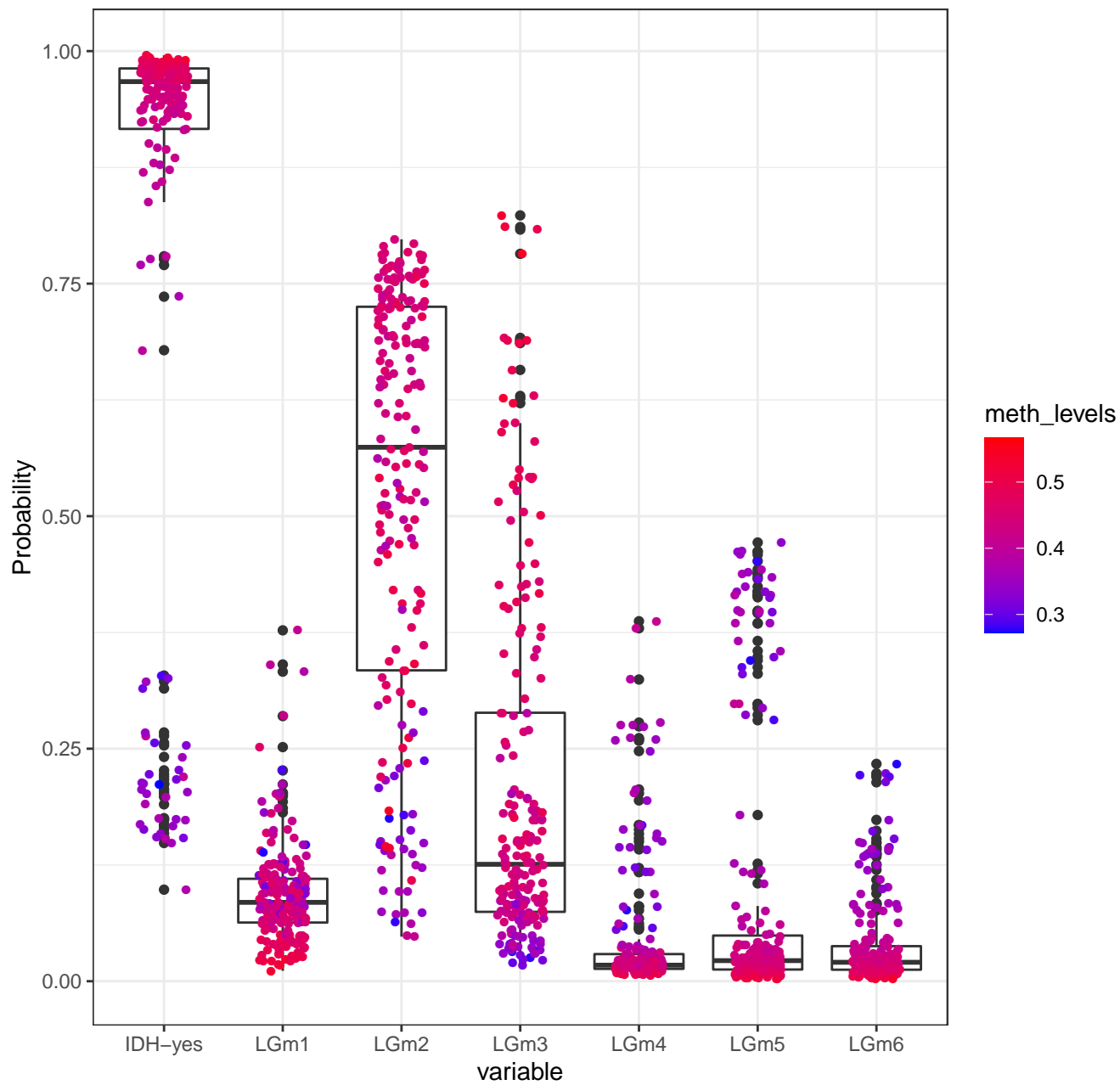
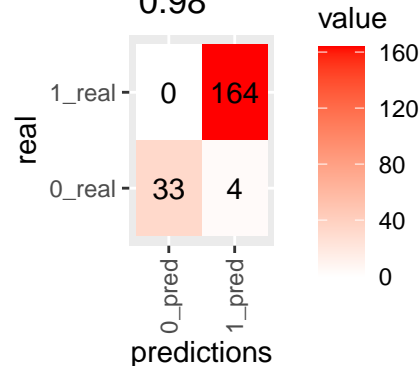


