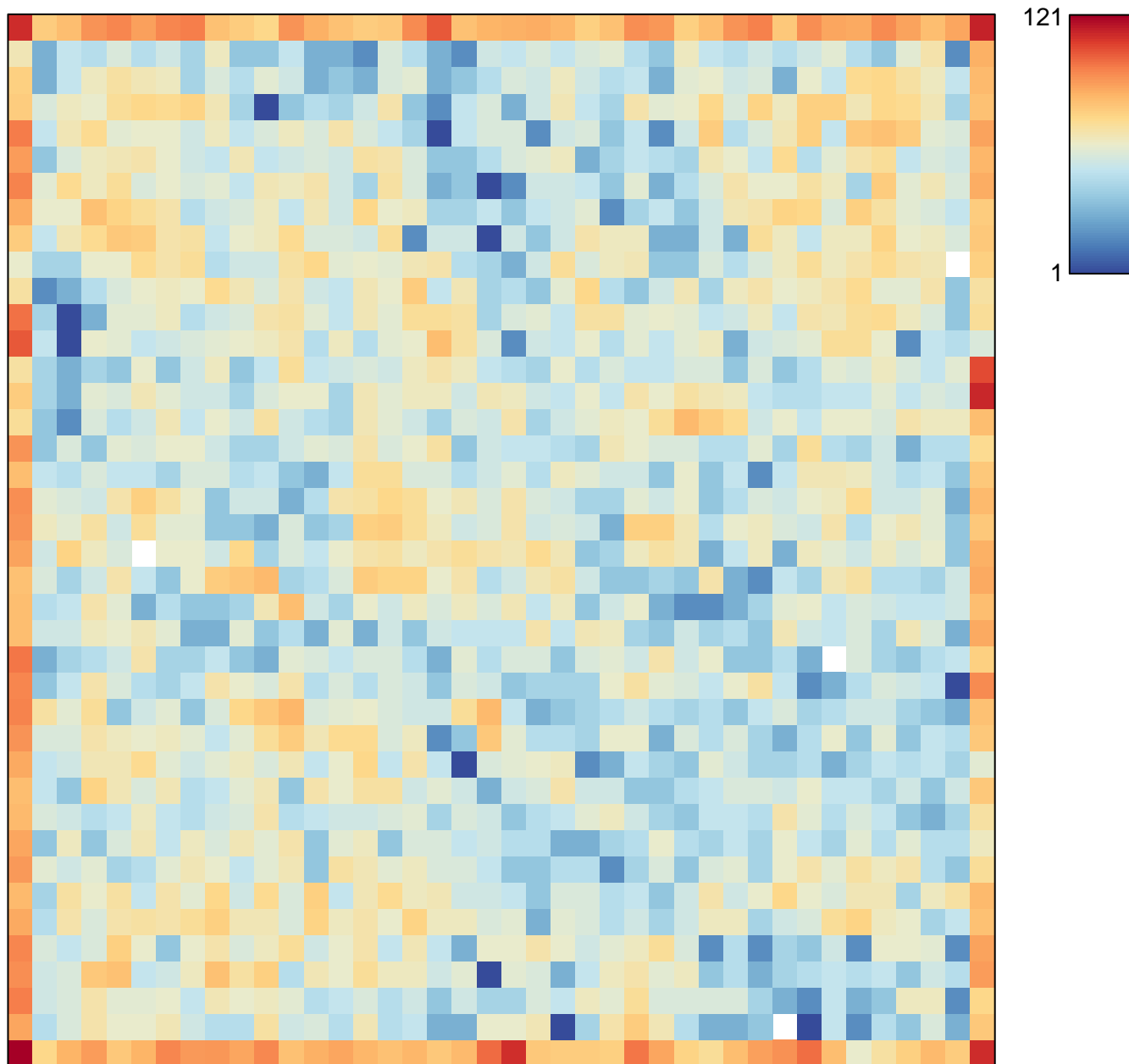
