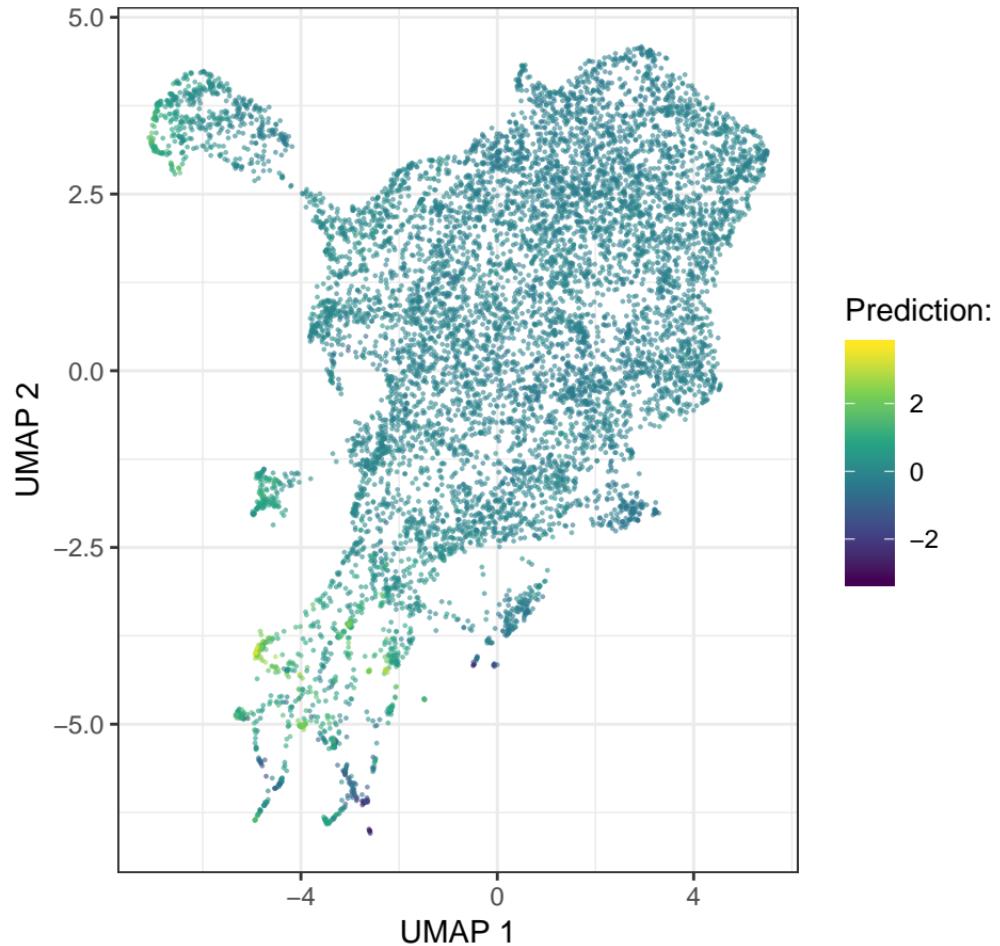
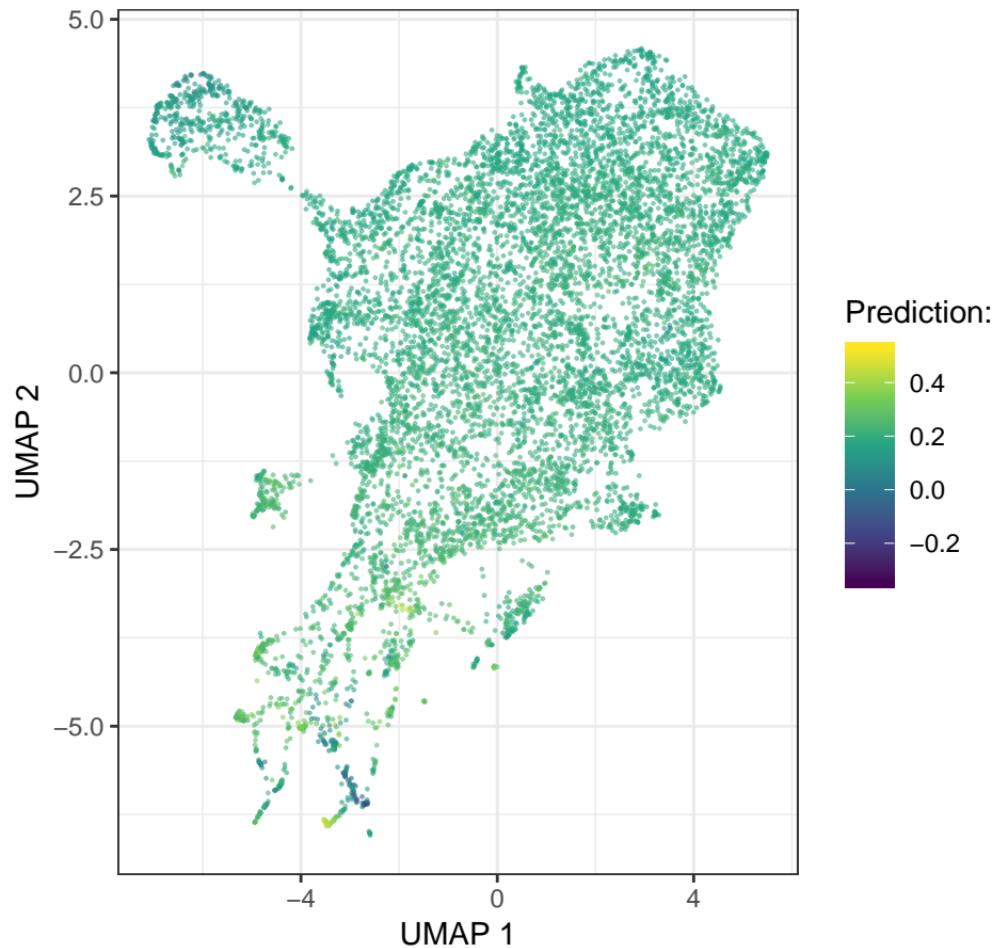


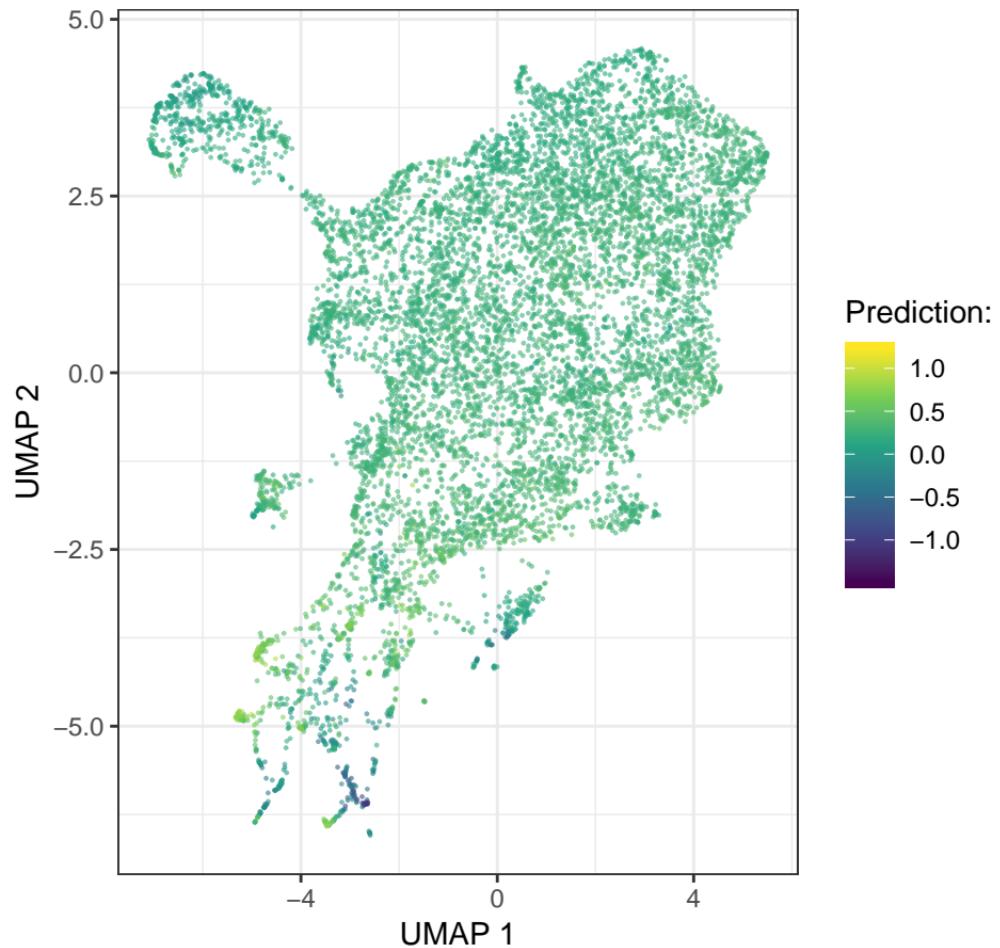
cc\_cc\_g1\_mean



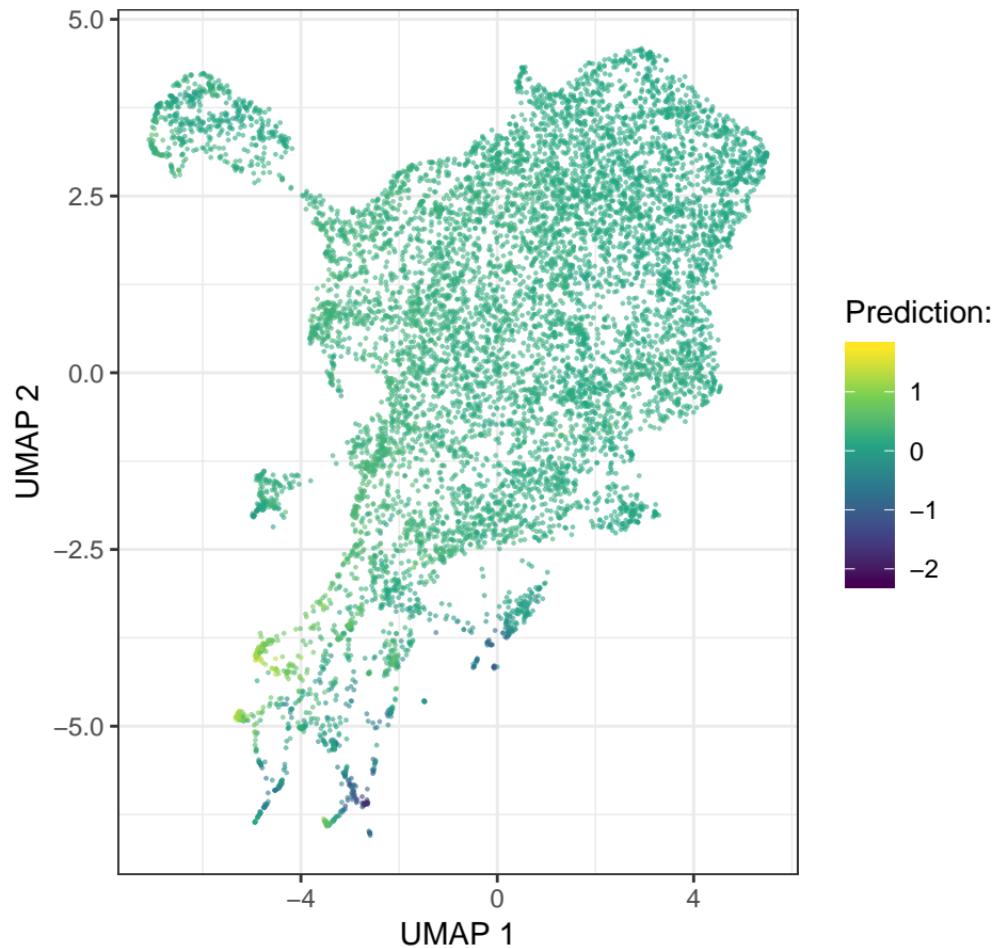
cc\_mitosis\_ph3\_neg\_n\_spots\_mean



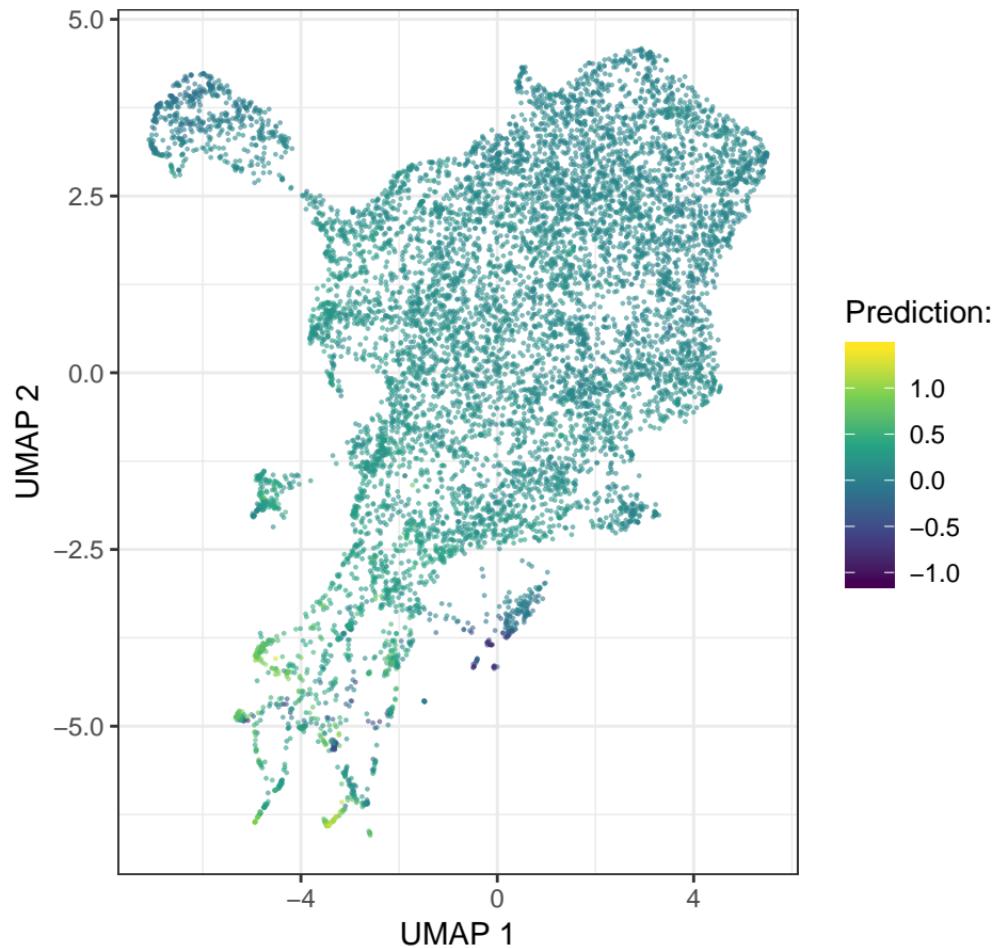
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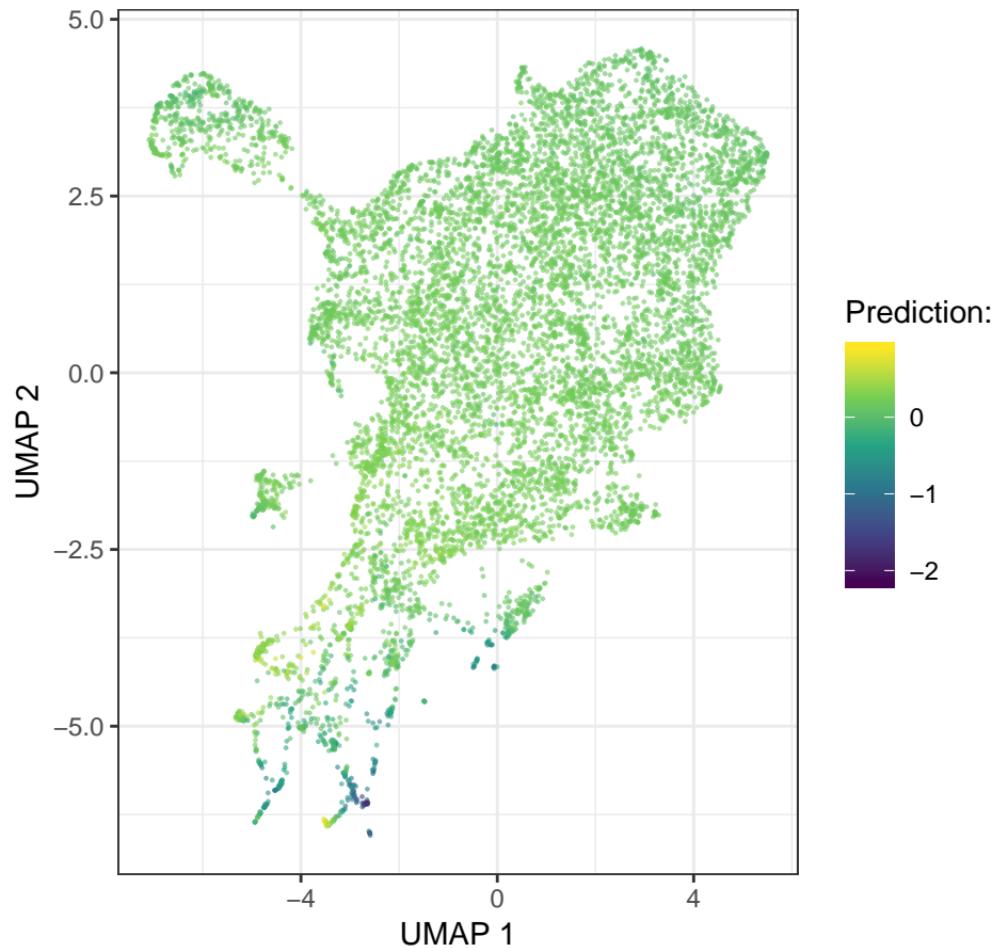
cc\_polyplloid\_n\_spots\_mean



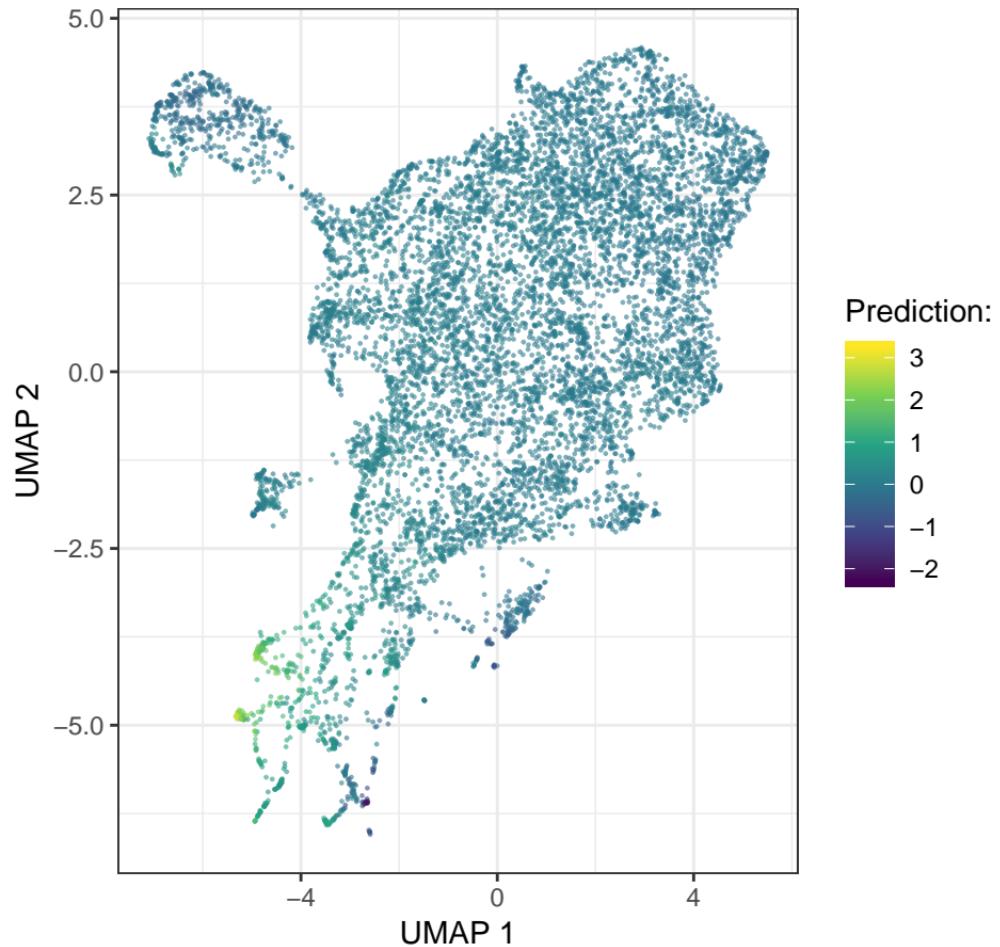
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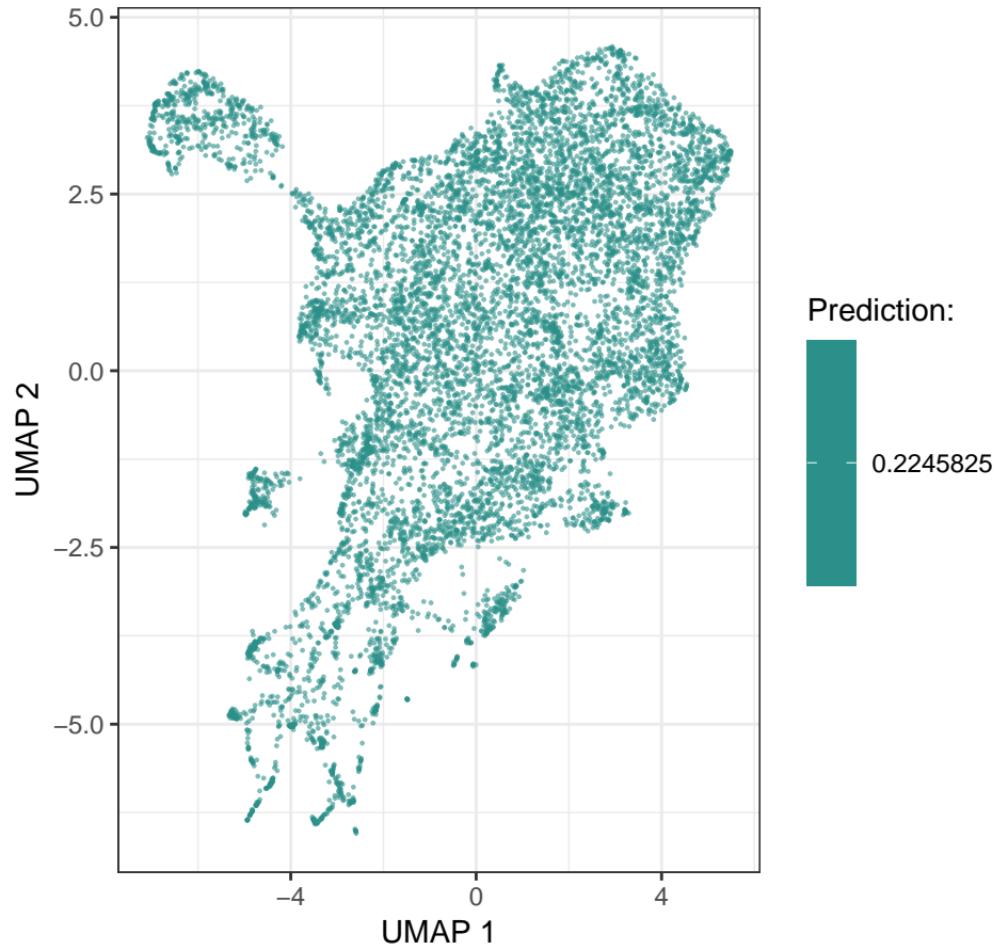
# cc\_polyplloid\_high\_n\_spots\_h2ax\_mean



cc\_cc\_n\_spots\_per\_nucleus\_area\_mean



# cc\_mitosis\_ph3\_neg\_high\_n\_spots\_h2ax\_mean



cc\_edu\_pos\_alexa647\_intensity\_nucleus\_area\_sum

UMAP 2

5.0

2.5

0.0

-2.5

-5.0

-4

0

4

UMAP 1

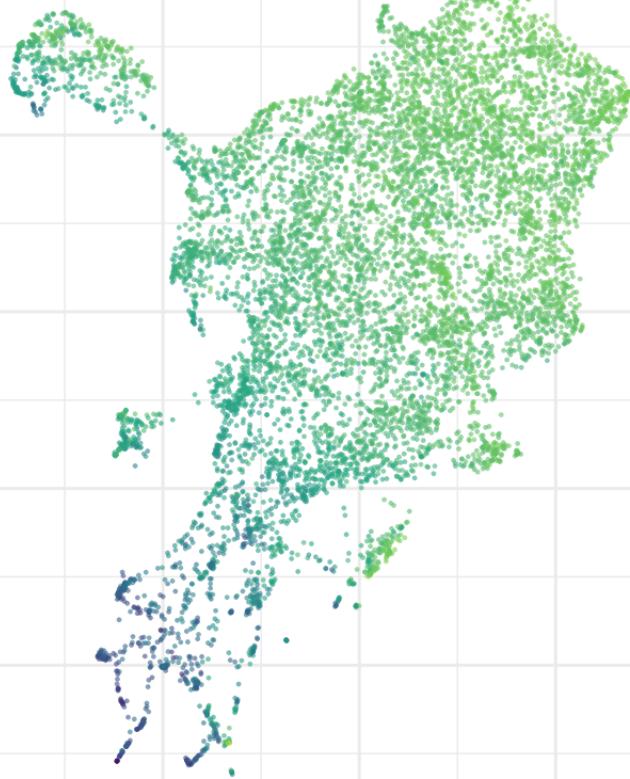
Prediction:

1

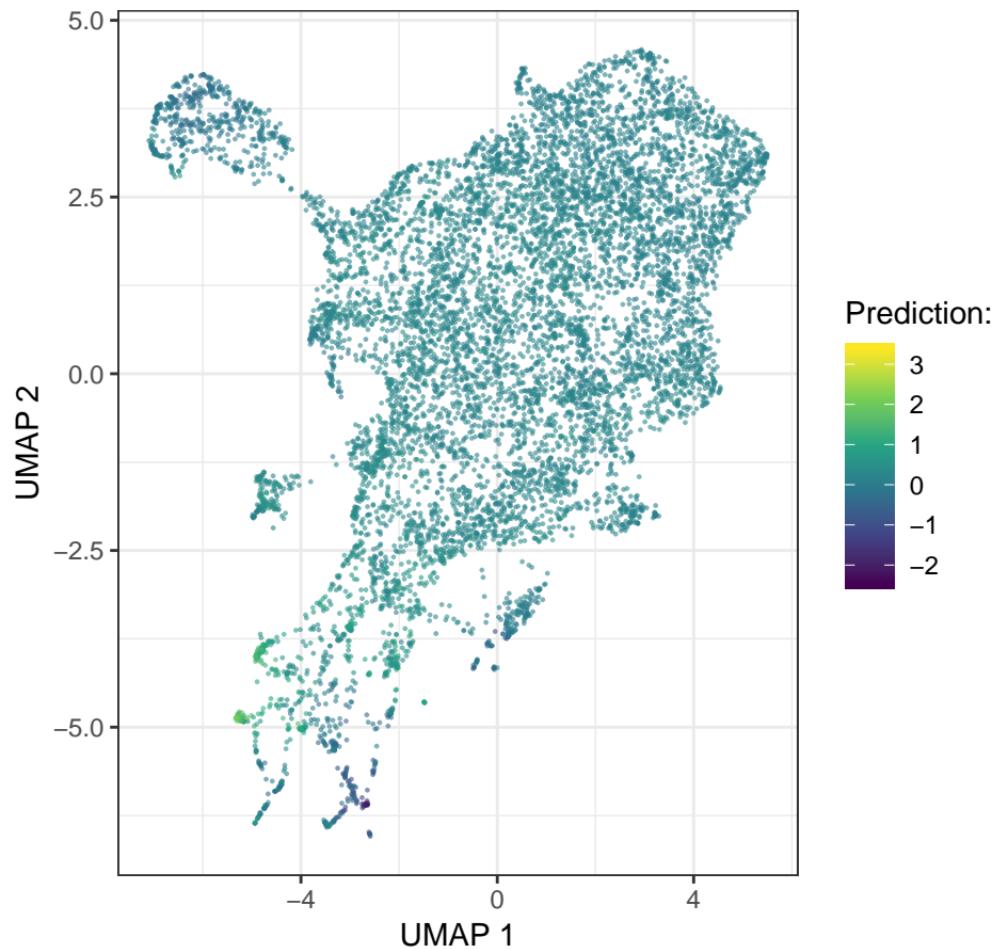
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-1

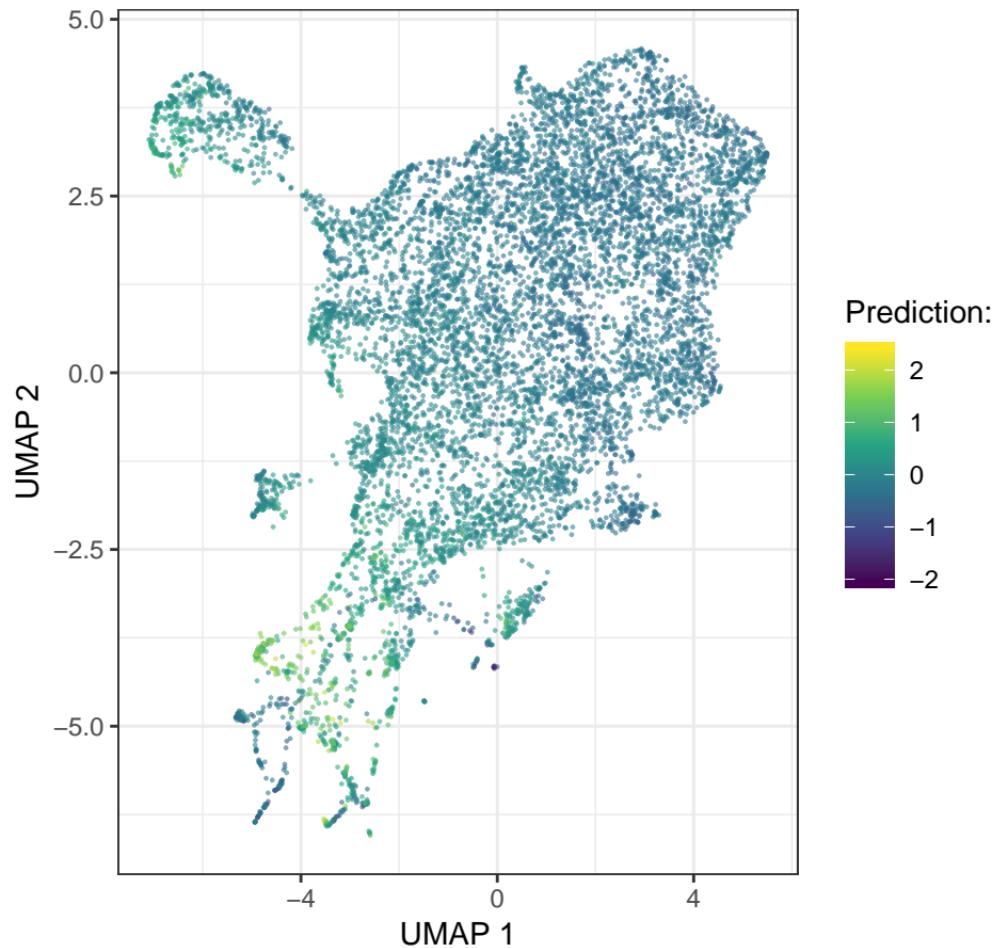
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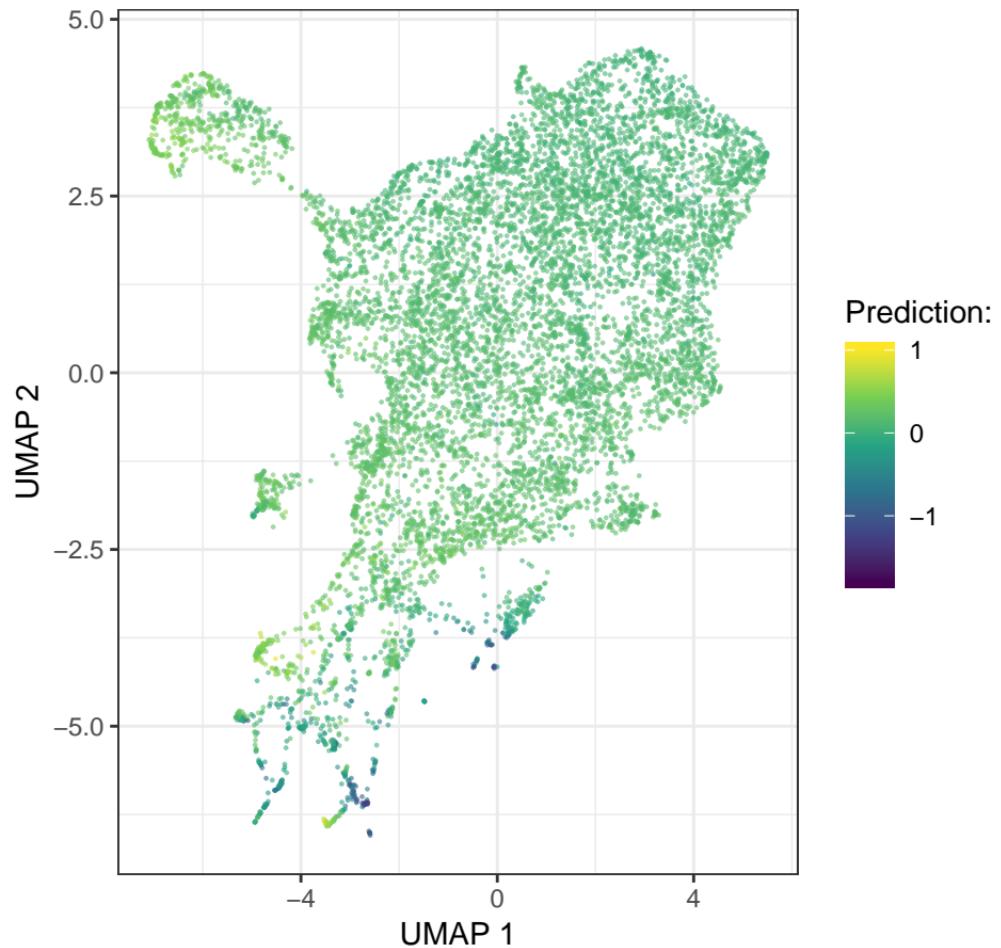
cc\_edu\_pos\_n\_spots\_per\_nucleus\_area\_mean



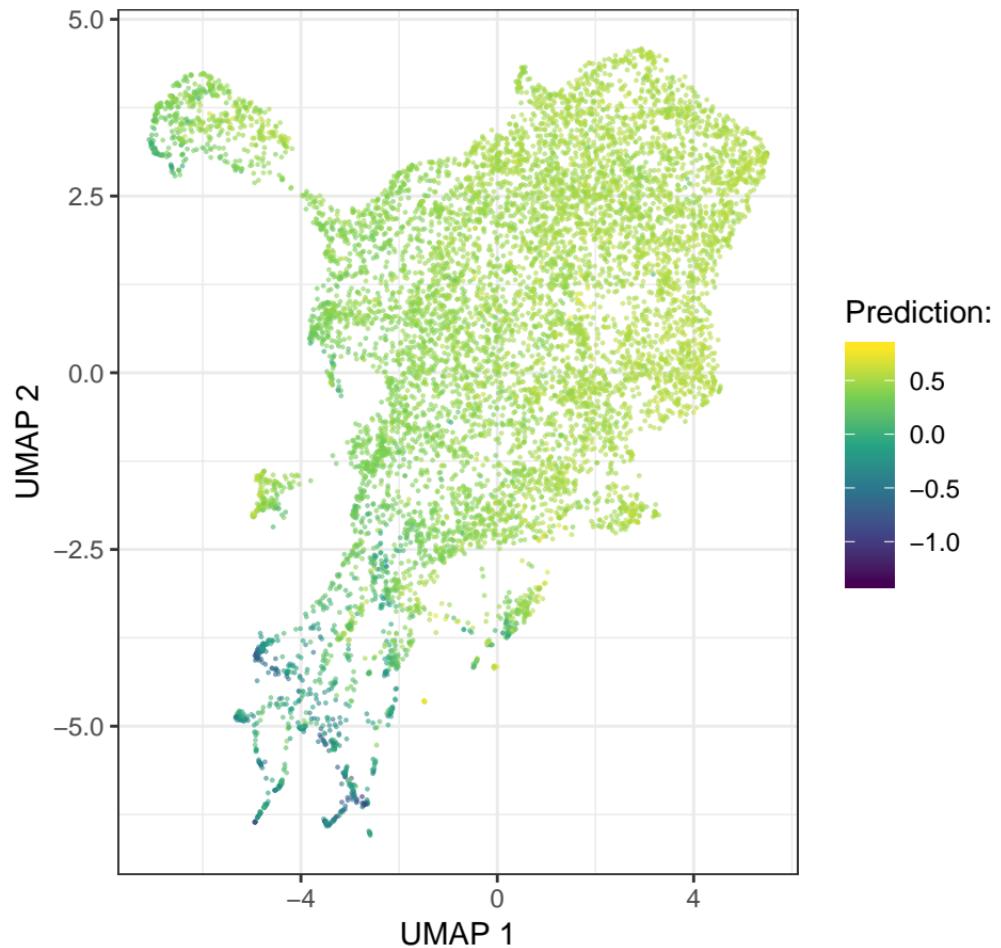
cc\_all\_nucleus\_roundness\_mean



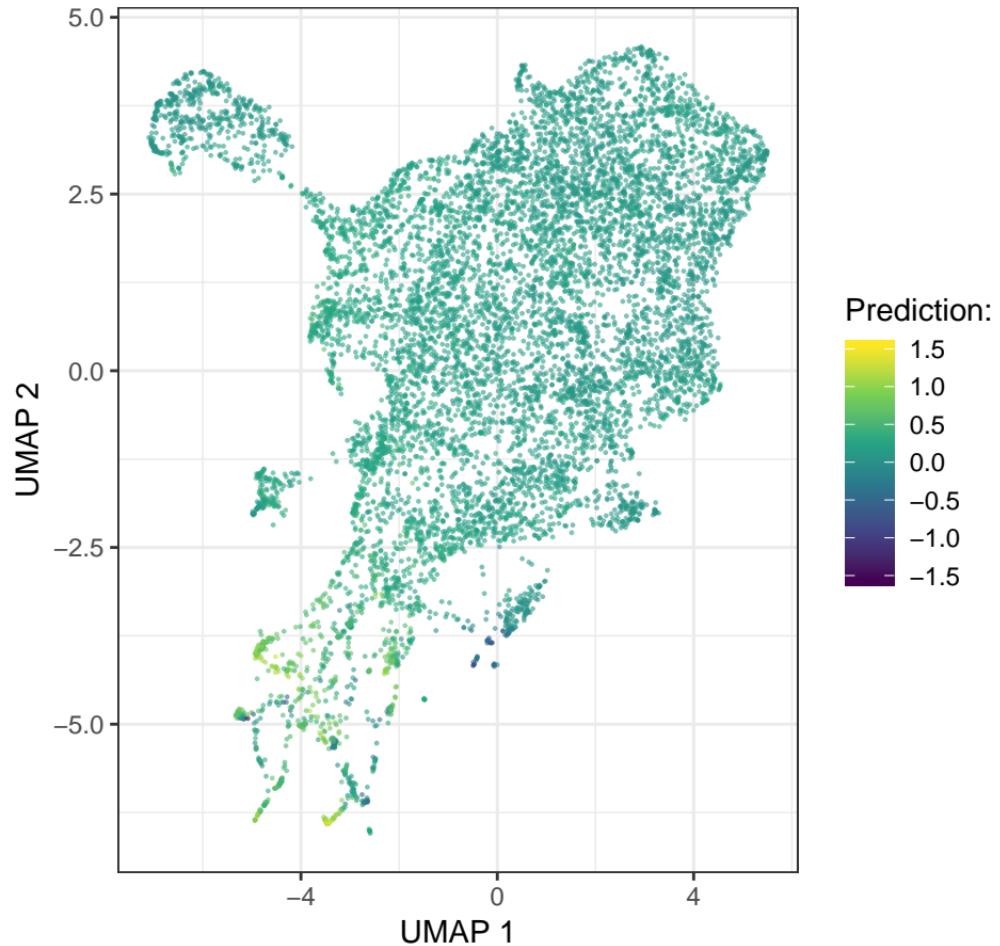
cc\_polynuclear\_n\_spots\_mean



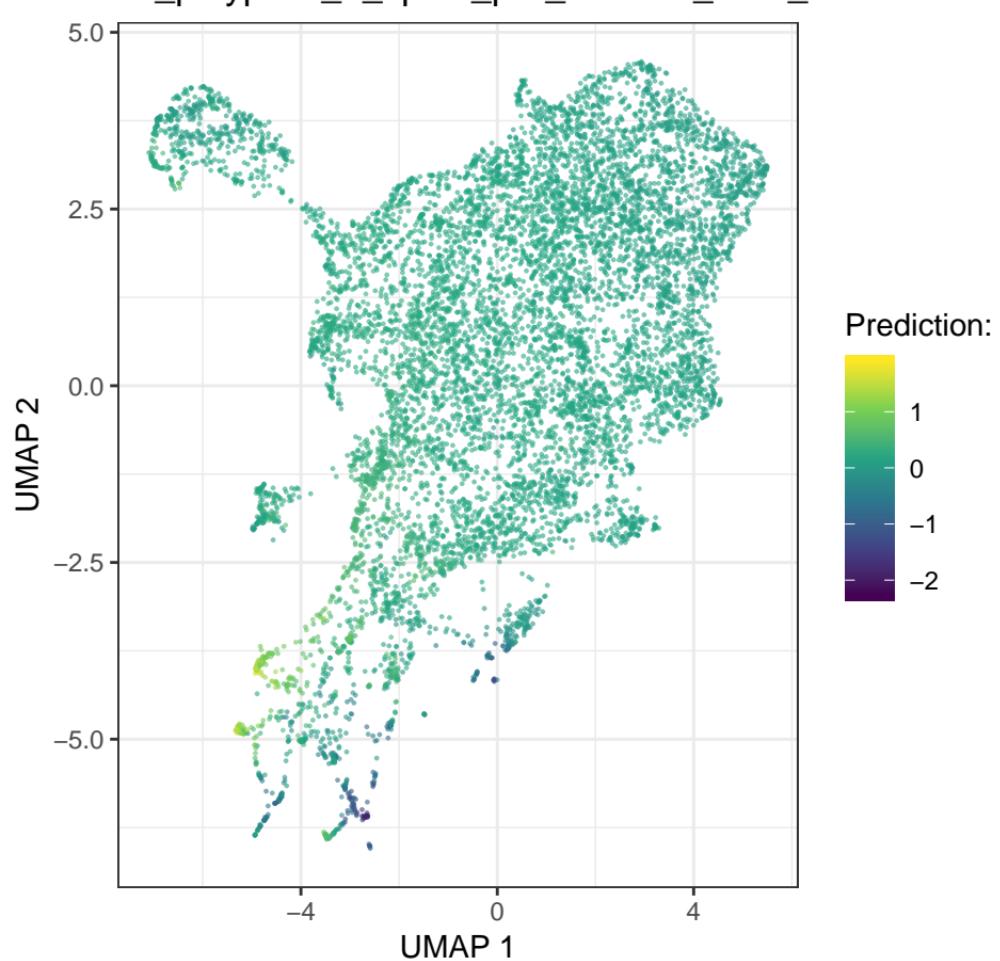
cc\_polynuclear\_n\_objects



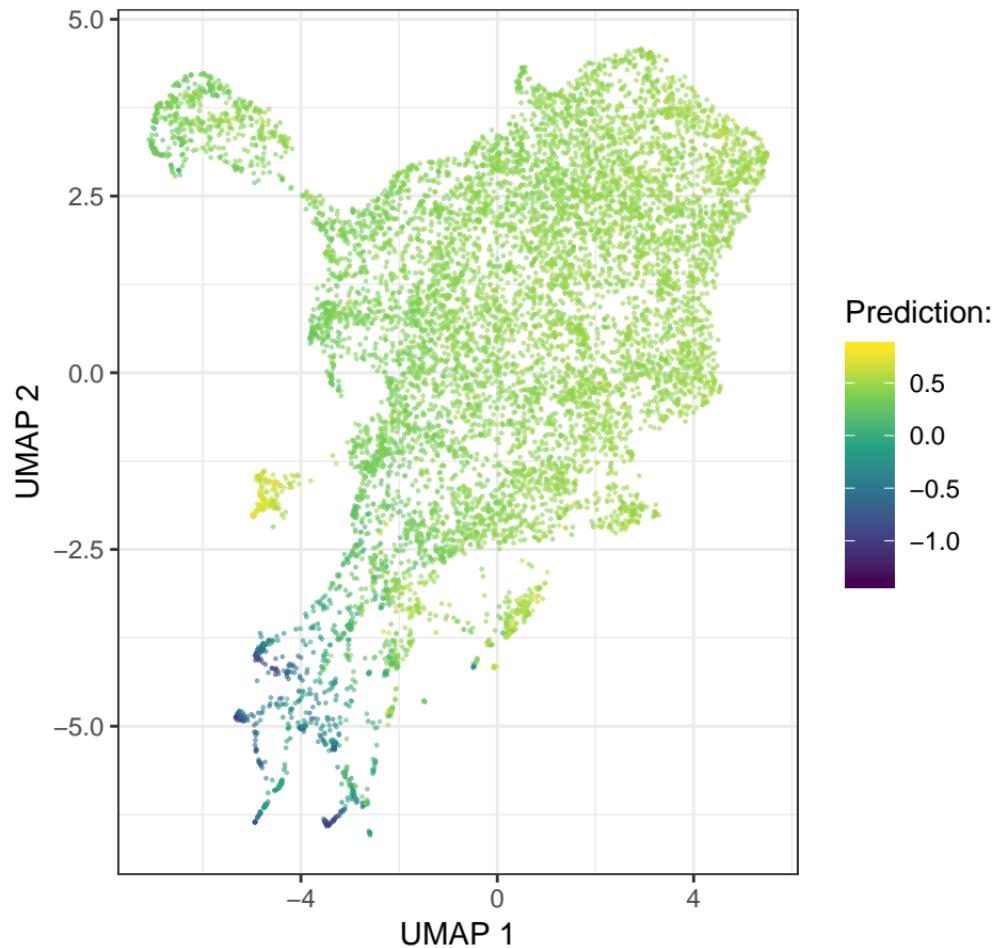
cc\_g2\_ph3\_neg\_high\_n\_spots\_h2ax\_mean



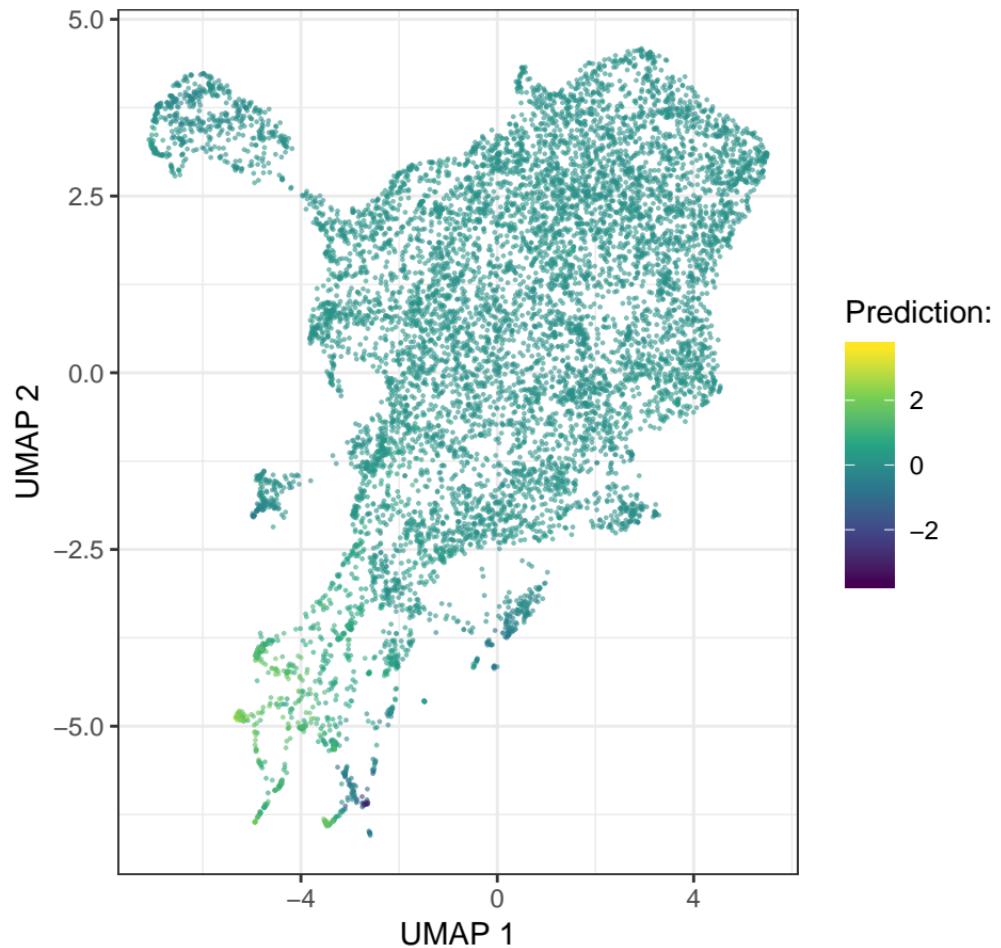
cc\_polyplloid\_n\_spots\_per\_nucleus\_area\_mean



