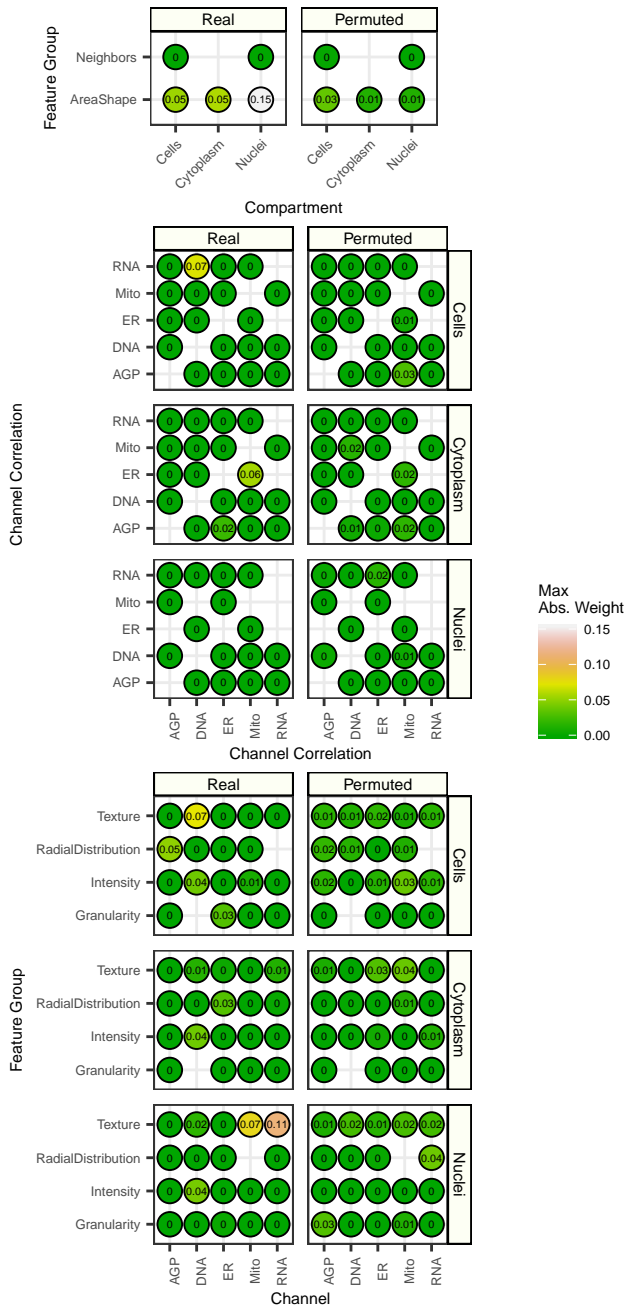
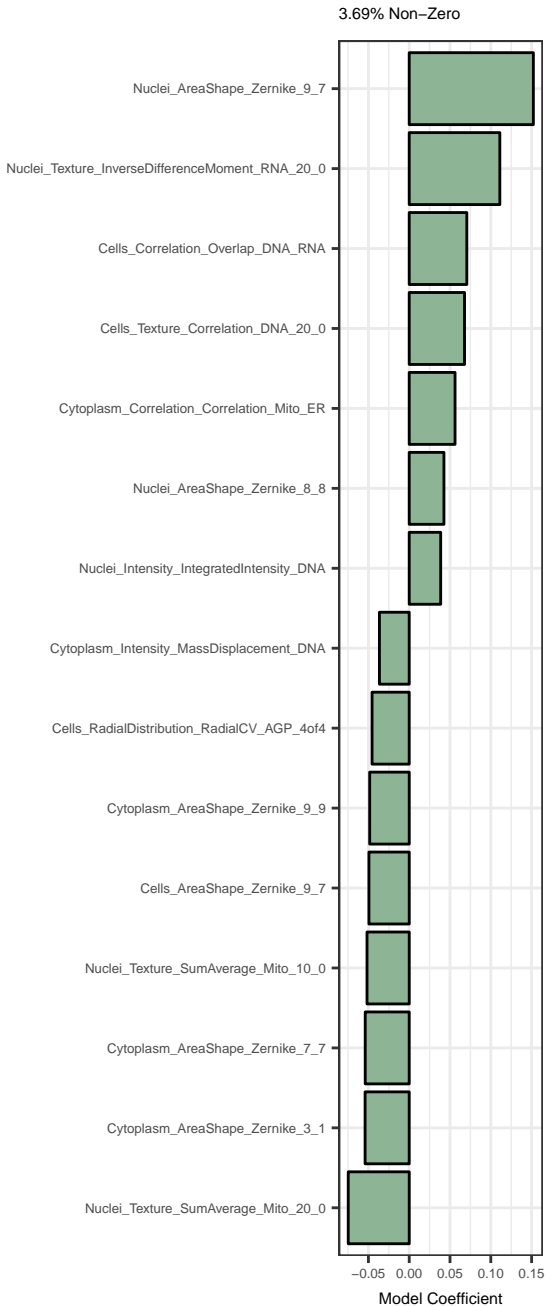
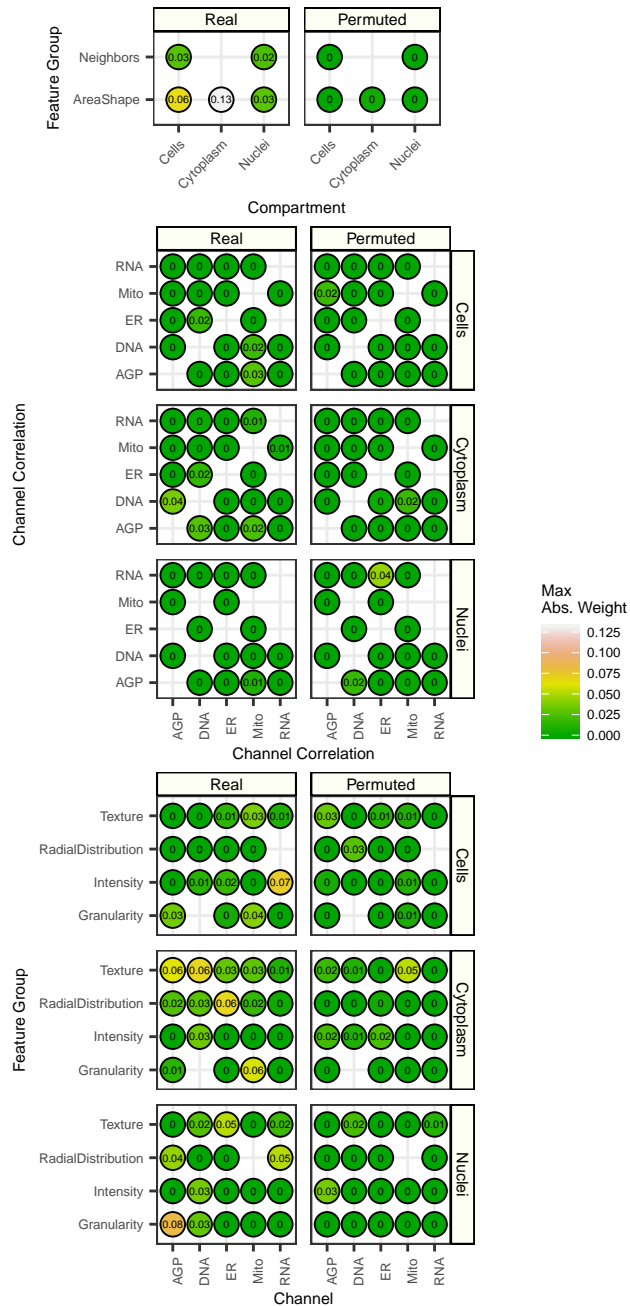
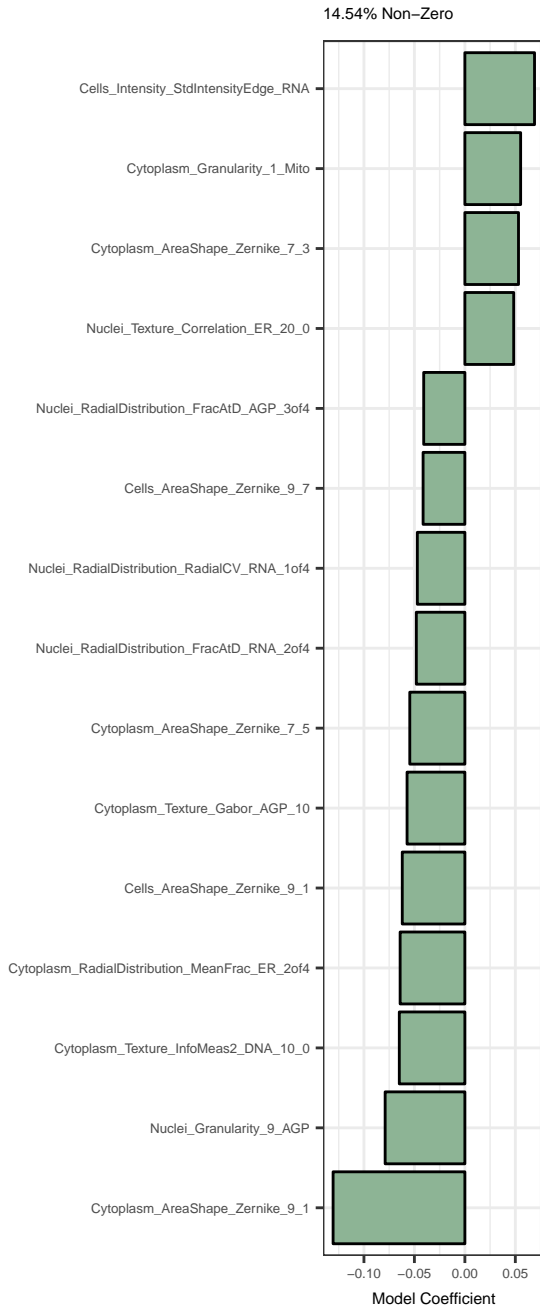


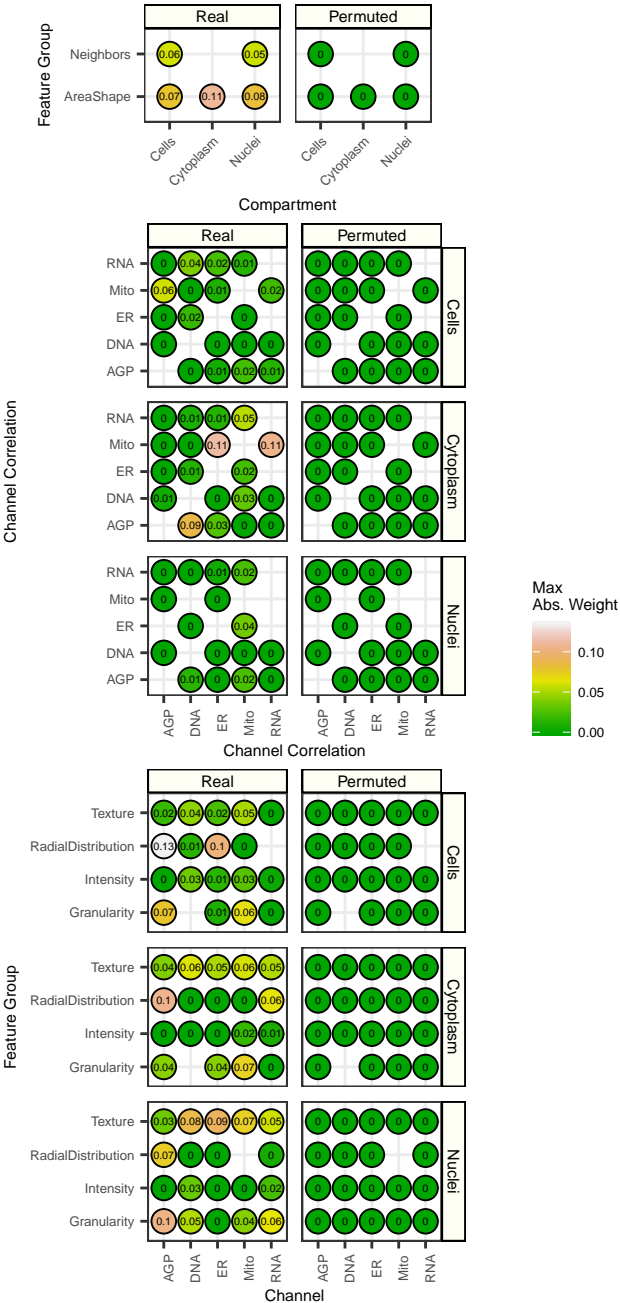
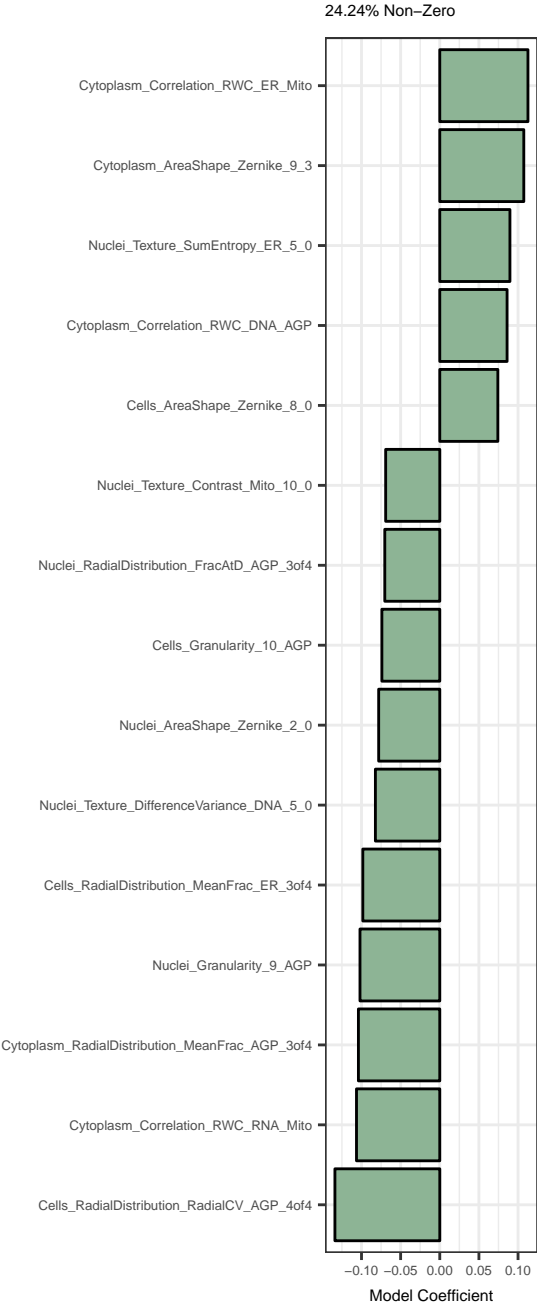
ALL – % High gH2AX Spots



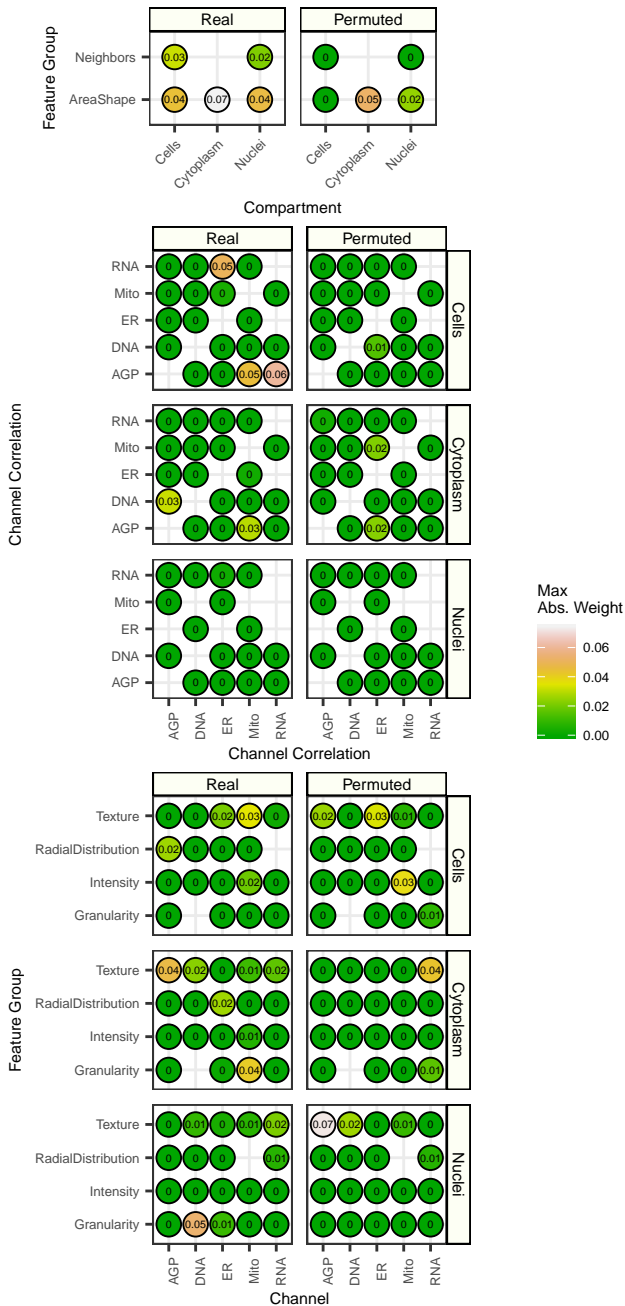
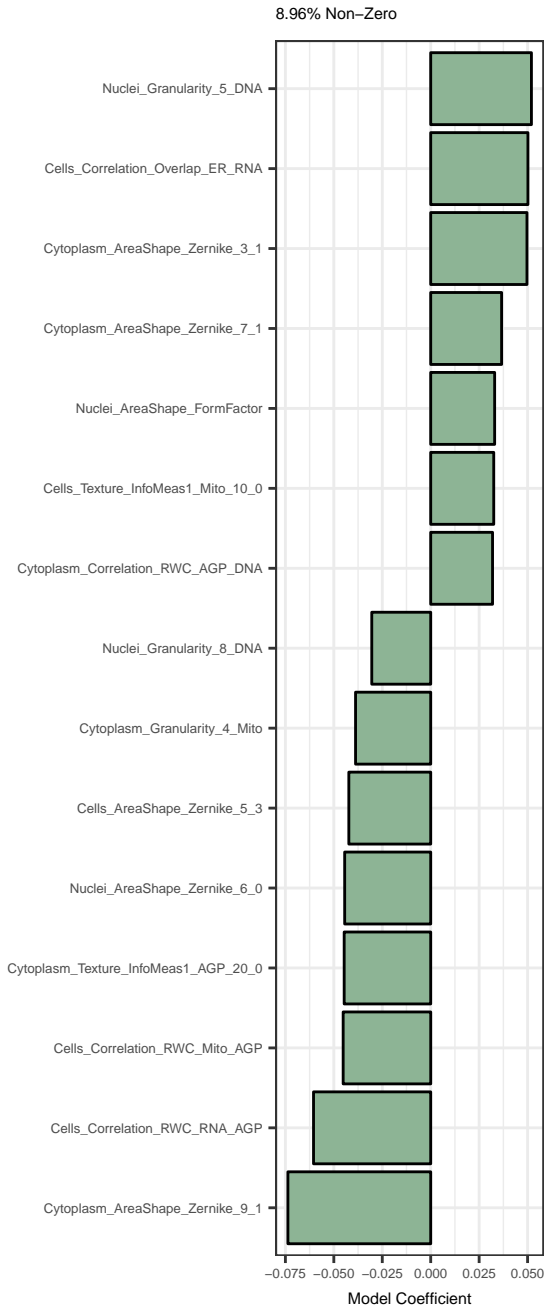
ALL – Polynuclear



ALL – Polyploid

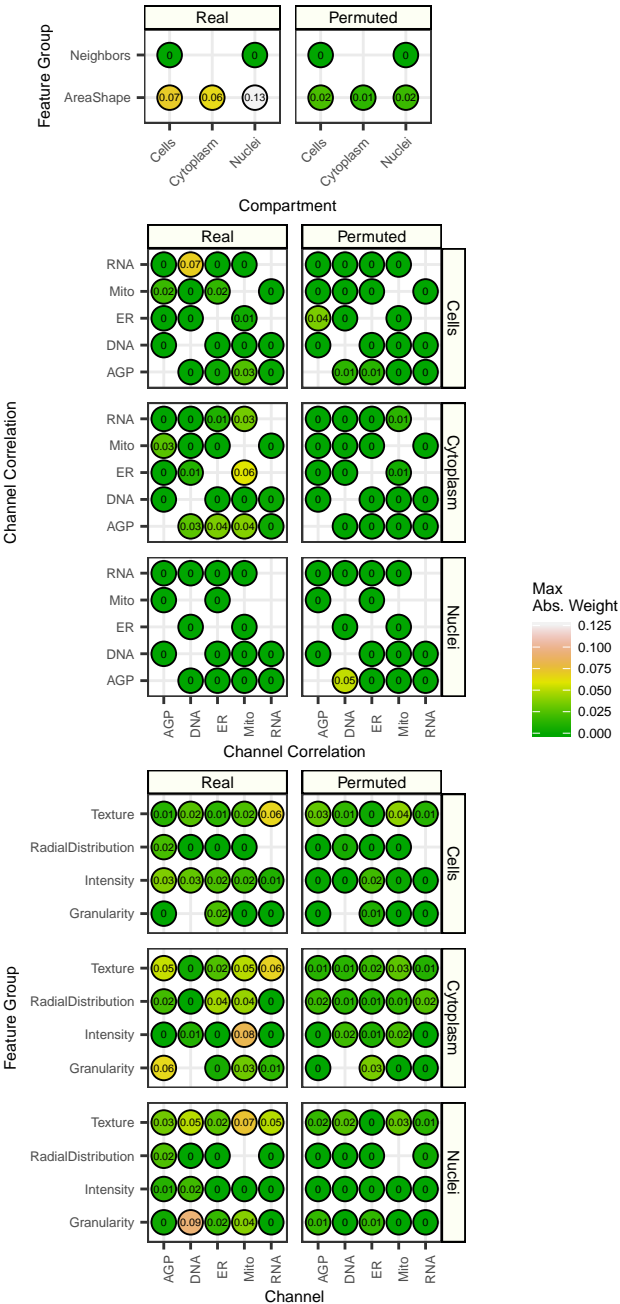
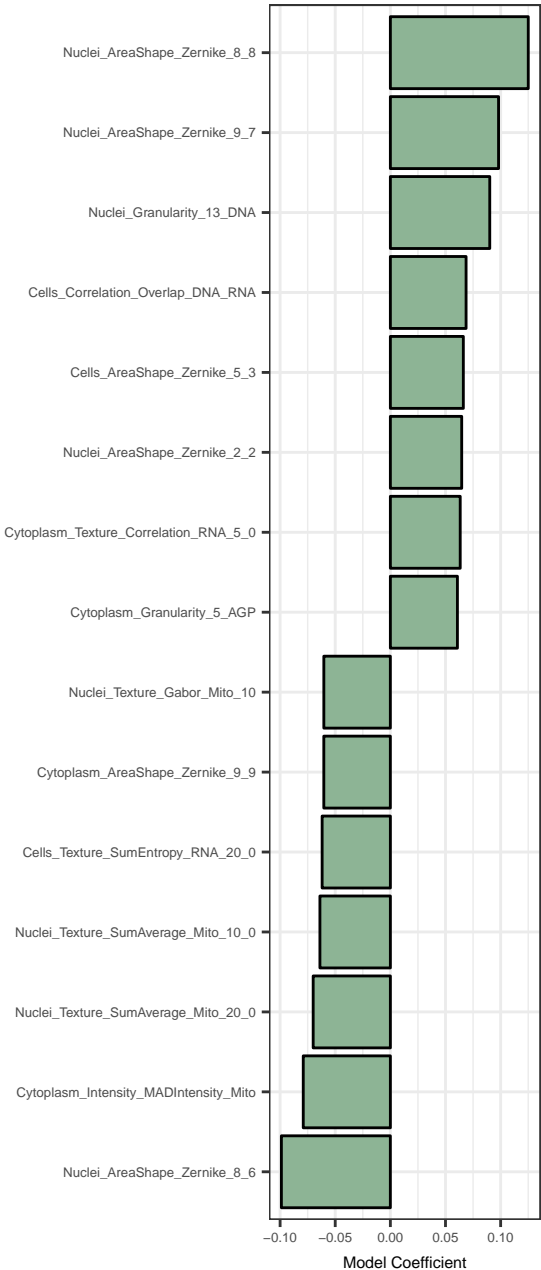


ALL – # cells



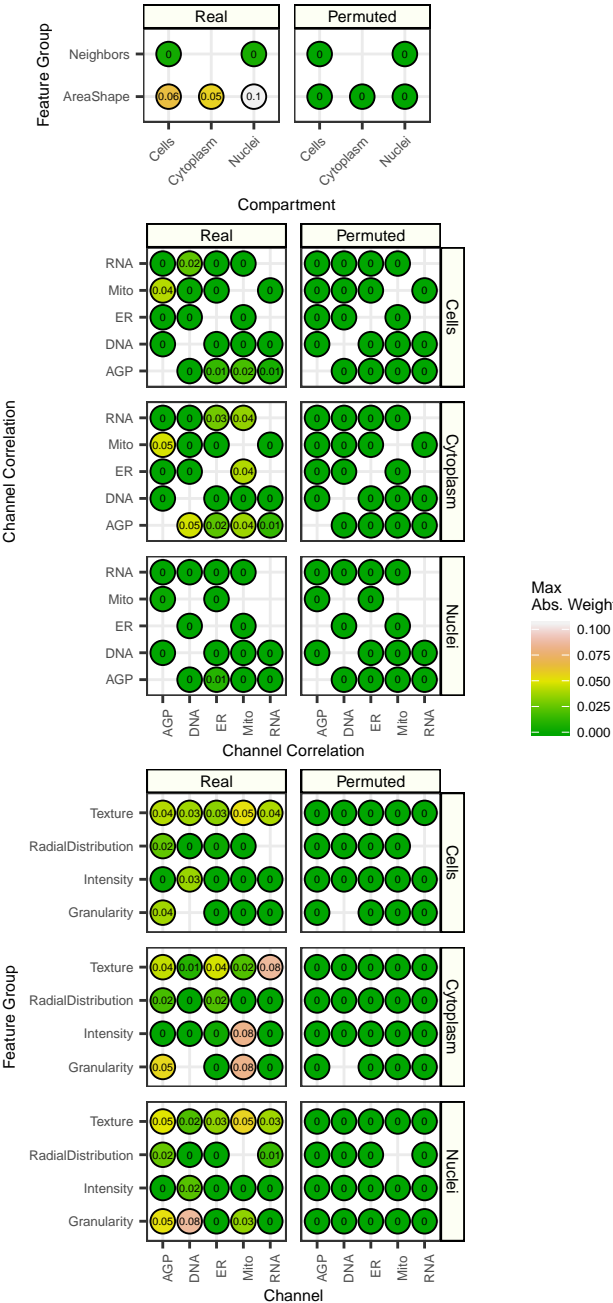
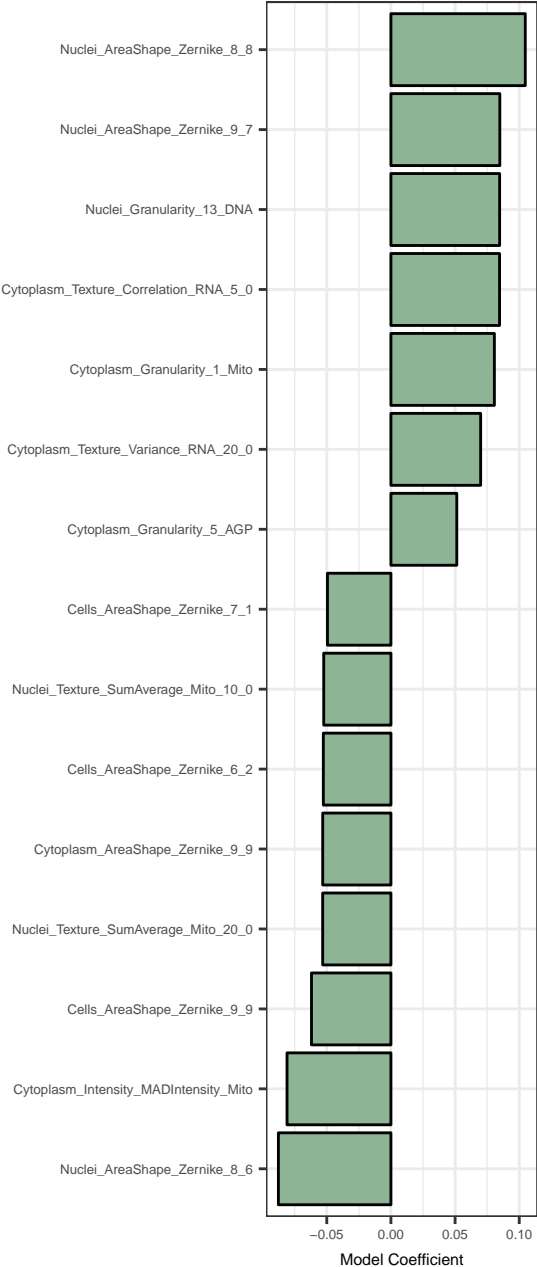
ALL – # of gH2AX Spots

17.18% Non-Zero

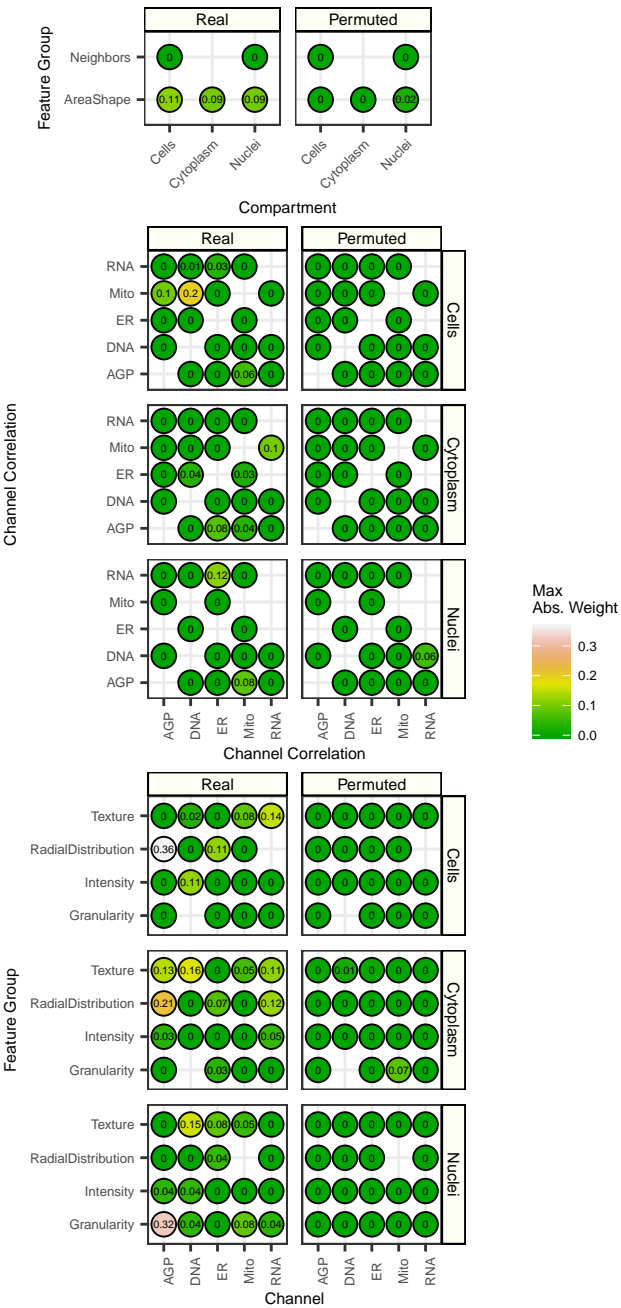
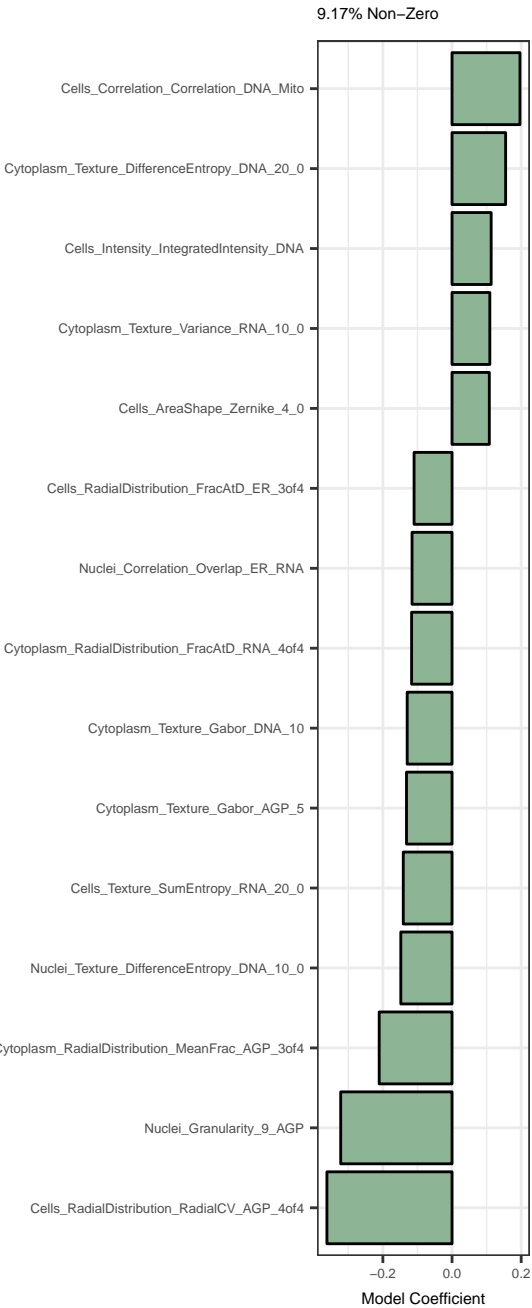


ALL – # of gH2AX Spots per Area of Nucleus

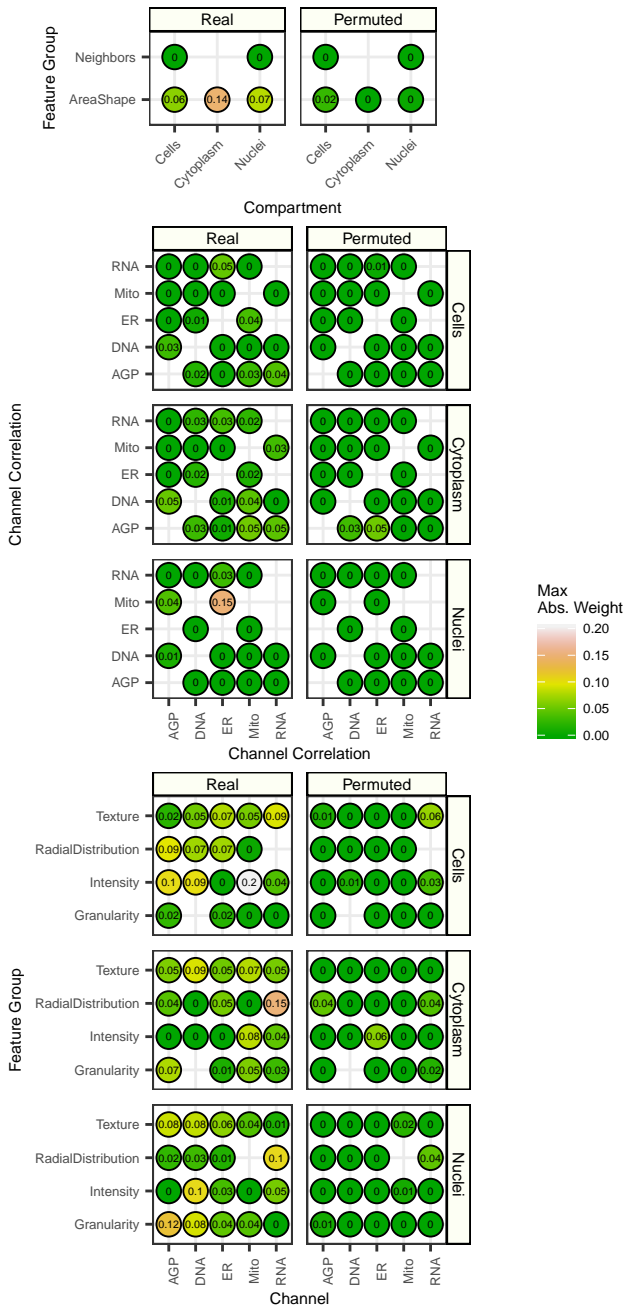
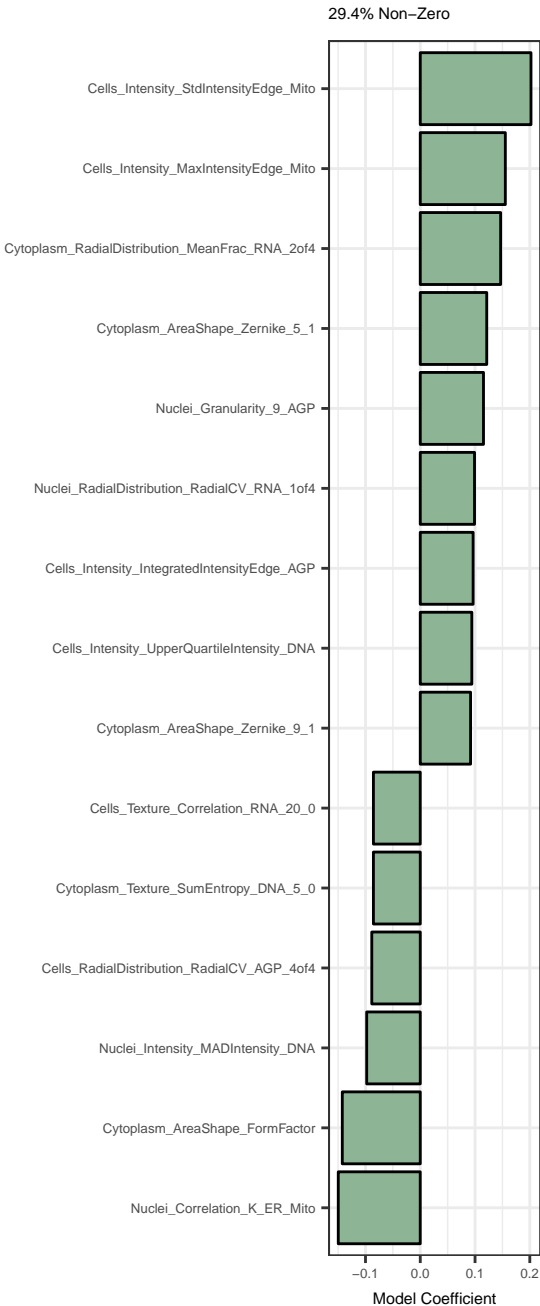
17.39% Non-Zero



