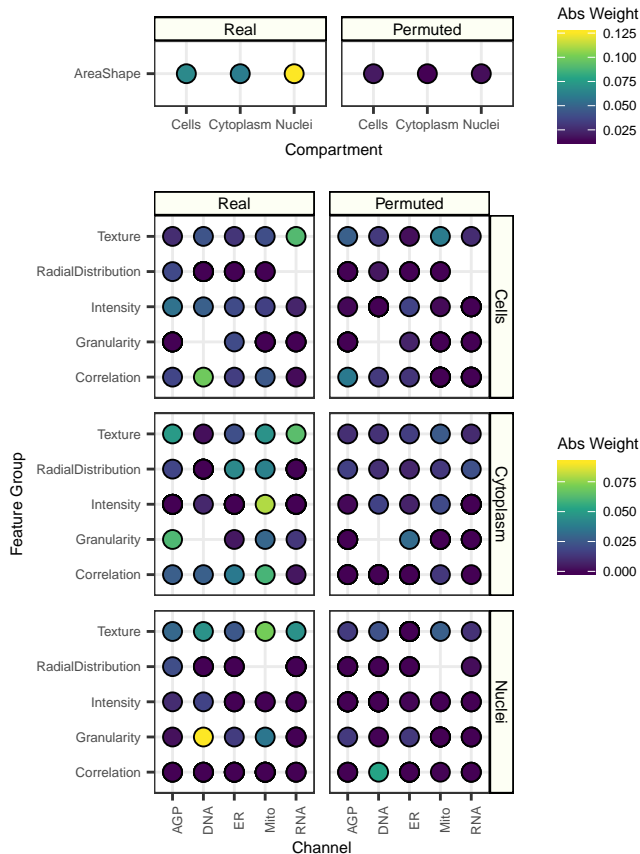
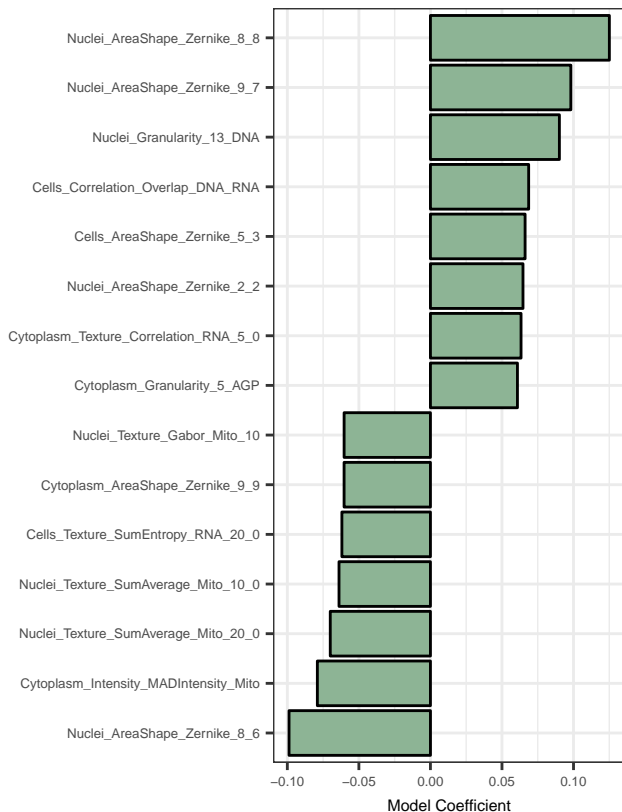


# ALL – # cells



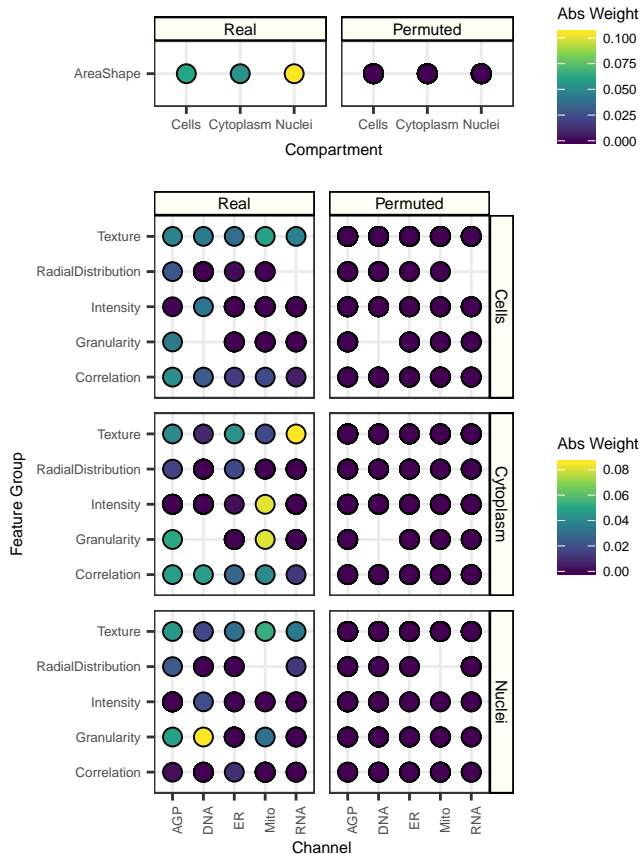
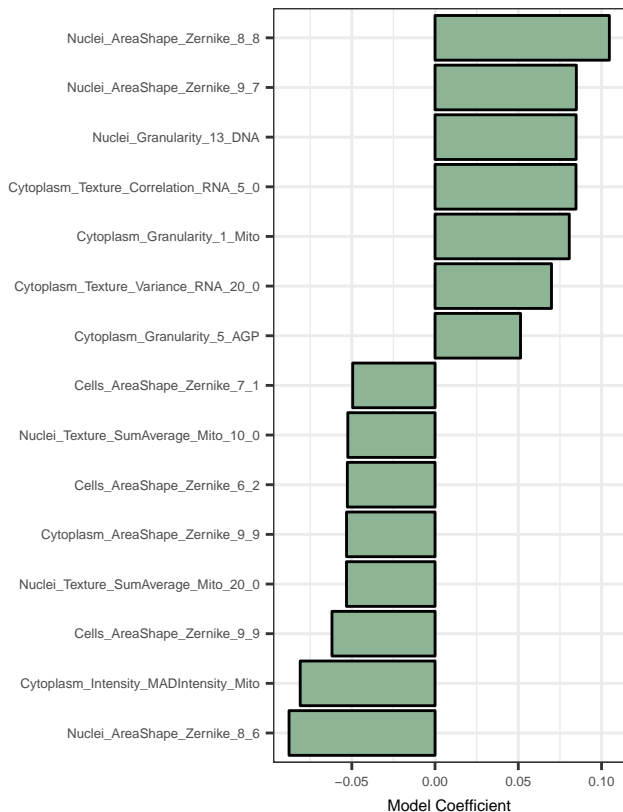
# ALL – # of gH2AX Spots

17.18% Non-Zero

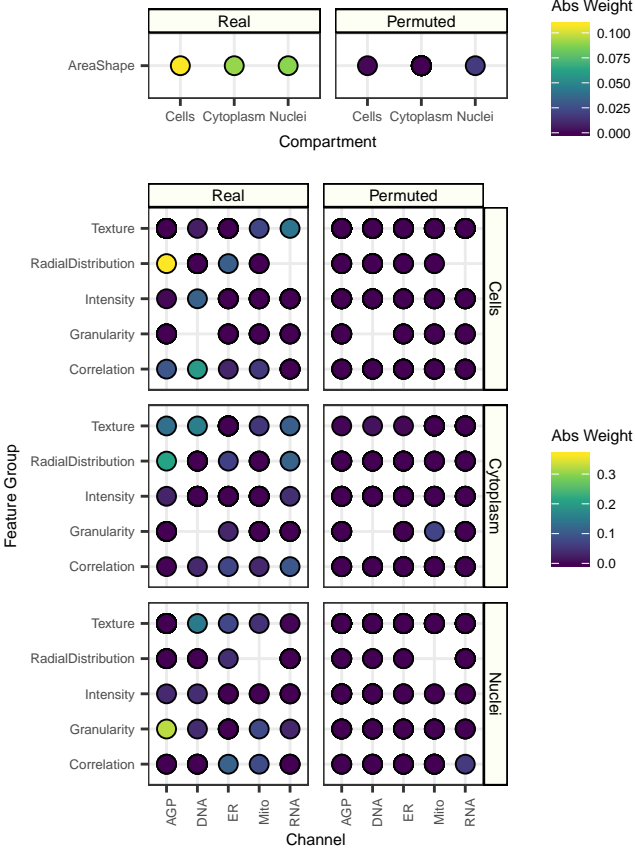
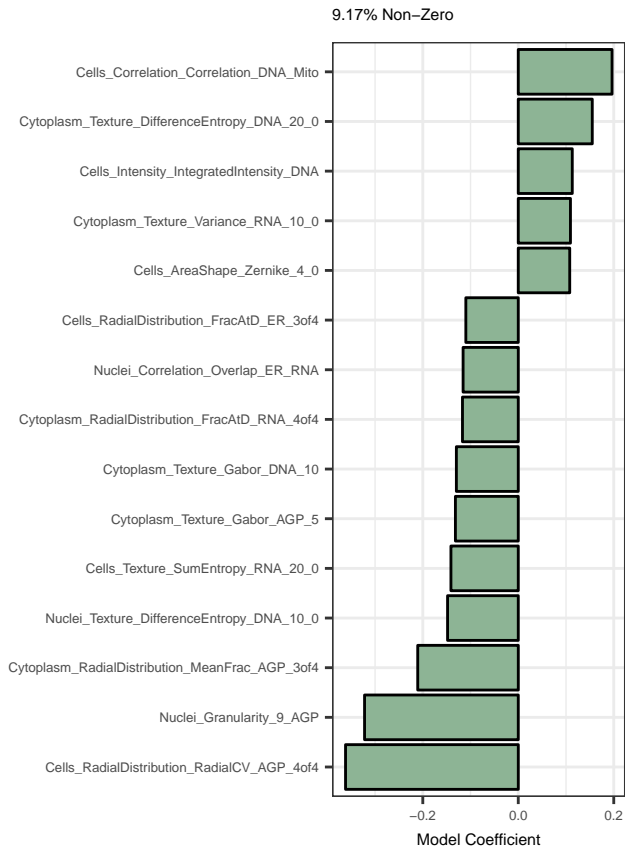


# ALL – # of gH2AX Spots per Area of Nucleus

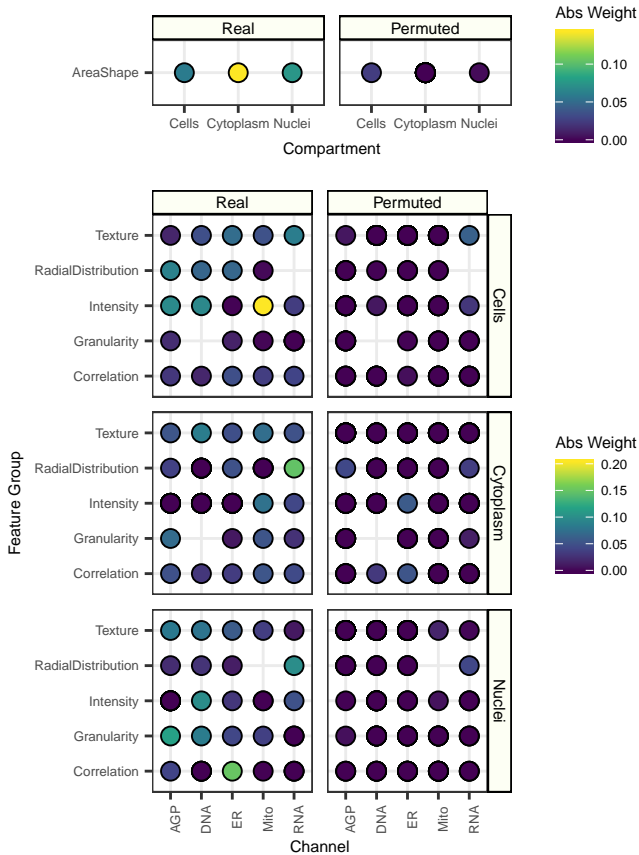
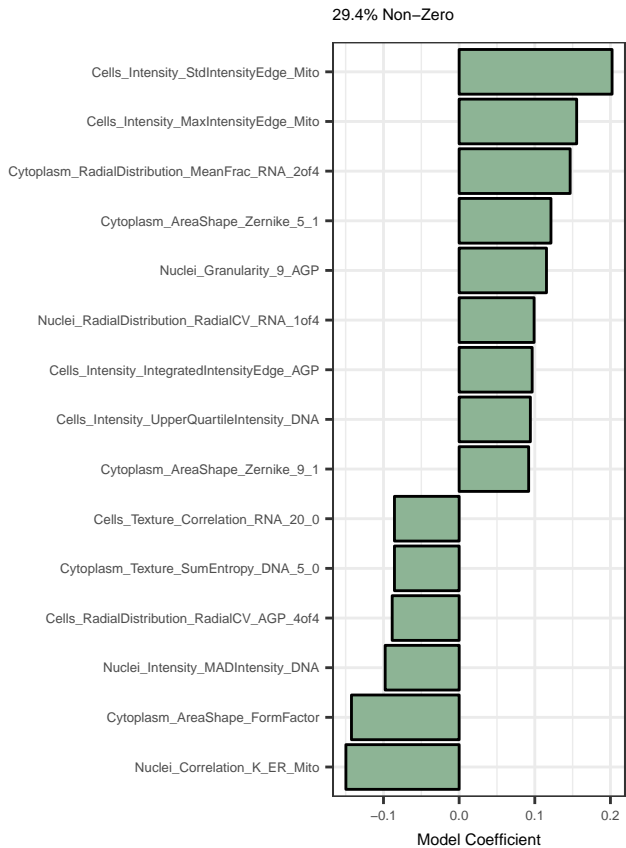
17.39% Non-Zero



# ALL – Nucleus Area

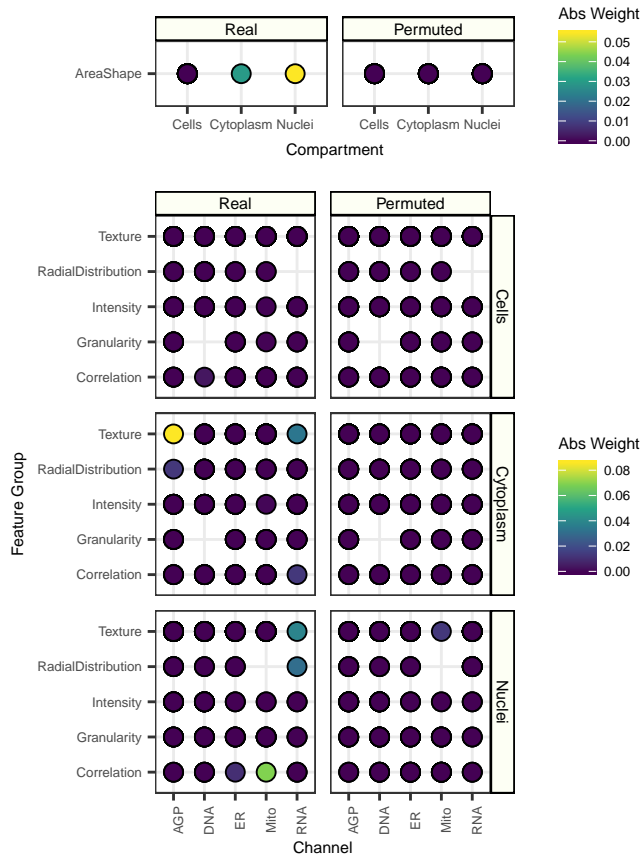
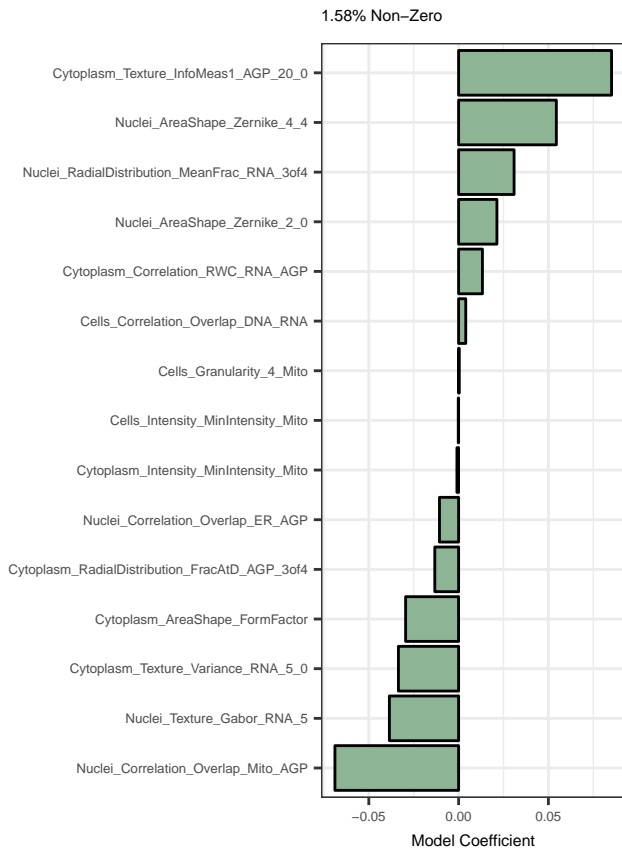


# ALL – Nucleus Roundness



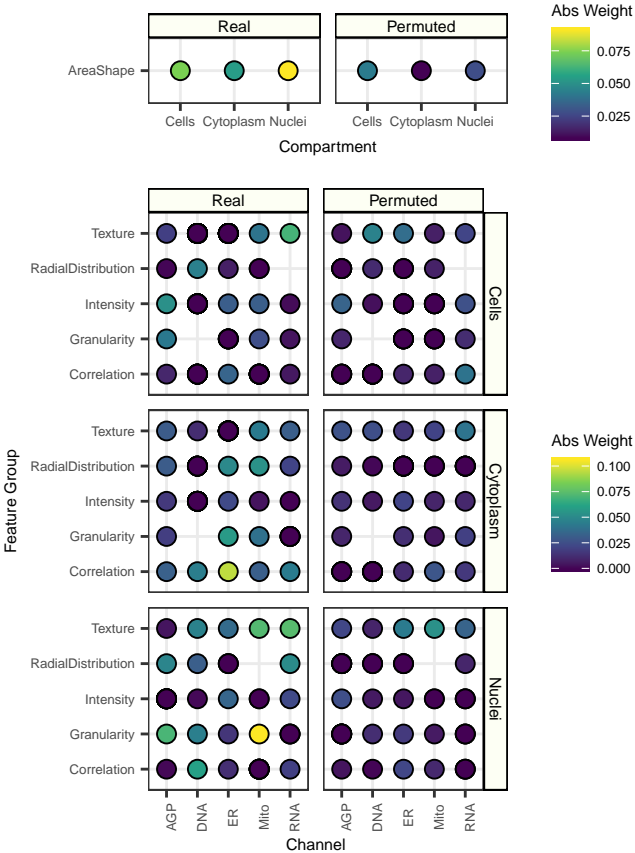
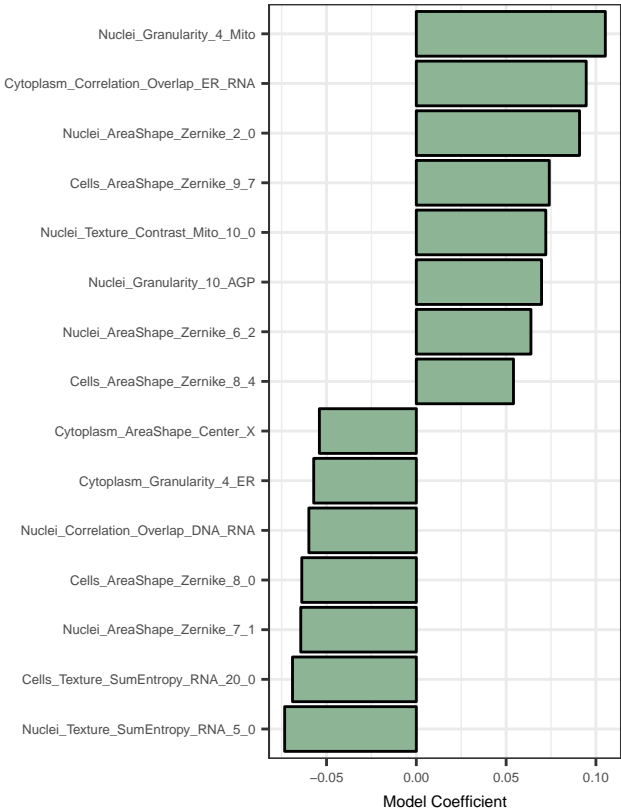


## CC – % early M

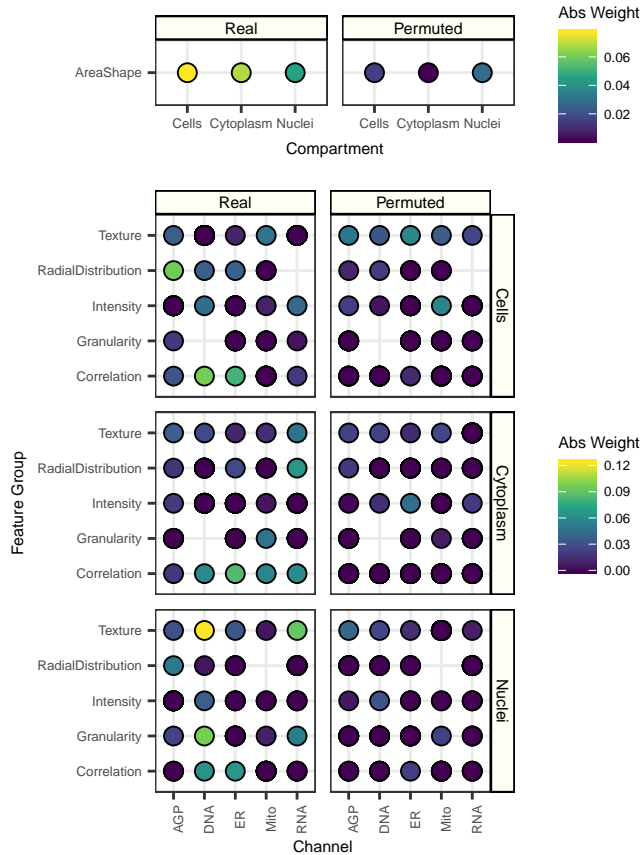
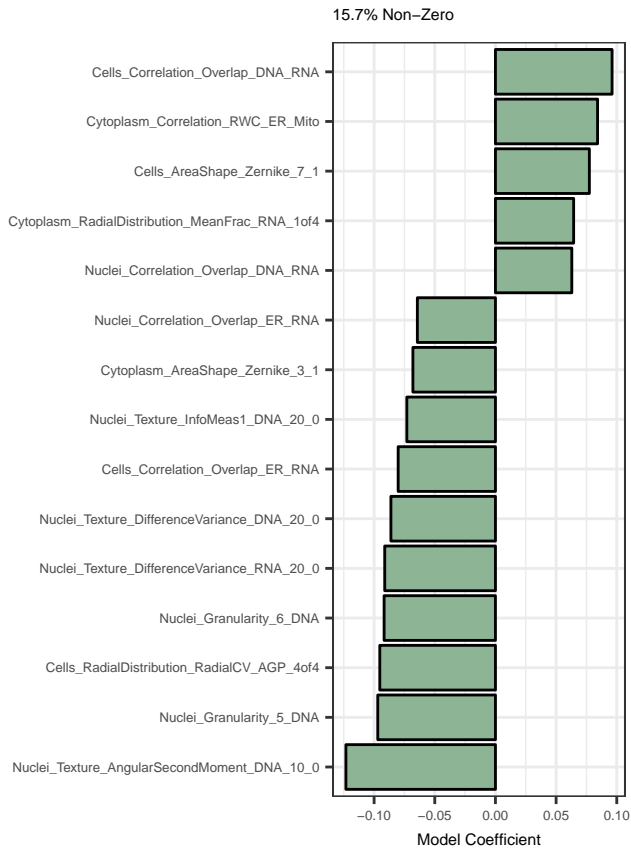