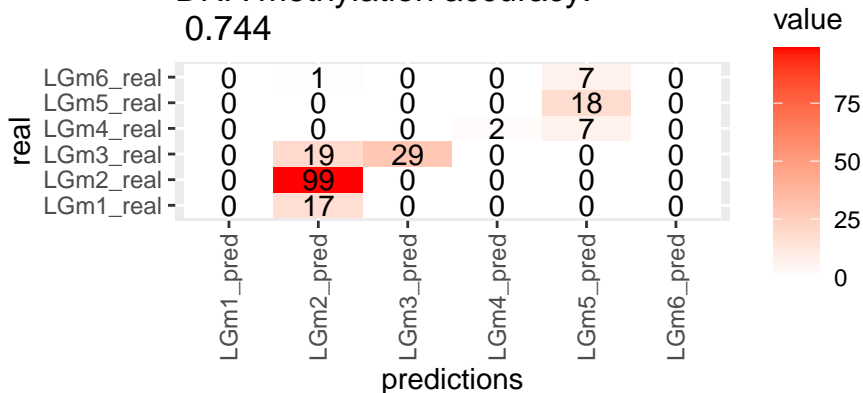
N.patients: 199–201



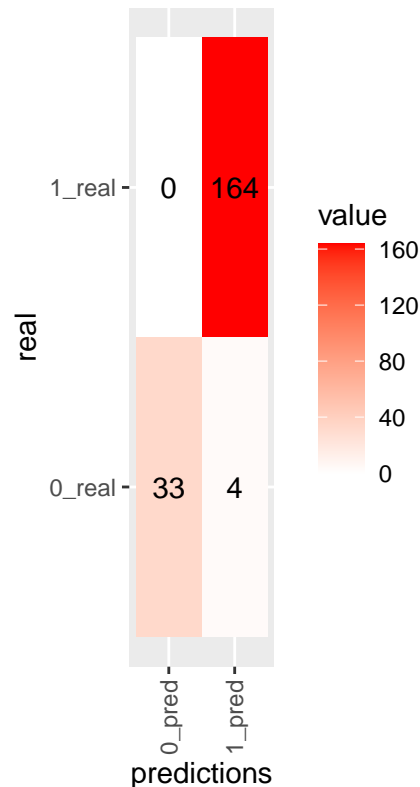
IDH model accuracy:  
0.98



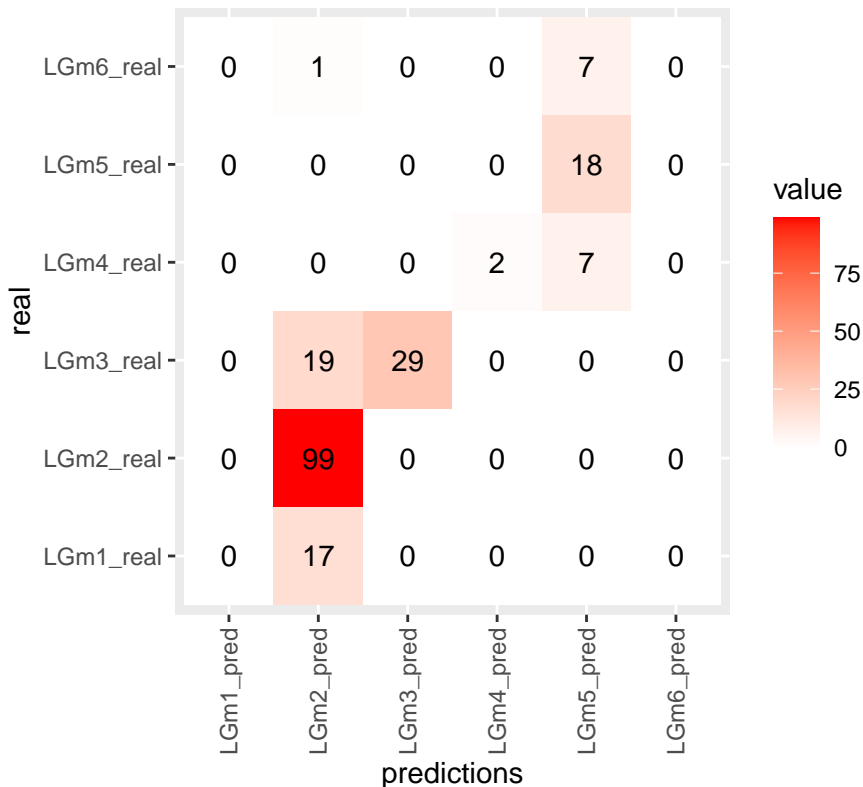
