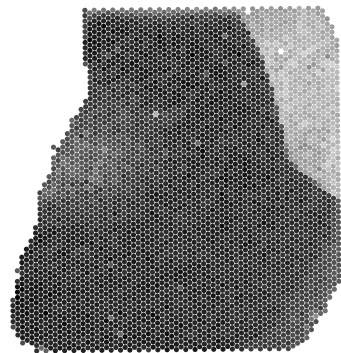
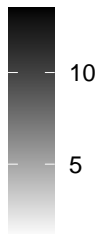


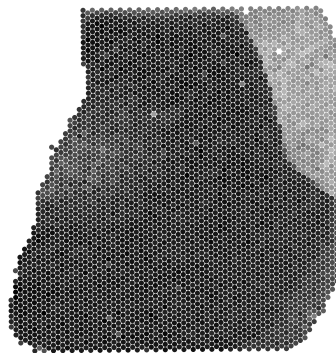
Library size



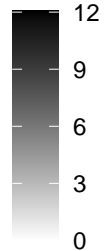
sum_umi_log



Unique genes



sum_gene_log



Mito percent



expr_chrM_ratio

