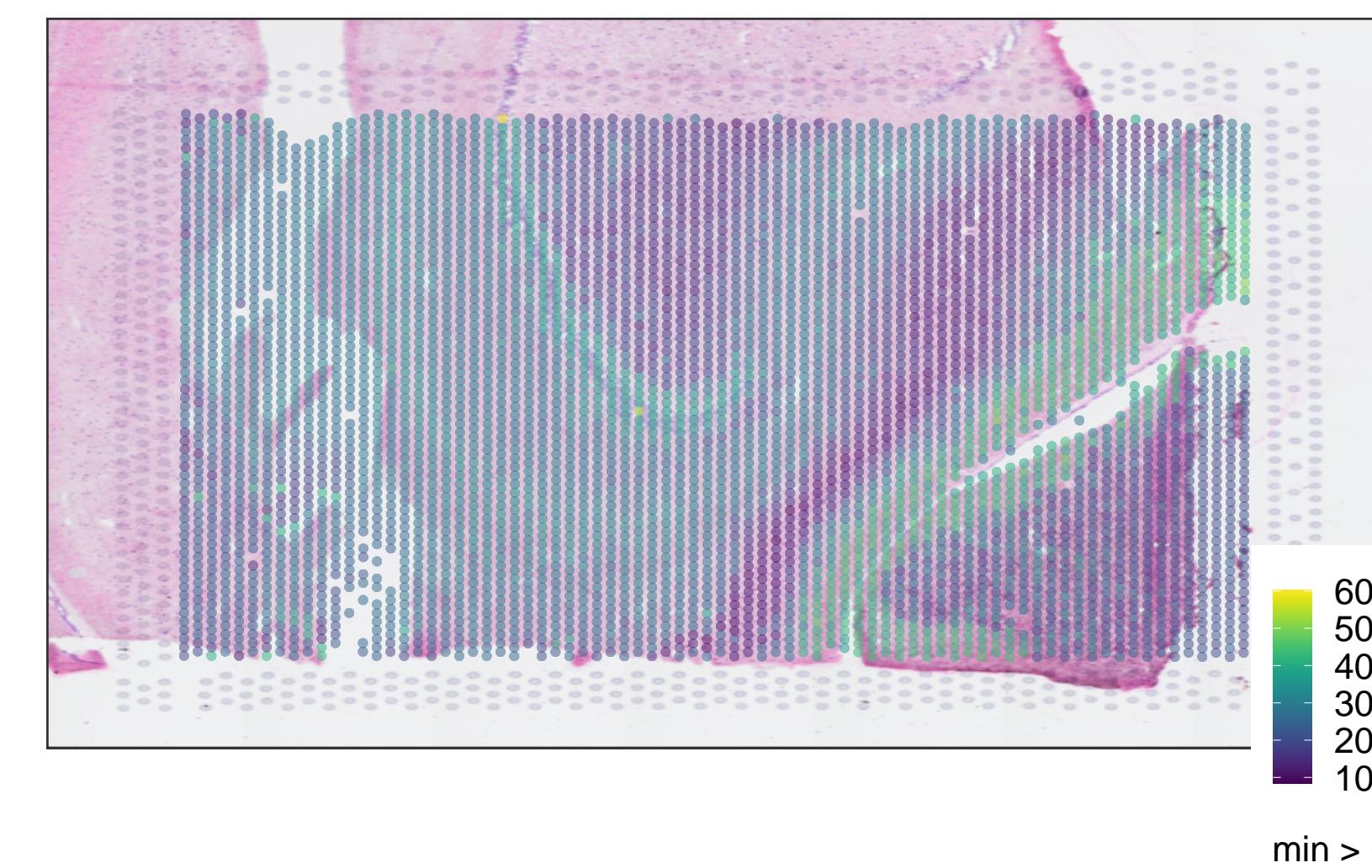
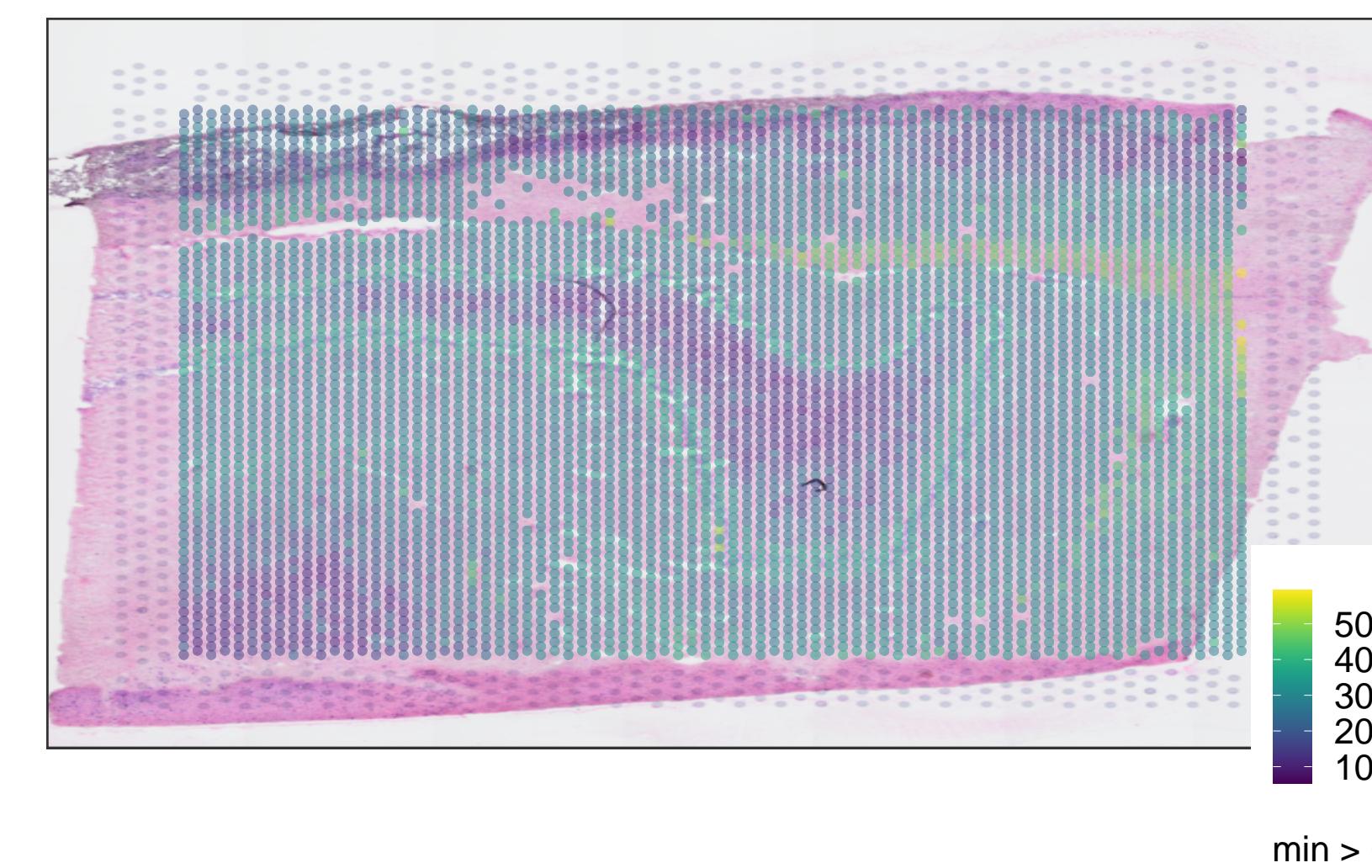


