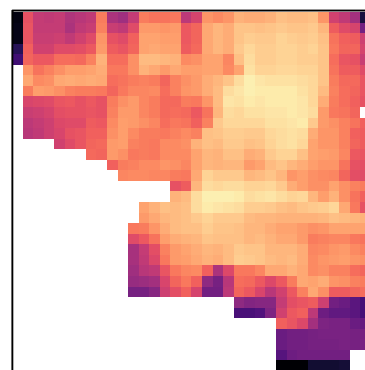
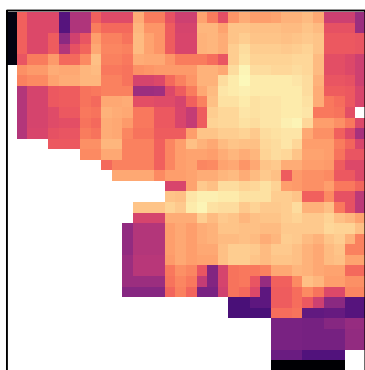
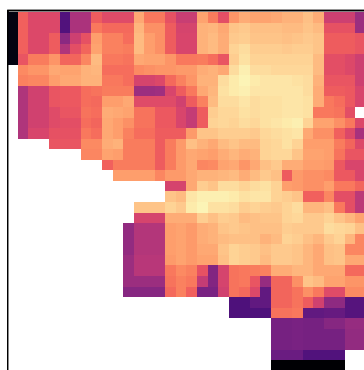
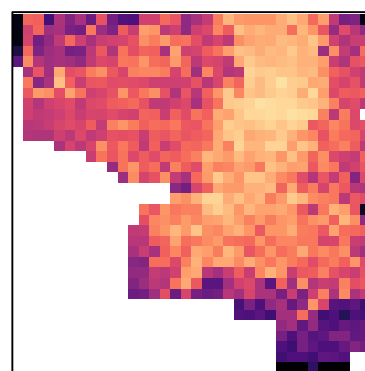
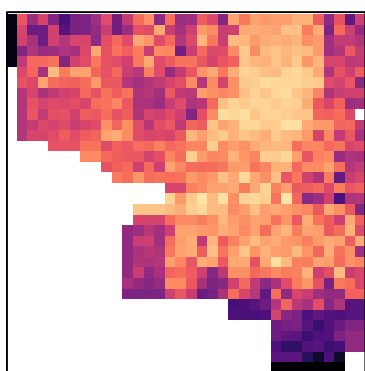
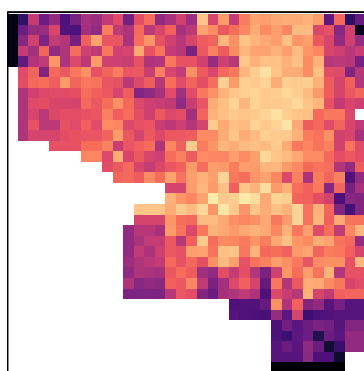
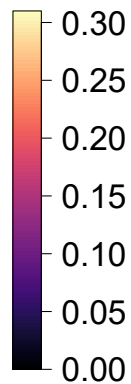


