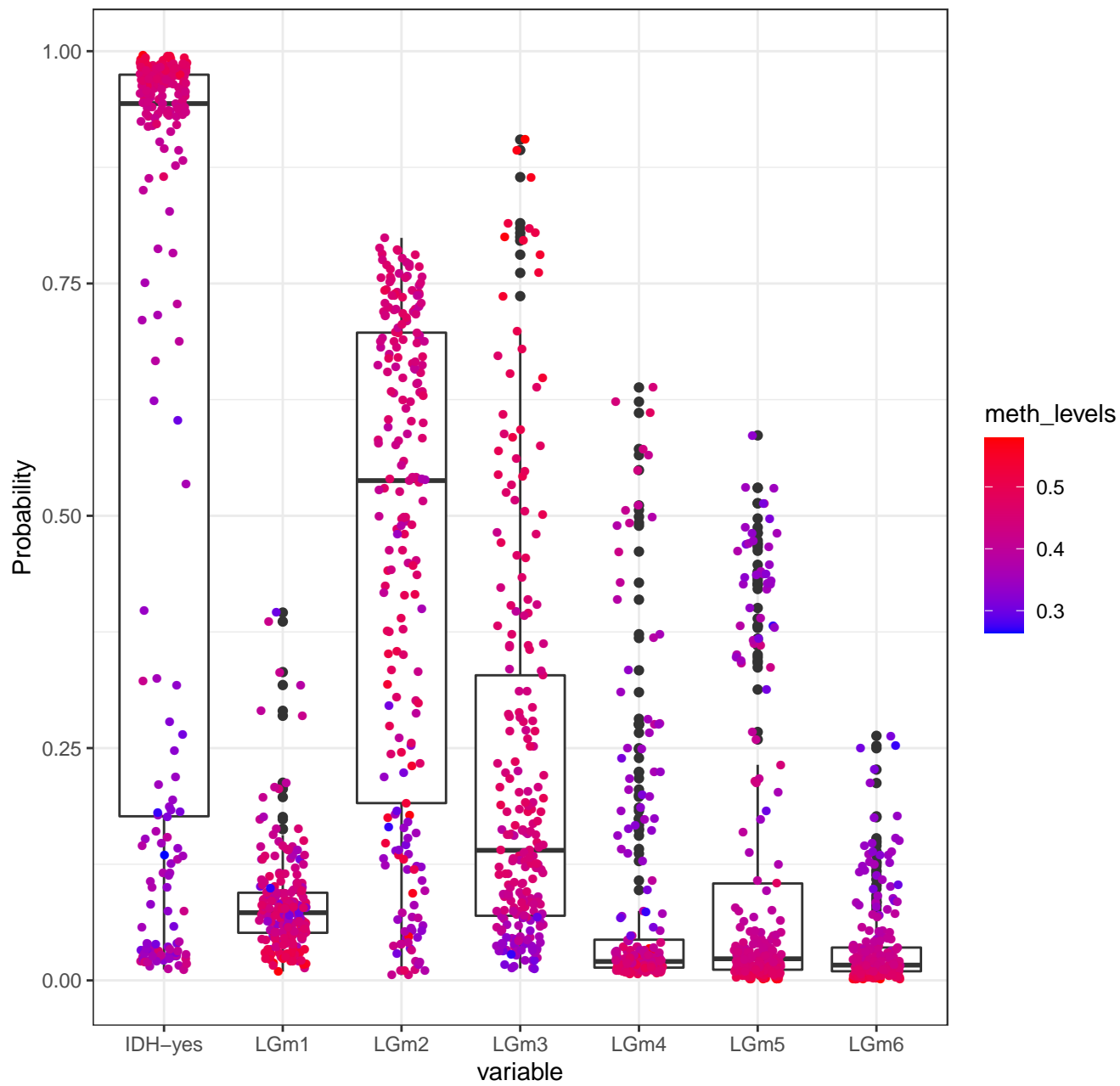
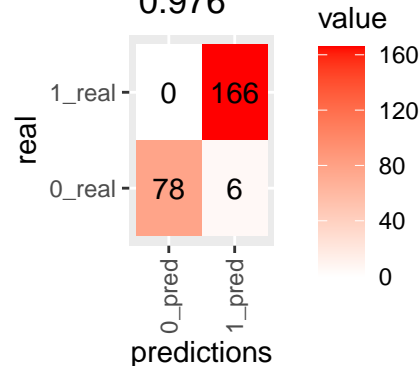


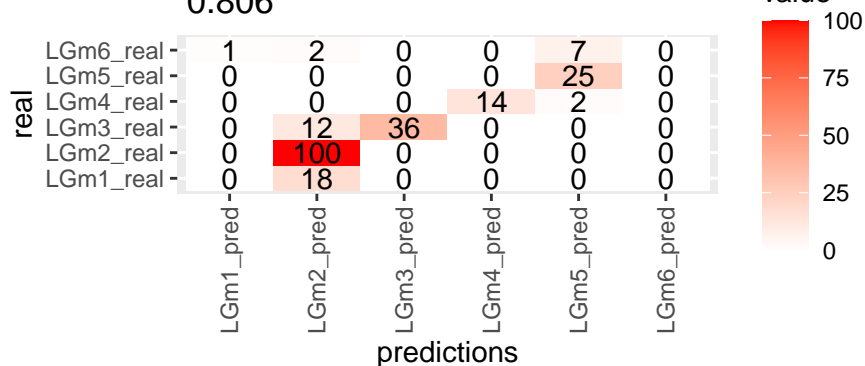