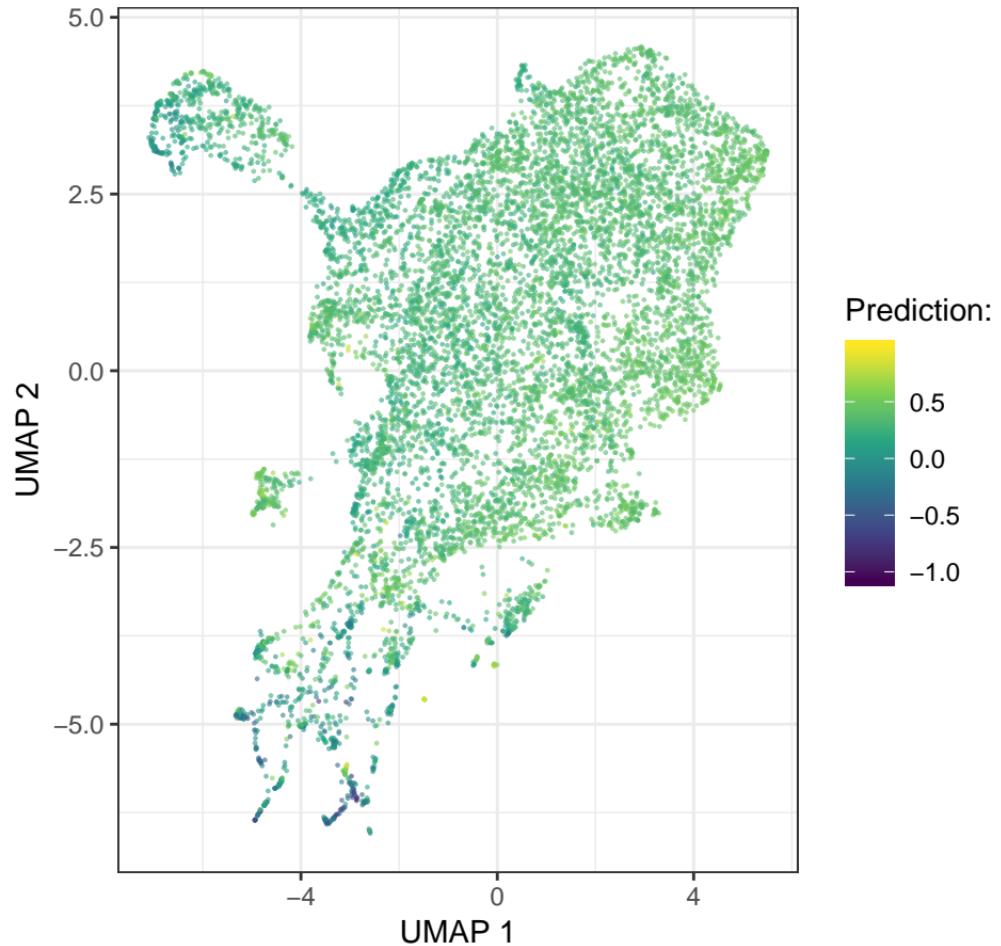
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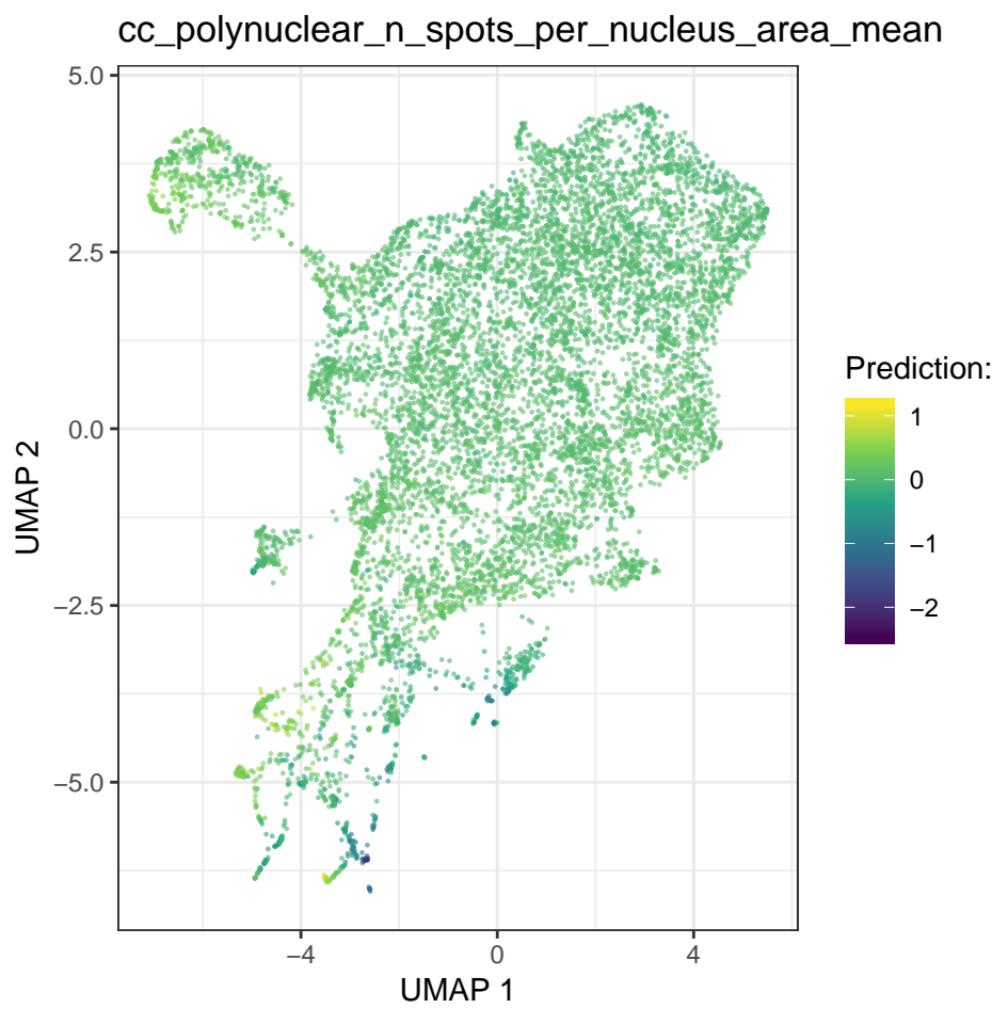


cc\_all\_n\_spots\_mean

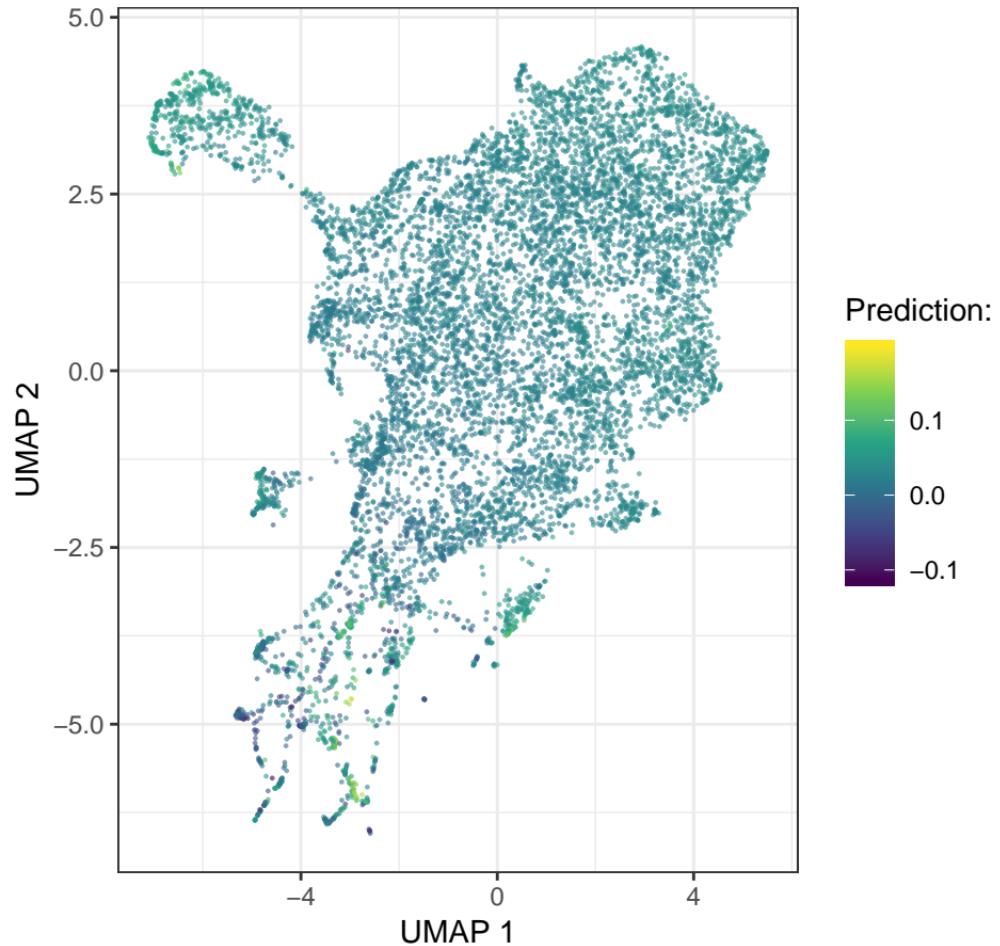


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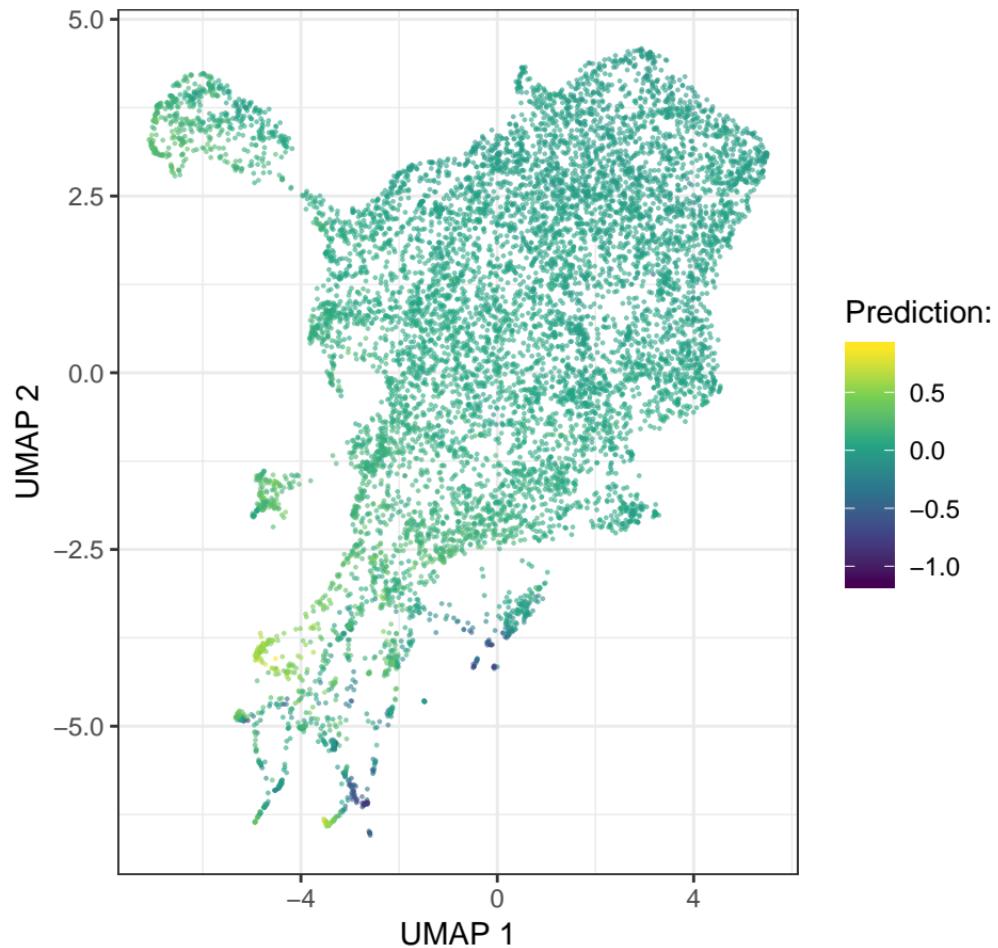




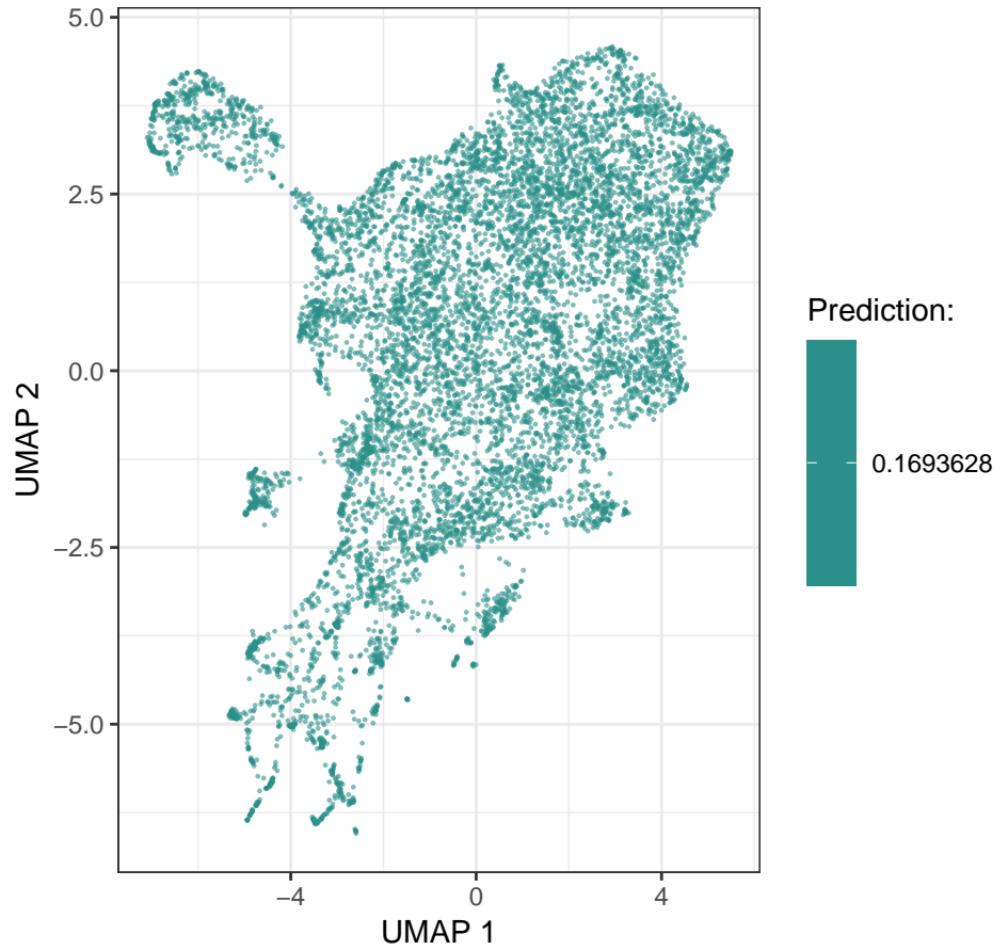
cc\_infection\_percentage



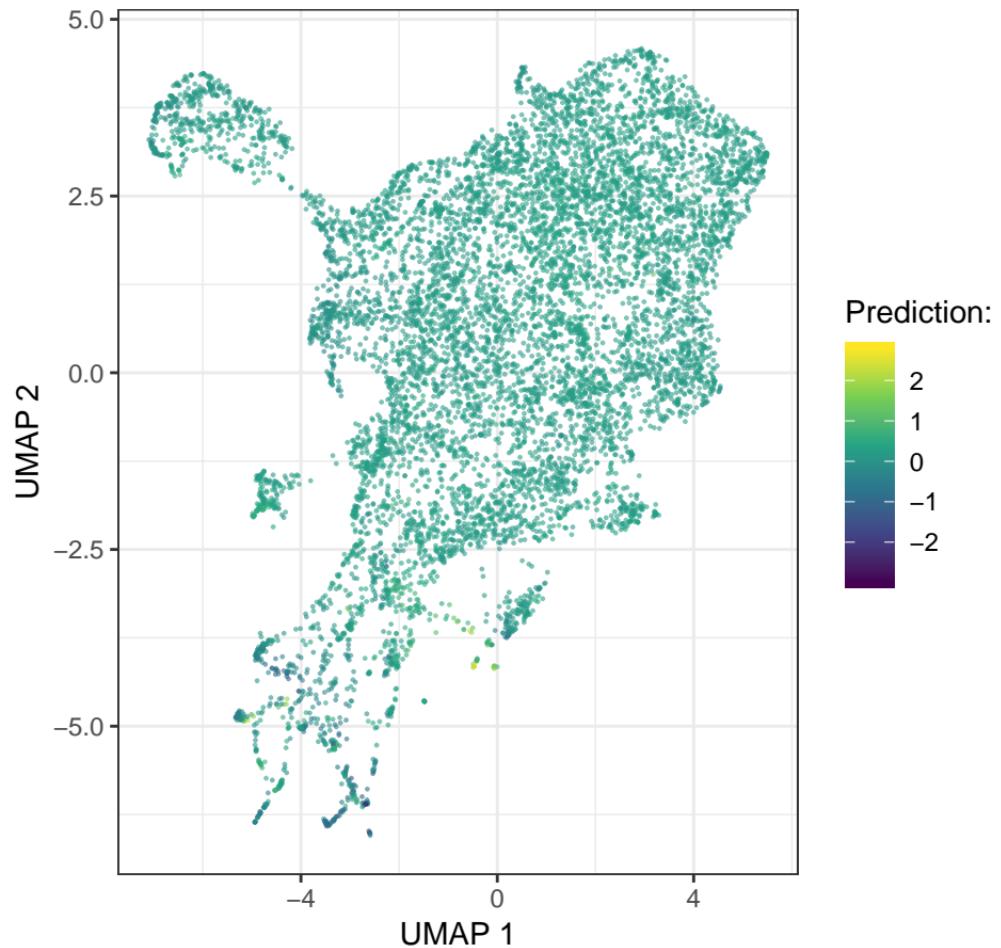
cc\_polynuclear\_high\_n\_spots\_h2ax\_mean



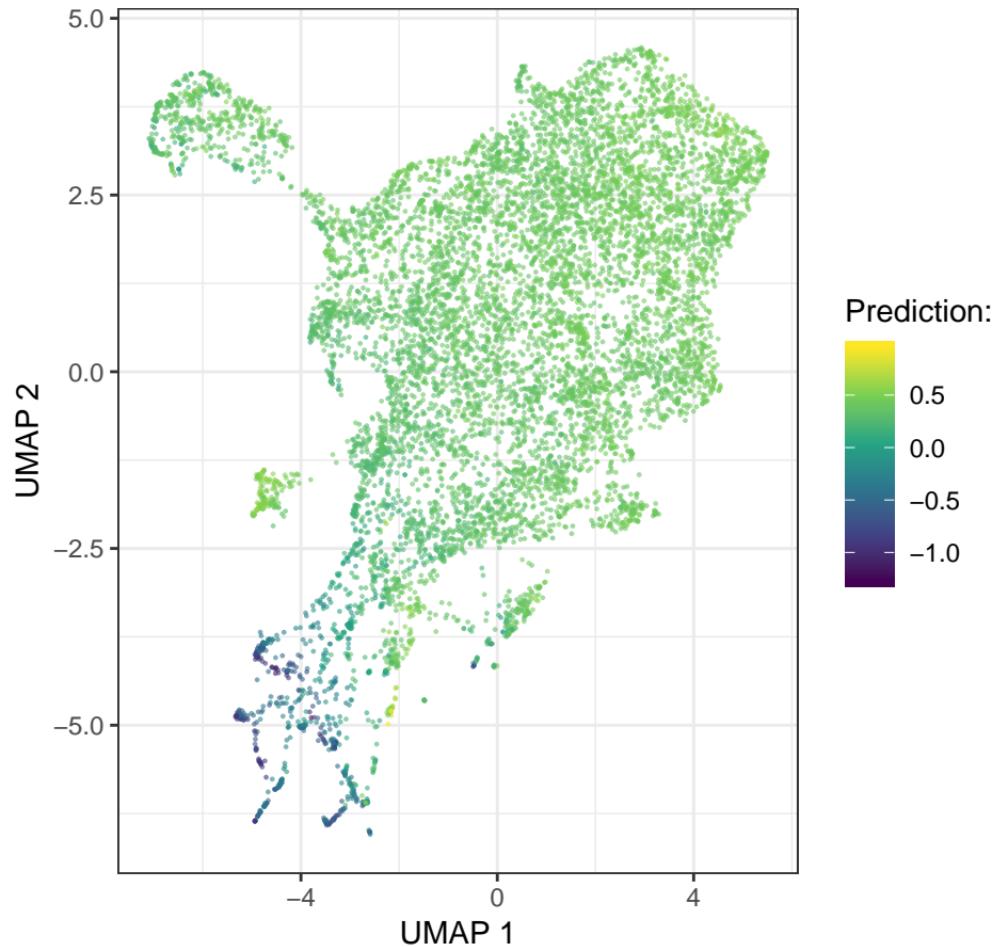
# cc\_mitosis\_ph3\_pos\_high\_n\_spots\_h2ax\_mean



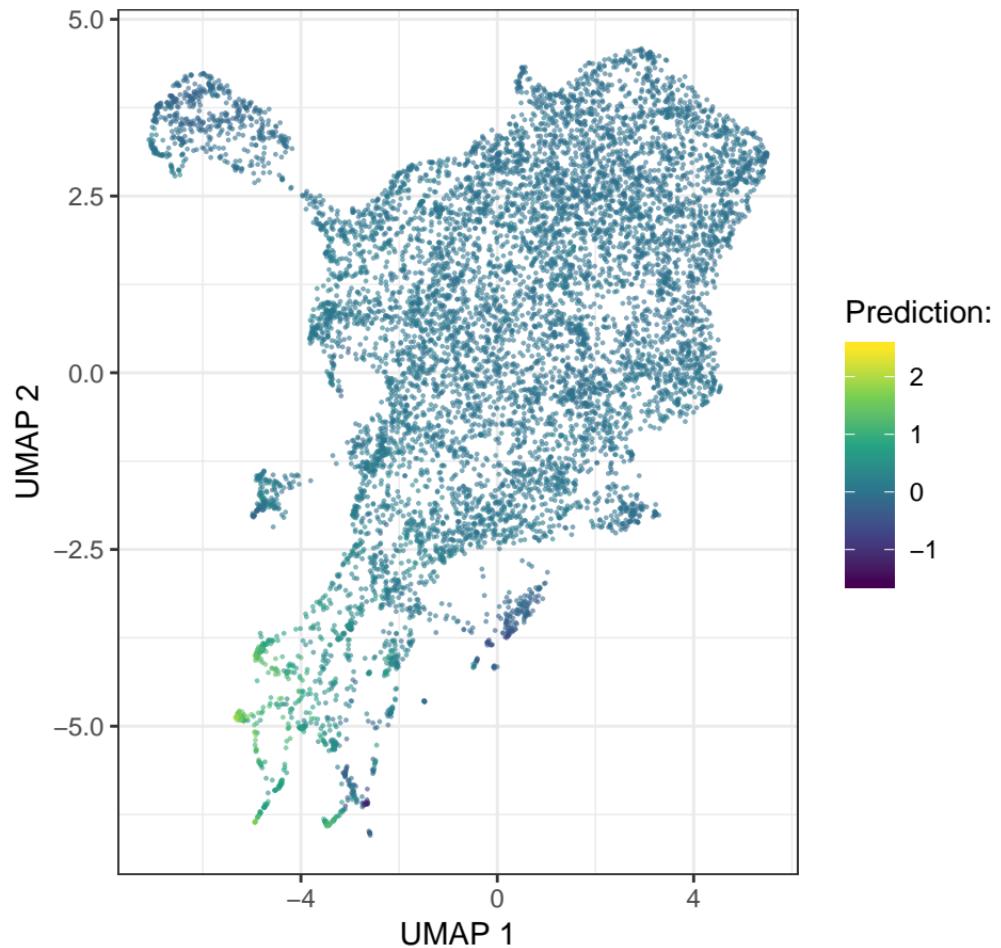
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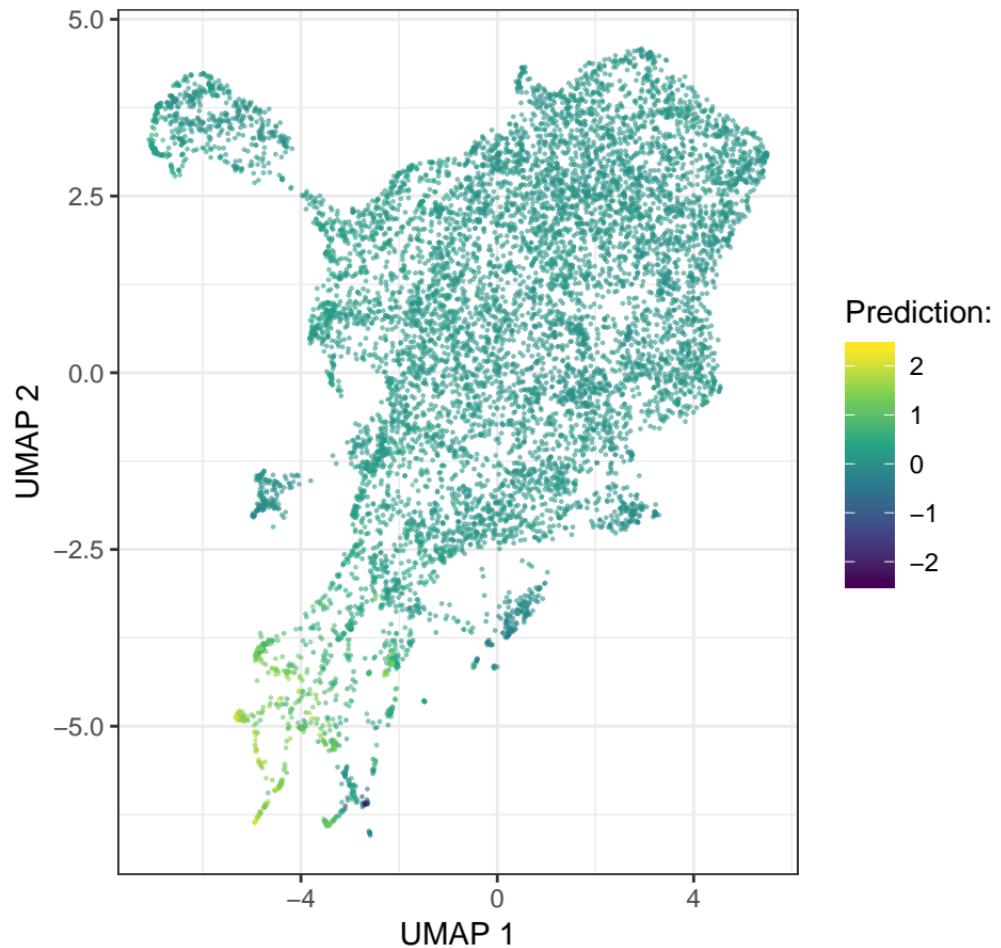
cc\_g2\_ph3\_pos\_n\_objects



