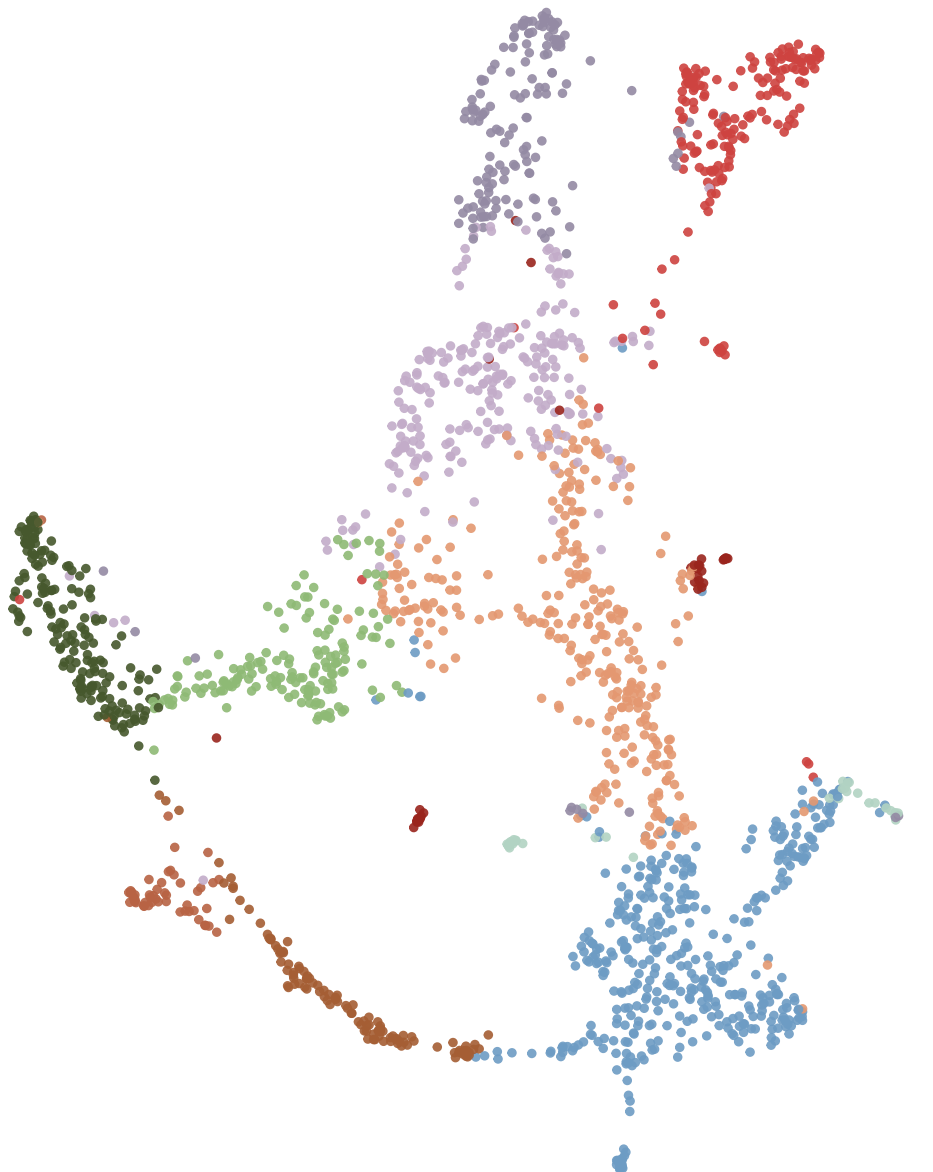


UMAP_2

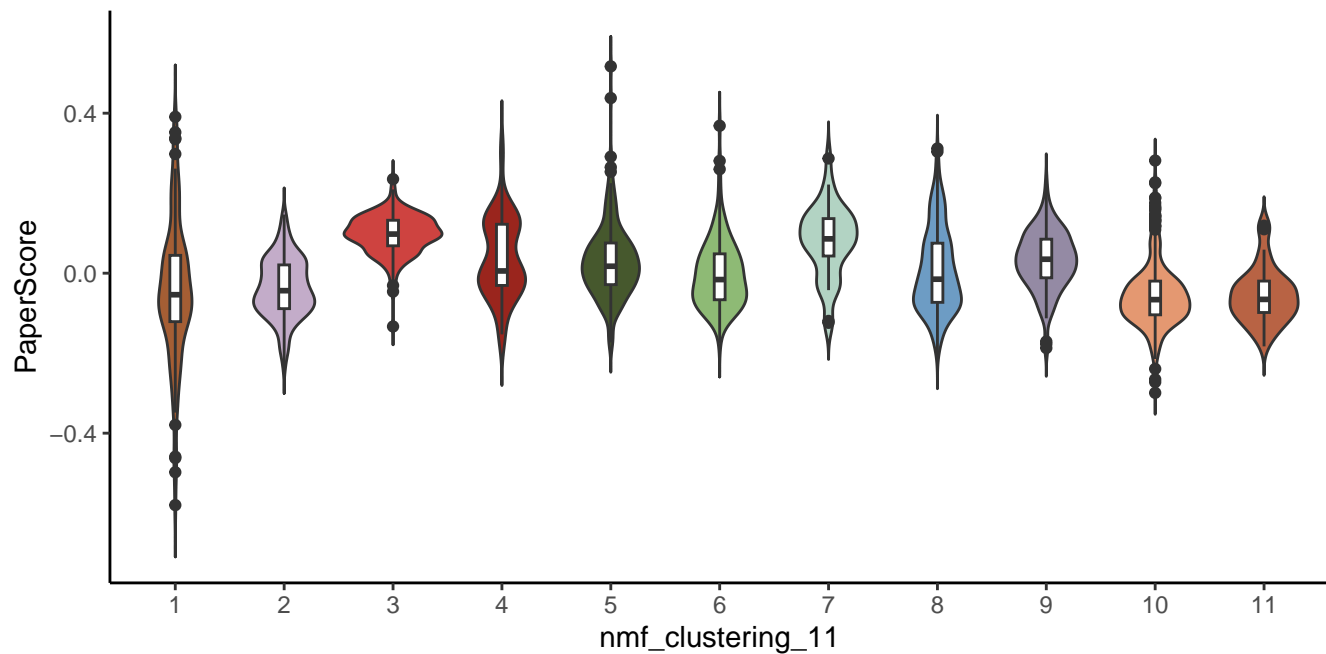
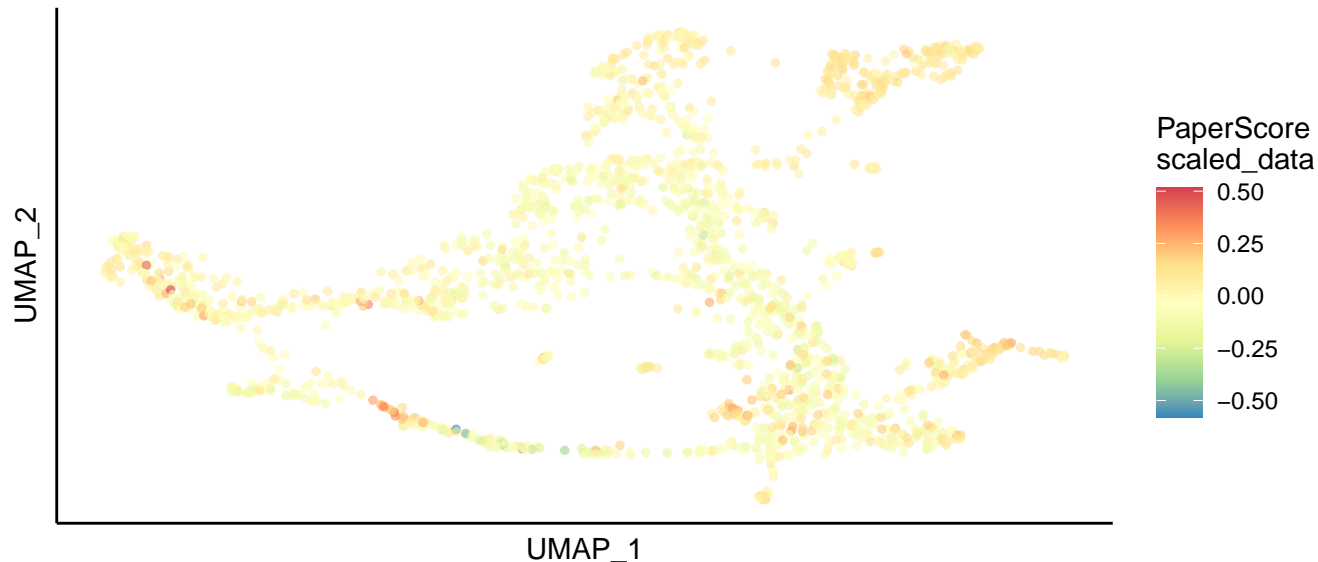
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nmf_clustering_11