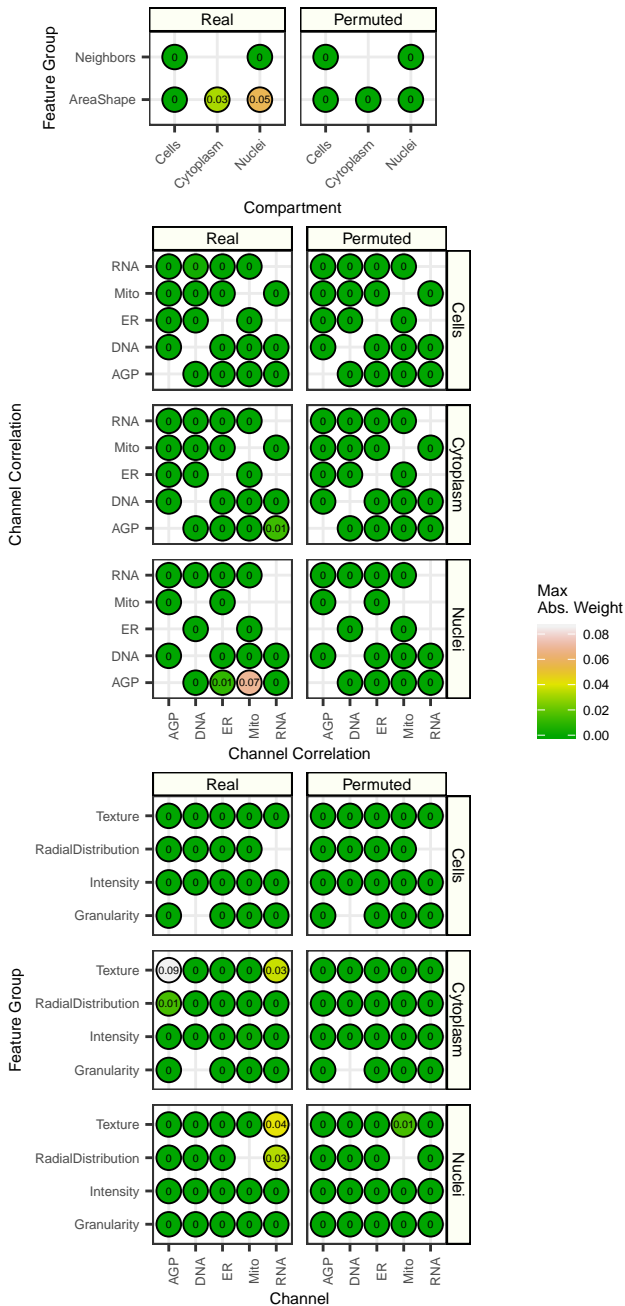
ALL – Nucleus Area



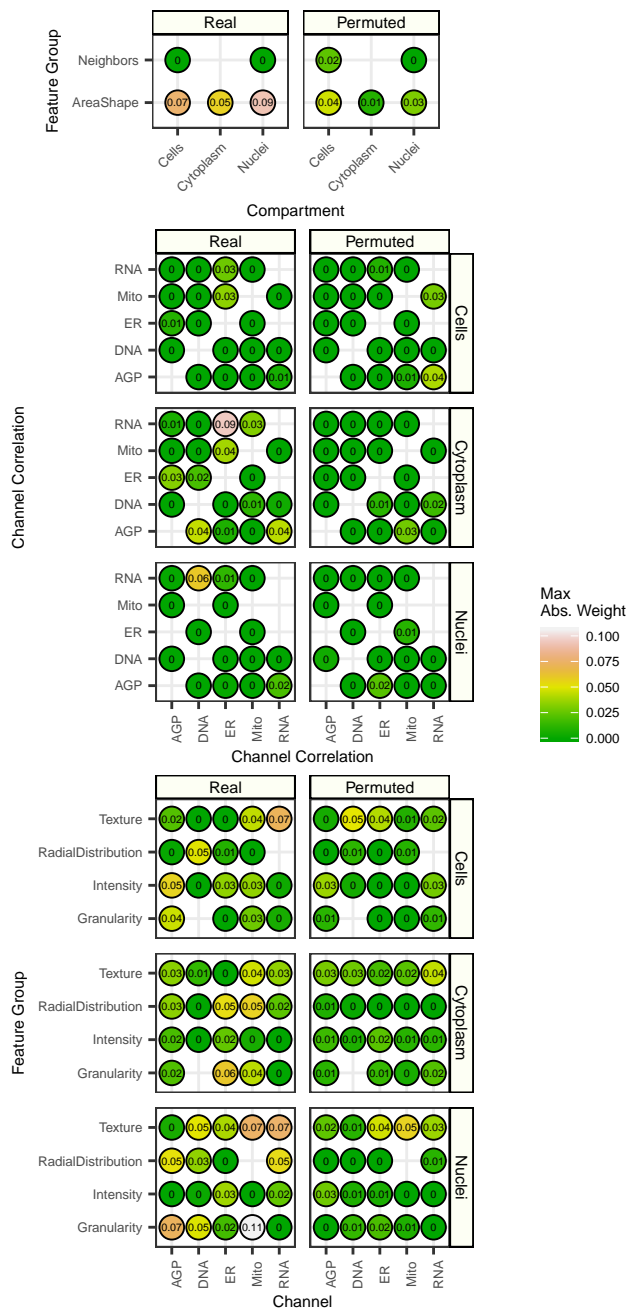
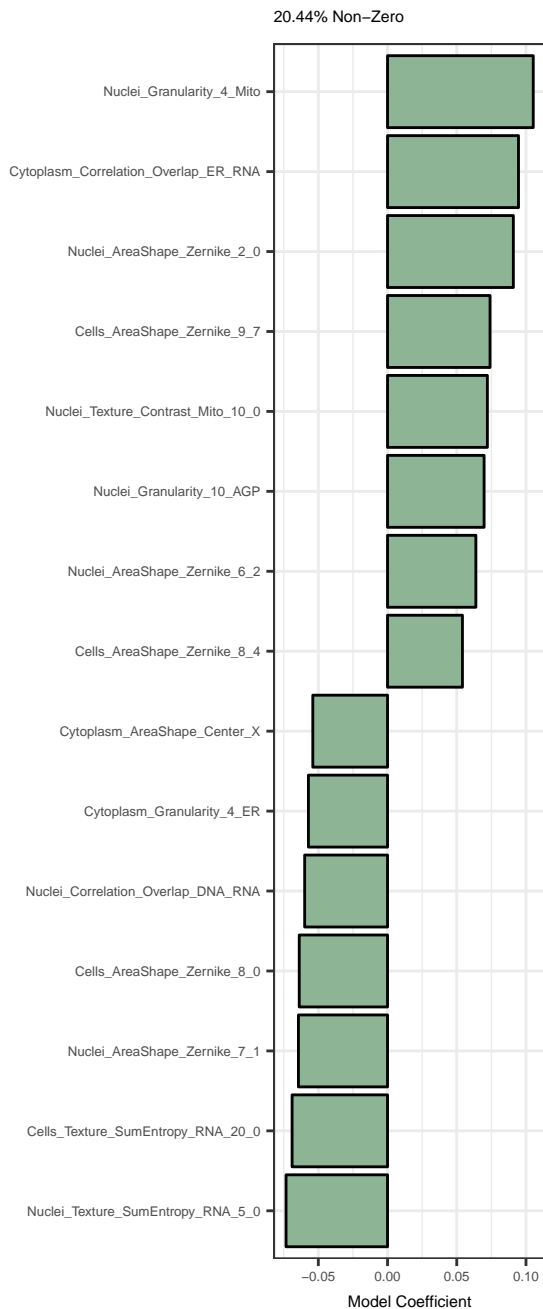
ALL – Nucleus Roundness



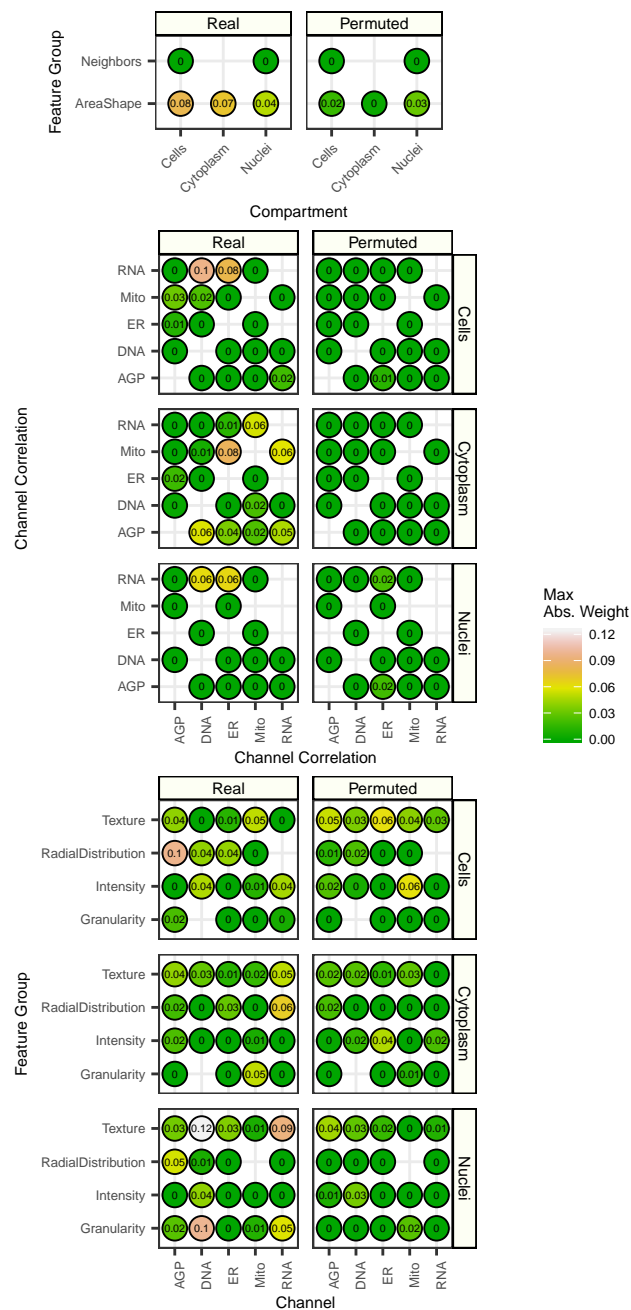
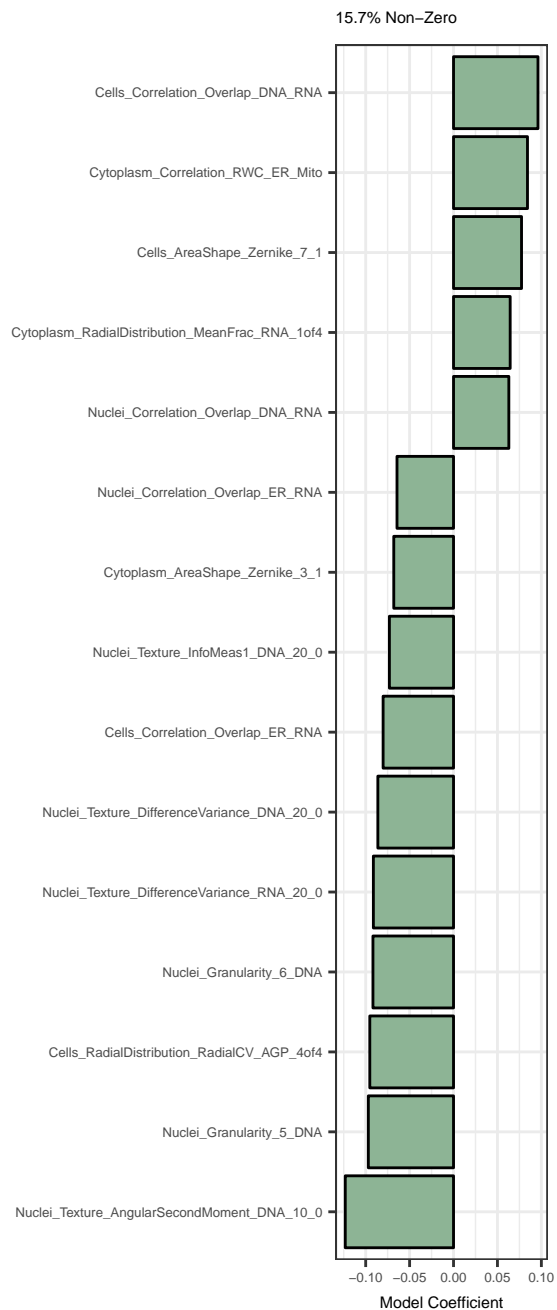
CC – % early M



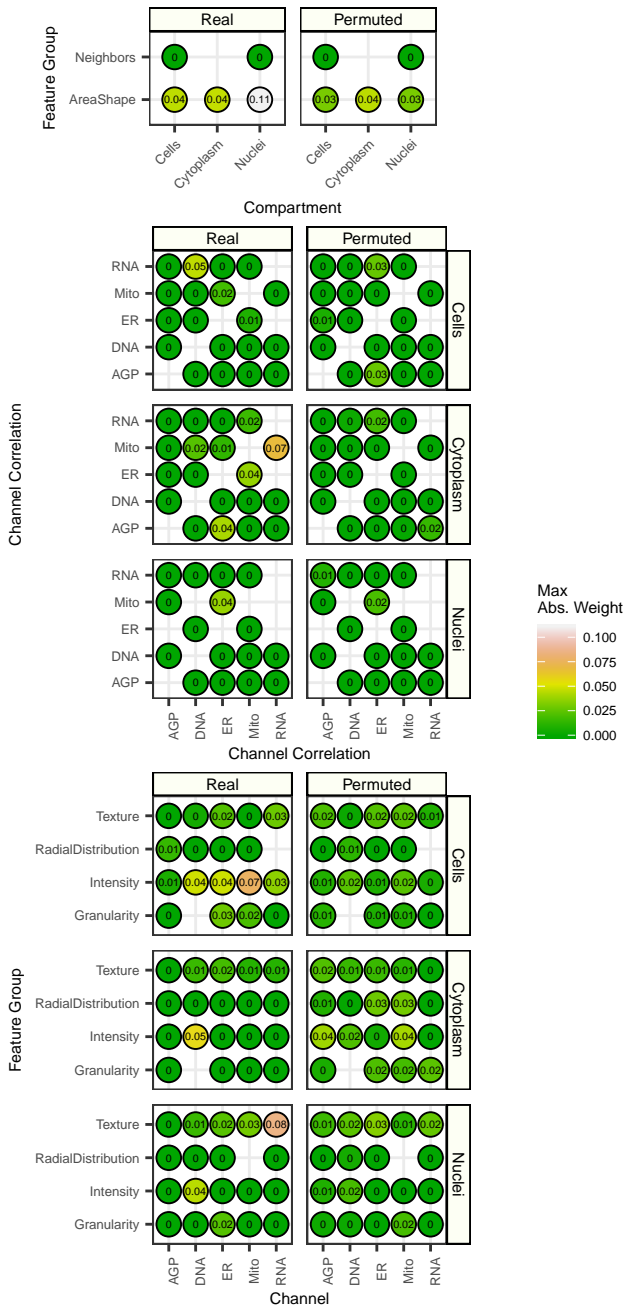
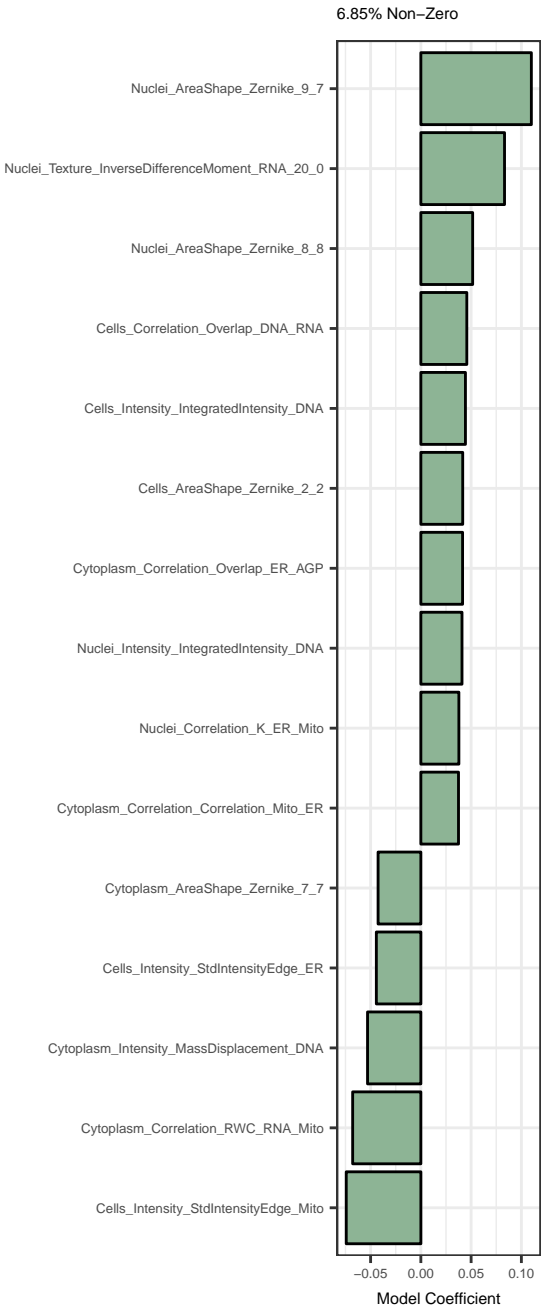
CC – % G1



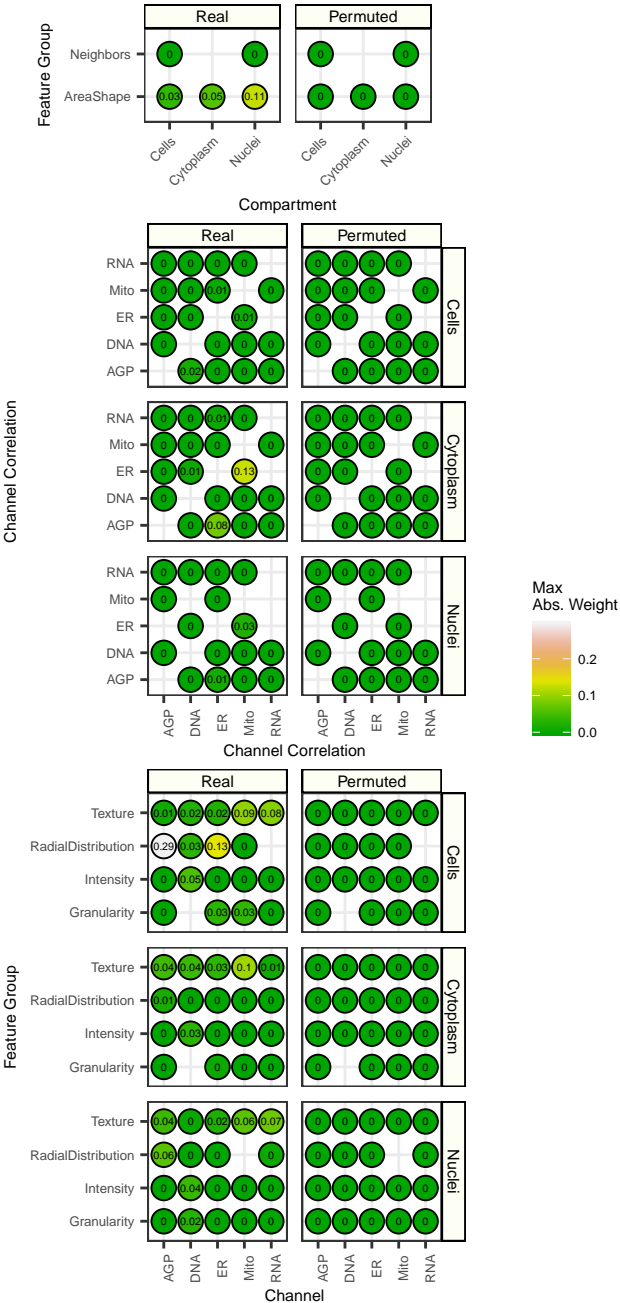
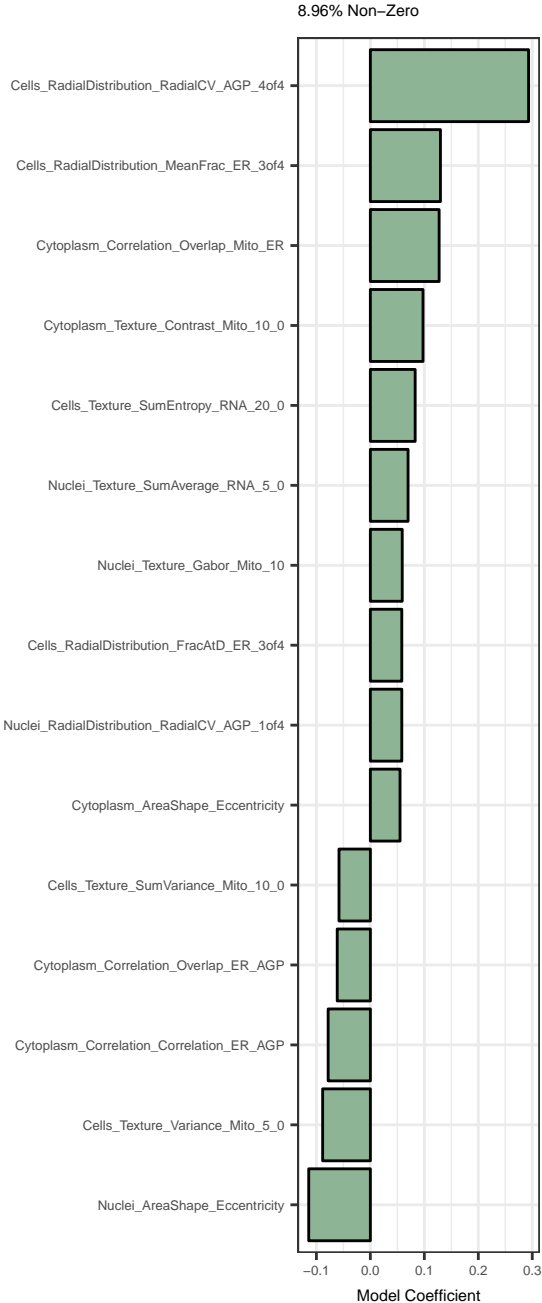
CC - % G2



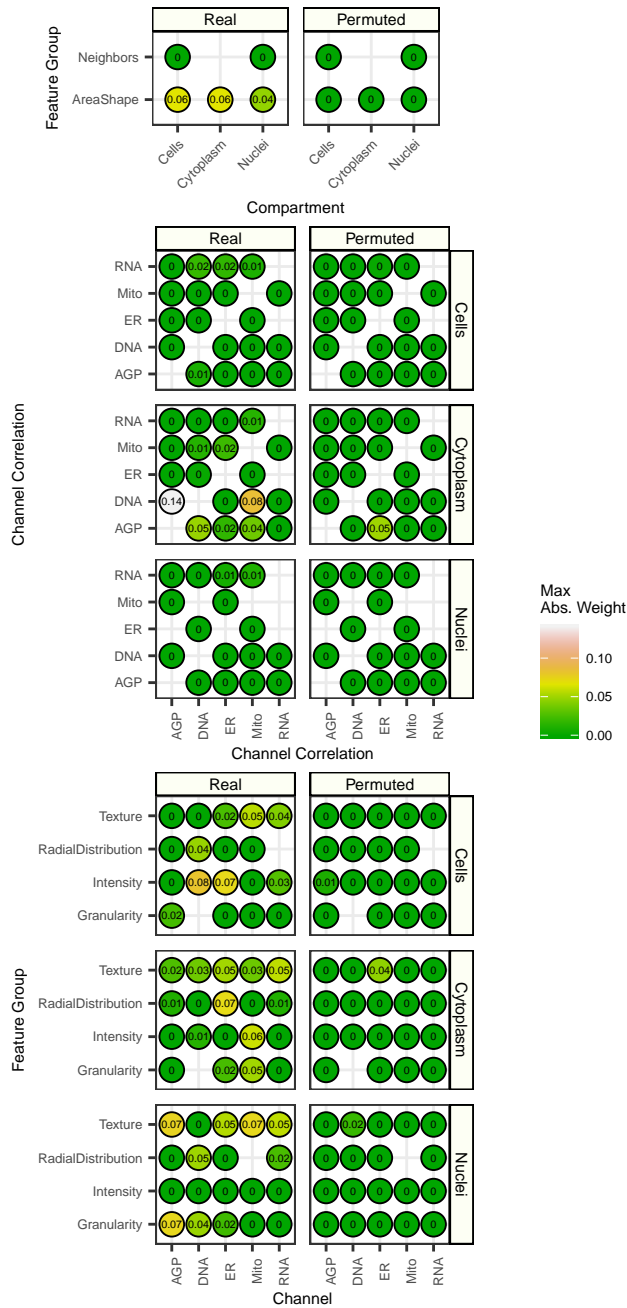
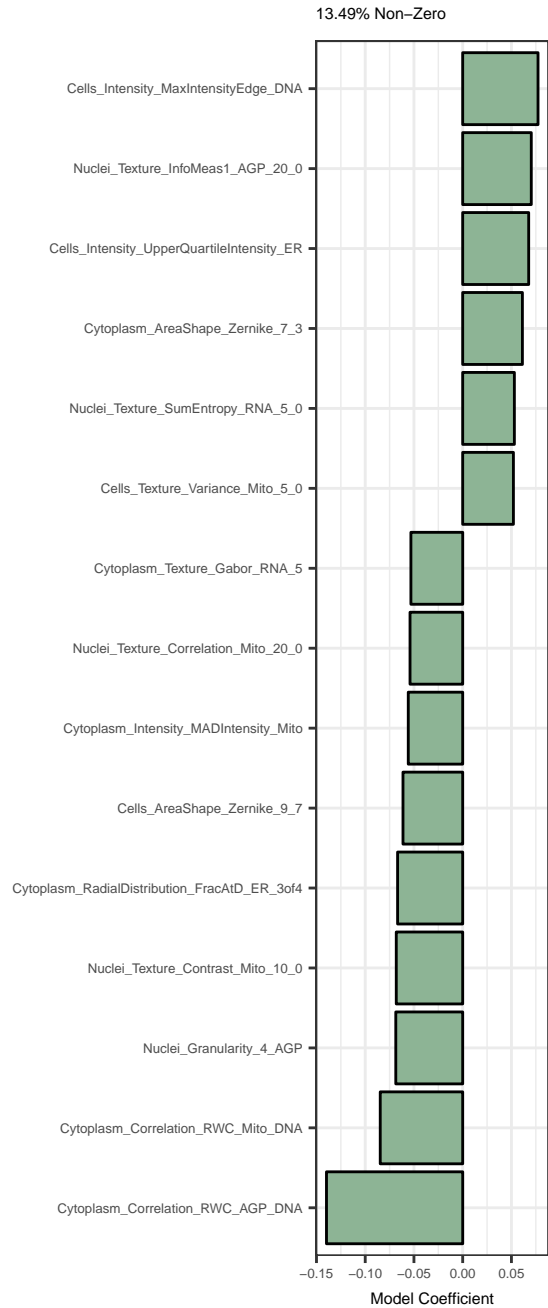
CC – % High gH2AX spots



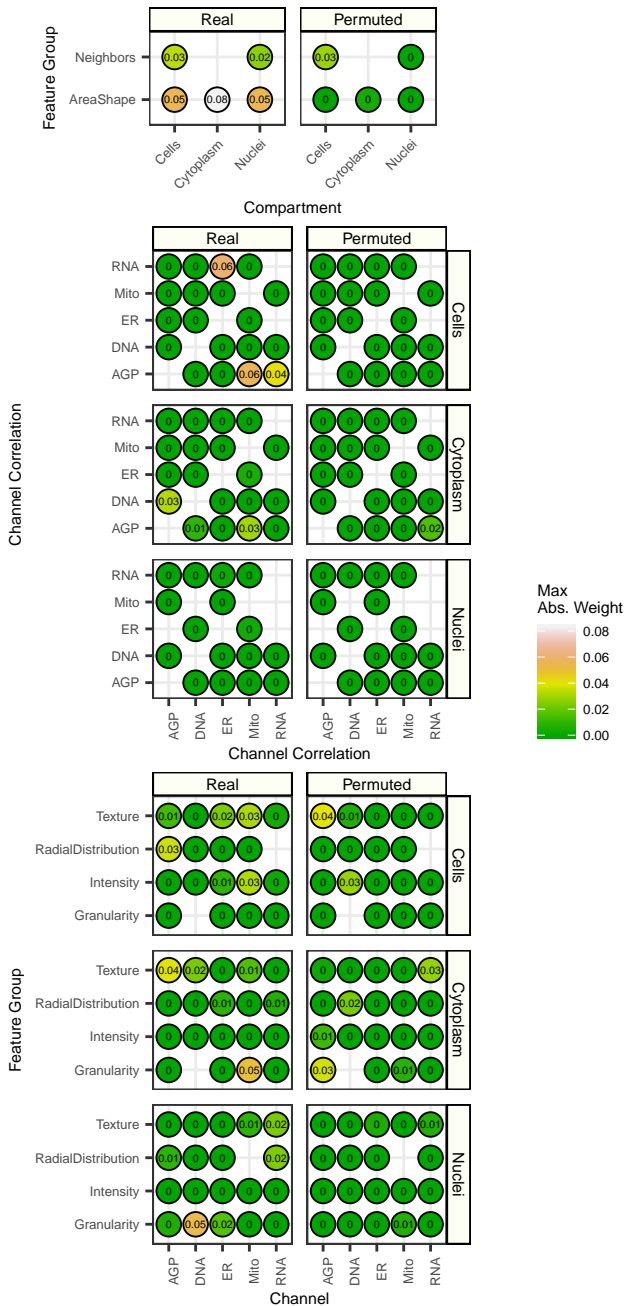
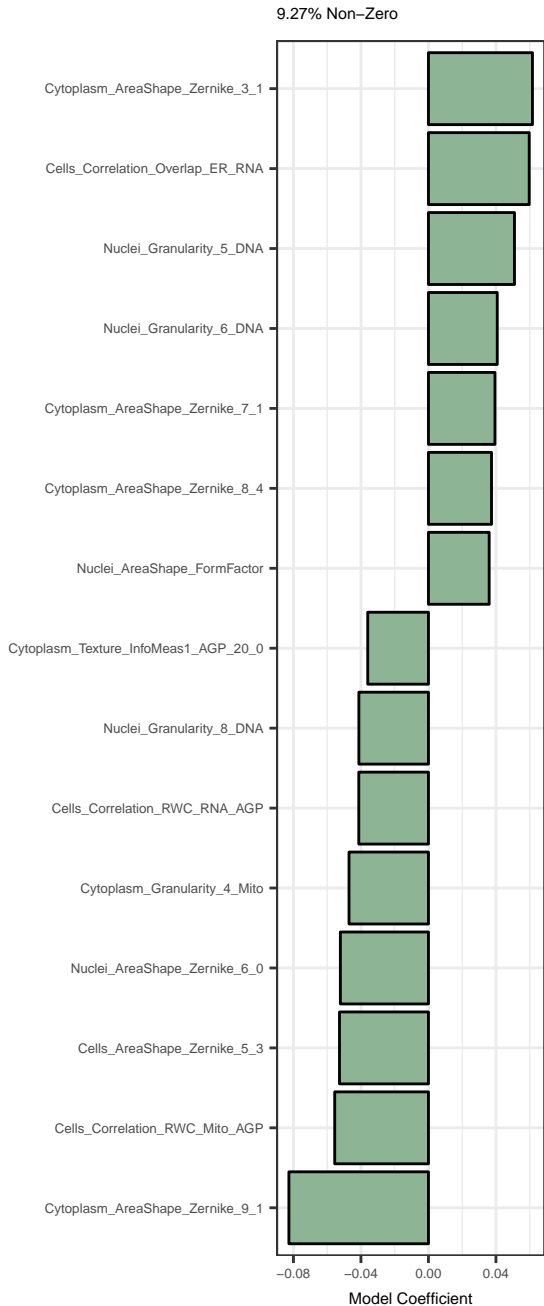
CC - % Late M



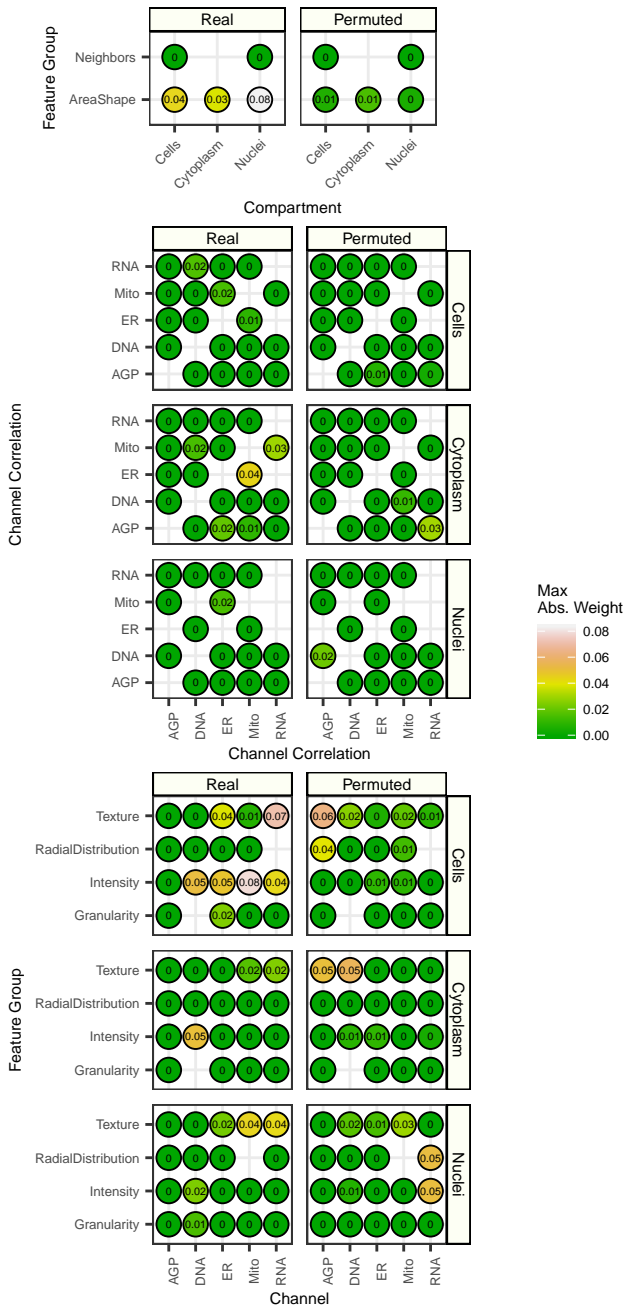
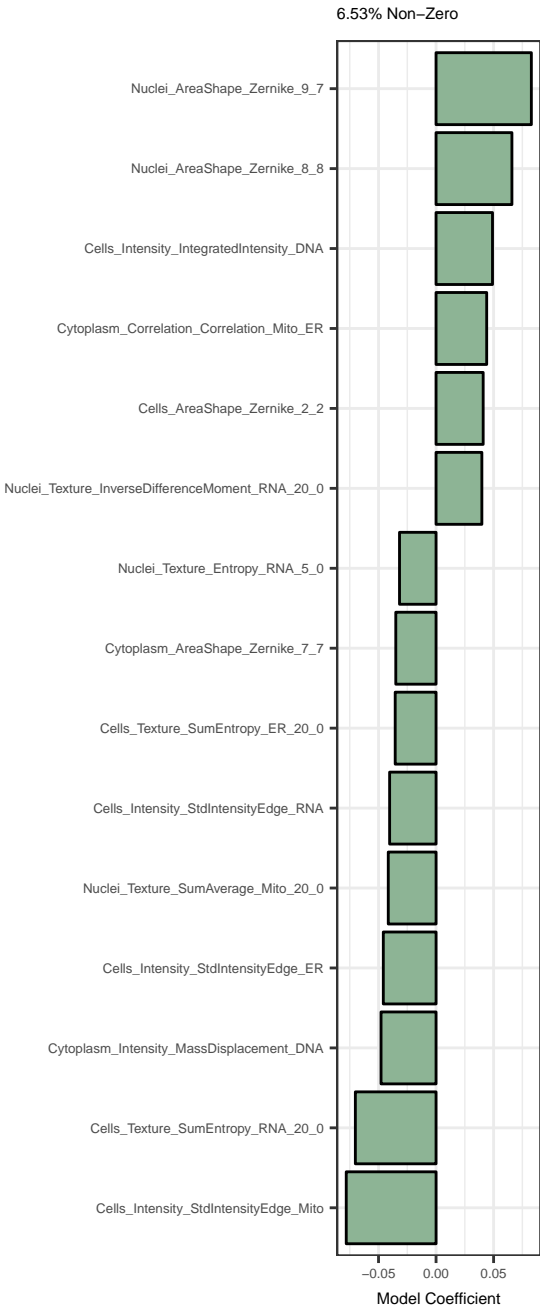
CC - % M



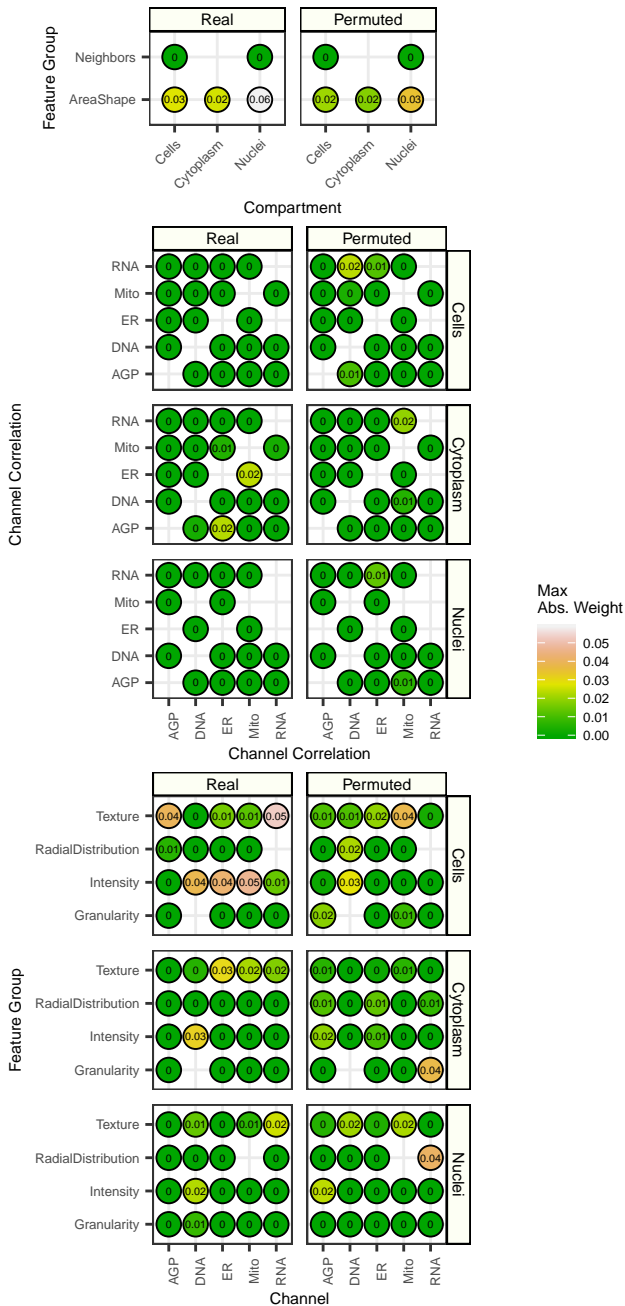
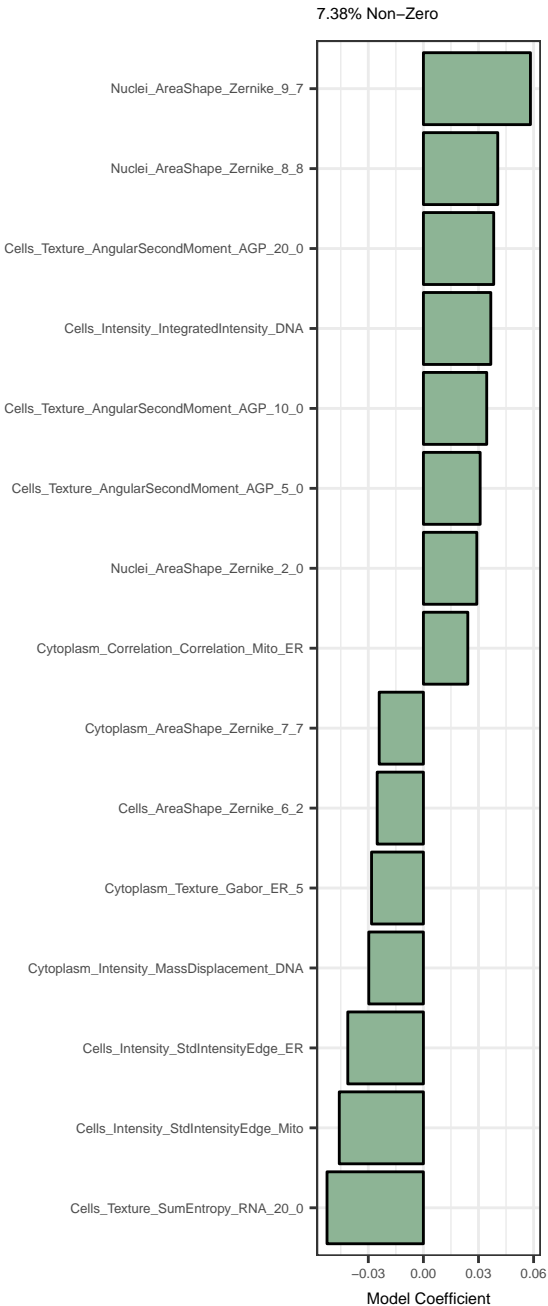
CC – # cells