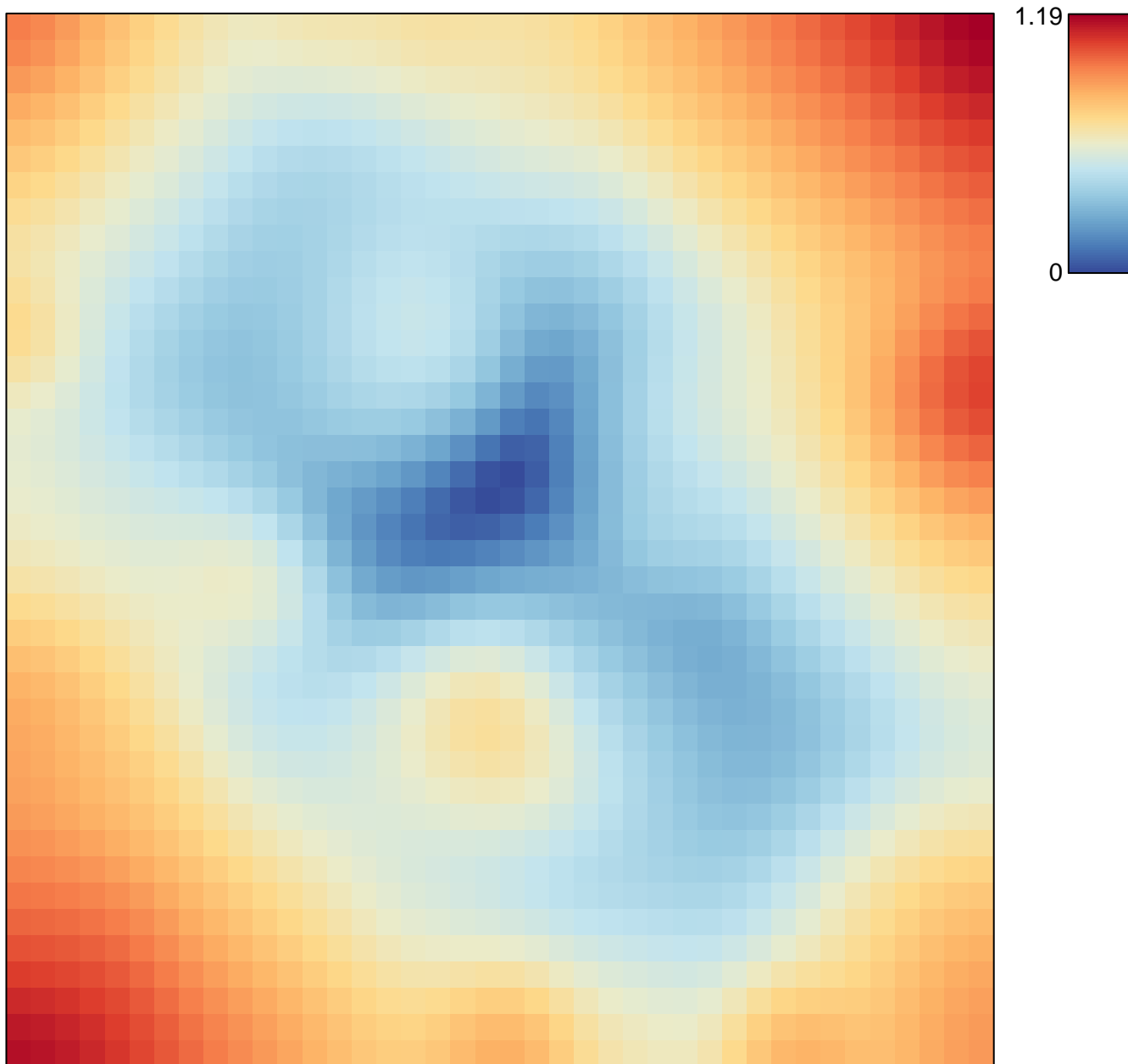