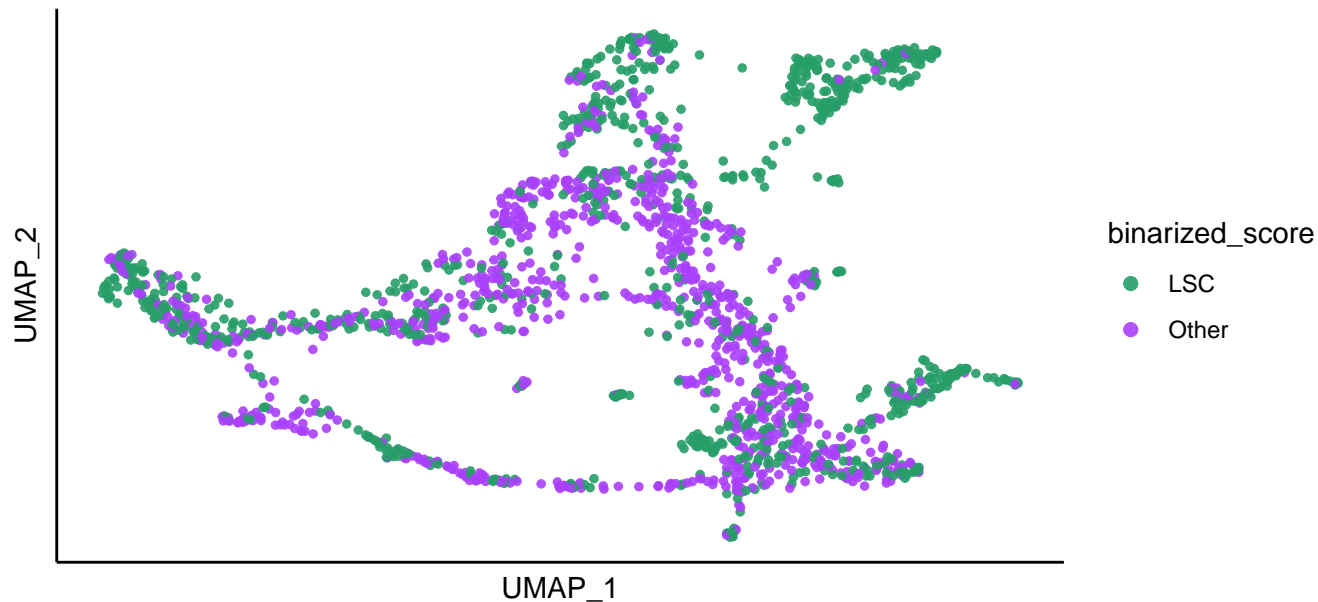
- 1
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- 3
- 4
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- 7
- 8
- 9
- 10
- 11



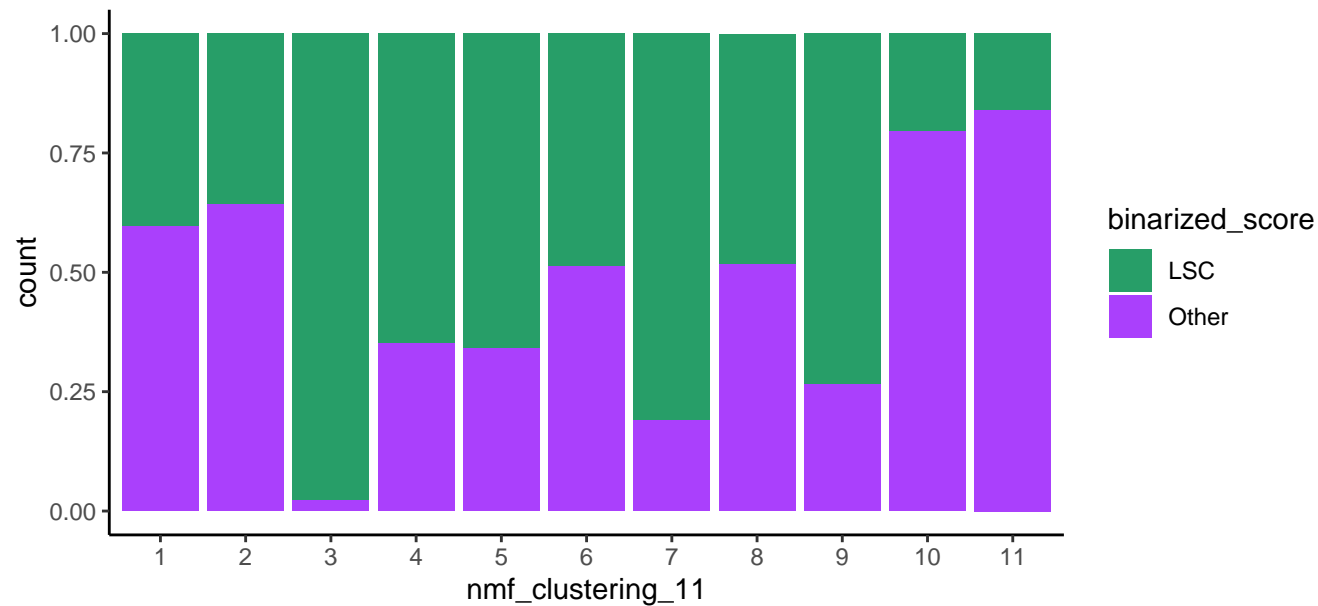
PaperScore
scaled_data



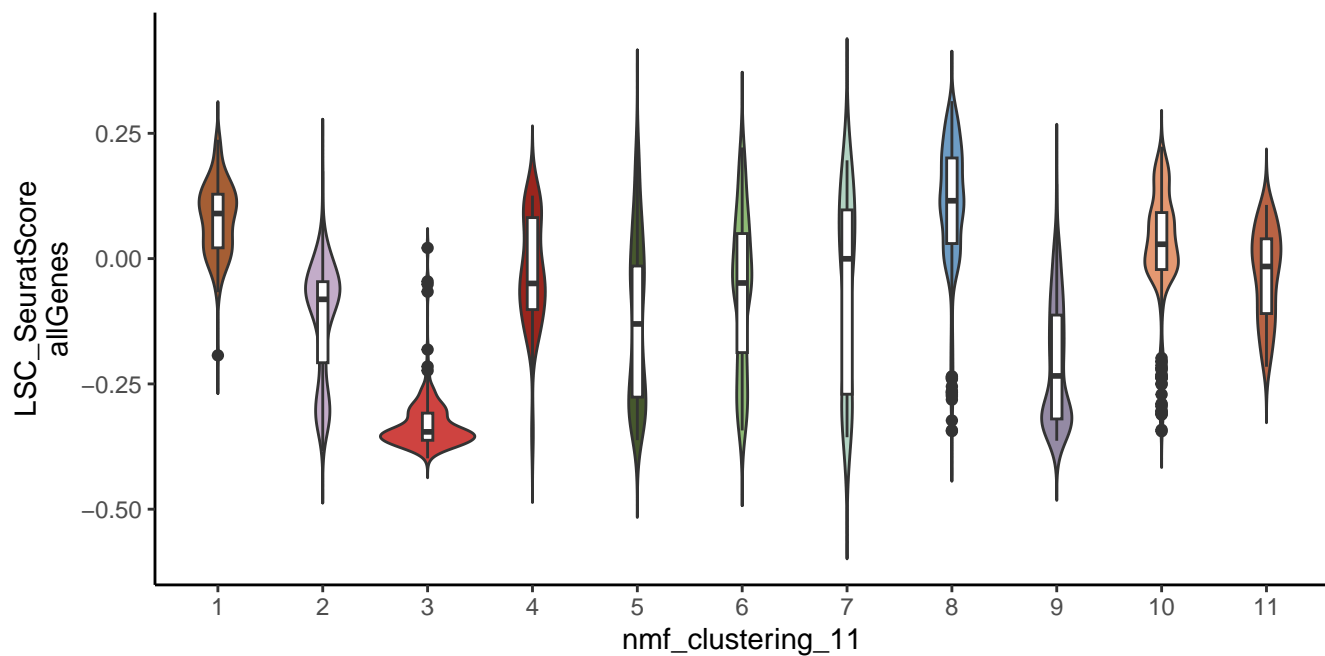
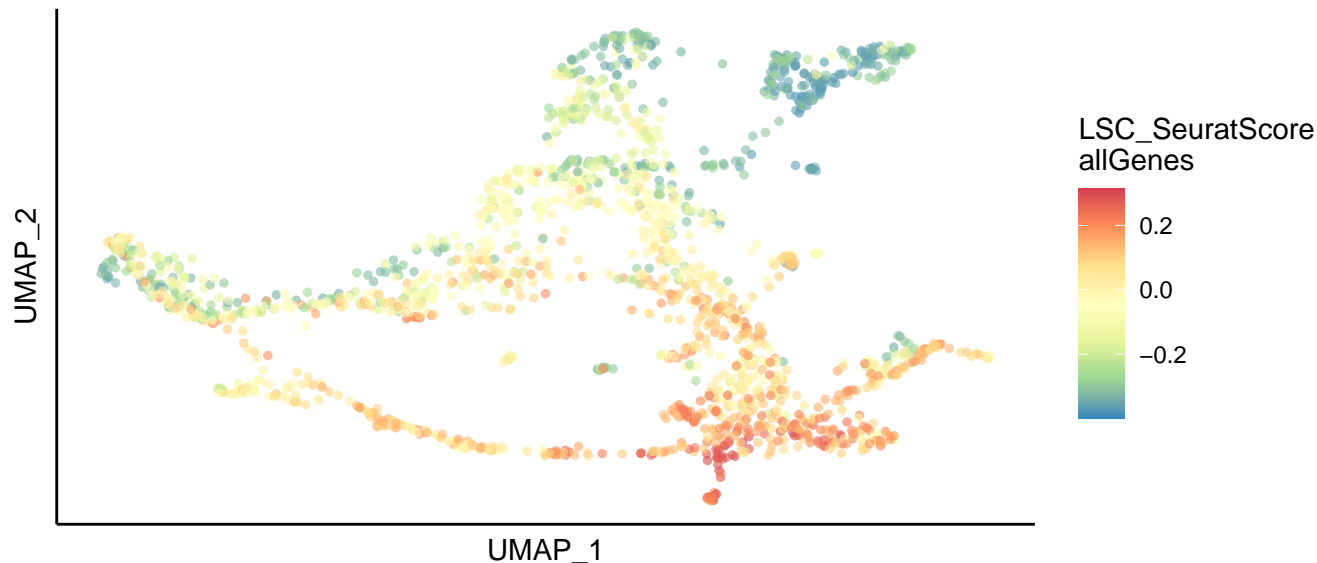
Median Binarized PaperScore