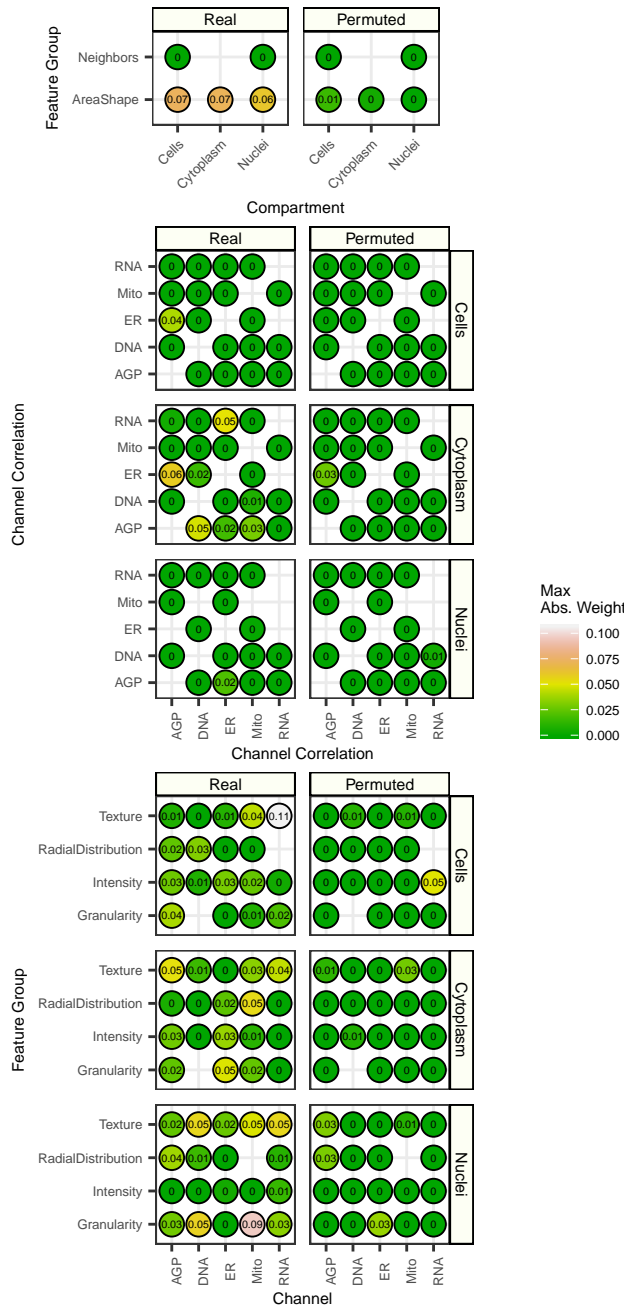
CC – # of gH2AX Spots



CC – # of gH2AX Spots per Area of Nucleus

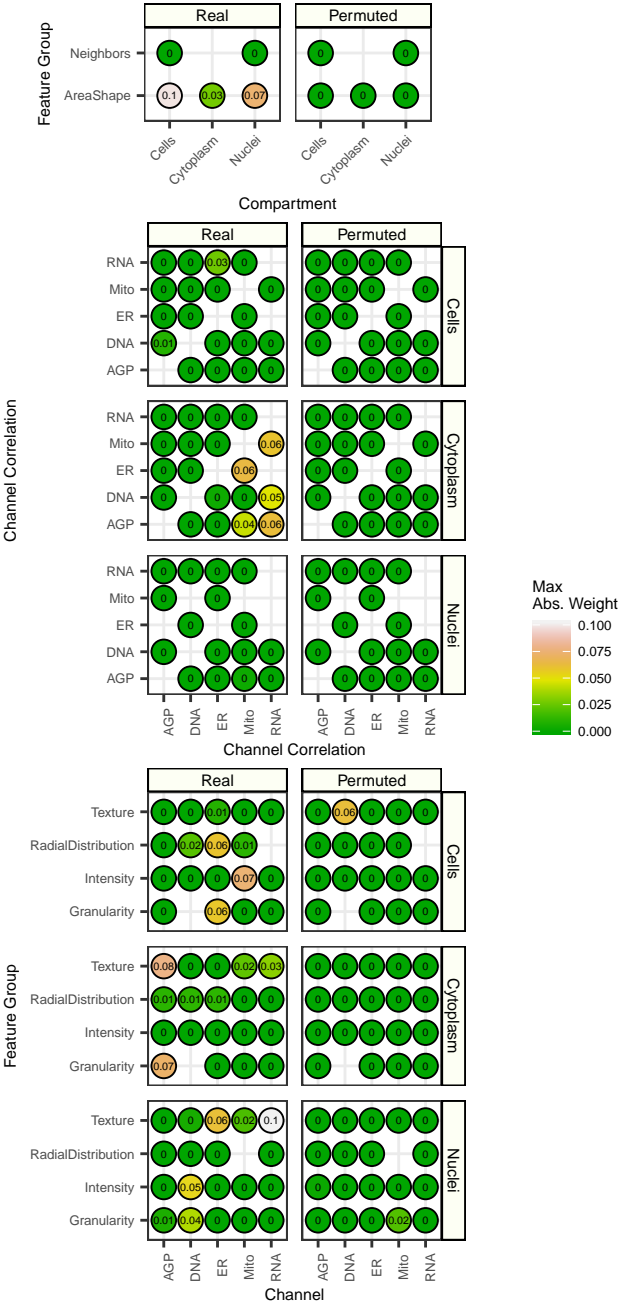
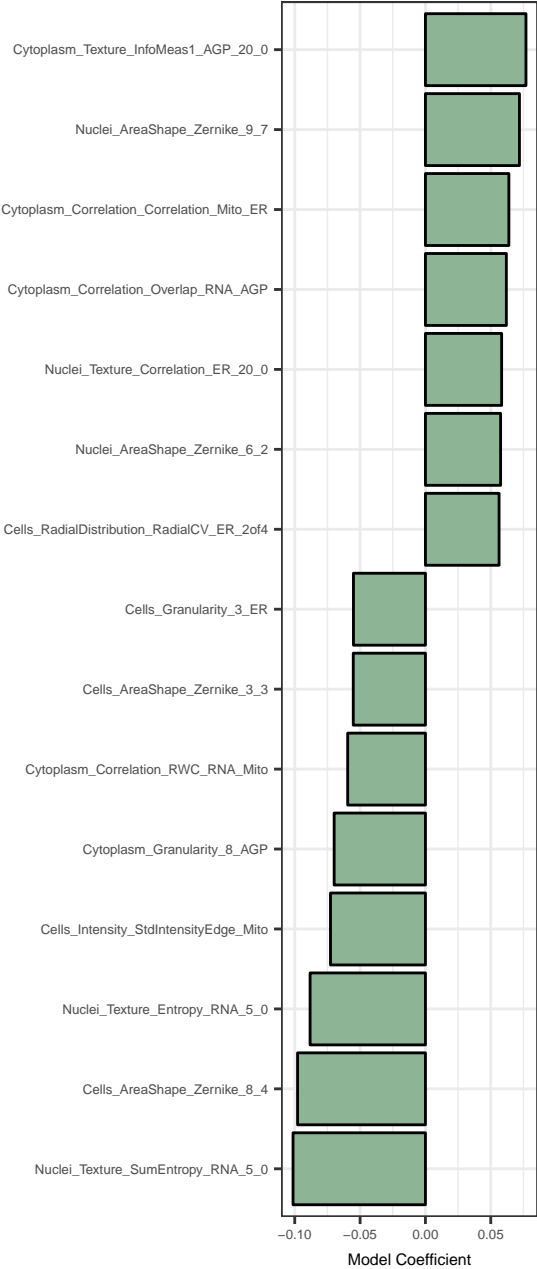


CC - % S

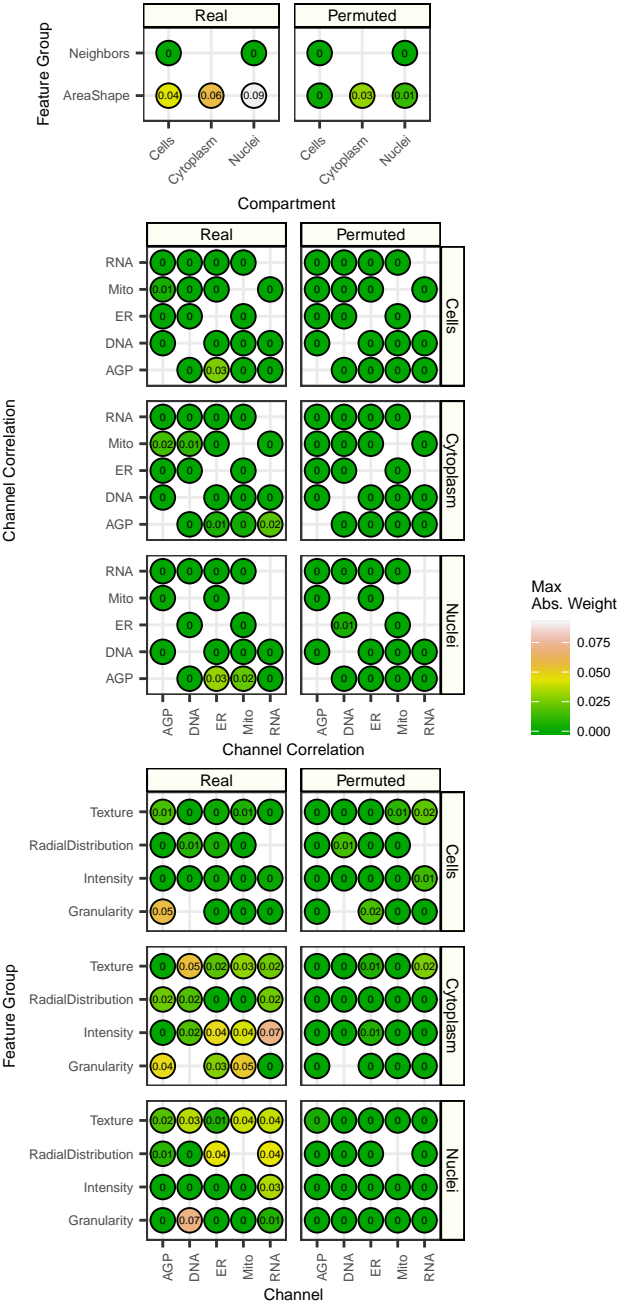
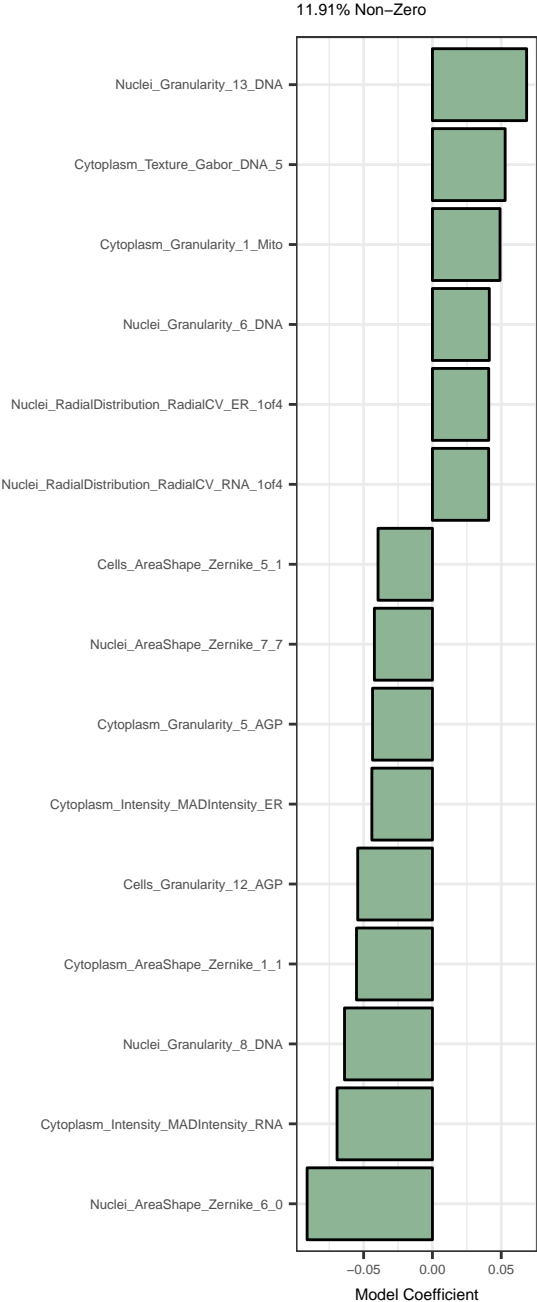


early M – % High gH2AX Spots

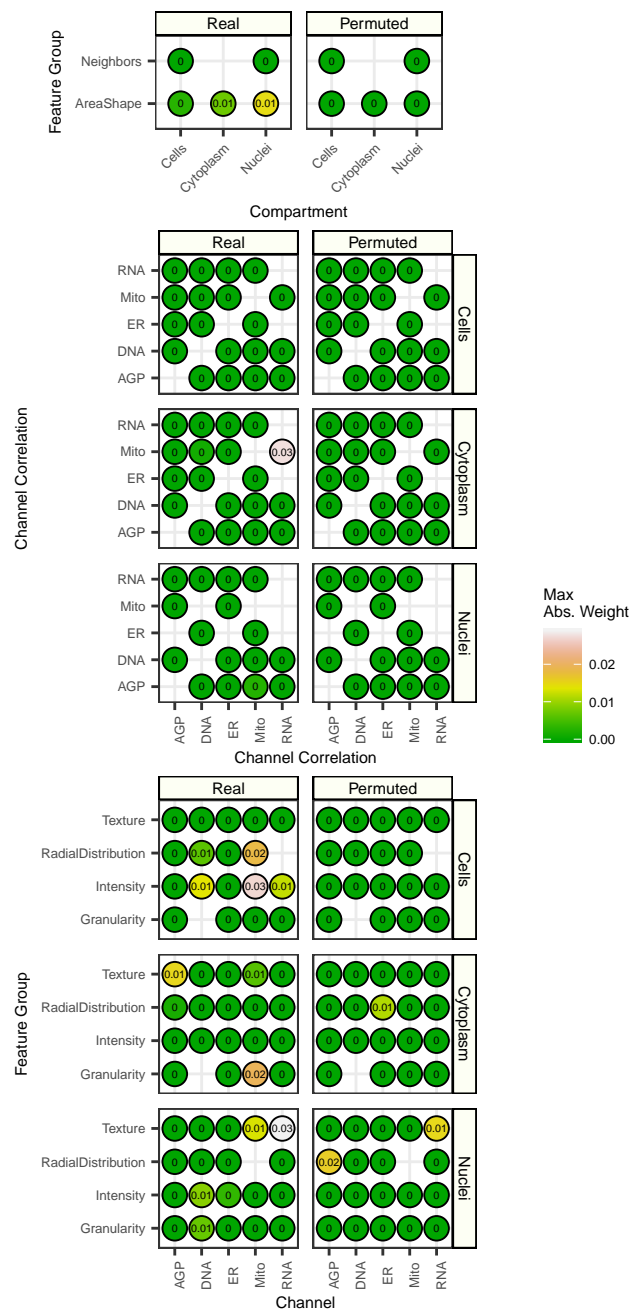
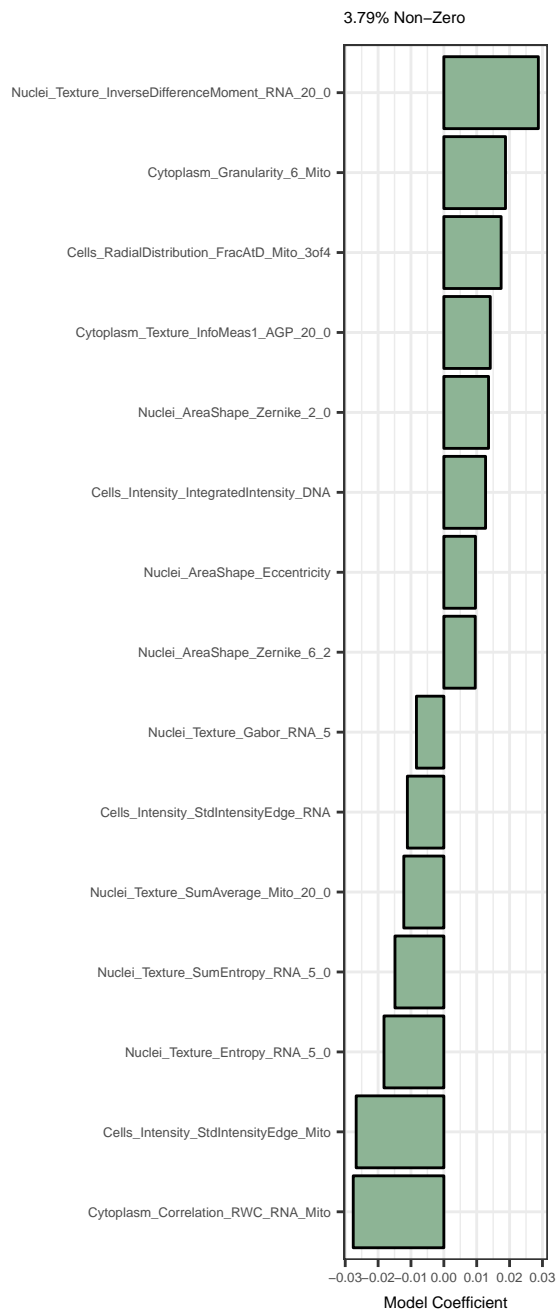
7.06% Non-Zero



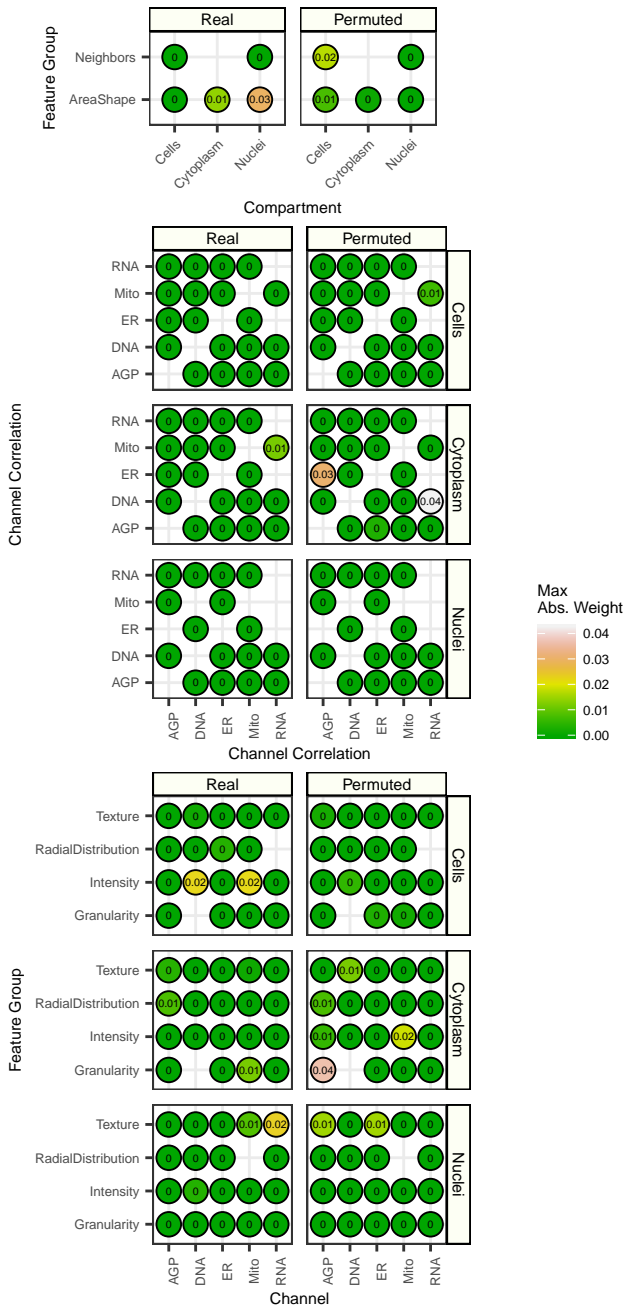
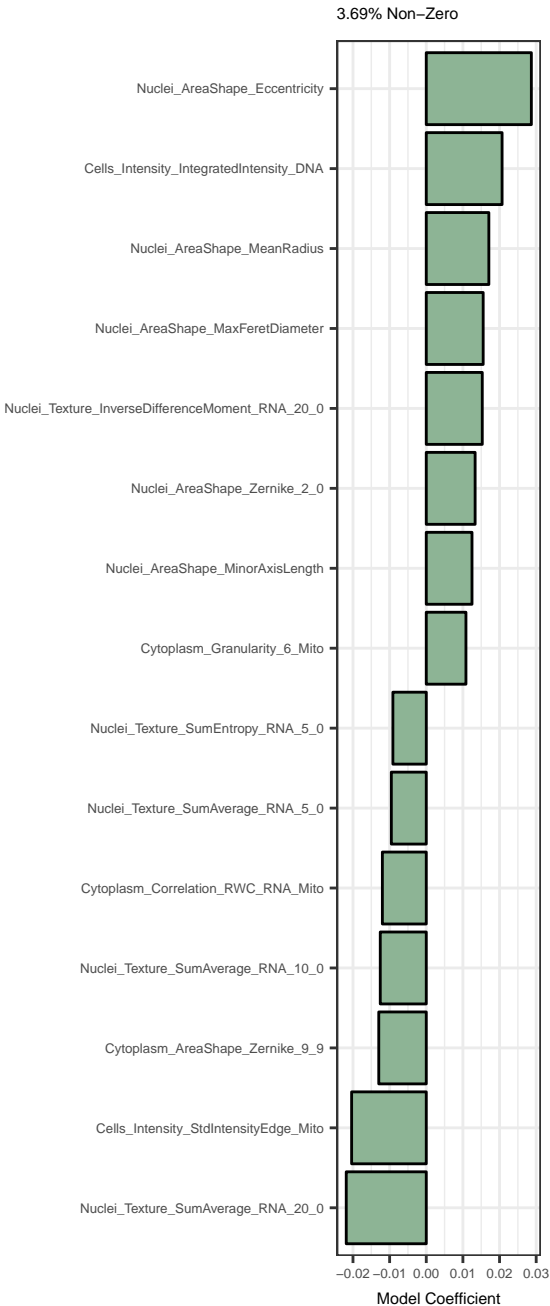
early M – # cells



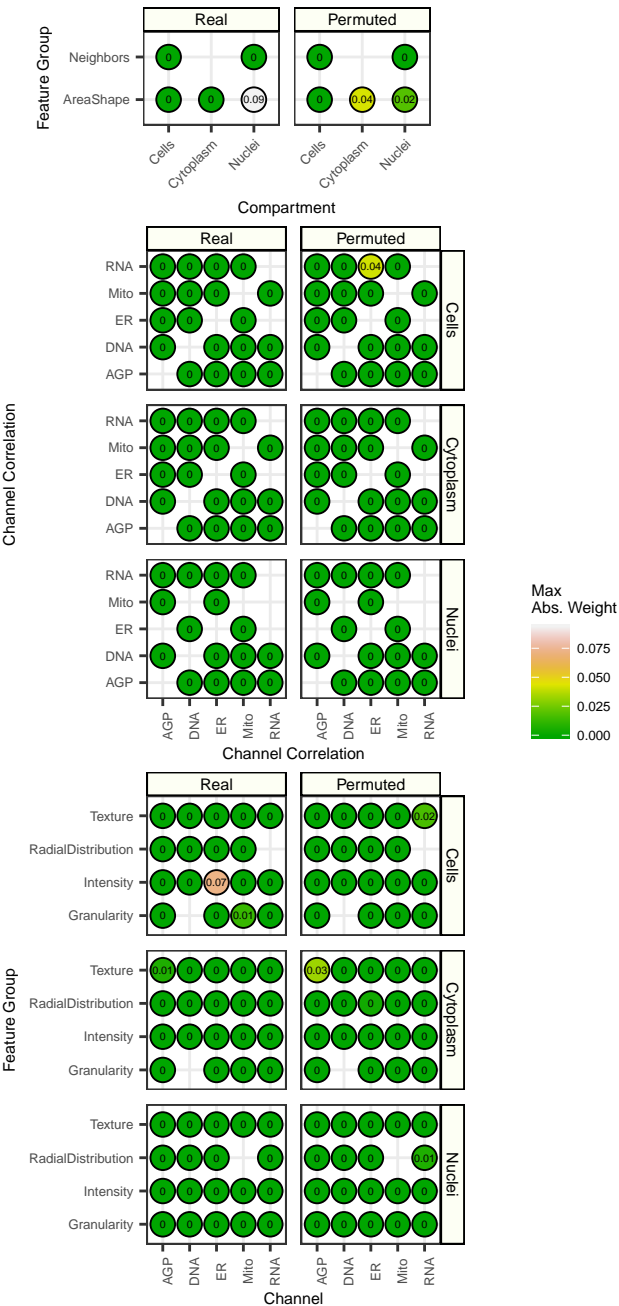
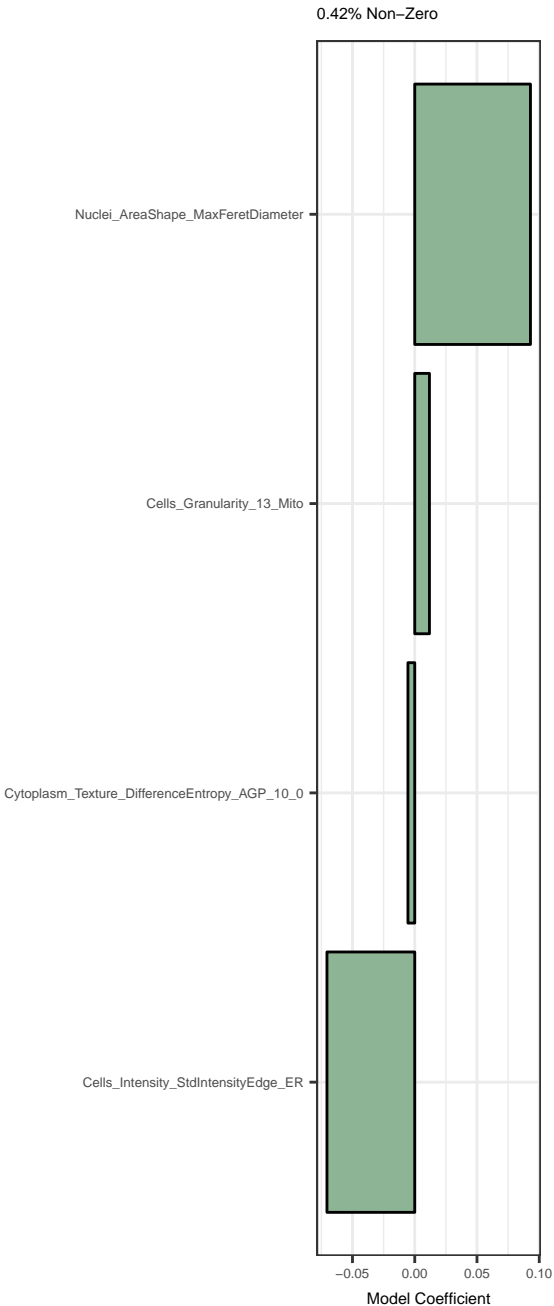
early M – # of gH2AX Spots



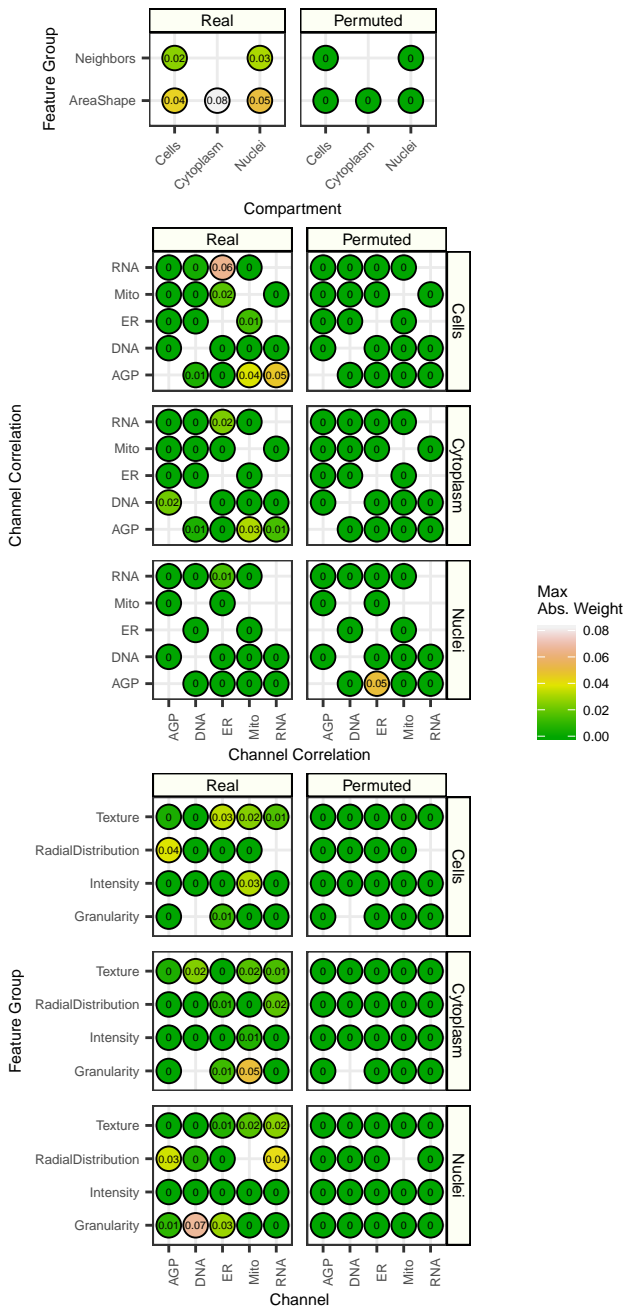
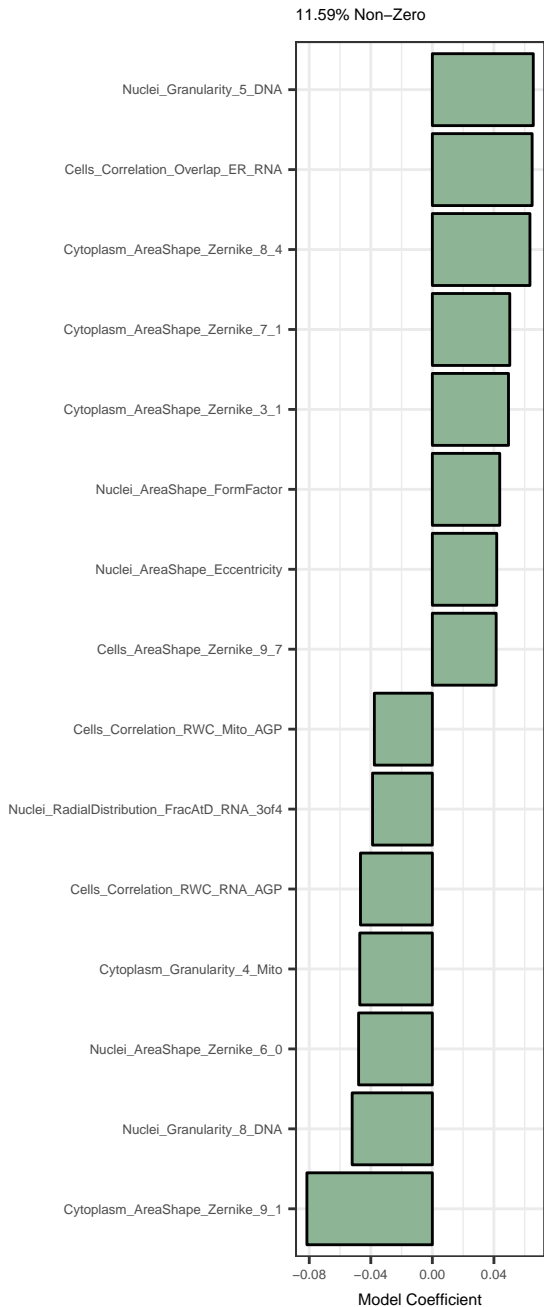
early M – # of gH2AX Spots per Area of Nucleus



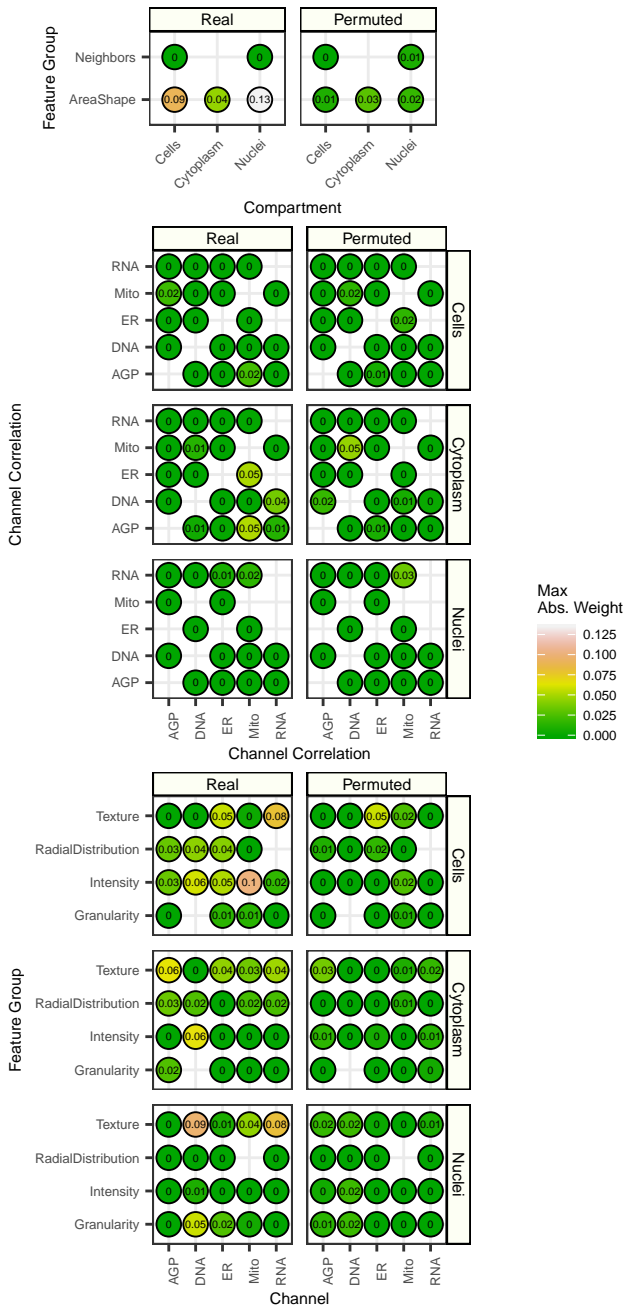
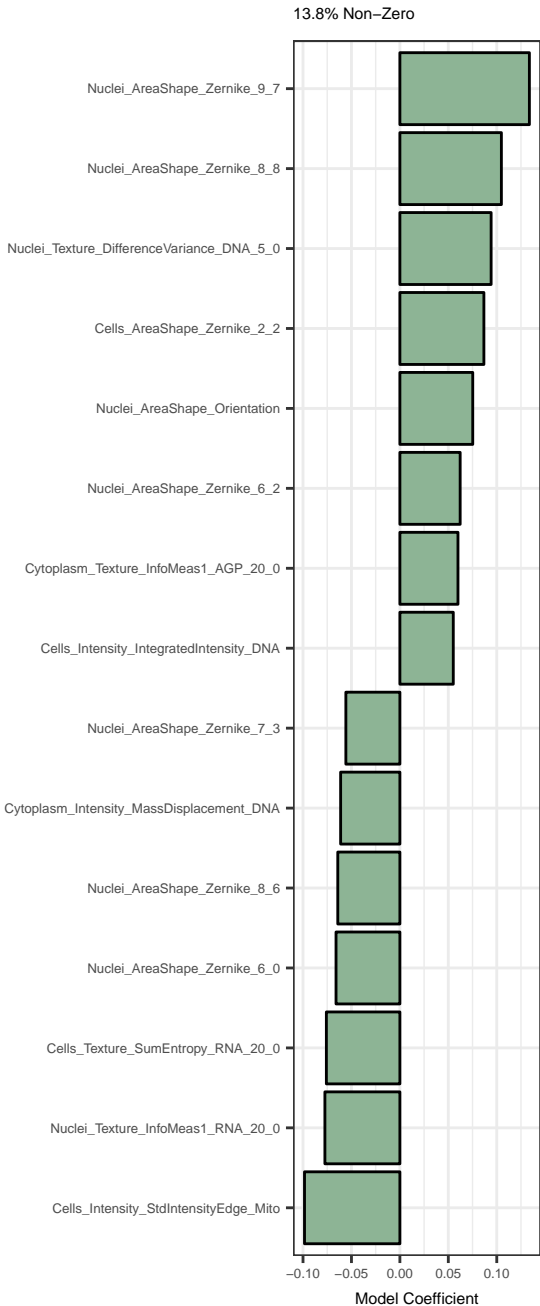
G1 – % High gH2AX Spots



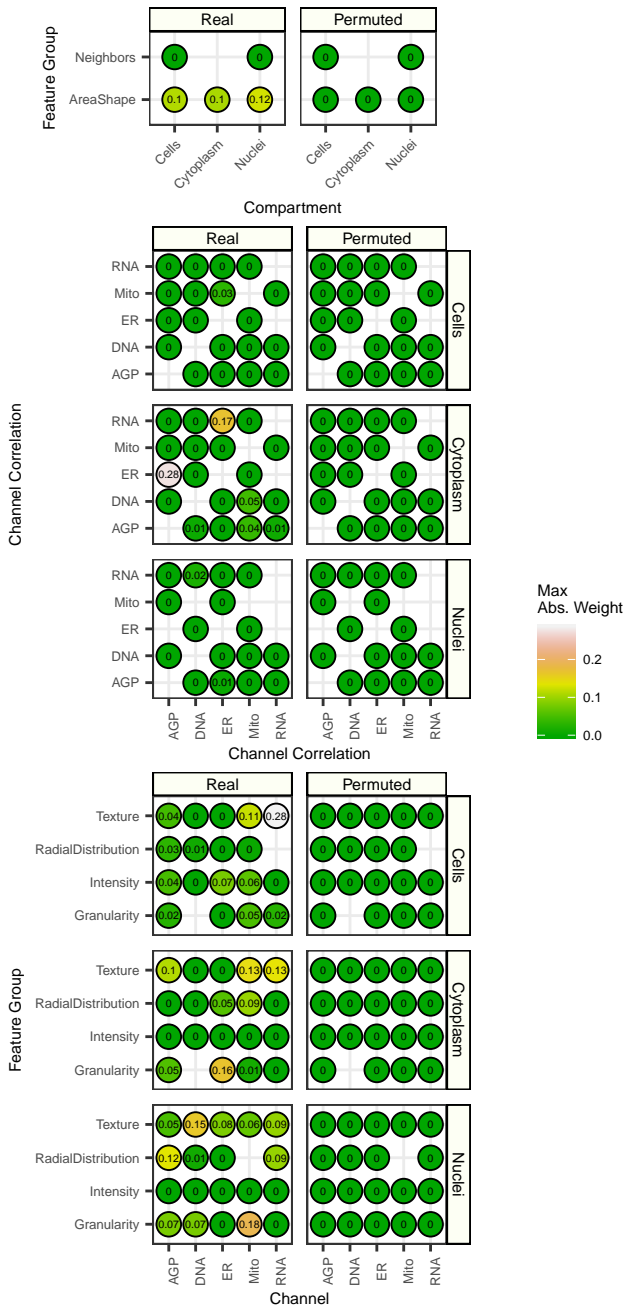
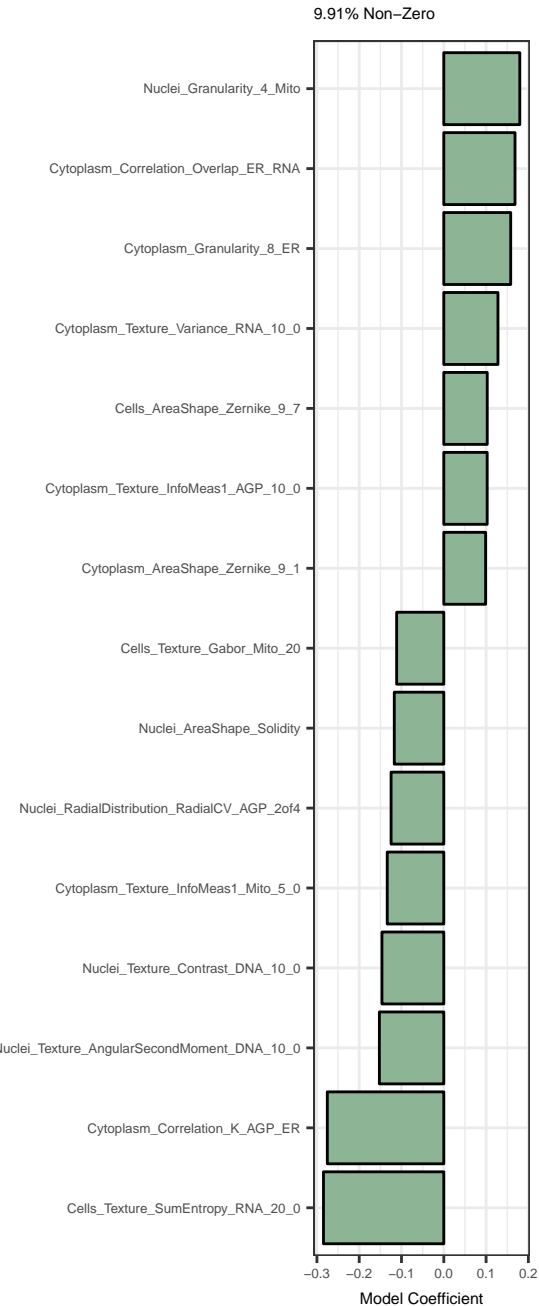
G1 – # cells