# Population Map



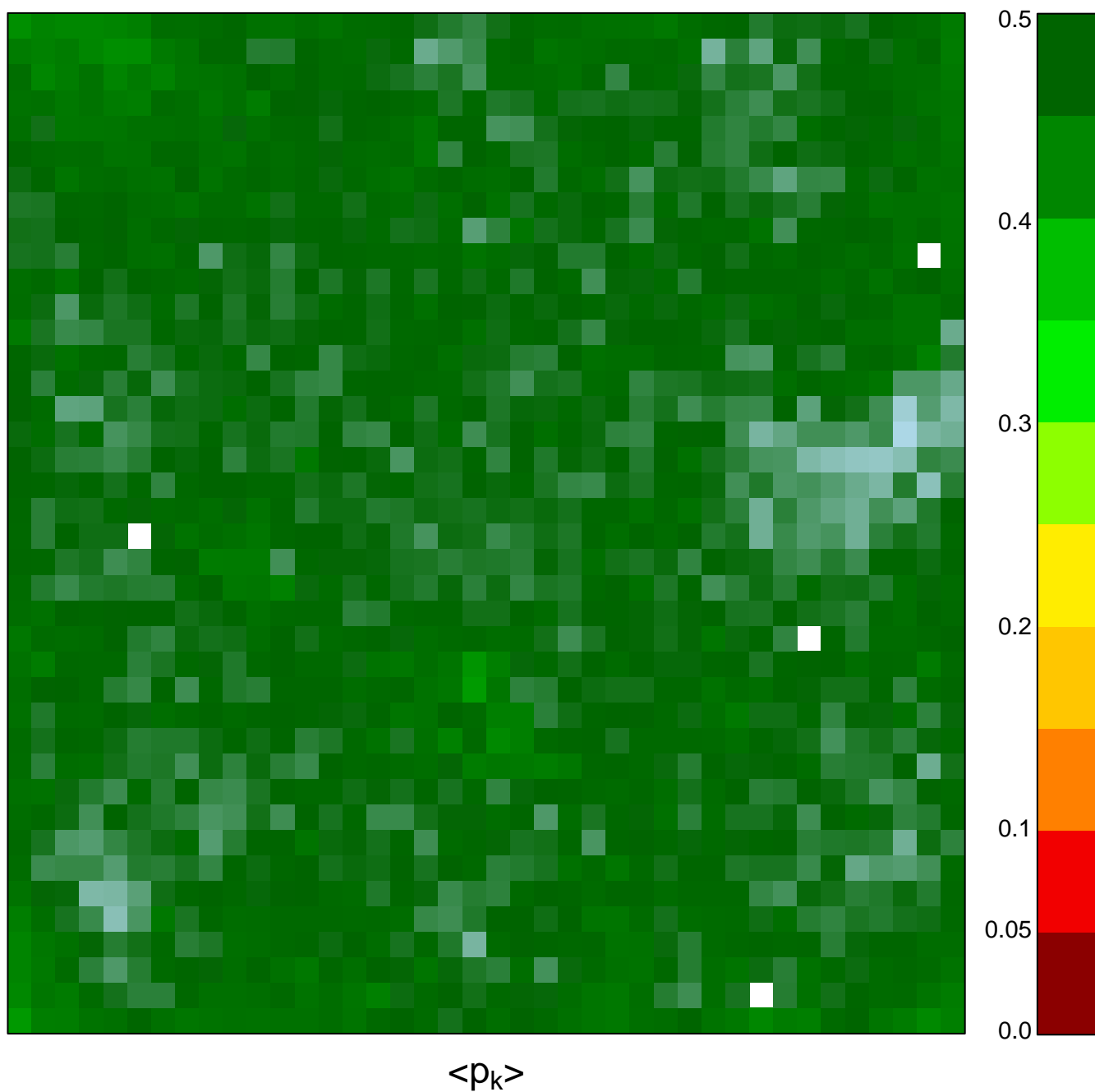
log ( # genes in metagene )