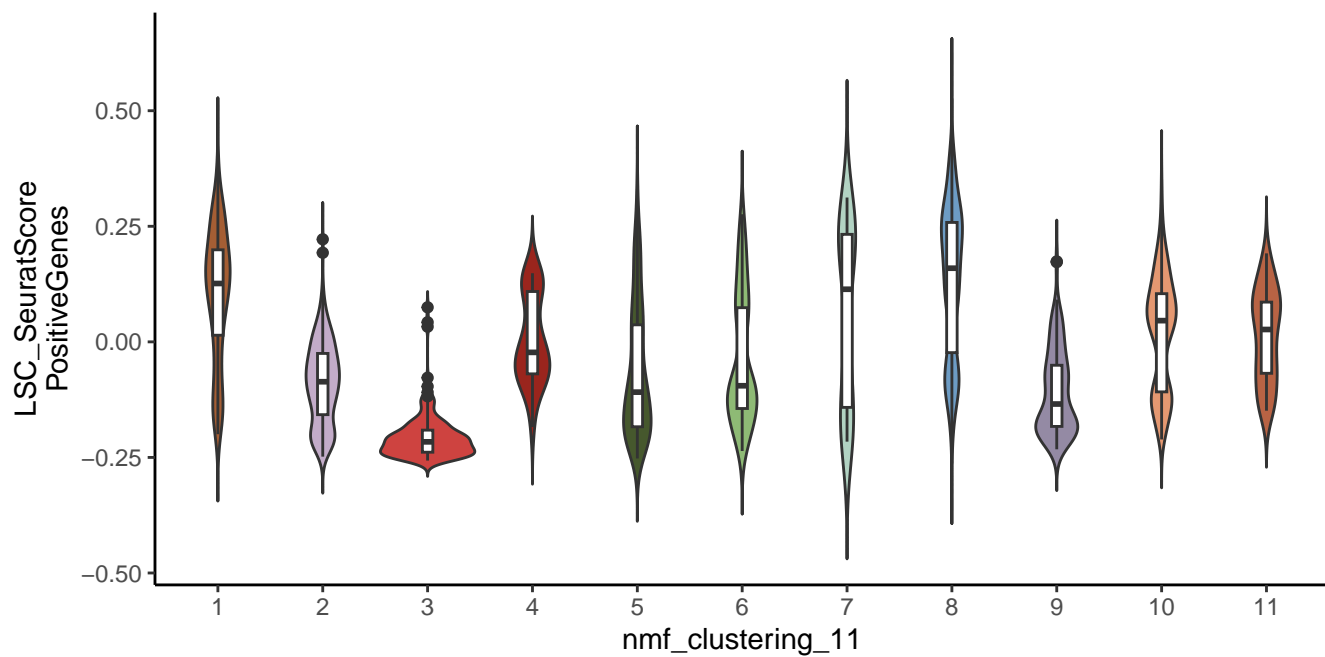
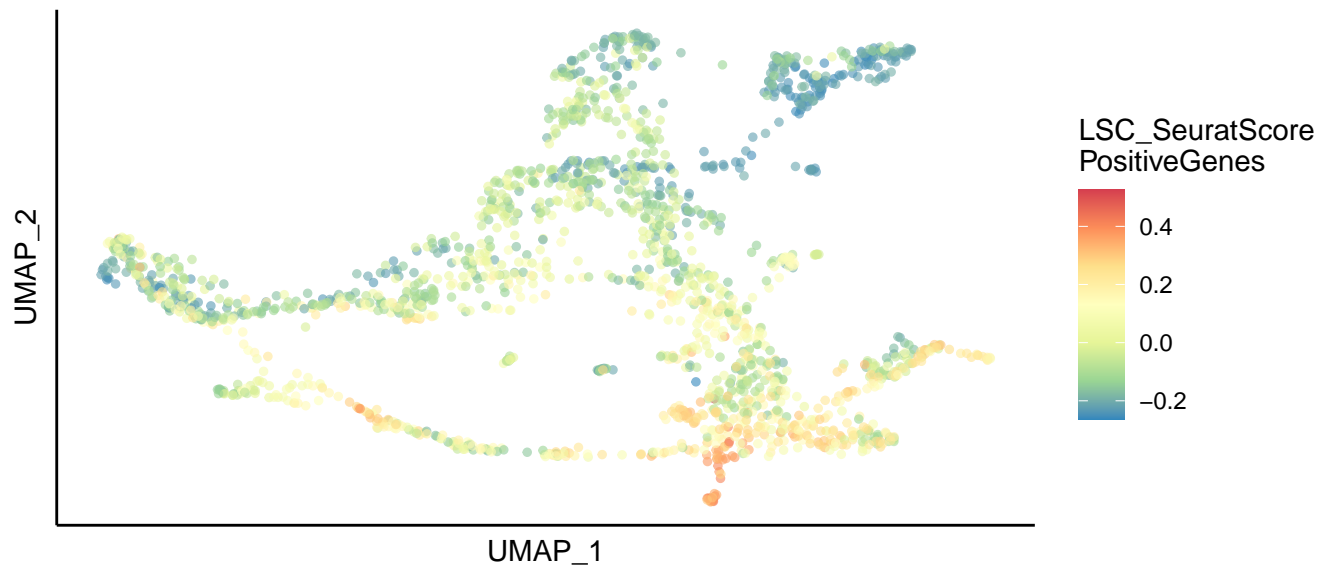
Median Binarized PaperScore Distribution