G1 – # of gH2AX Spots

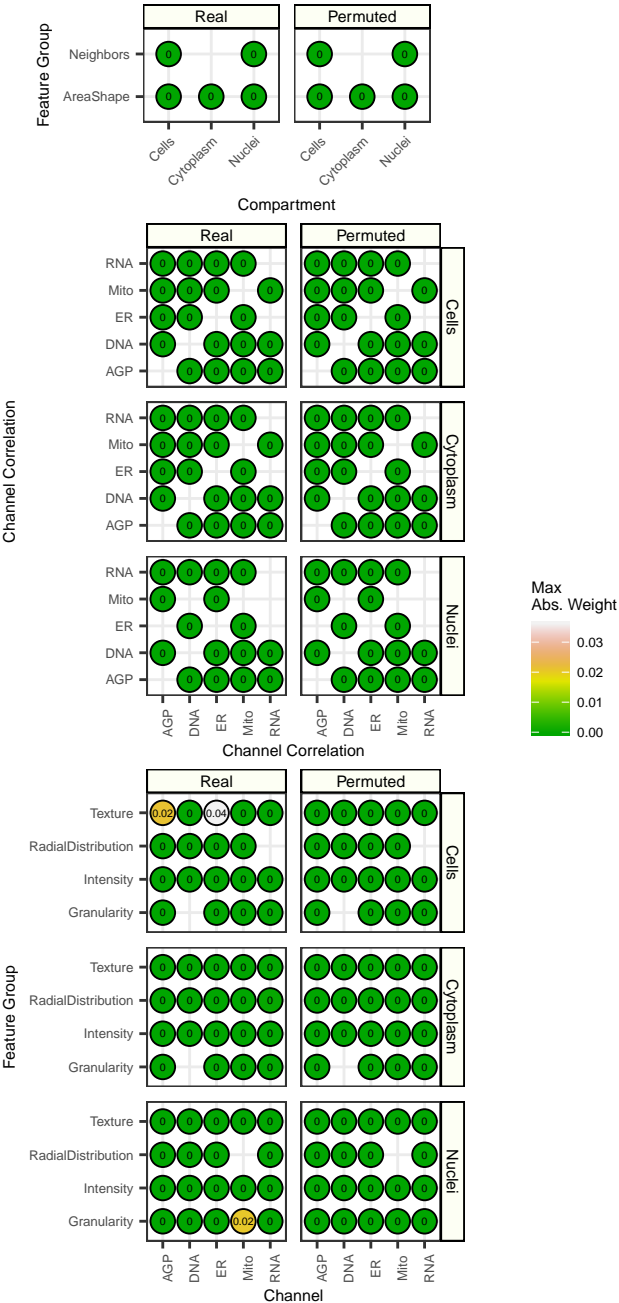
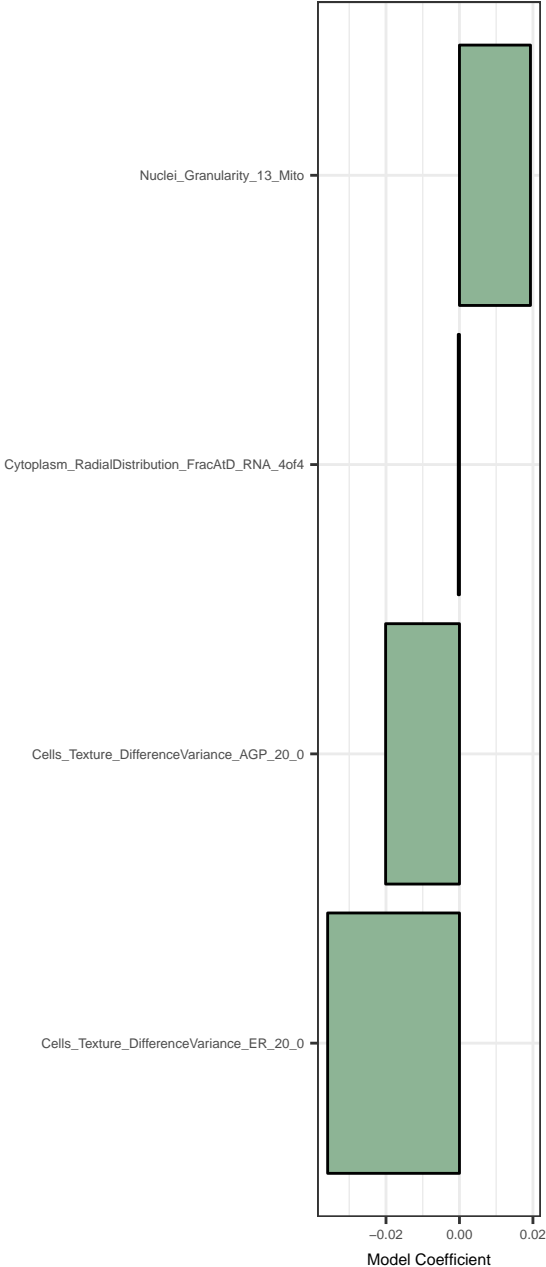


Cell Count – G1+G2

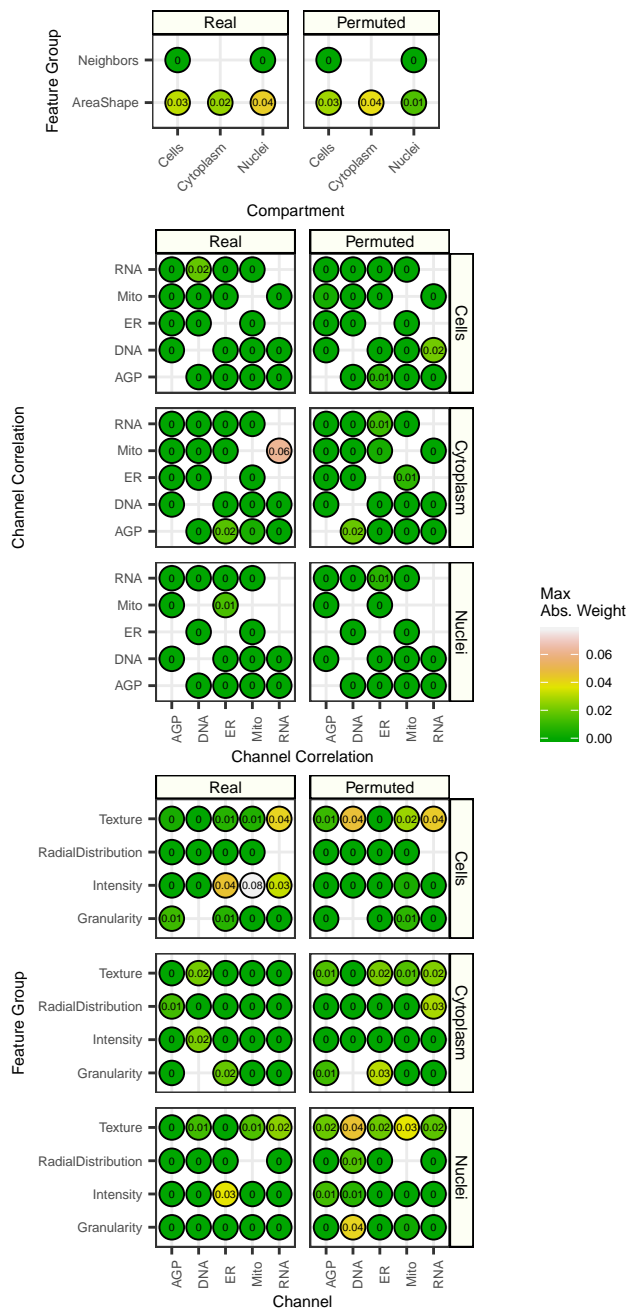
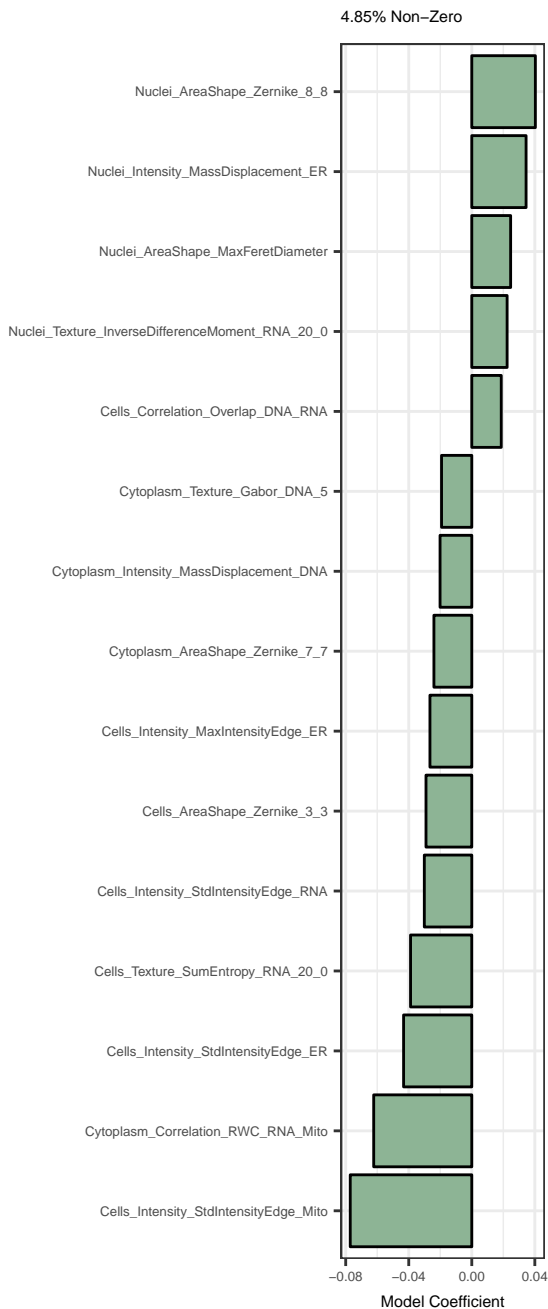


Cell Count – G2/G1

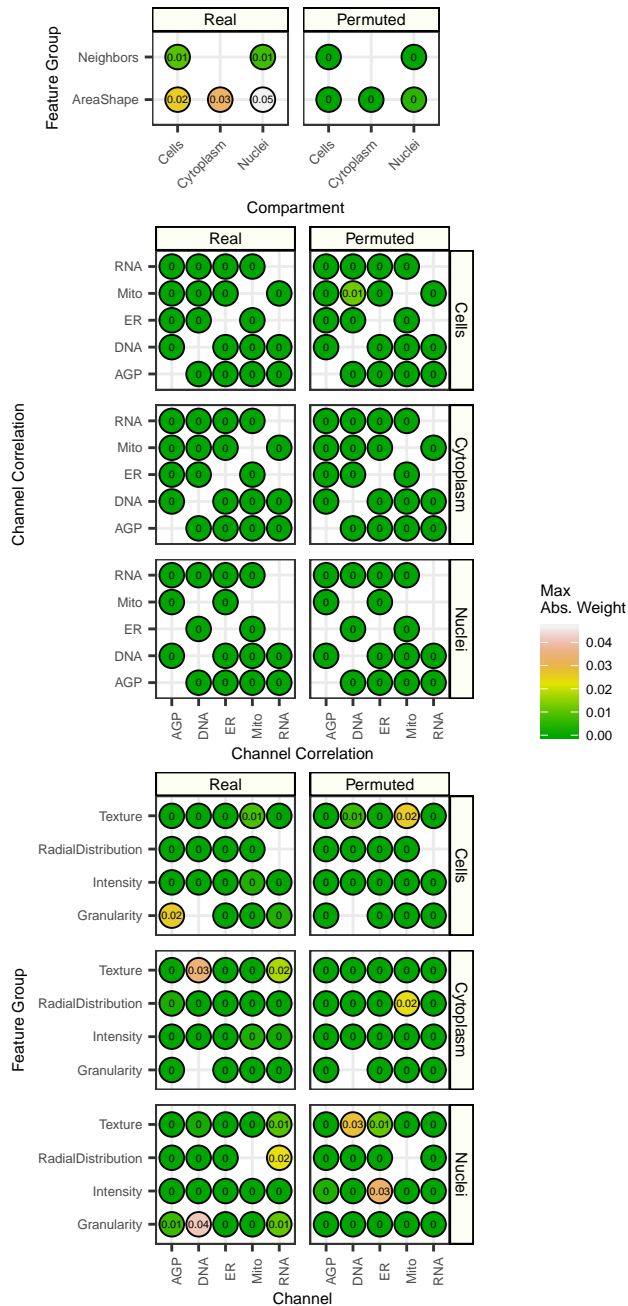
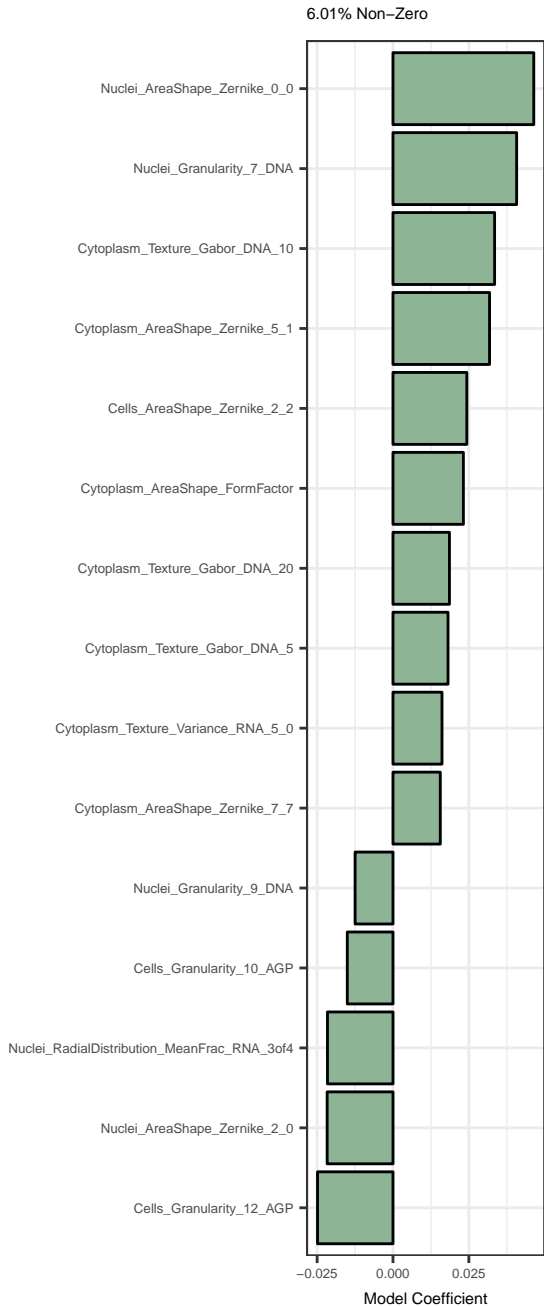
0.42% Non-Zero



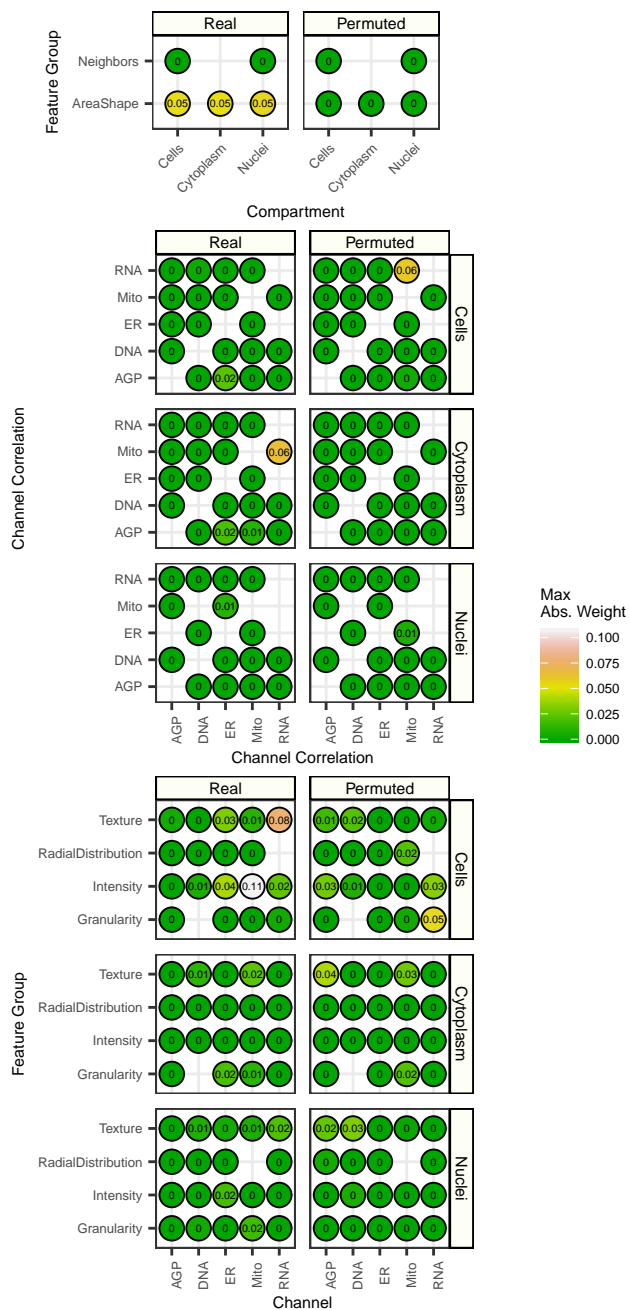
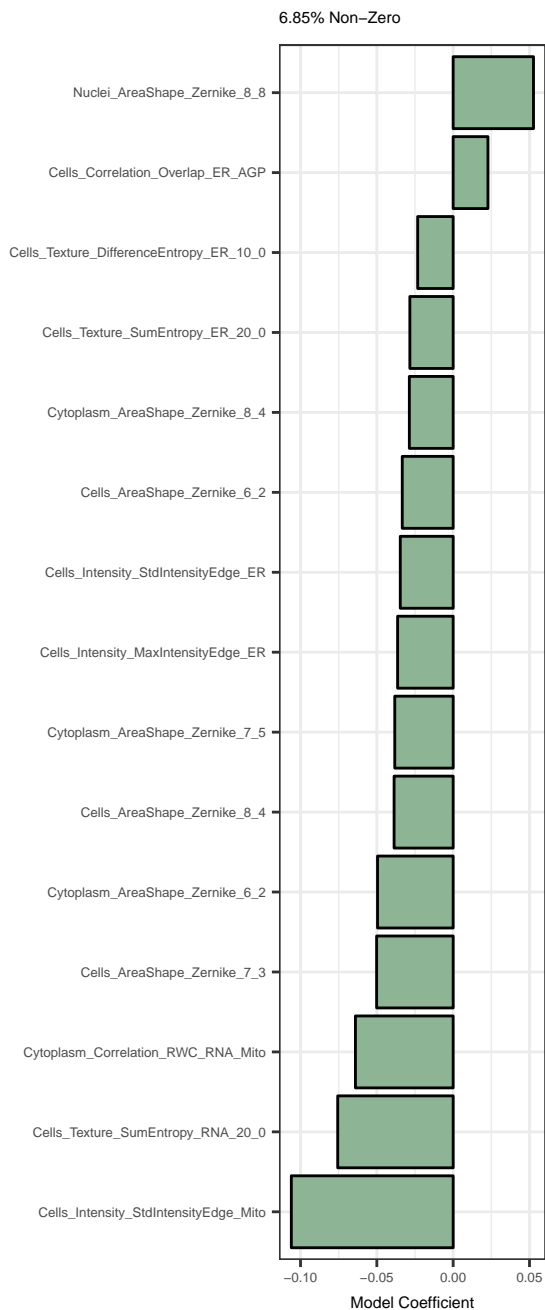
G2 – % High gH2AX Spots



G2 - # cells

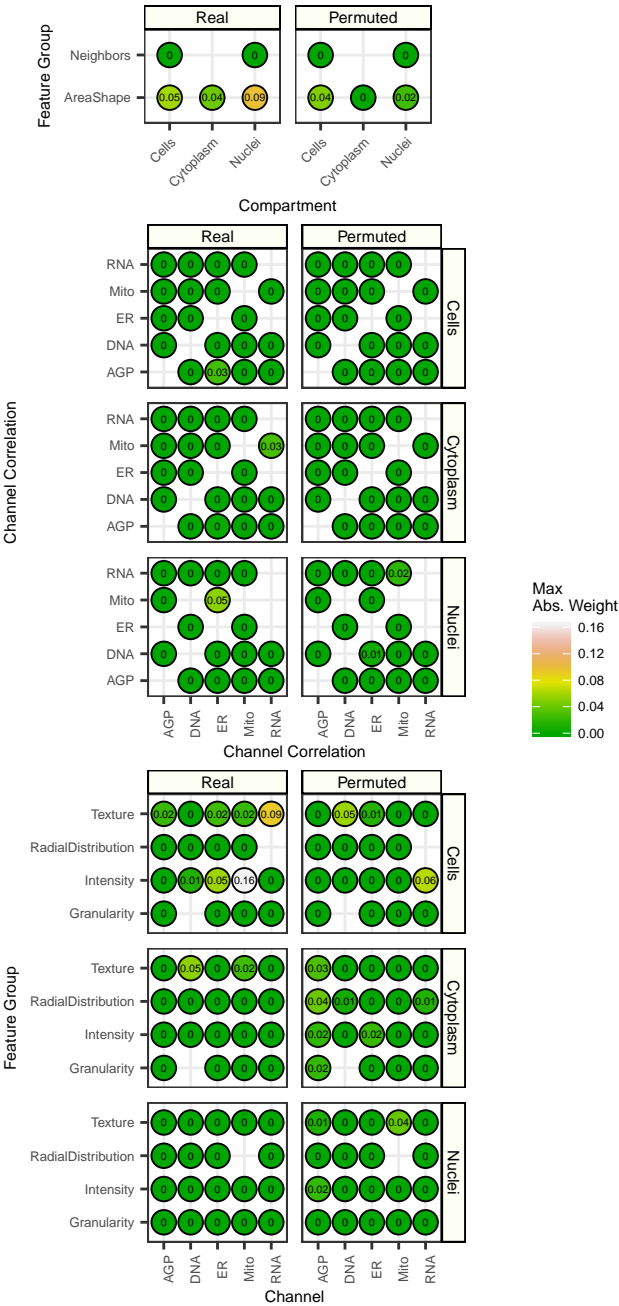
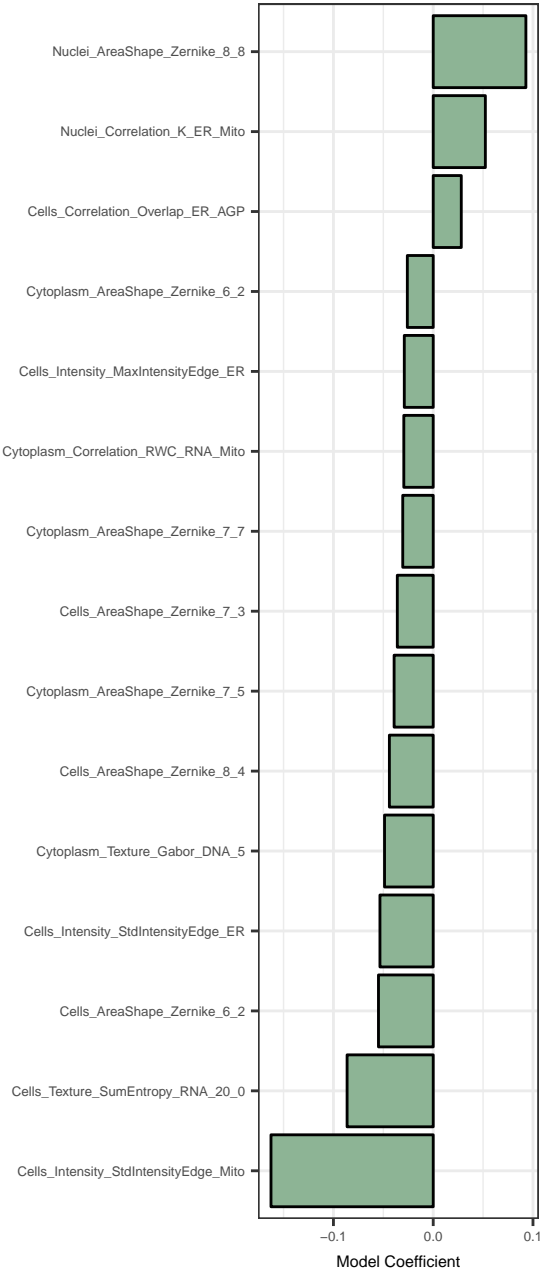


G2 – # of gH2AX Spots

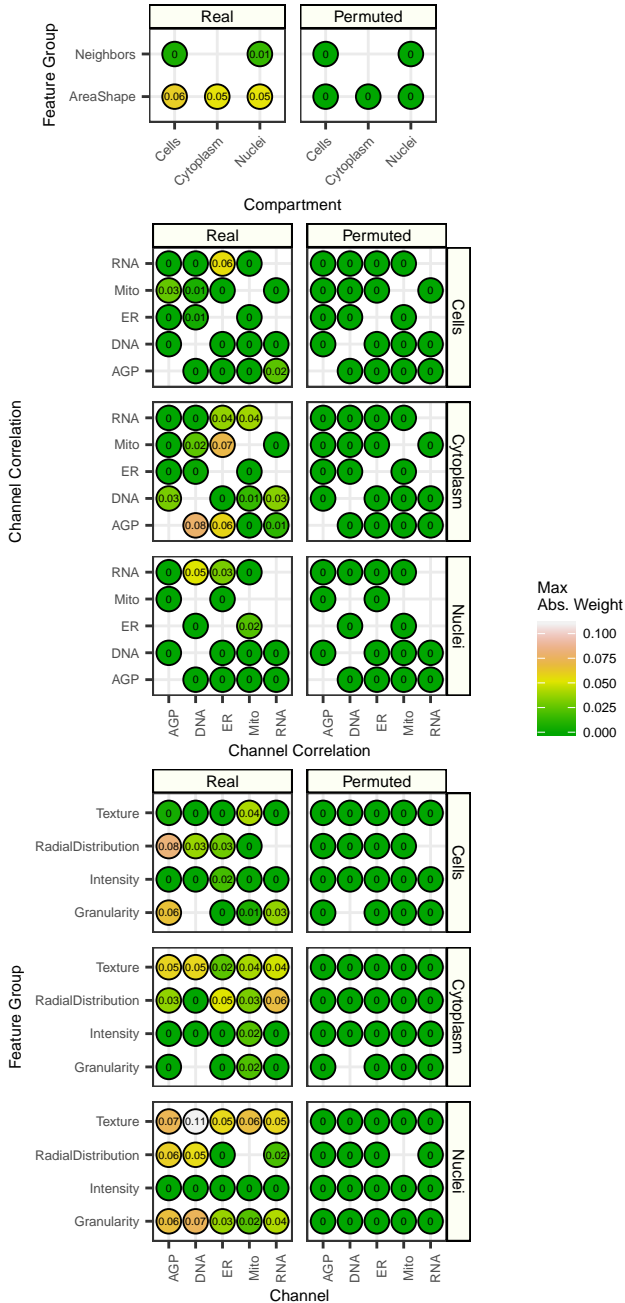
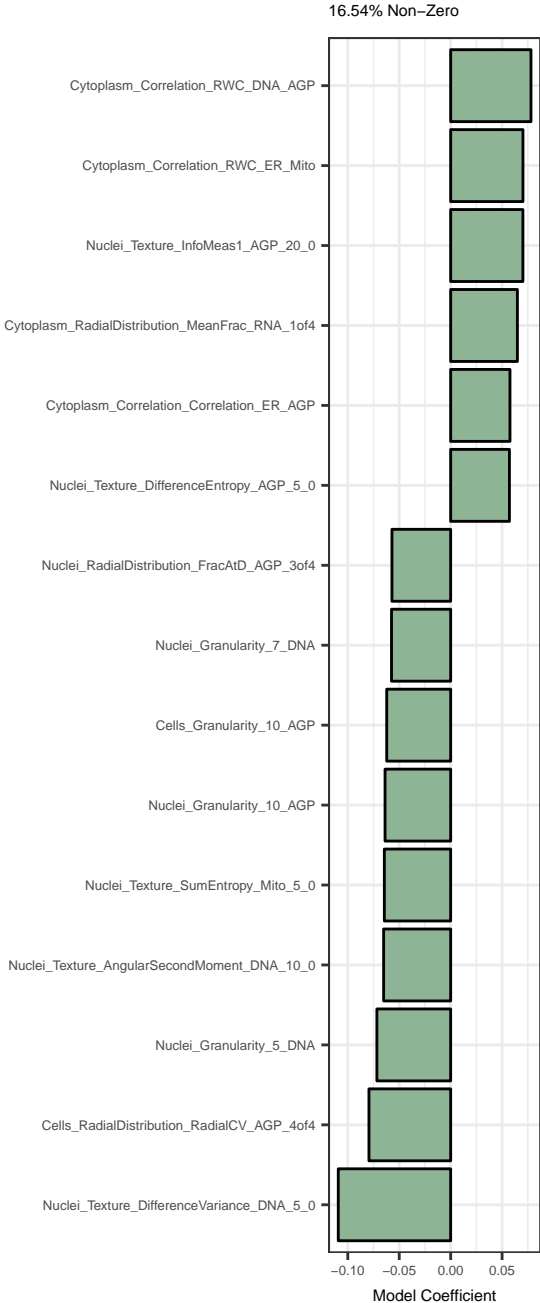


G2 – # of gH2AX Spots per Area of Nucleus

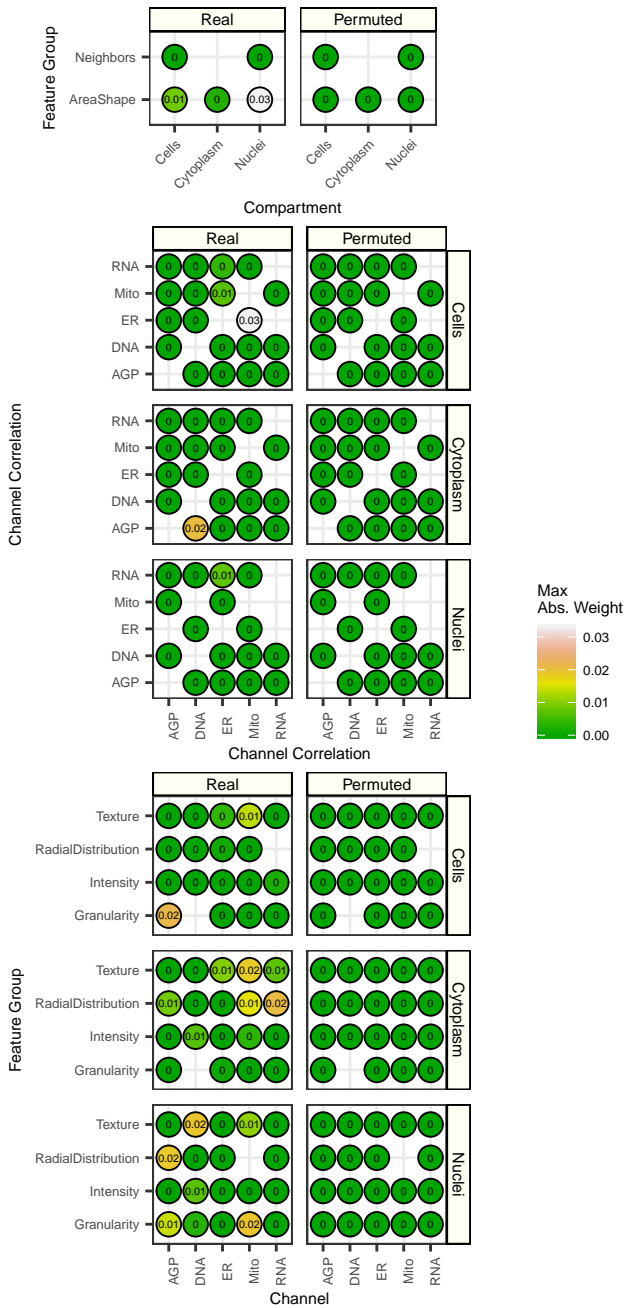
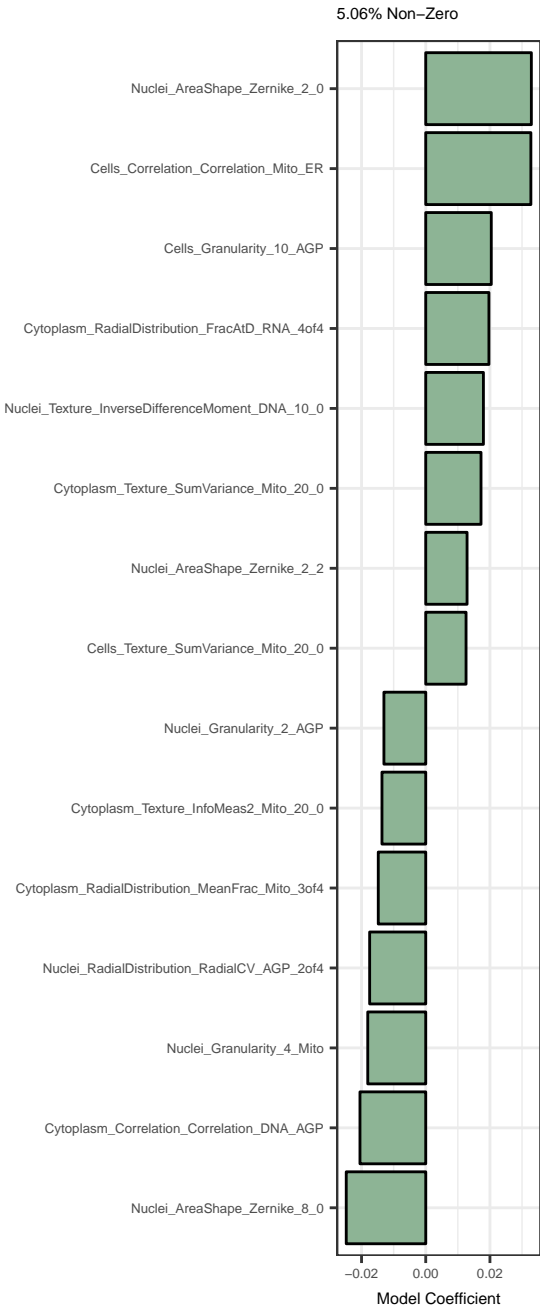
2.63% Non-Zero