DNA methylation accuracy:  
0.744

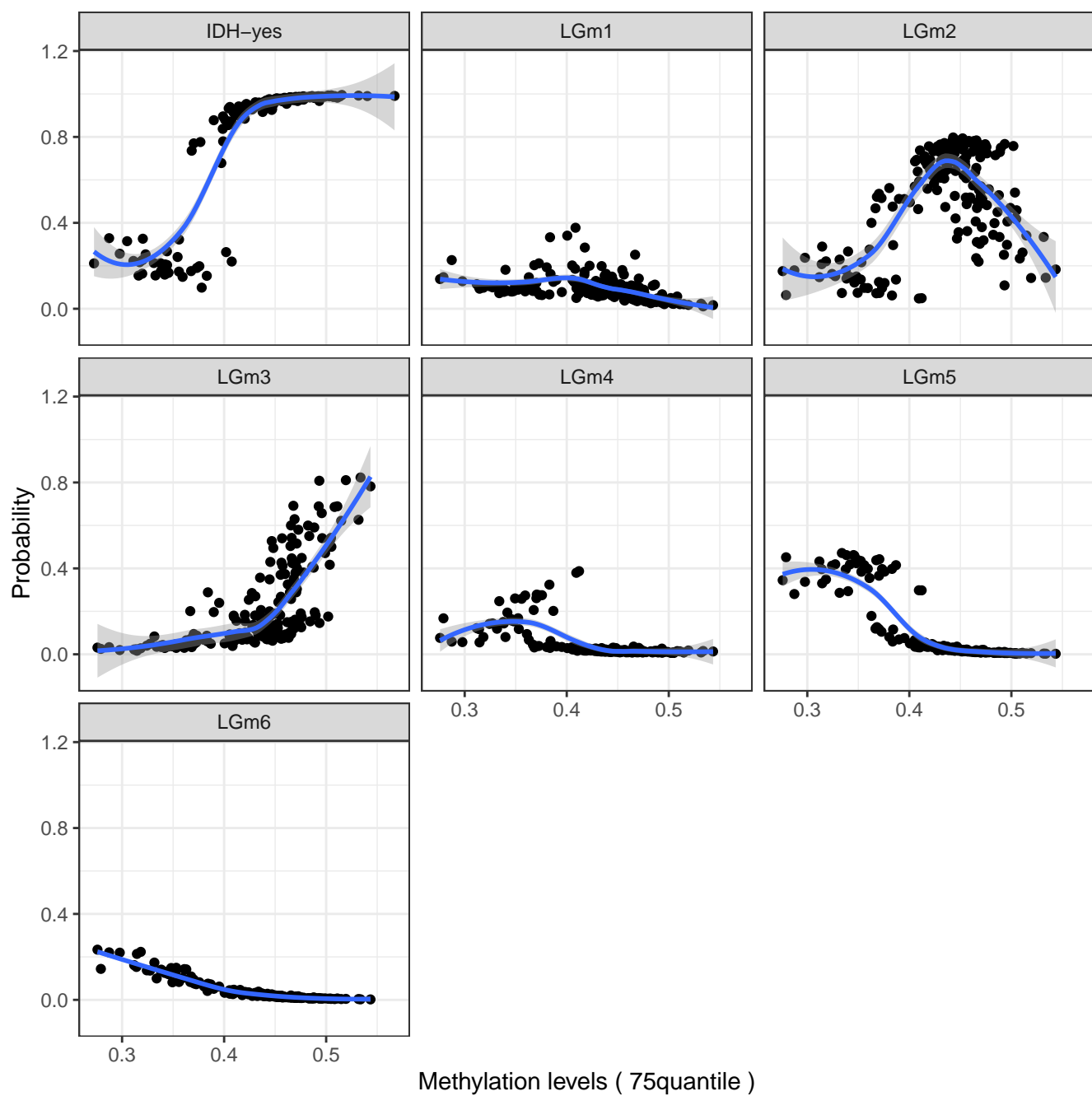


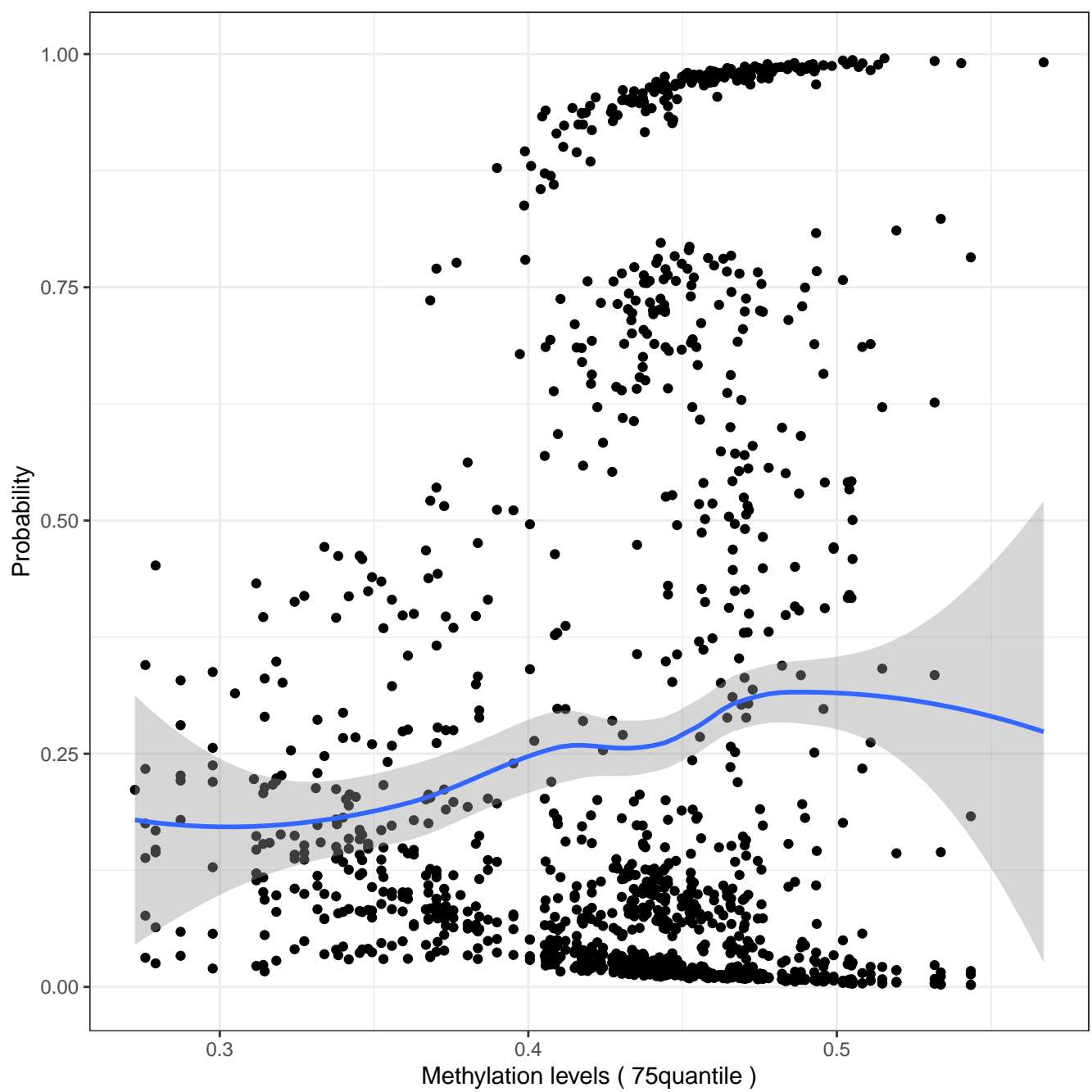
IDH model accuracy:  
0.98

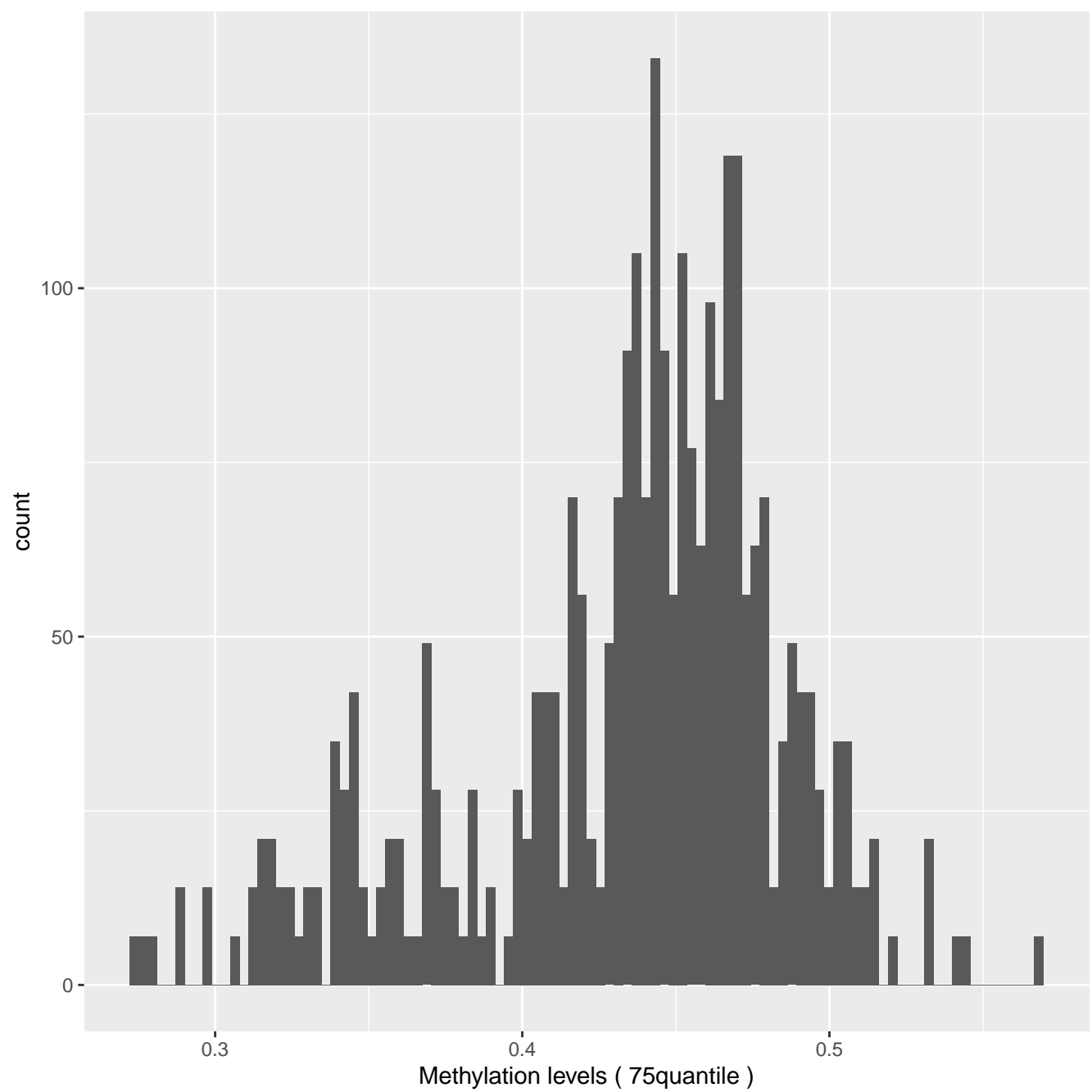