10,000 SNPs
200 samples
RADseq-type dataset

100,000 SNPs
200 samples
WGS-type dataset

142,129 SNPs
1,688 samples
Full dataset

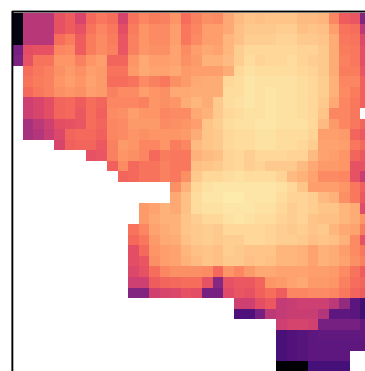
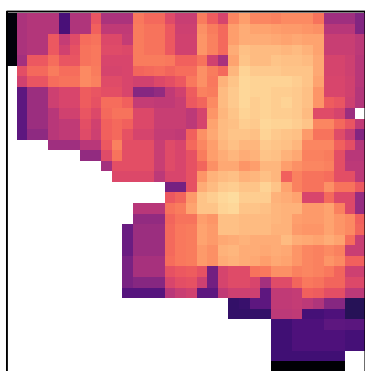
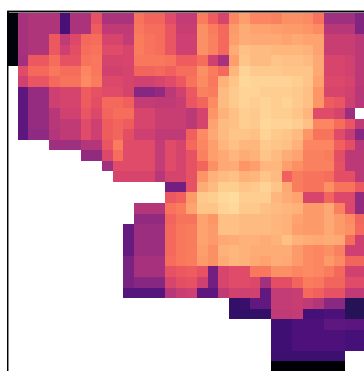
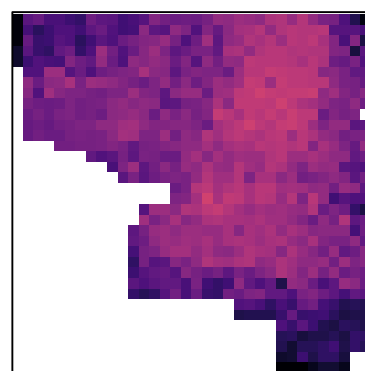
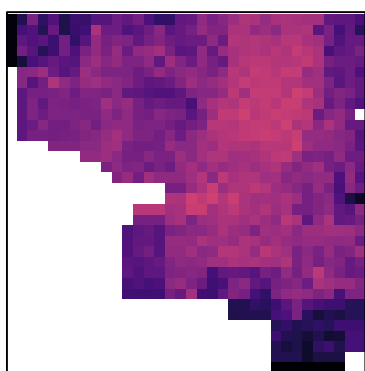
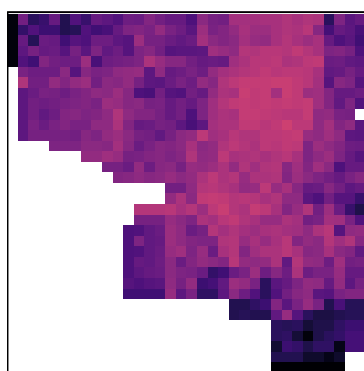
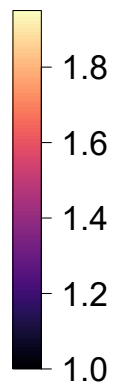
(a) π



Rarified

Not rarified

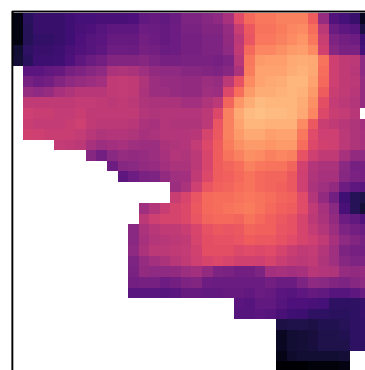
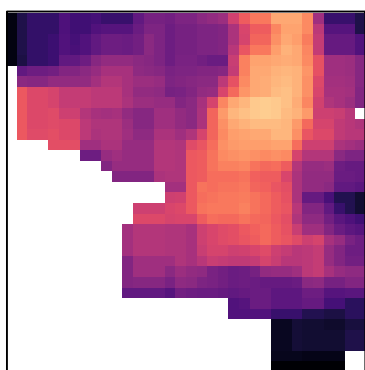
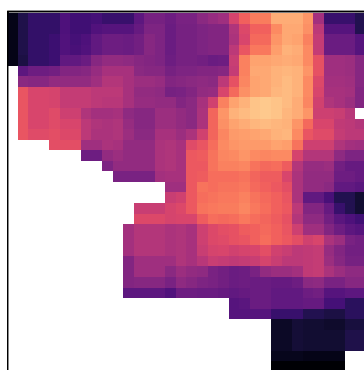
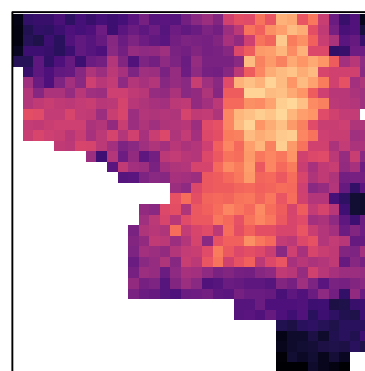
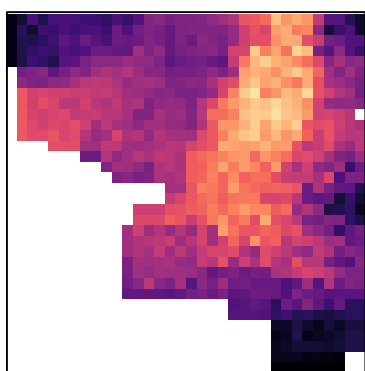
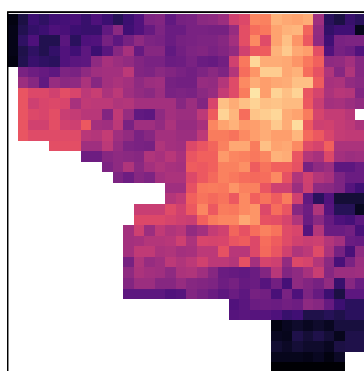
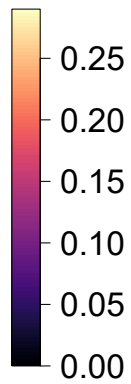
(b) AR



Rarified

Not rarified

(c) H



Rarified

Not rarified