# Metagene Variance Map

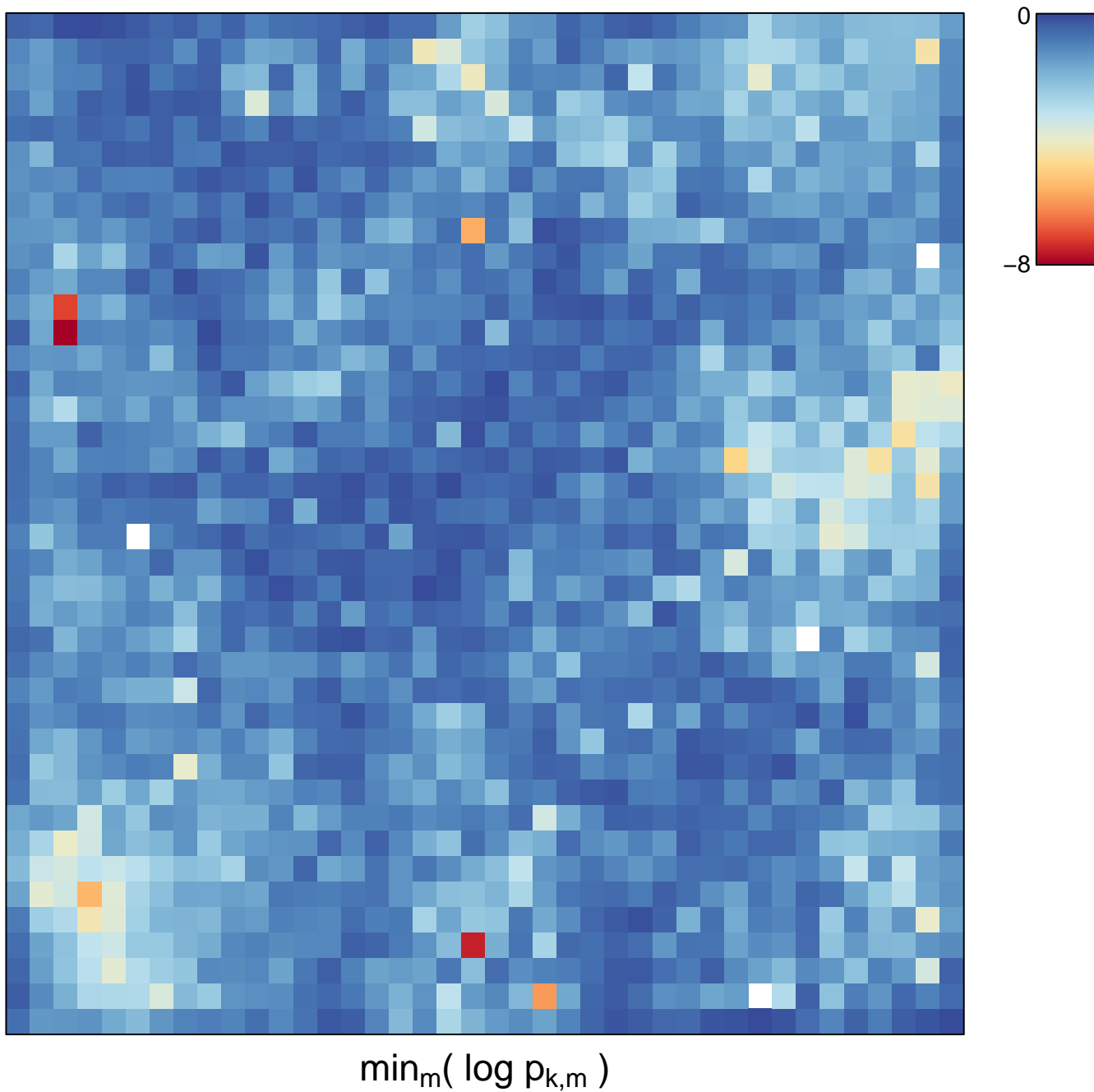


log ( metagene variance )