DNA methylation accuracy:  
0.744

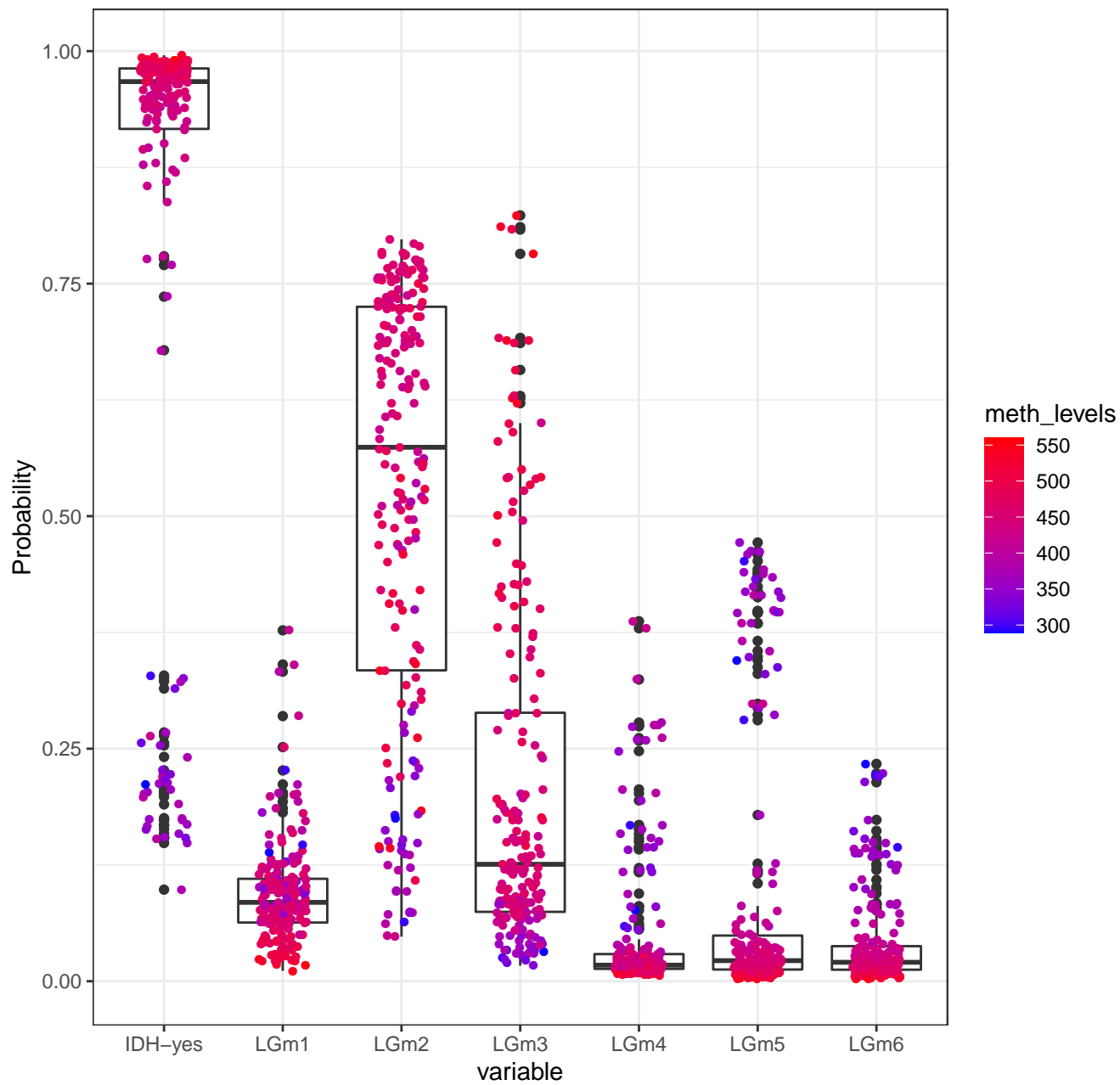




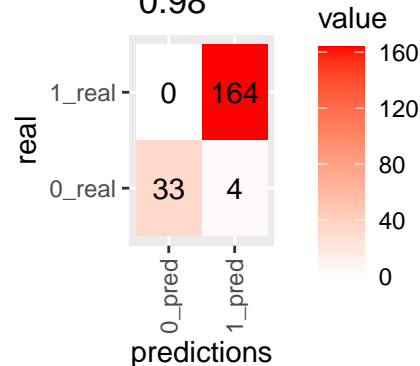




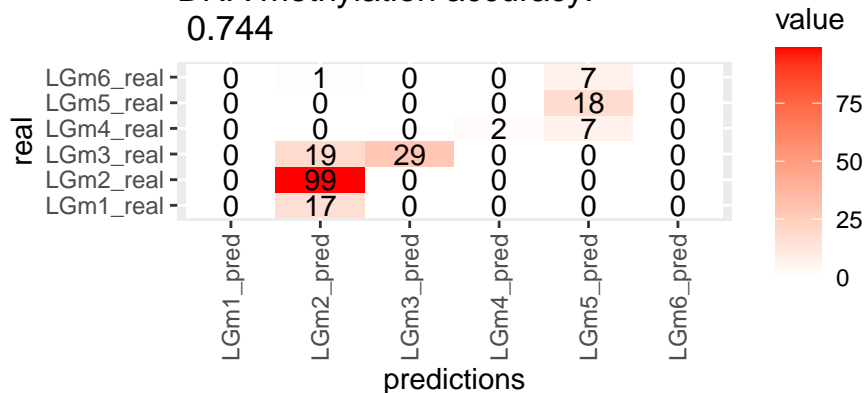