Cell Count – G2 + M

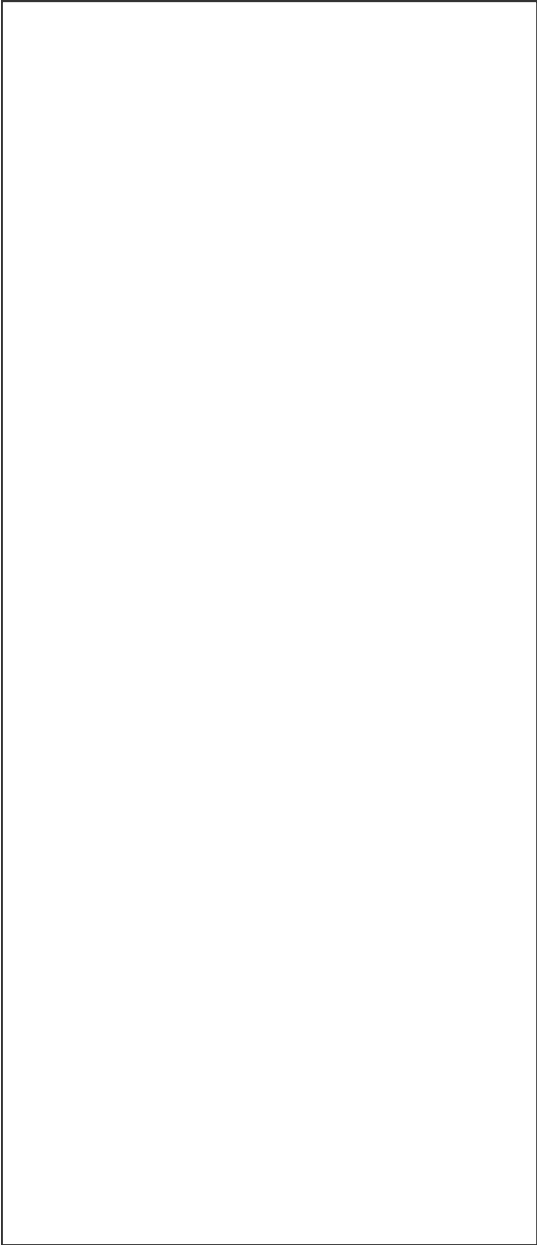


Inf %

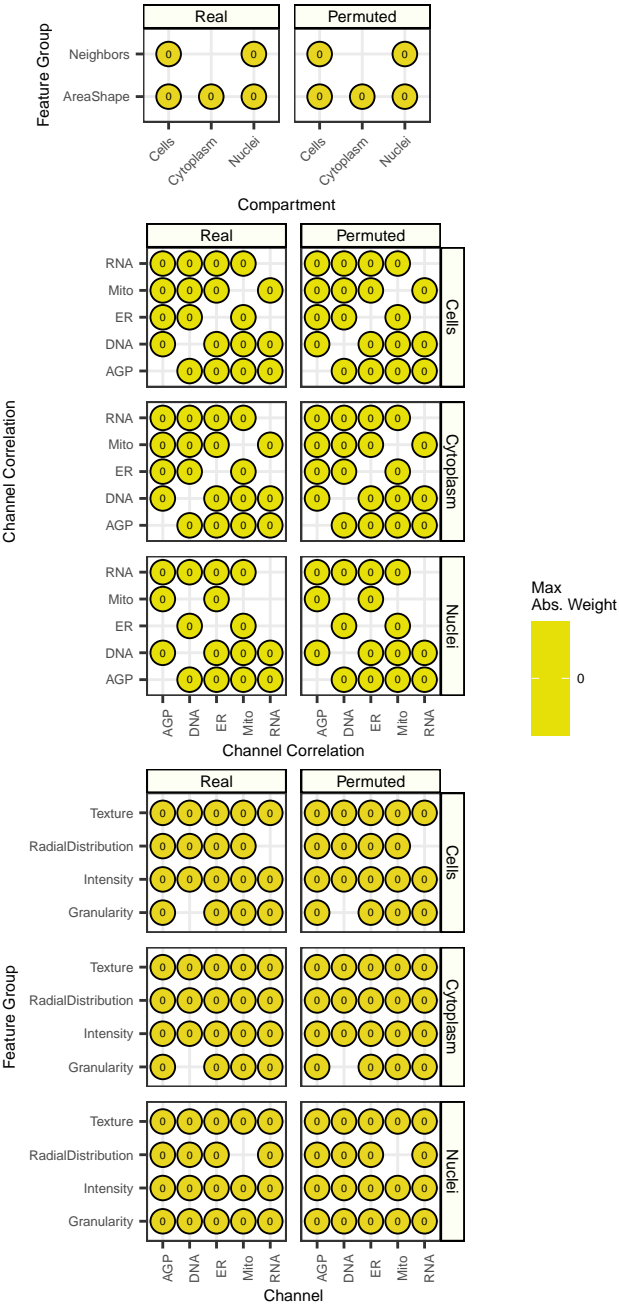


Late M – % High gH2AX Spots

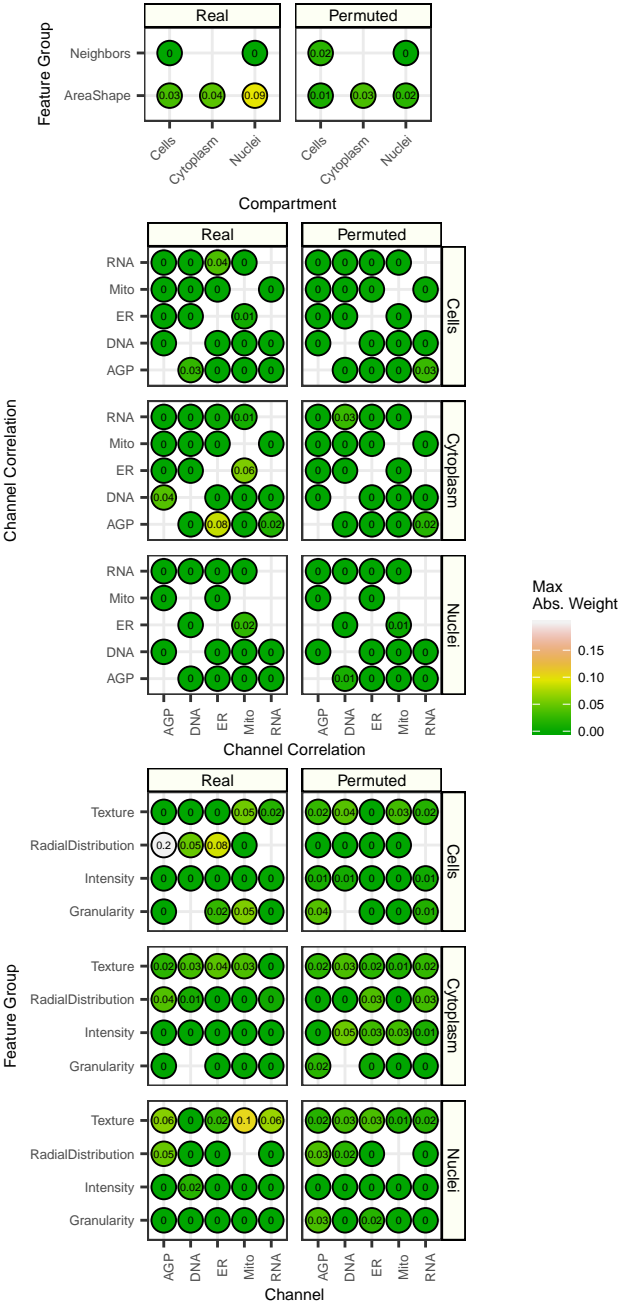
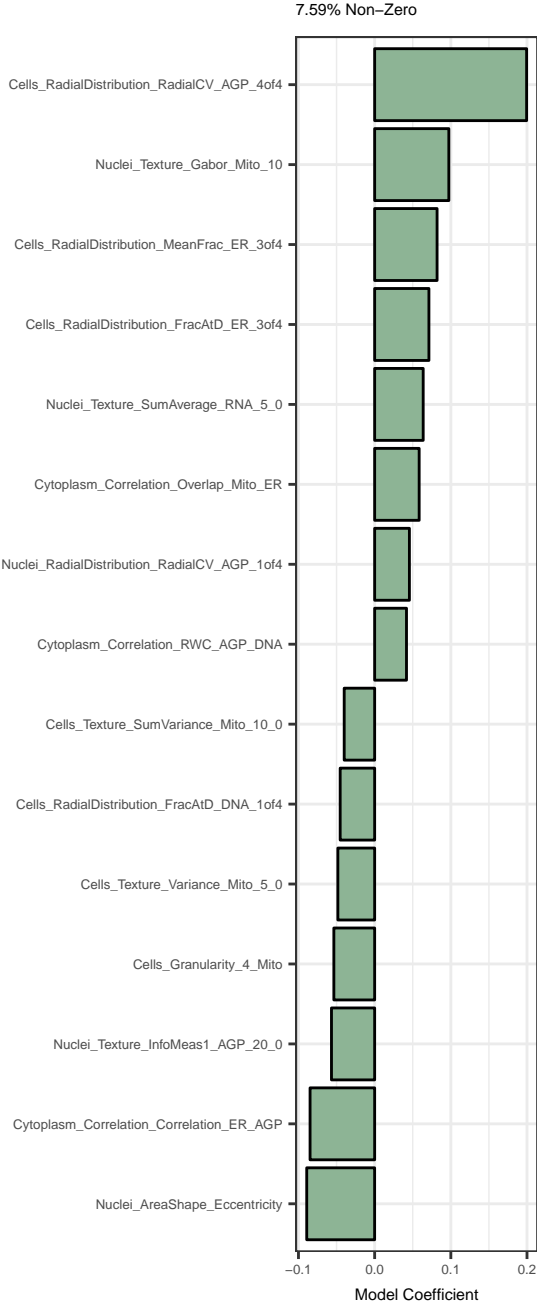
0% Non-Zero



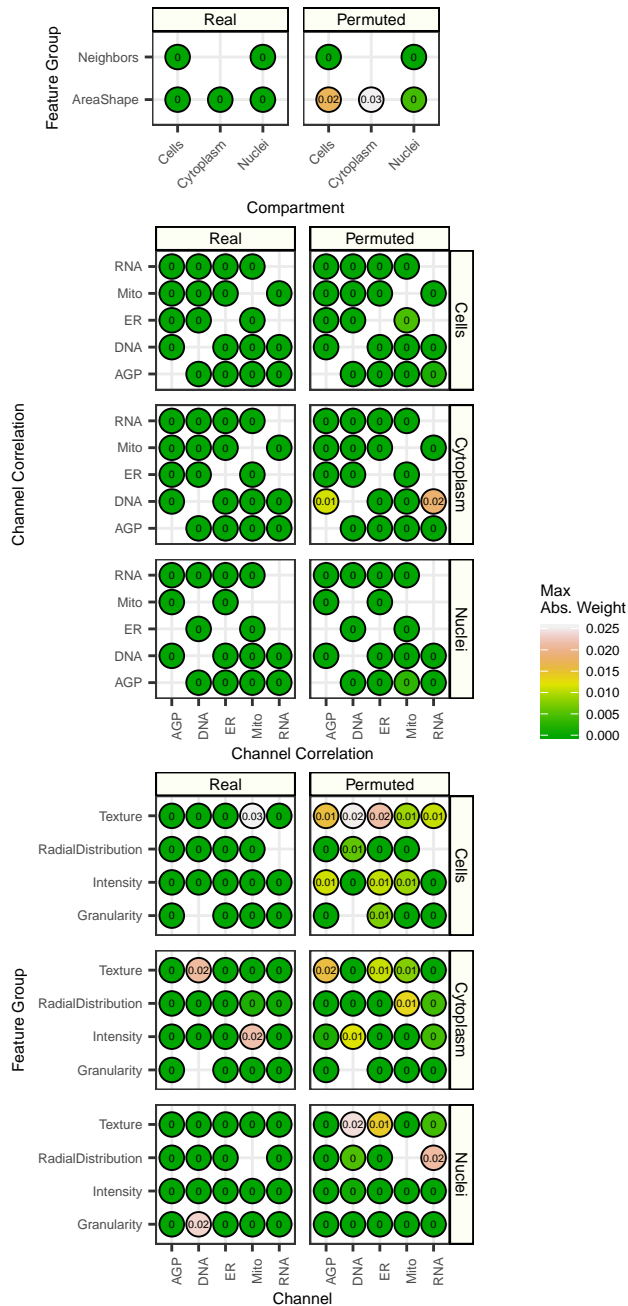
Model Coefficient



Late M – # cells

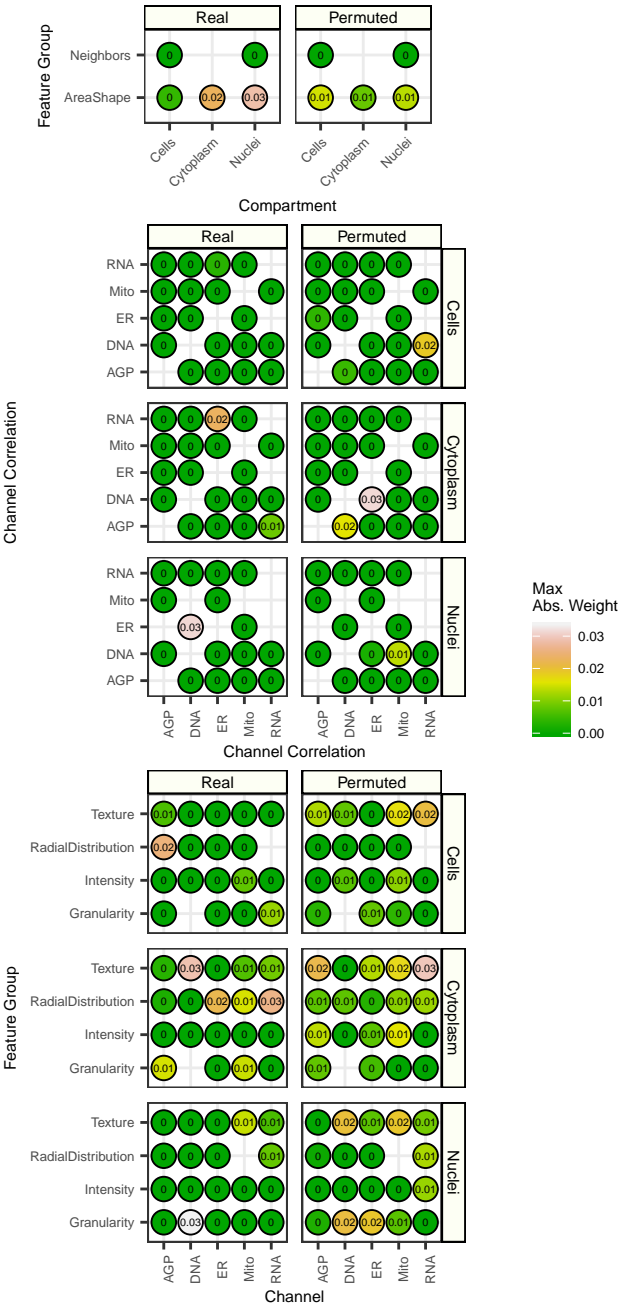
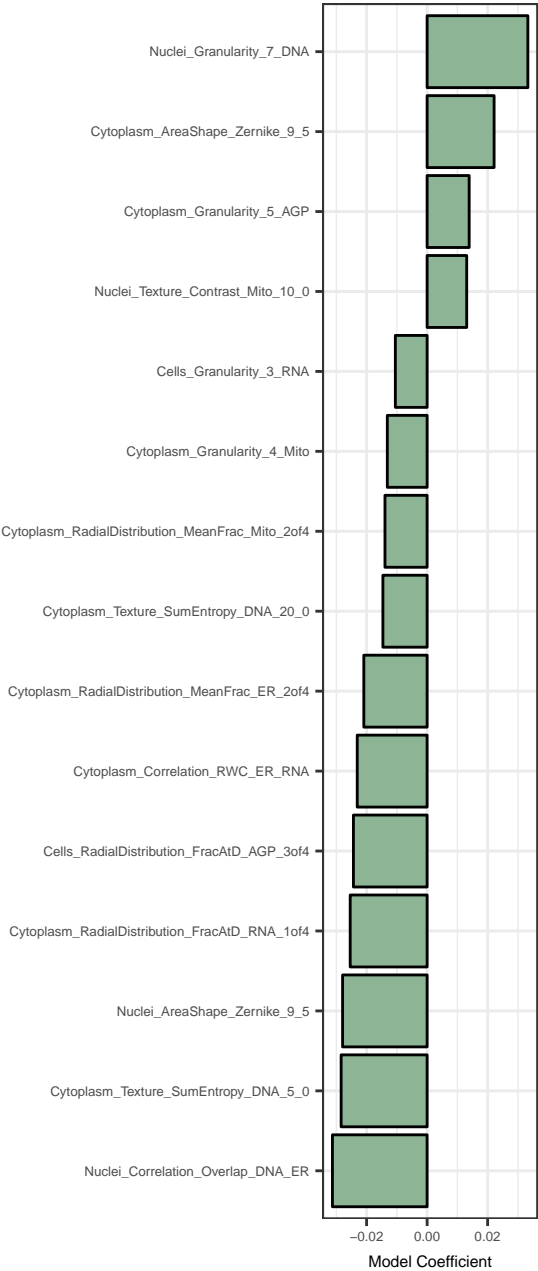


Late M – # of gH2AX Spots



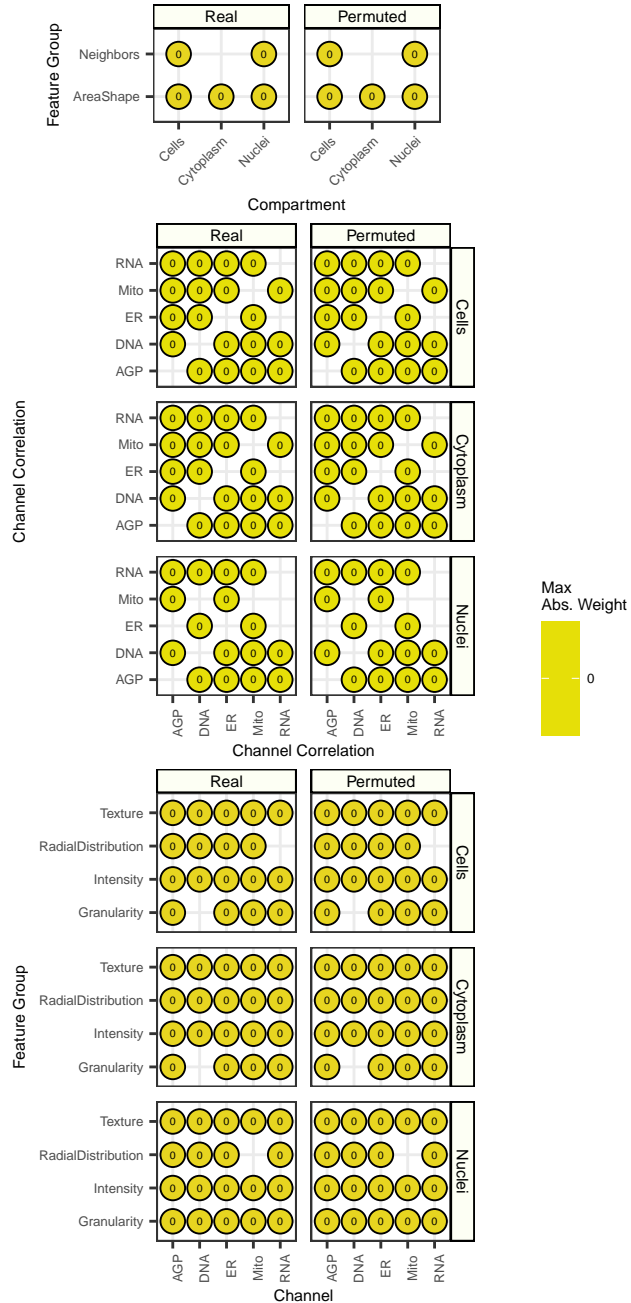
Late M – # of gH2AX Spots per Area of Nucleus

4.21% Non-Zero

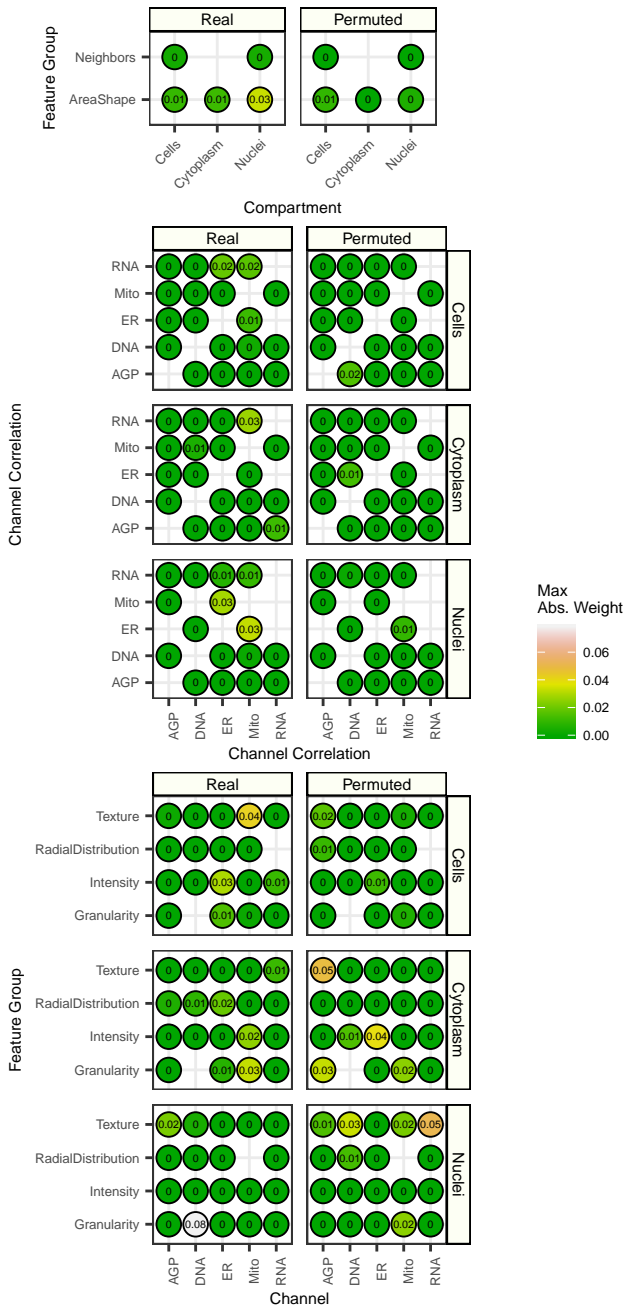
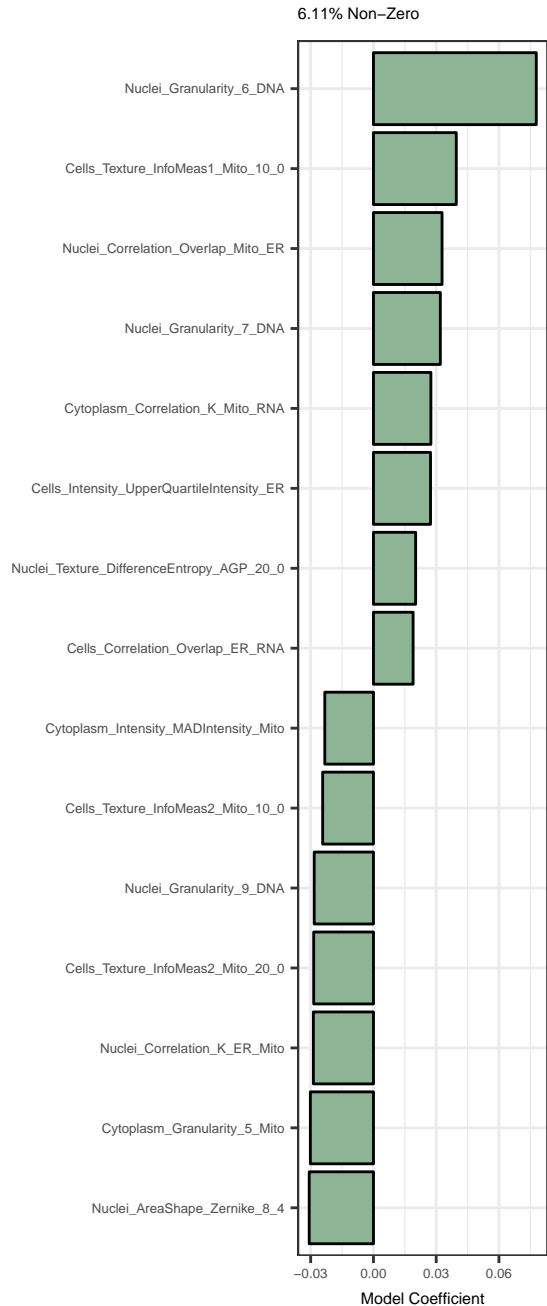


M – % High gH2AX Spots

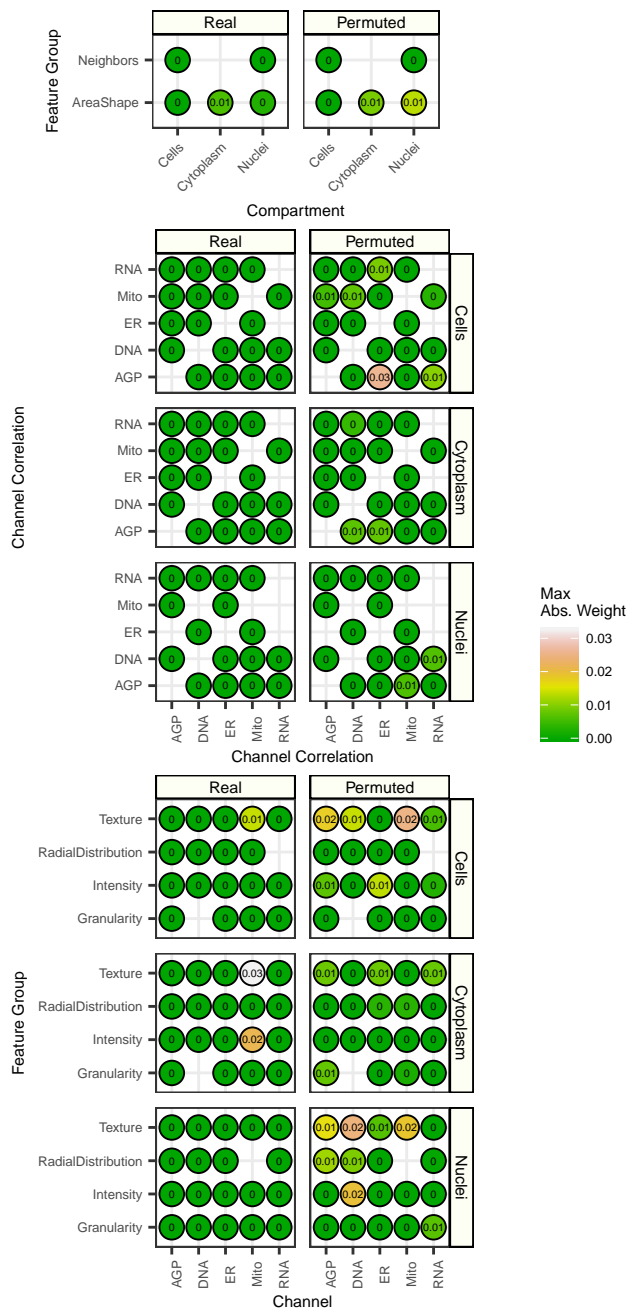
0% Non-Zero



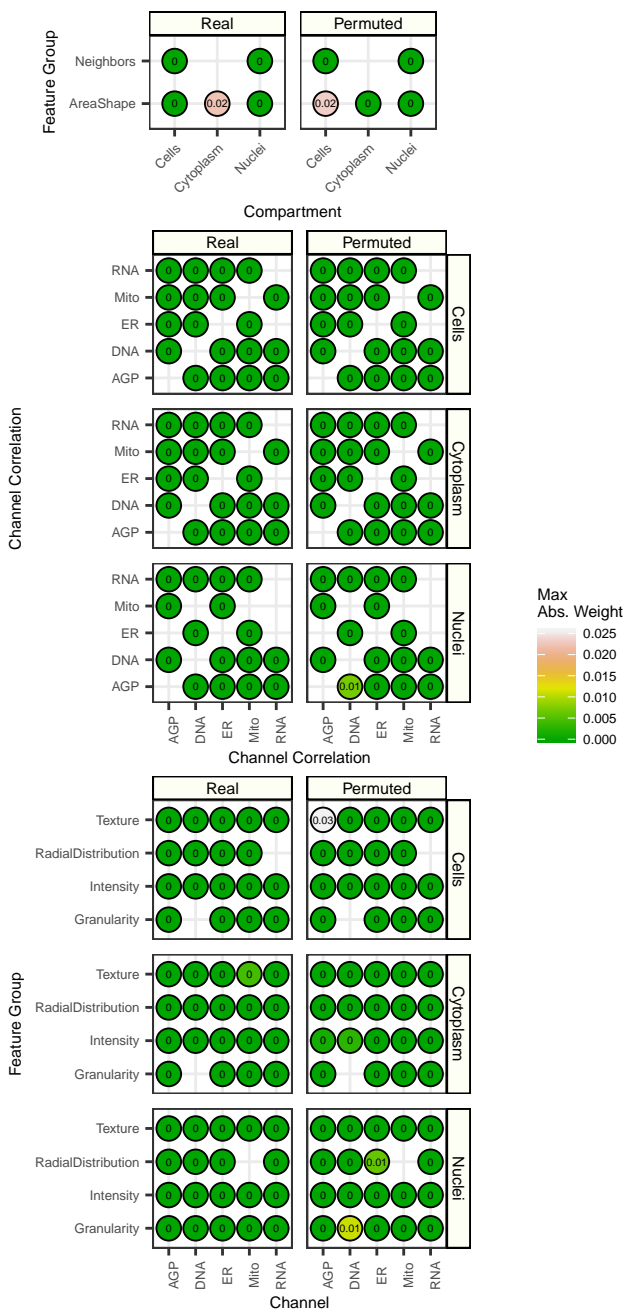
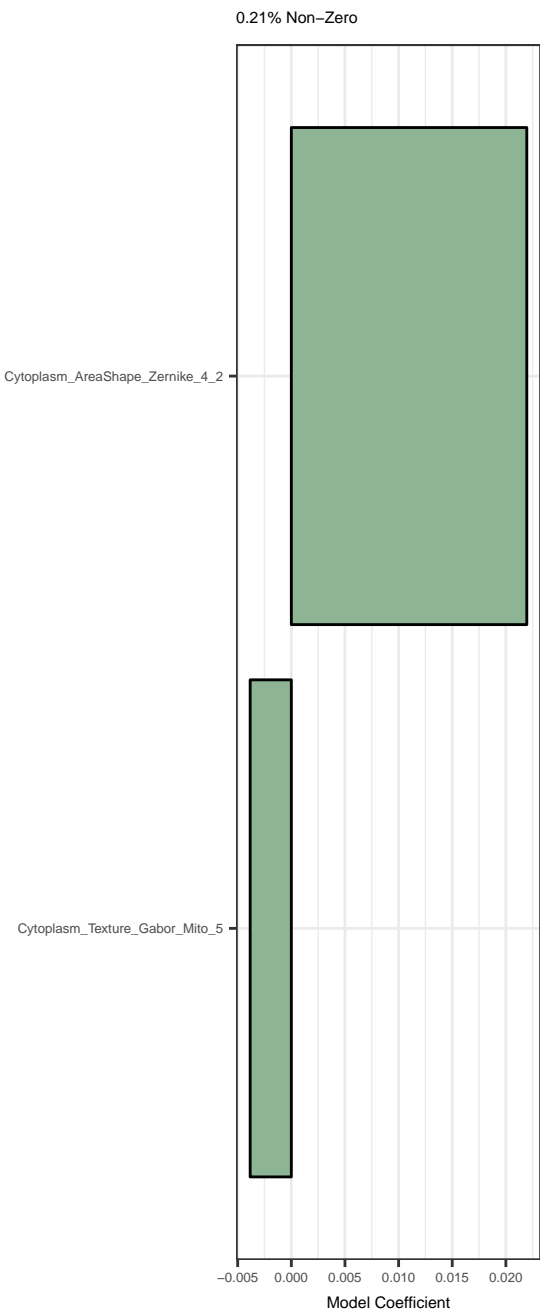
M – # cells



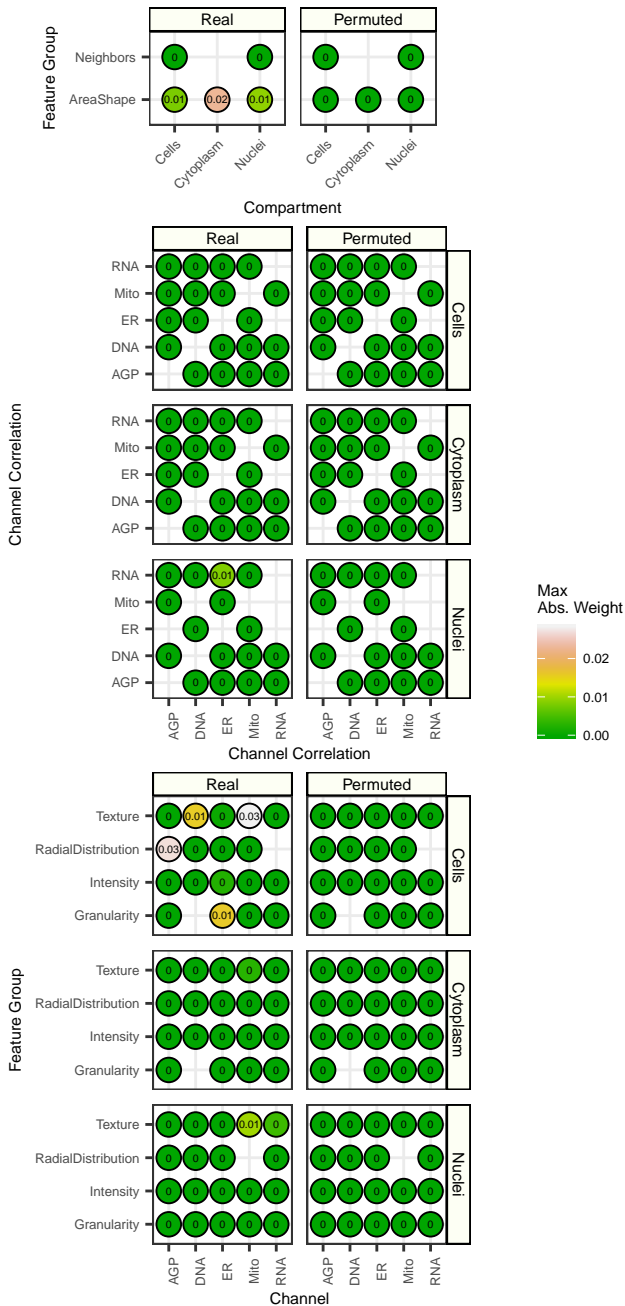
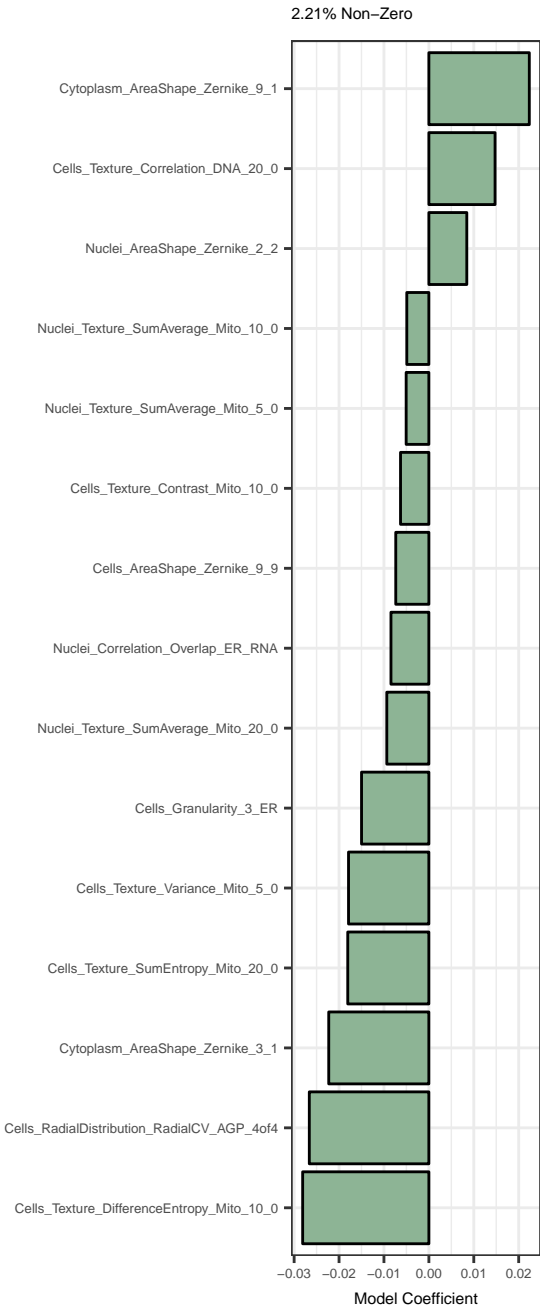
M – # of gH2AX Spots



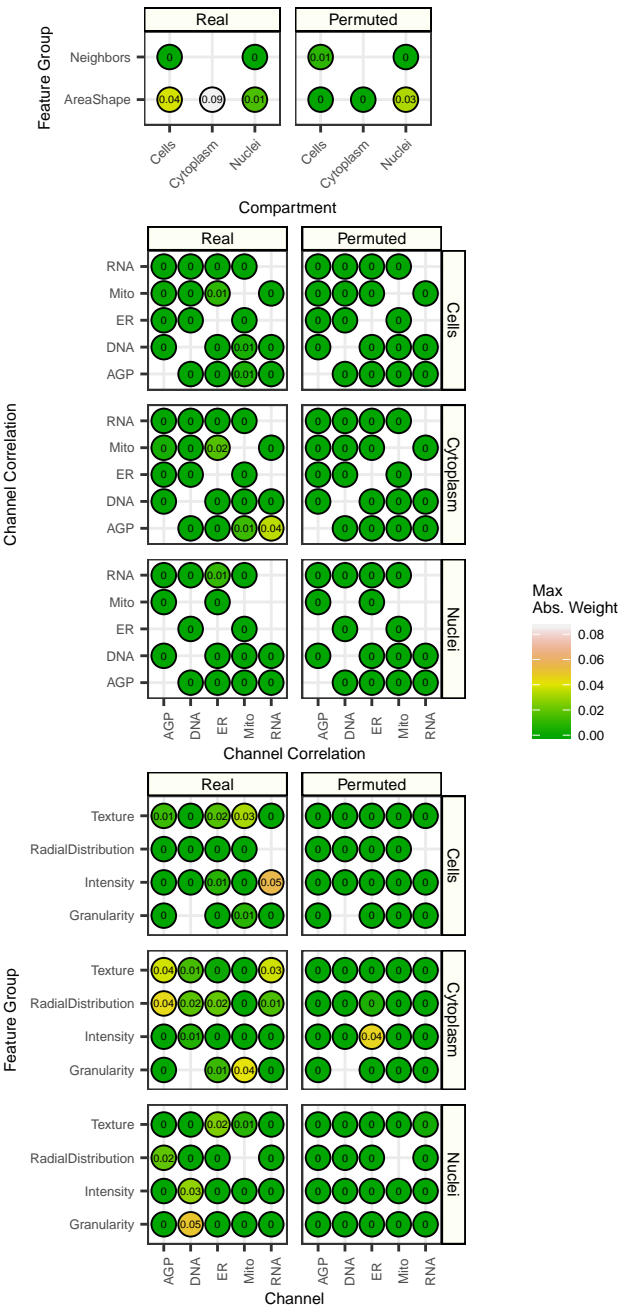
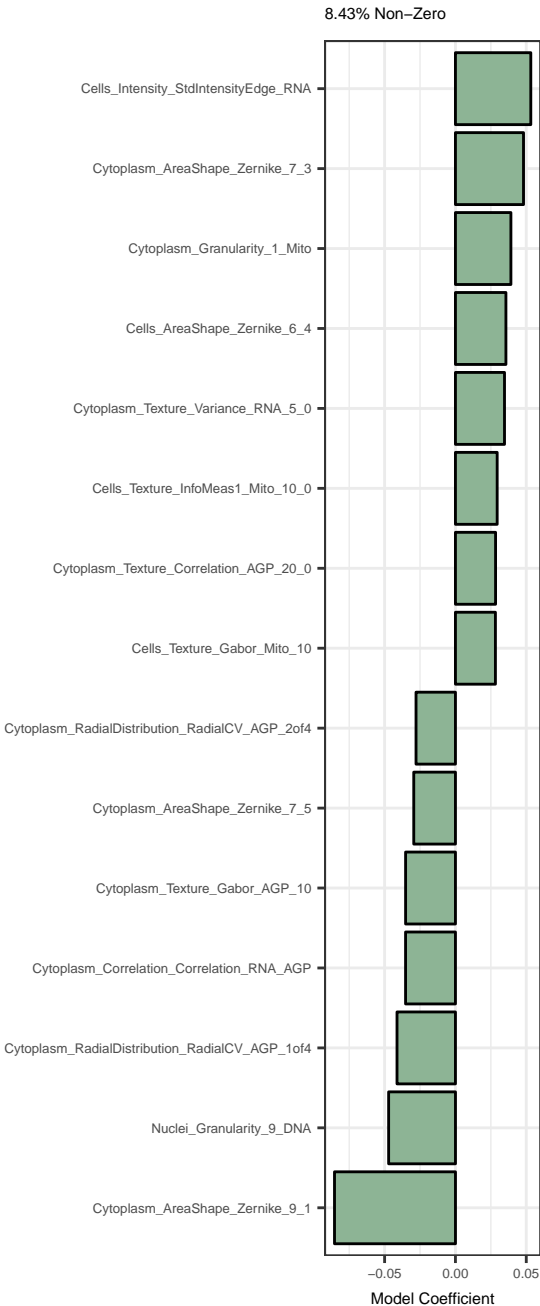
M – # of gH2AX Spots per Area of Nucleus