# CC - % G1

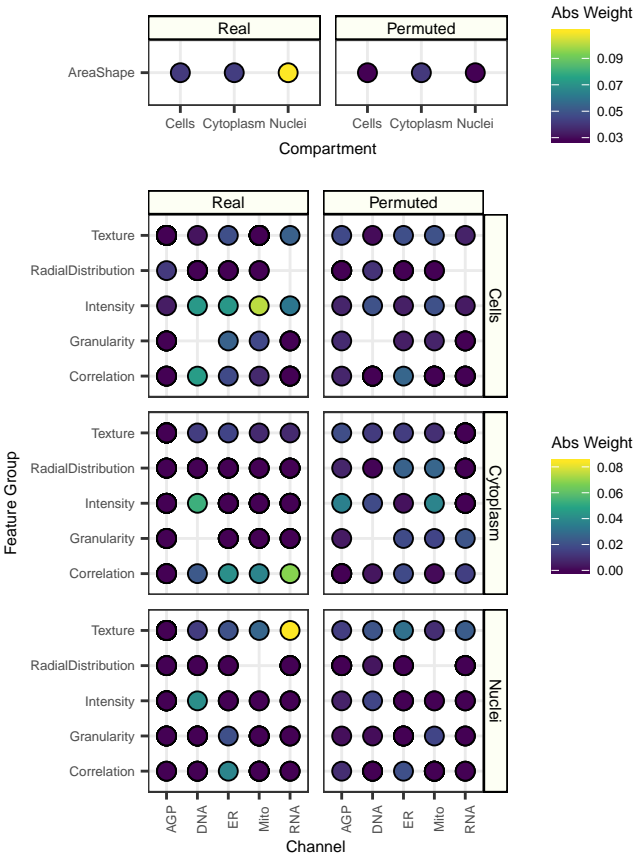
20.44% Non-Zero



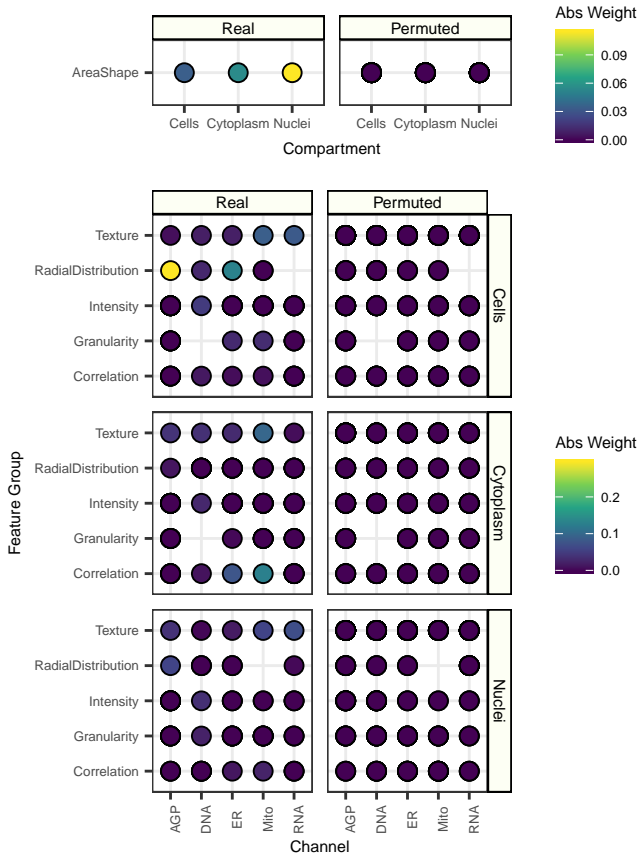
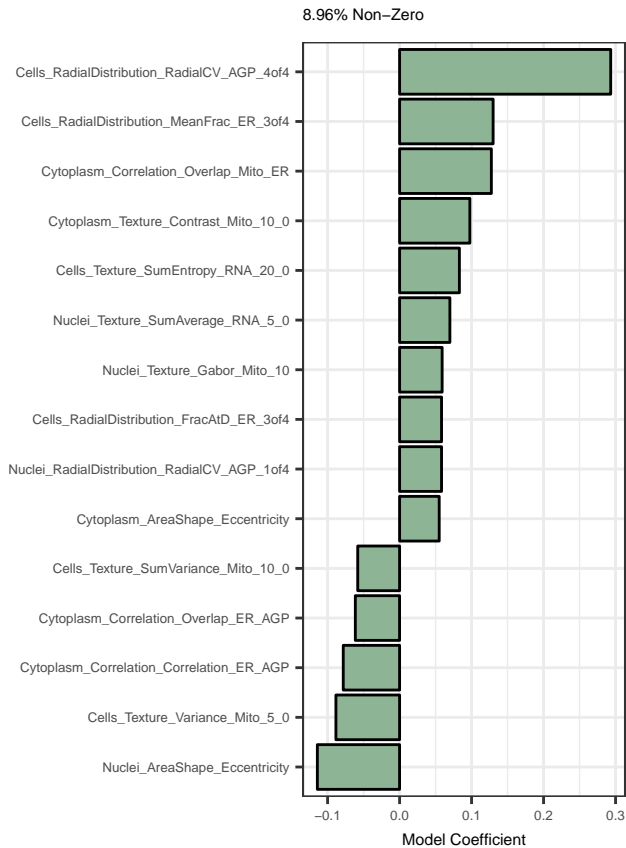
# CC - % G2



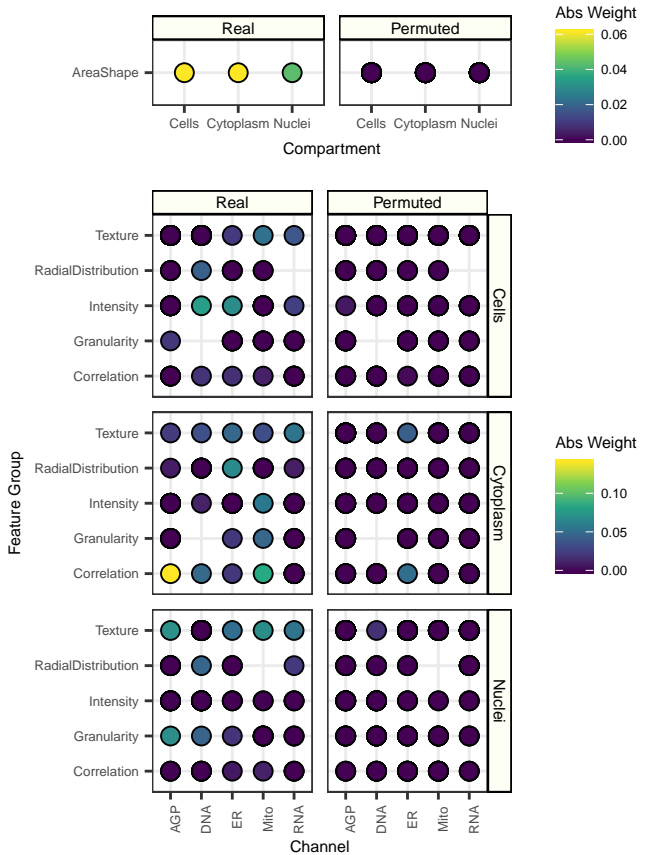
# CC - % High gH2AX spots



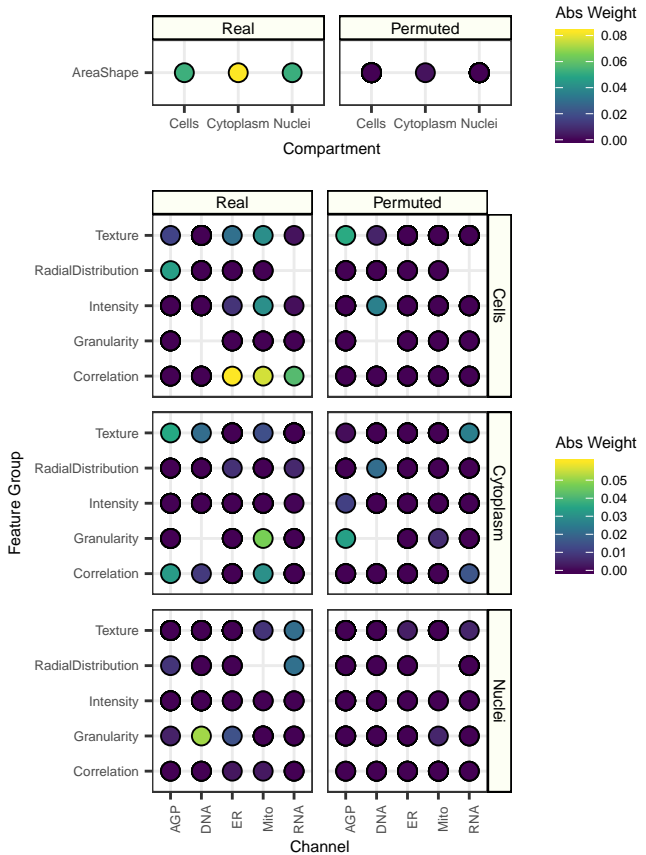
# CC – % Late M



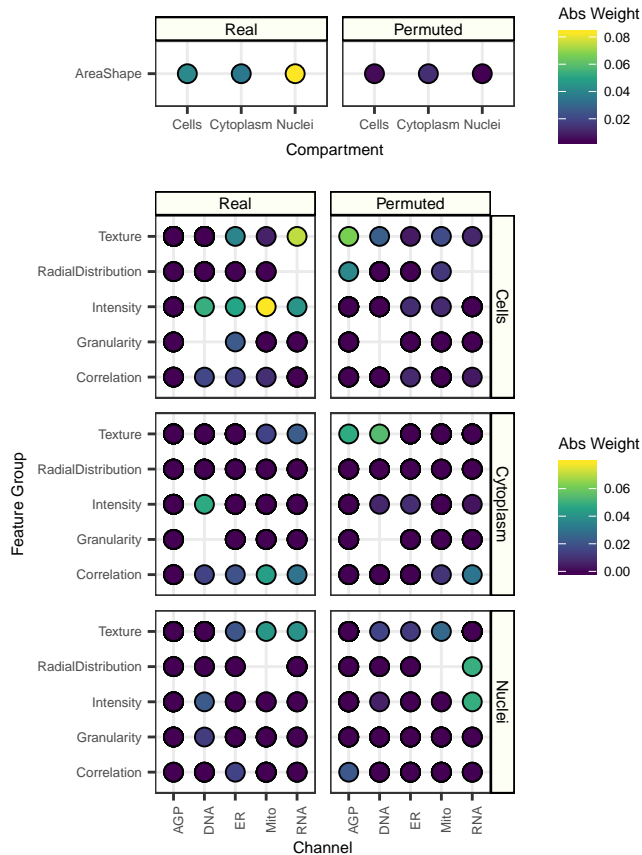
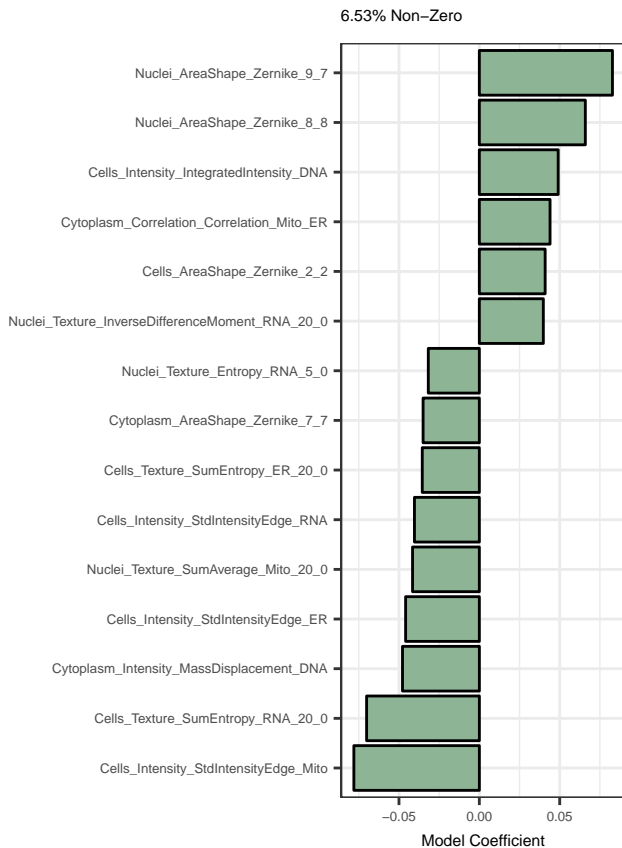
# CC - % M



# CC - # cells



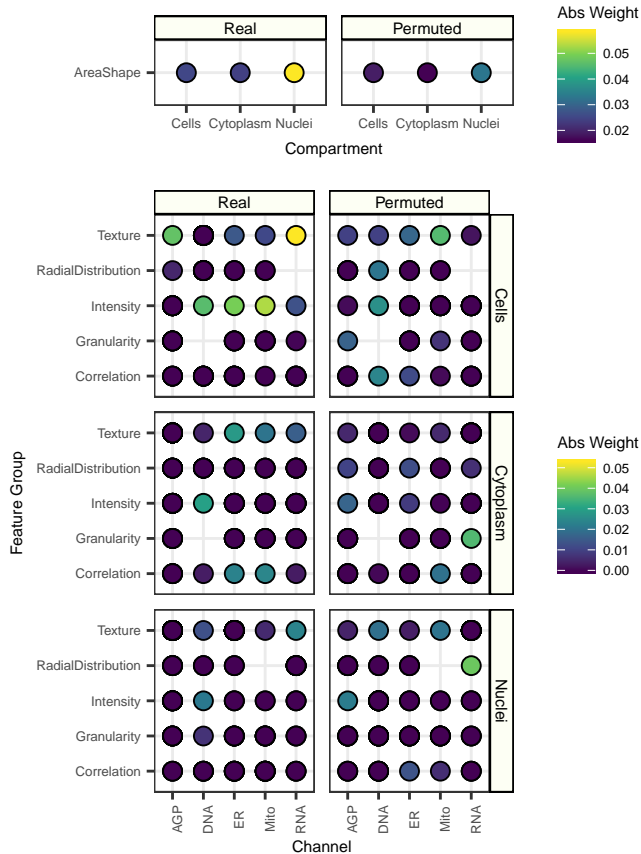
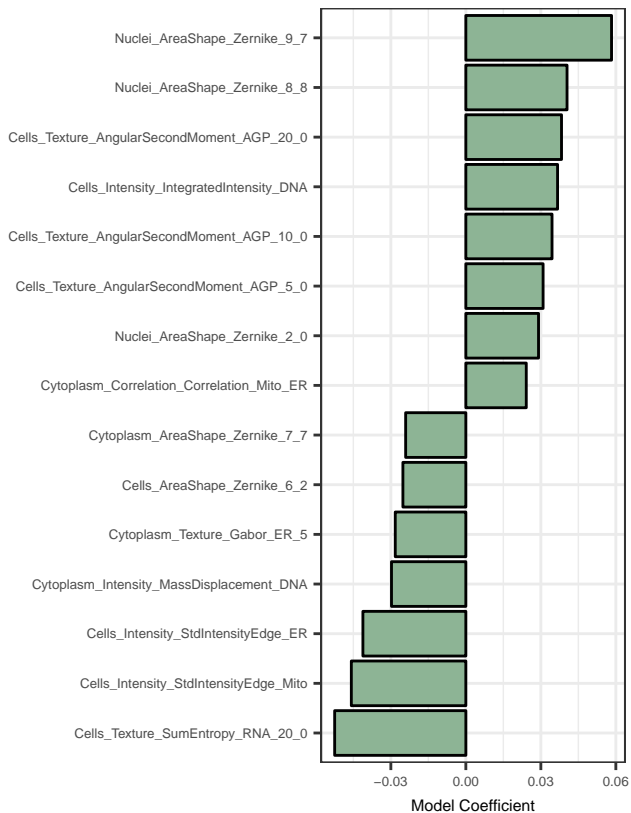
# CC - # of gH2AX Spots



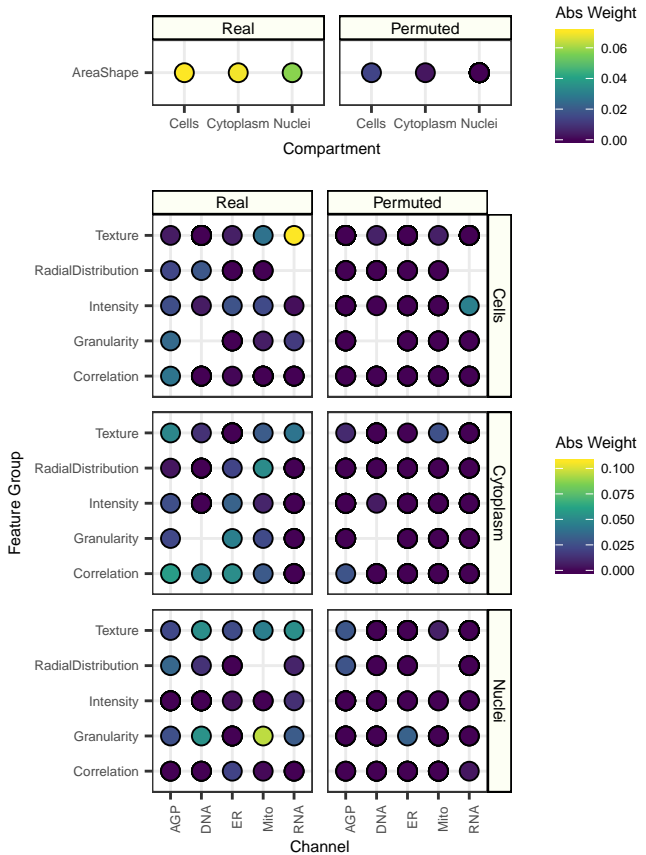
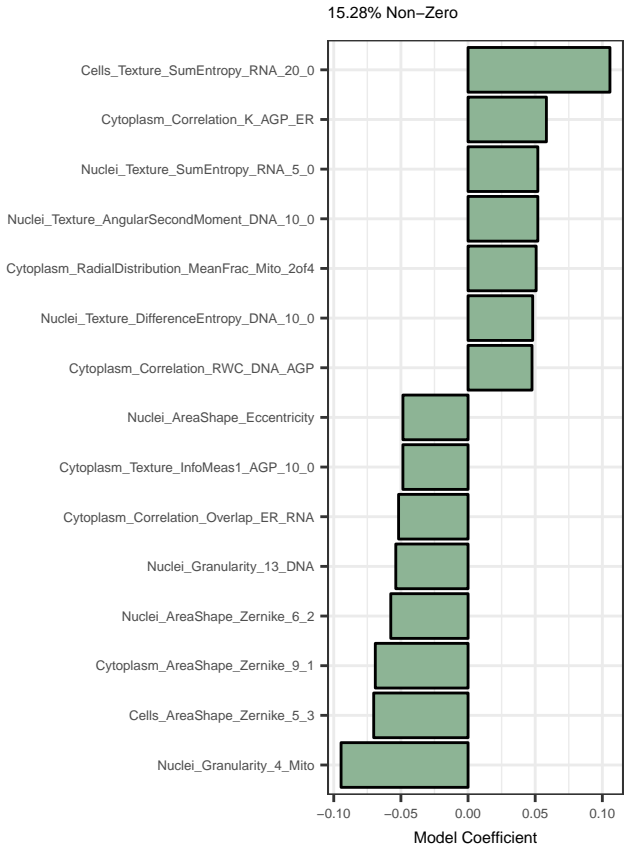


# CC – # of gH2AX Spots per Area of Nucleus

7.38% Non-Zero

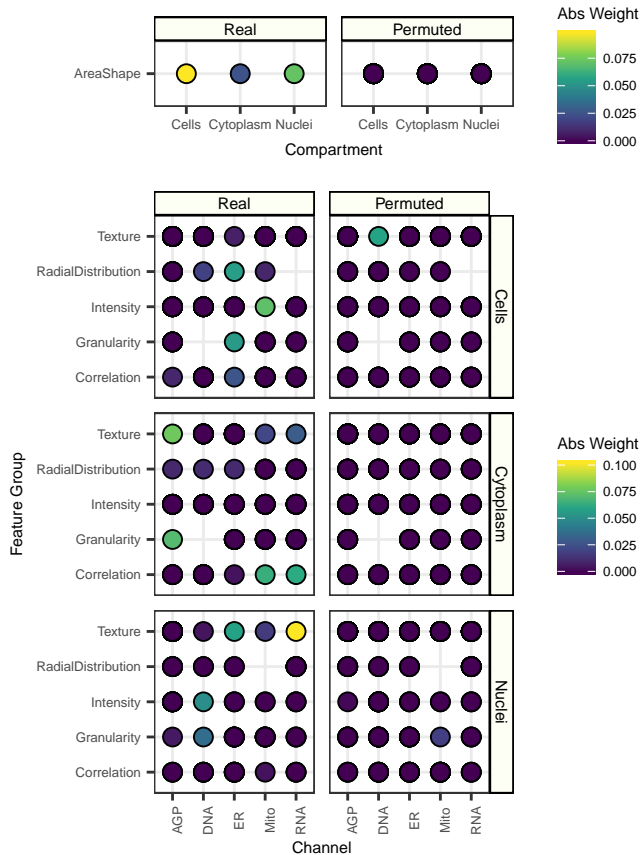
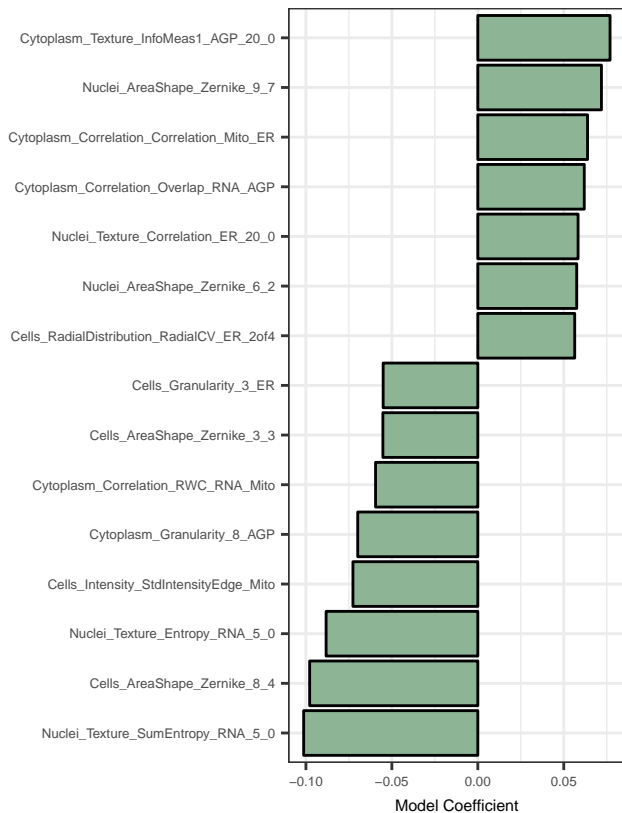


# CC - % S



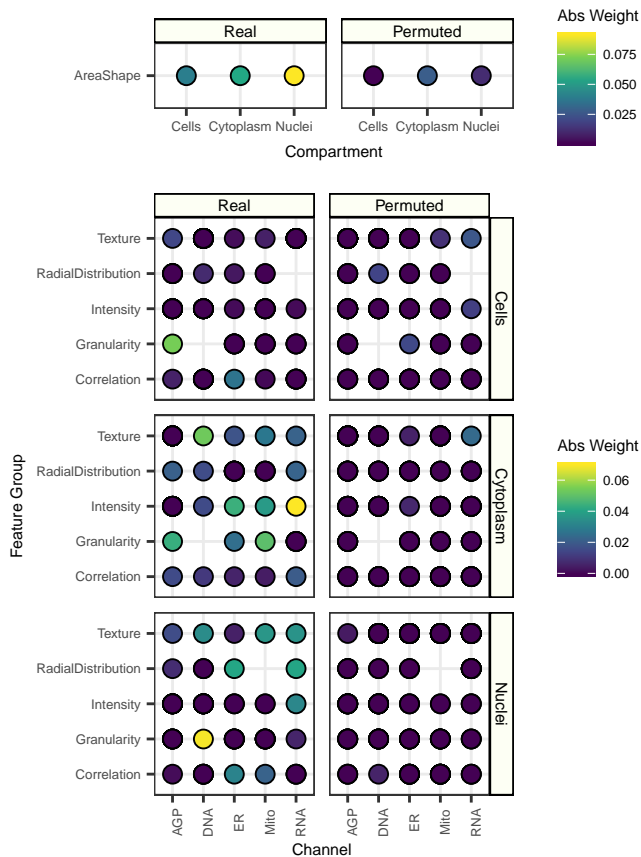
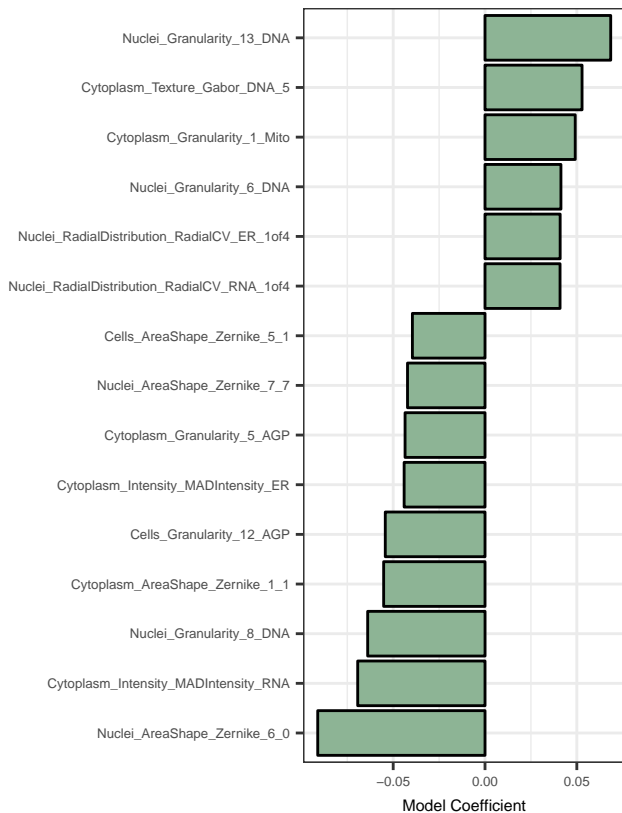
# early M – % High gH2AX Spots

7.06% Non-Zero

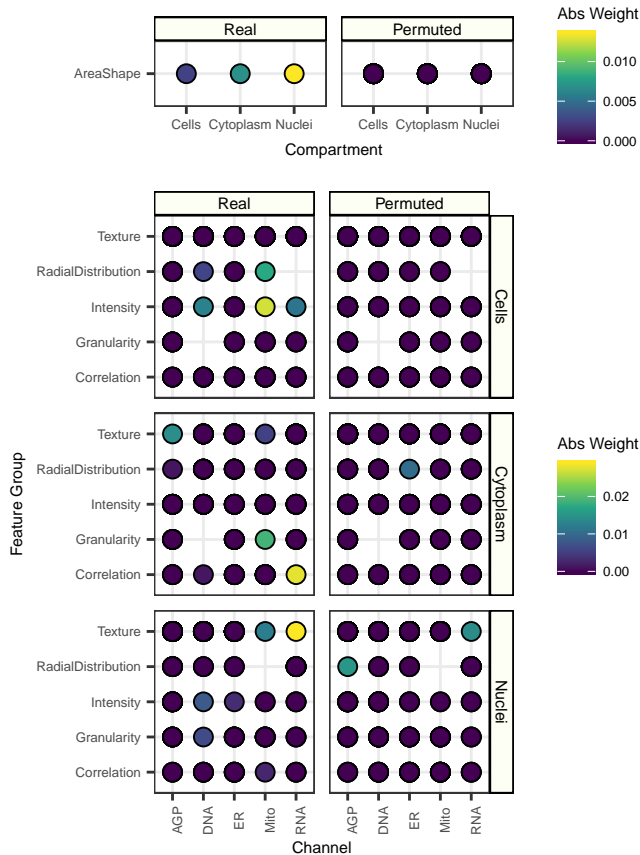
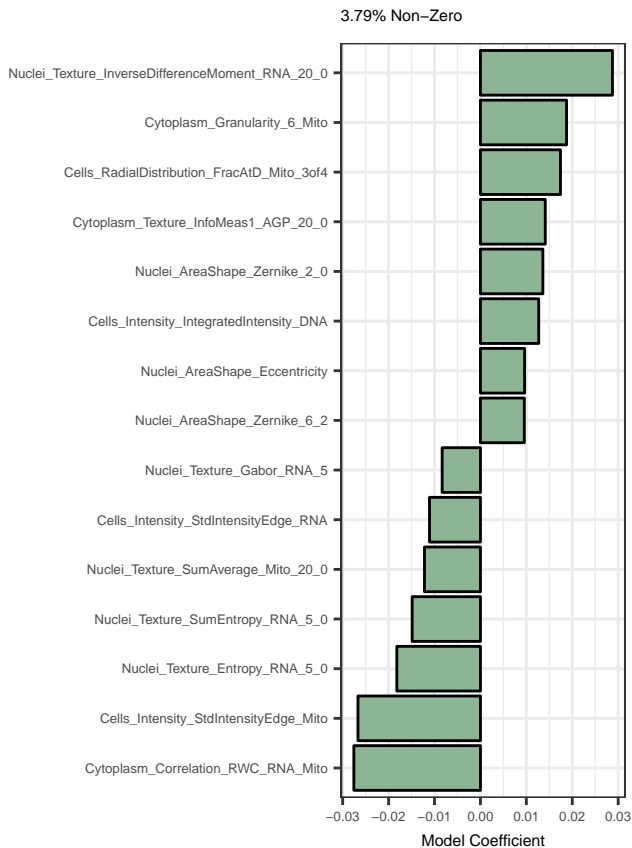