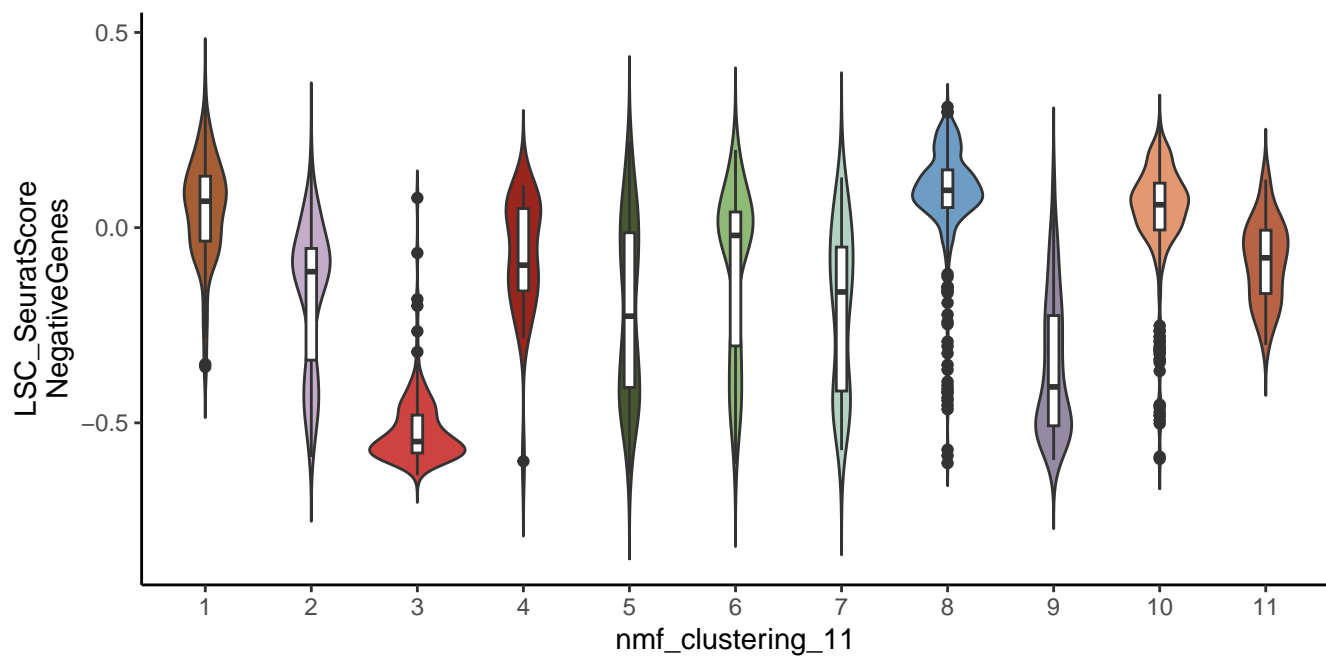
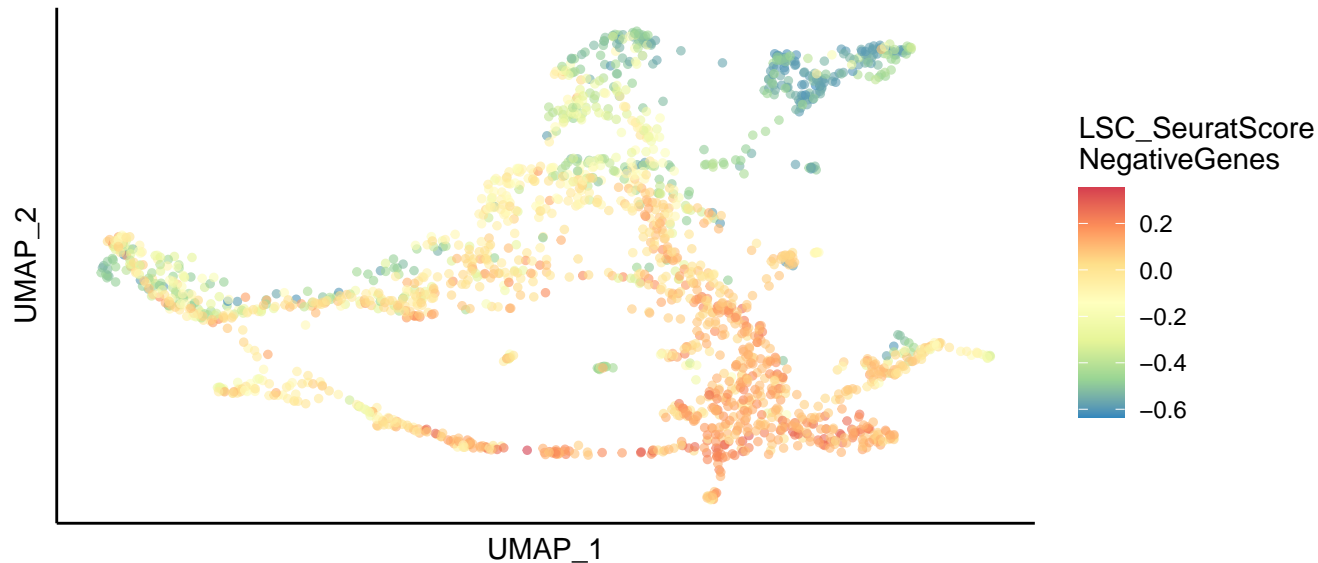
LSC_SeuratScore
allGenes