N.patients: 199–201



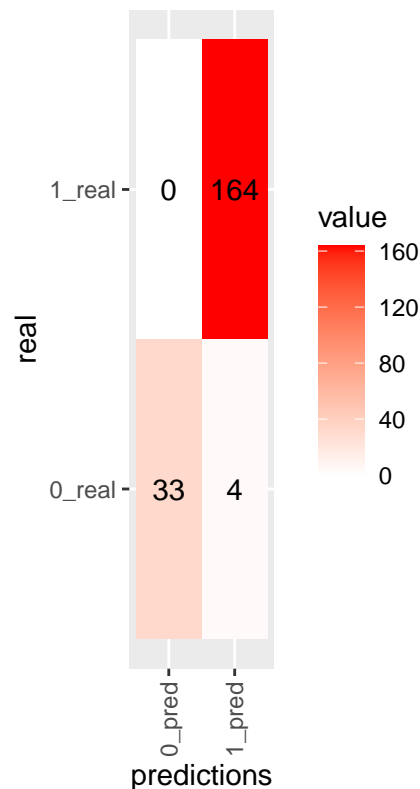
IDH model accuracy:  
0.98



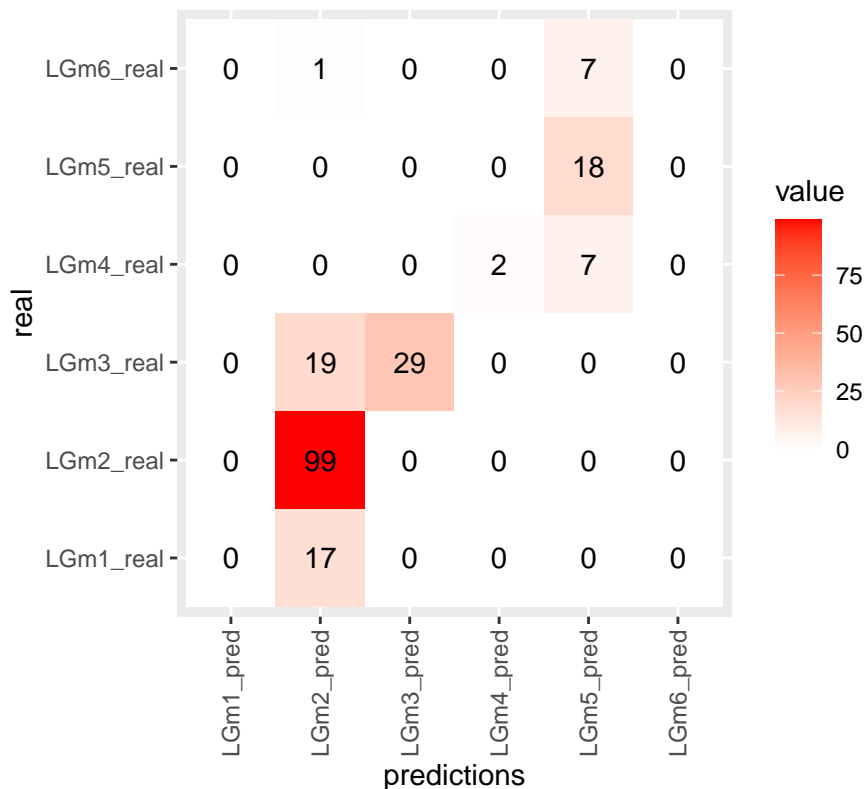
DNA methylation accuracy:  