# early M – # cells

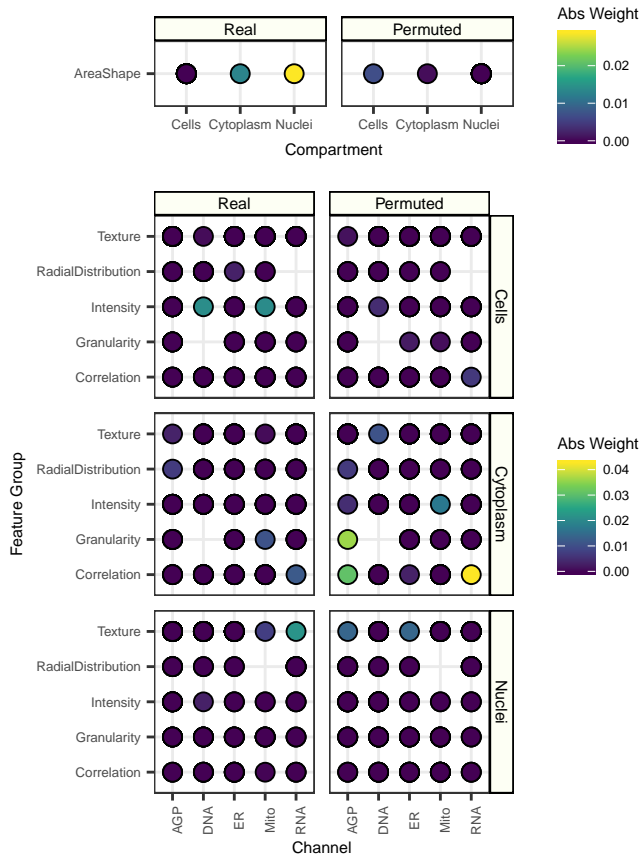
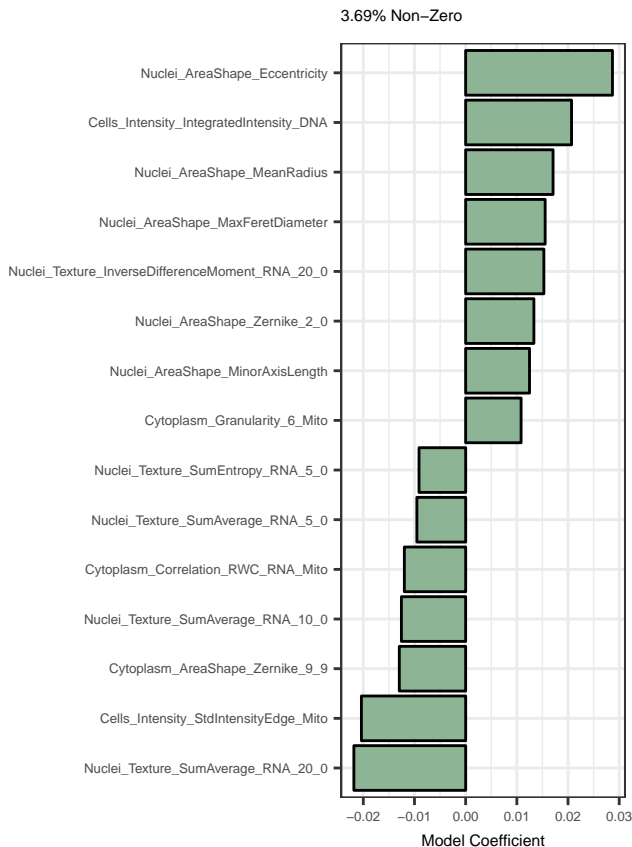
11.91% Non-Zero



# early M – # of gH2AX Spots

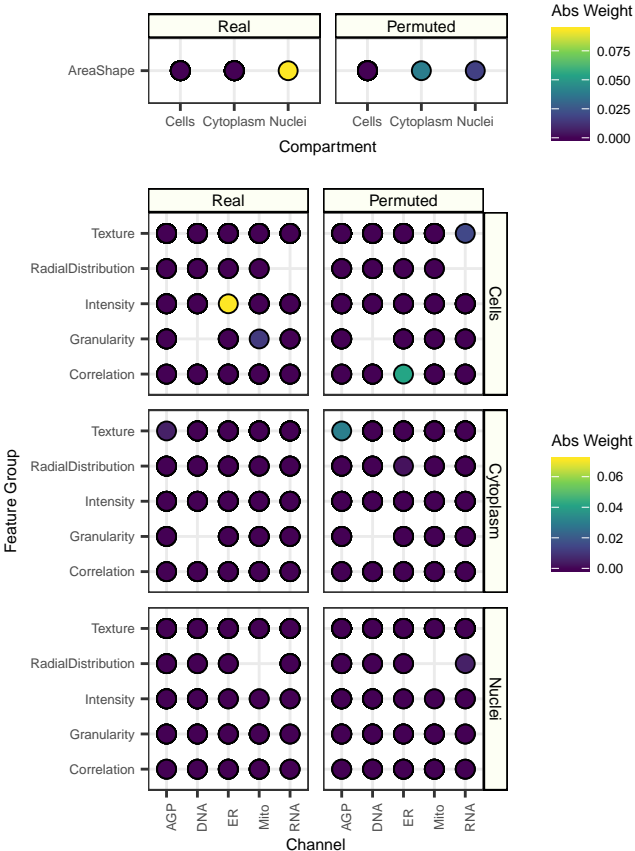
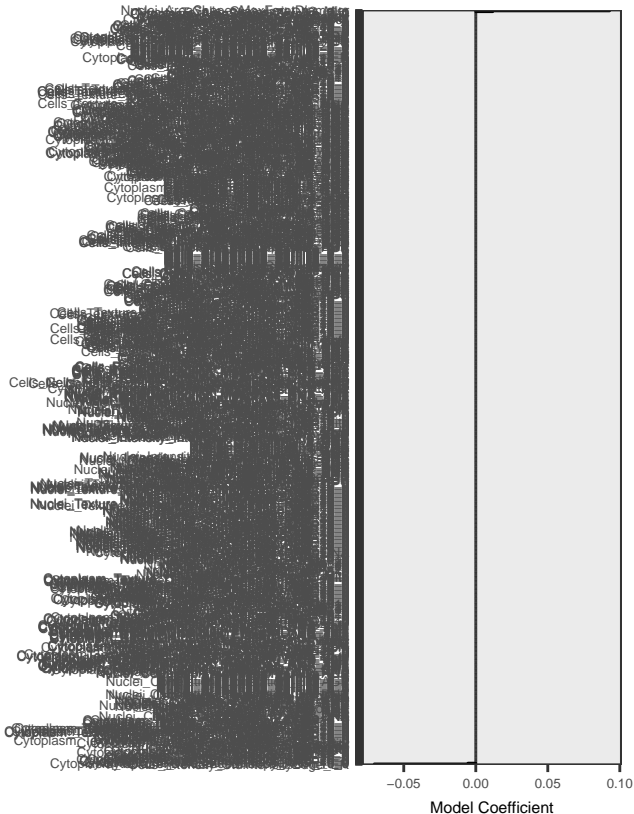


# early M – # of gH2AX Spots per Area of Nucleus

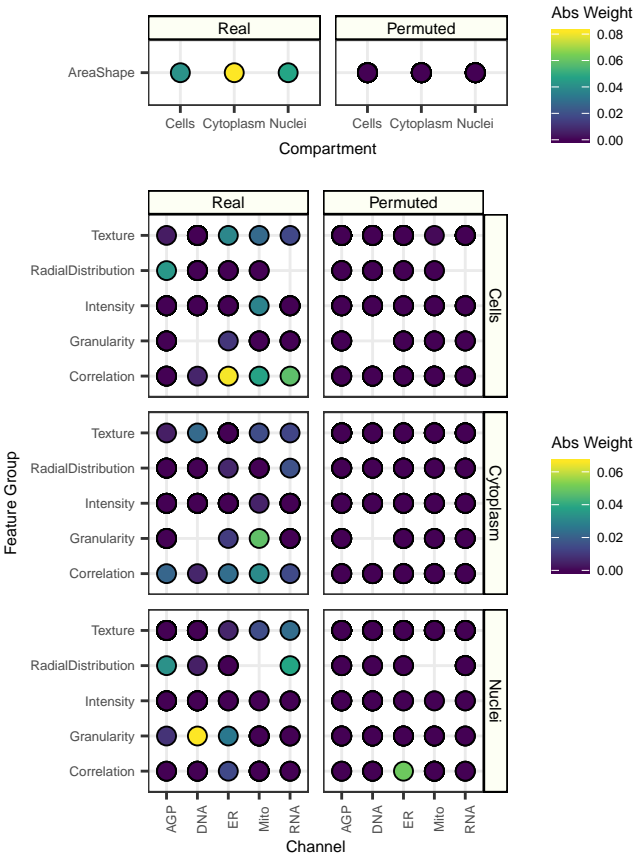
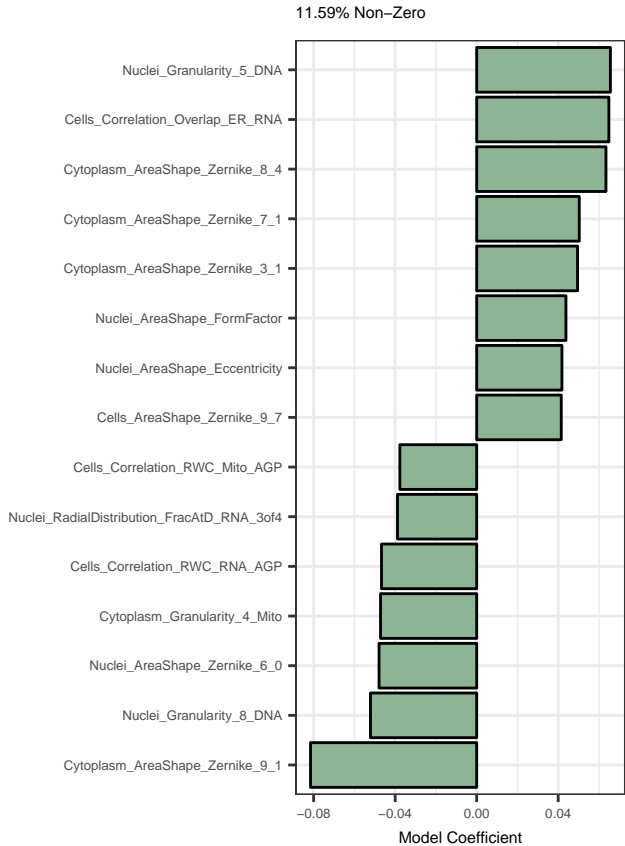


# G1 – % High gH2AX Spots

0.42% Non-Zero

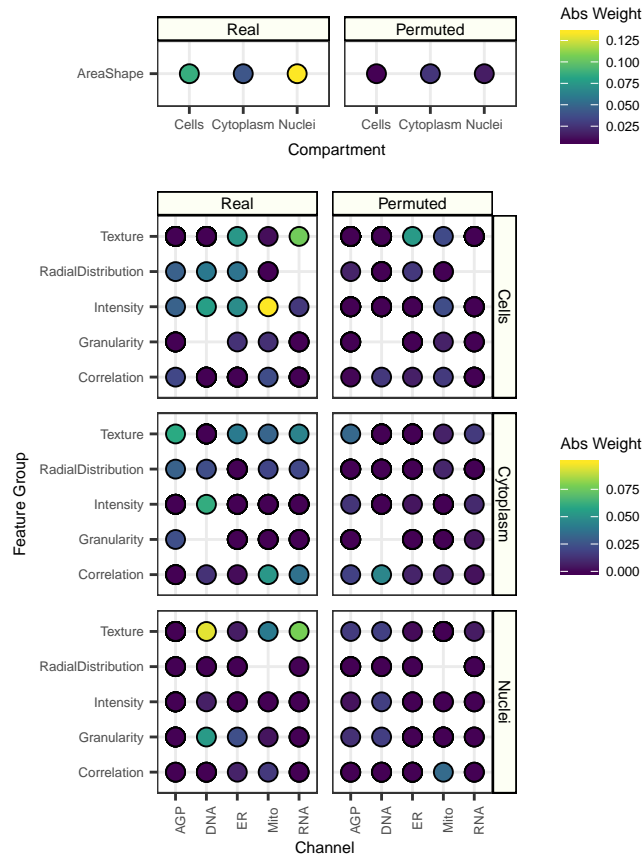


# G1 – # cells

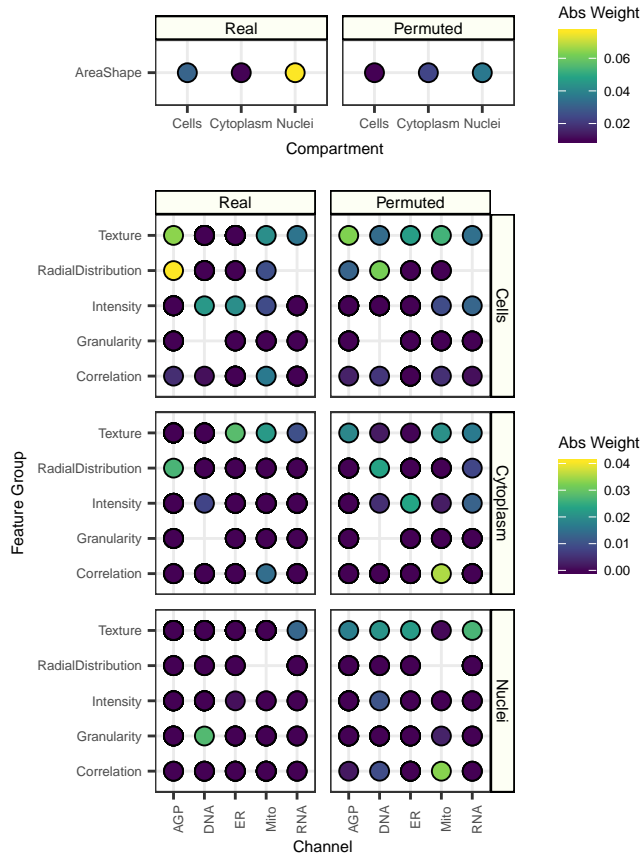
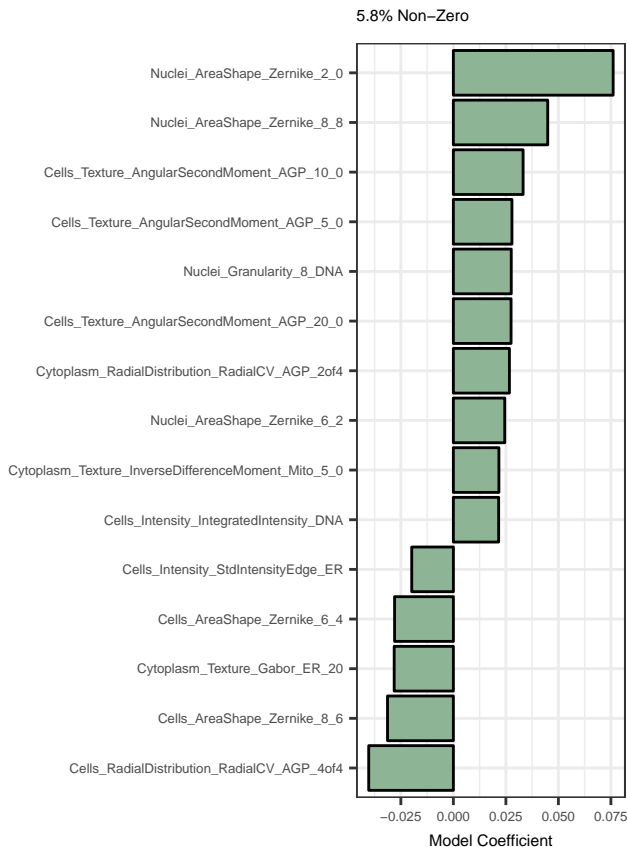




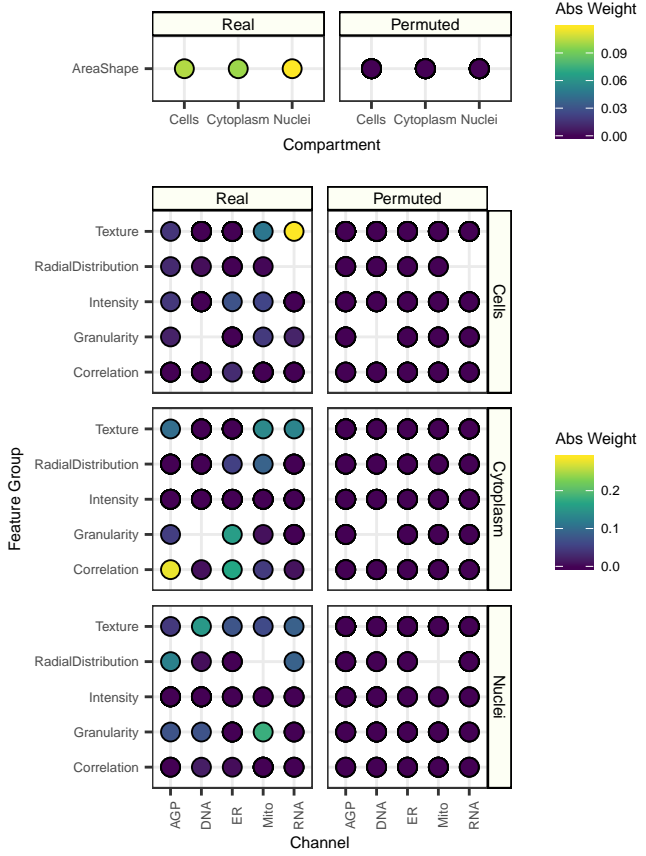
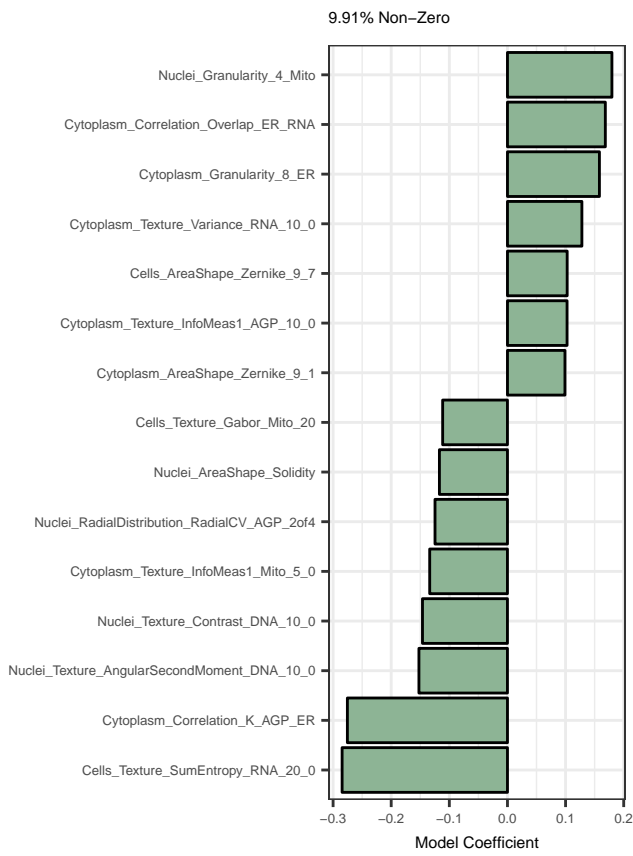
# G1 – # of gH2AX Spots



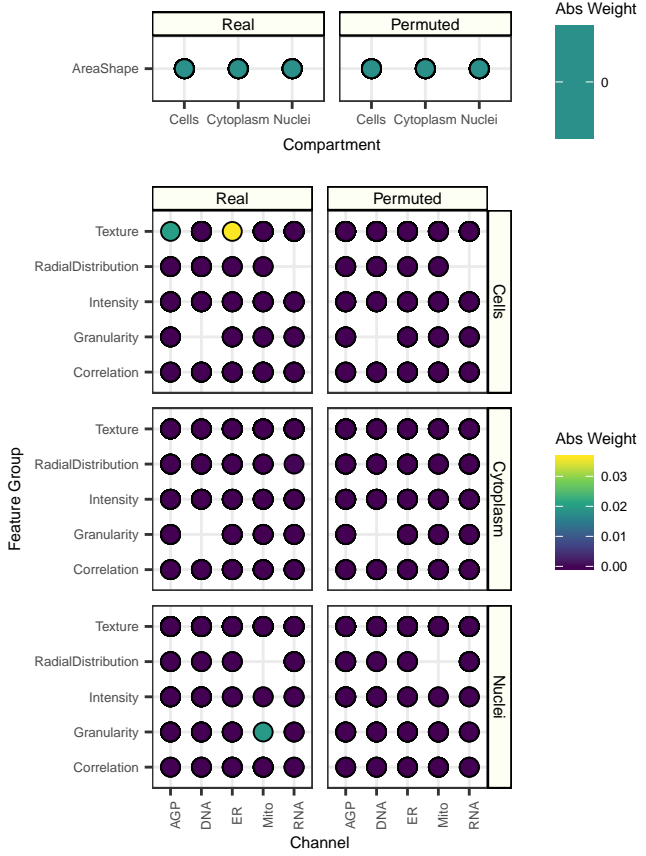
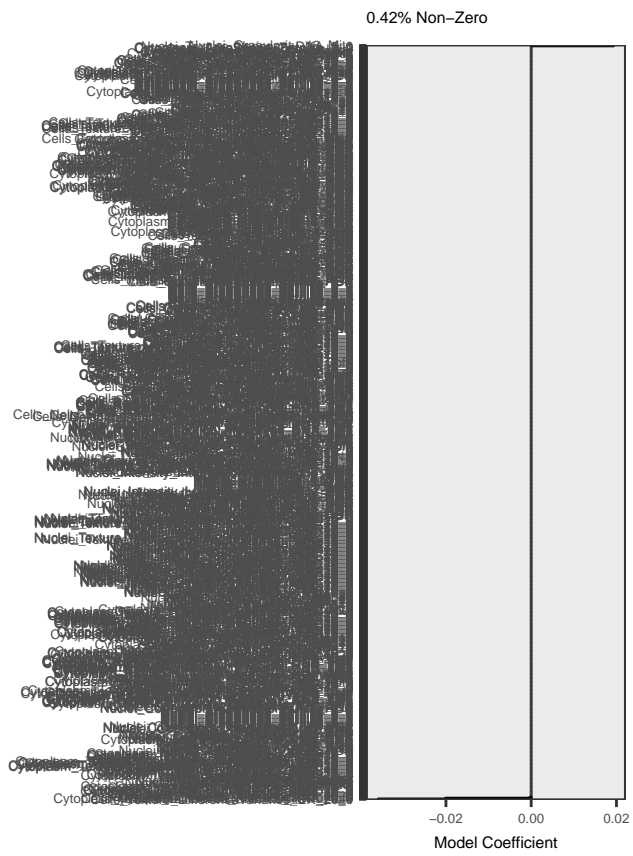
# G1 – # of gH2AX Spots per Area of Nucleus



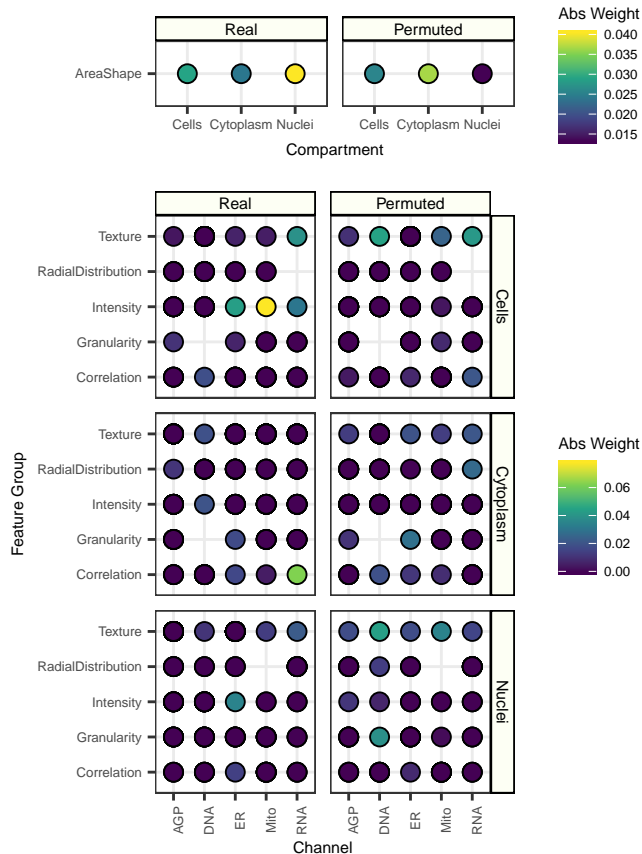
# Cell Count – G1+G2



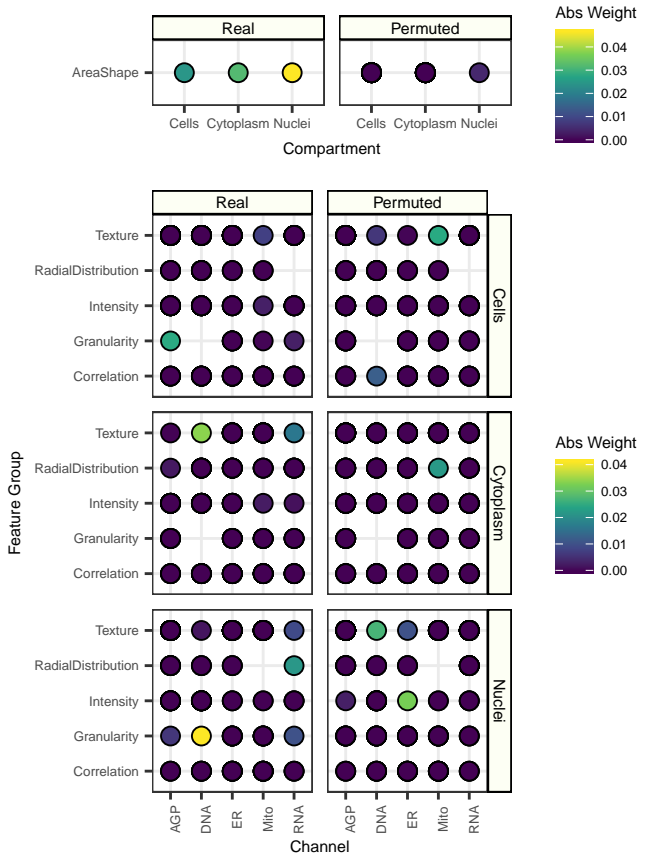
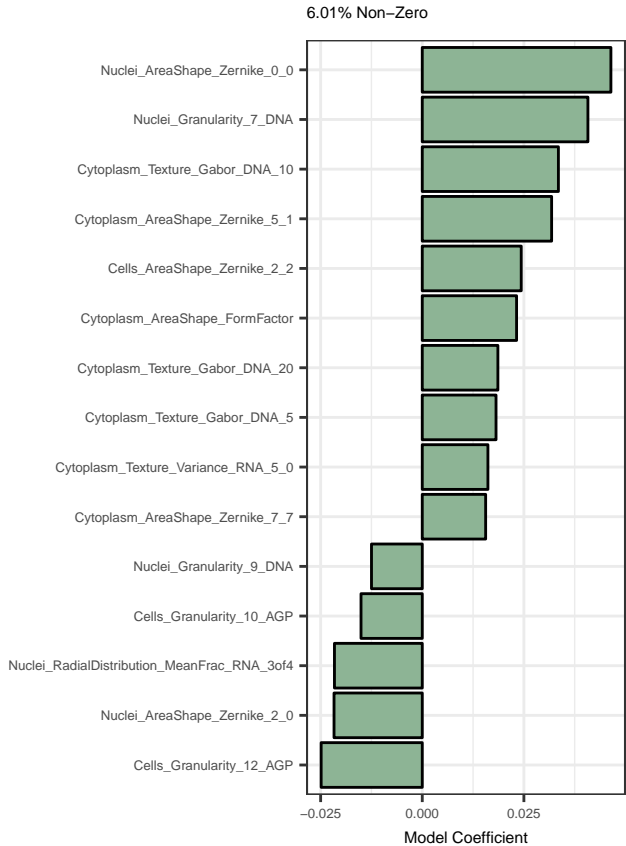
# Cell Count – G2/G1