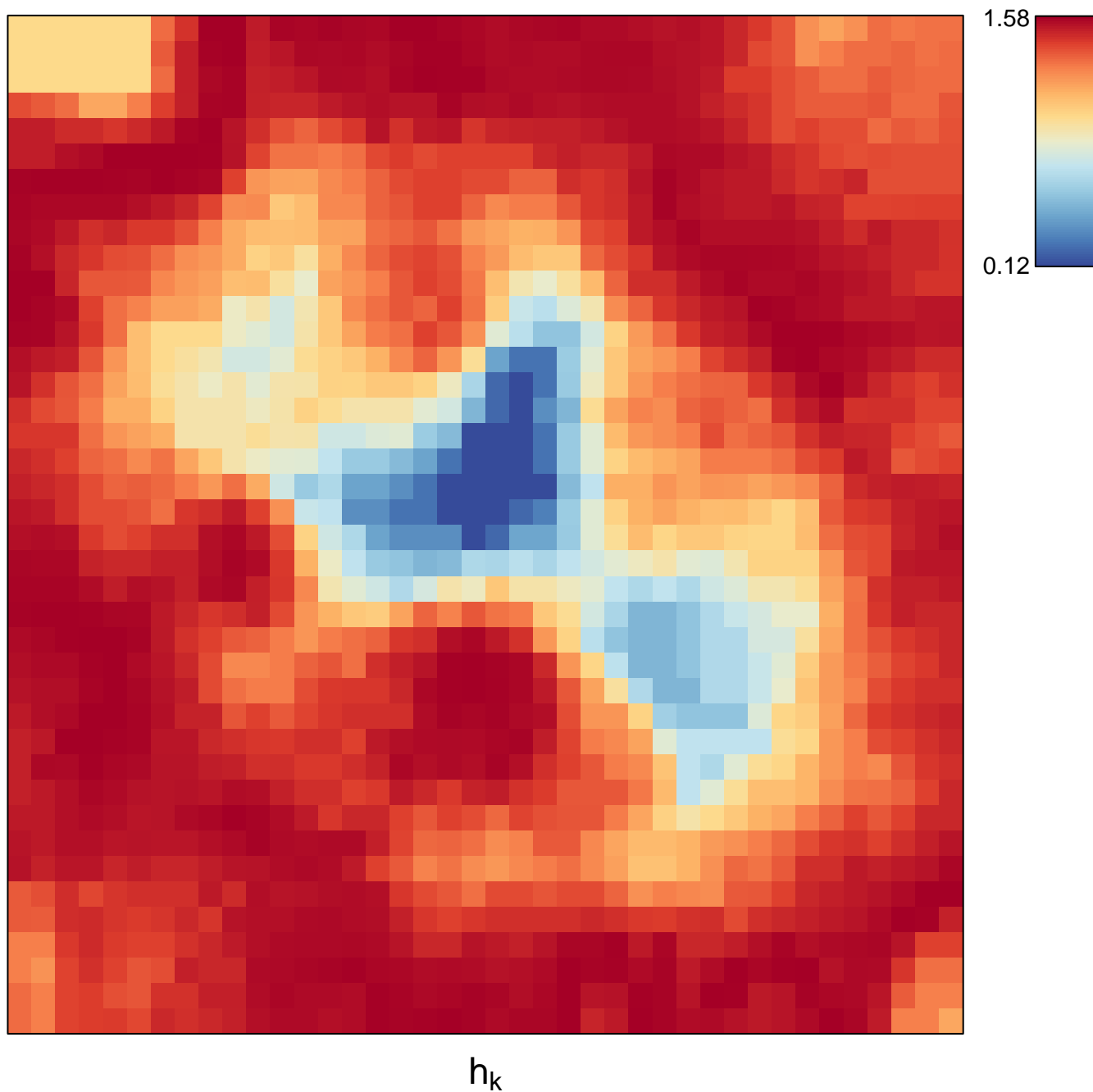
# Metagene Significance Map



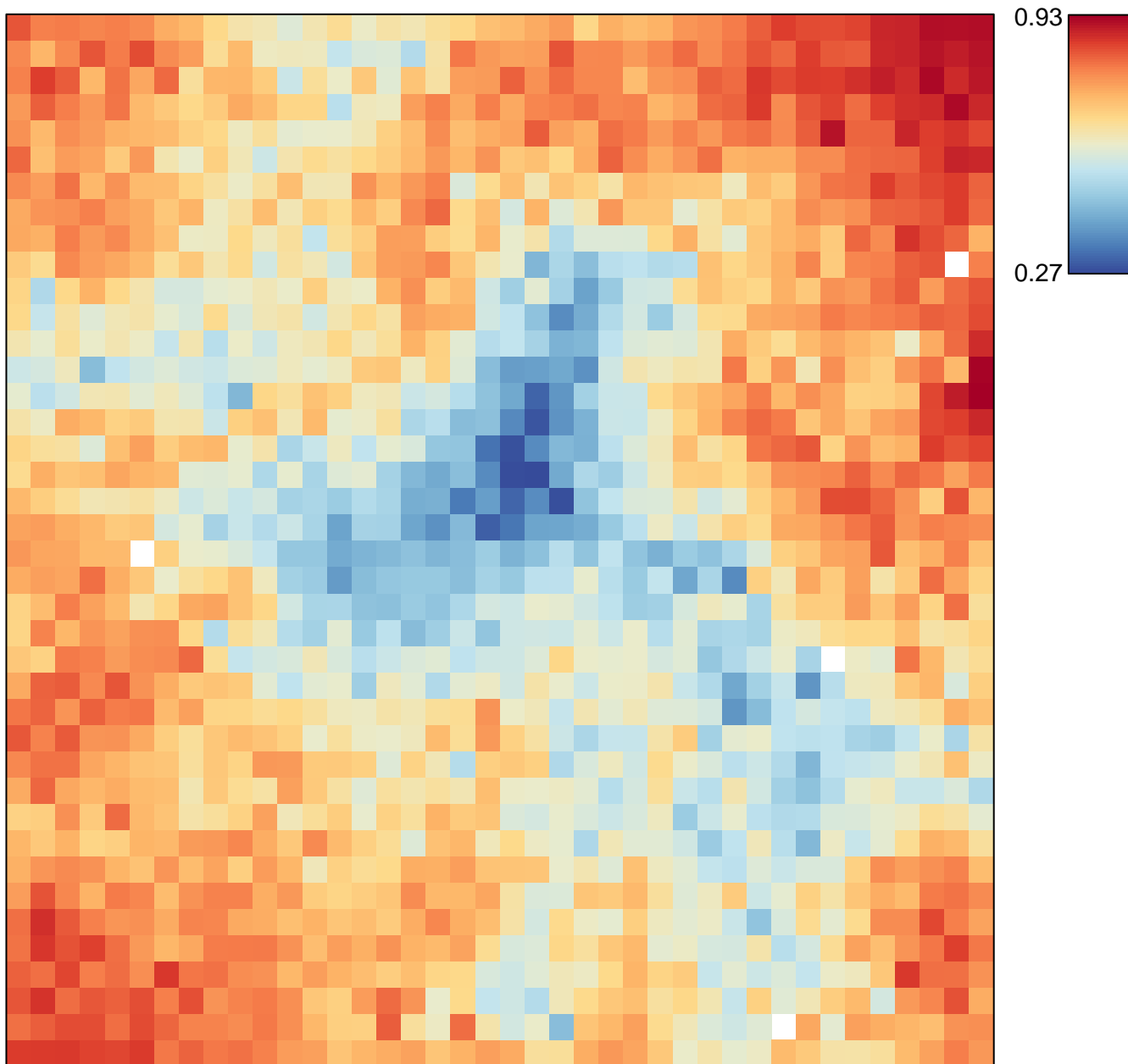