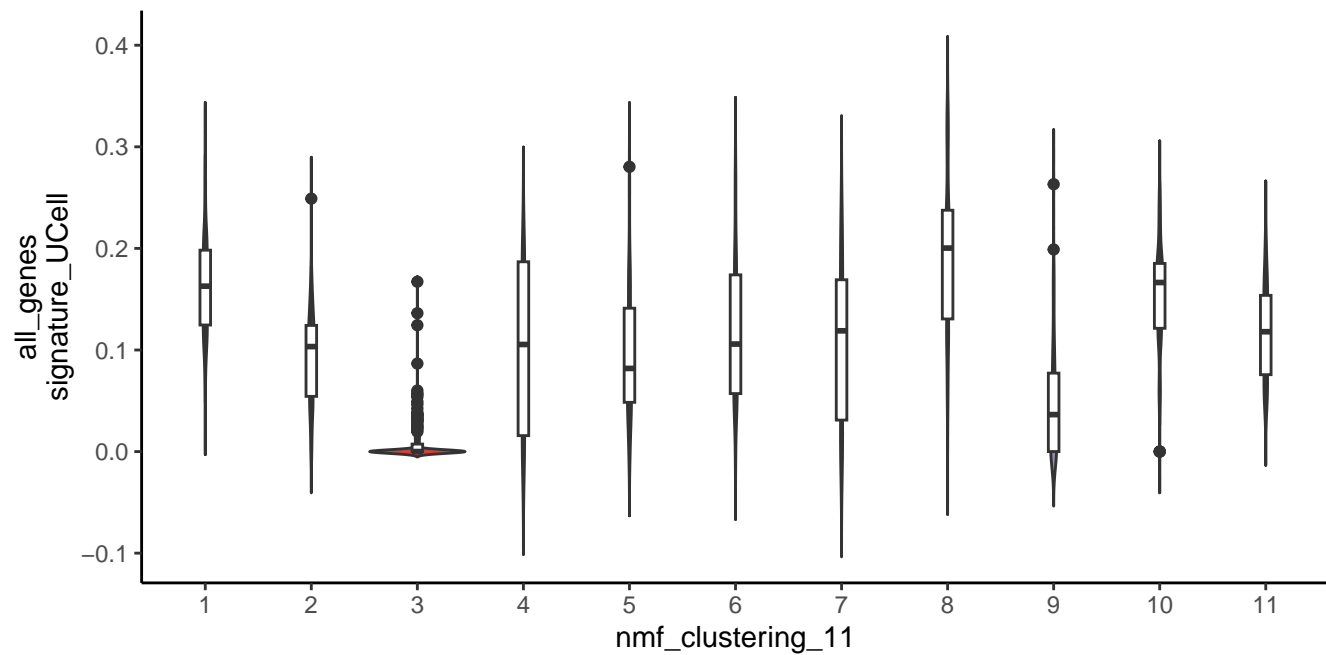
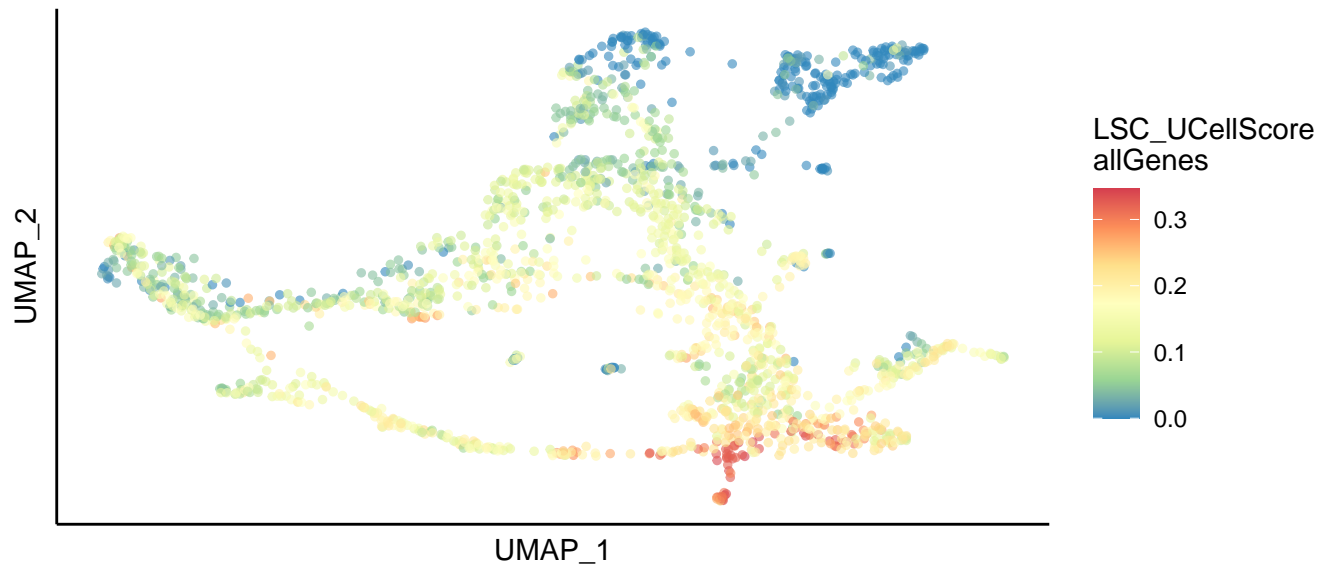
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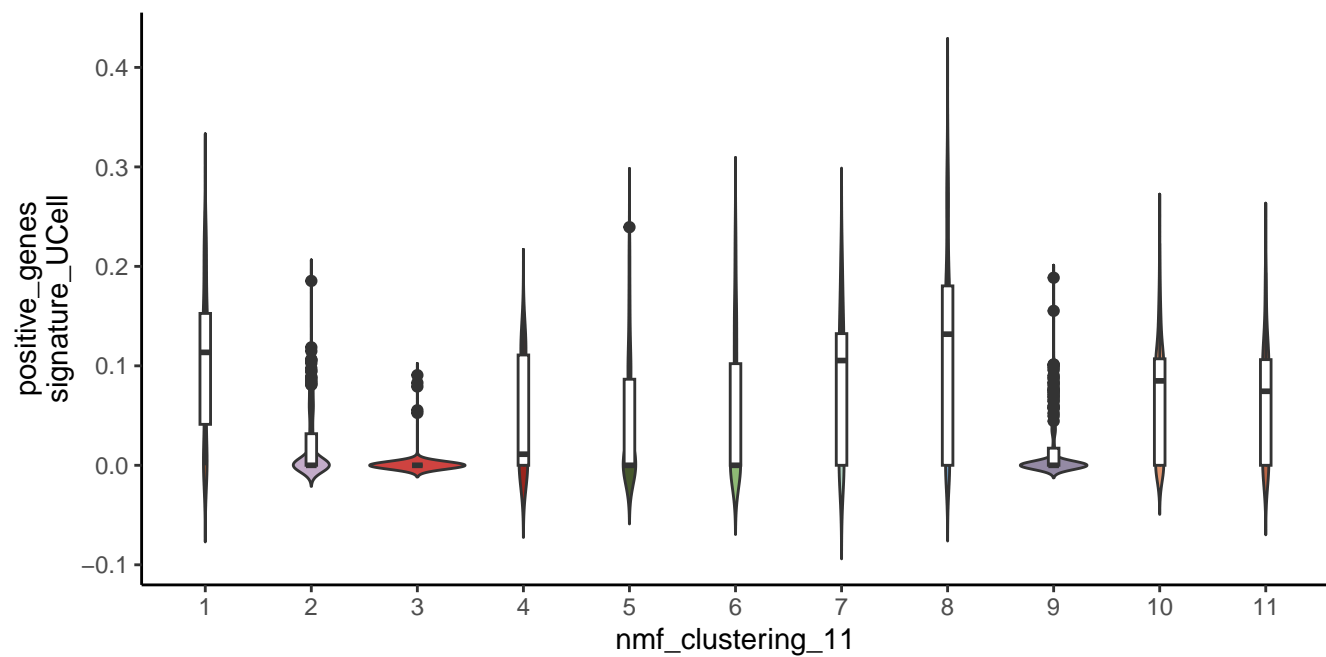
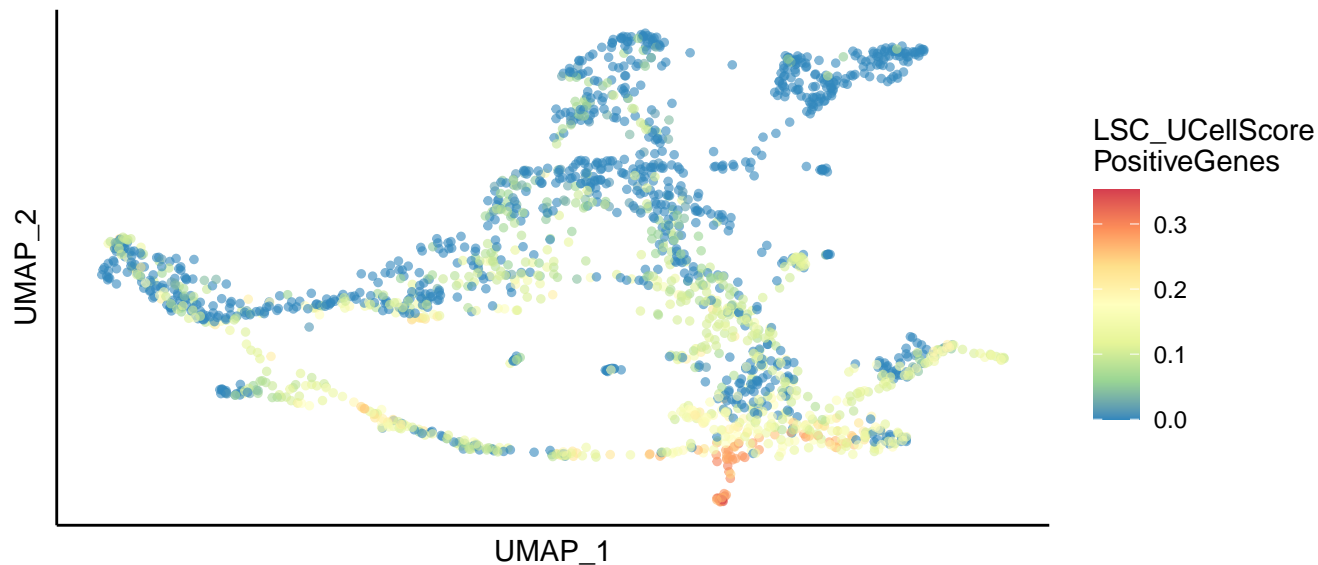
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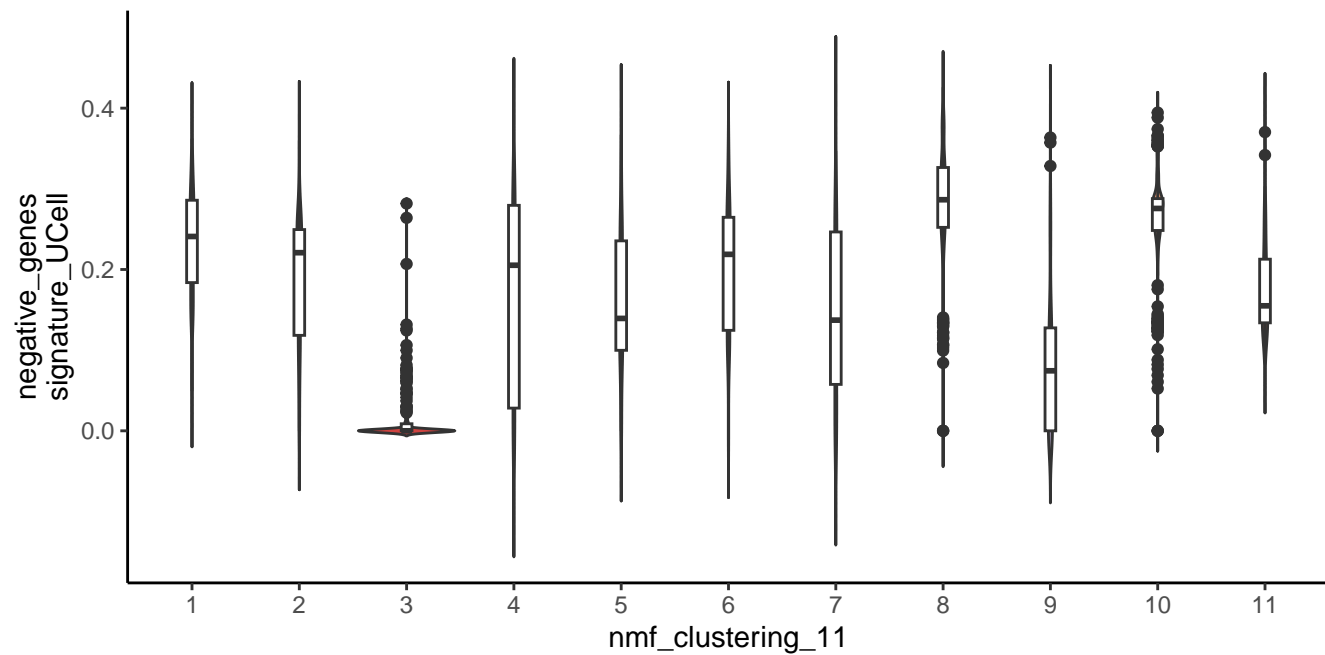
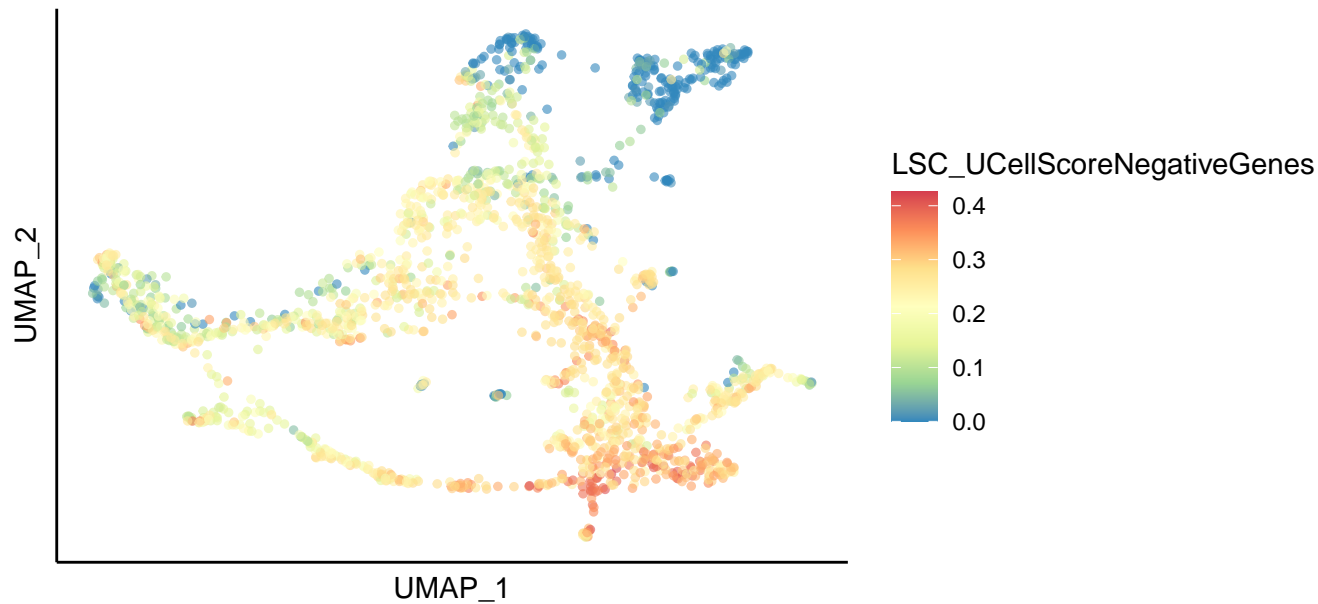
LSC_UCellScore
allGenes