# G2 - % High gH2AX Spots

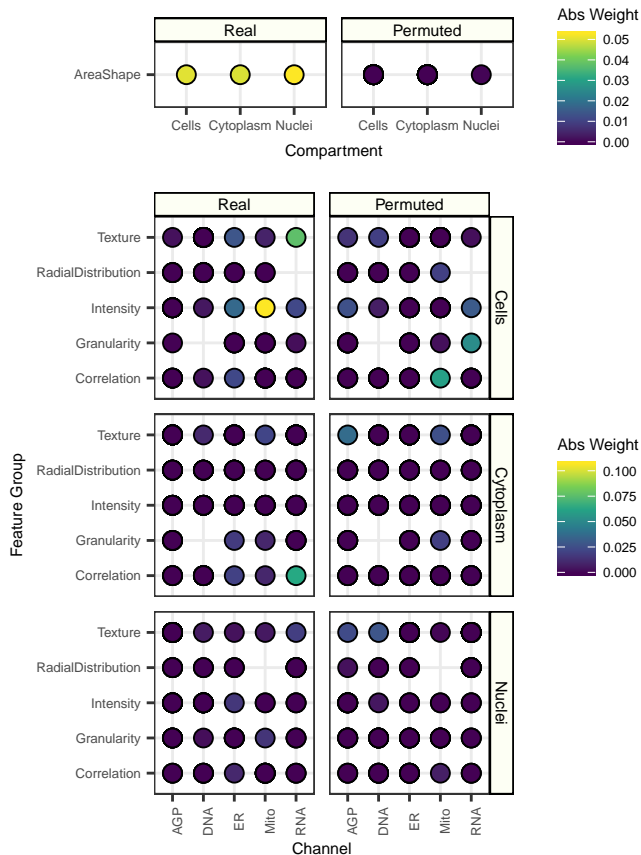
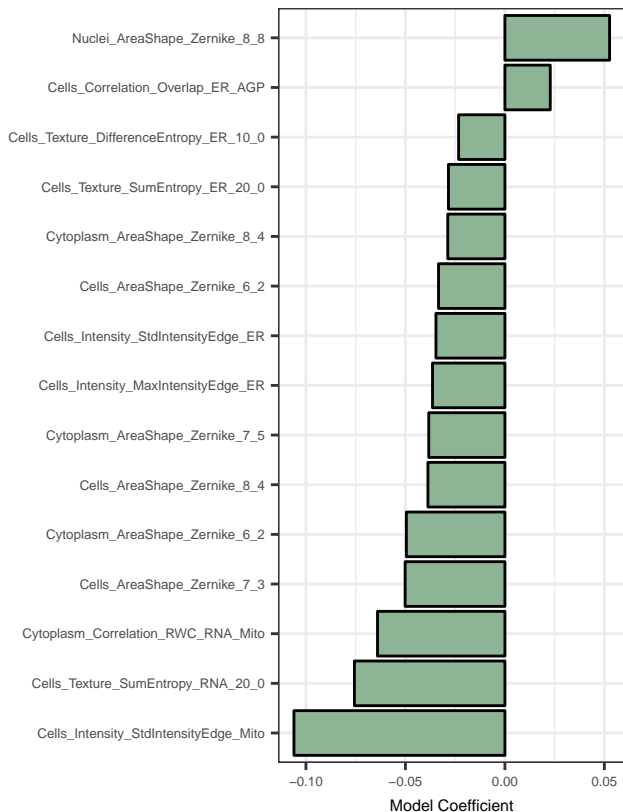


# G2 - # cells



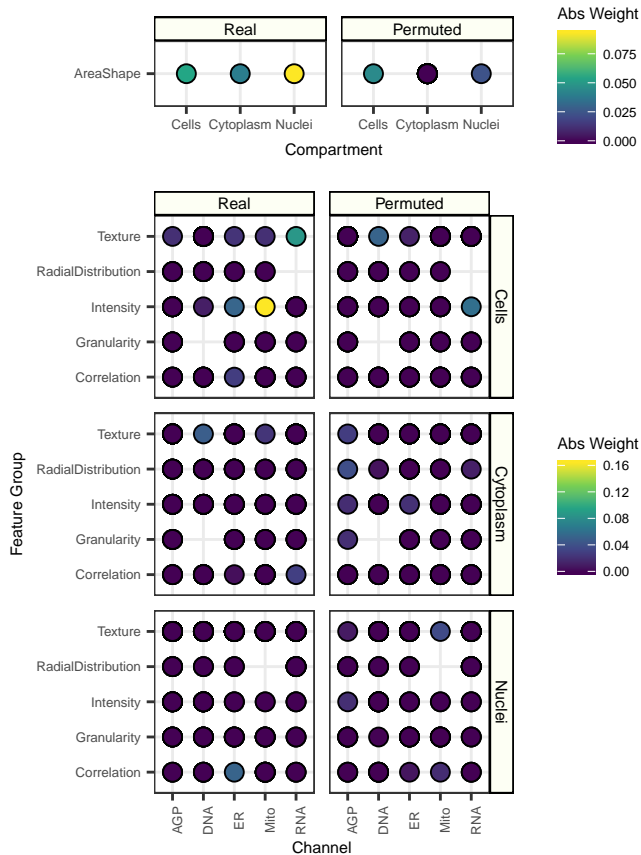
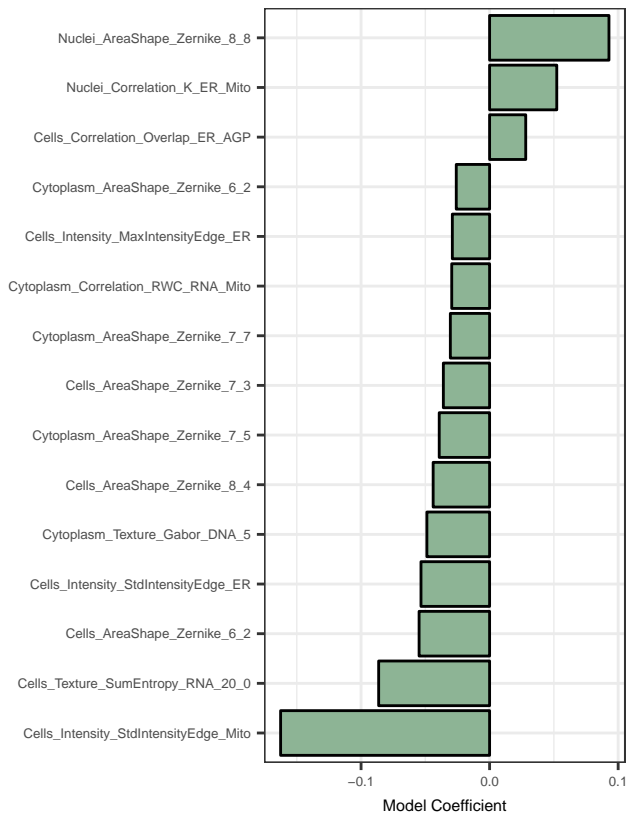
# G2 – # of gH2AX Spots

6.85% Non-Zero



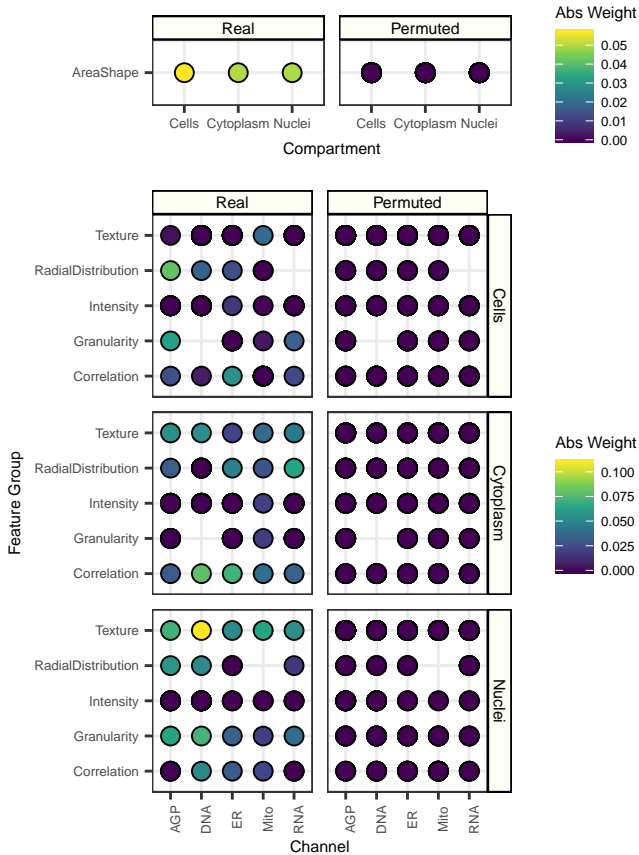
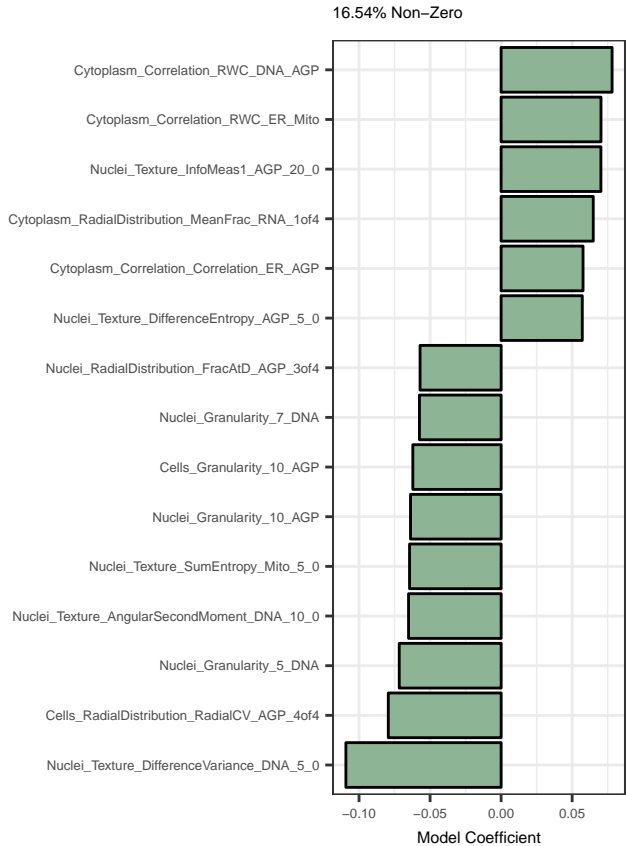
# G2 – # of gH2AX Spots per Area of Nucleus

2.63% Non-Zero

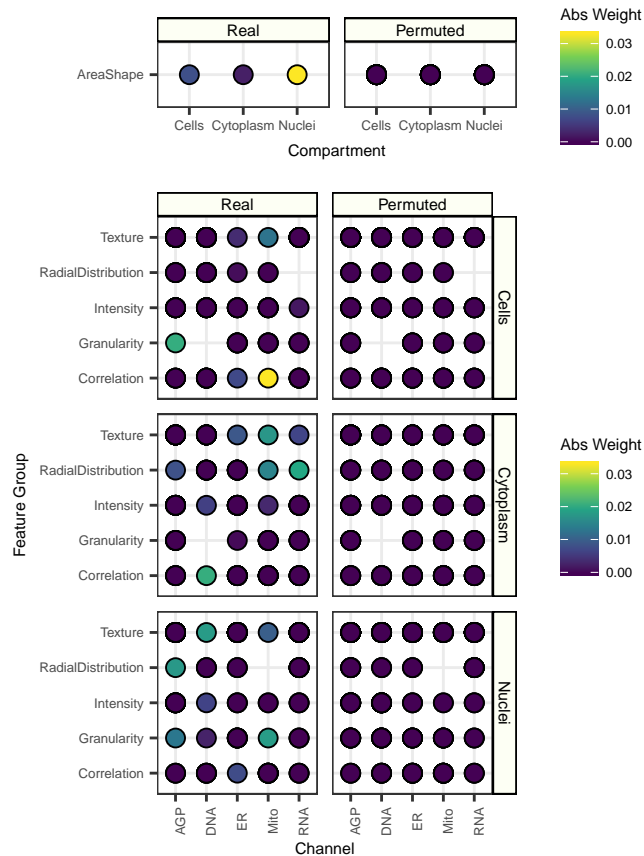
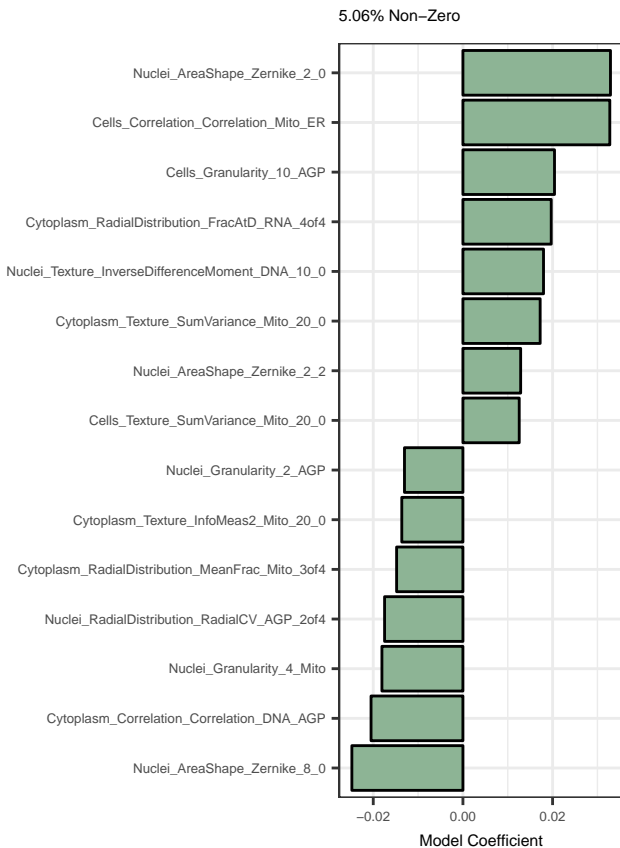




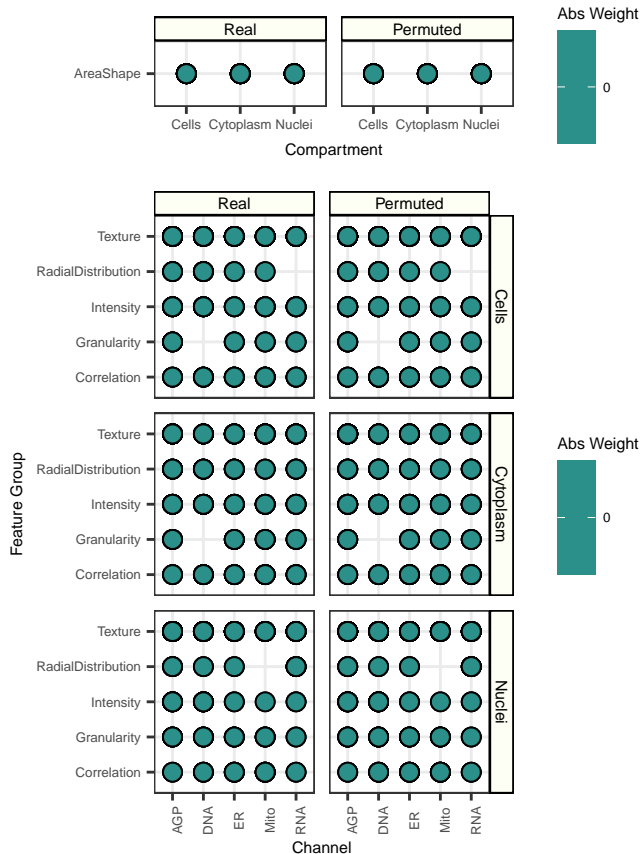
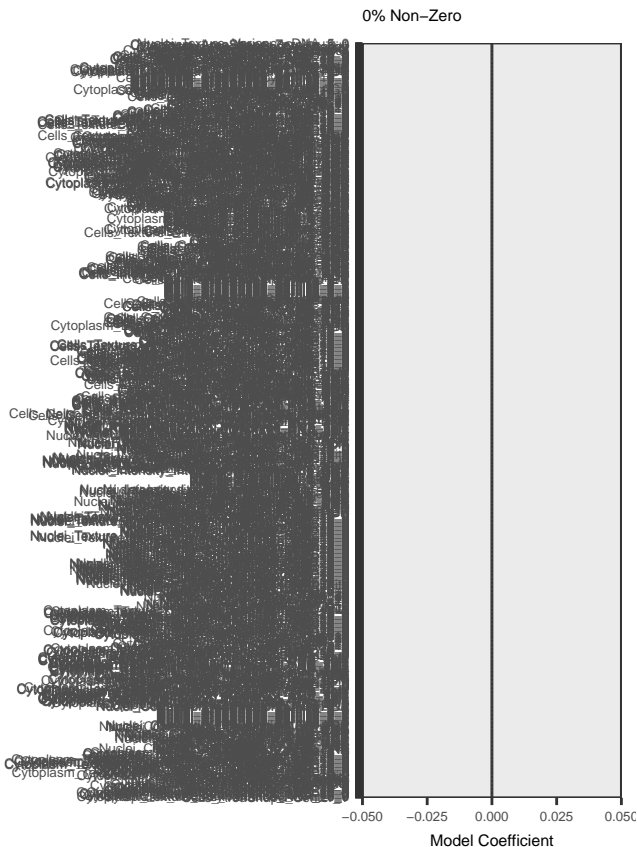
# Cell Count – G2 + M



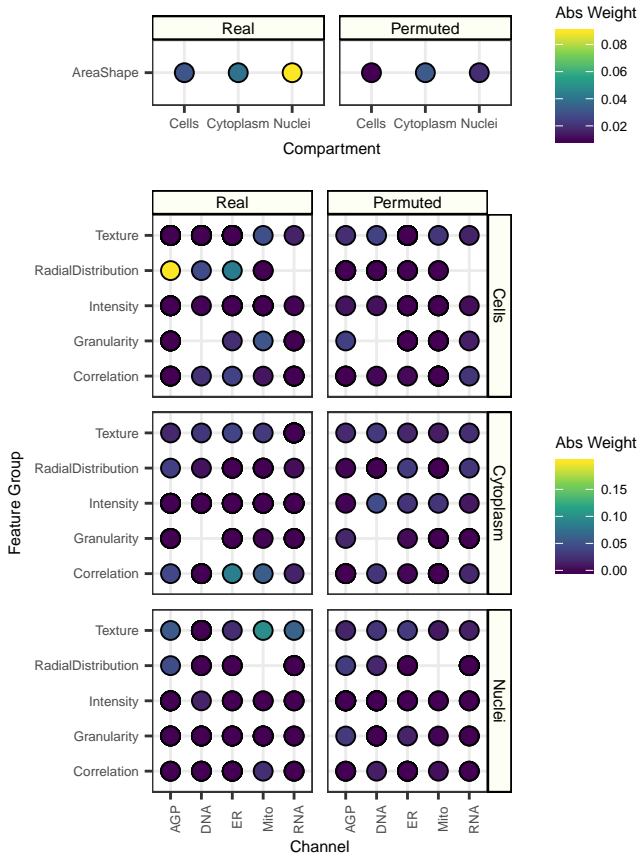
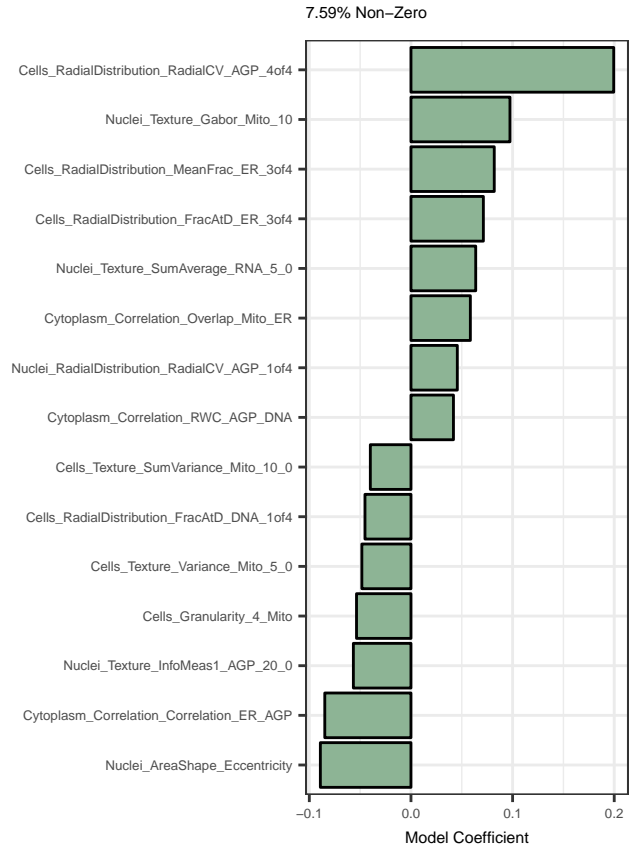
# Inf %



# Late M – % High gH2AX Spots

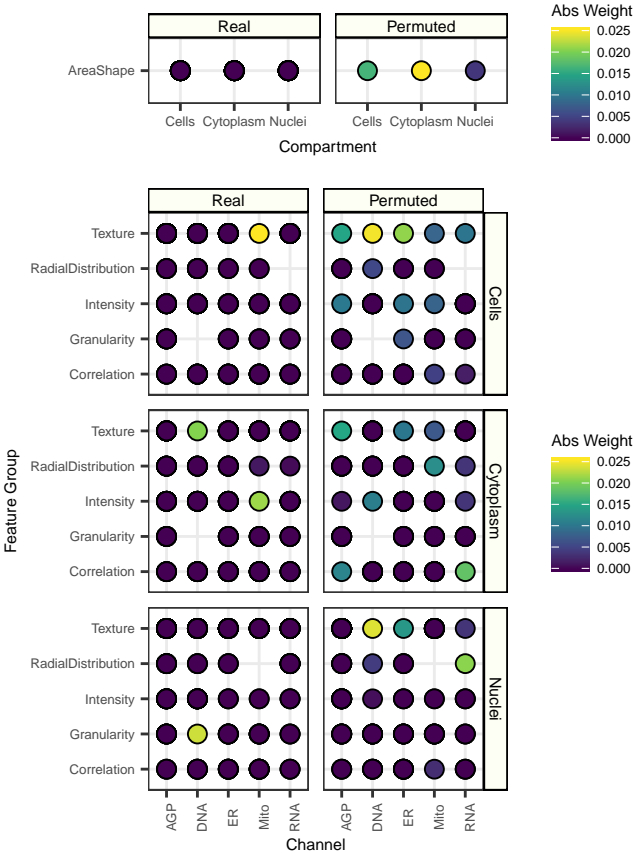
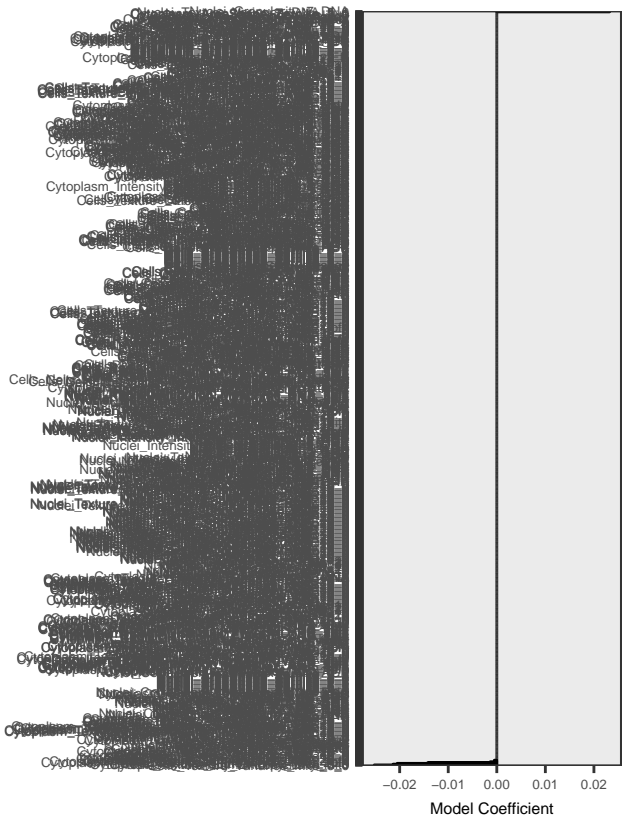