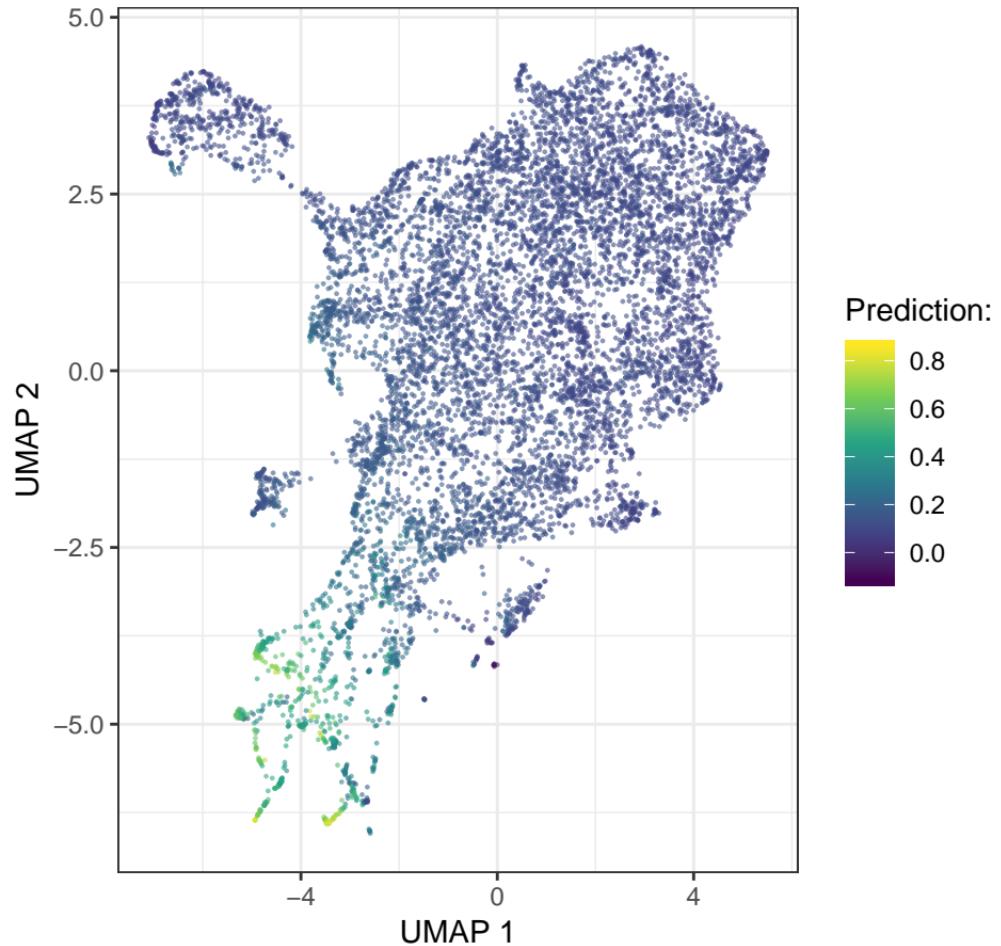
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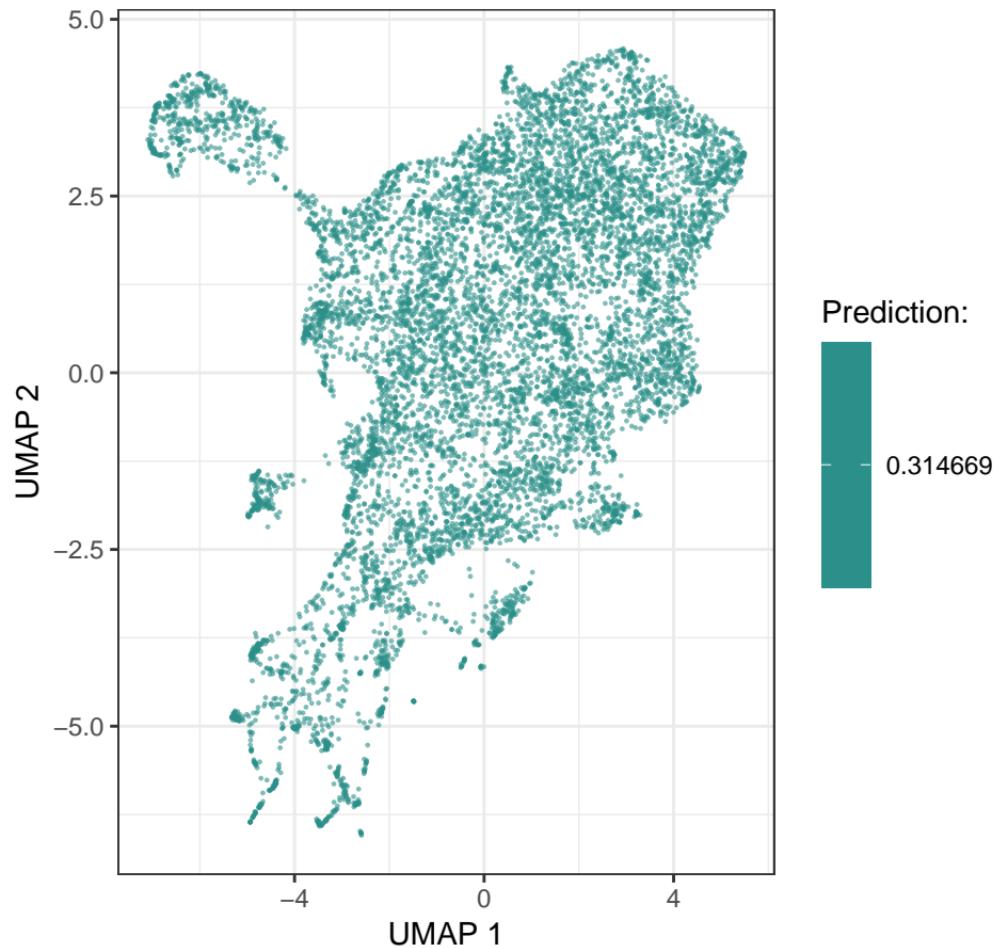
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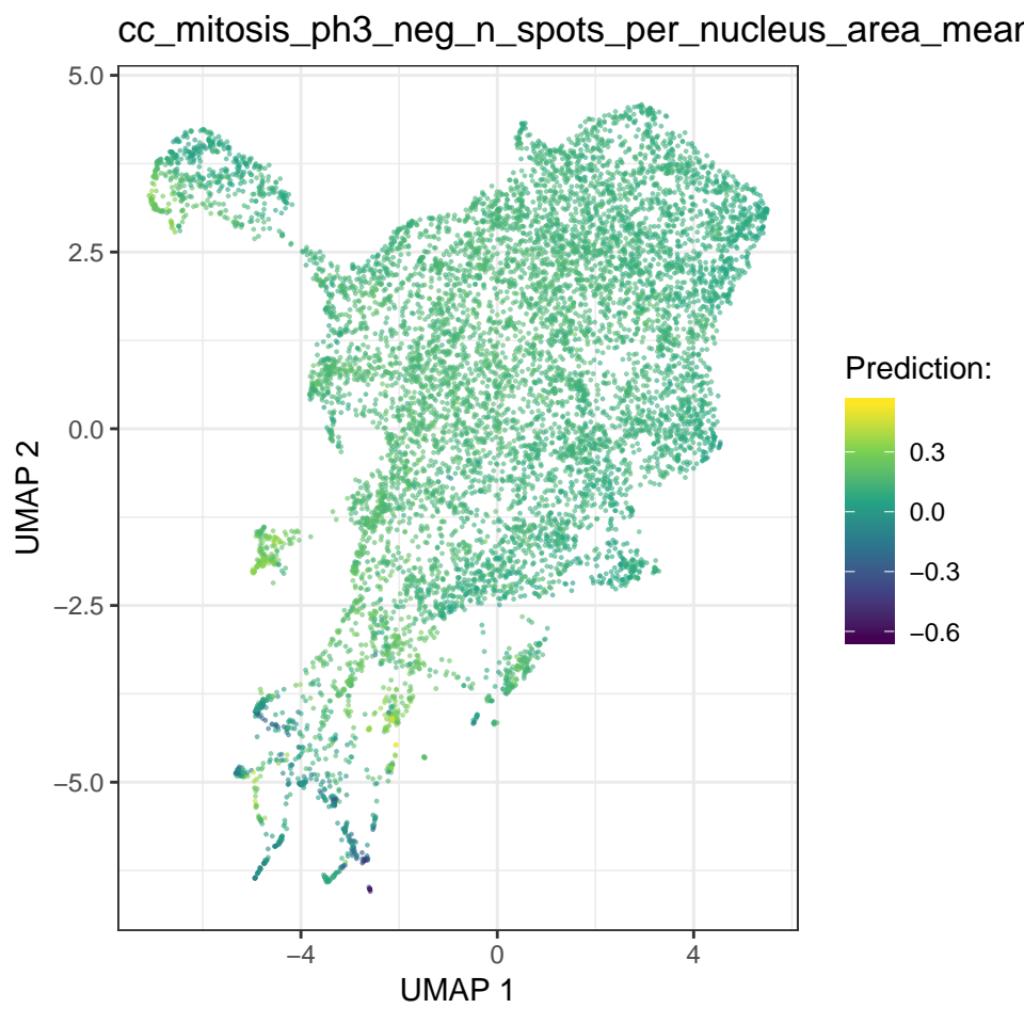


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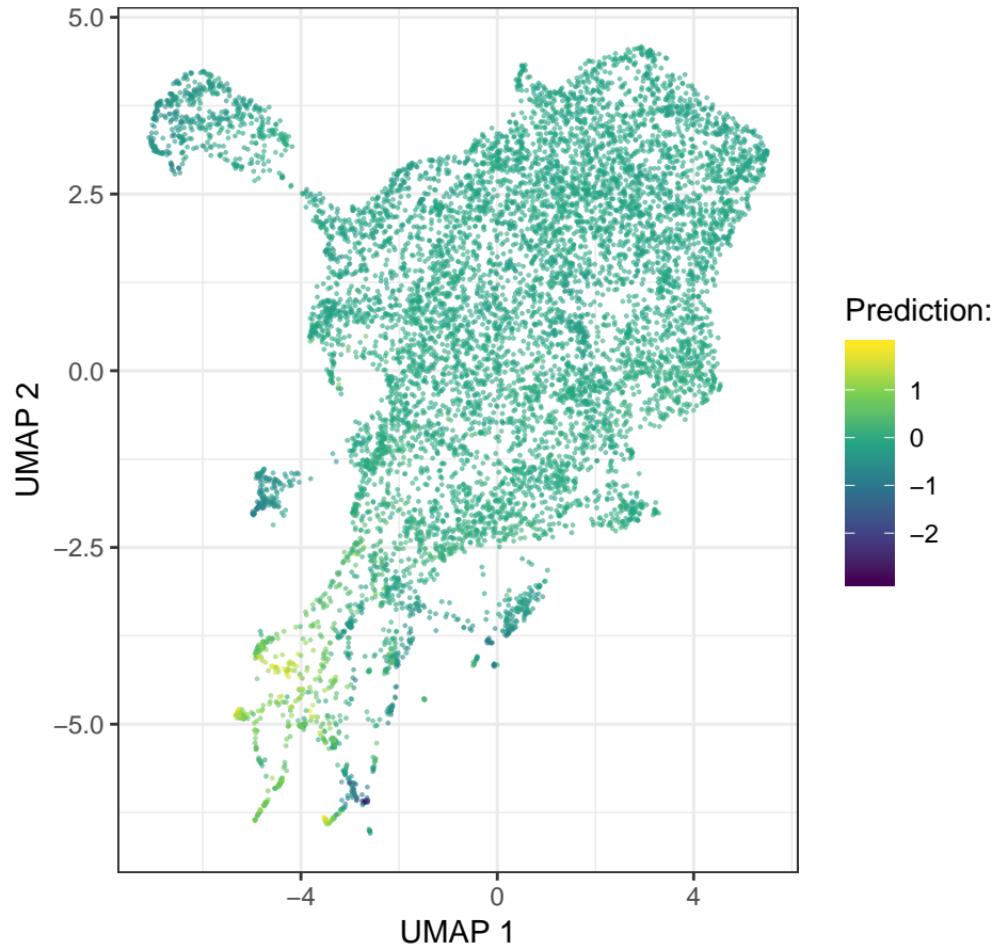


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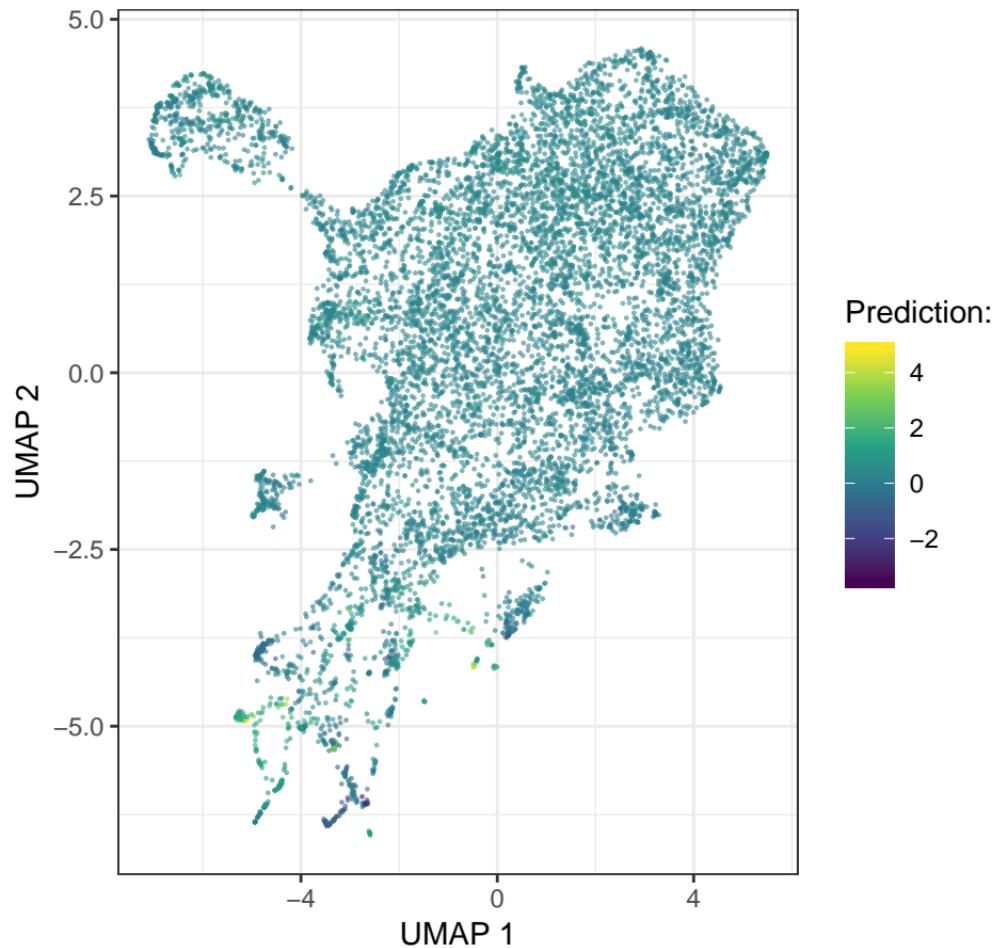




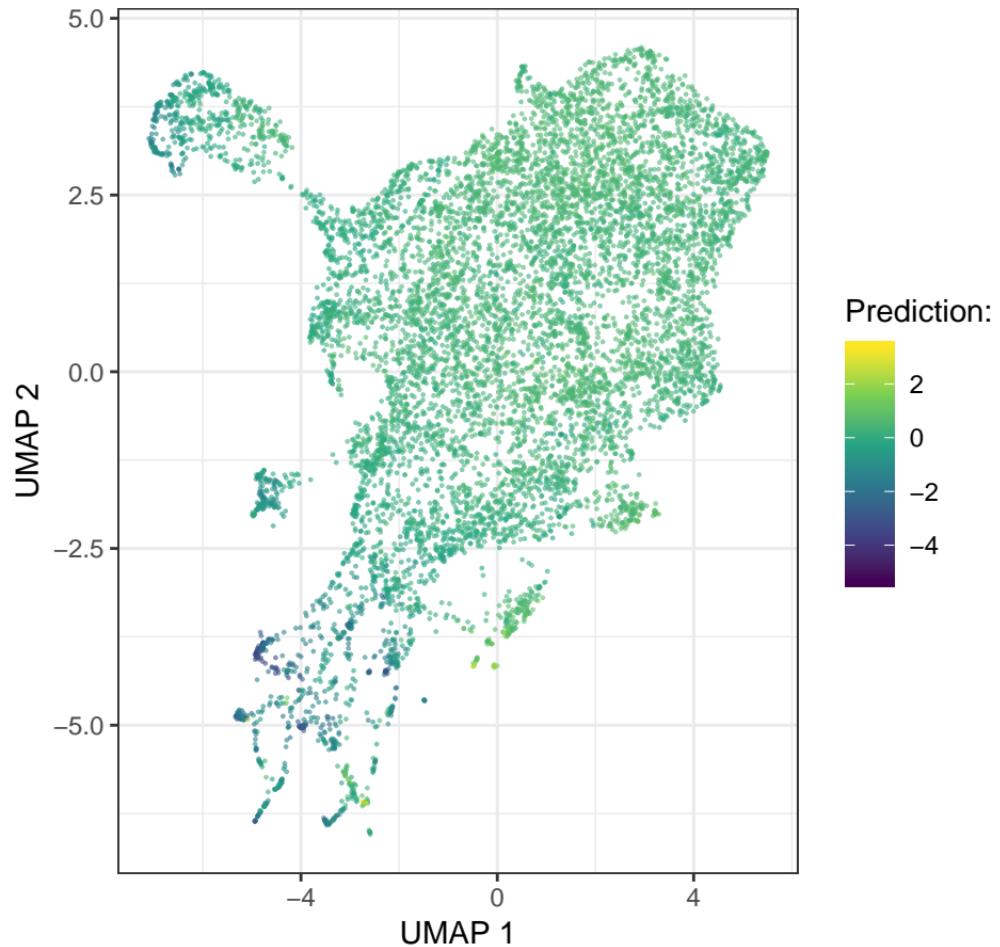
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cc\_all\_large\_notround\_polinuclear\_mean