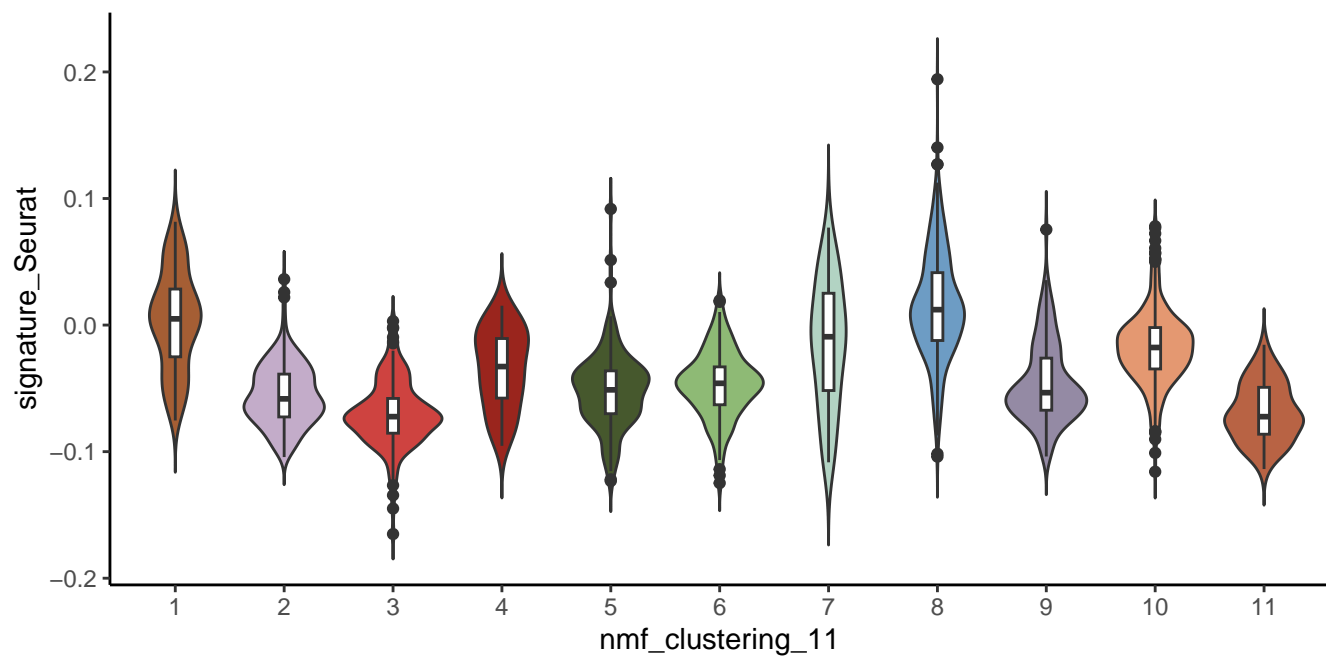
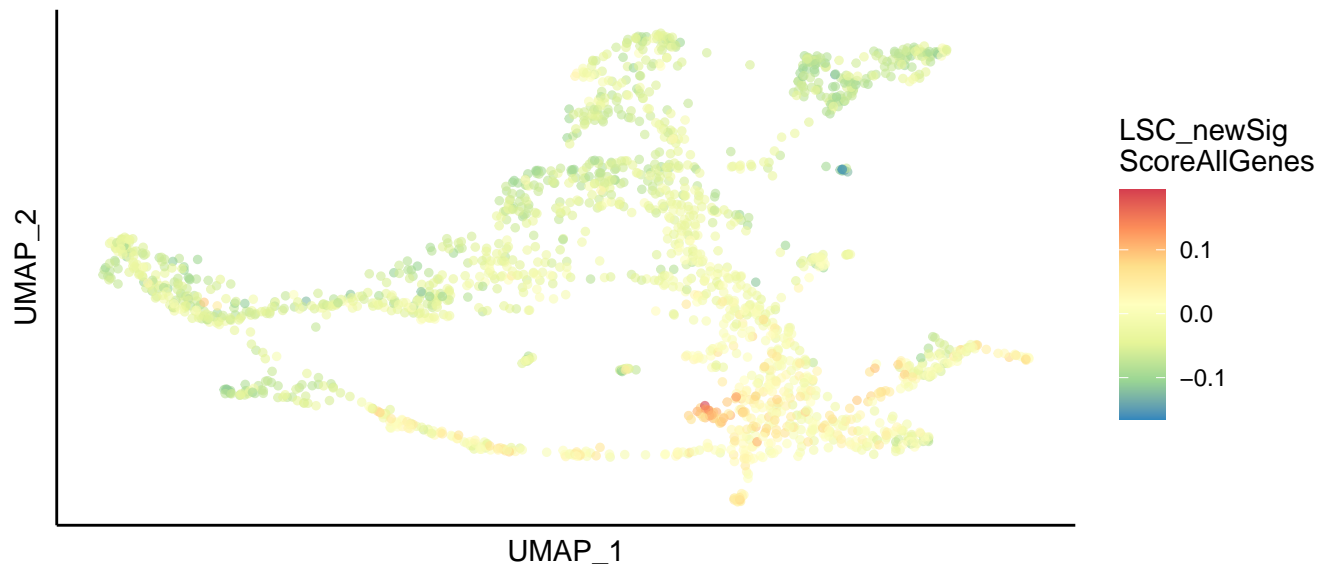
LSC_UCellScore
PositiveGenes