# Late M – # cells

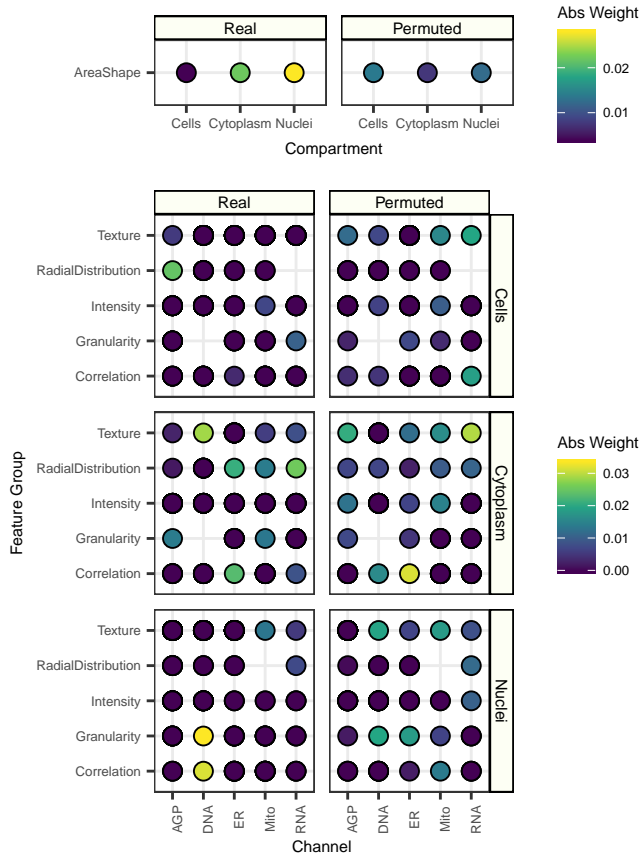
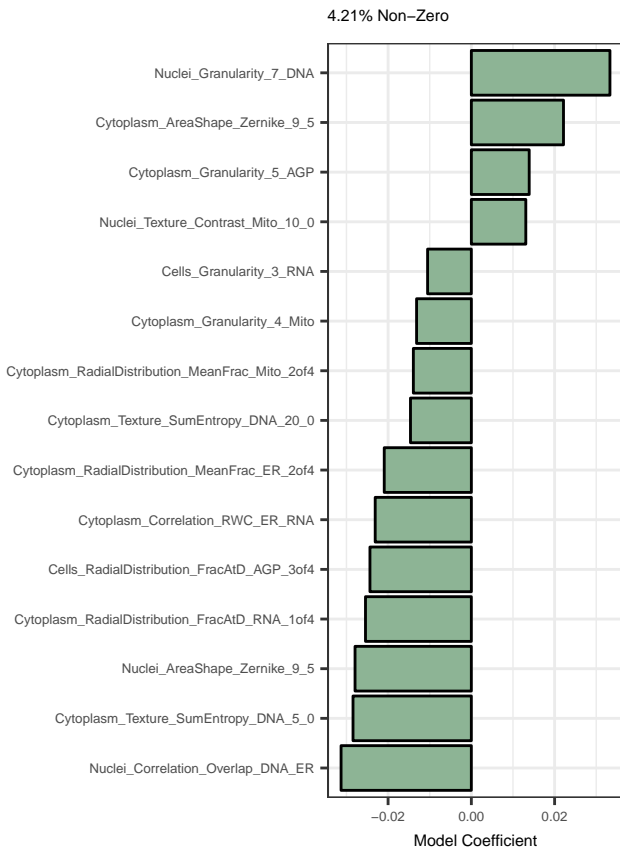


# Late M – # of gH2AX Spots

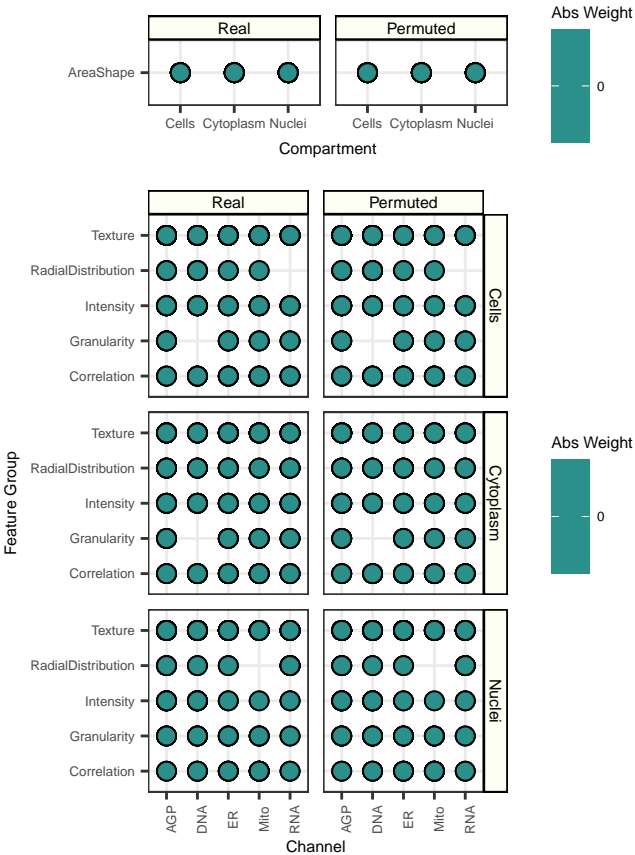
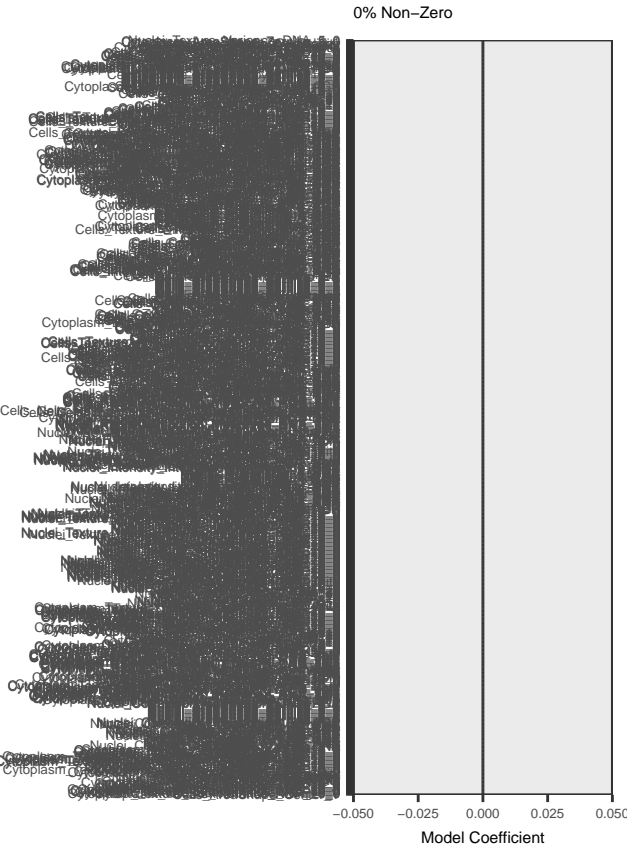
0.84% Non-Zero



# Late M – # of gH2AX Spots per Area of Nucleus

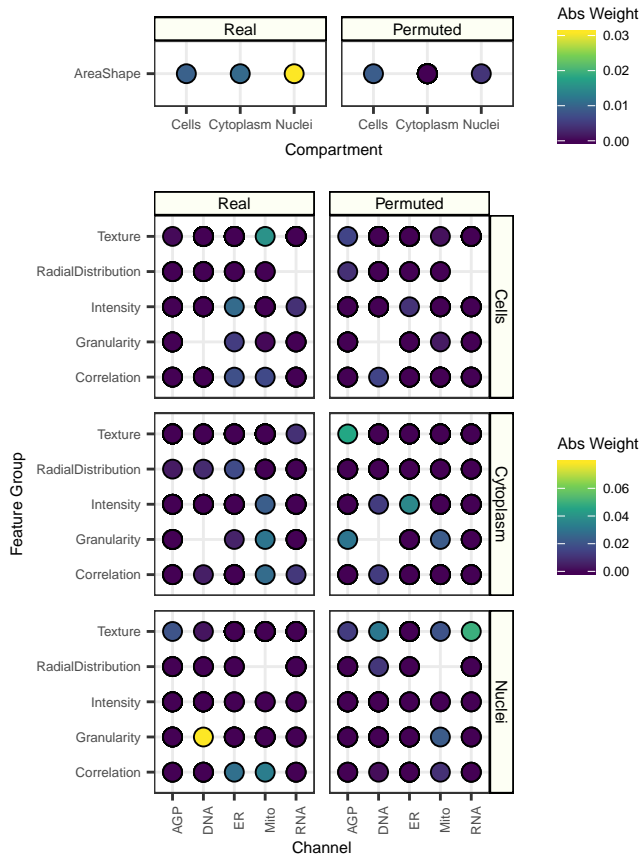
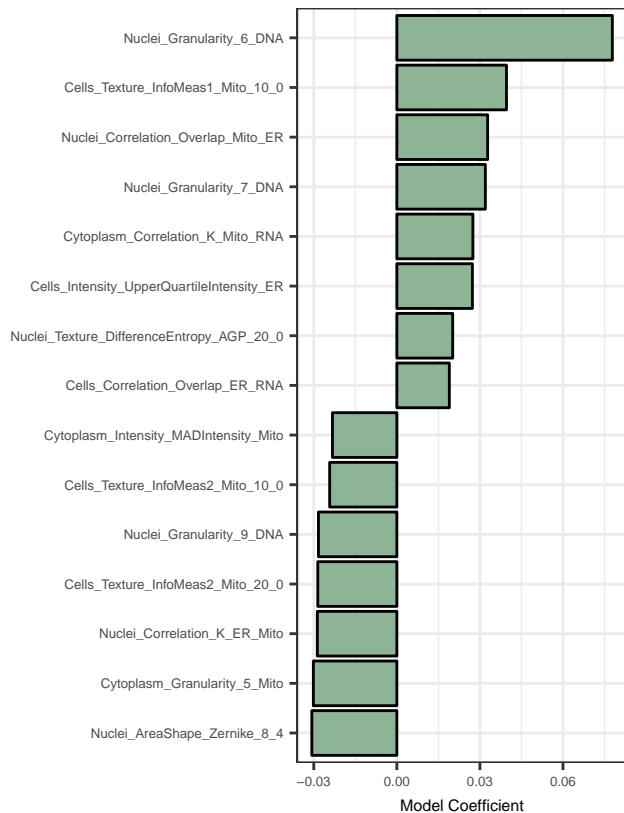


# M – % High gH2AX Spots



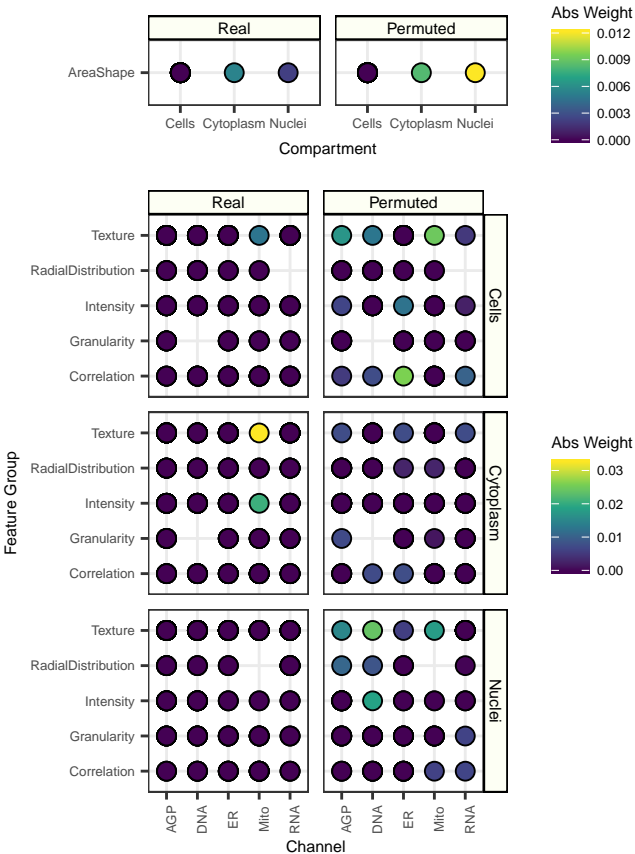
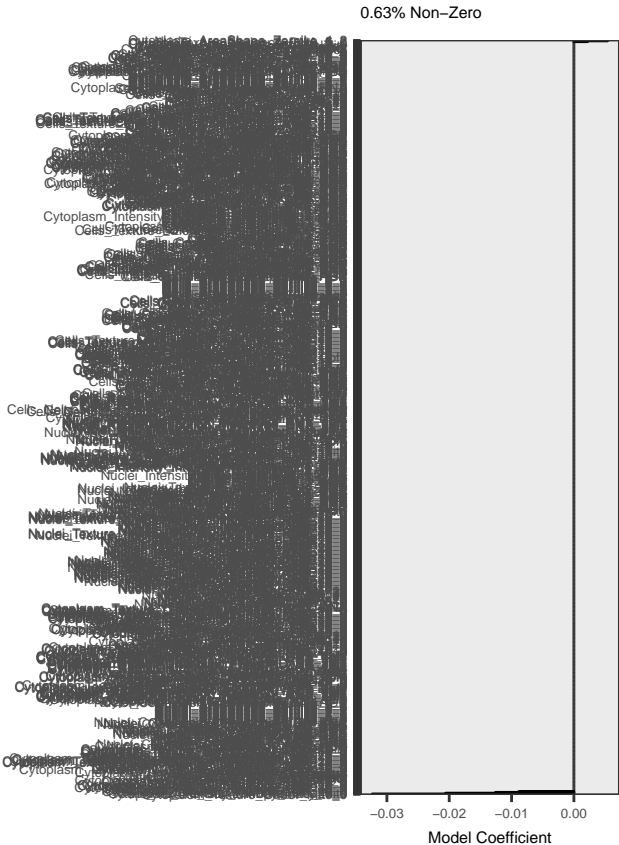
# M - # cells

6.11% Non-Zero



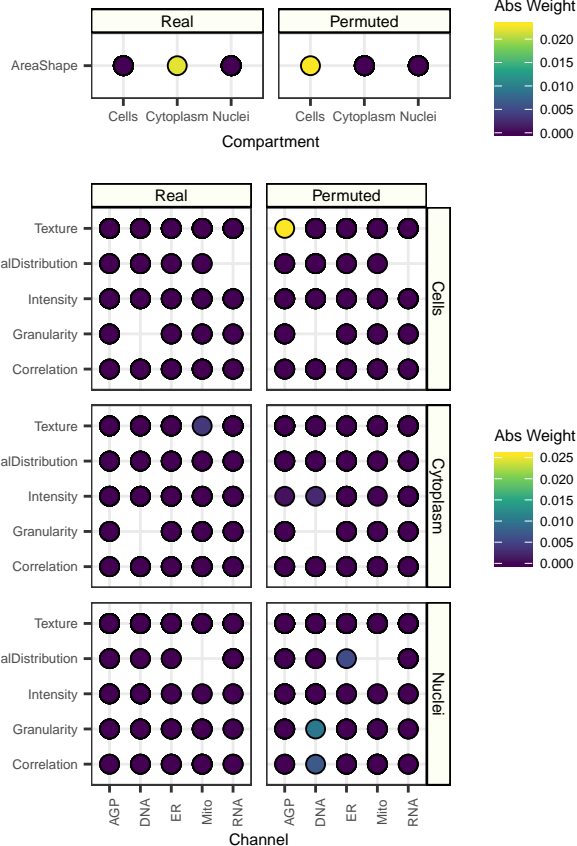
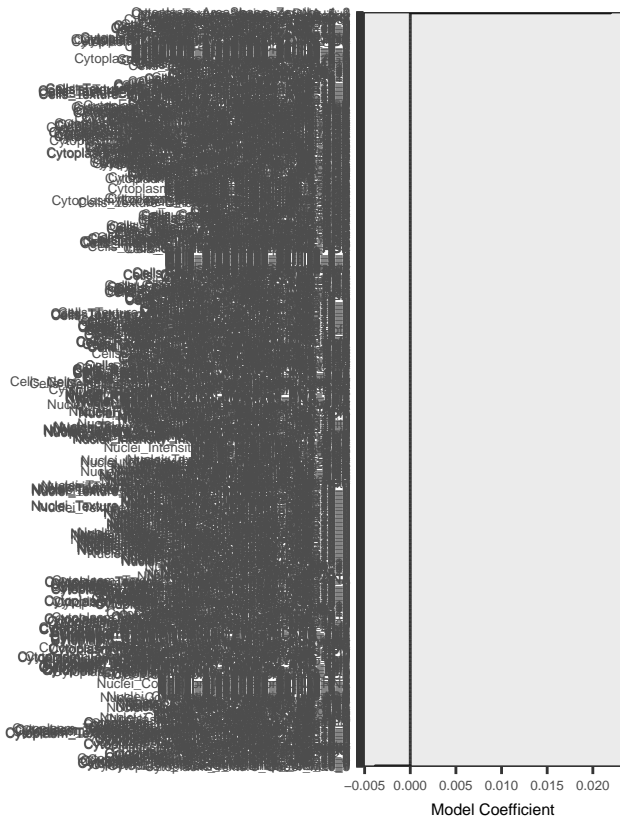


# M – # of gH2AX Spots



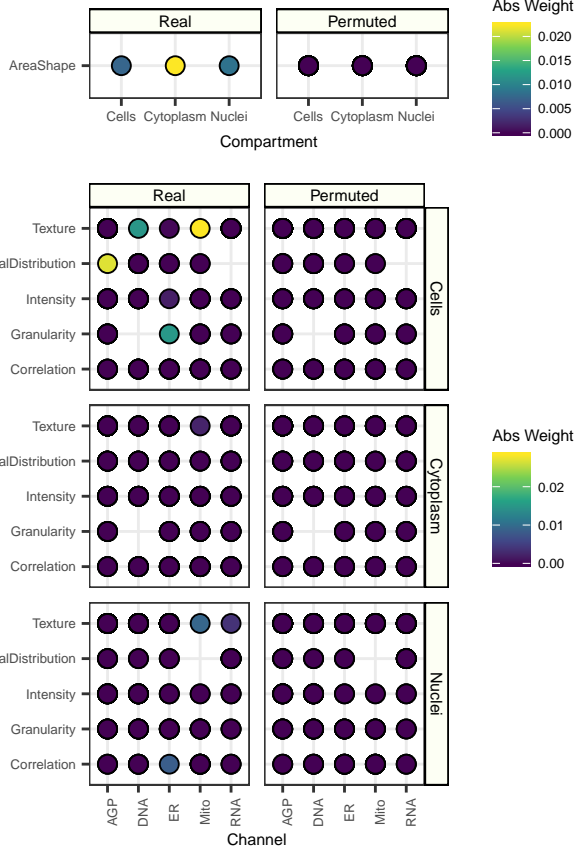
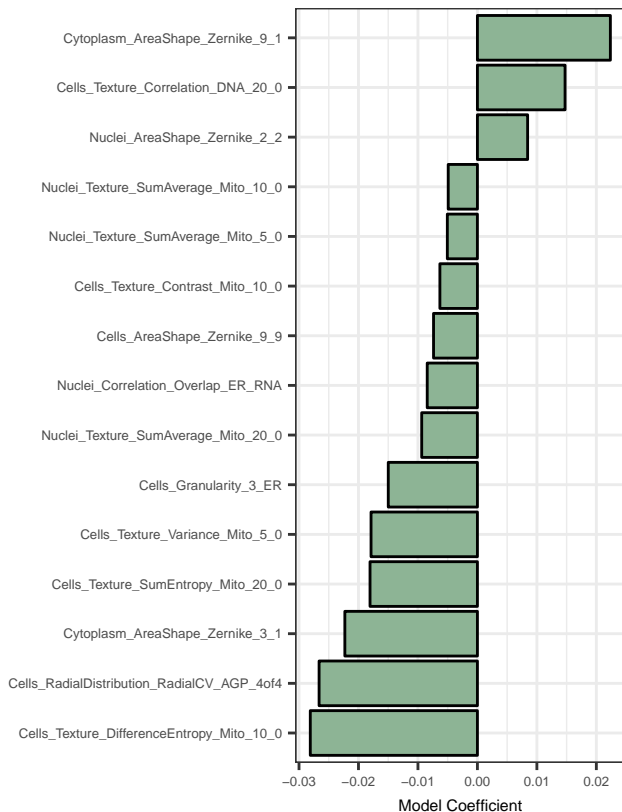
# M – # of gH2AX Spots per Area of Nucleus

0.21% Non-Zero

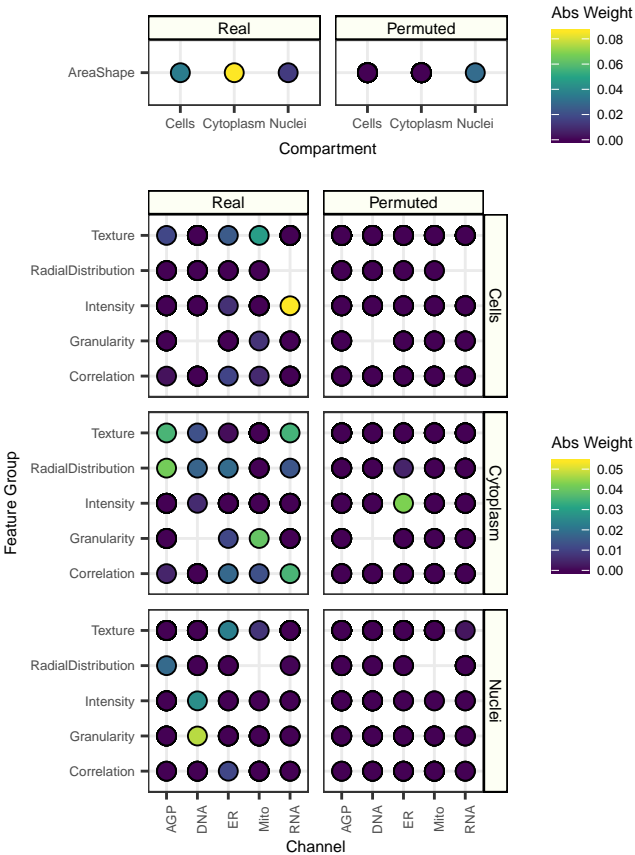
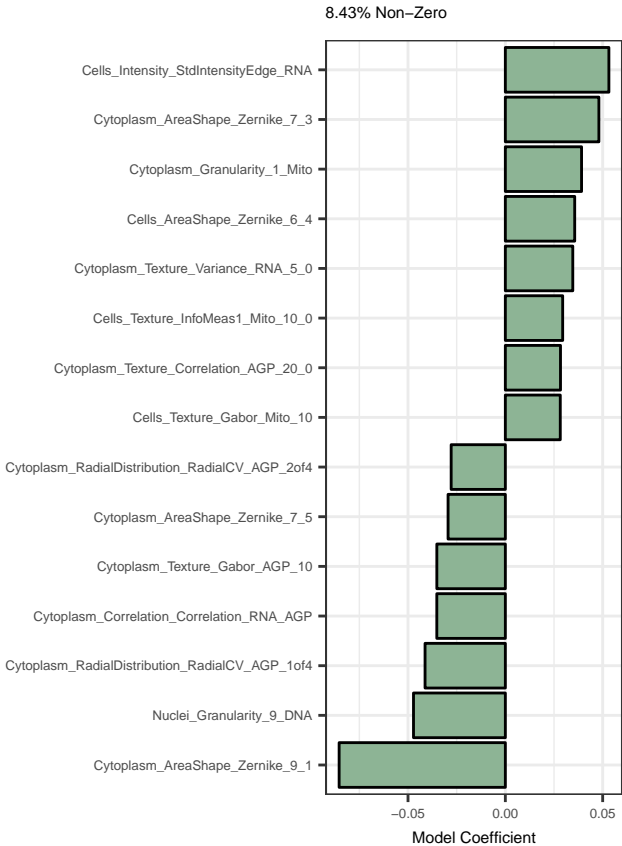


# Polynuclear – % High gH2AX Spots

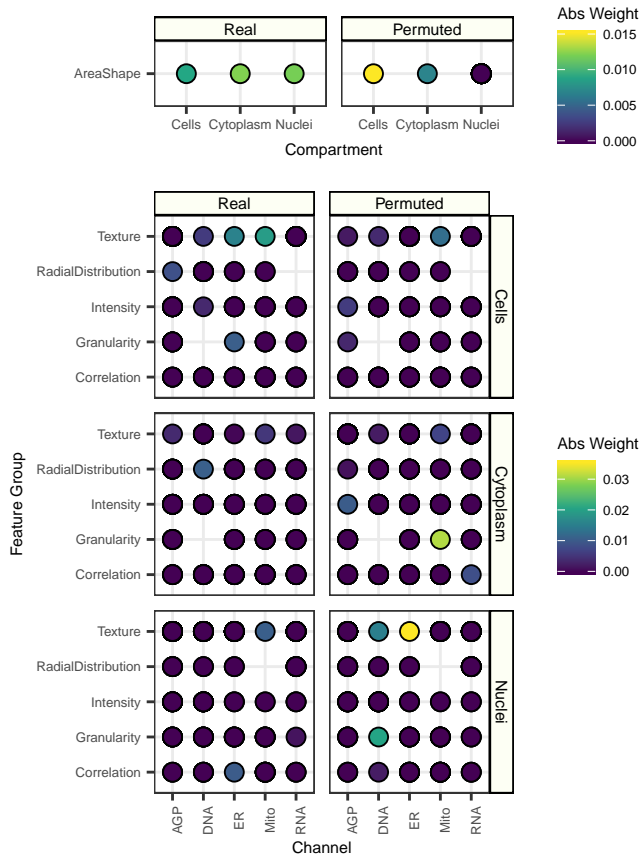
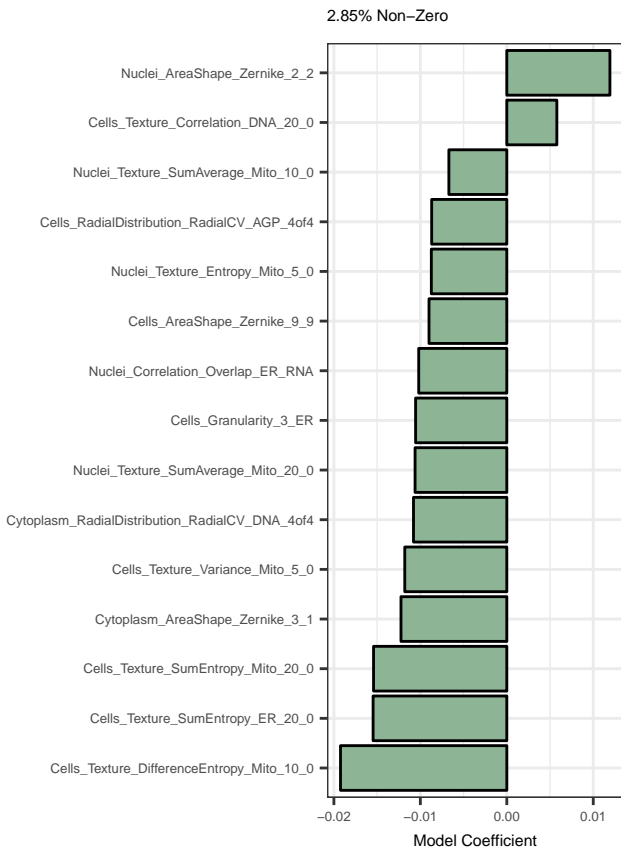
2.21% Non-Zero