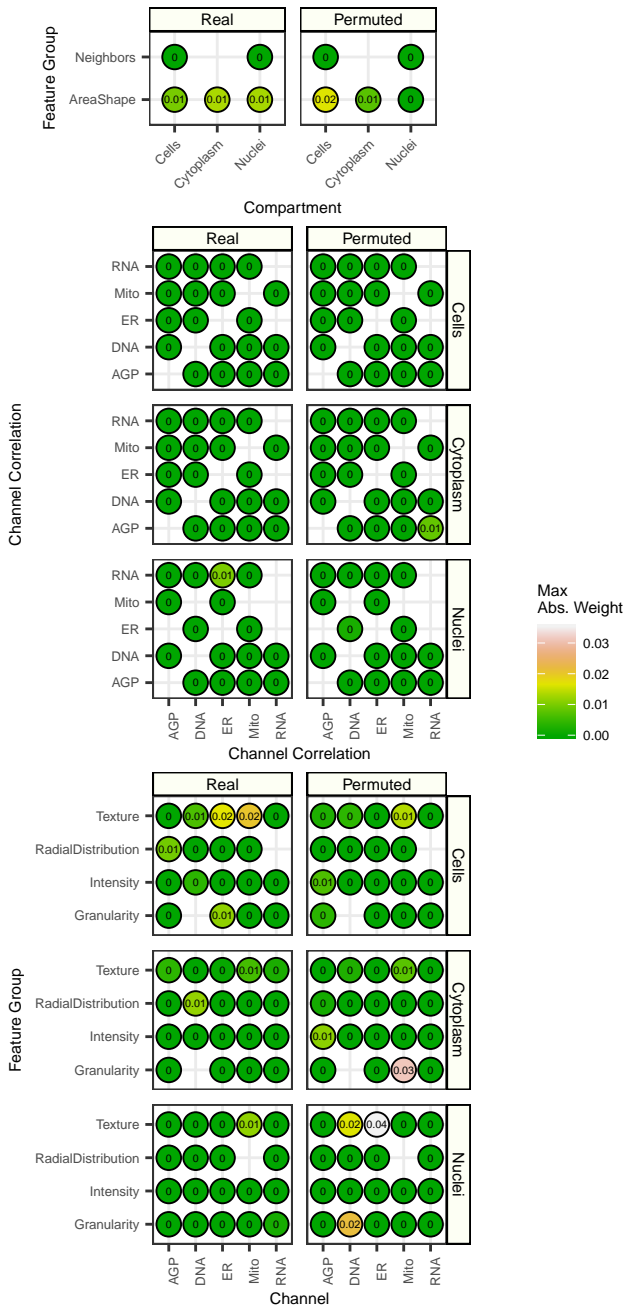
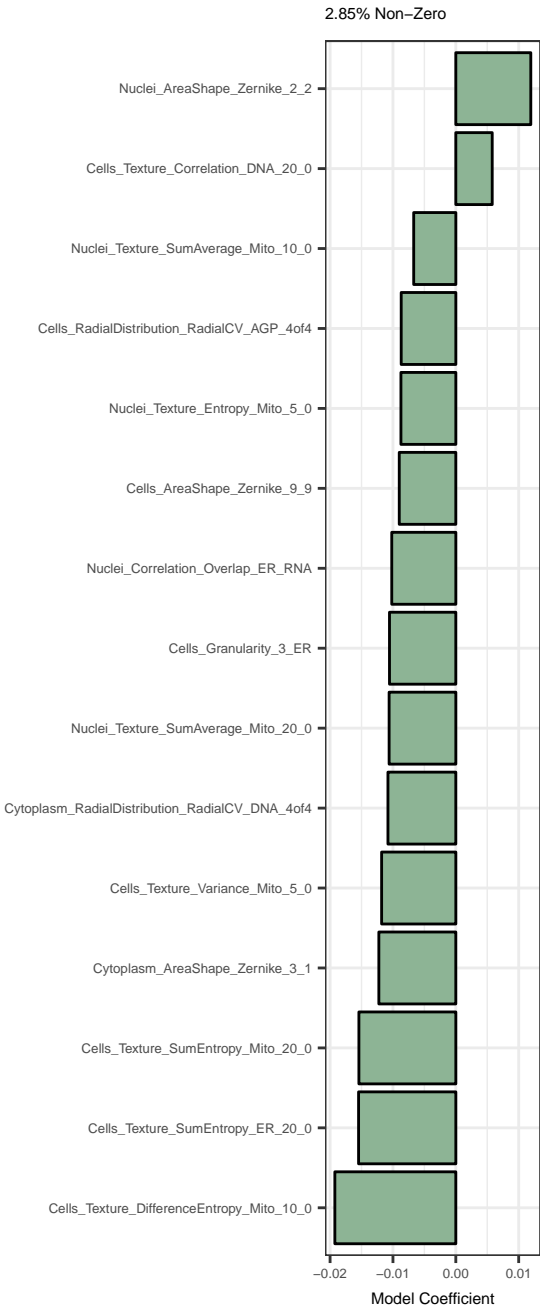
Polynuclear – % High gH2AX Spots



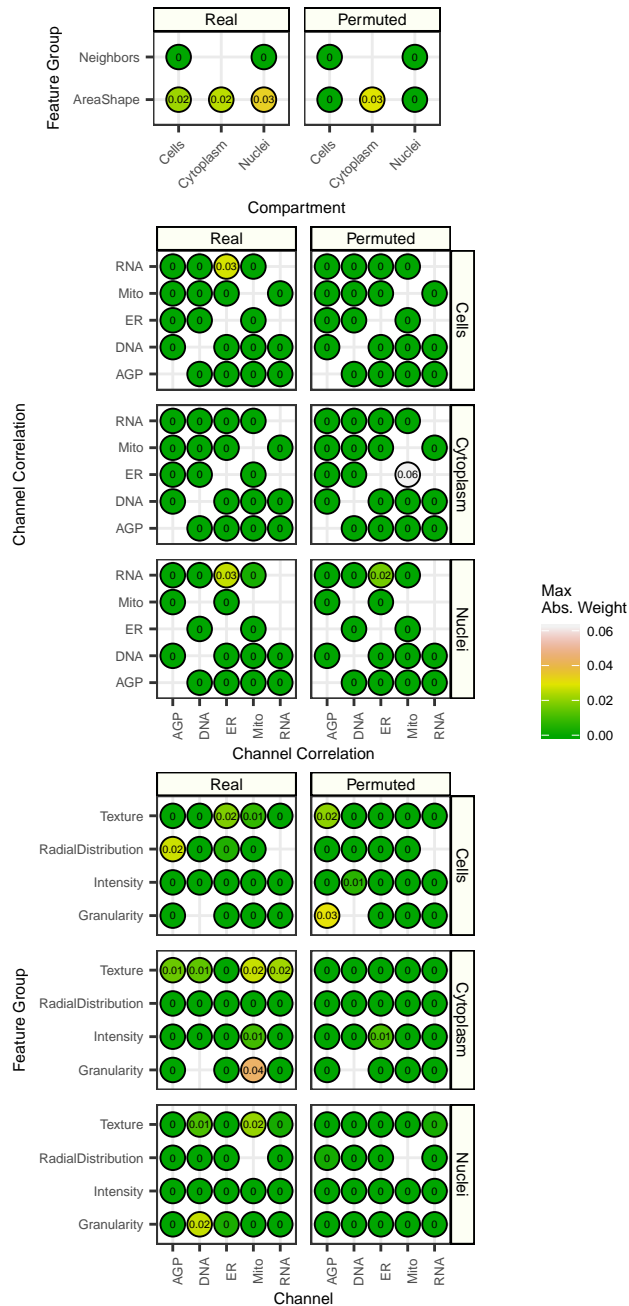
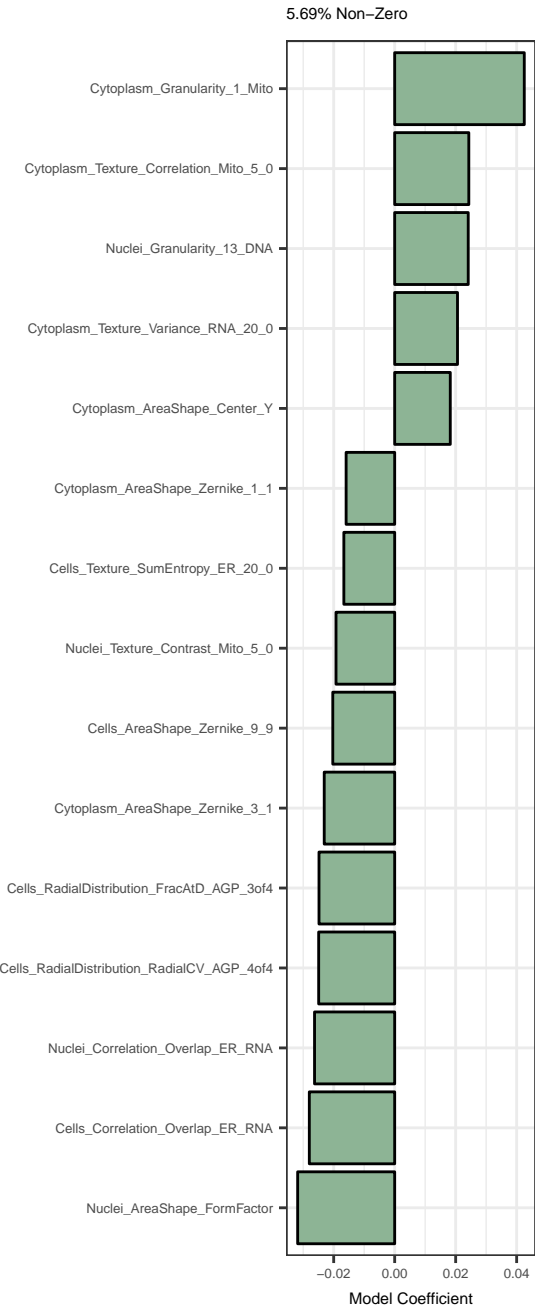
Polynuclear – # cells



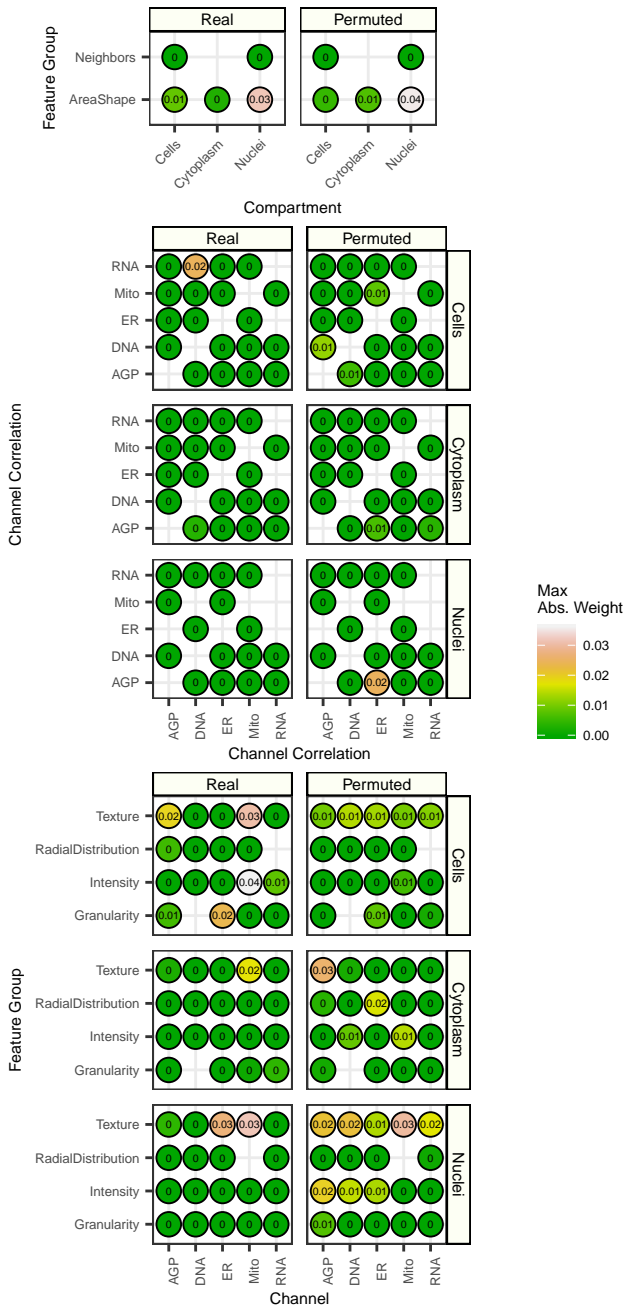
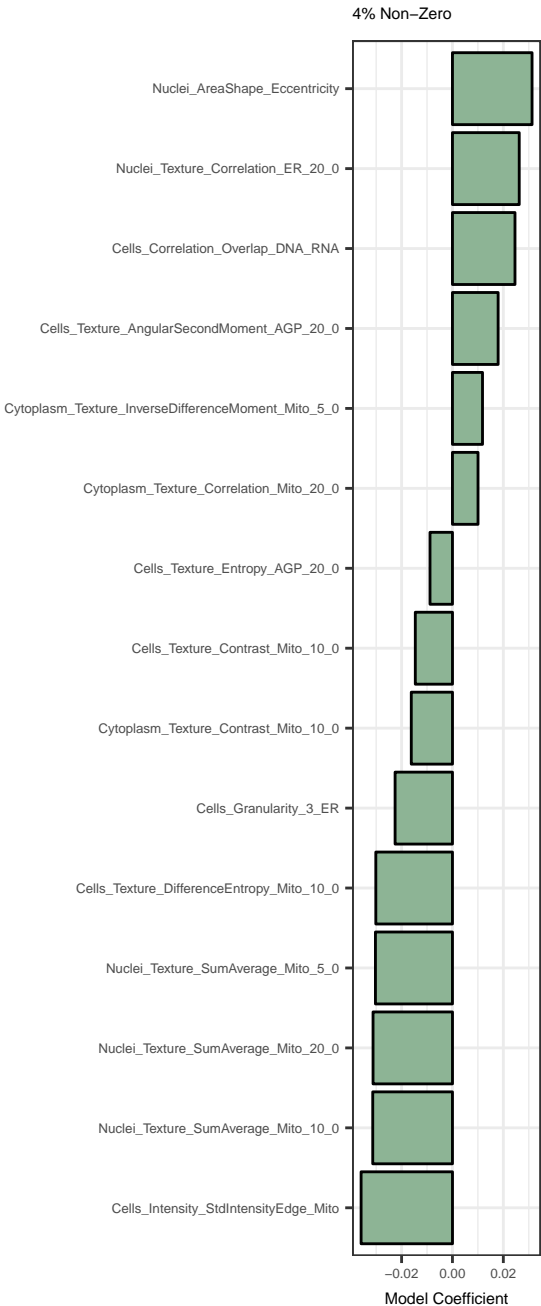
Polynuclear – # of gH2AX Spots



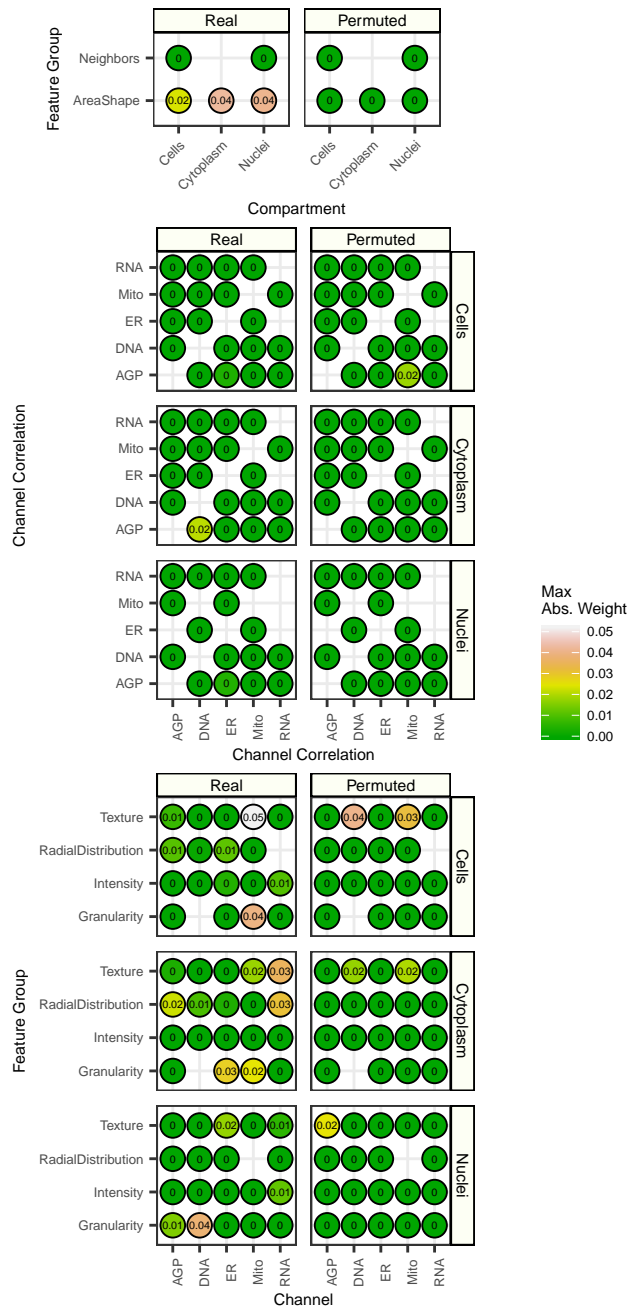
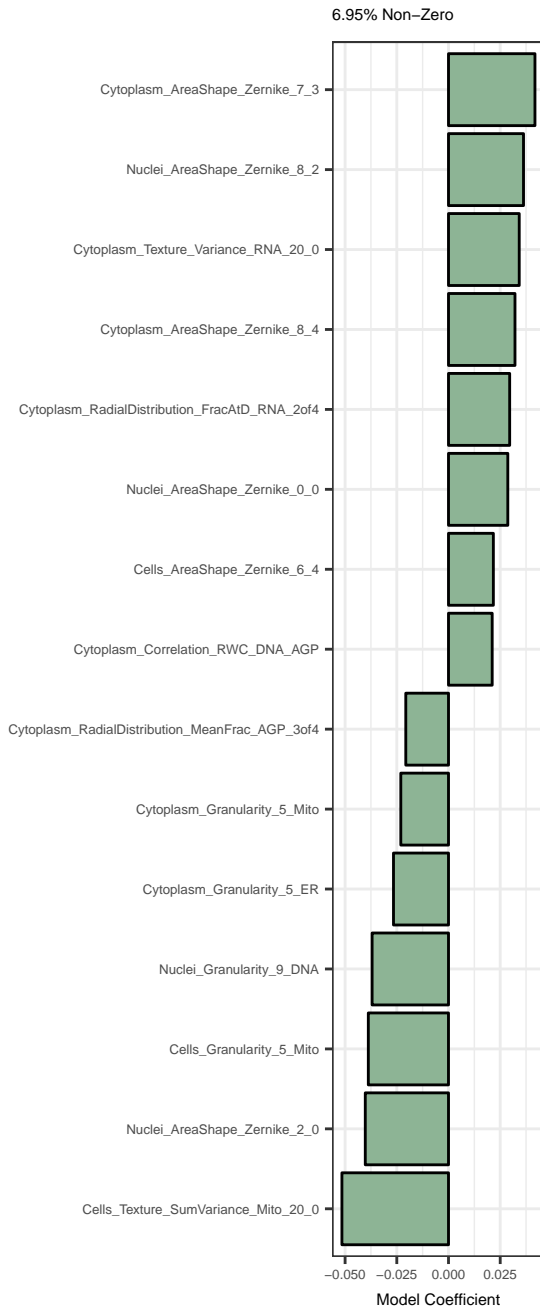
Polynuclear – # of gH2AX Spots per Area of Nucleus



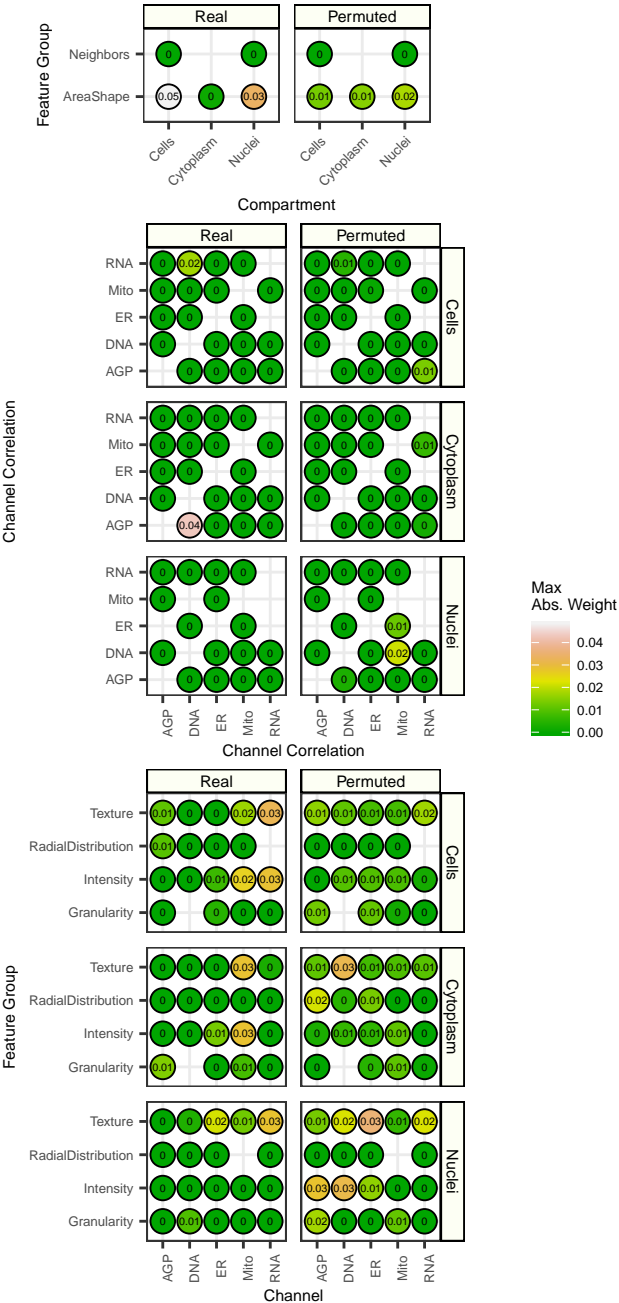
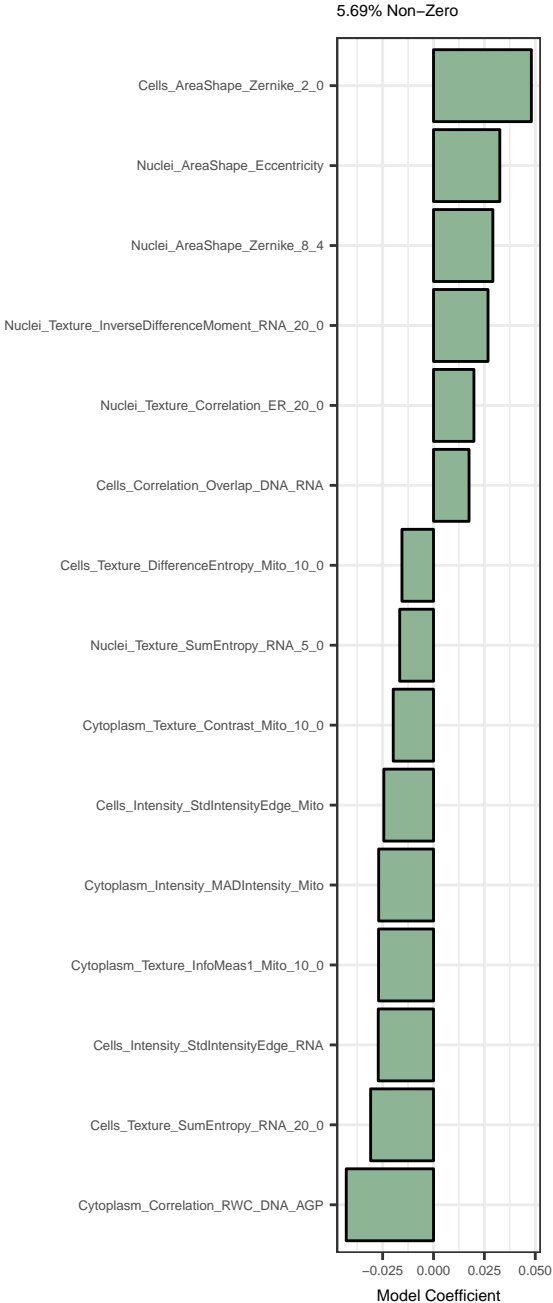
Polyploid – % High gH2AX Spots



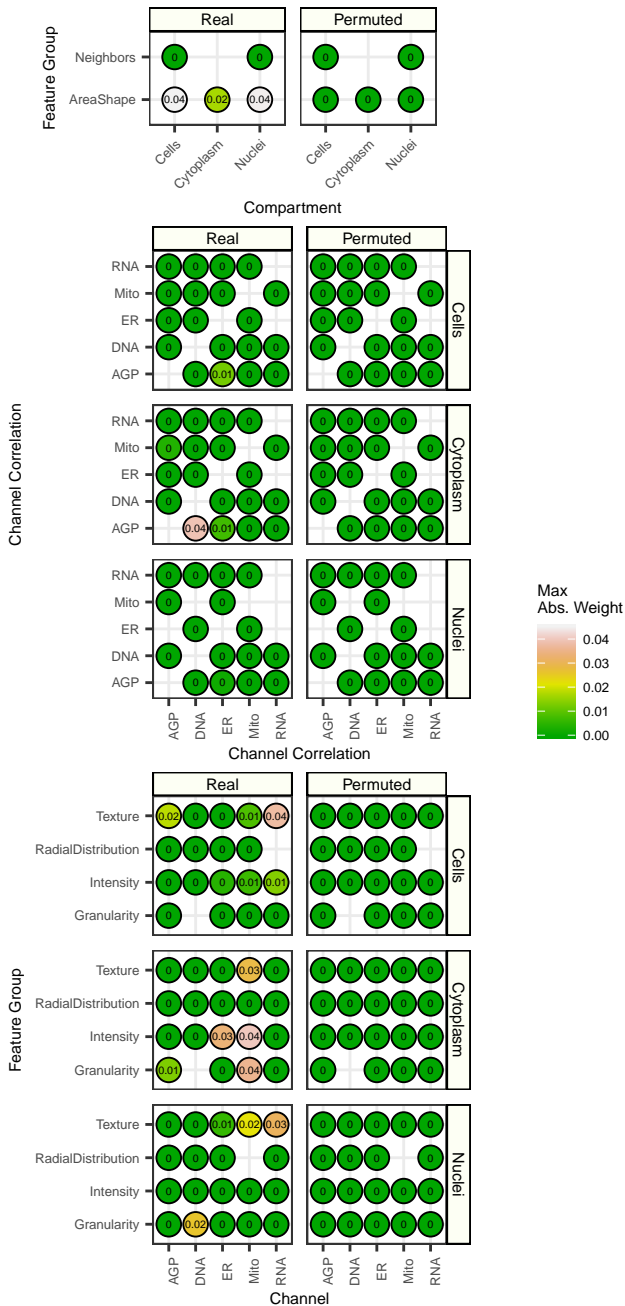
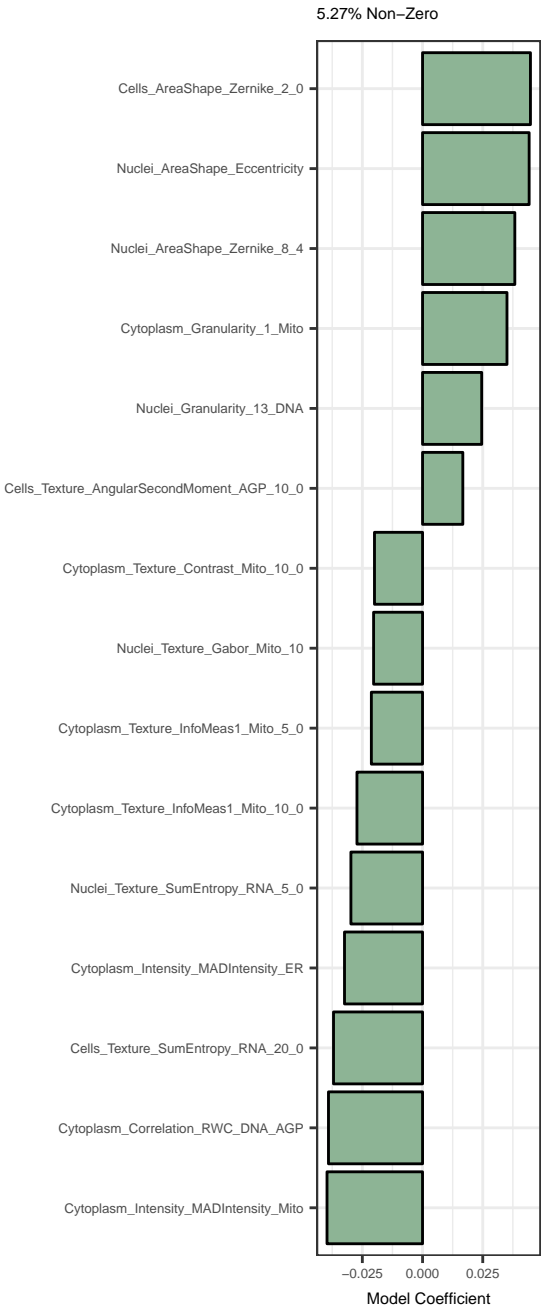
Polyploid – # cells



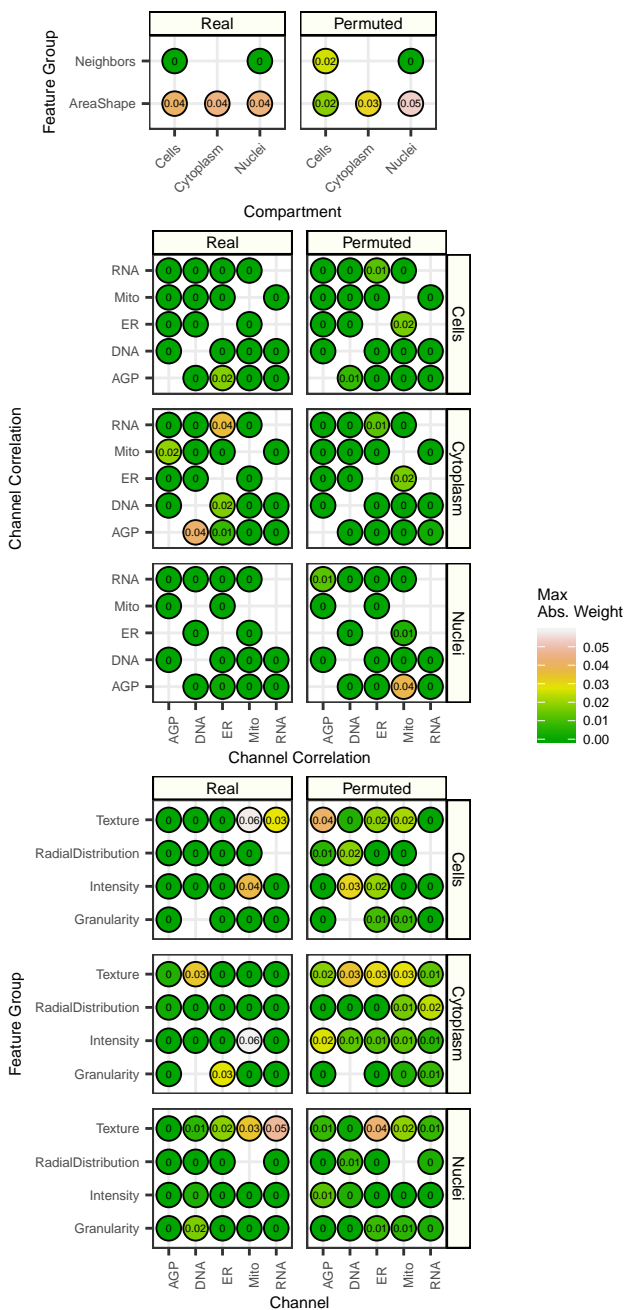
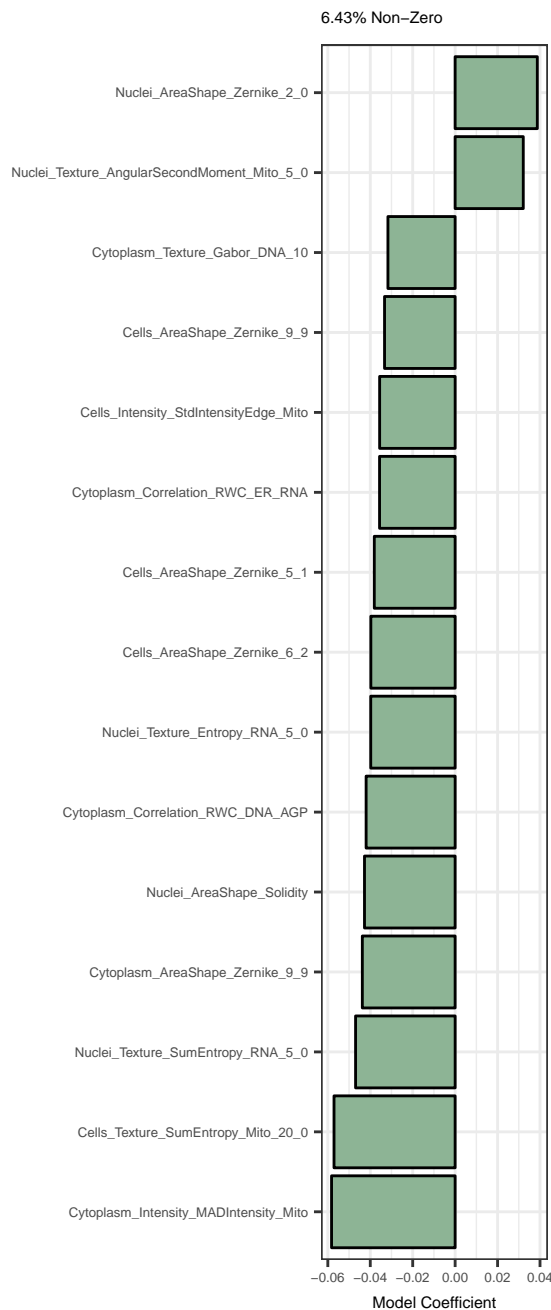
Polyloid – # of gH2AX Spots



Polyploid – # of gH2AX Spots per Area of Nucleus

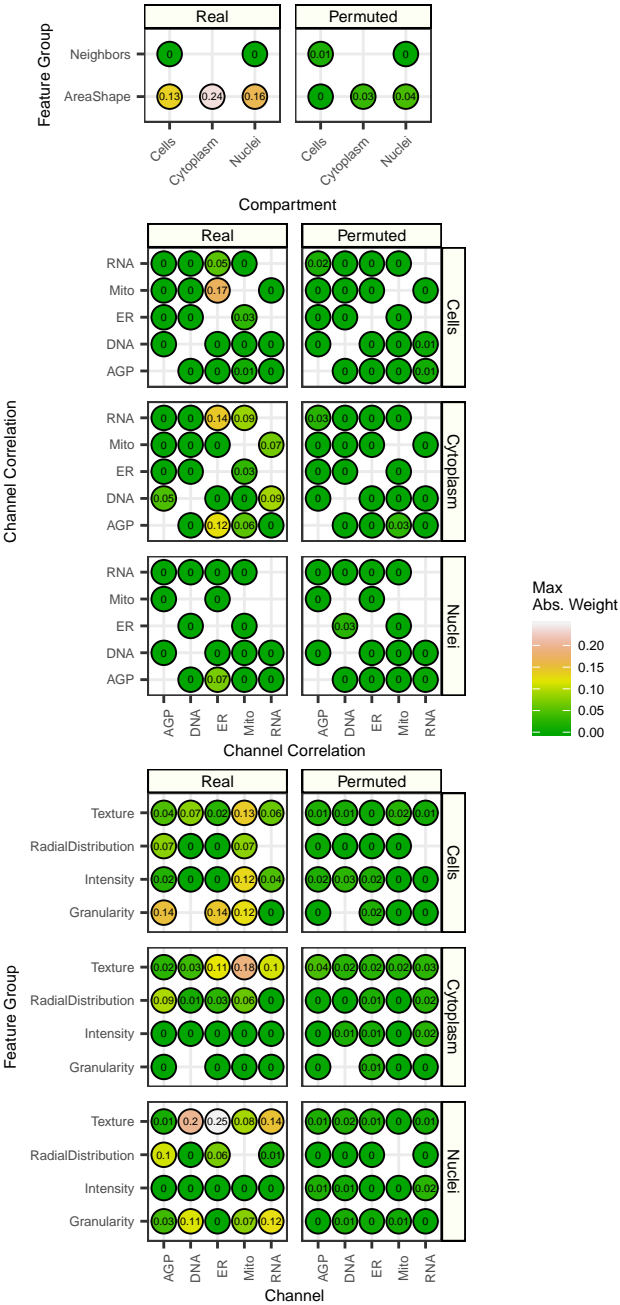
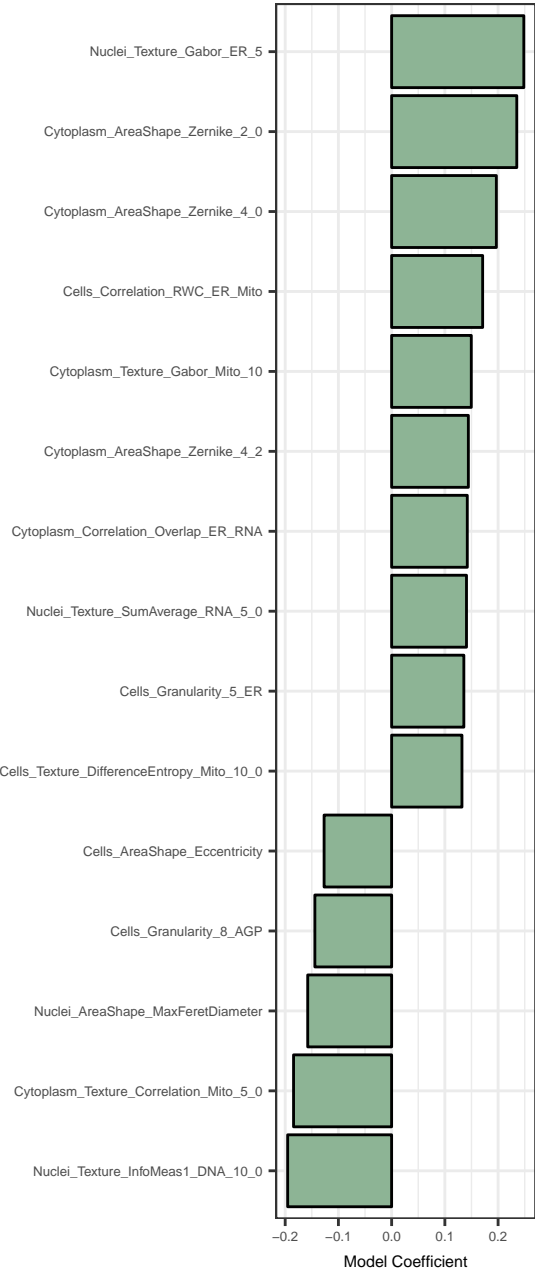


S – % High gH2AX Spots

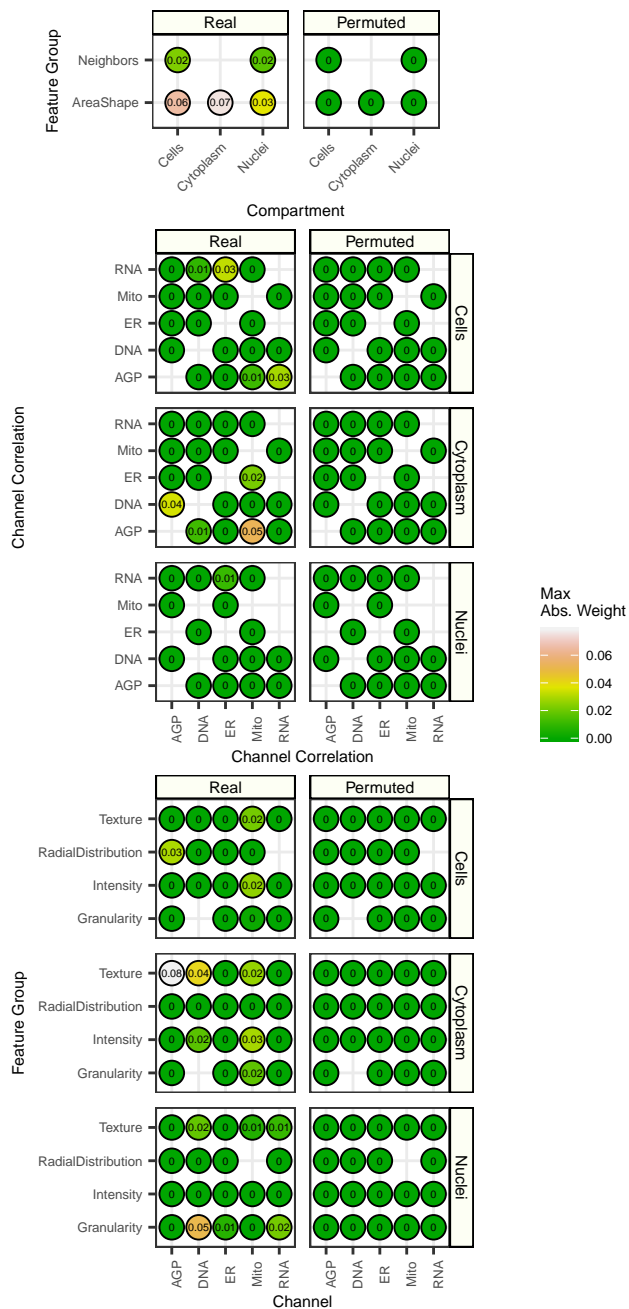
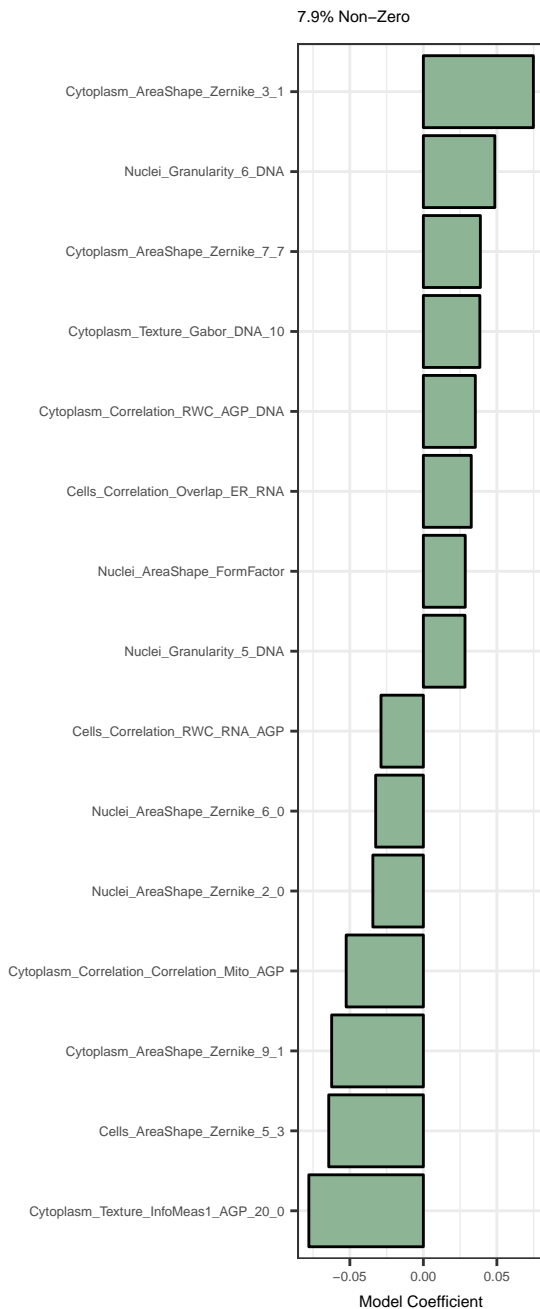