0.744

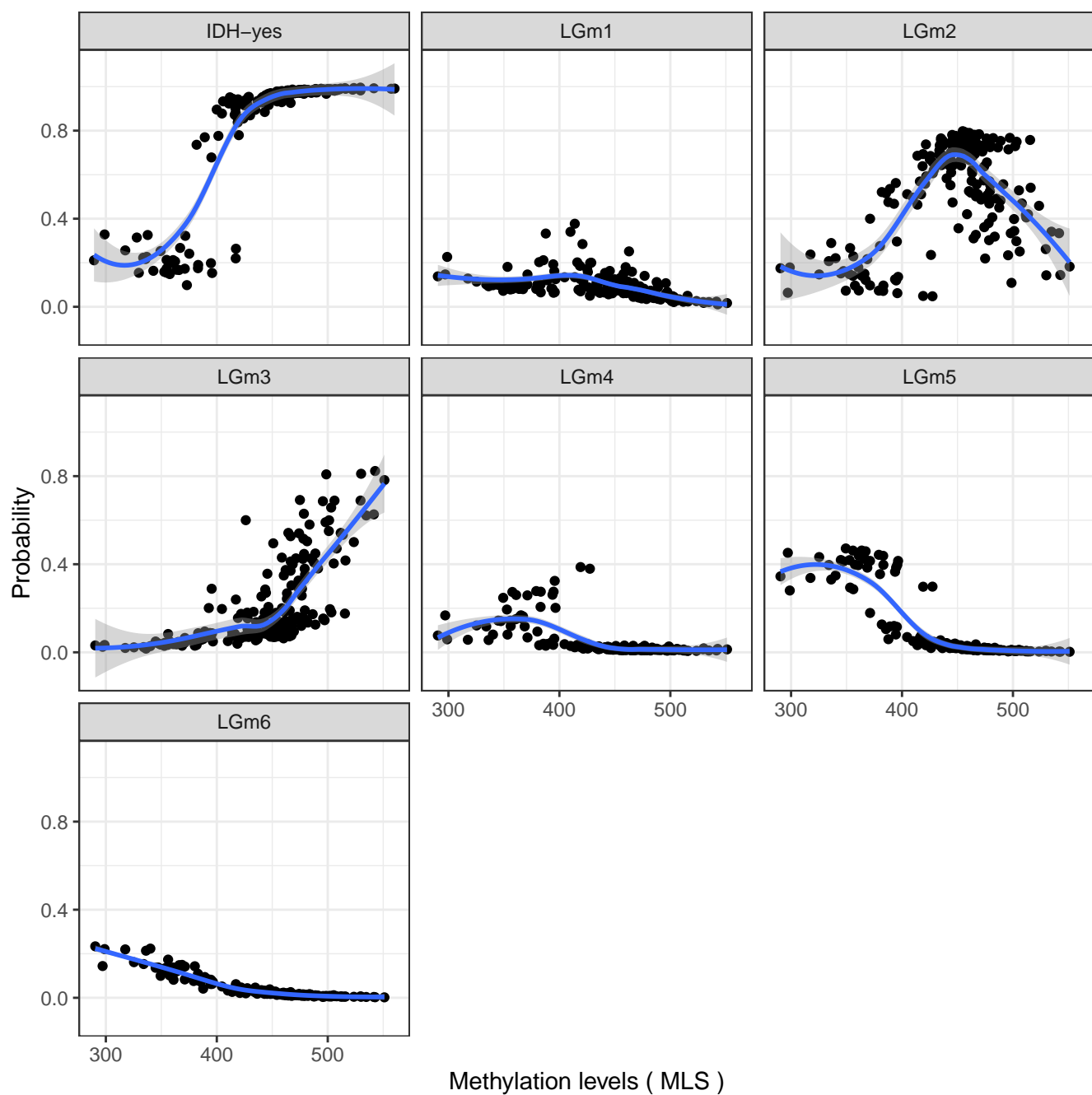


IDH model accuracy:  
0.98

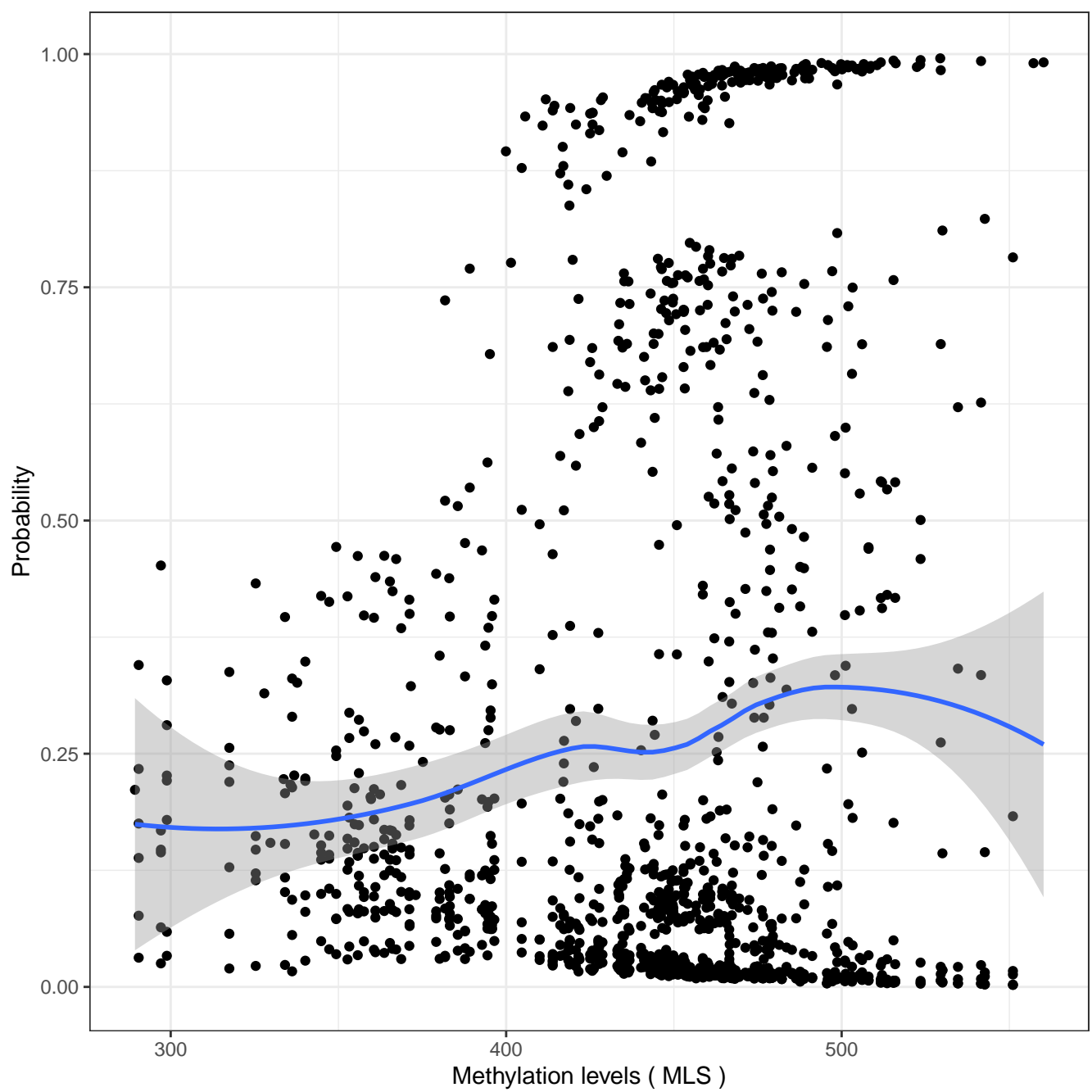


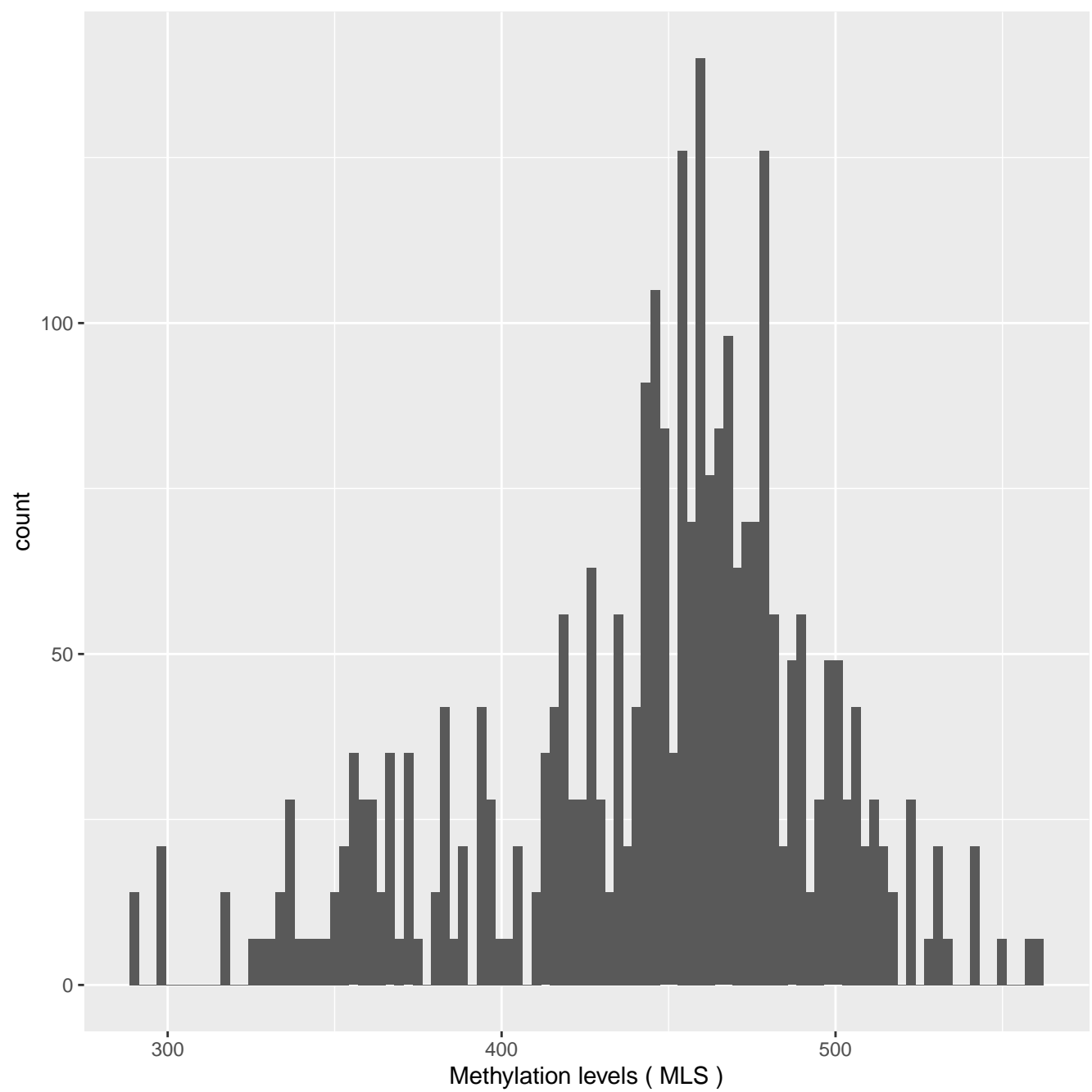
DNA methylation accuracy:  