# Metagene Significance Map



# Standard Metagene Entropy Map

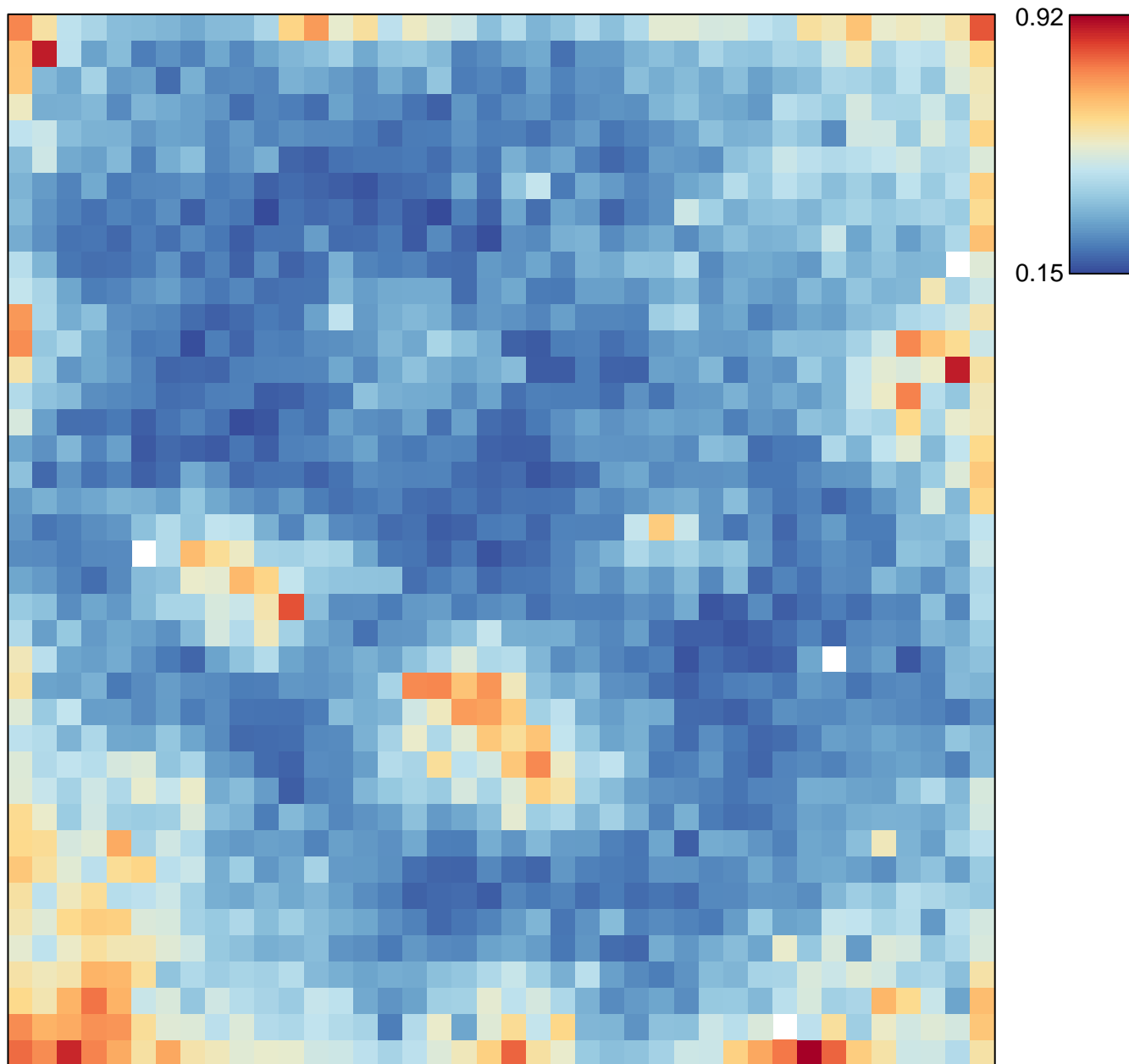