S – Intensity Nucleus EdU Mean

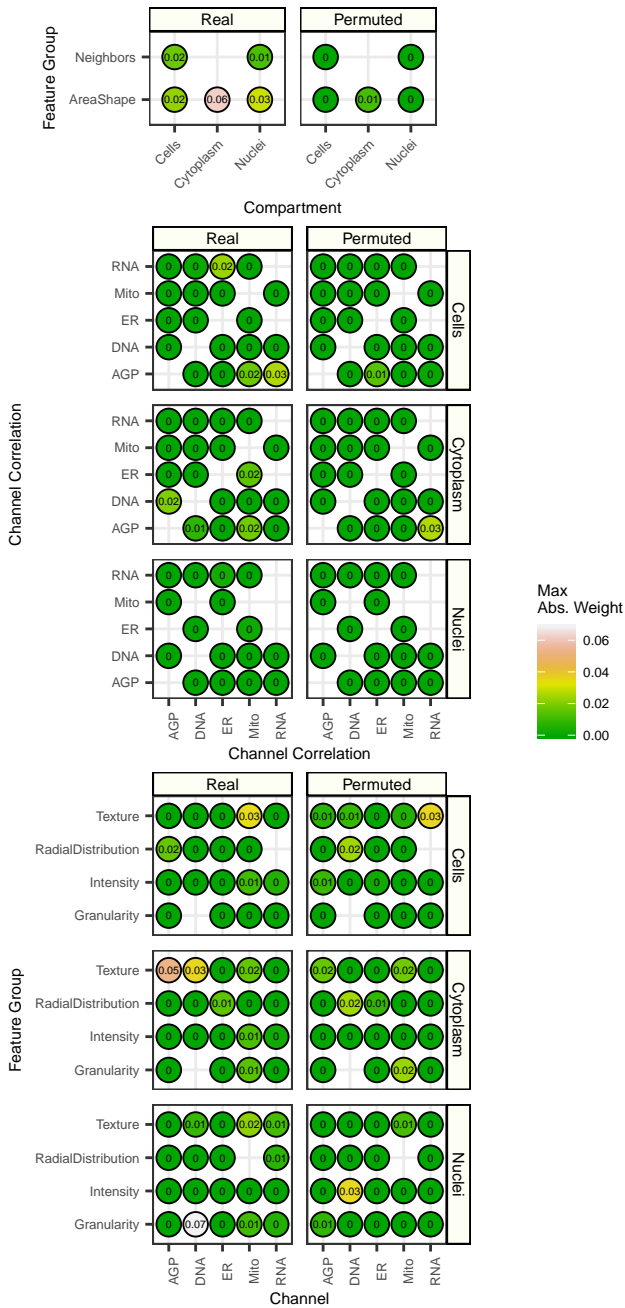
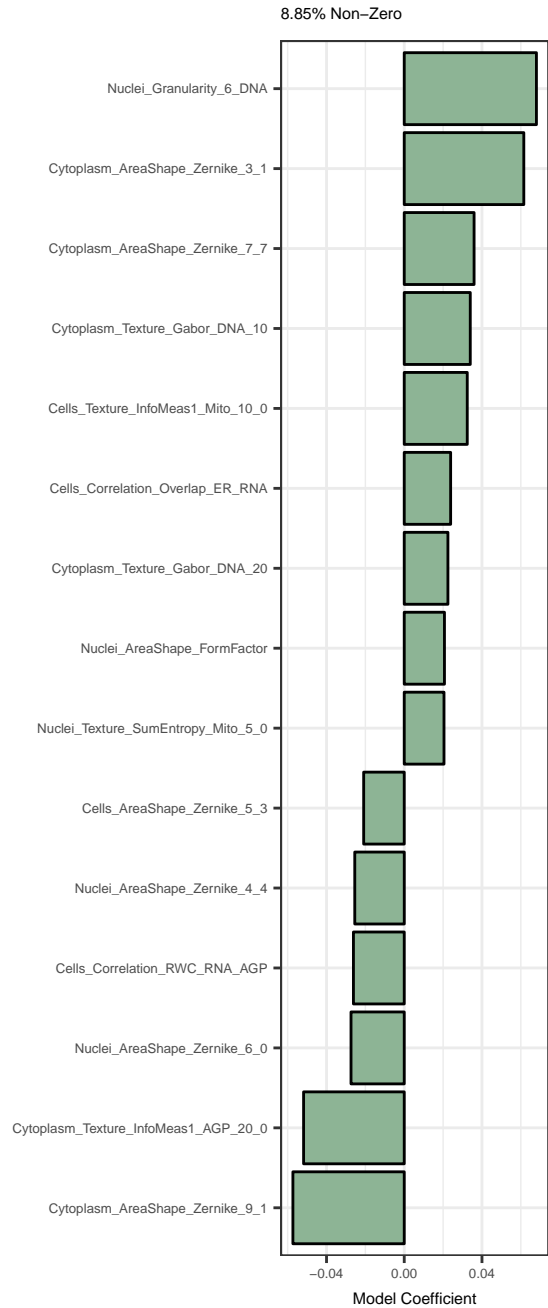
11.28% Non-Zero



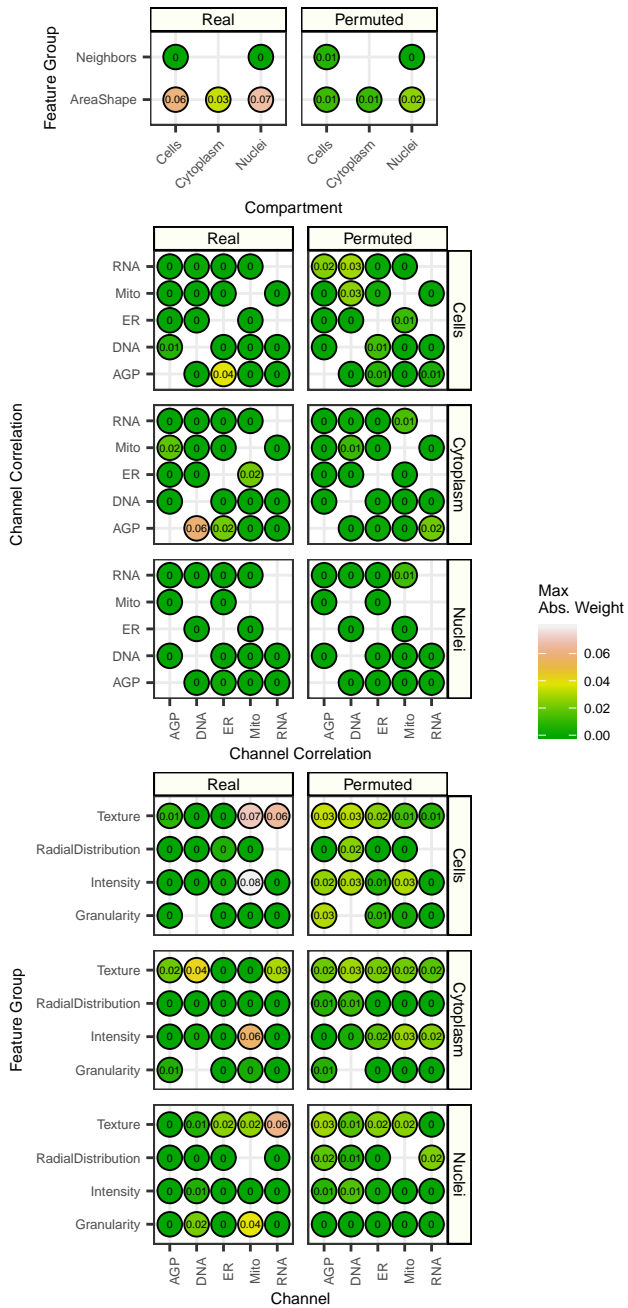
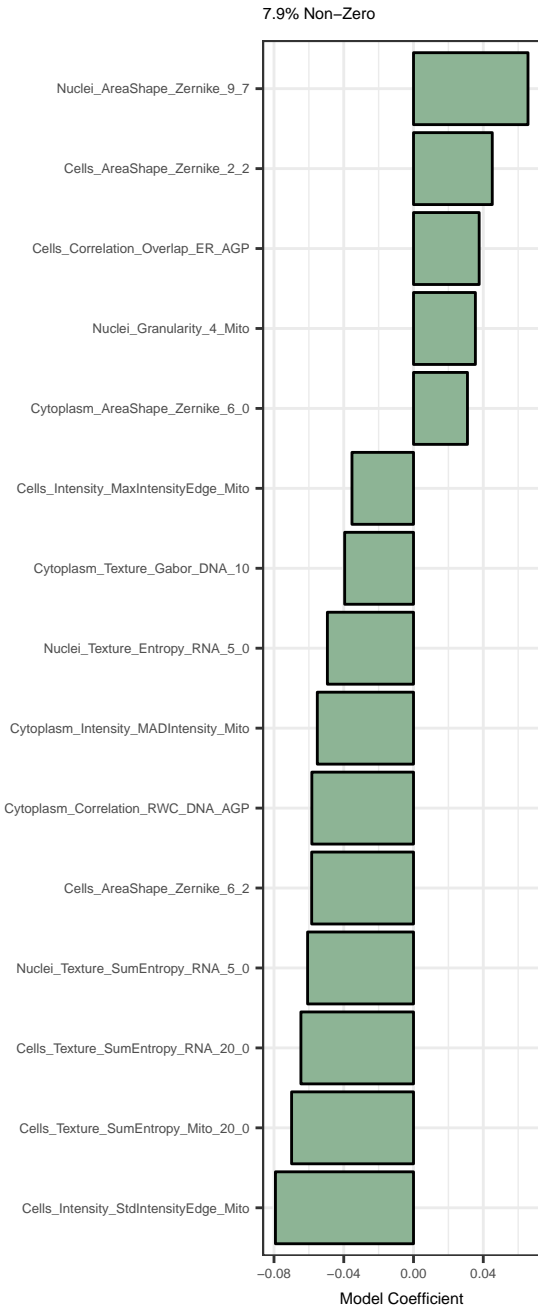
S – Intensity Nucleus EdU Sum



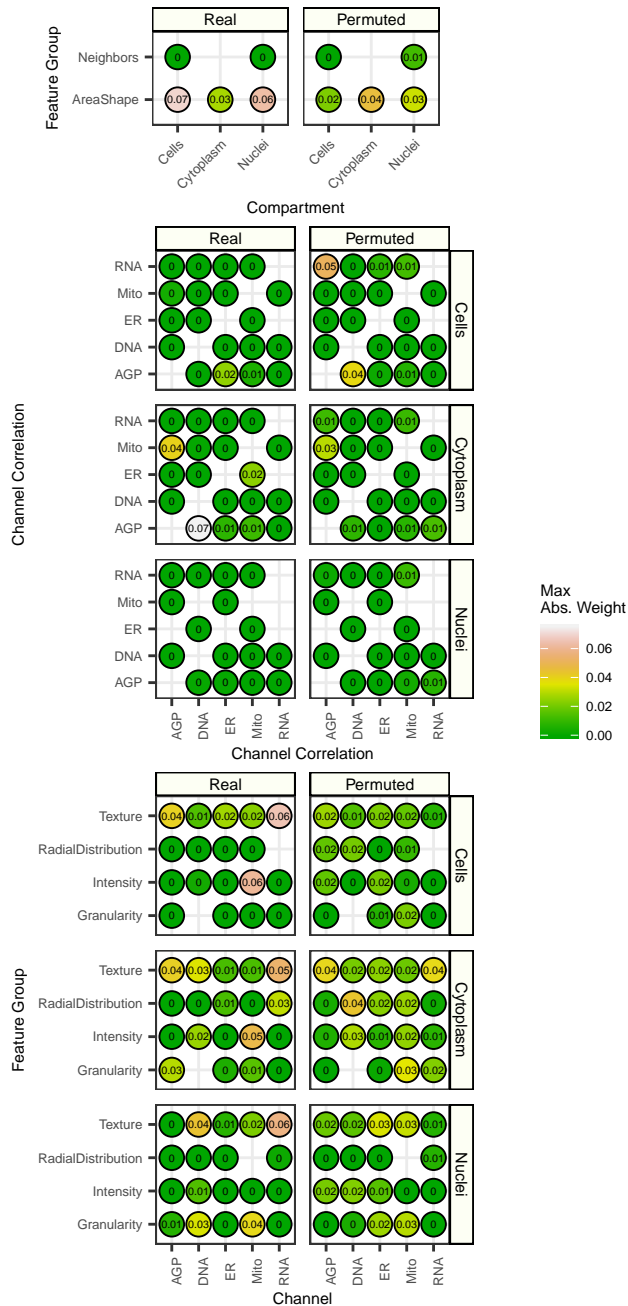
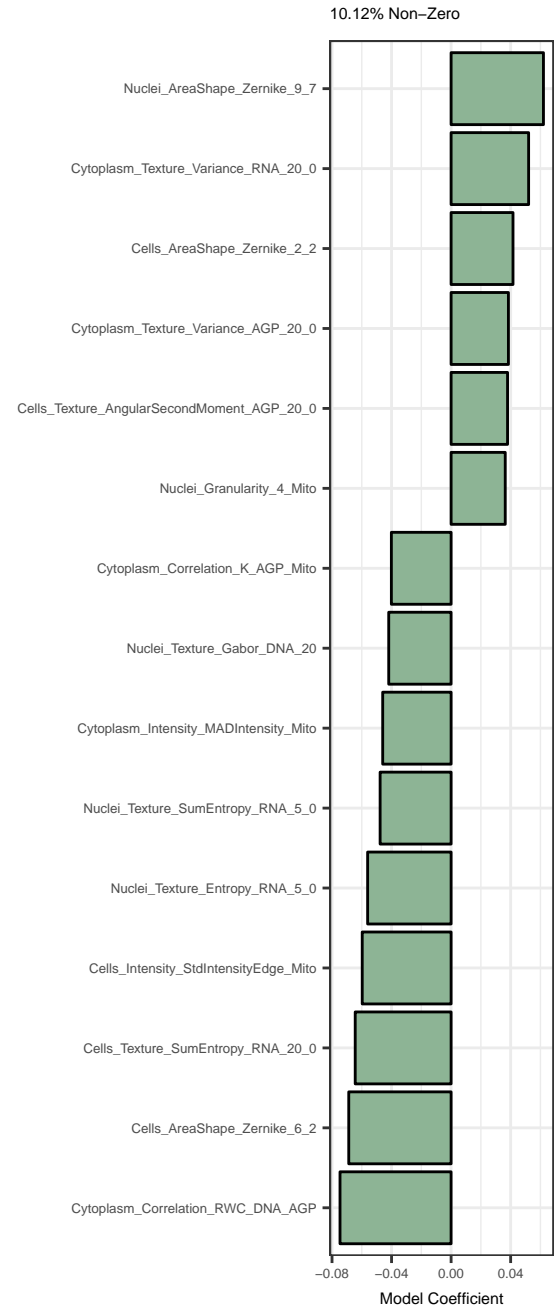
S – # cells



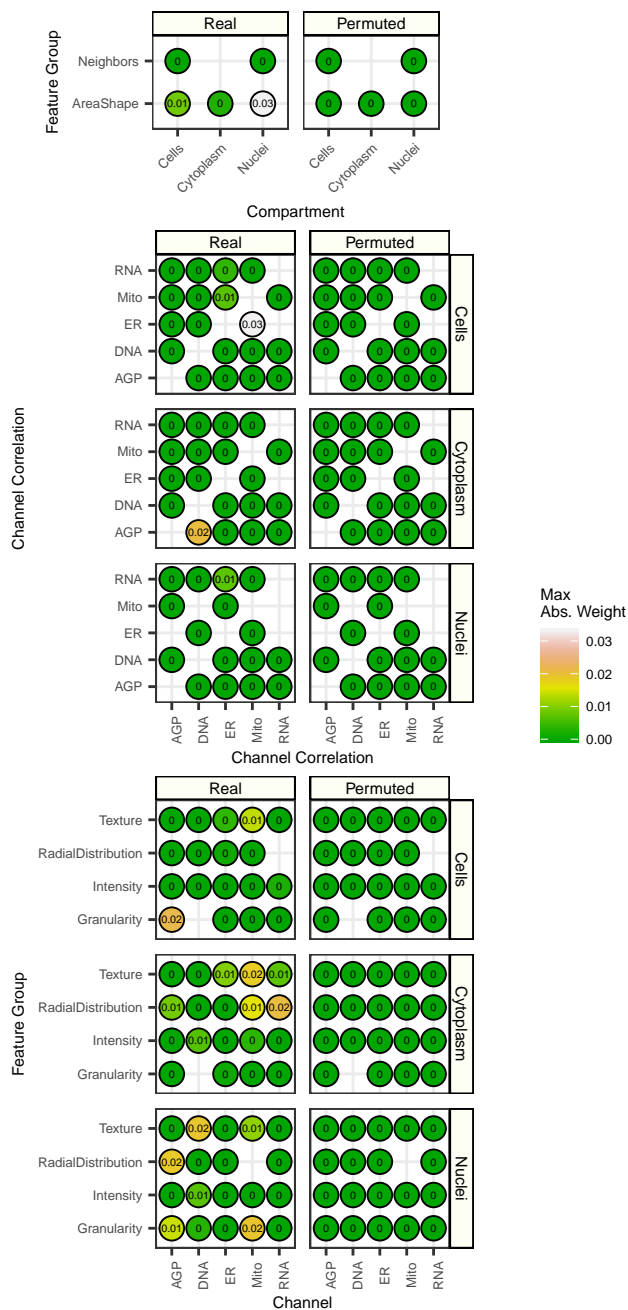
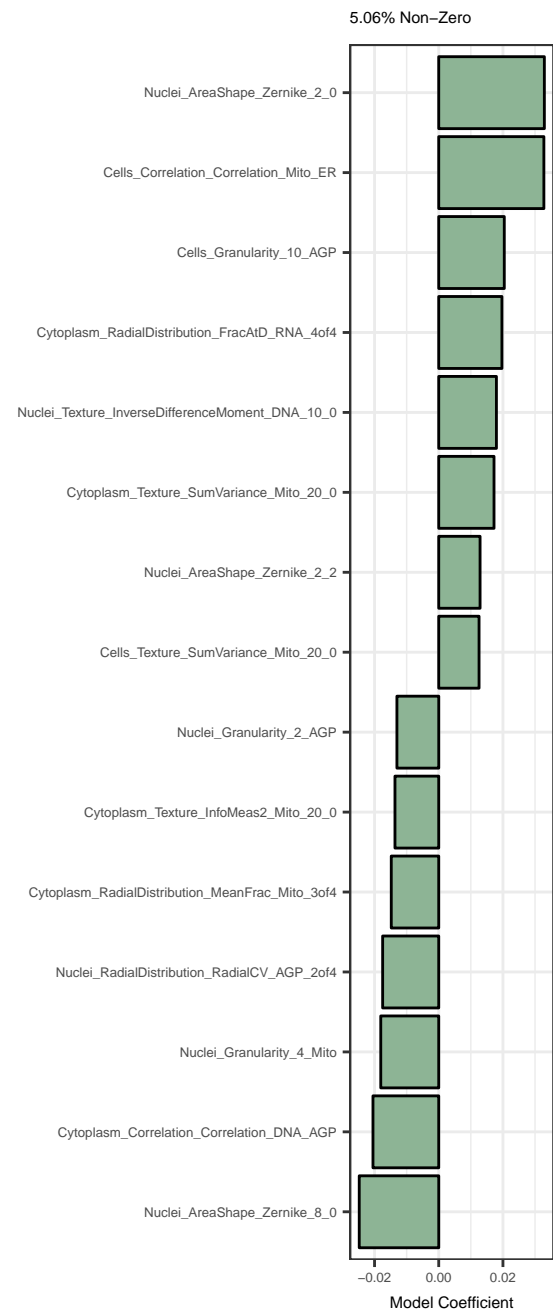
S – # of gH2AX Spots



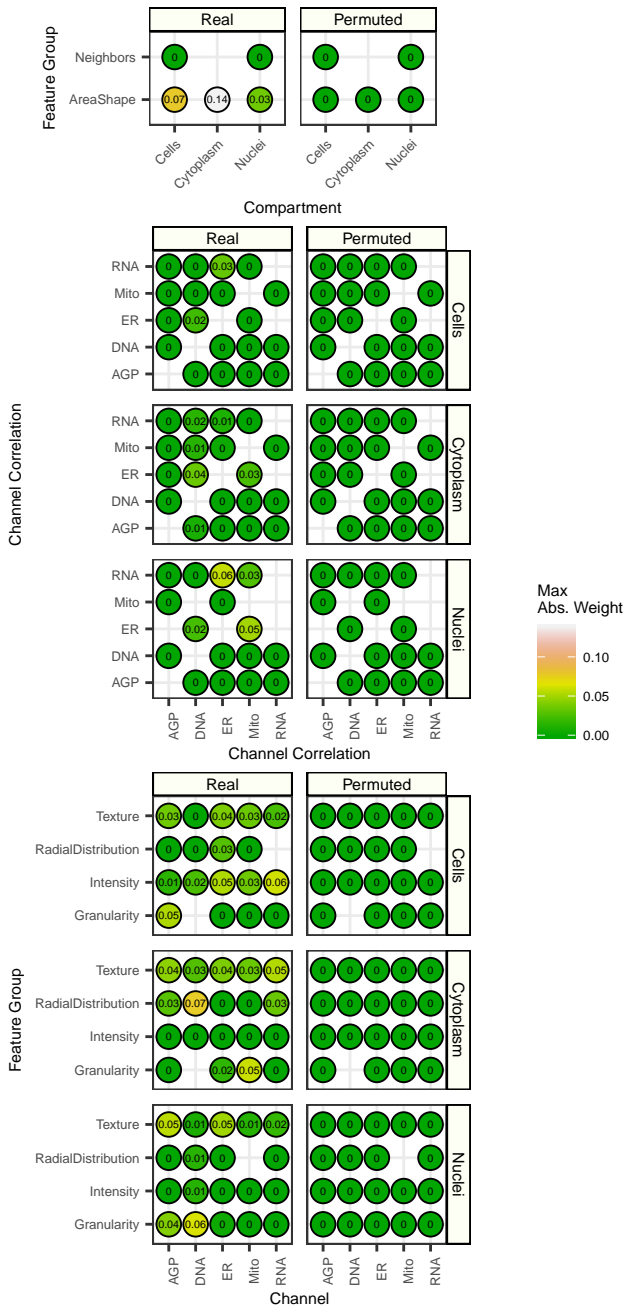
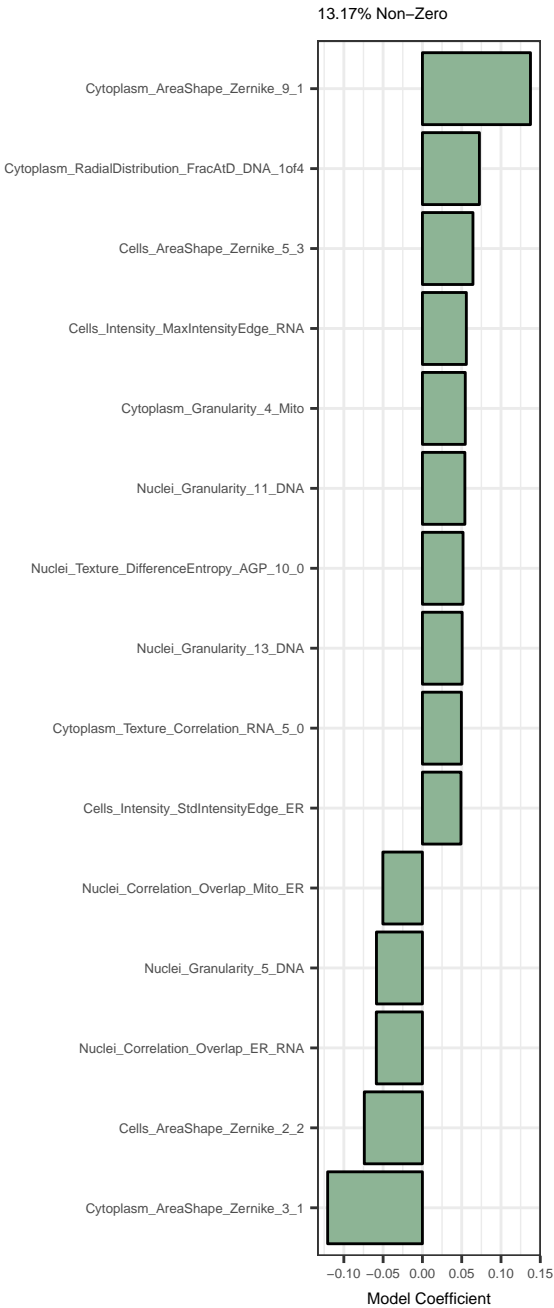
S – # of gH2AX Spots per Area of Nucleus



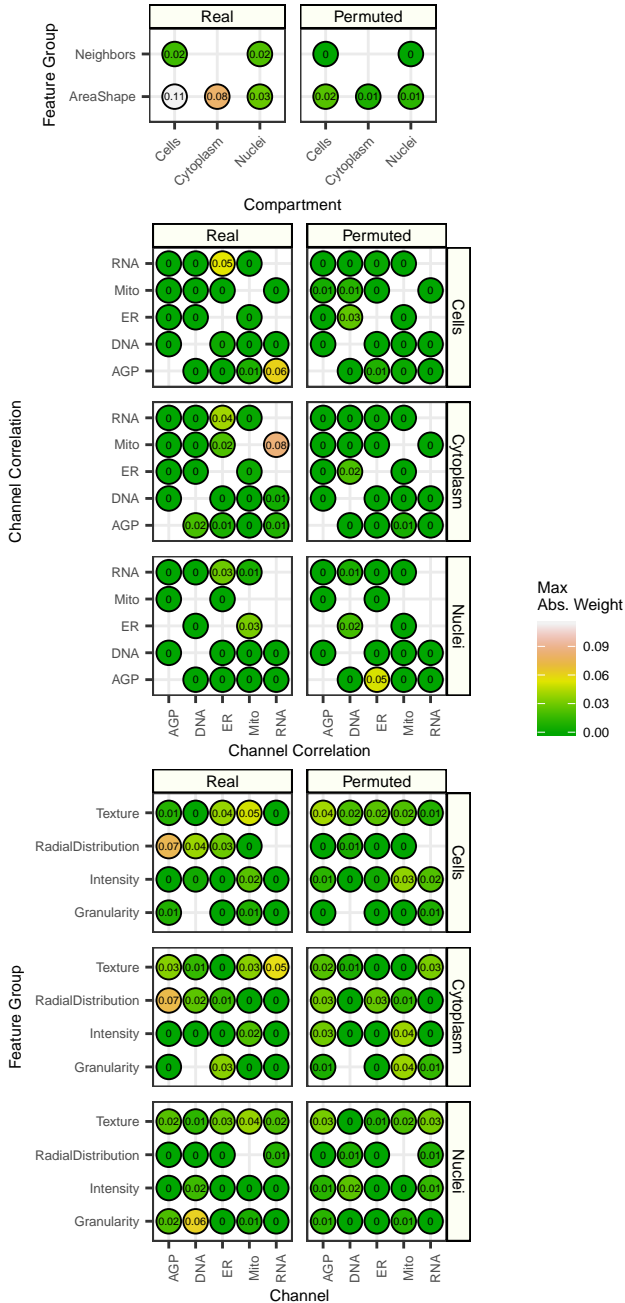
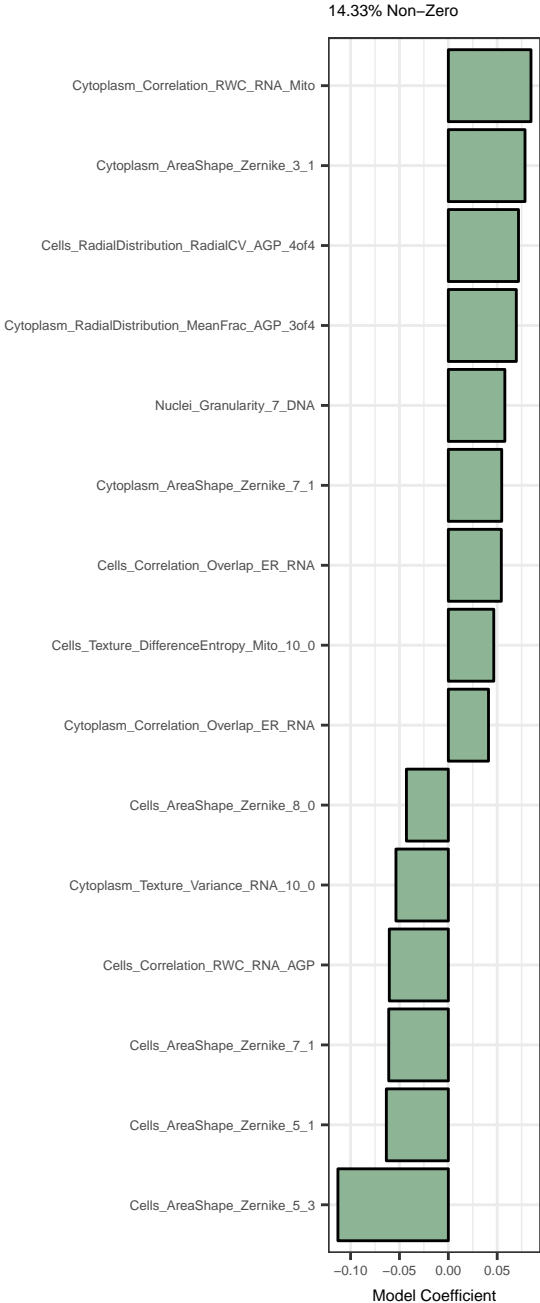
% Inf



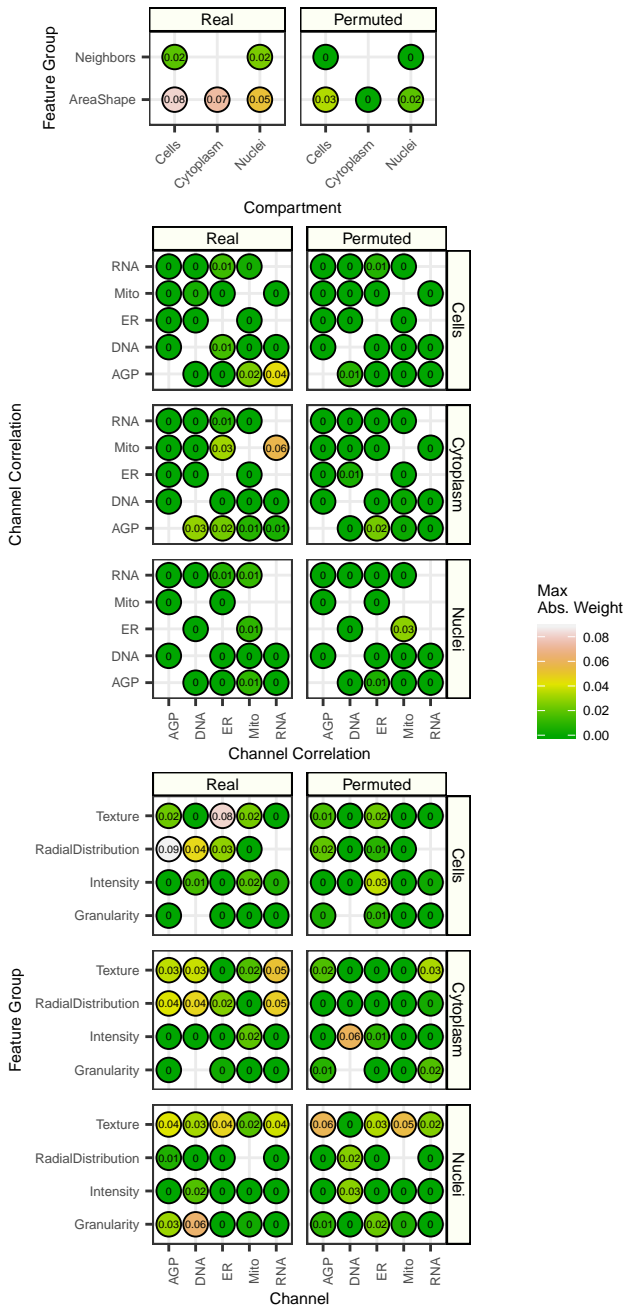
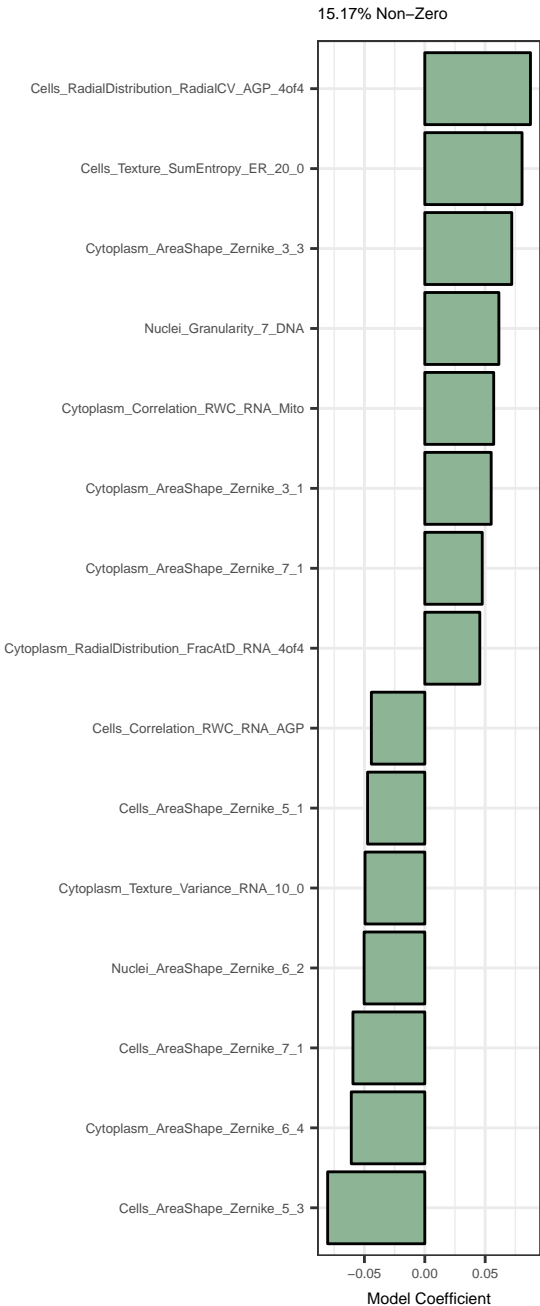
Live Cell Area



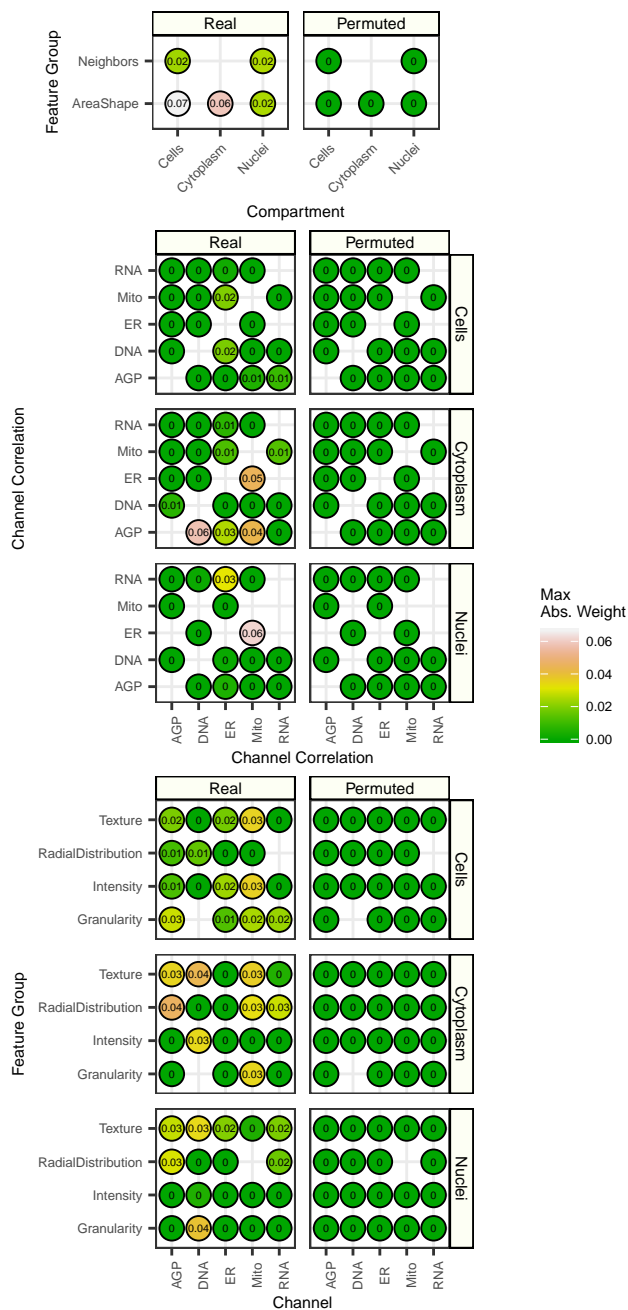
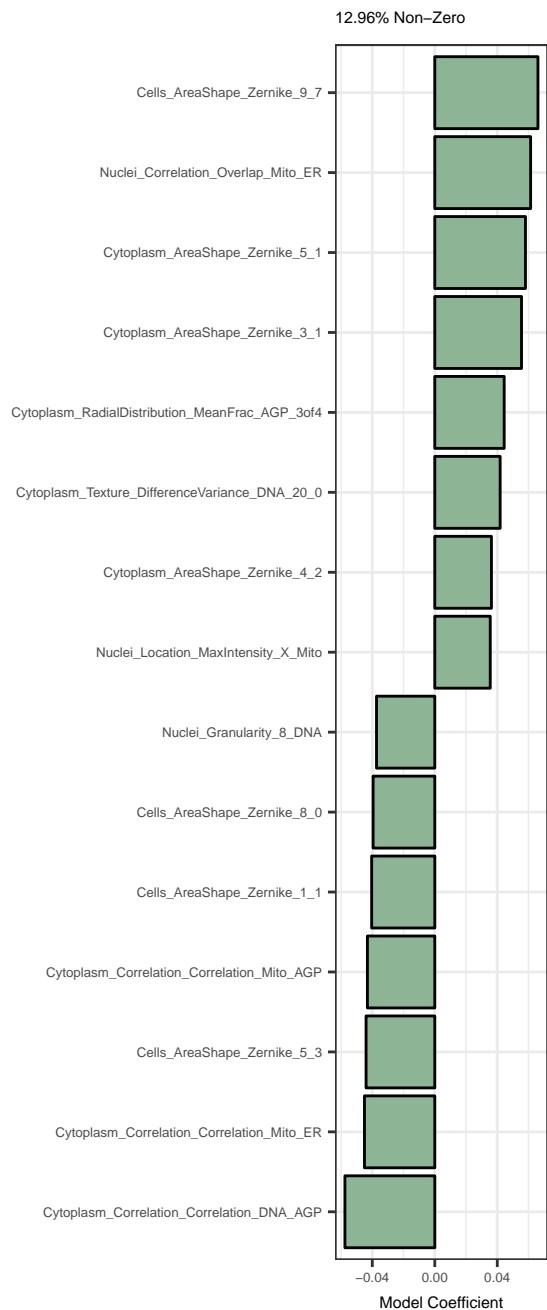
Live Cell Roundness