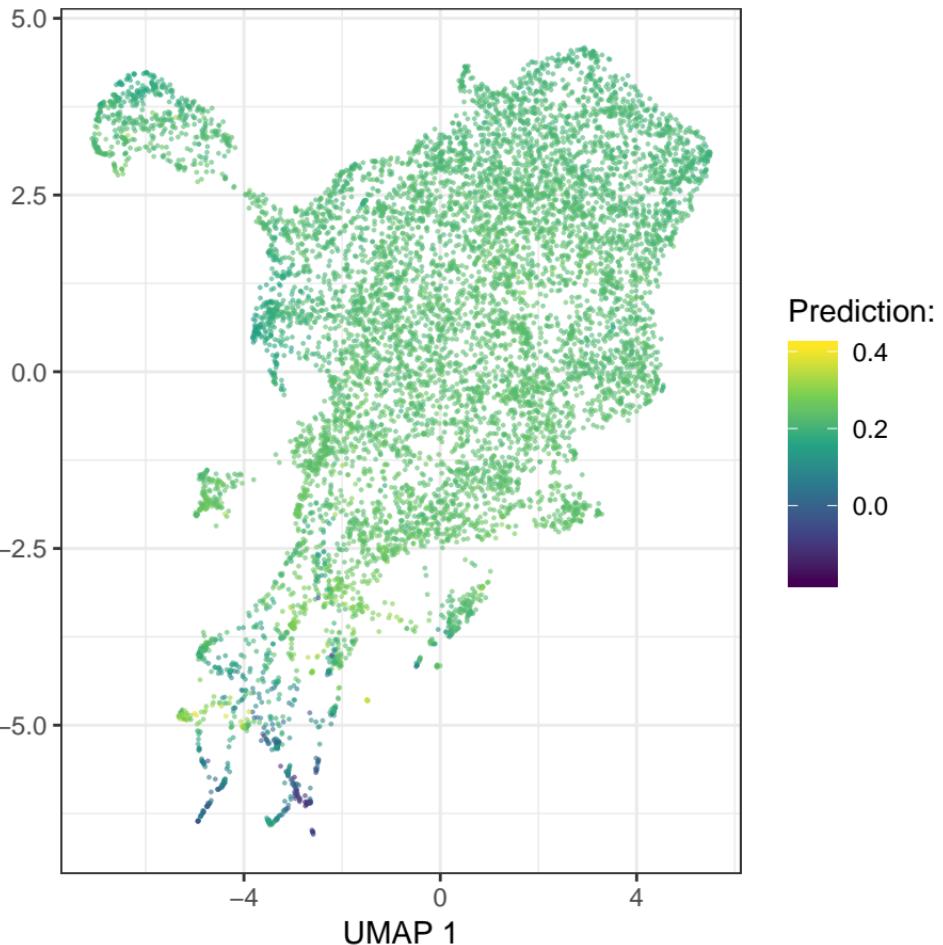


cc\_cc\_edu\_pos\_mean

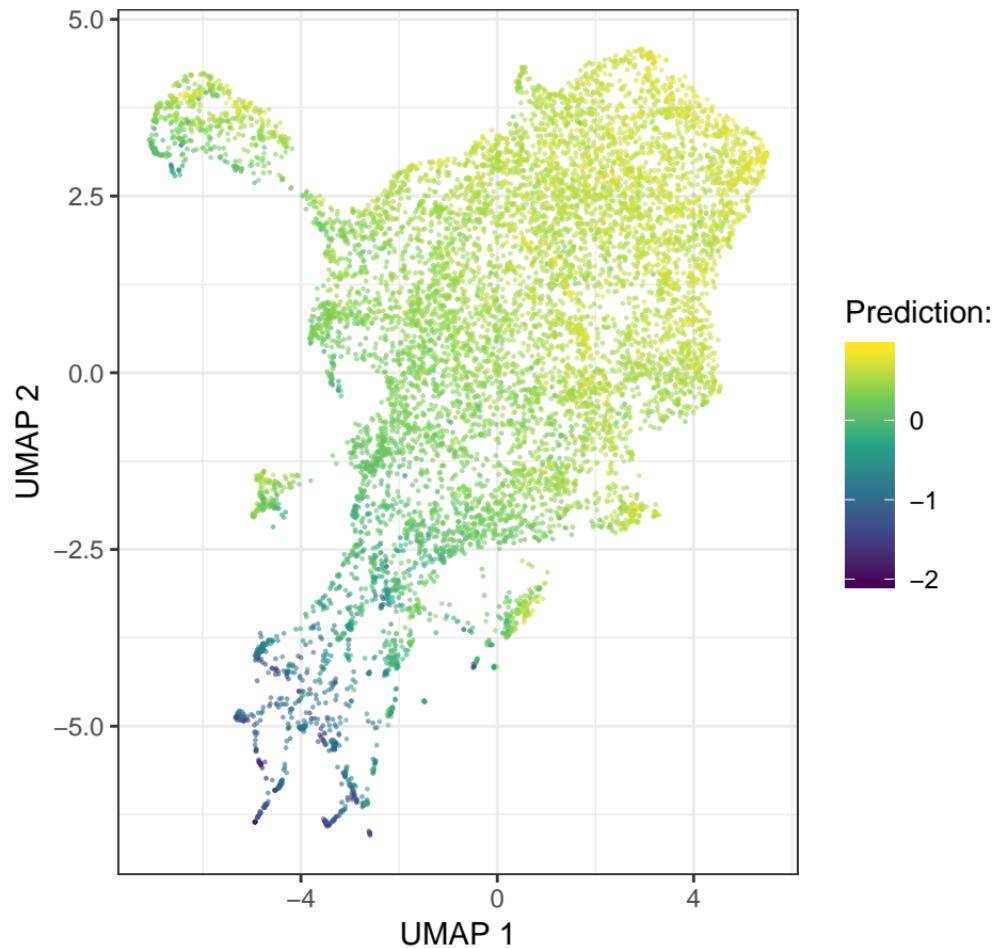


cc\_mitosis\_ph3\_pos\_n\_spots\_per\_nucleus\_area\_mean

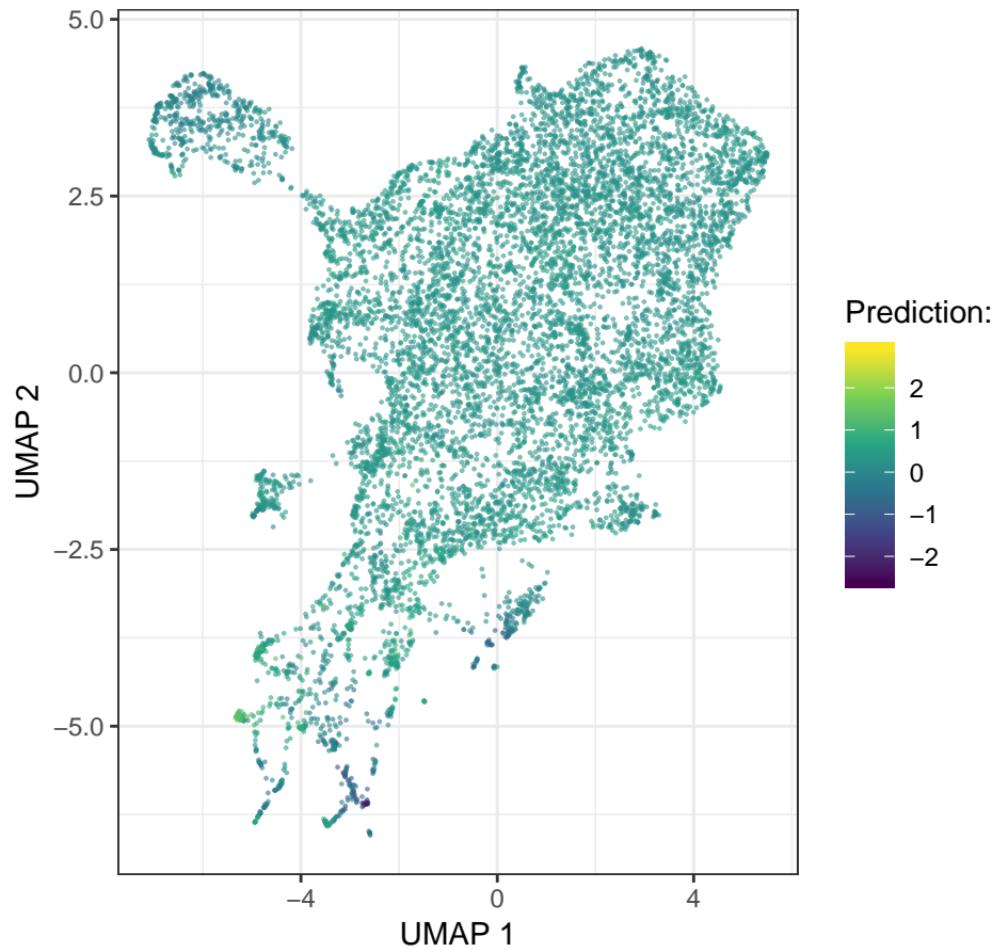
UMAP 2



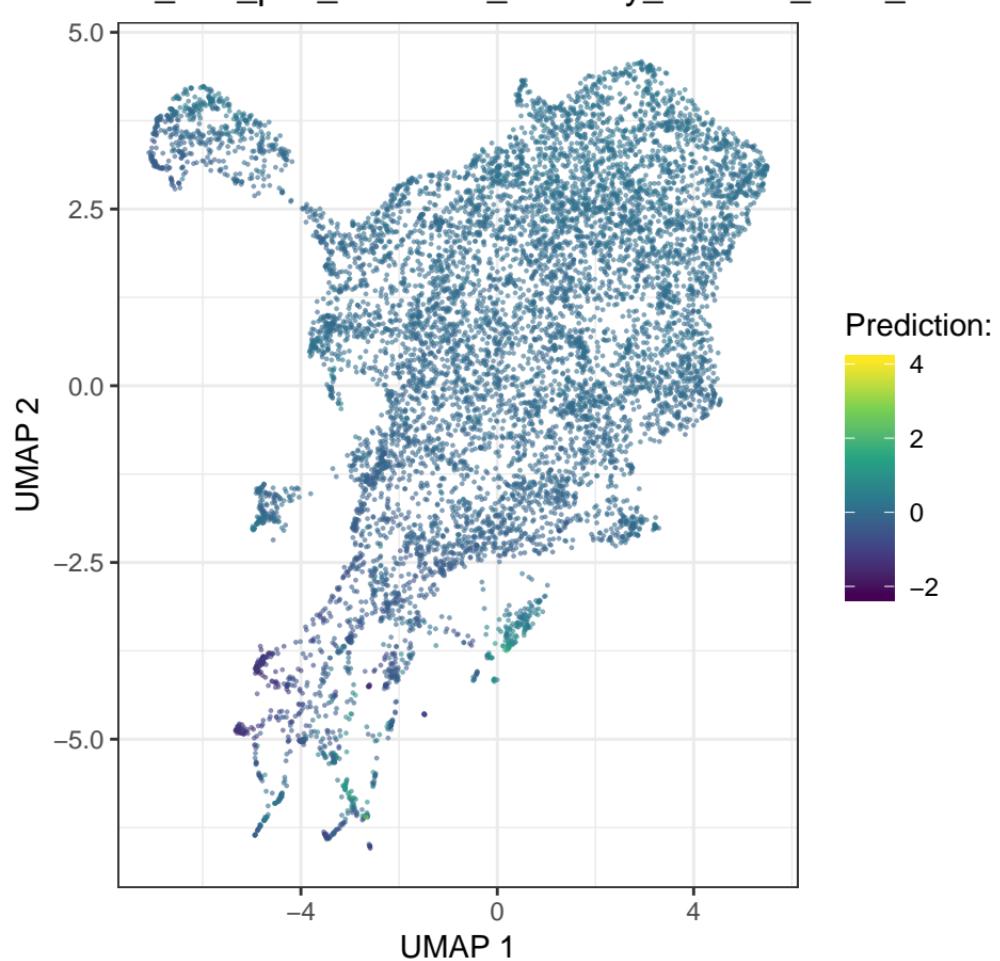
cc\_cc\_n\_objects



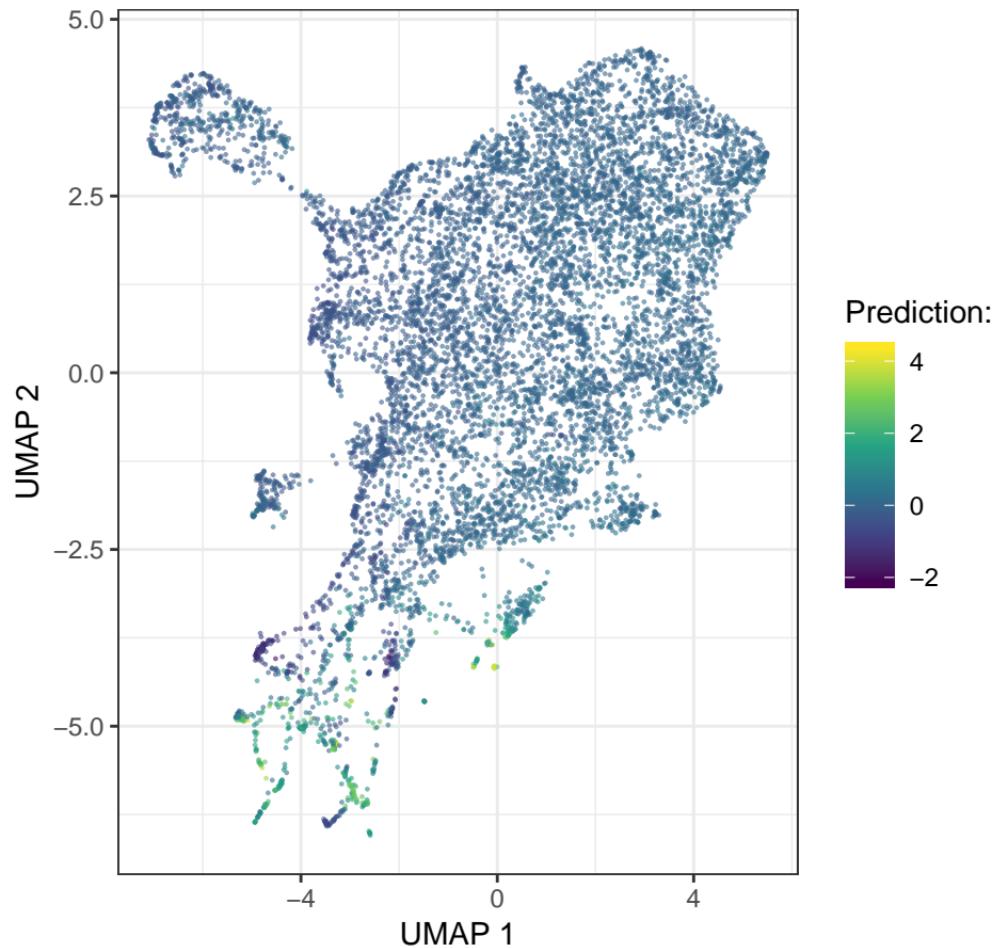
cc\_edu\_pos\_n\_spots\_mean



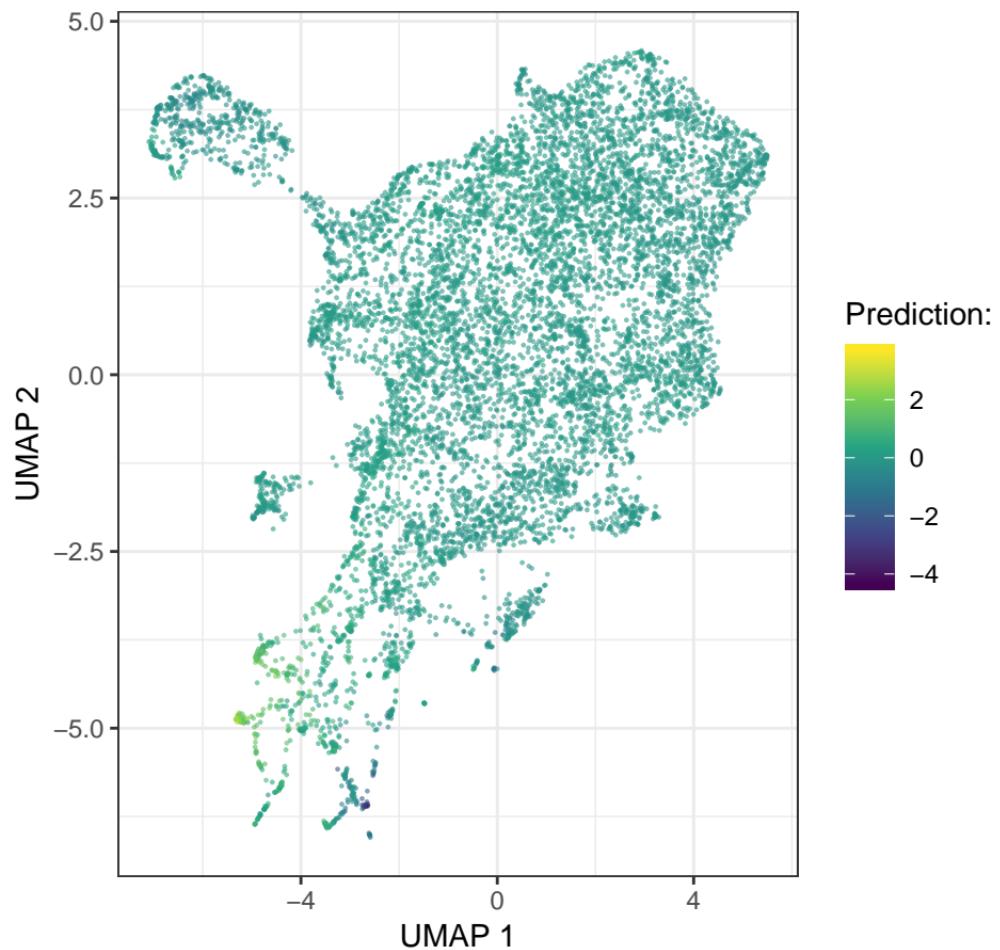
cc\_edu\_pos\_alexa647\_intensity\_nucleus\_area\_mean



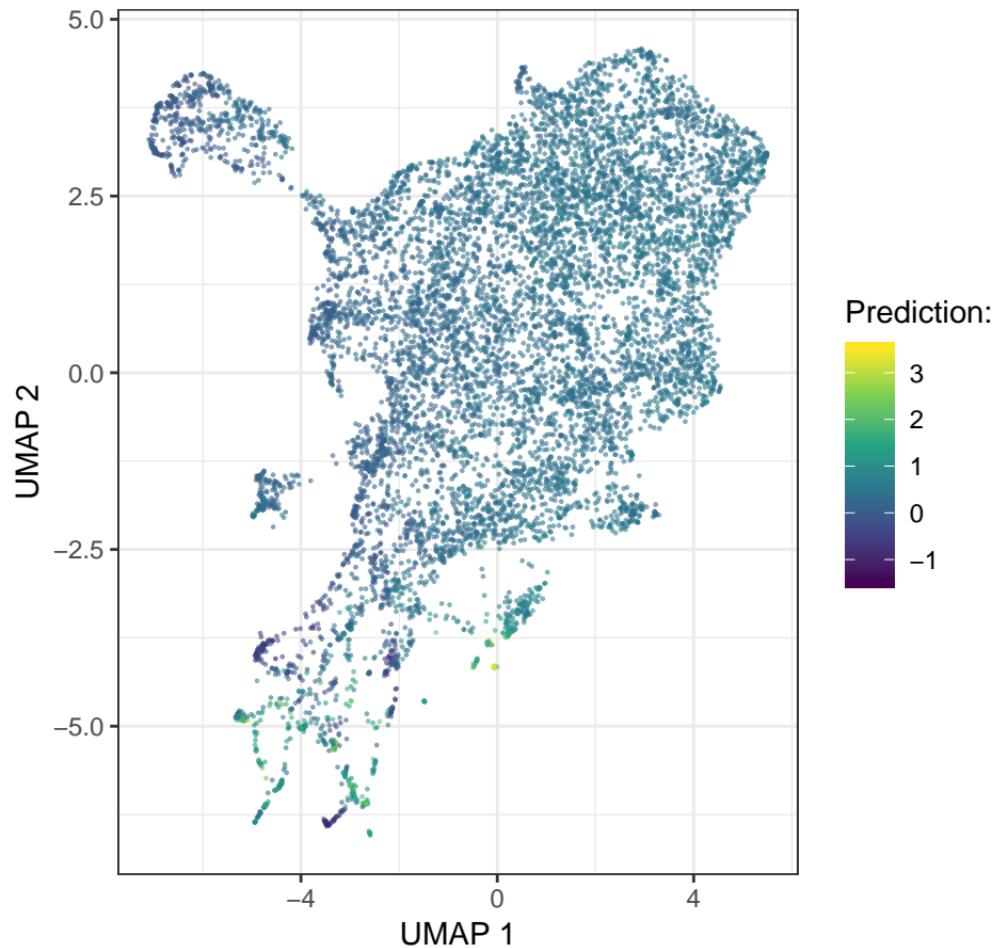
cc\_cc\_ph3\_neg\_hoechst\_late\_mitosis\_mean



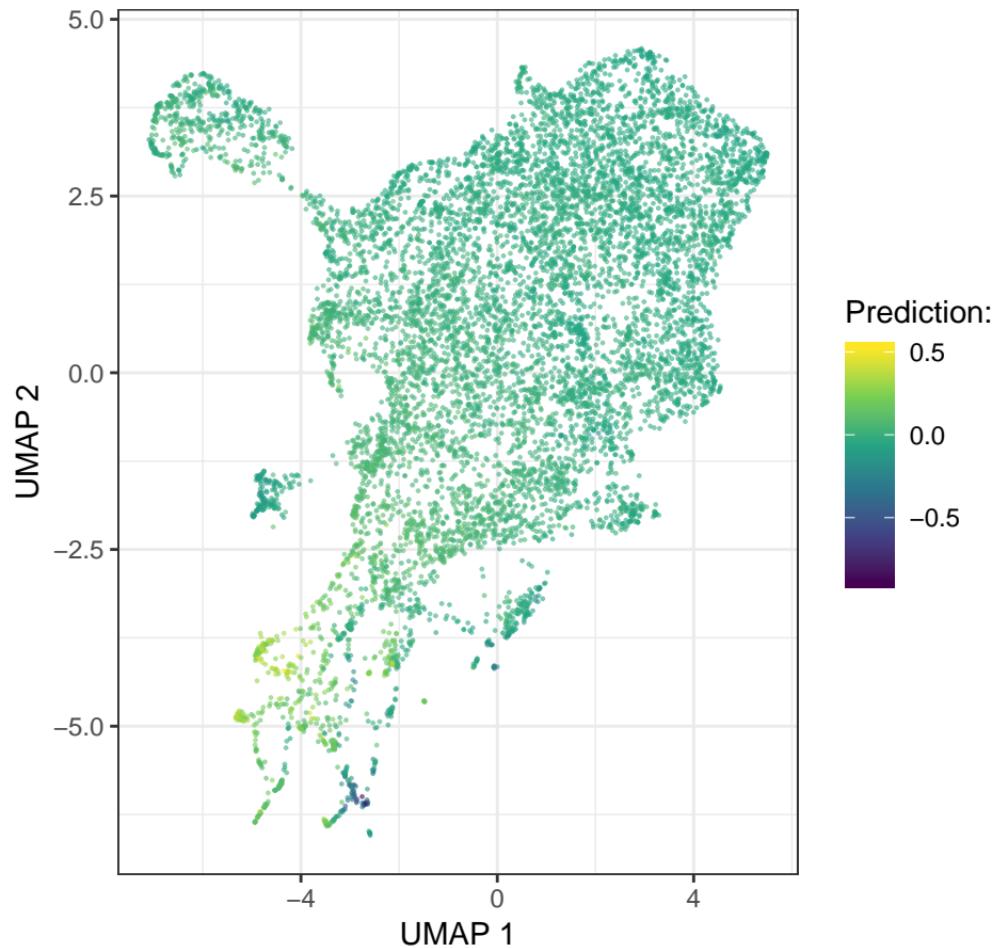
cc\_all\_n\_spots\_per\_nucleus\_area\_mean



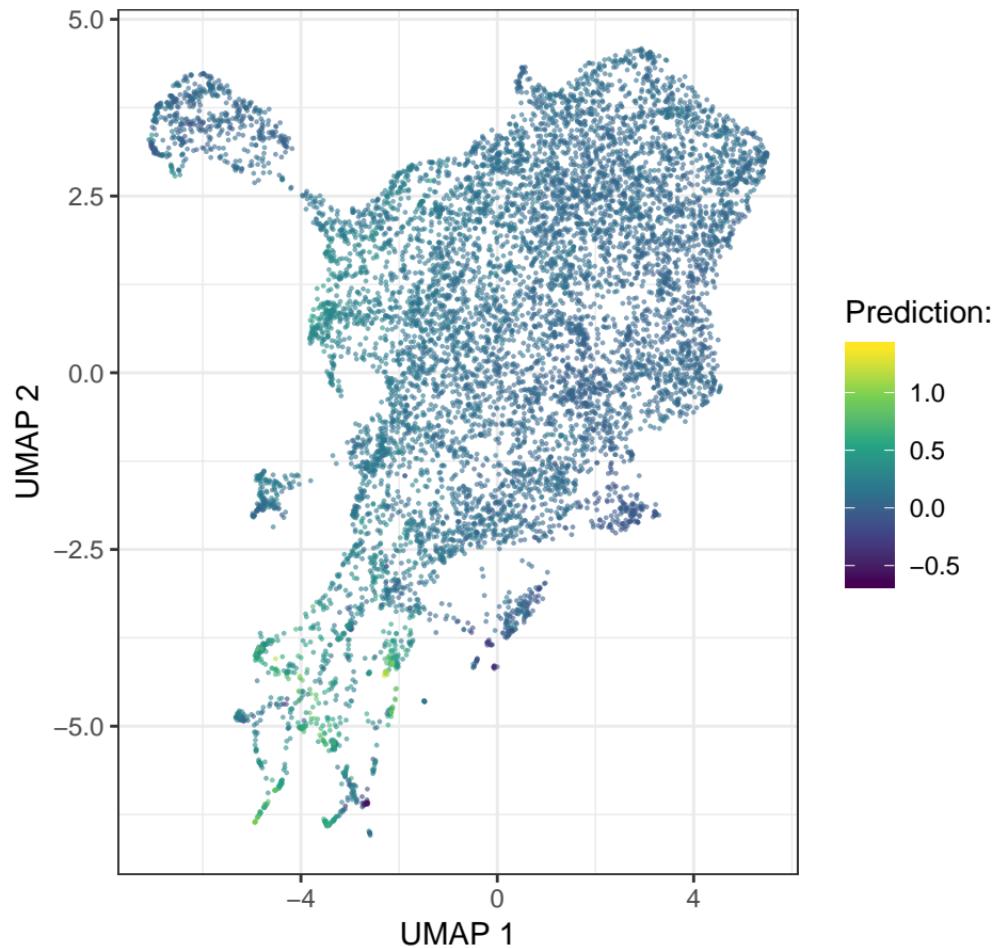
cc\_mitosis\_ph3\_neg\_n\_objects



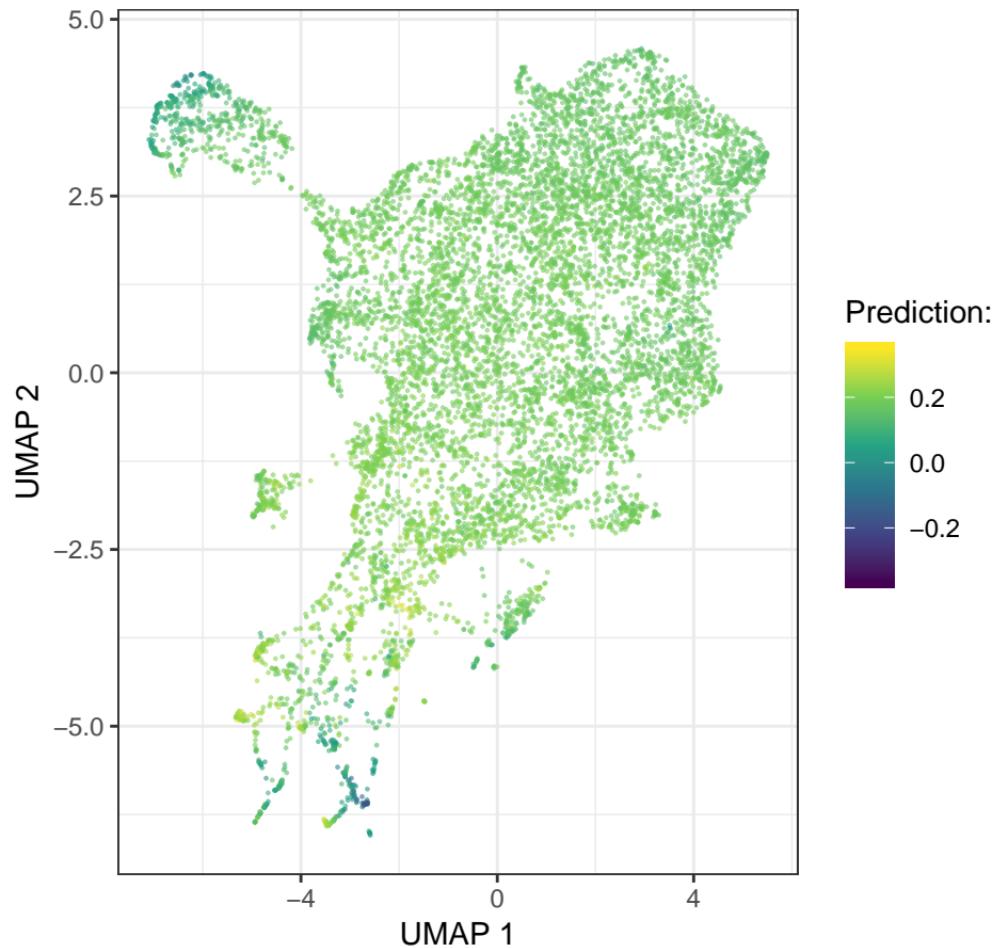
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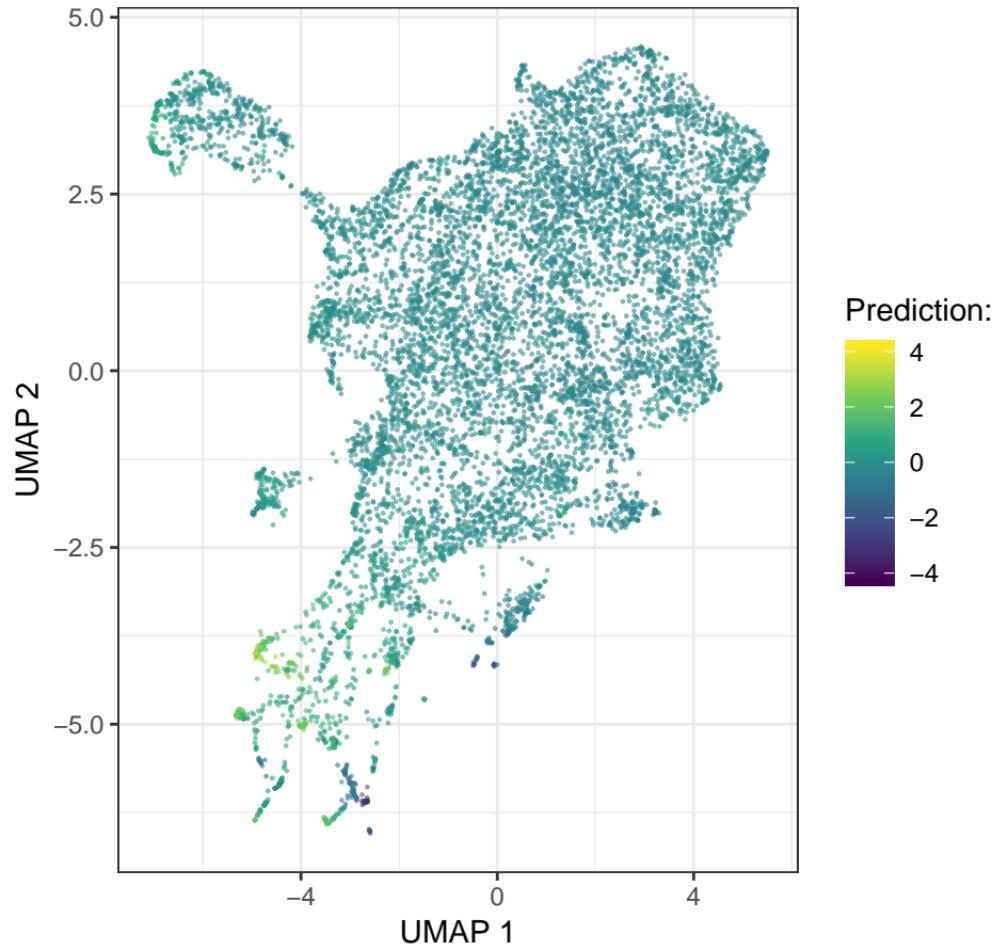
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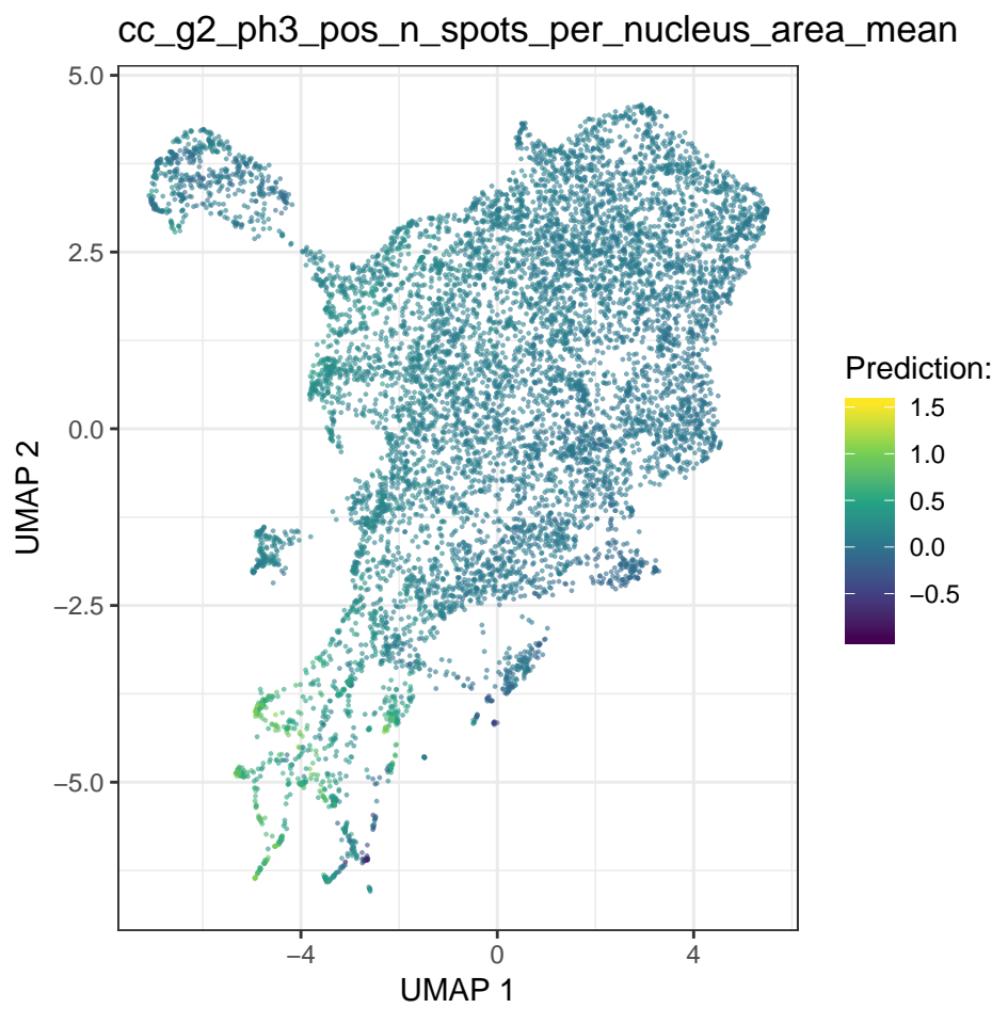