# Polynuclear – # cells

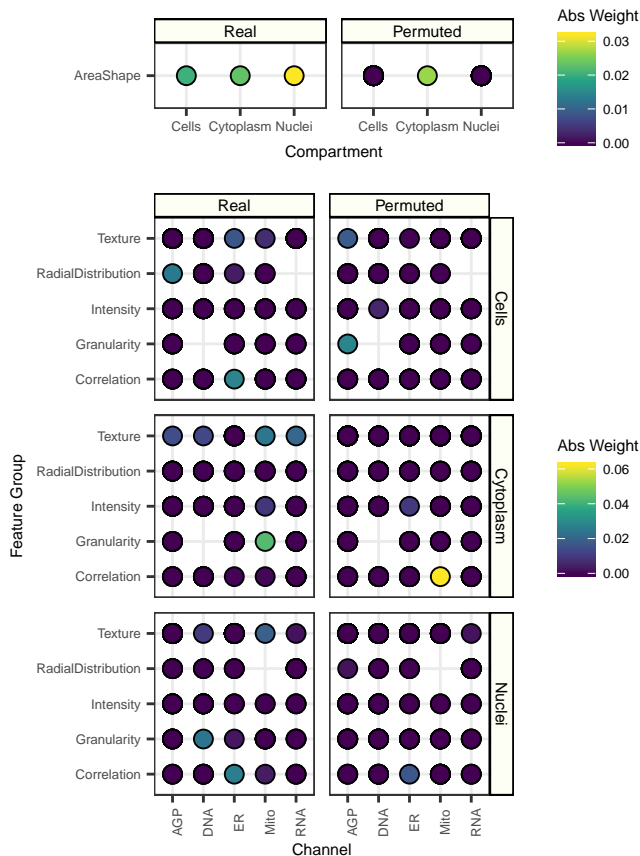
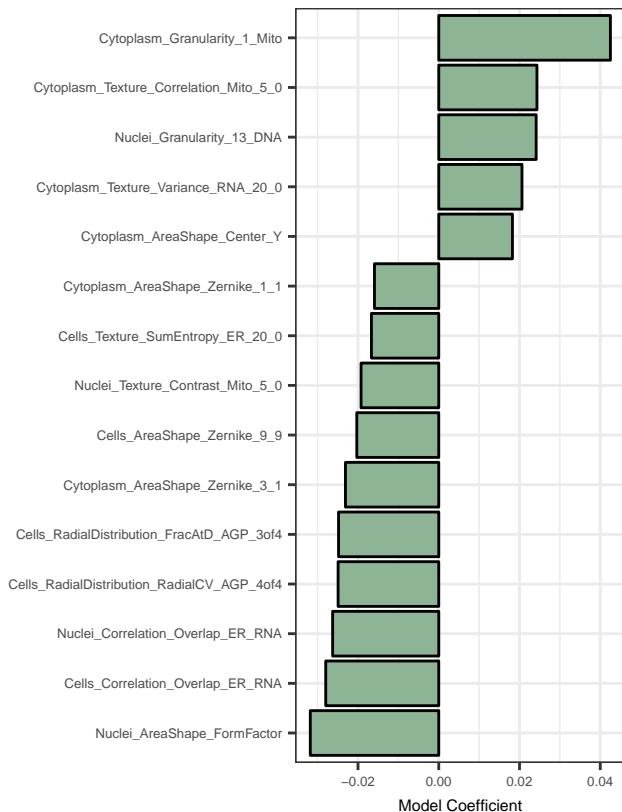


# Polynuclear – # of gH2AX Spots

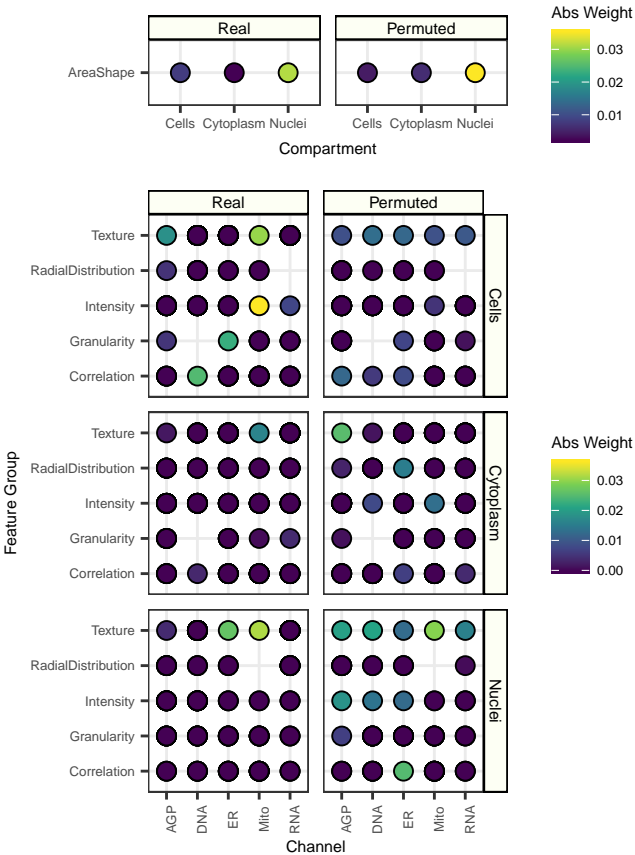
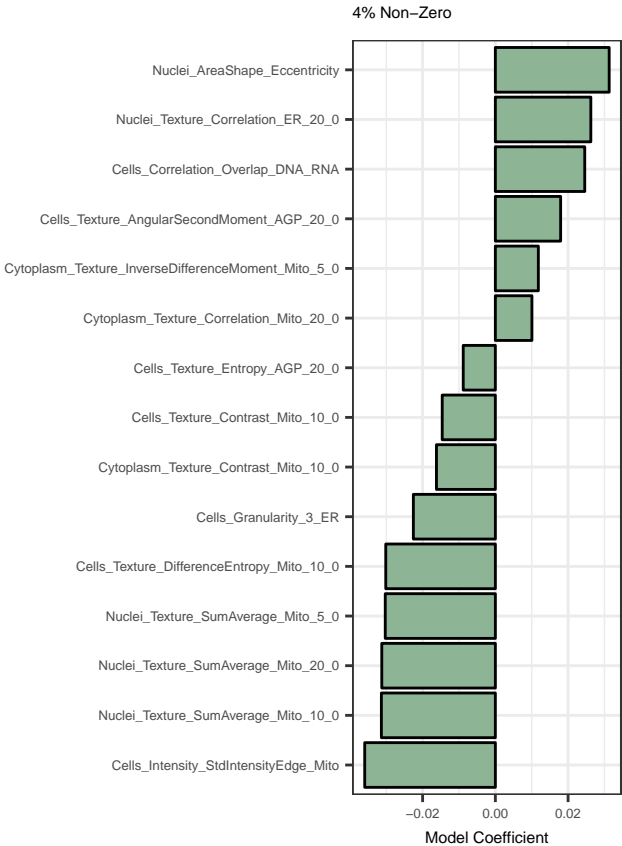


# Polynuclear – # of gH2AX Spots per Area of Nucleus

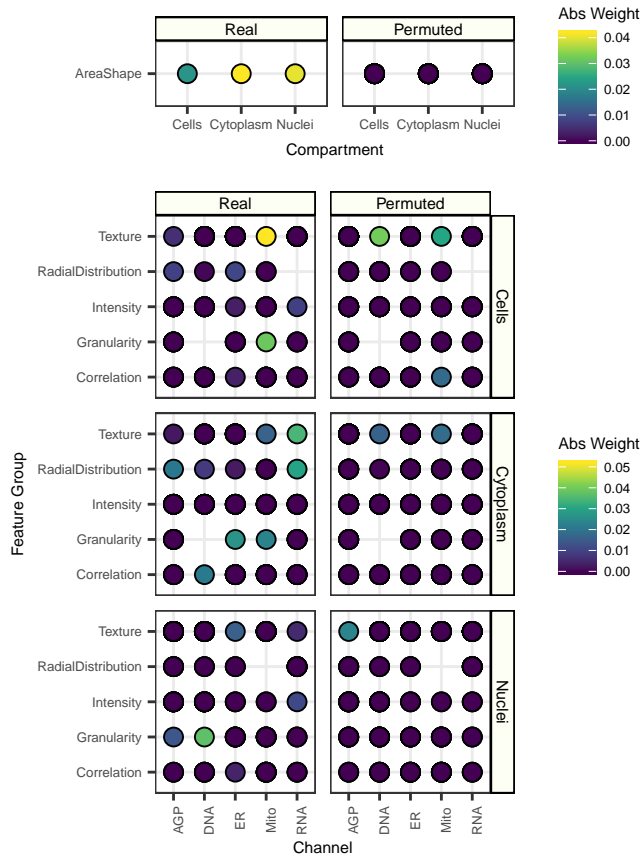
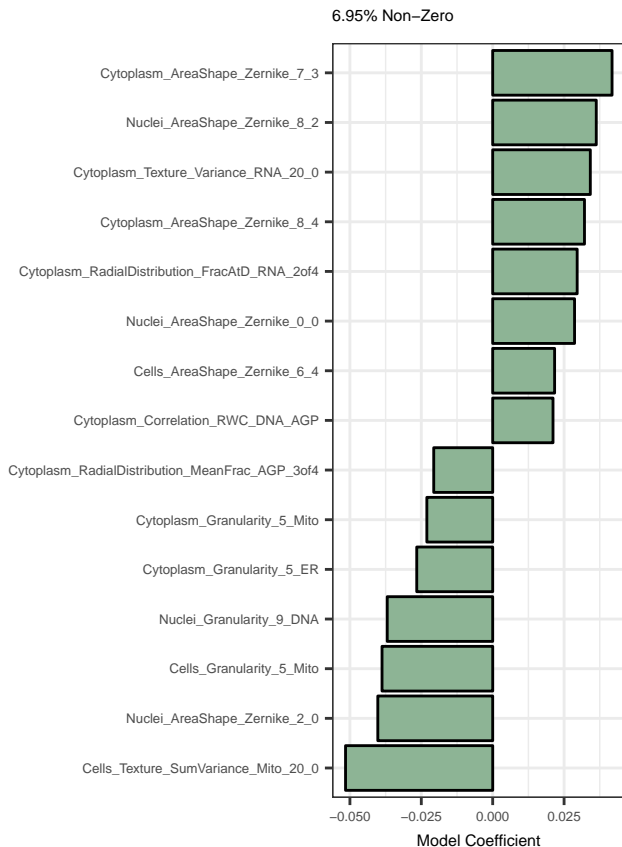
5.69% Non-Zero



# Polyploid – % High gH2AX Spots

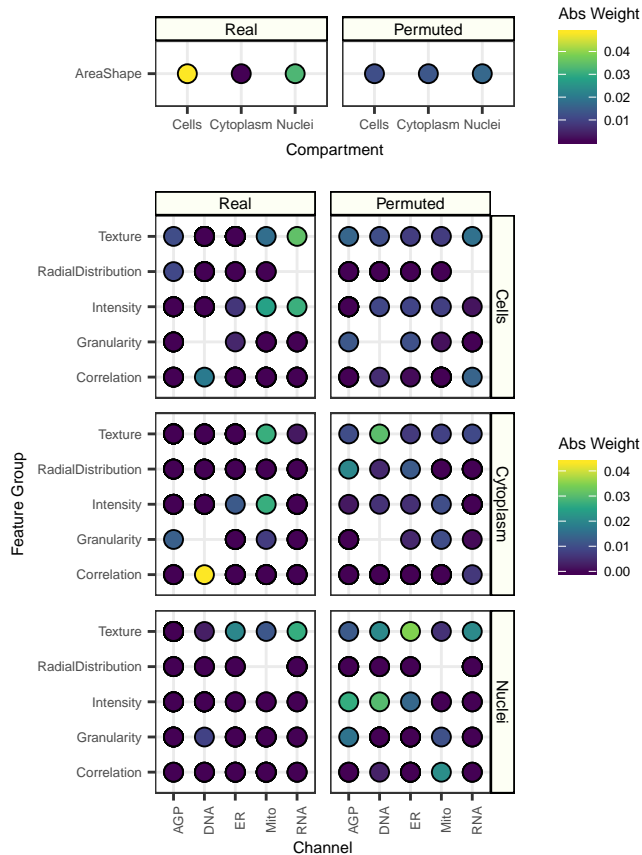


# Polyploid – # cells

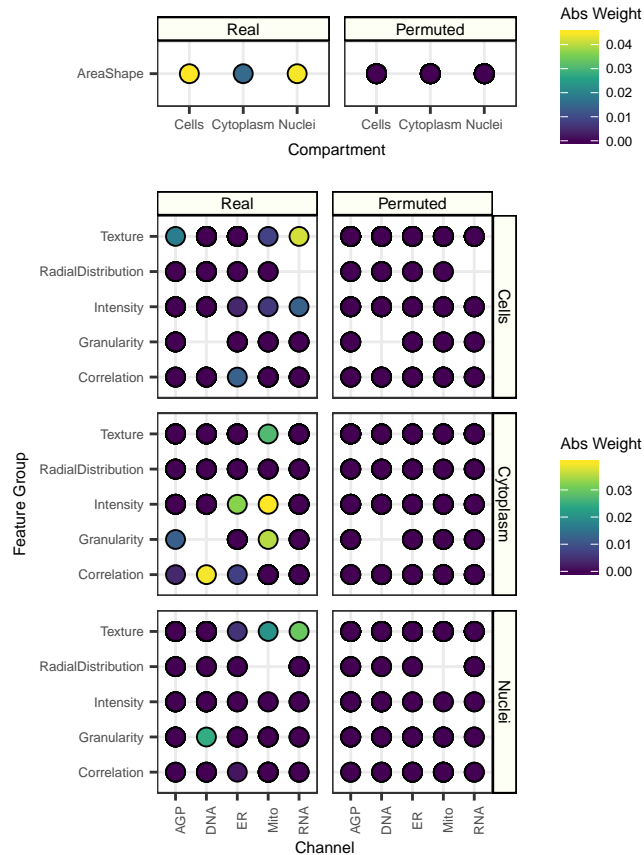




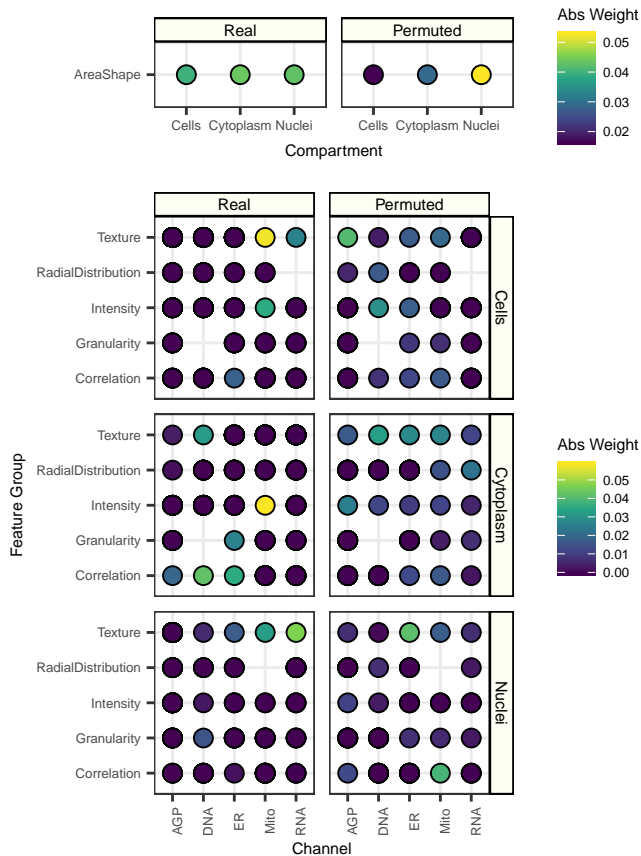
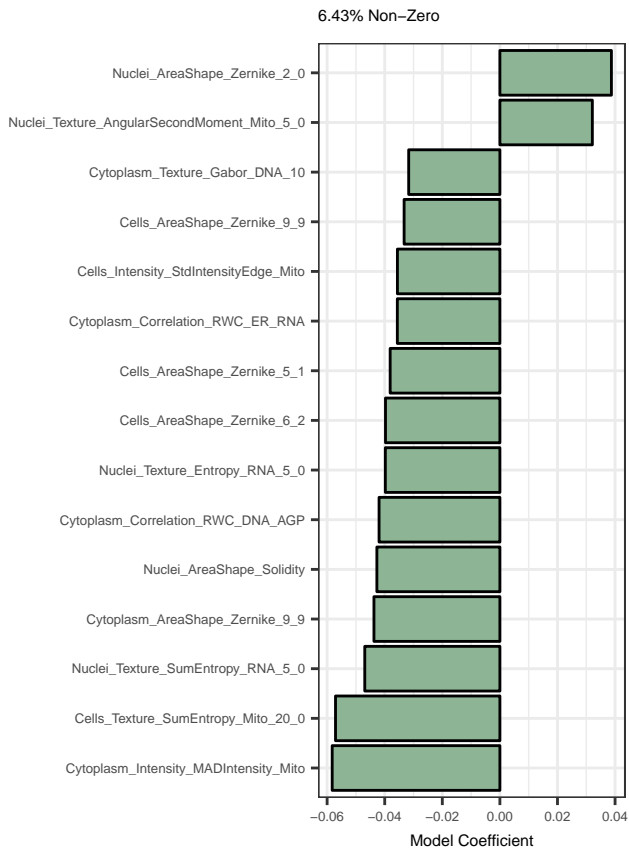
# Polyploid – # of gH2AX Spots



# Polyploid – # of gH2AX Spots per Area of Nucleus

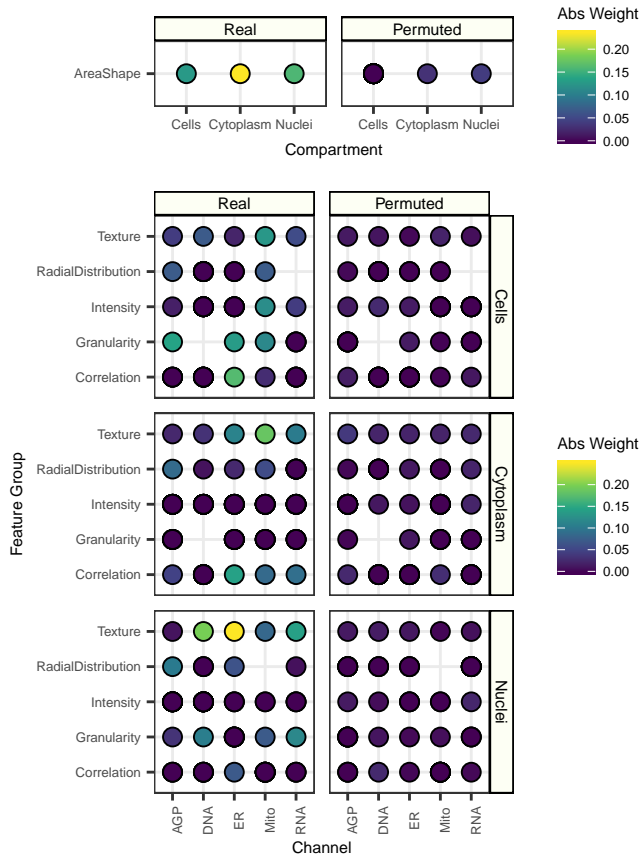
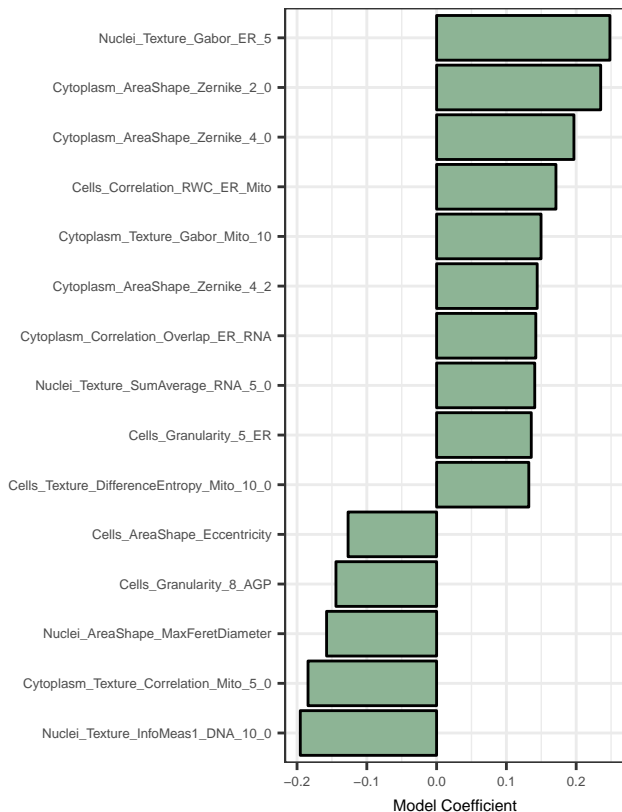


# S – % High gH2AX Spots



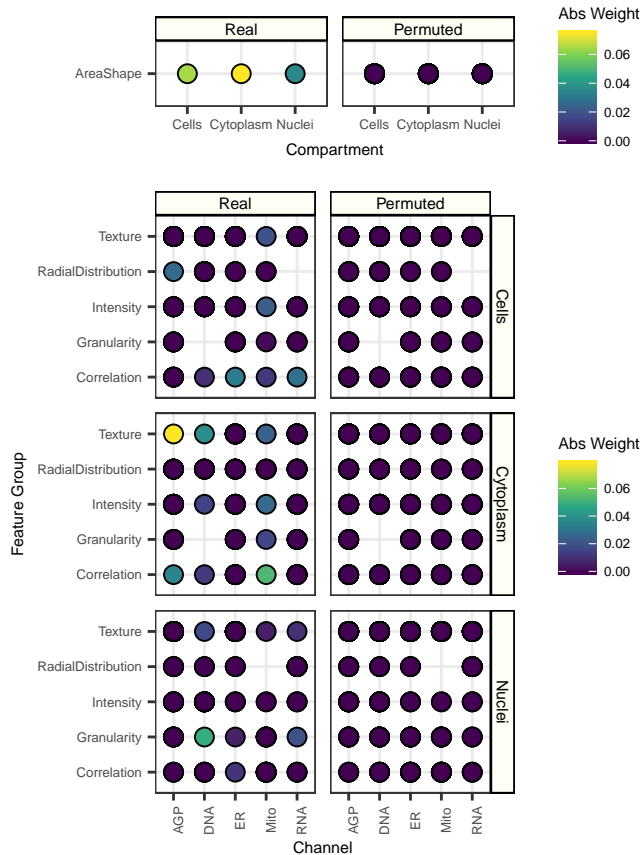
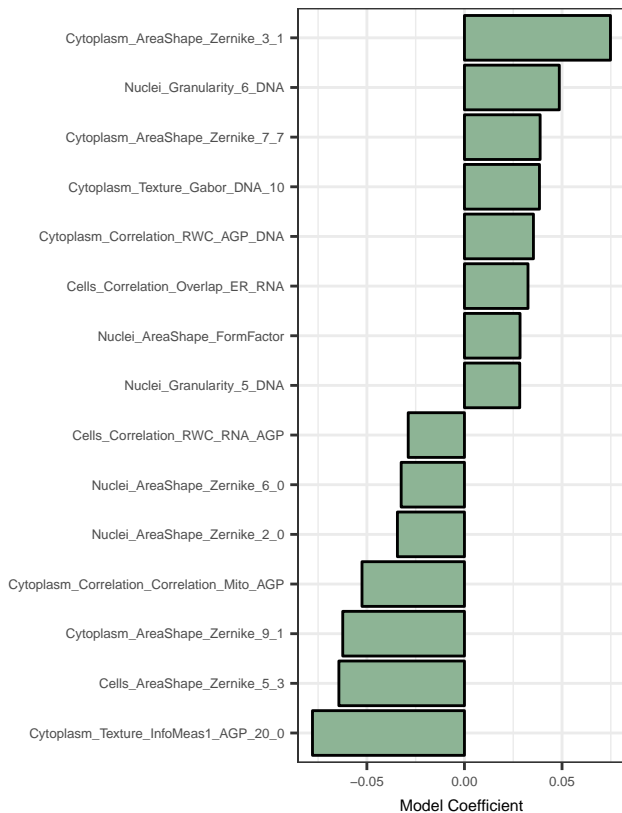
# S – Intensity Nucleus EdU Mean