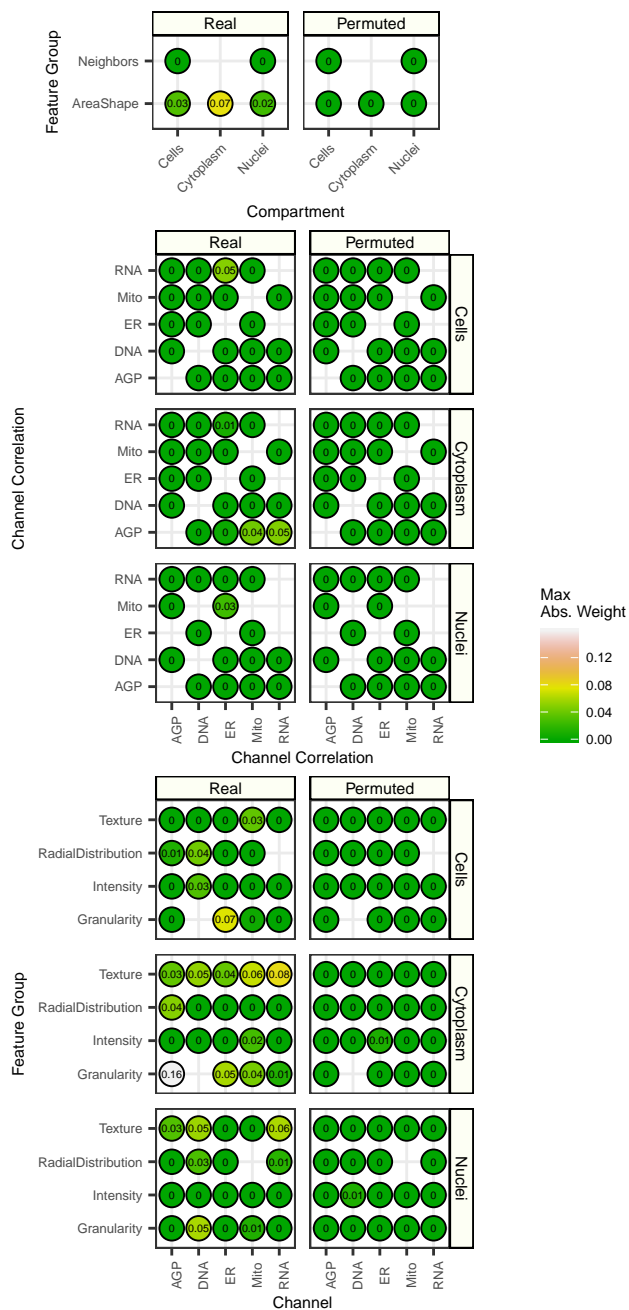
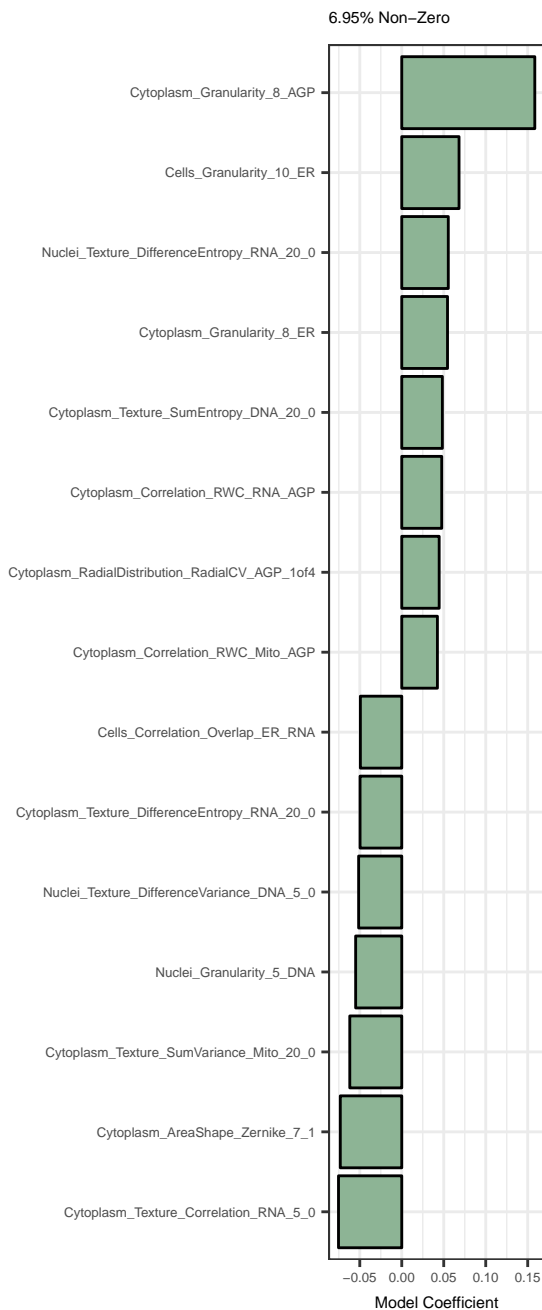
Live Width / Length



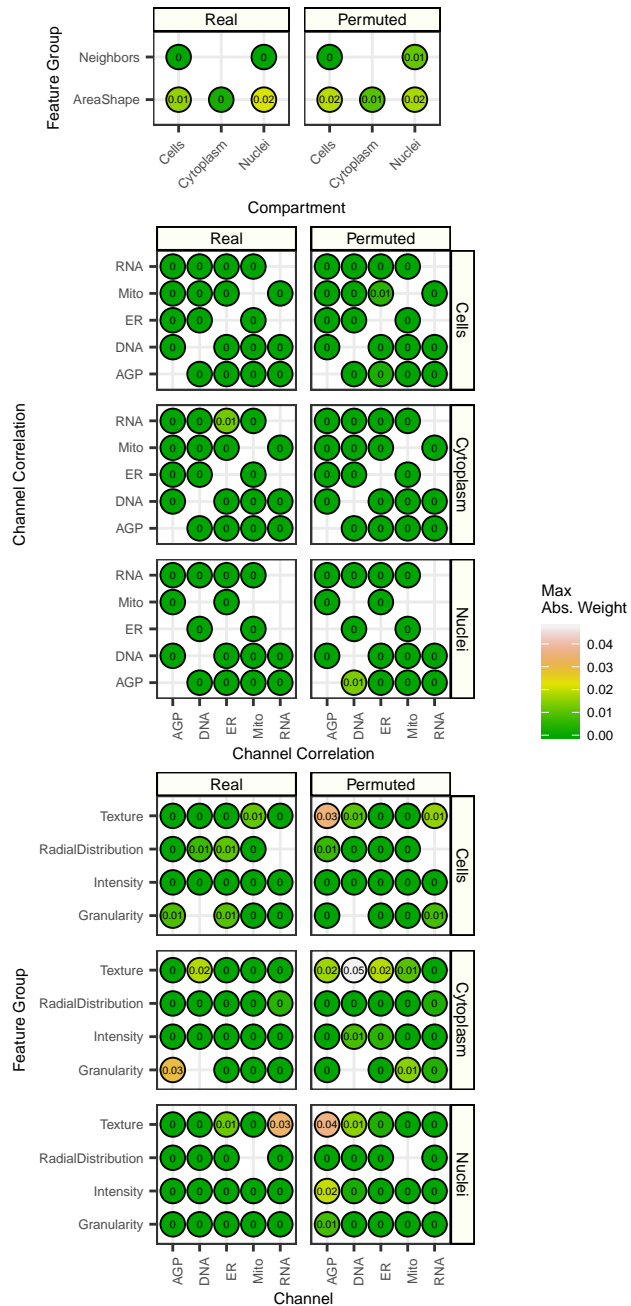
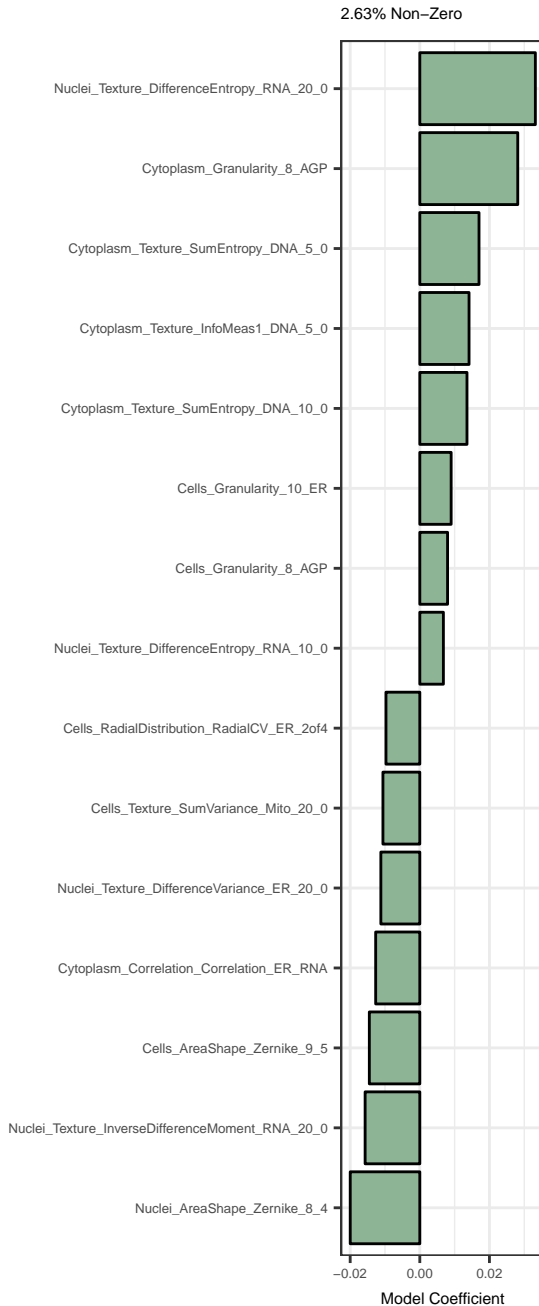
Live Cells



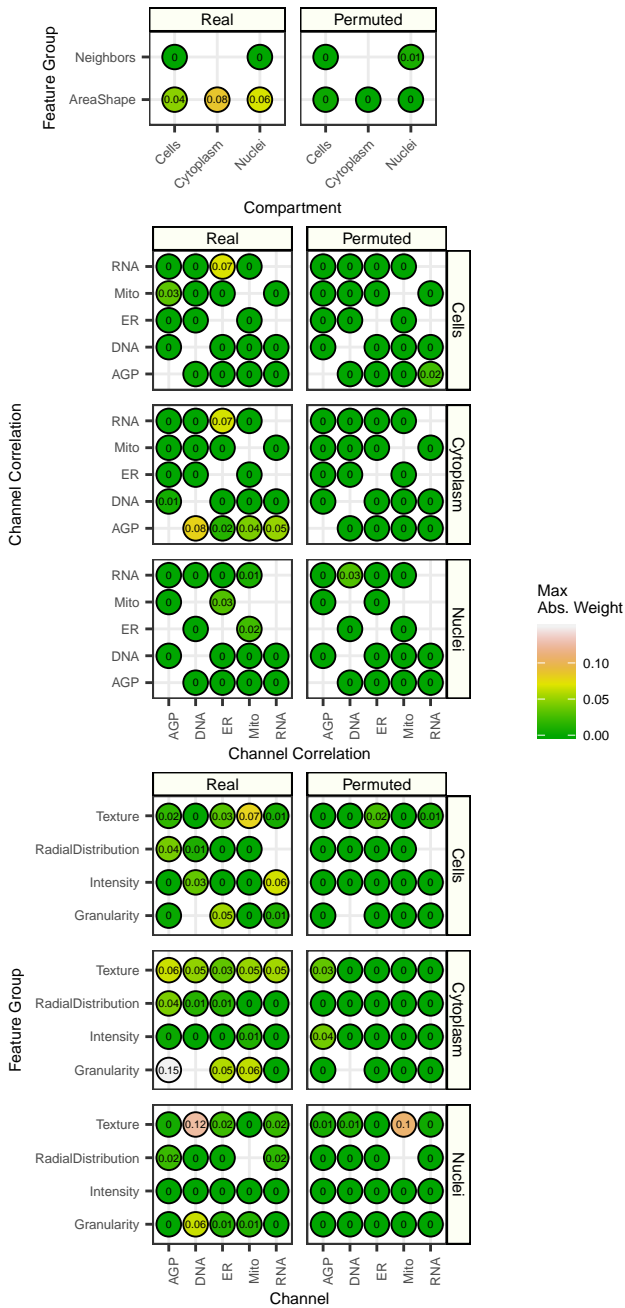
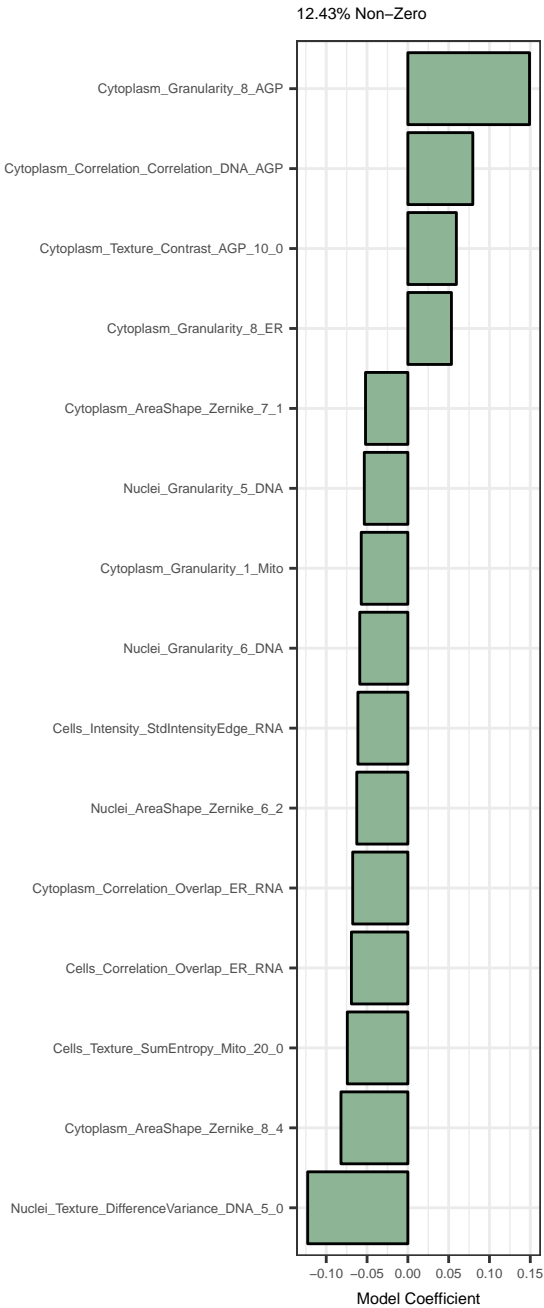
% All Apoptosis (CASP+)



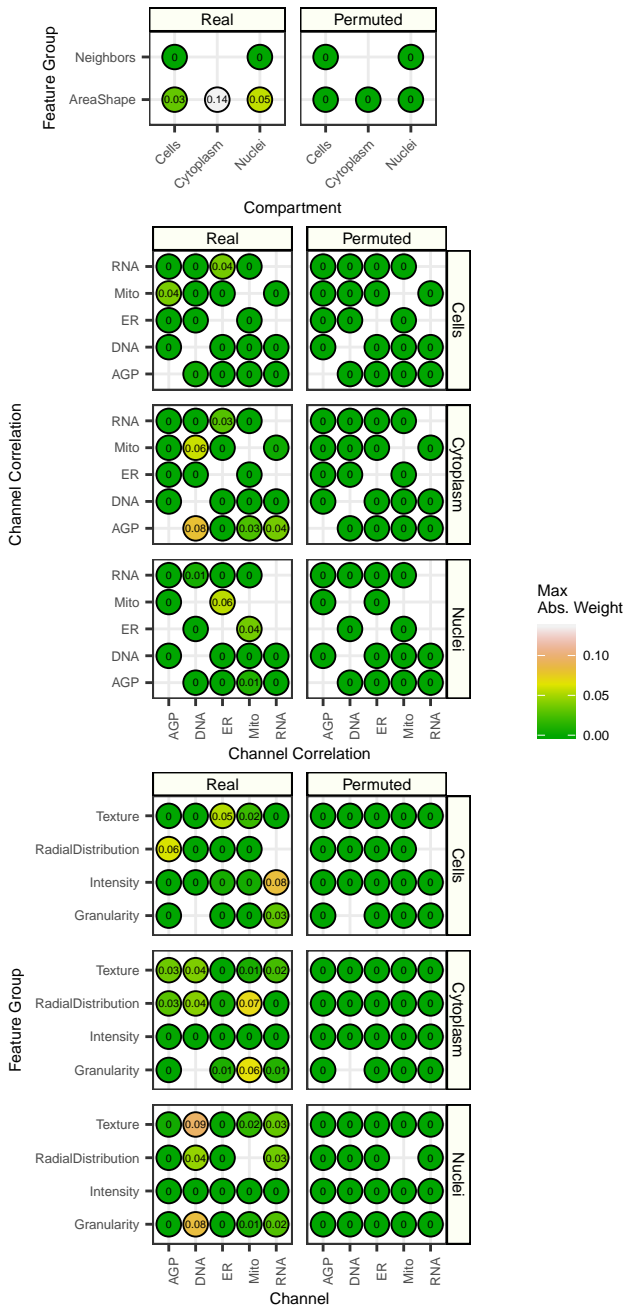
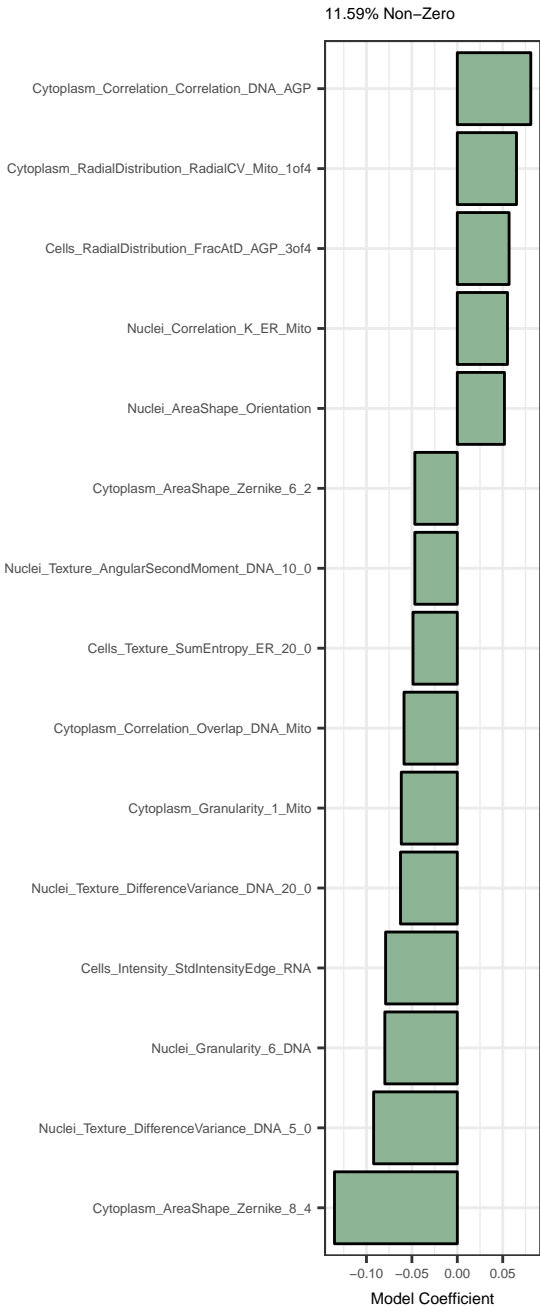
% Caspase / % Dead Only



% All Dead (DRAQ7+)

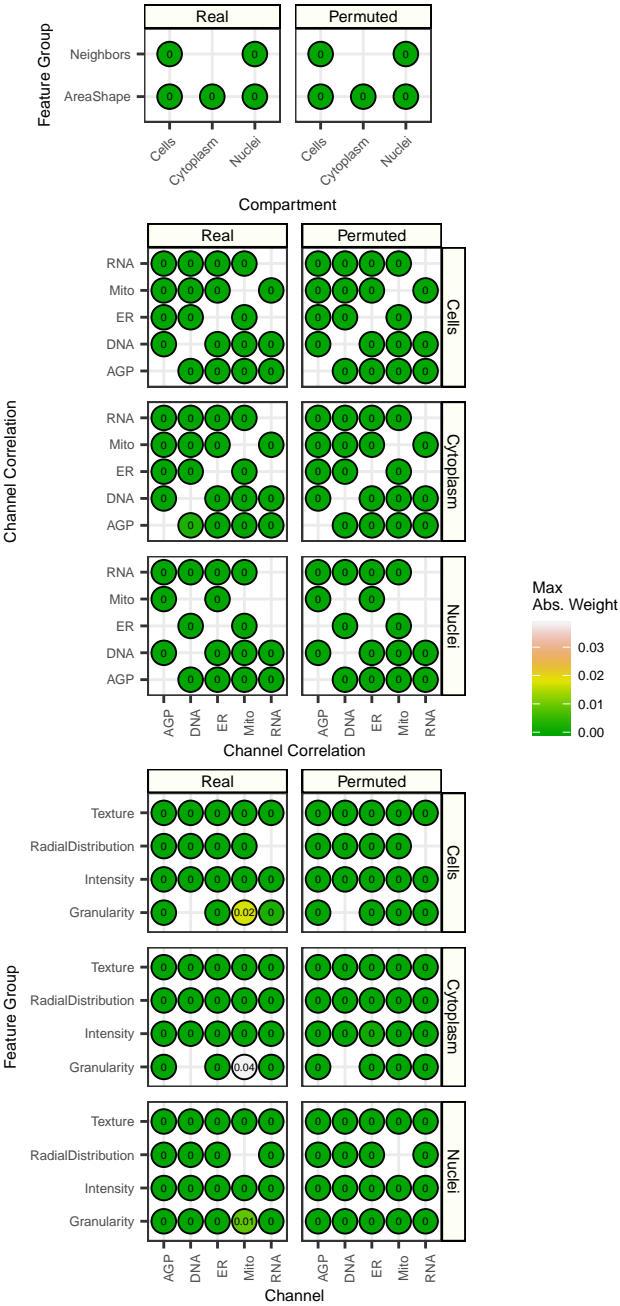
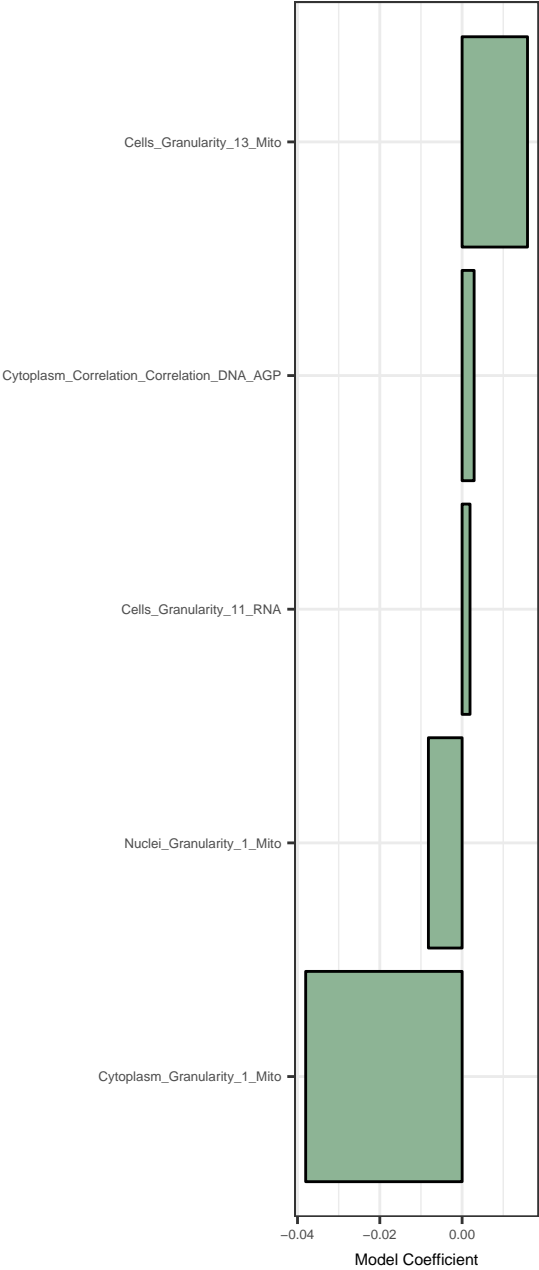


% Dead Only (CASP-; DRAQ7+)



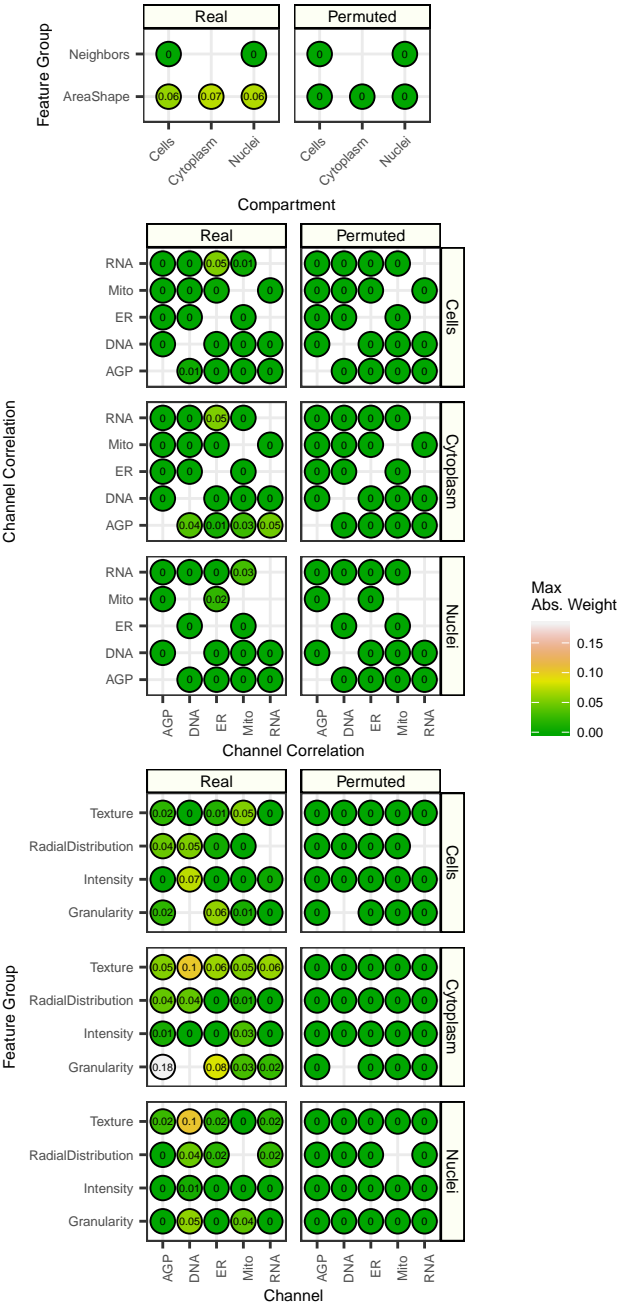
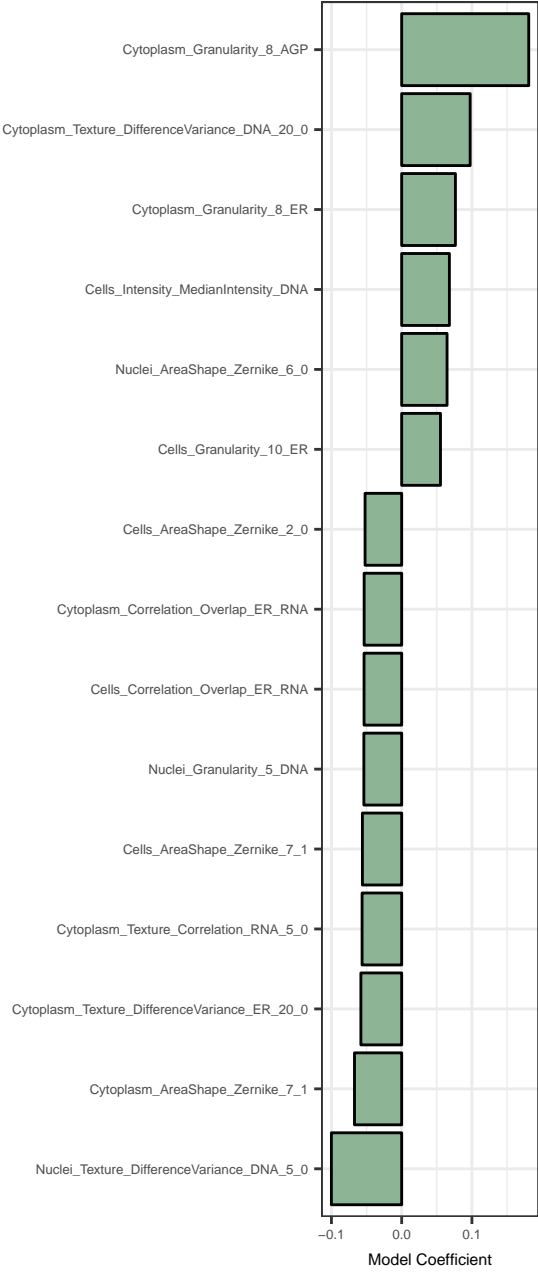
% Early Apoptosis (CASP+; DRAQ7-)

0.53% Non-Zero

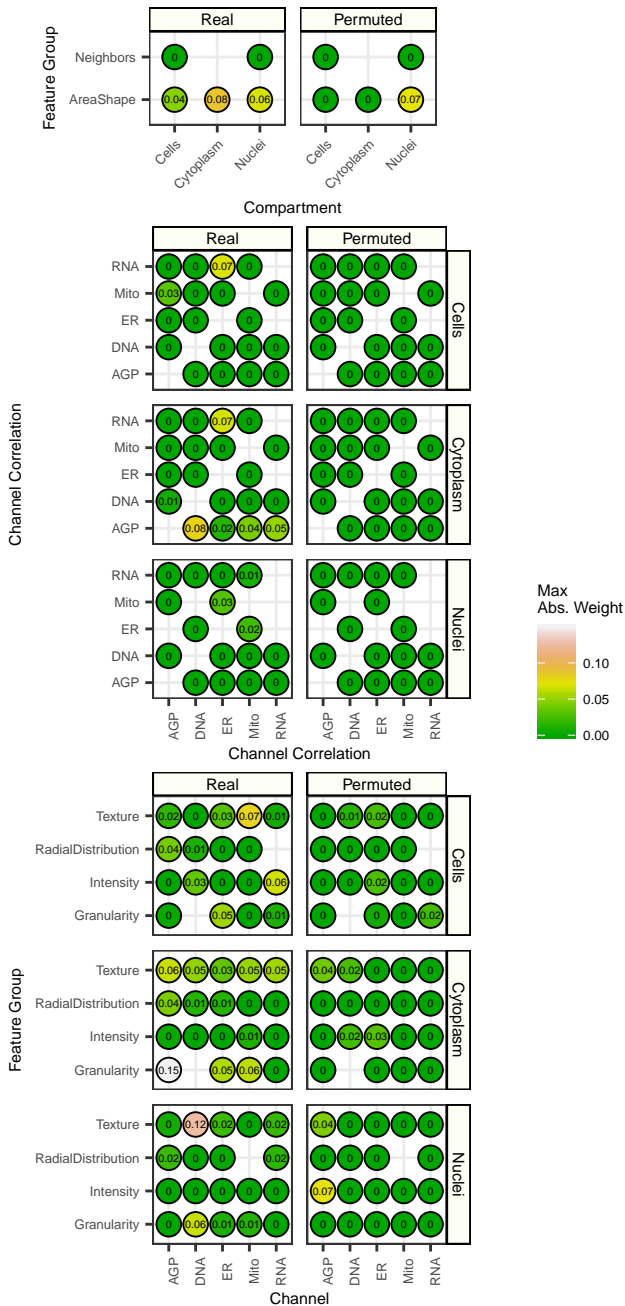
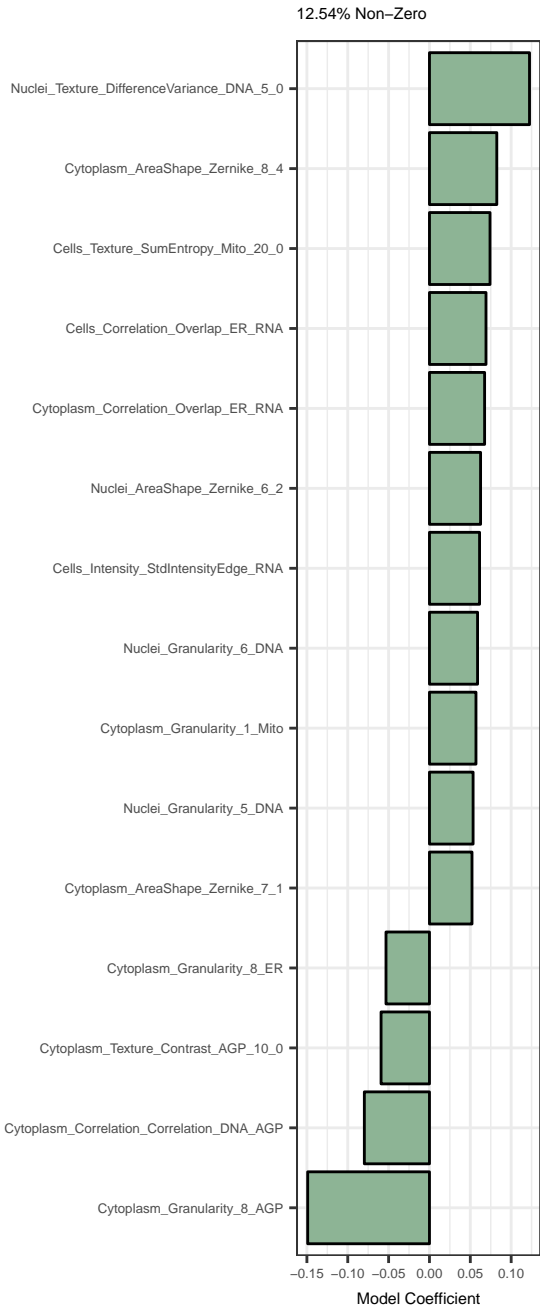


% Late Apoptosis (CASP+; DRAQ7+)

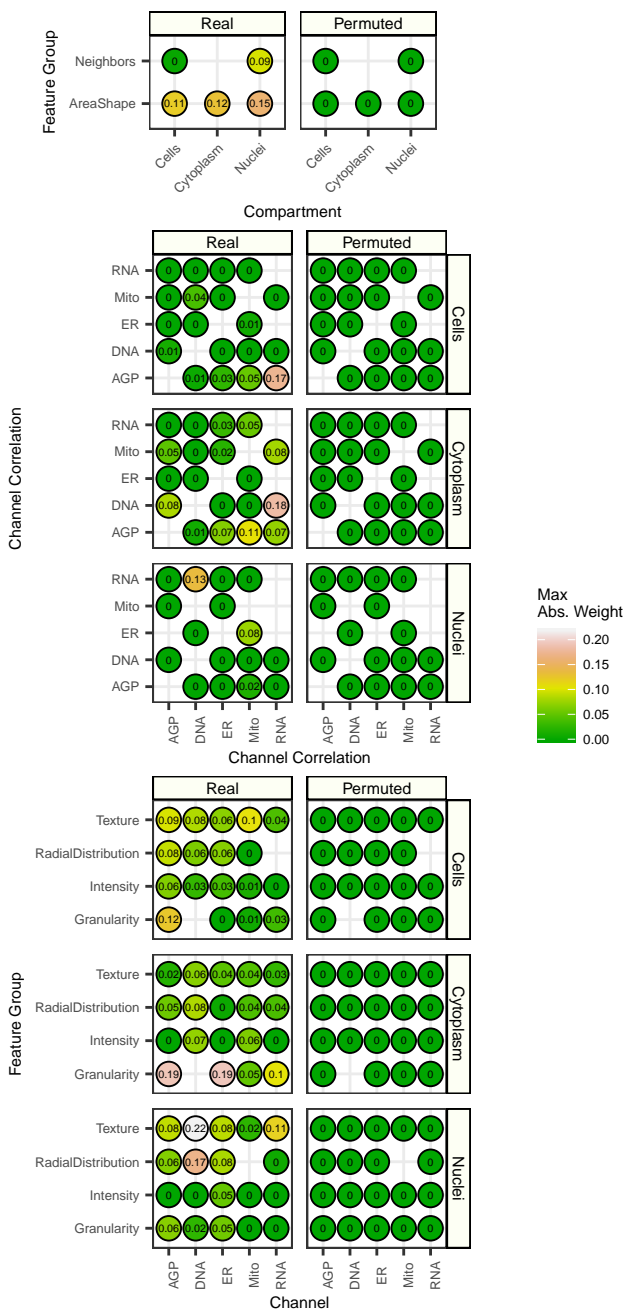
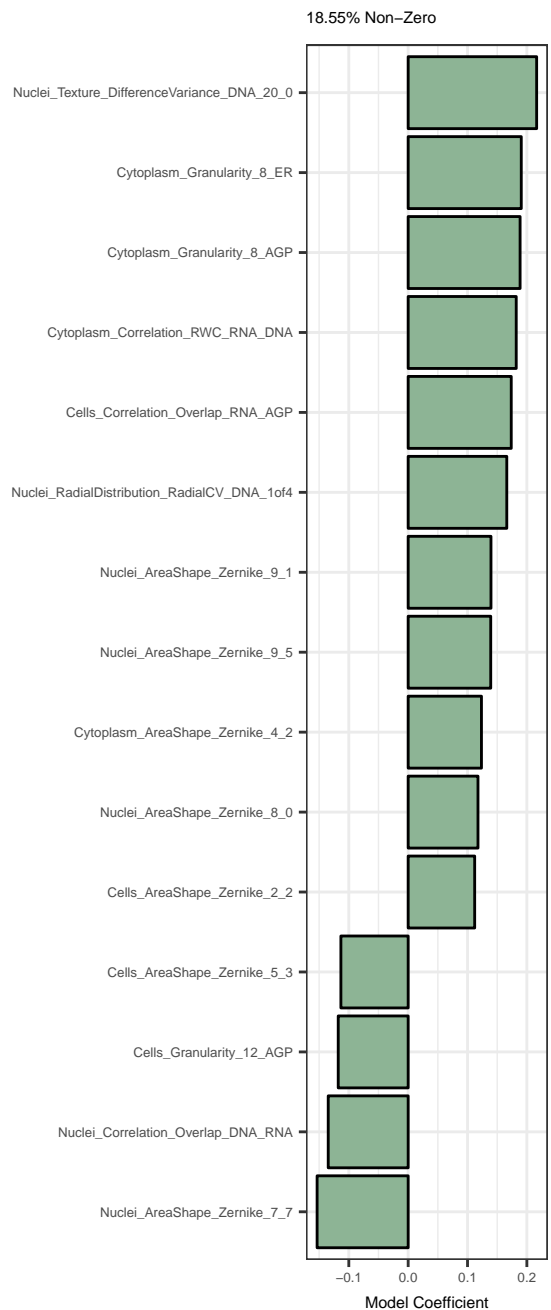
13.28% Non-Zero



% Live (DRAQ7-; CASP-)



ROS-back



ROS