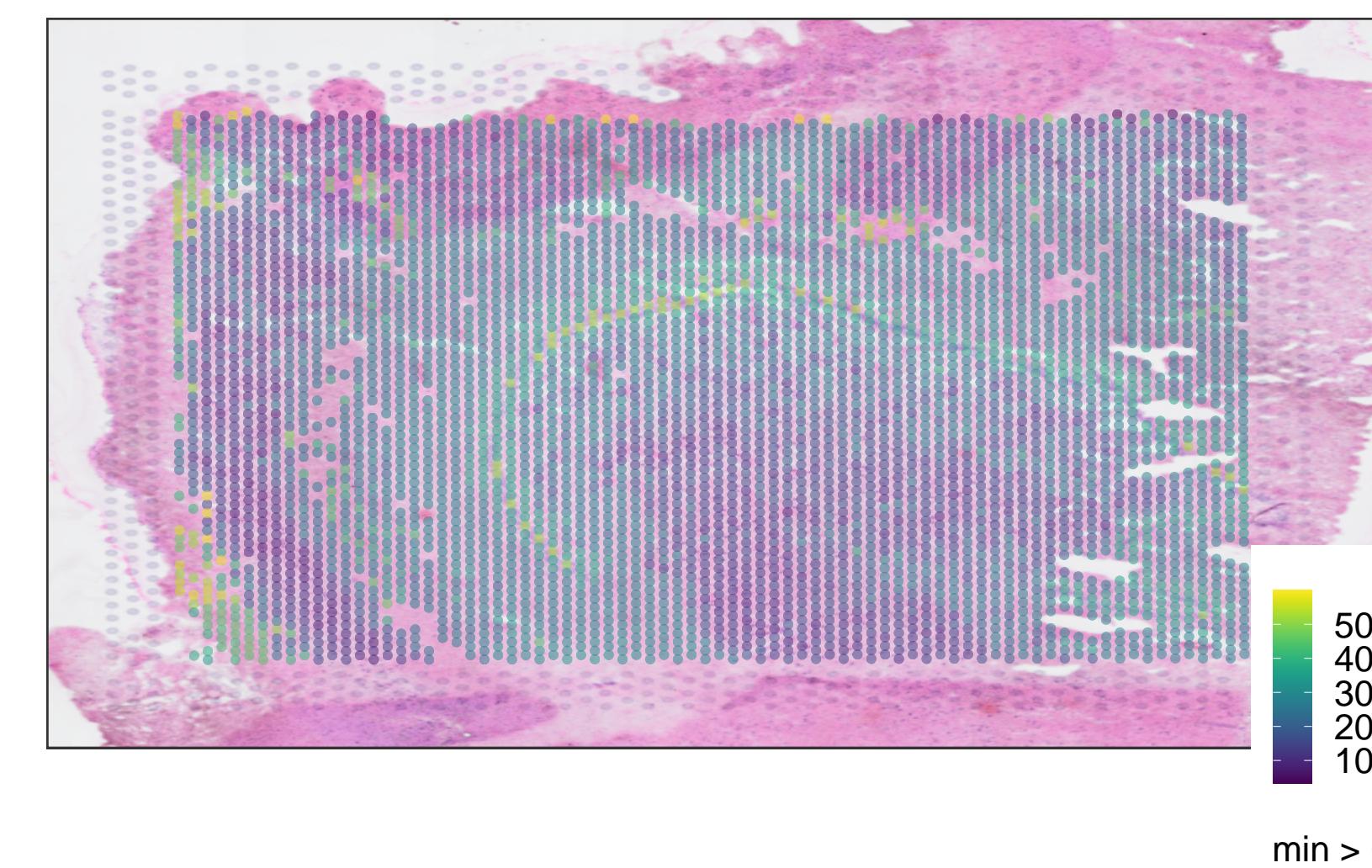
# Br6129\_new pseudotime TRUE



# Br6299\_new pseudotime TRUE

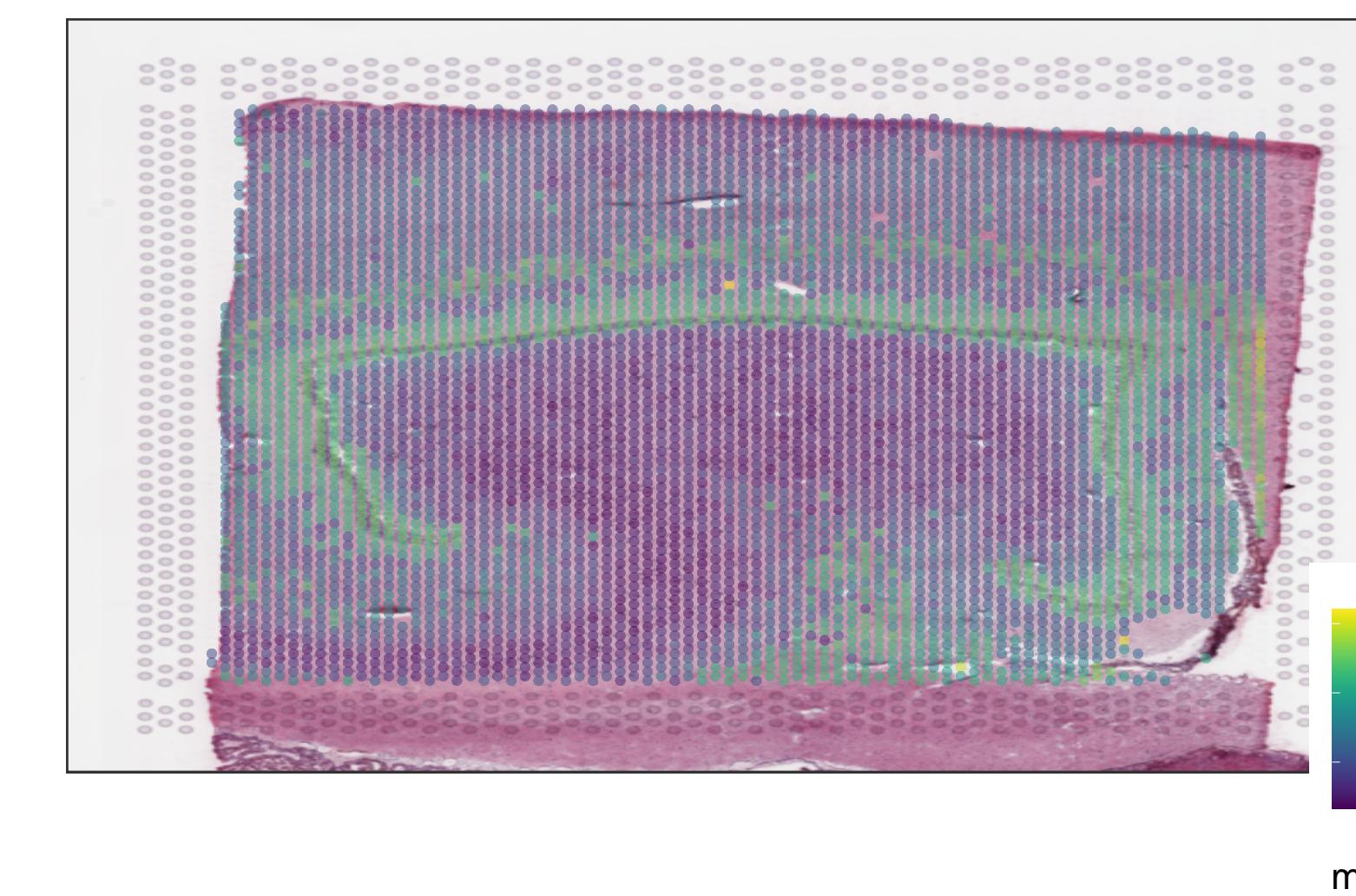


Br2720 pseudotime TRUE

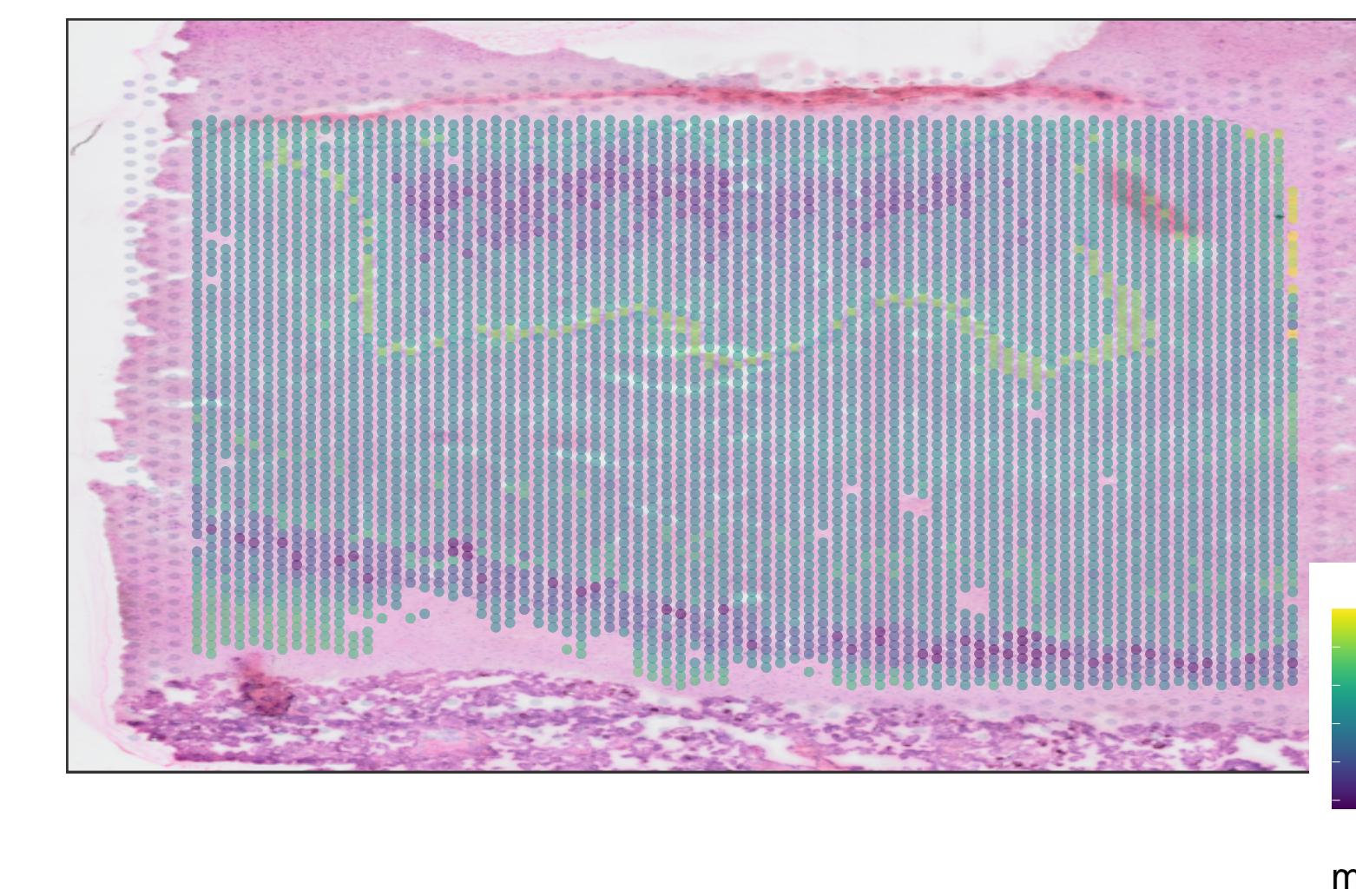