11.28% Non-Zero

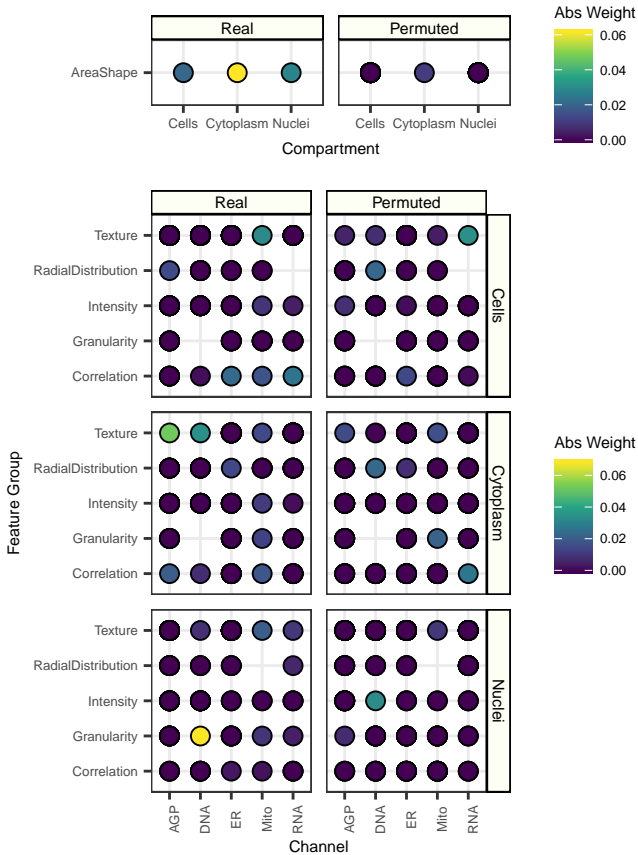
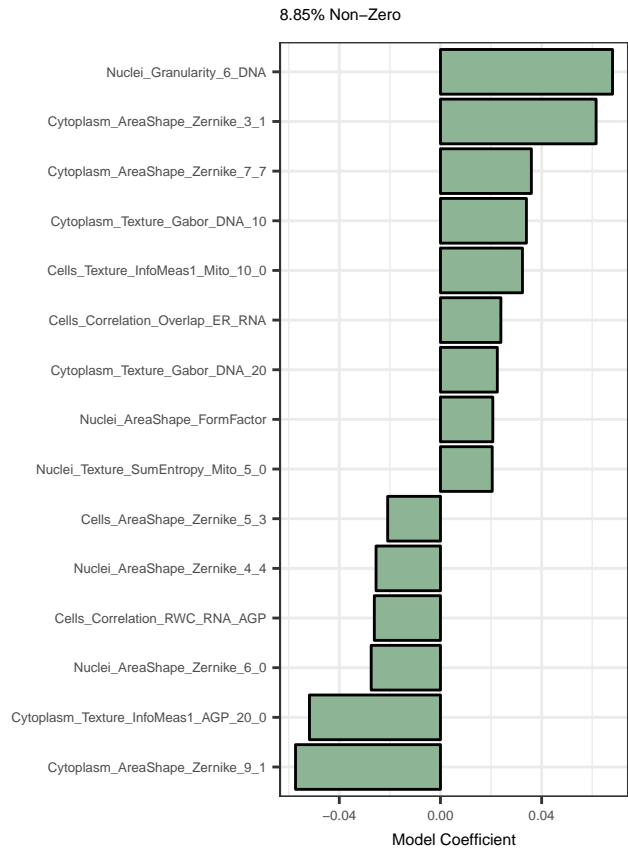


# S – Intensity Nucleus EdU Sum

7.9% Non-Zero

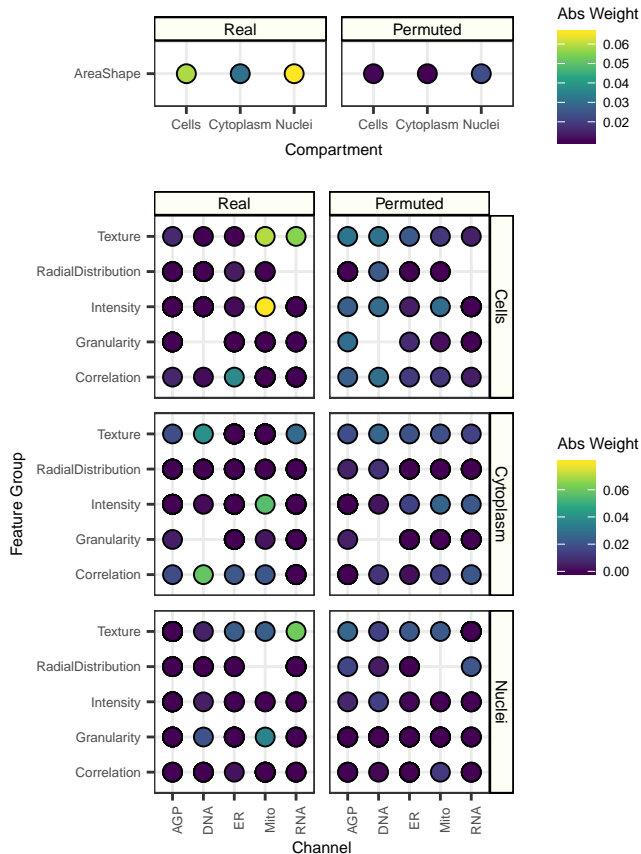
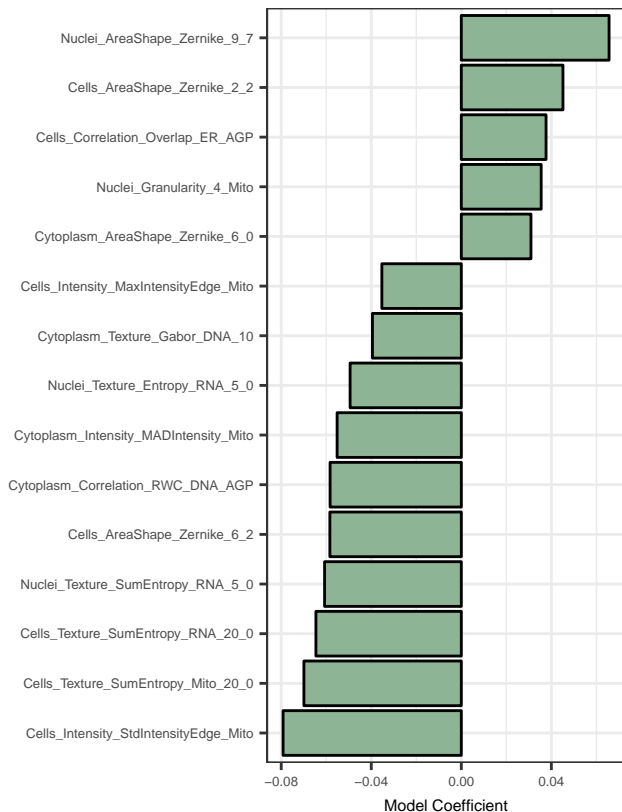


# S - # cells

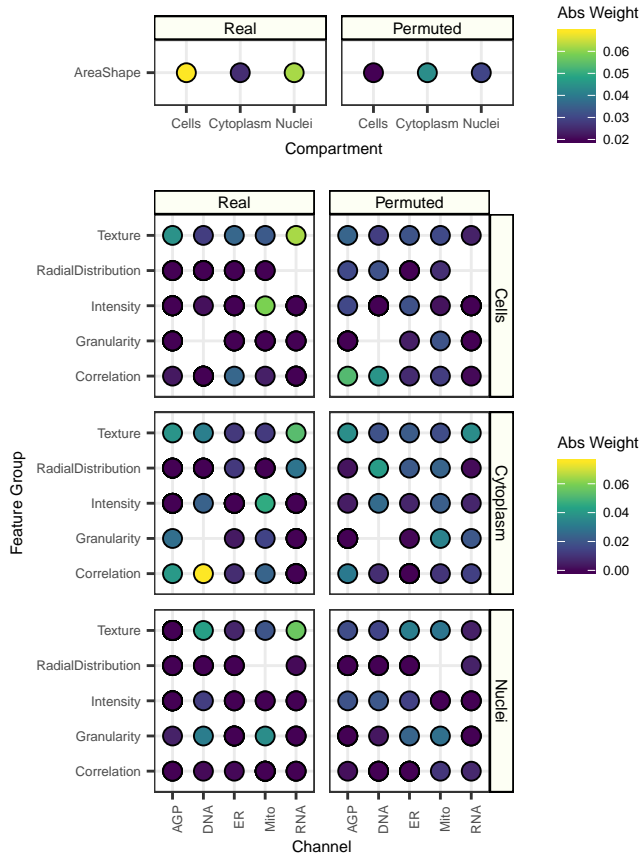
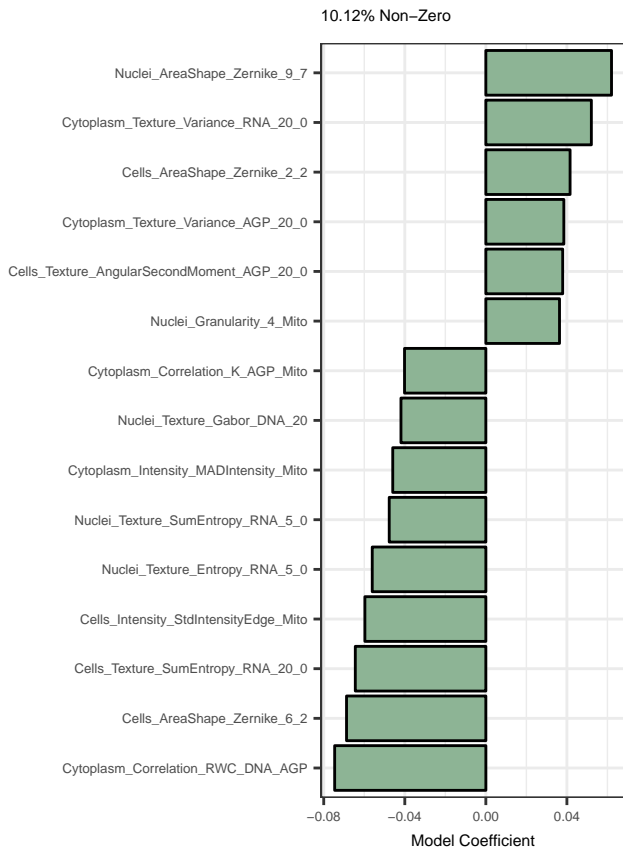


# S - # of gH2AX Spots

7.9% Non-Zero

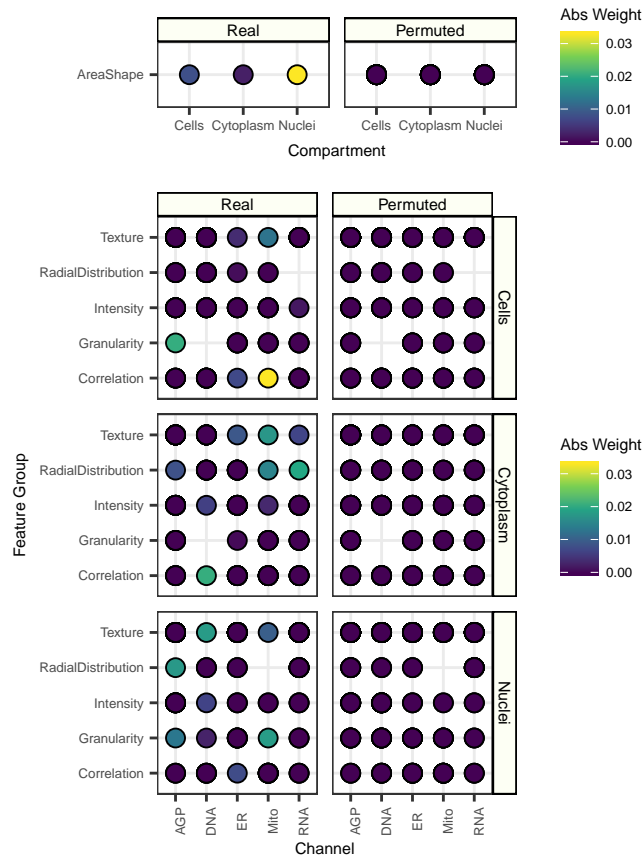
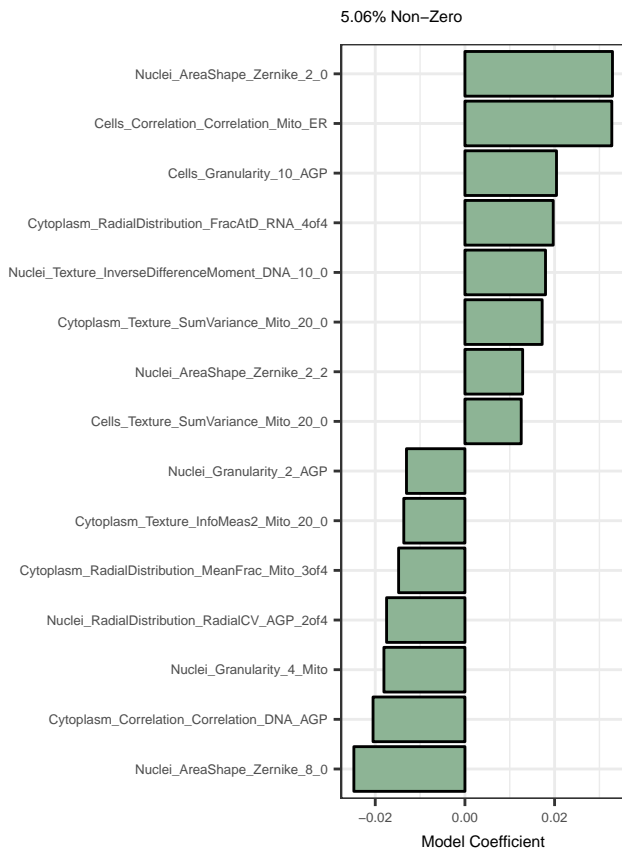


# S – # of gH2AX Spots per Area of Nucleus

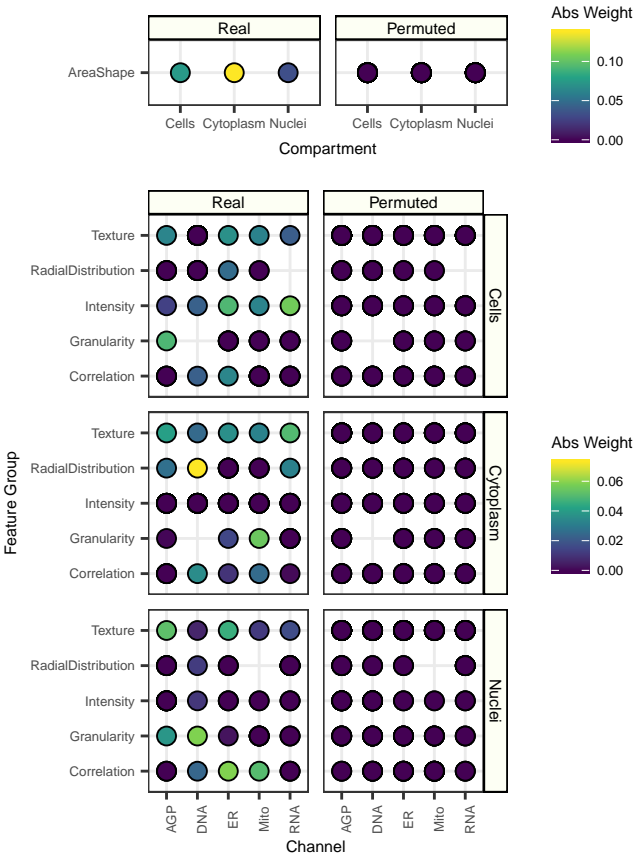
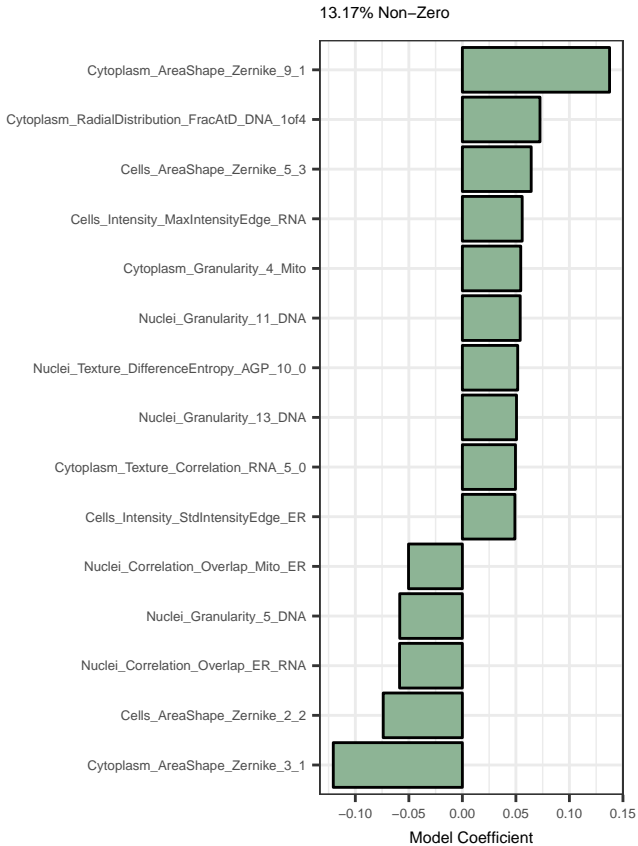




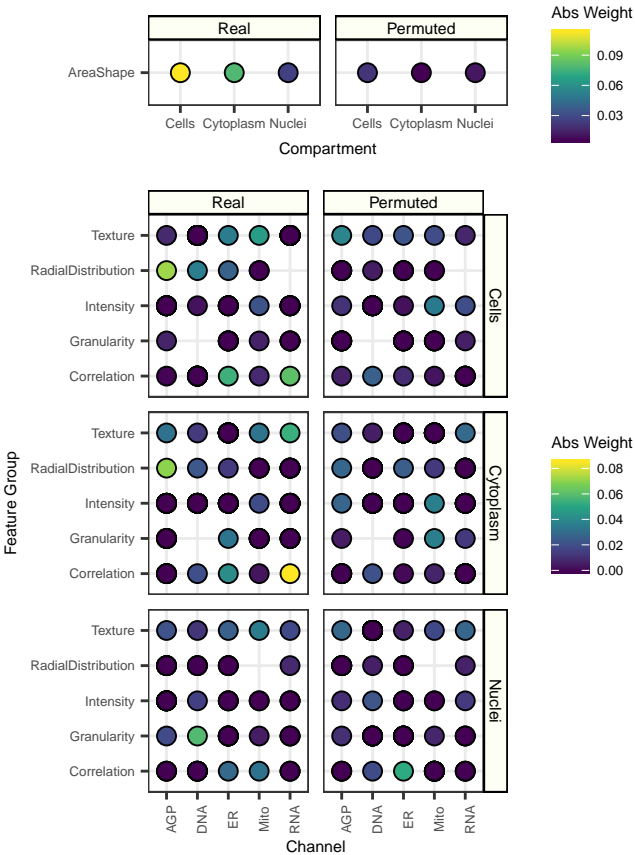
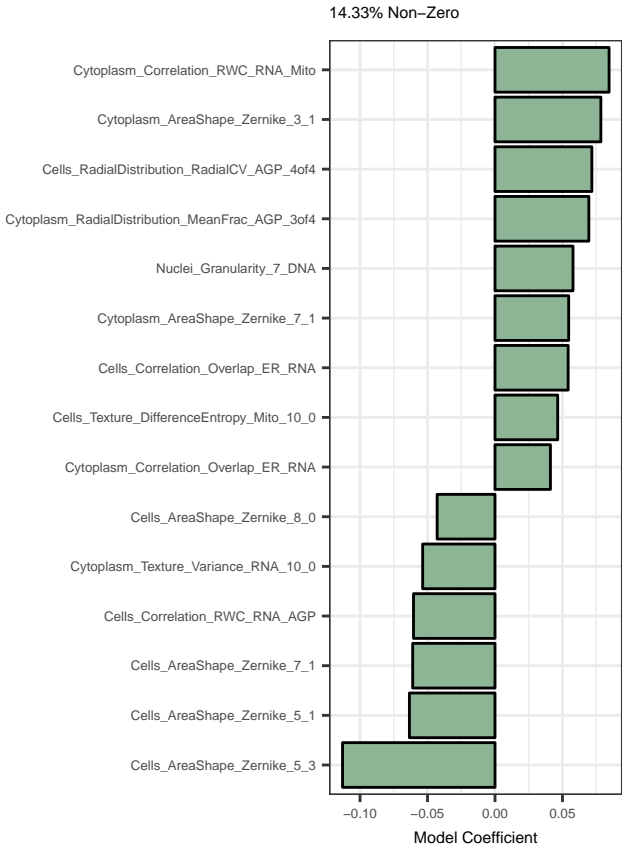
# % Inf



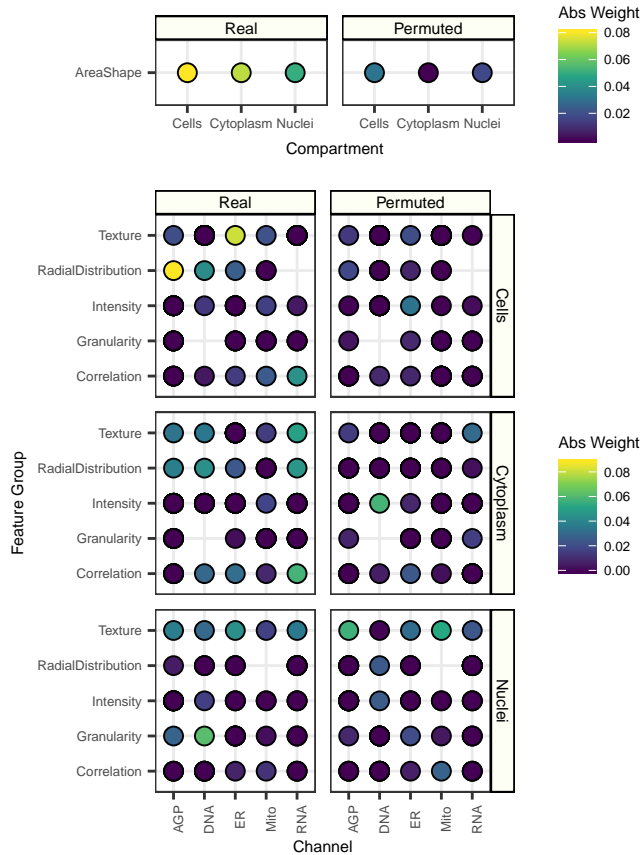
# Live Cell Area



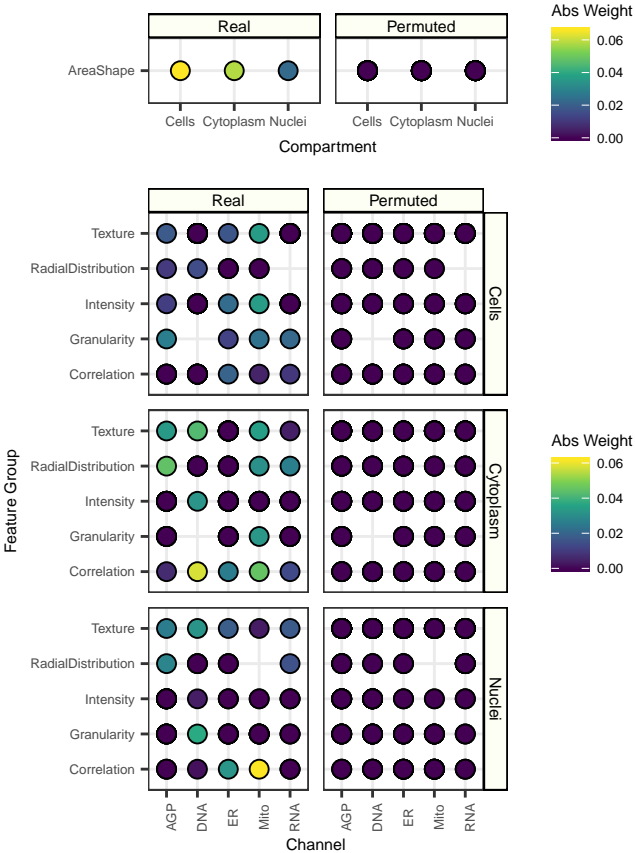
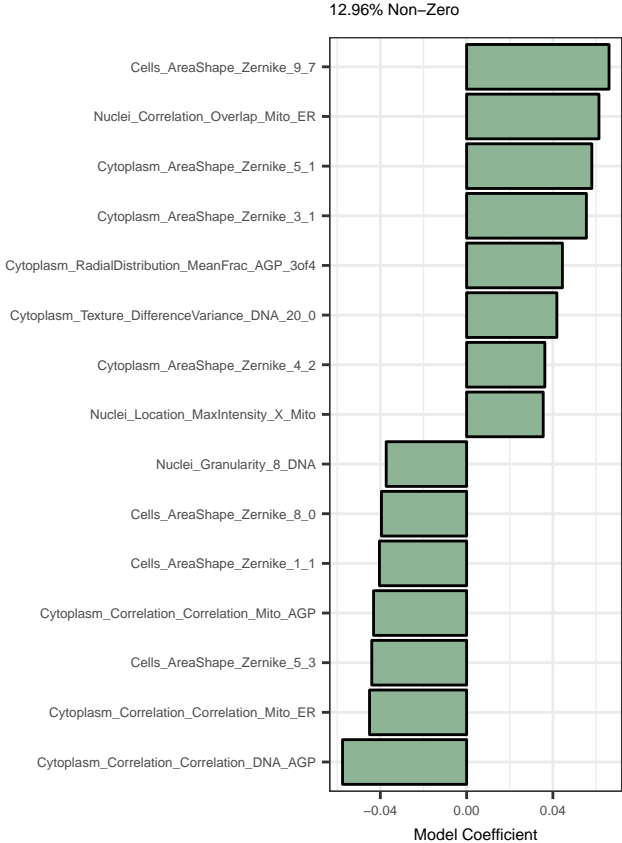
# Live Cell Roundness