N.patients: 217–250



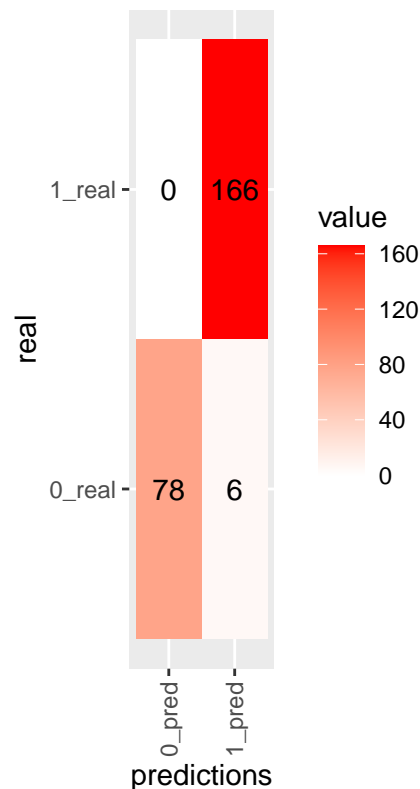
IDH model accuracy:  
0.976



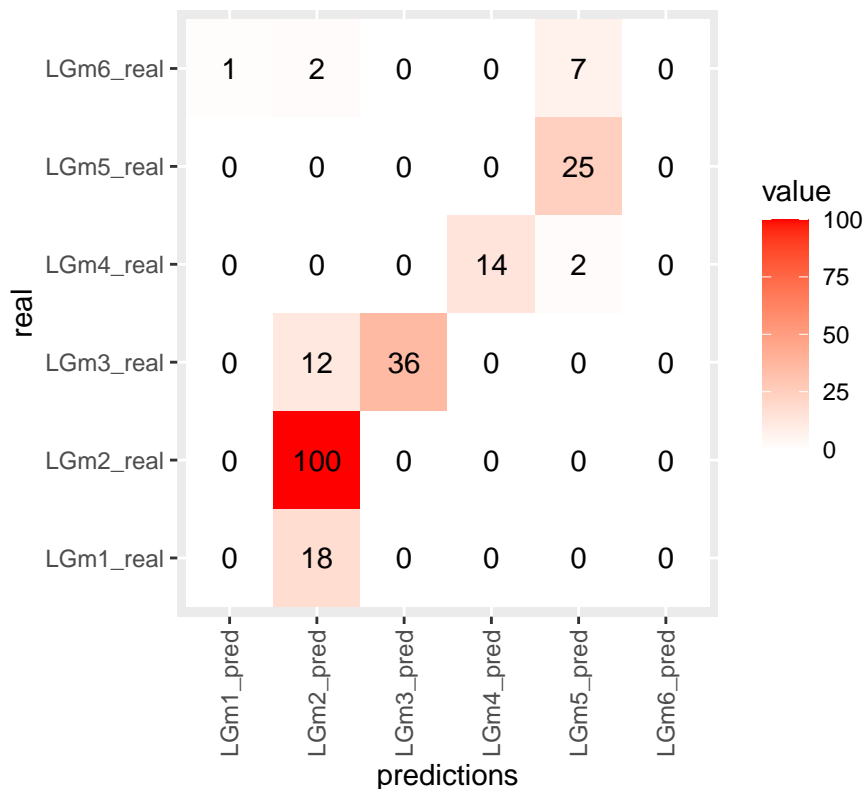
DNA methylation accuracy:  
0.806

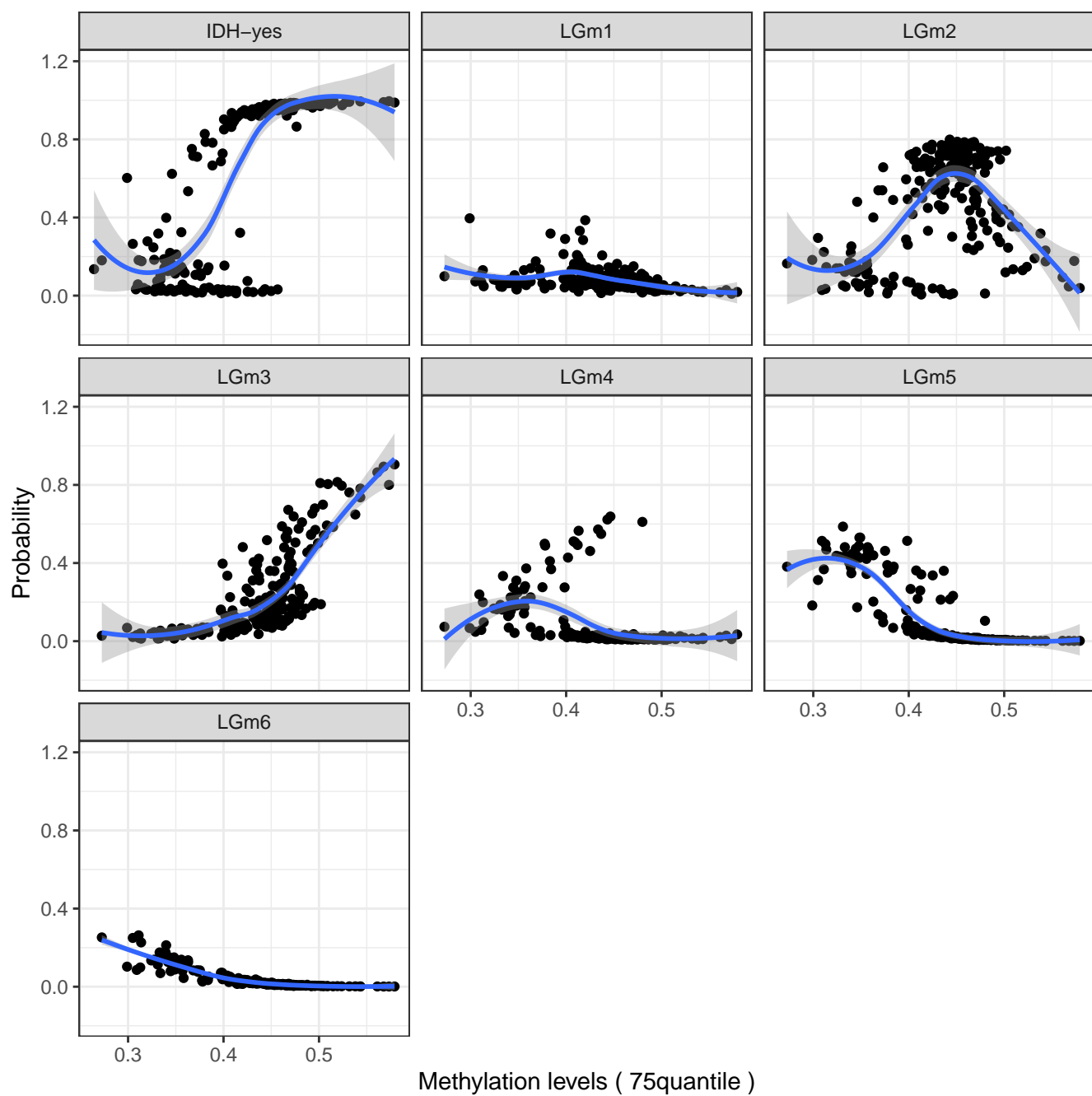