# Br8195 pseudotime TRUE

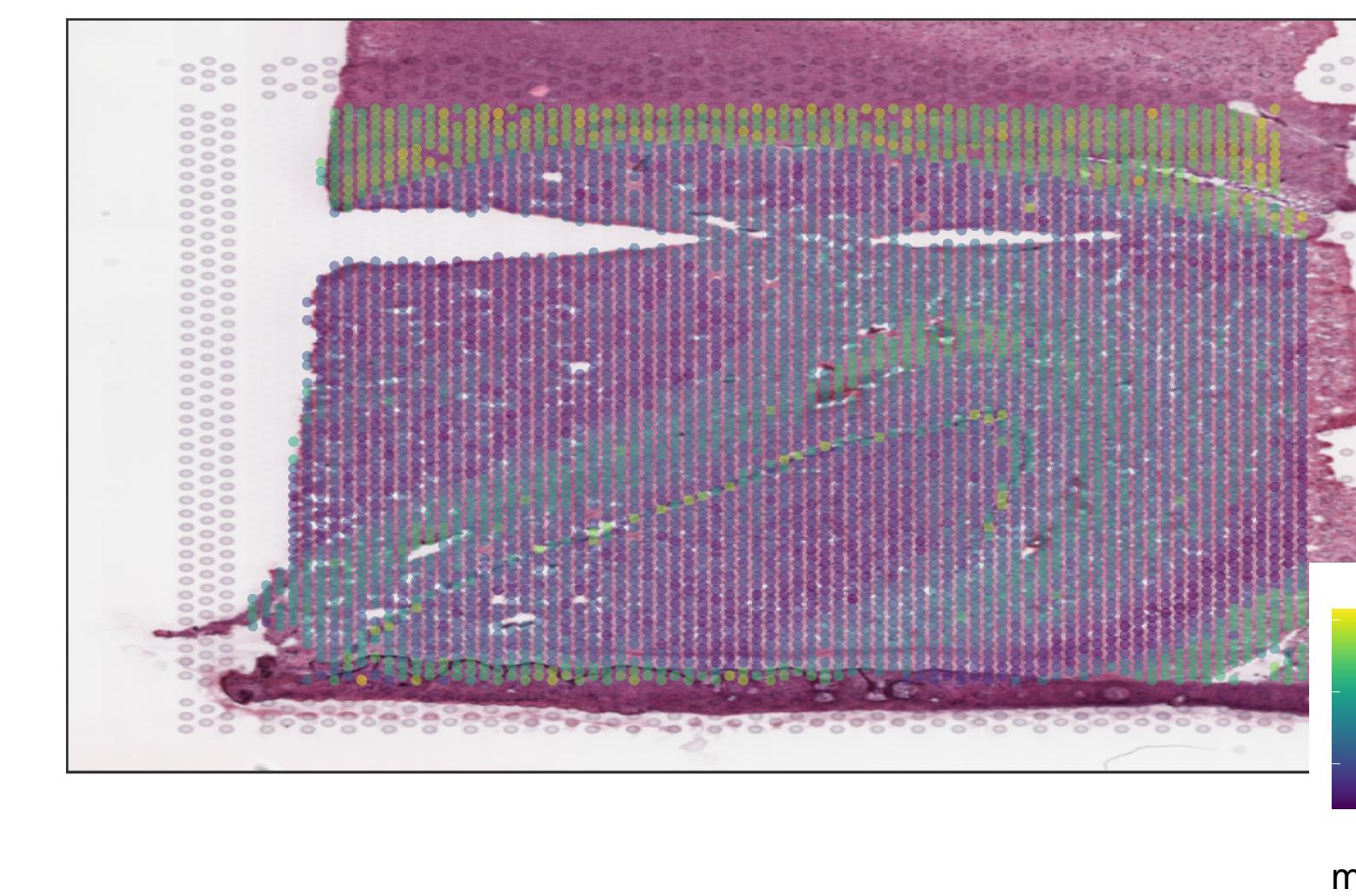
# Br1412 pseudotime TRUE



# Br6522 pseudotime TRUE



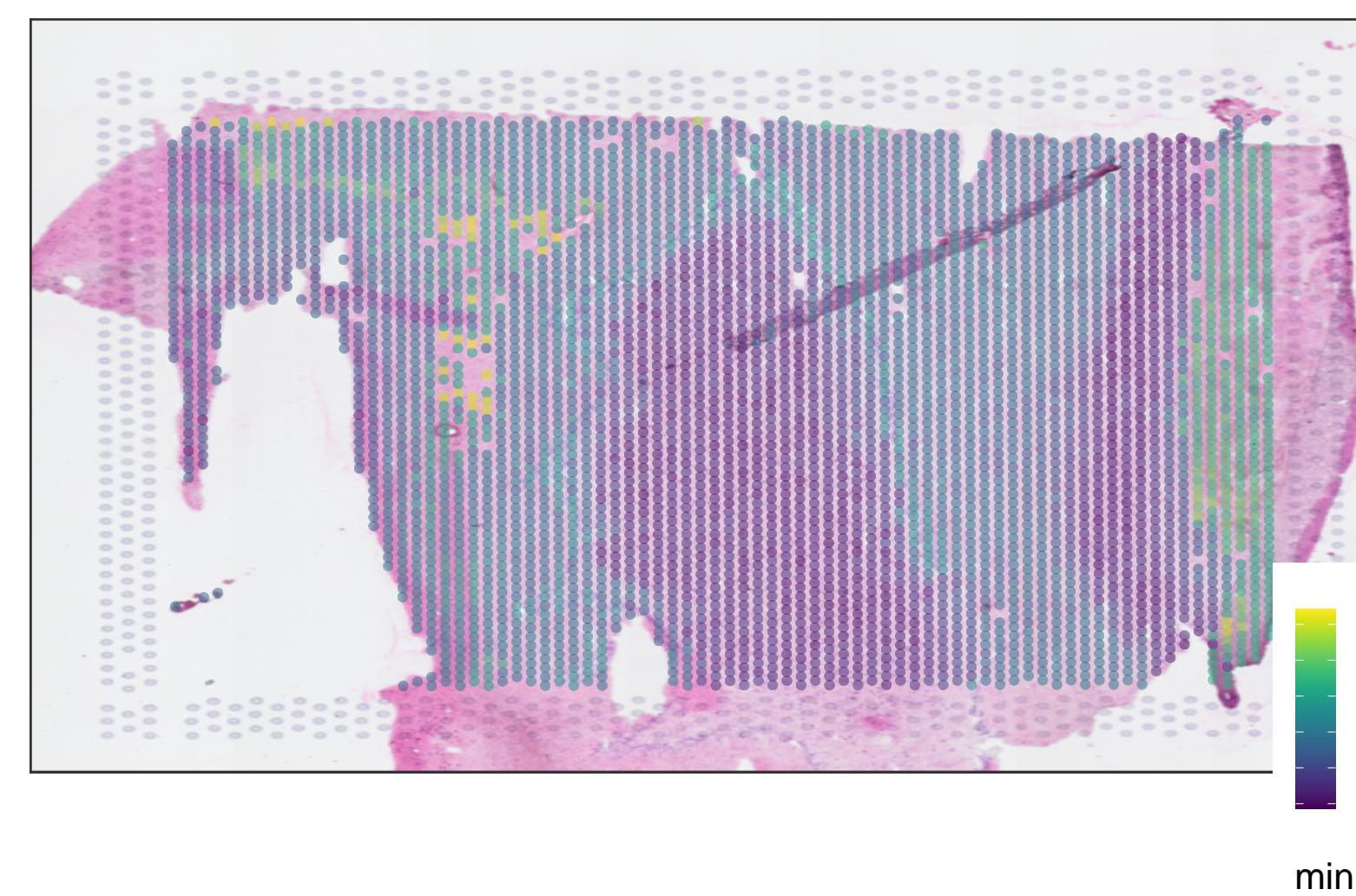