cc\_mitosis\_ph3\_pos\_n\_spots\_mean

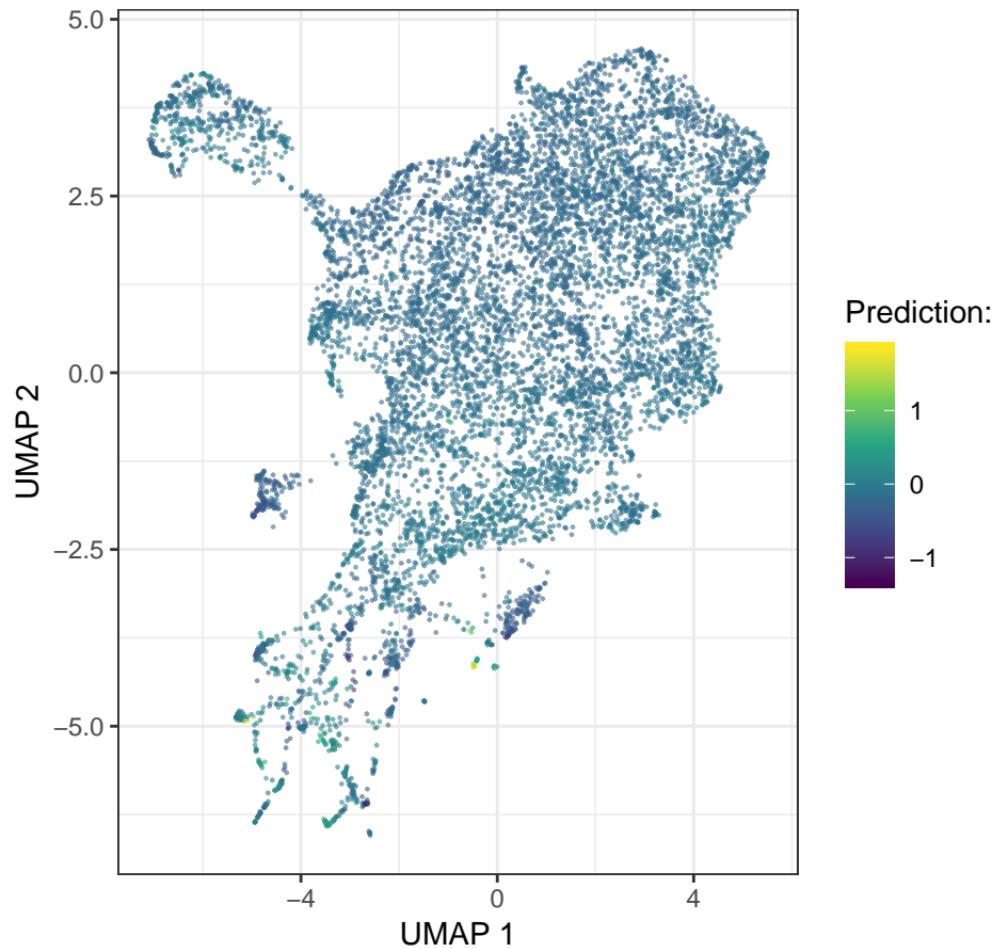


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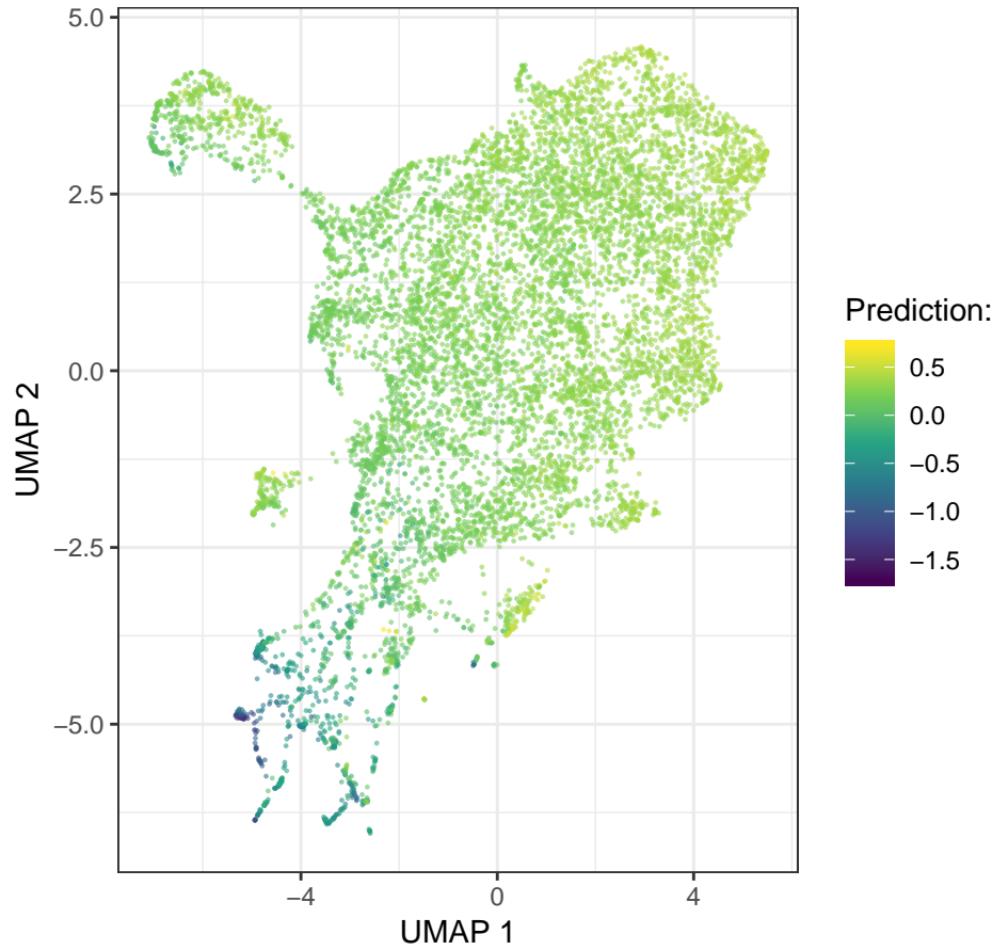




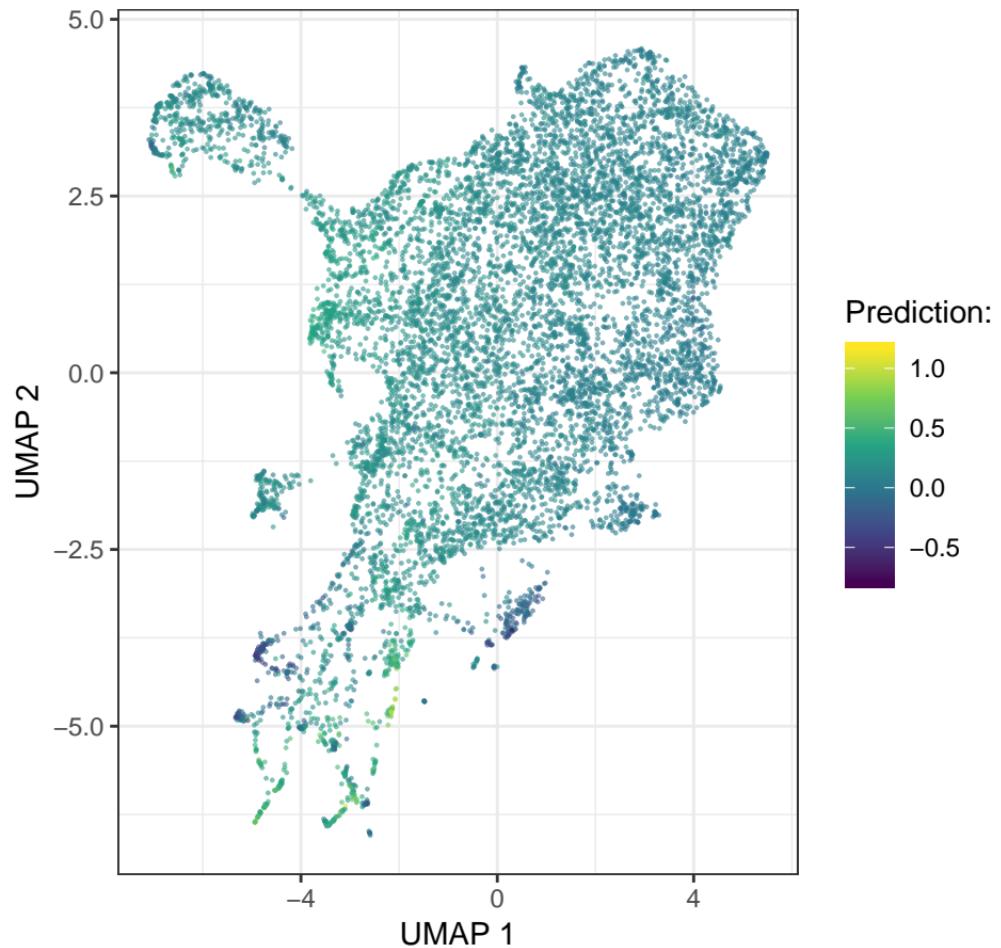
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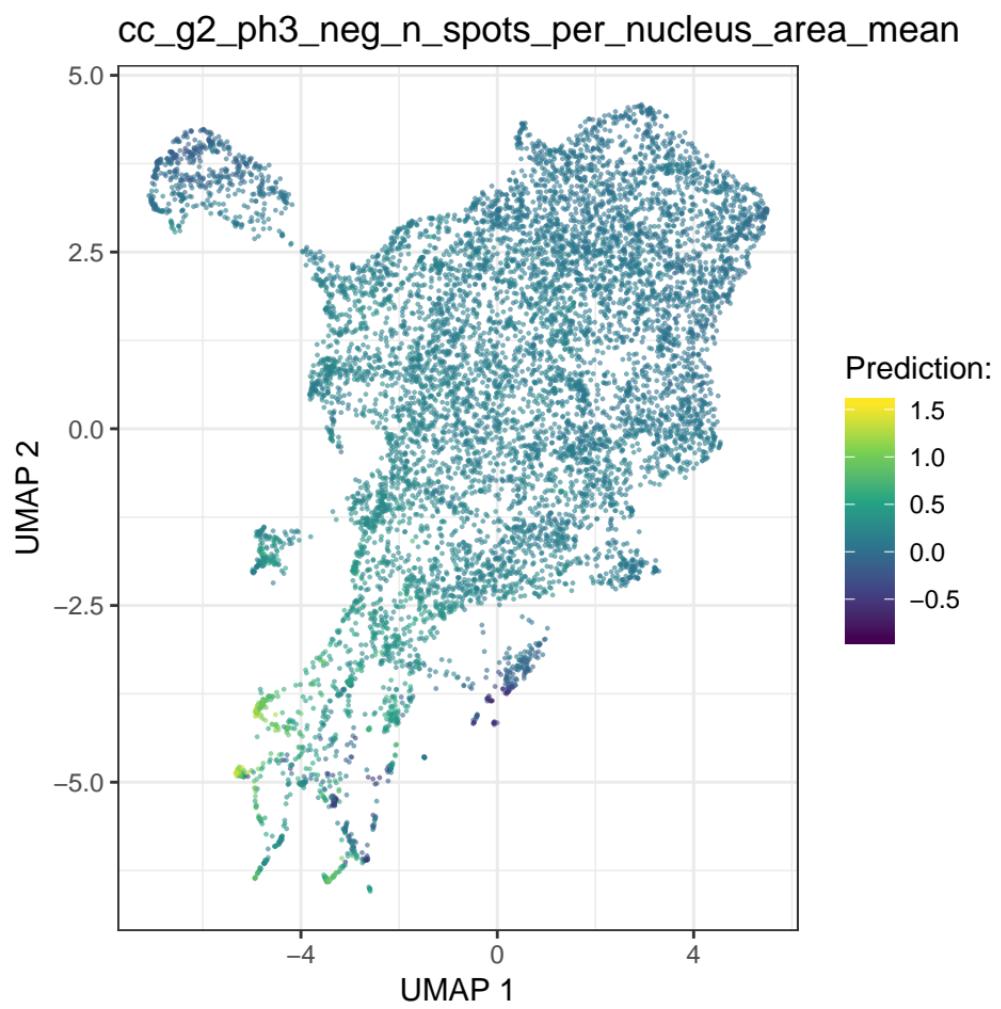