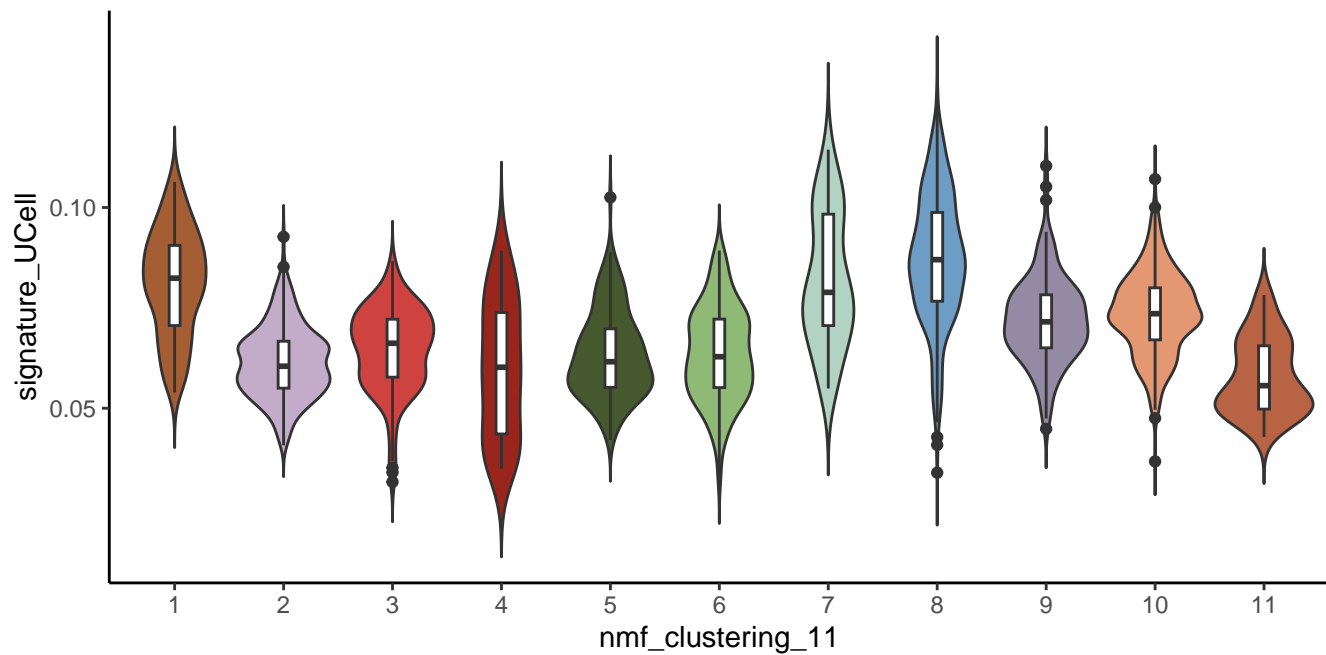
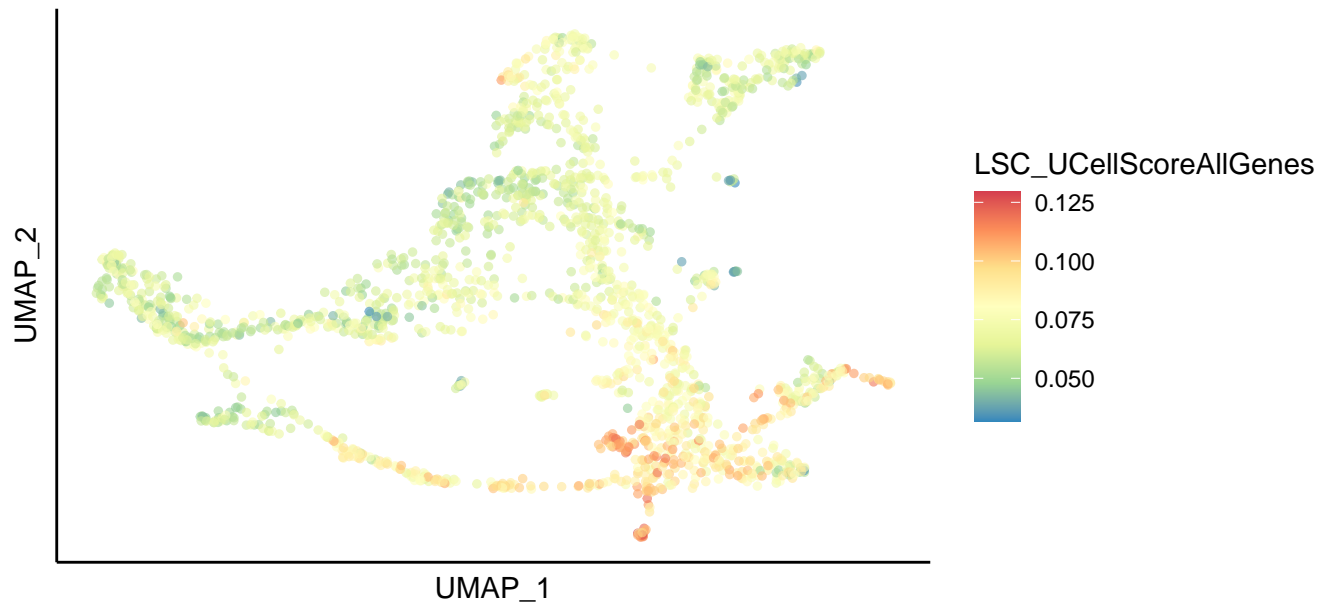
LSC_UCellScoreNegativeGenes