# Br5242 pseudotime TRUE



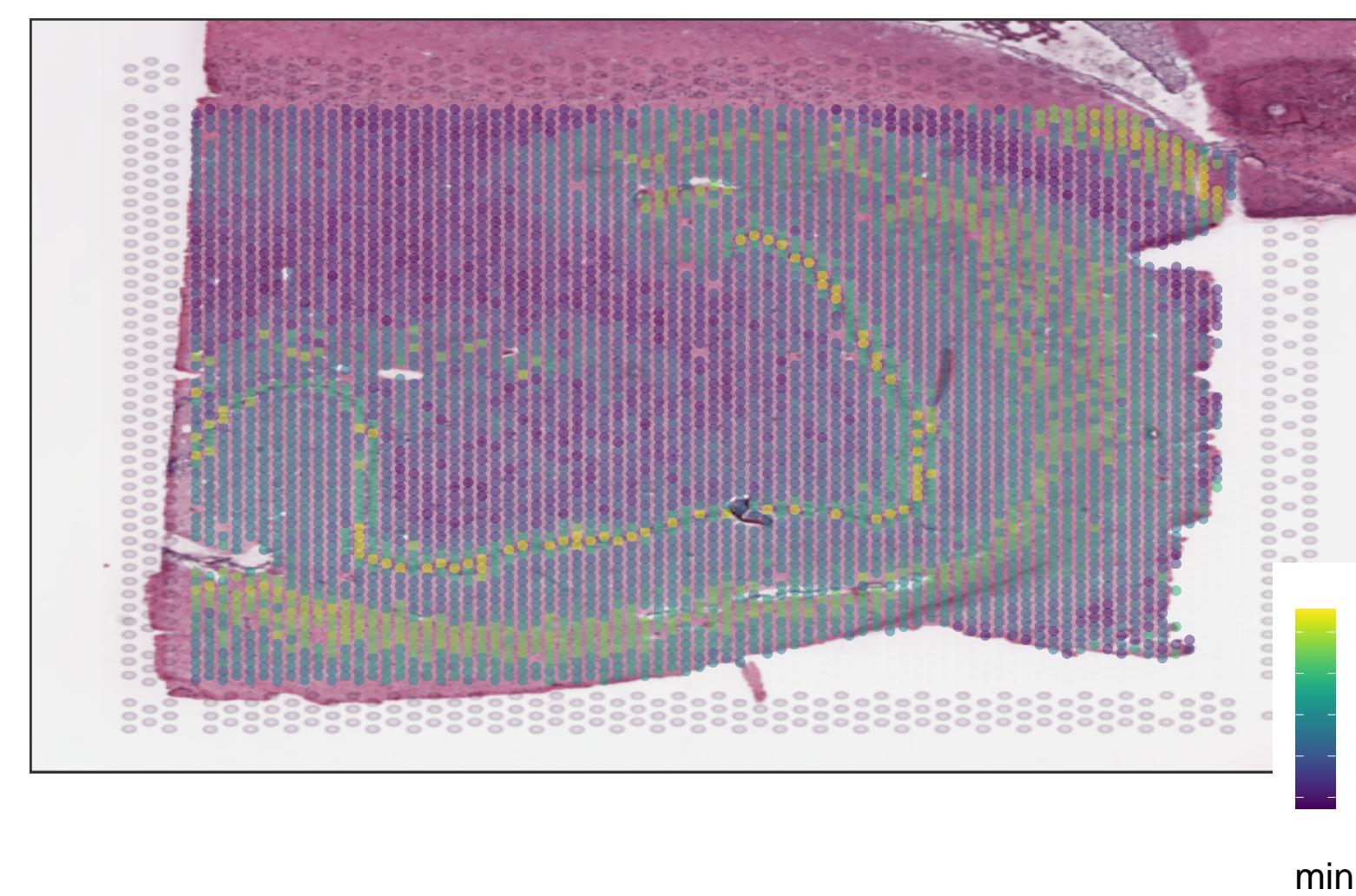
# Br8533 pseudotime TRUE

This figure displays a spatial transcriptomics analysis of a tissue sample, likely a breast cancer specimen (Br8533). The background image shows a histological section with pink staining. Superimposed on this is a grid of colored dots representing gene expression levels over pseudotime. The color scale ranges from dark purple (low expression) to yellow (high expression). A color bar on the right is labeled "min". The expression pattern shows distinct clusters and gradients, indicating spatial gene regulation.