# Gene–Metagene Covariance Map



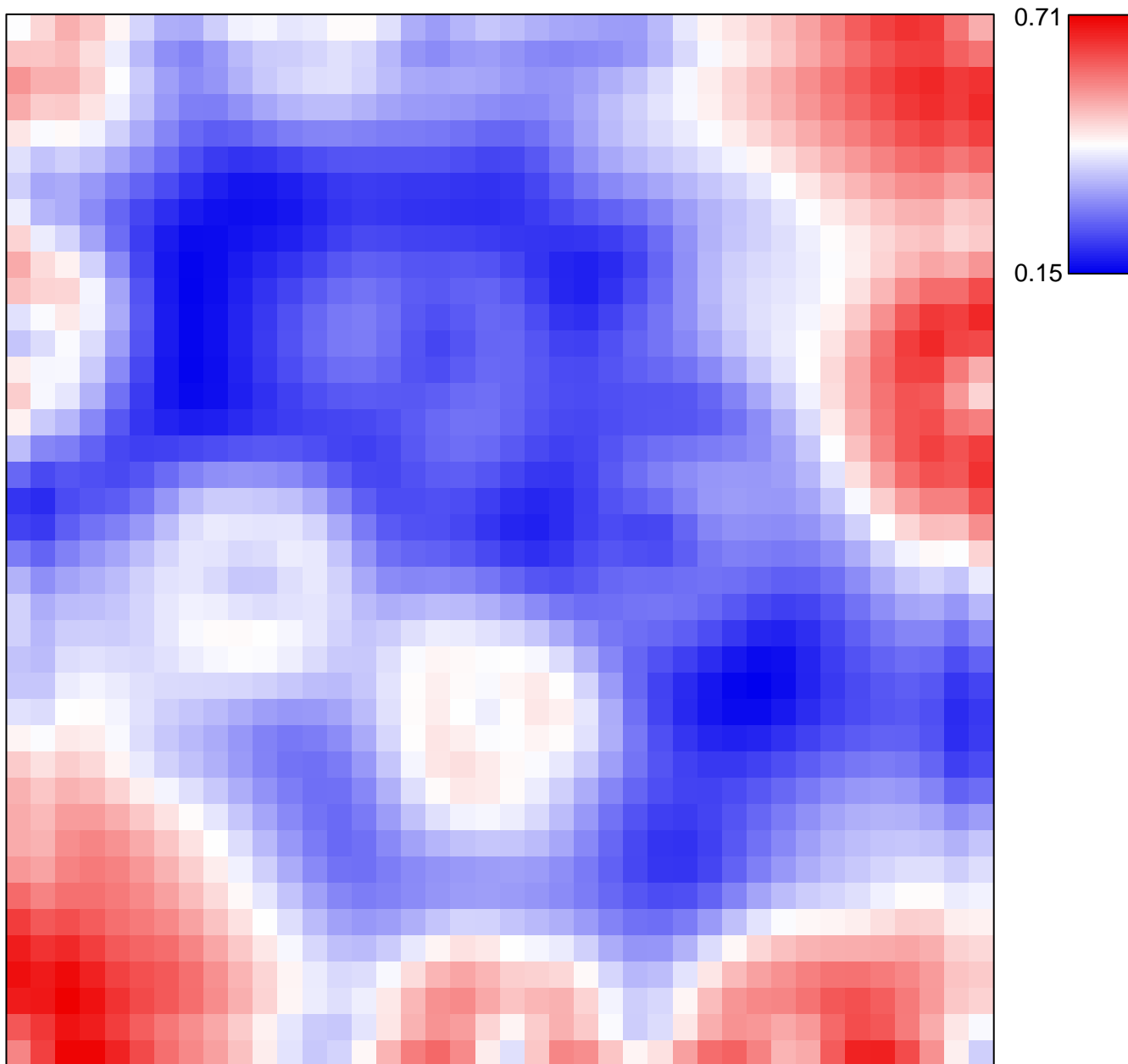
correlation genes – metagene

# Deviation Map



deviation genes – metagene

# Distance Map



deviation of adjacent metagenes