cc\_g2\_ph3\_neg\_n\_objects

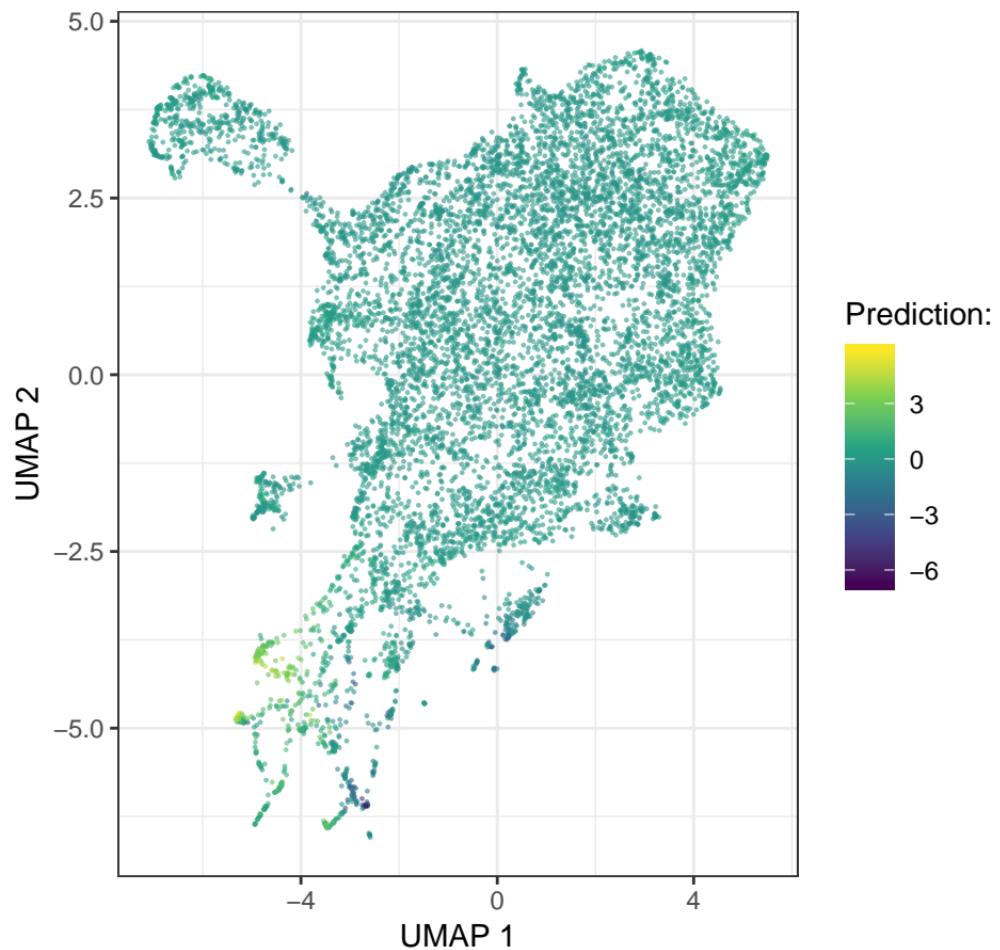


# cc\_cc\_g2\_ph3\_pos\_early\_mitosis\_mean

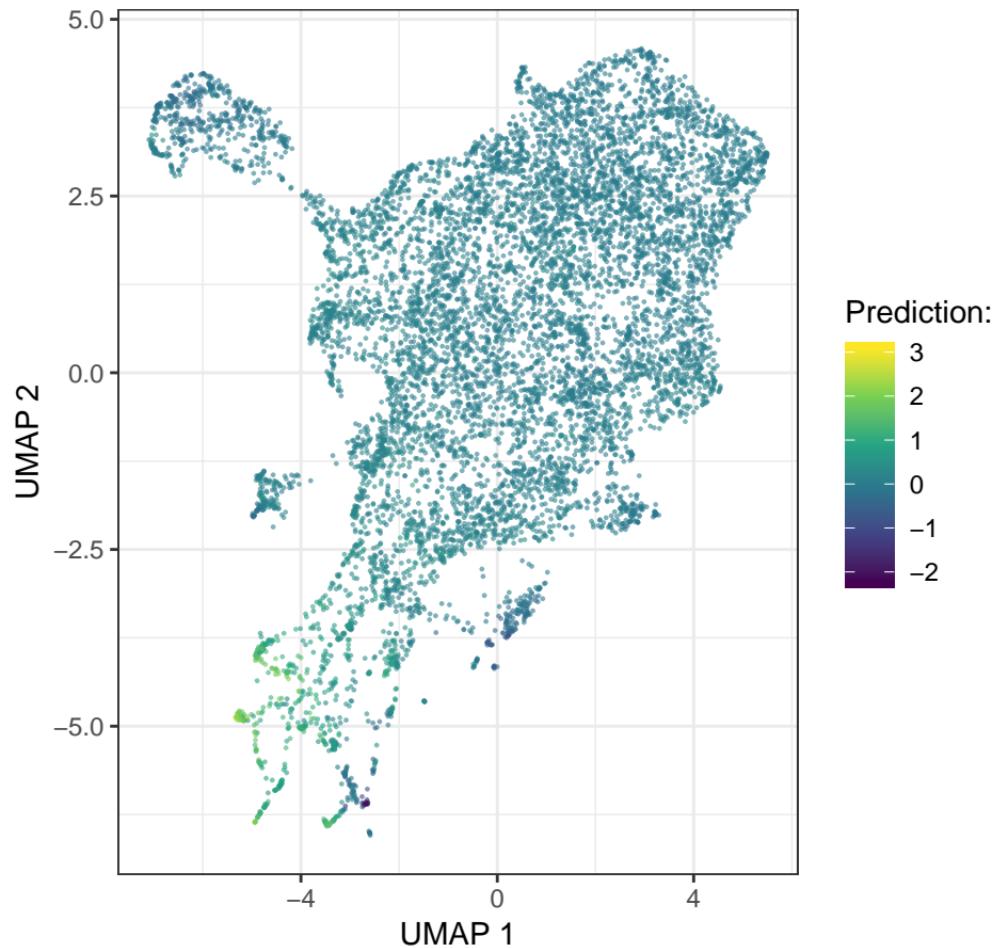




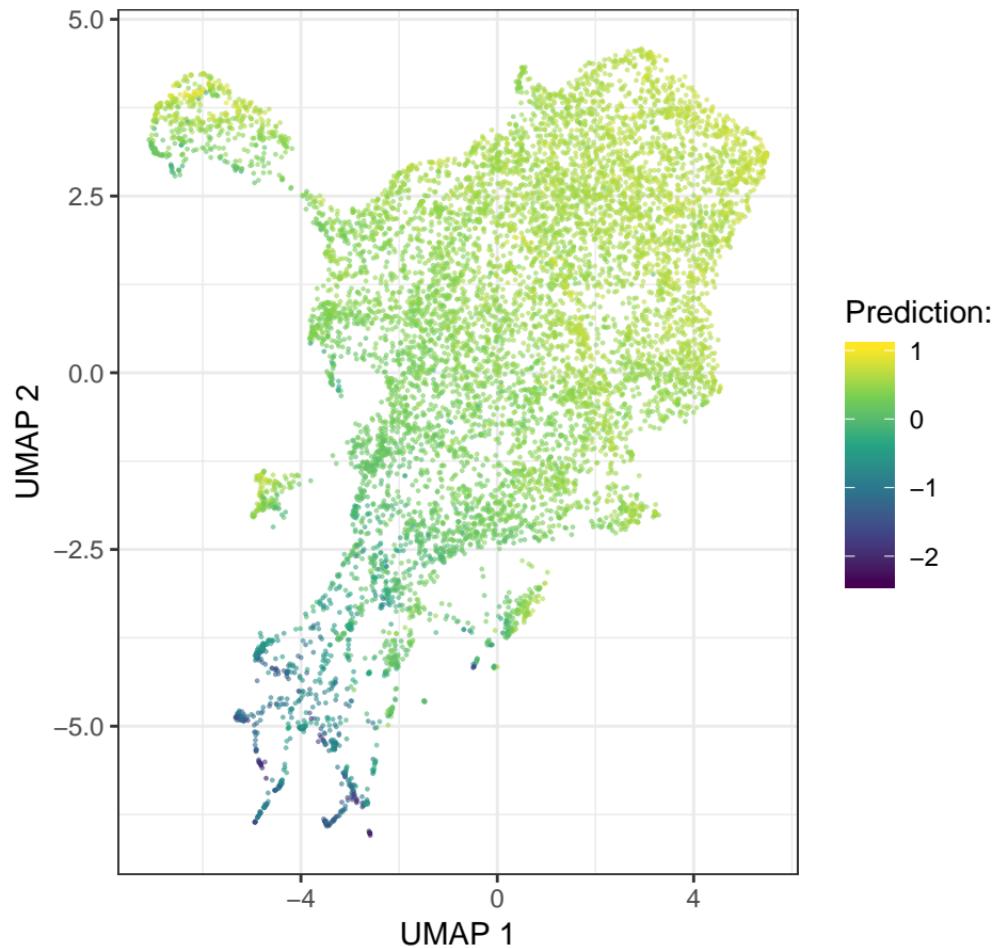
cc\_all\_nucleus\_area\_mean



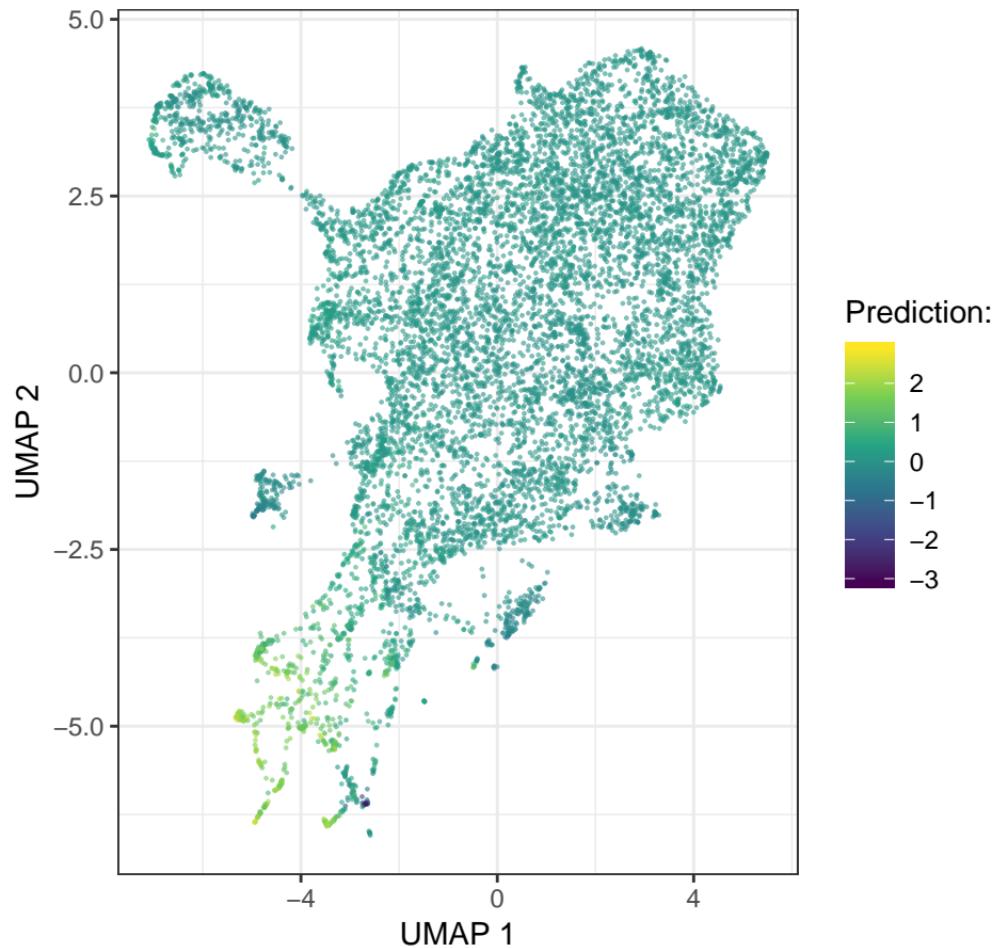
cc\_cc\_n\_spots\_mean



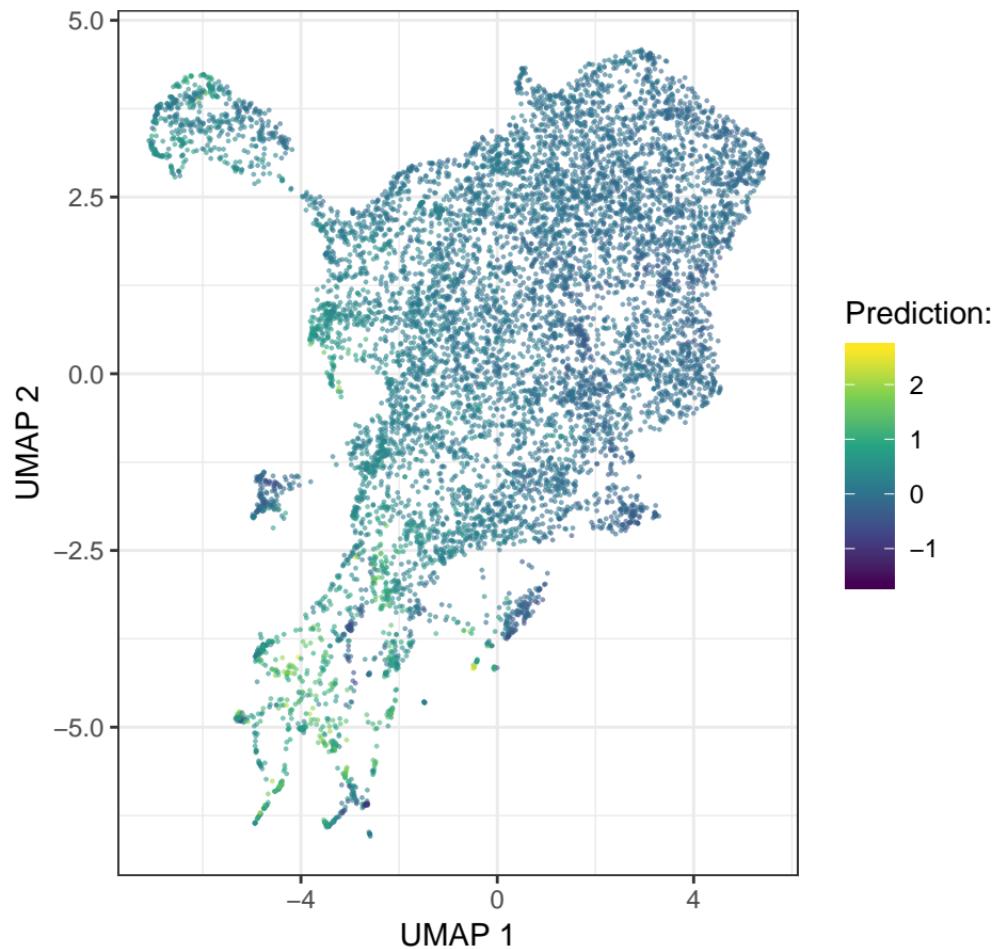
cc\_g1\_n\_objects



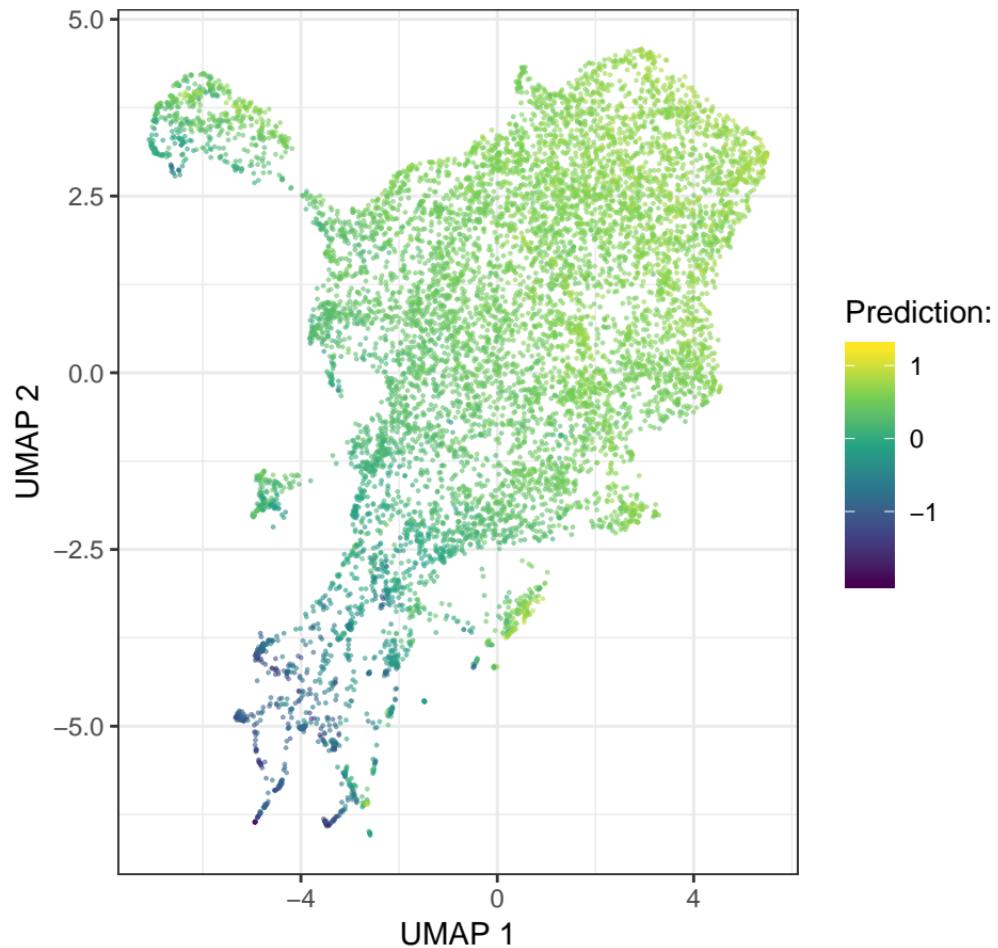
cc\_all\_high\_n\_spots\_h2ax\_mean



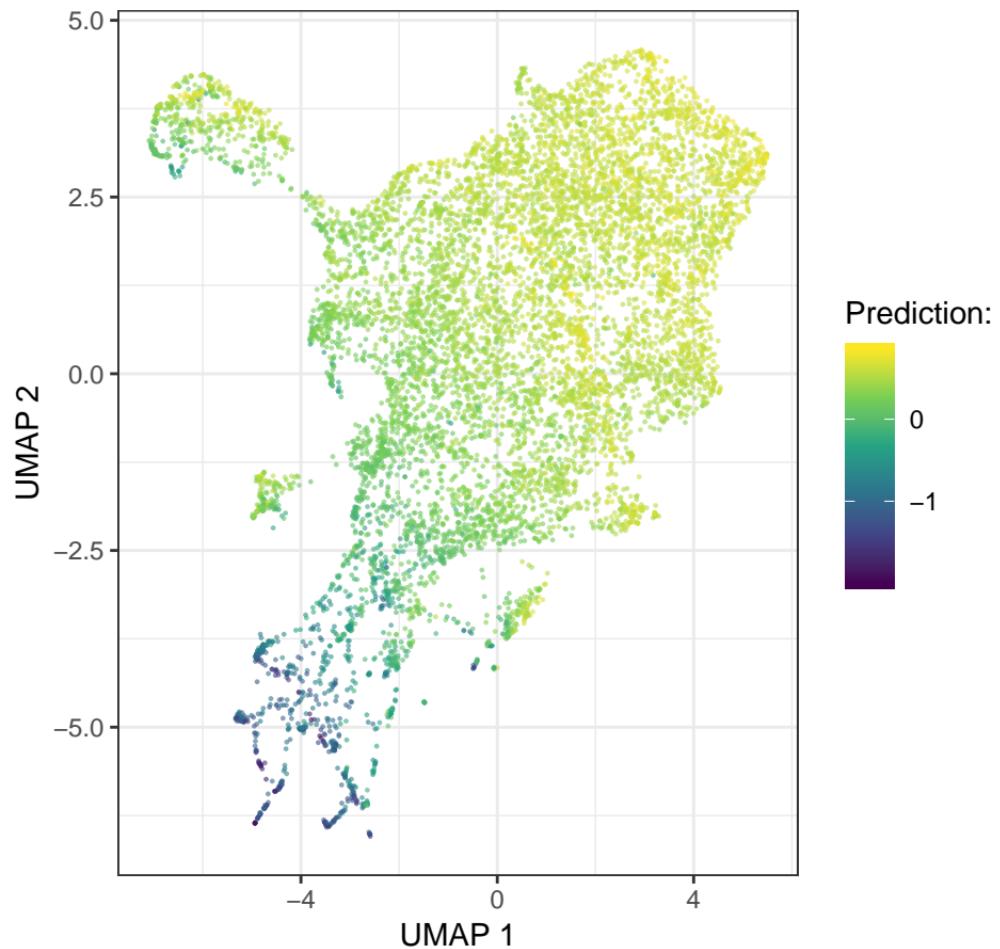
cc\_all\_large\_round\_polyplloid\_mean



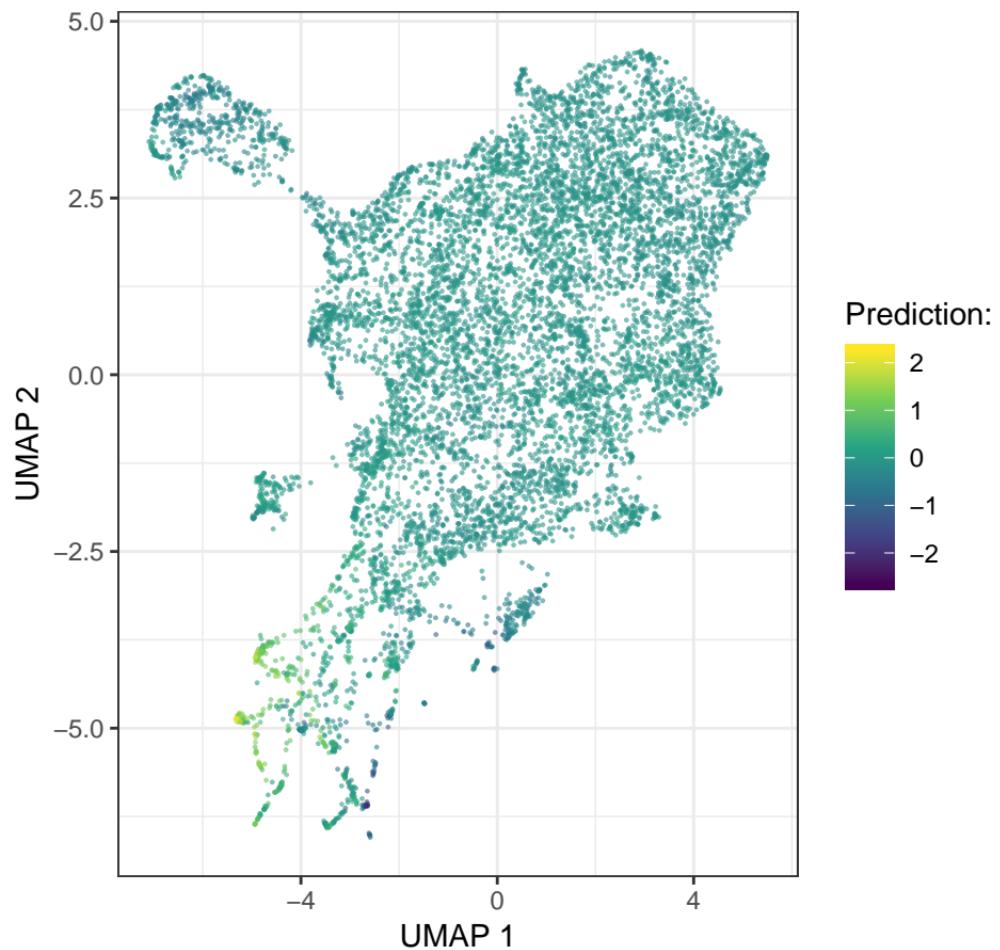
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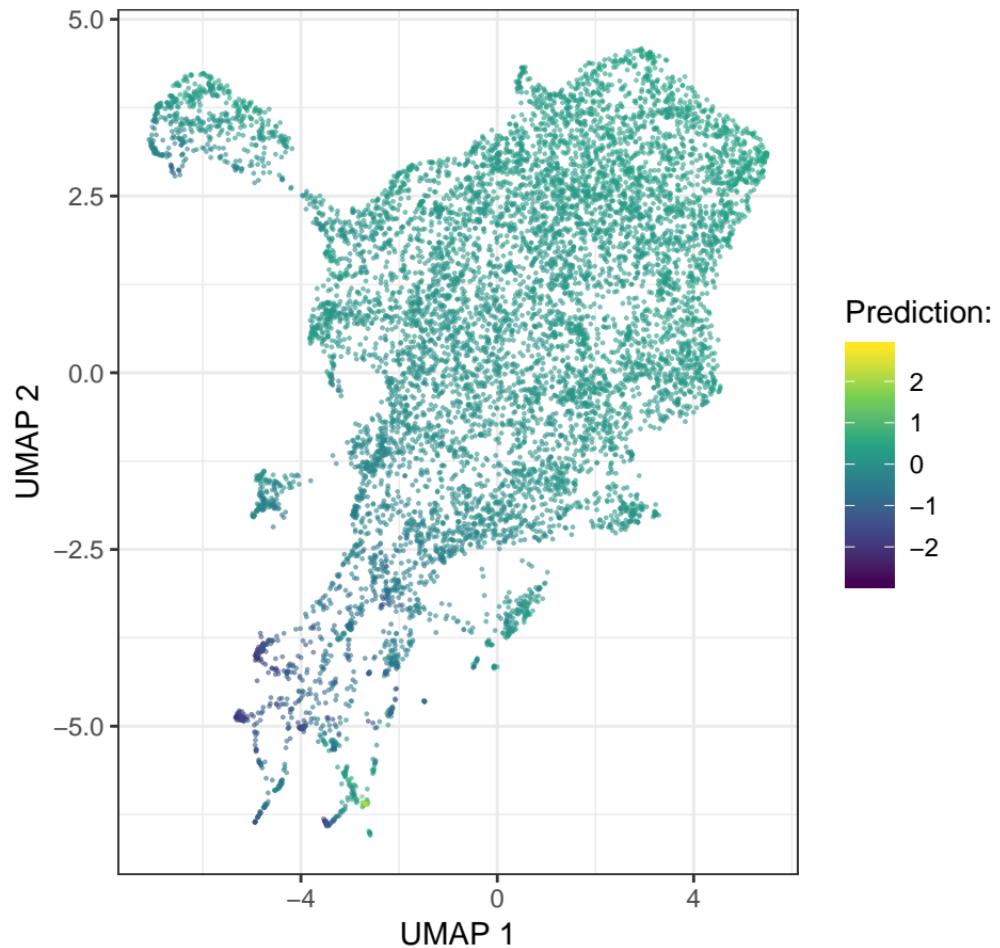
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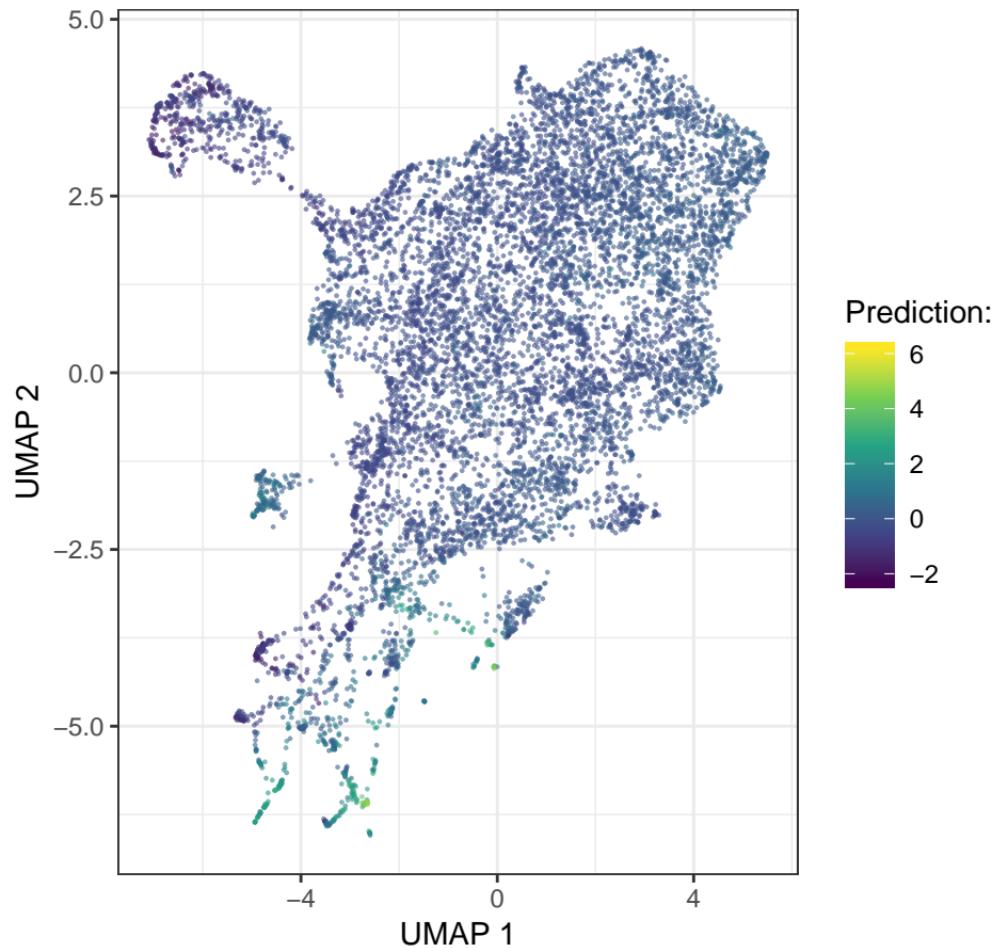
cc\_g1\_n\_spots\_per\_nucleus\_area\_mean