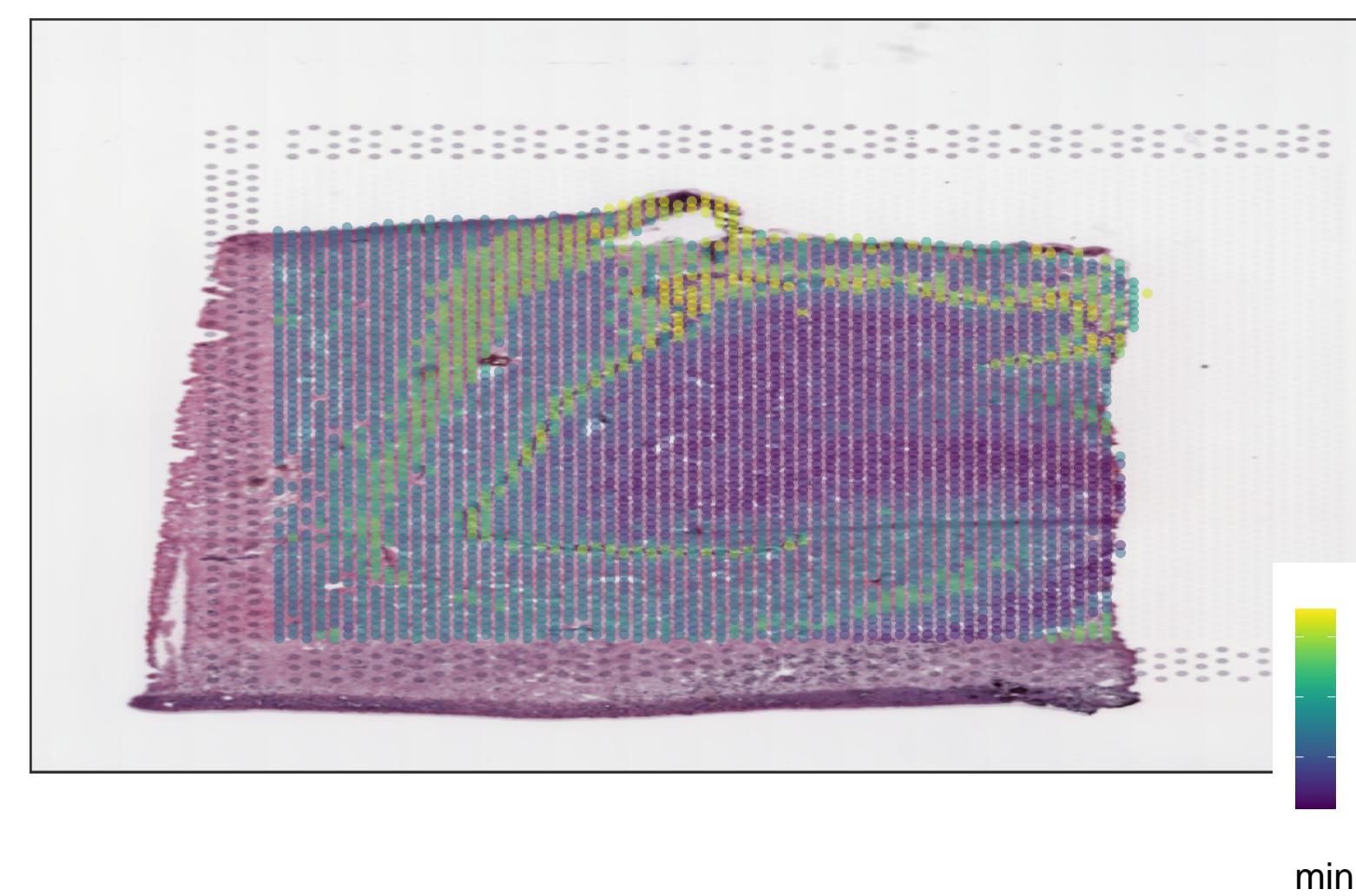
# Br8181 pseudotime TRUE



# Br8667 pseudotime TRUE

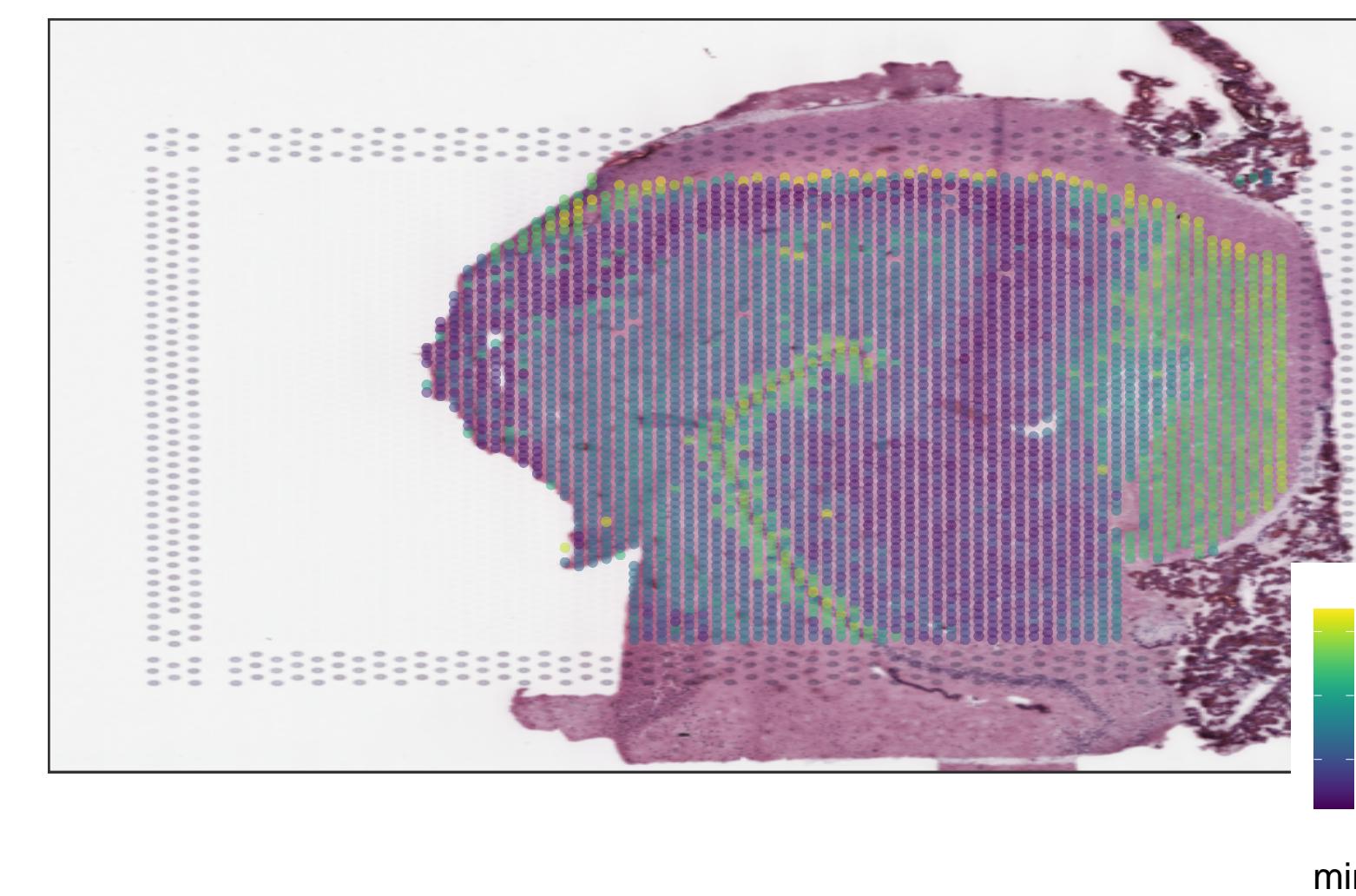