0.744



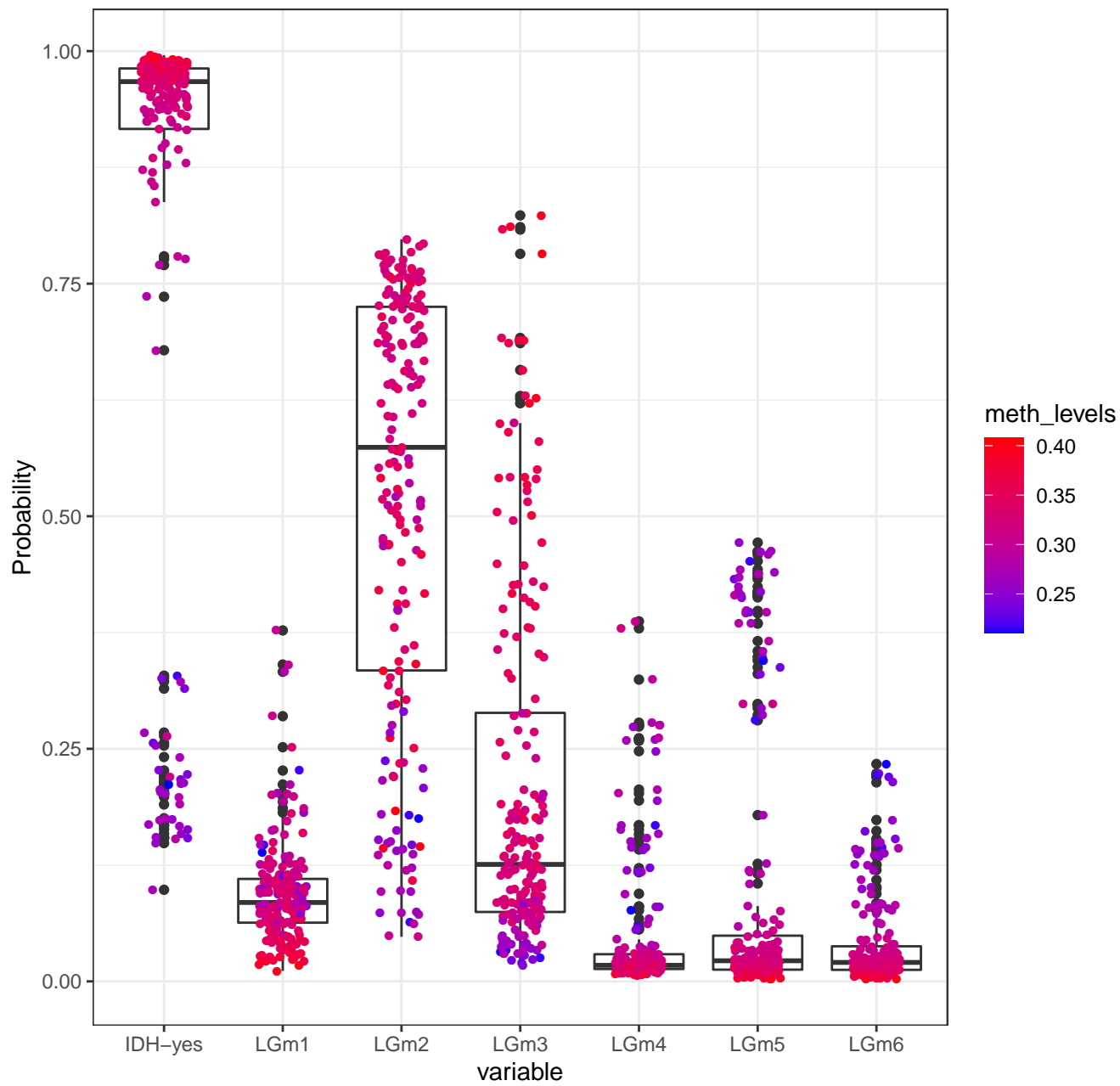




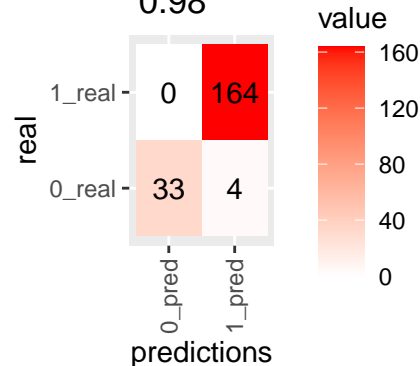




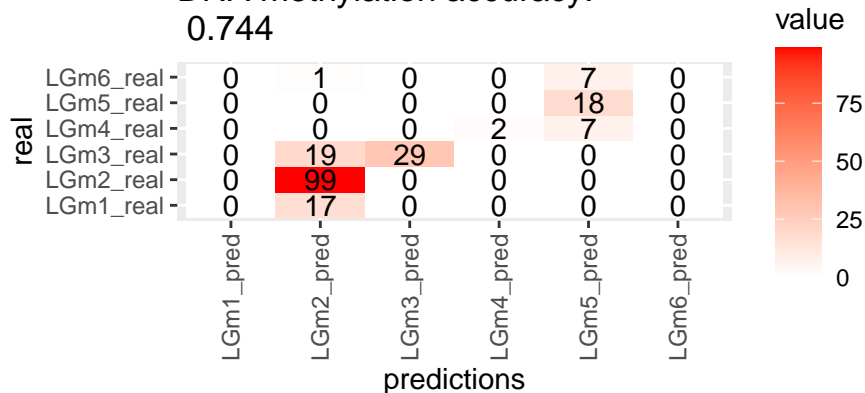
N.patients: 199–201



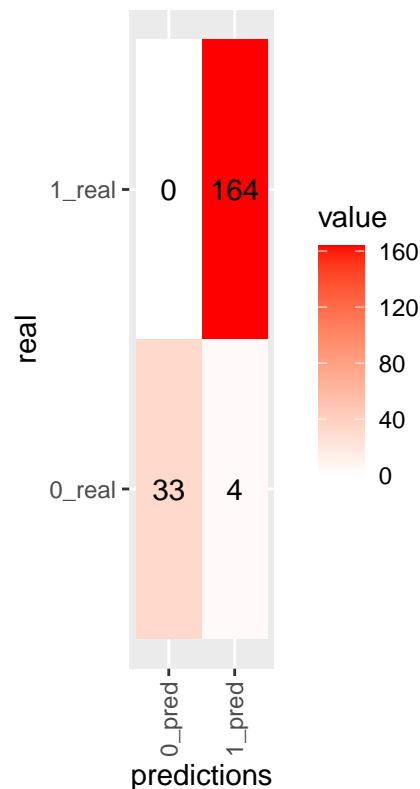
IDH model accuracy:  
0.98



DNA methylation accuracy:  
0.744



IDH model accuracy:  
0.98



DNA methylation accuracy:  
0.744