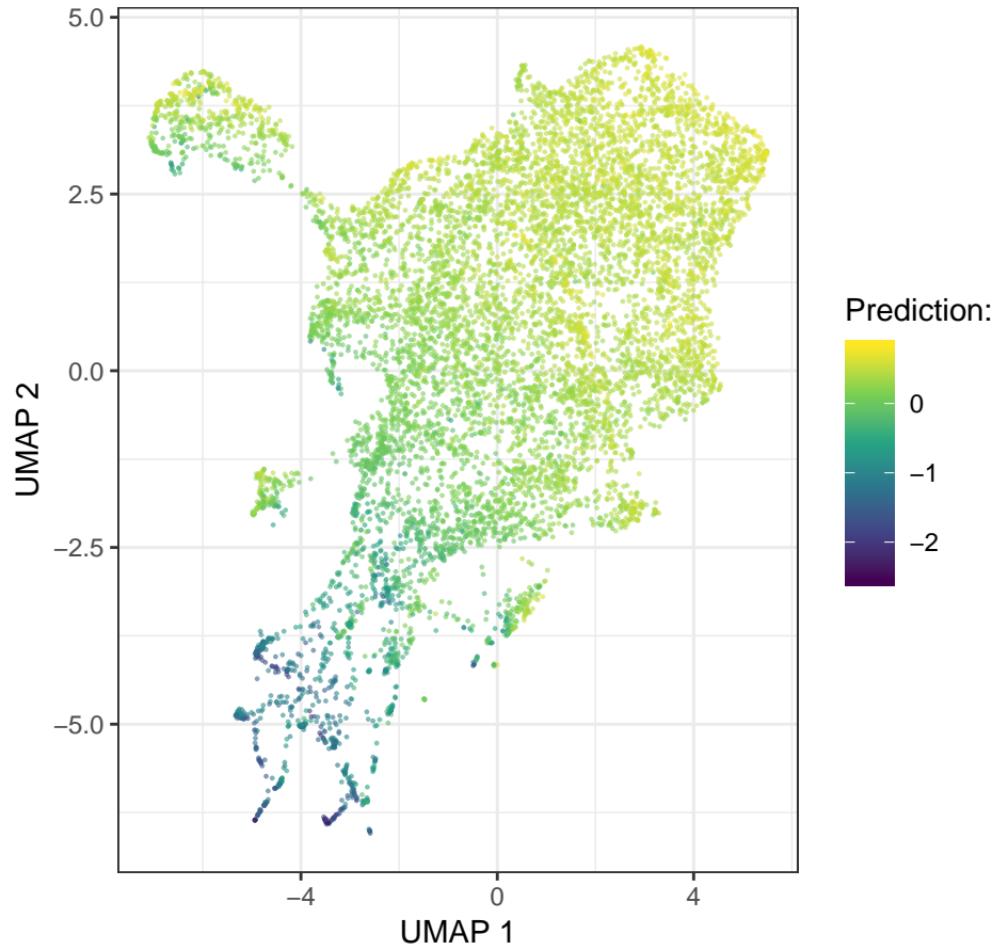
vb\_live\_cell\_roundness



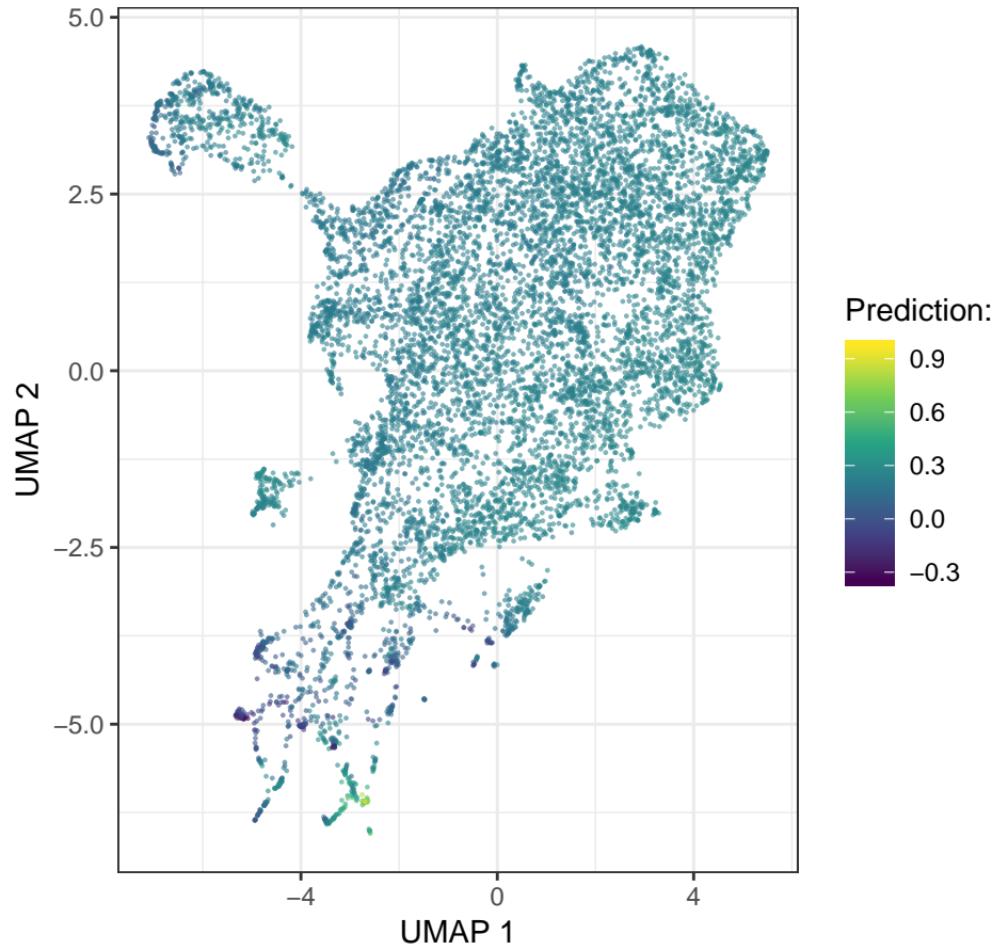
vb\_ros\_mean



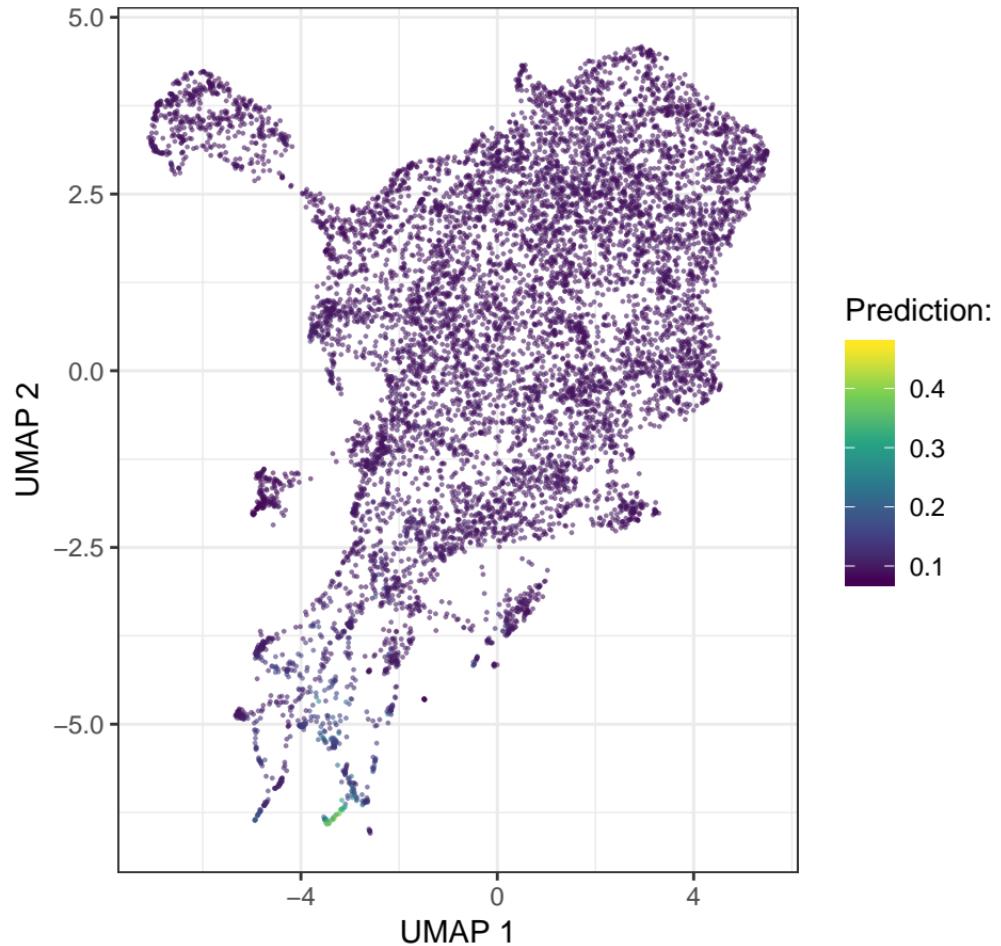
vb\_num\_live\_cells



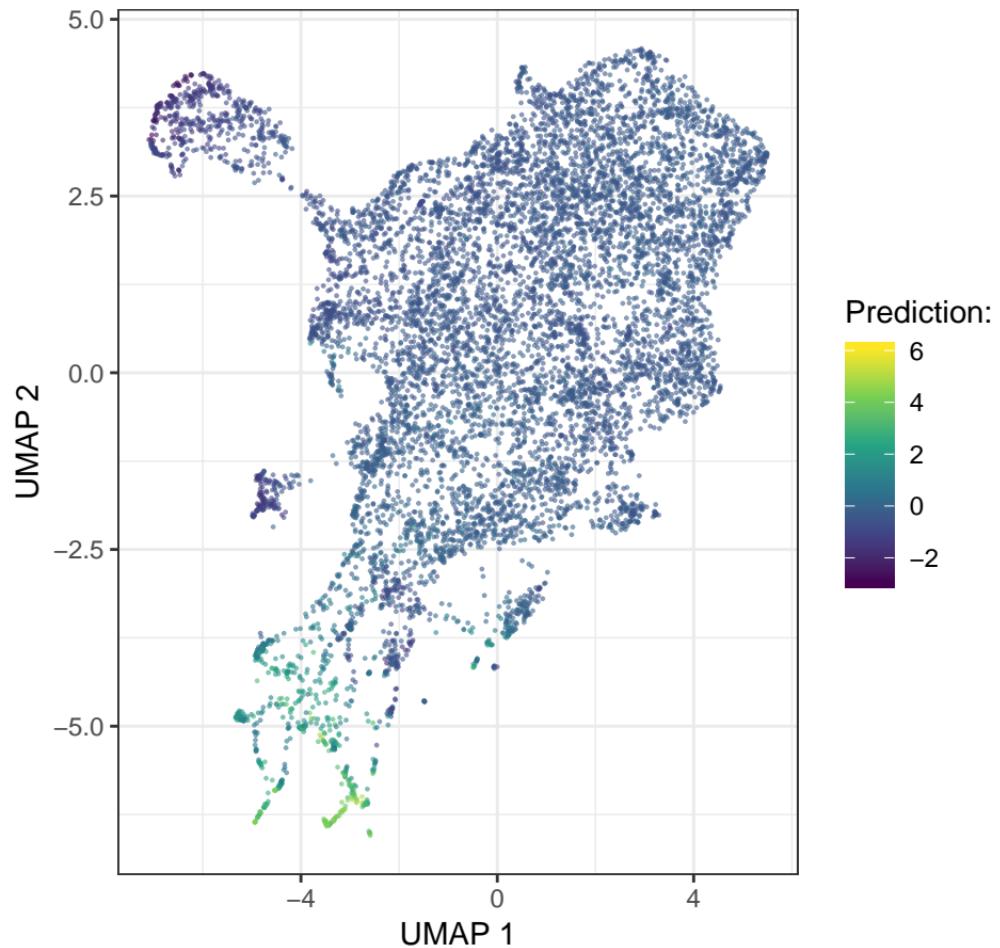
vb\_percent\_caspase\_dead\_only



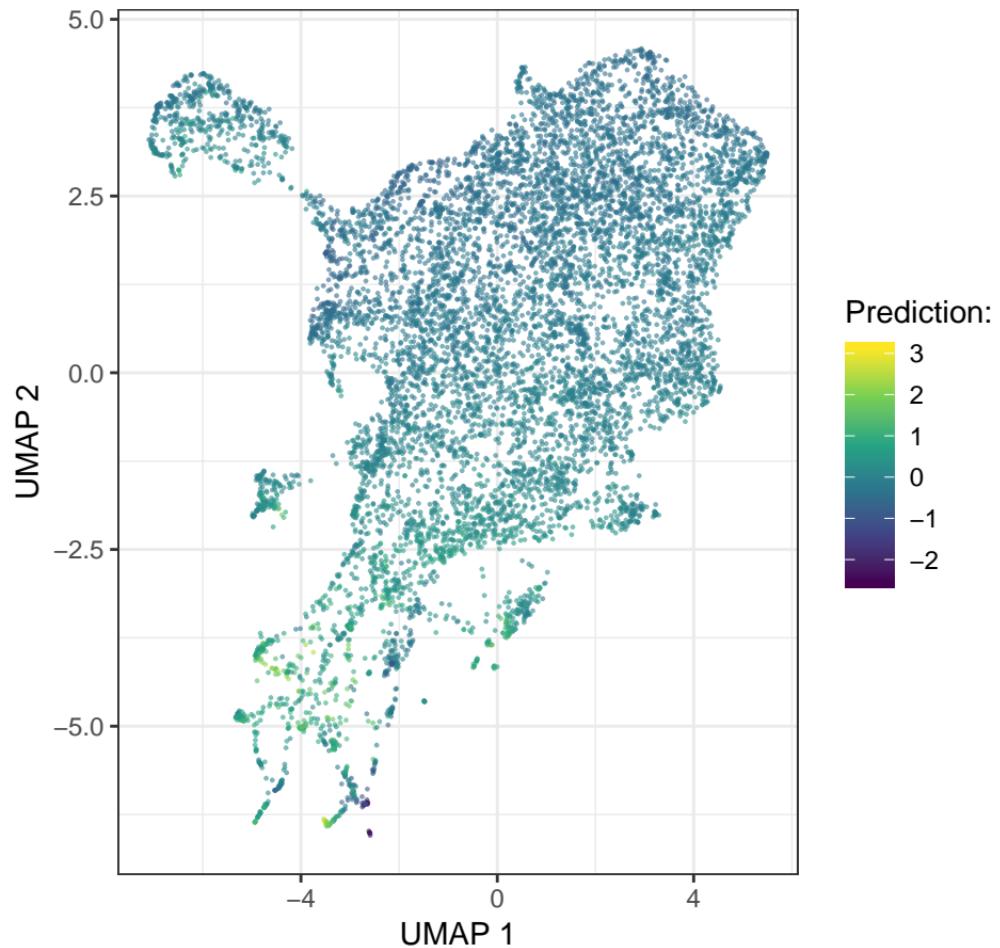
vb\_percent\_all\_early\_apoptosis



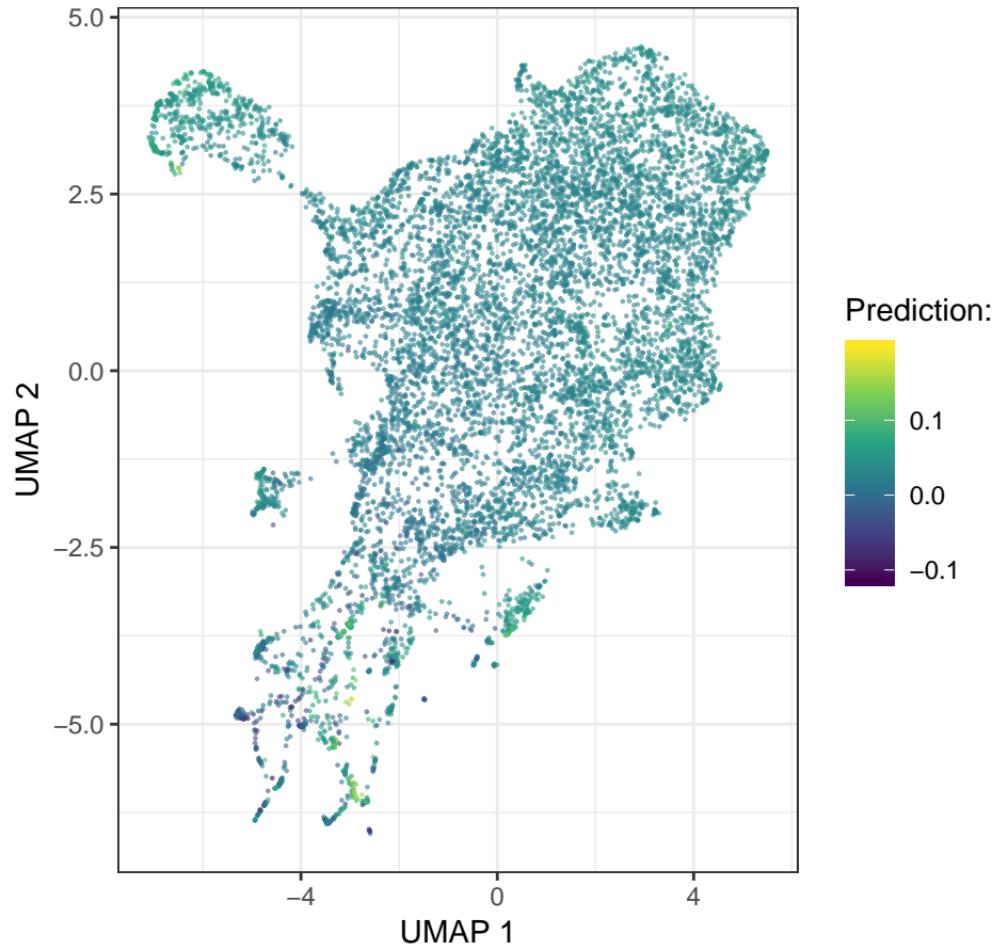
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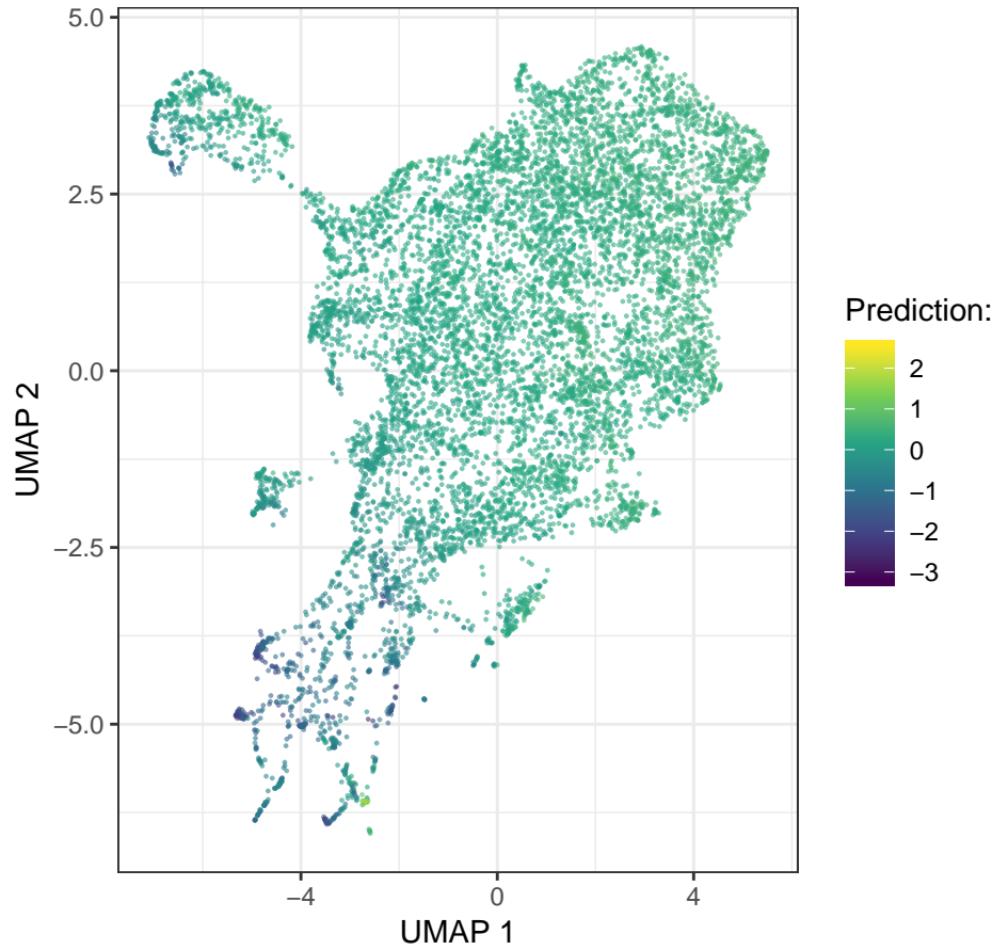
vb\_live\_cell\_area



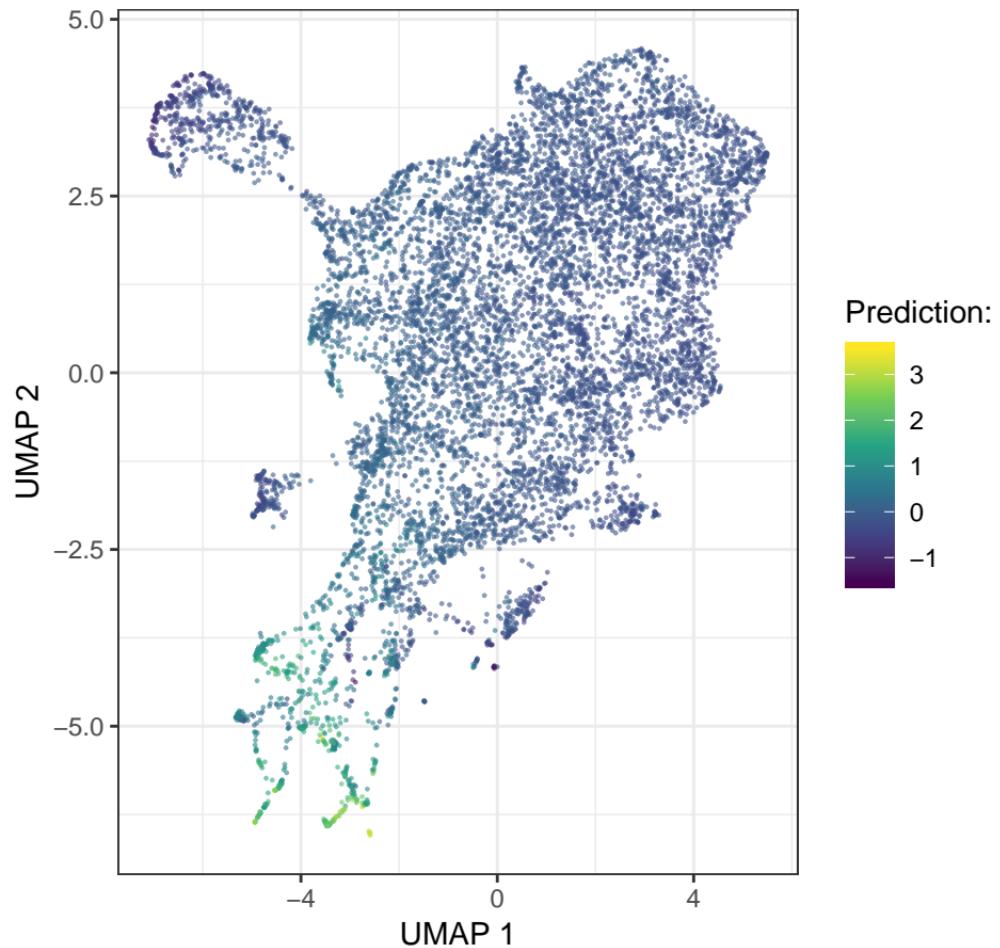
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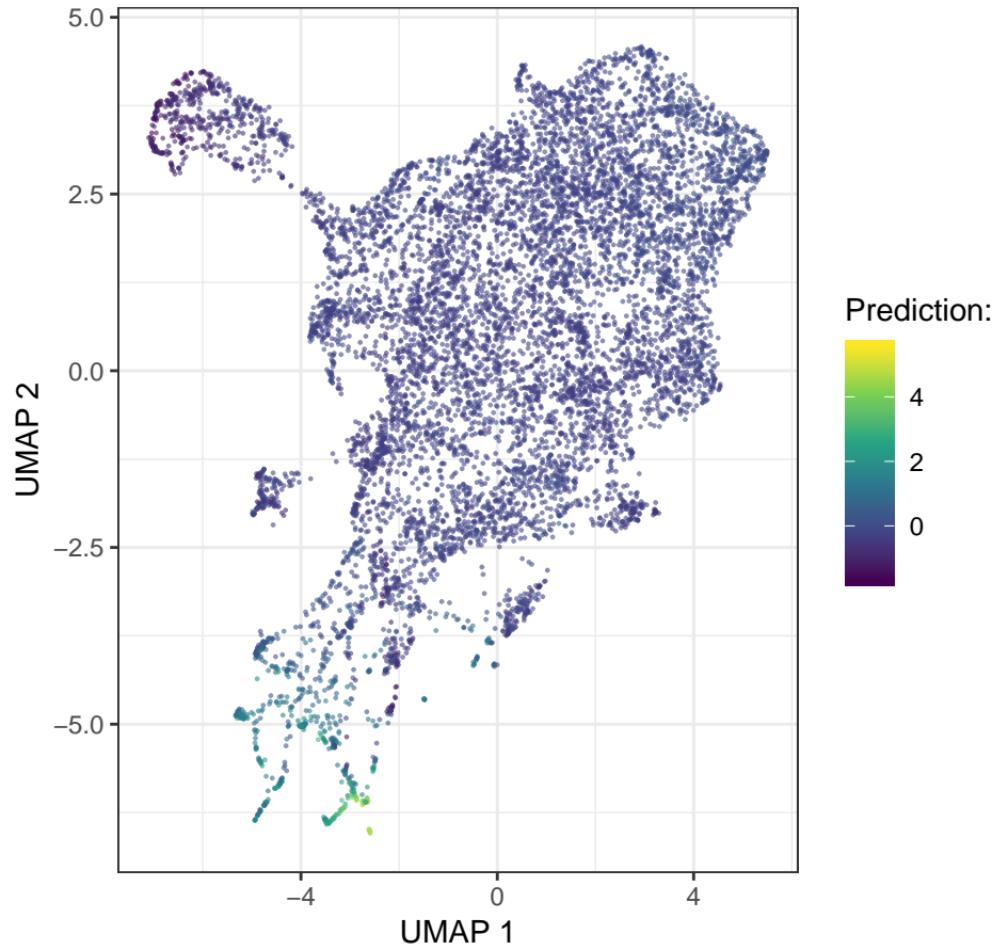
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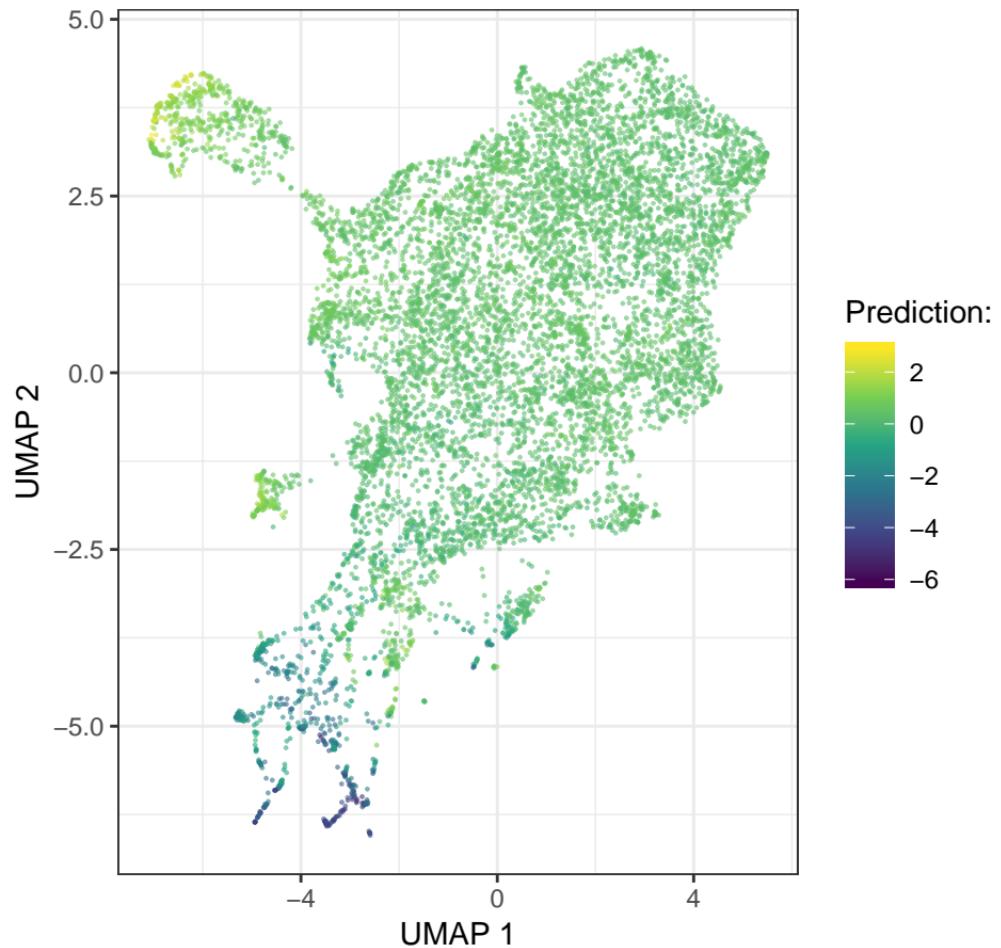
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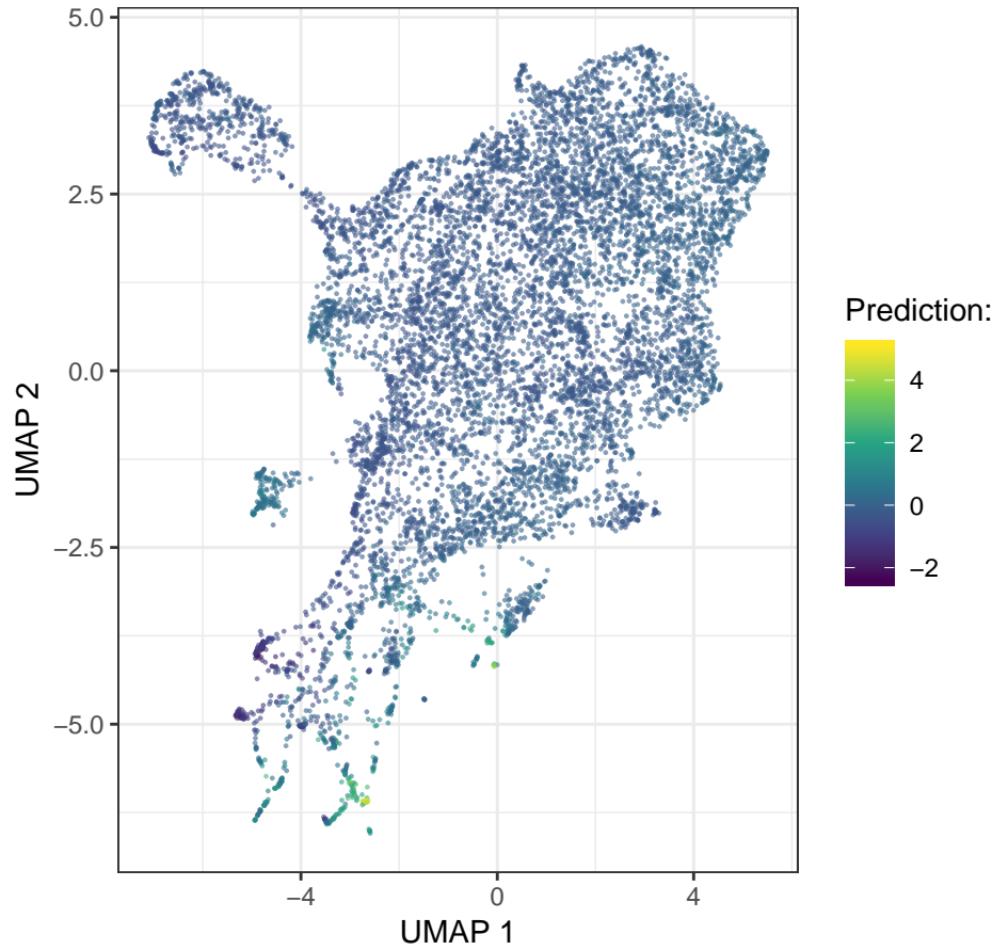
vb\_percent\_all\_late\_apoptosis



vb\_percent\_live



vb\_ros\_back\_mean



vb\_percent\_all\_apoptosis