# Br6023 pseudotime TRUE

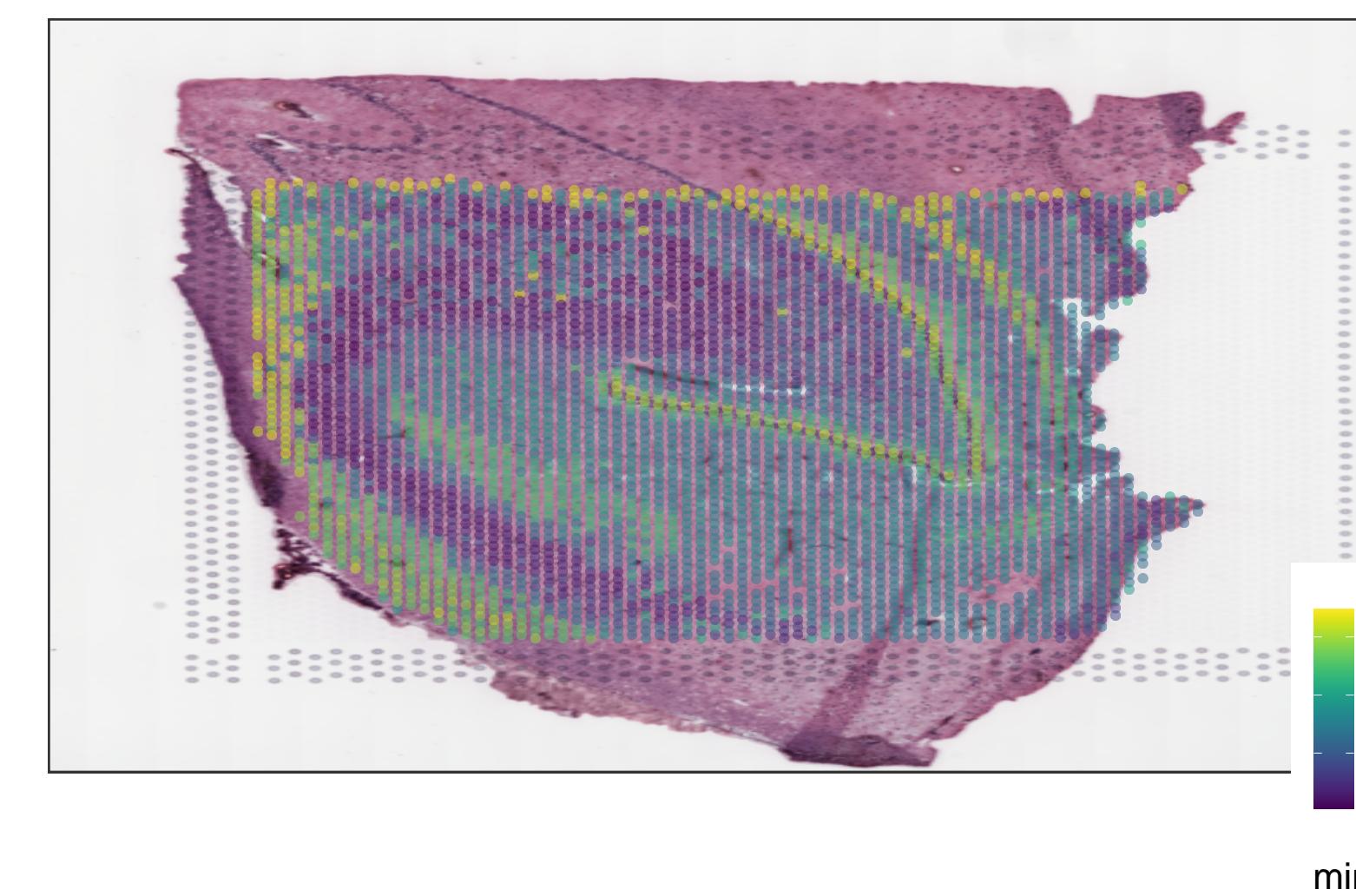


# Br8686 pseudotime TRUE

## Br2706 pseudotime TRUE



Br3942 pseudotime TRUE



Br5699\_new pseudotime TRUE