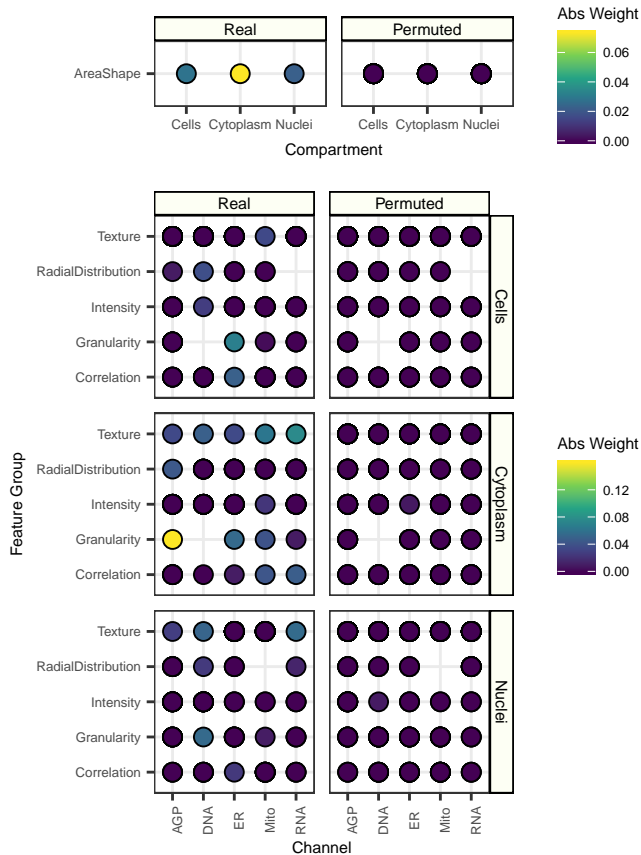
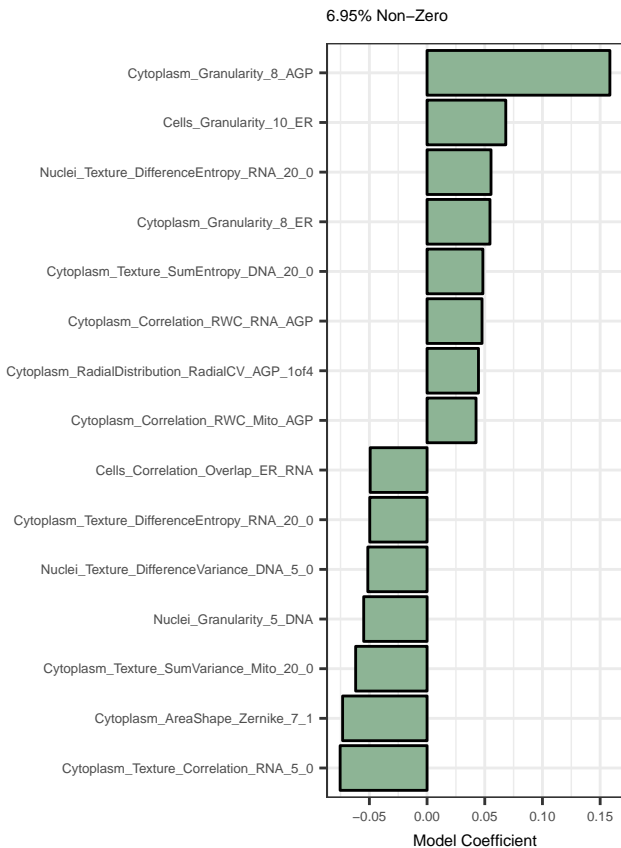
# Live Width / Length



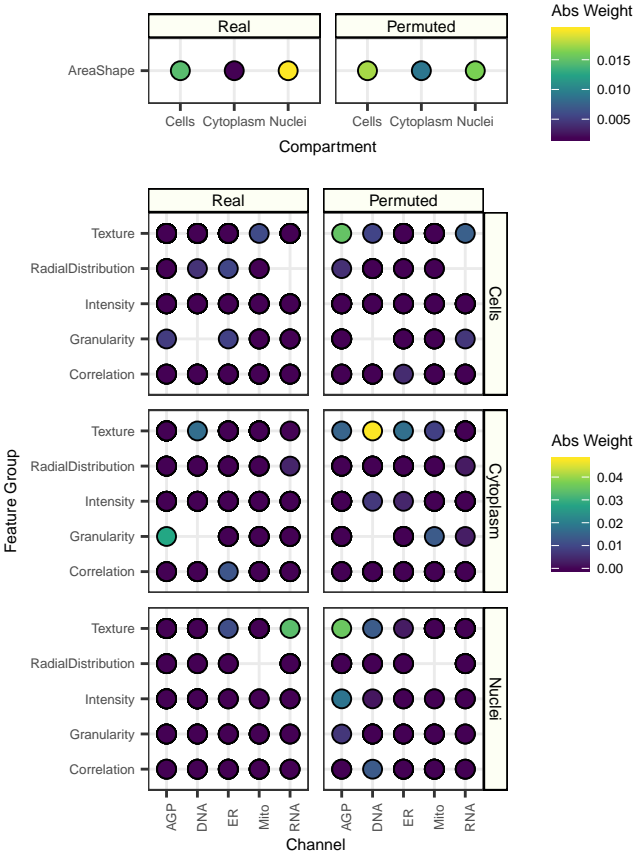
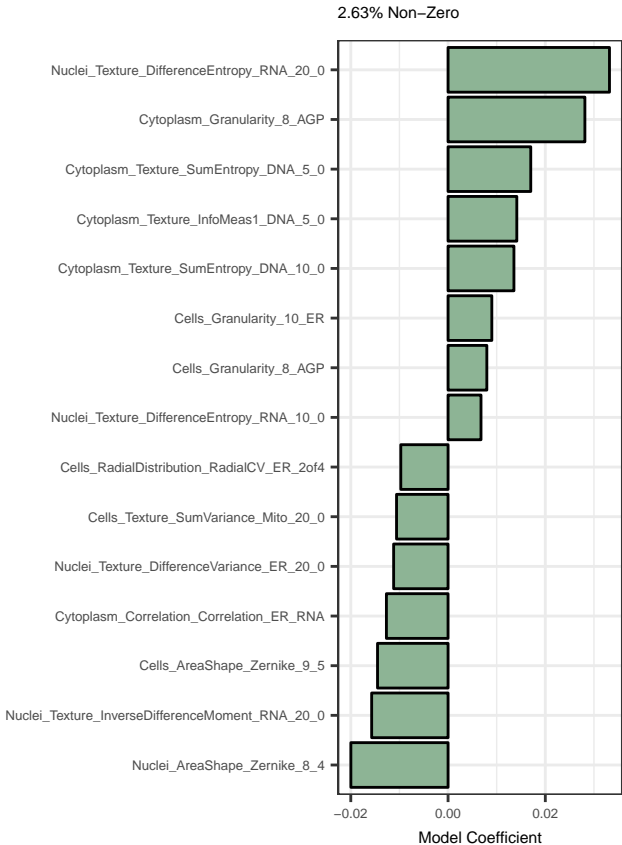
# # Live Cells



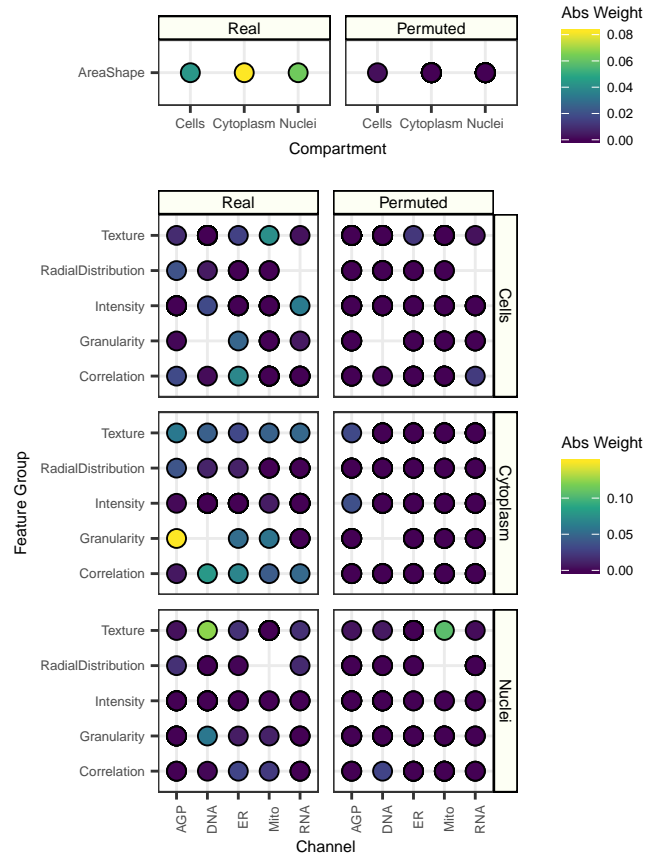
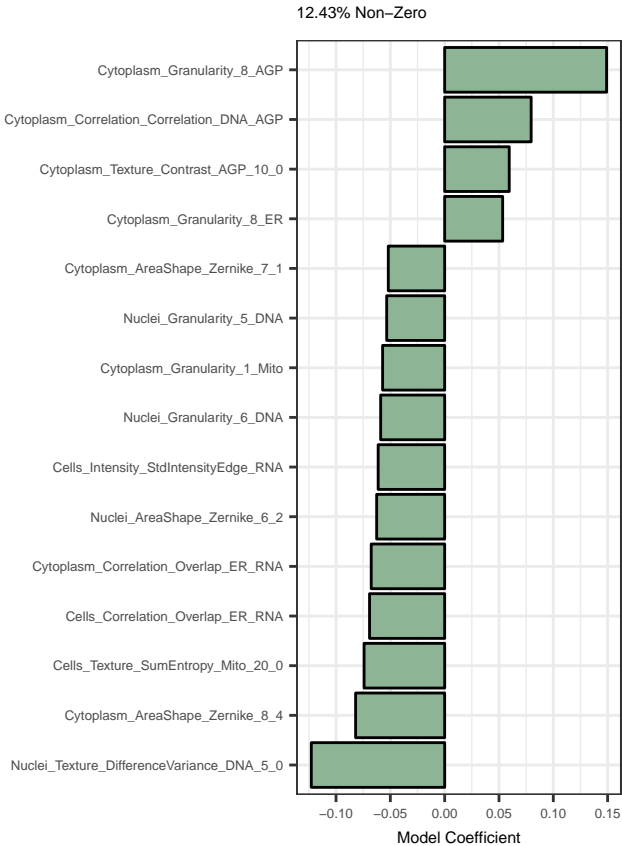
# % All Apoptosis (CASP+)



# % Caspase / % Dead Only



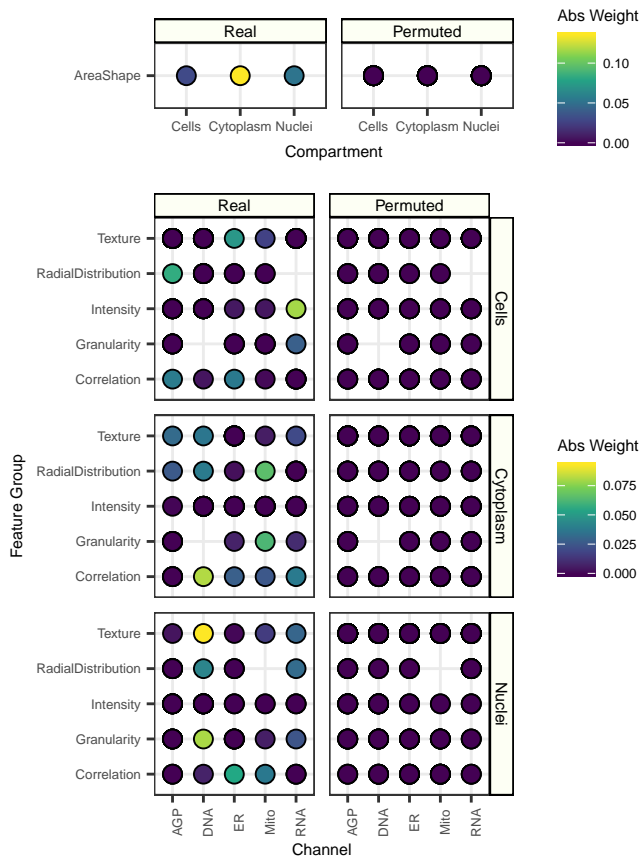
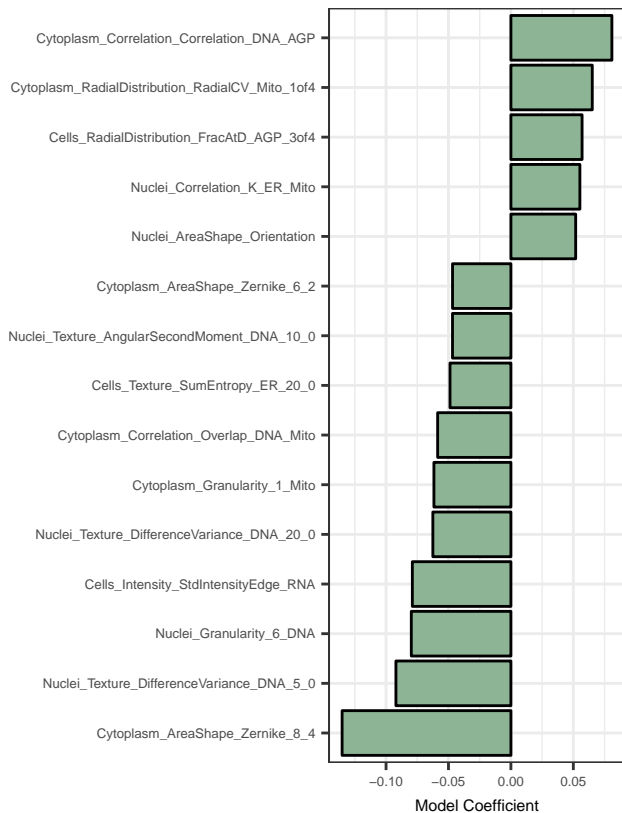
# % All Dead (DRAQ7+)





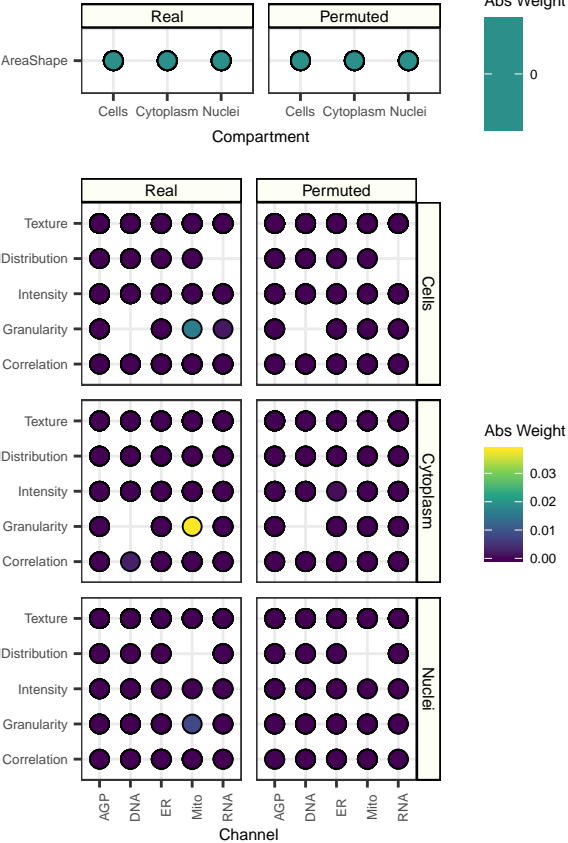
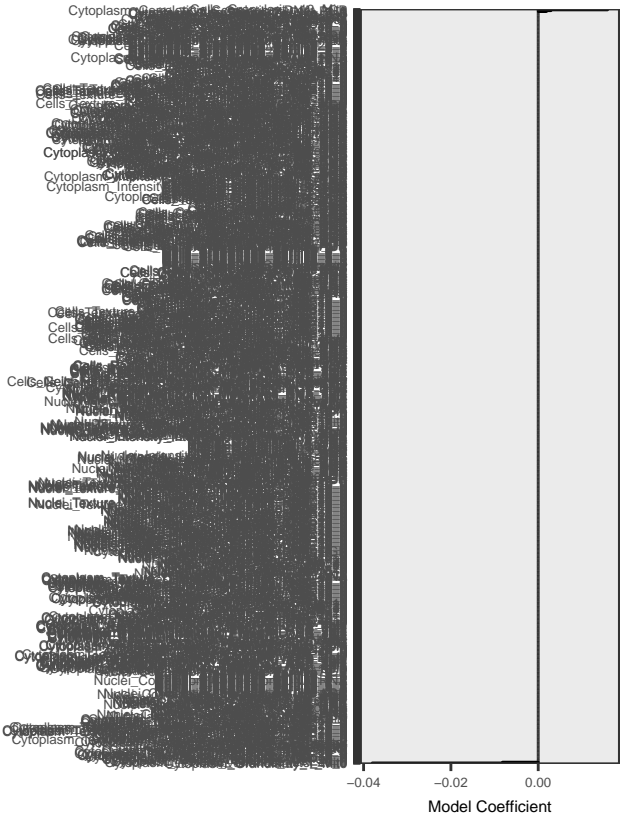
# % Dead Only (CASP-; DRAQ7+)

11.59% Non-Zero



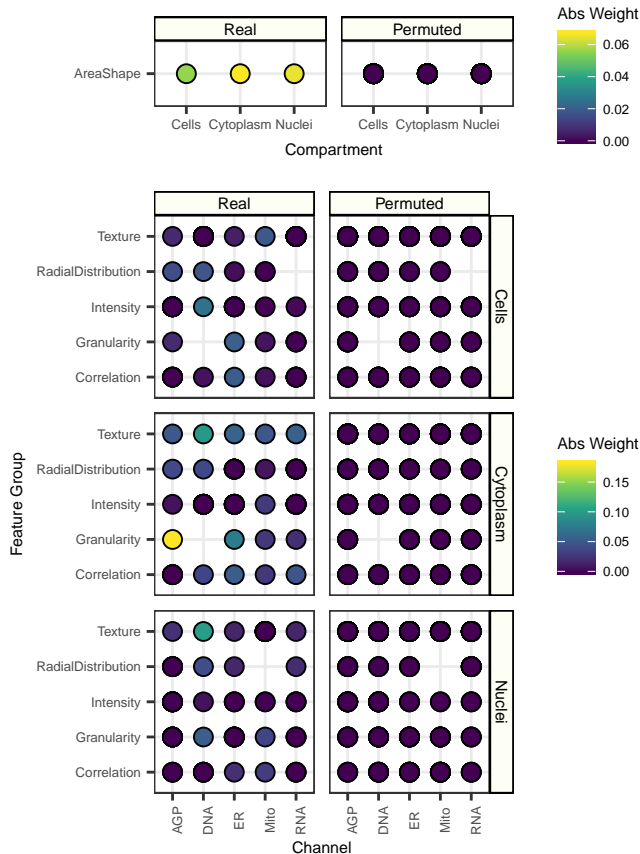
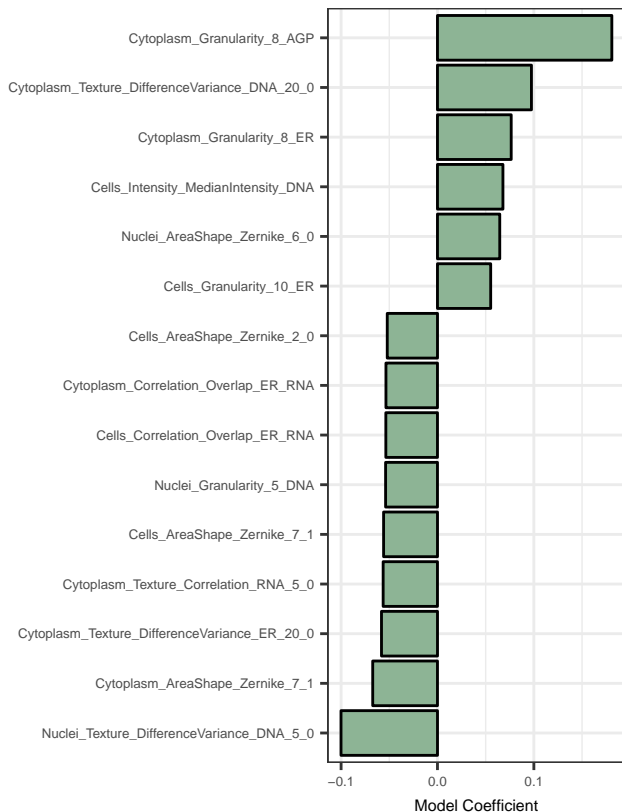
# % Early Apoptosis (CASP+; DRAQ7-)

0.53% Non-Zero



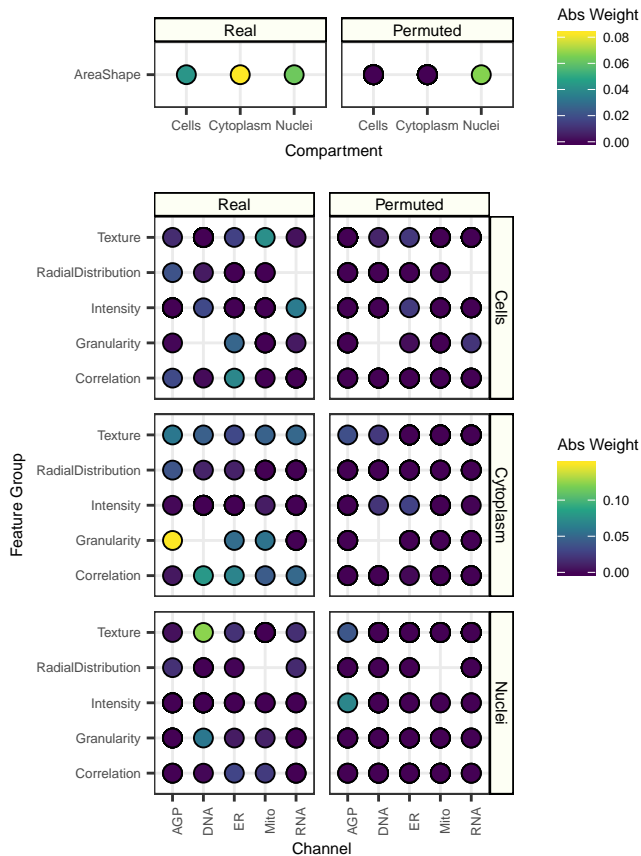
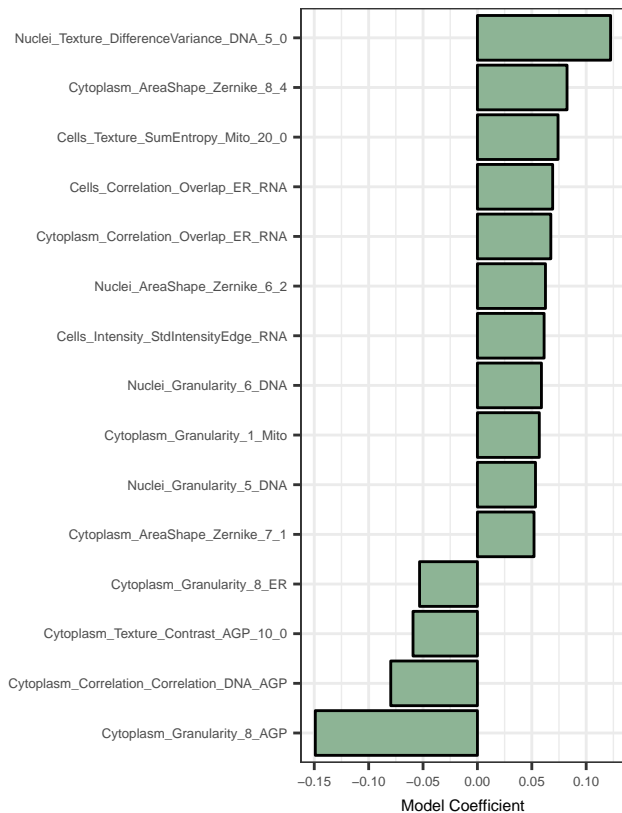
# % Late Apoptosis (CASP+; DRAQ7+)

13.28% Non-Zero

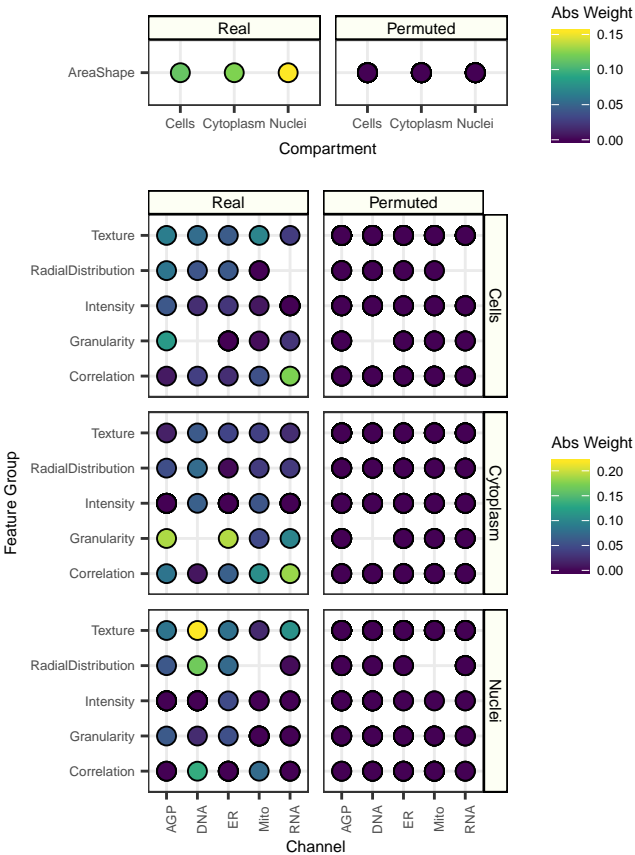
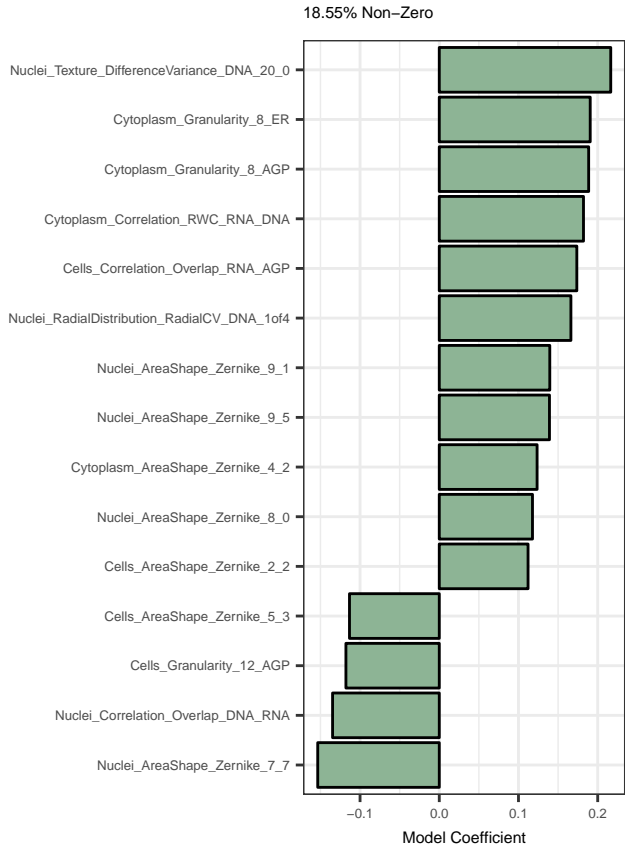


# % Live (DRAQ7-; CASP-)

12.54% Non-Zero



# ROS-back



# ROS

10.22% Non-Zero