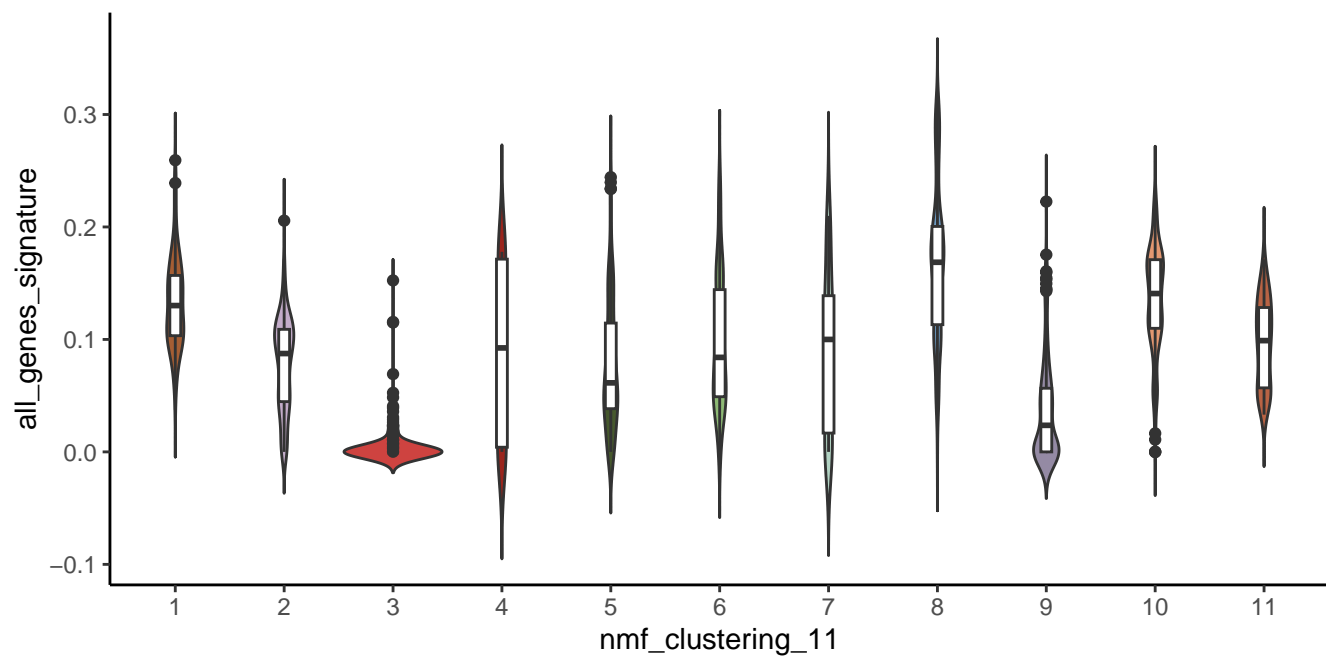
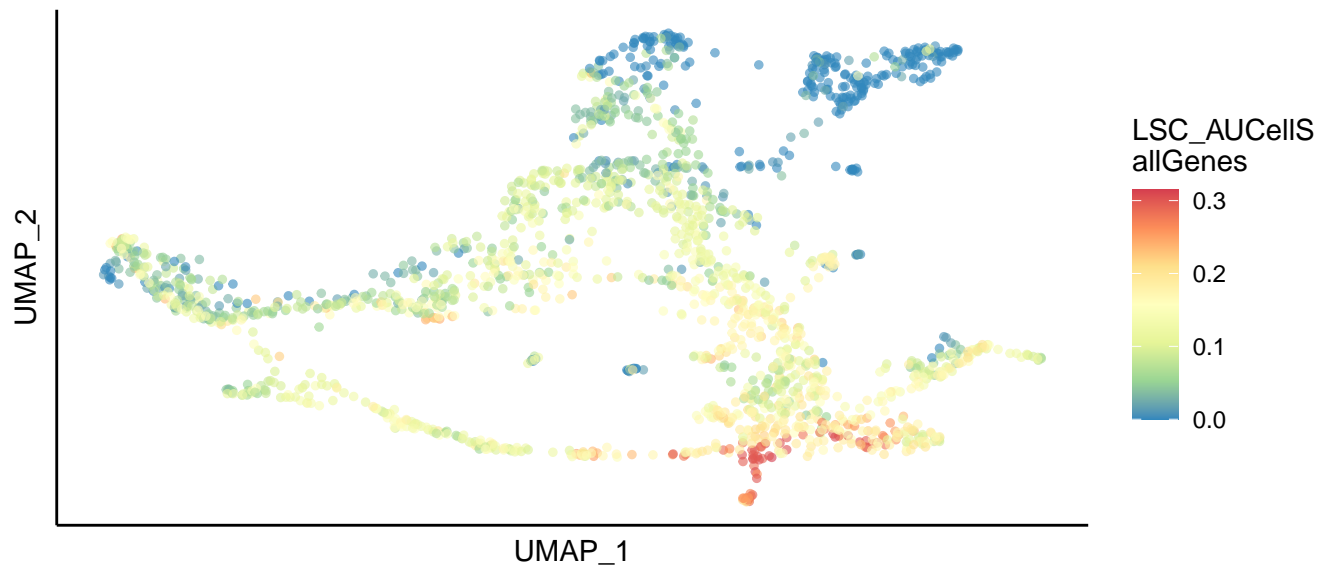
LSC_newSig
ScoreAllGenes



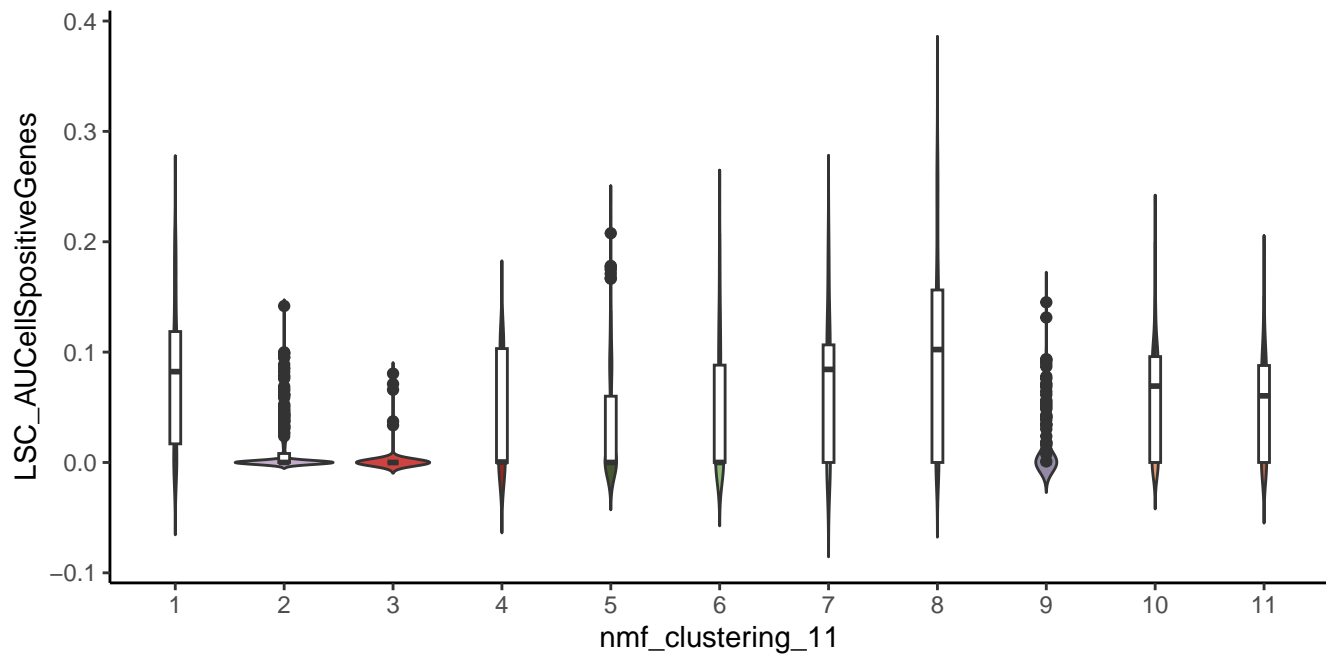
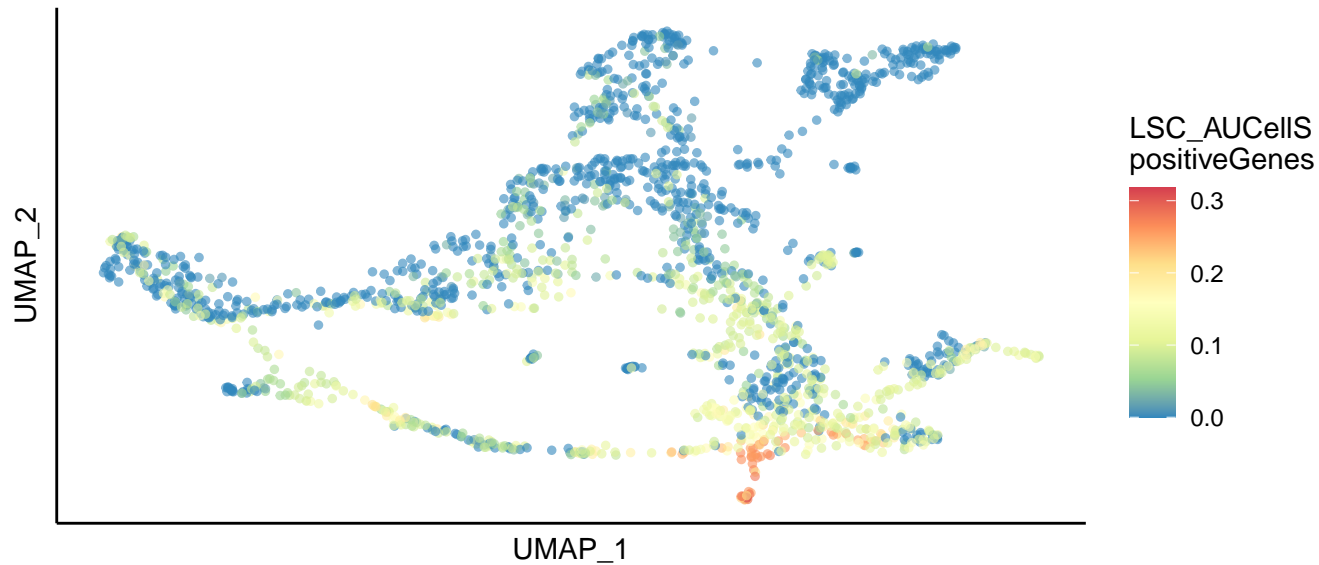
LSC_UCellScoreAllGenes



LSC_AUCeIS
allGenes



LSC_AUCelIS
positiveGenes



LSC_AUCellS
negativeGenes