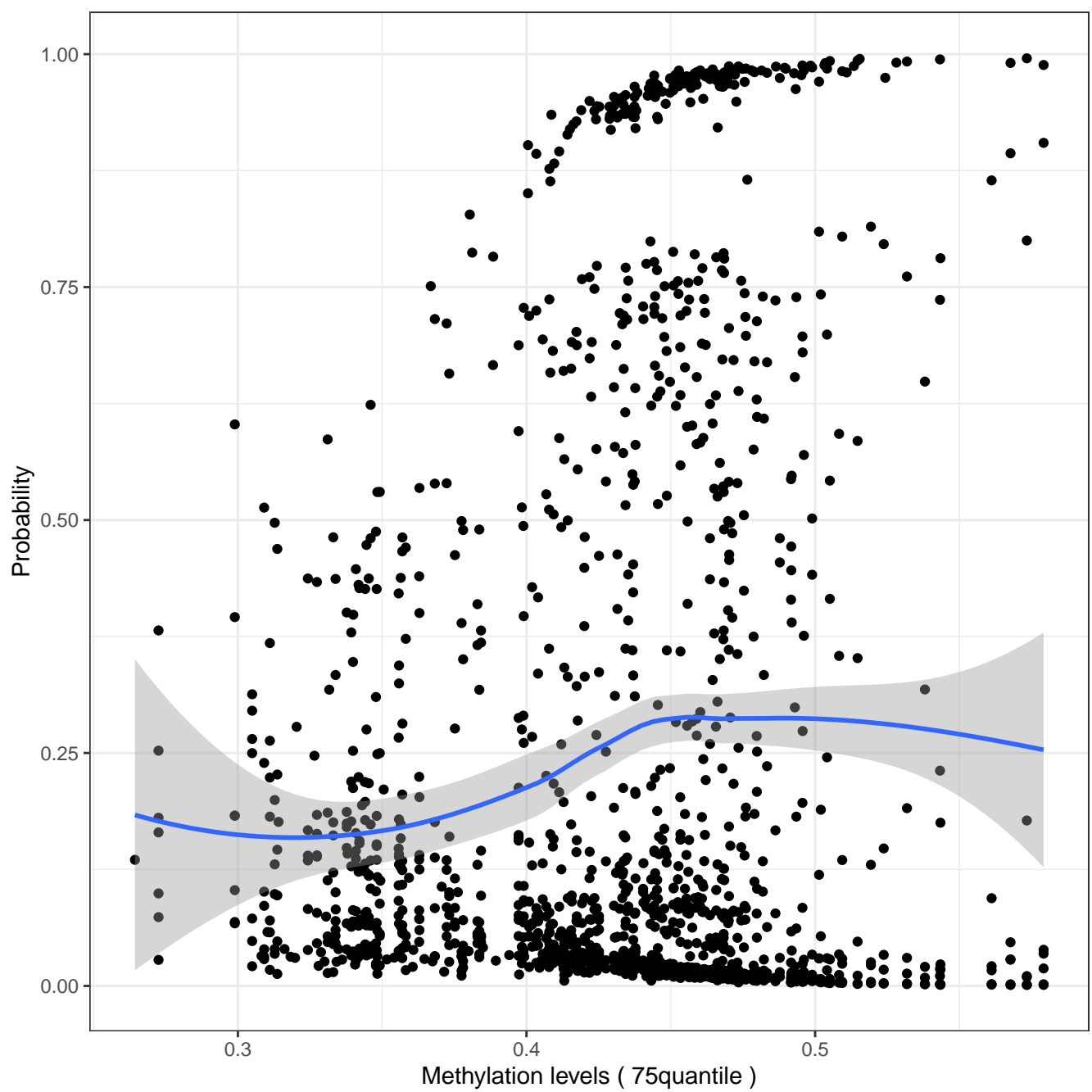
IDH model accuracy:  
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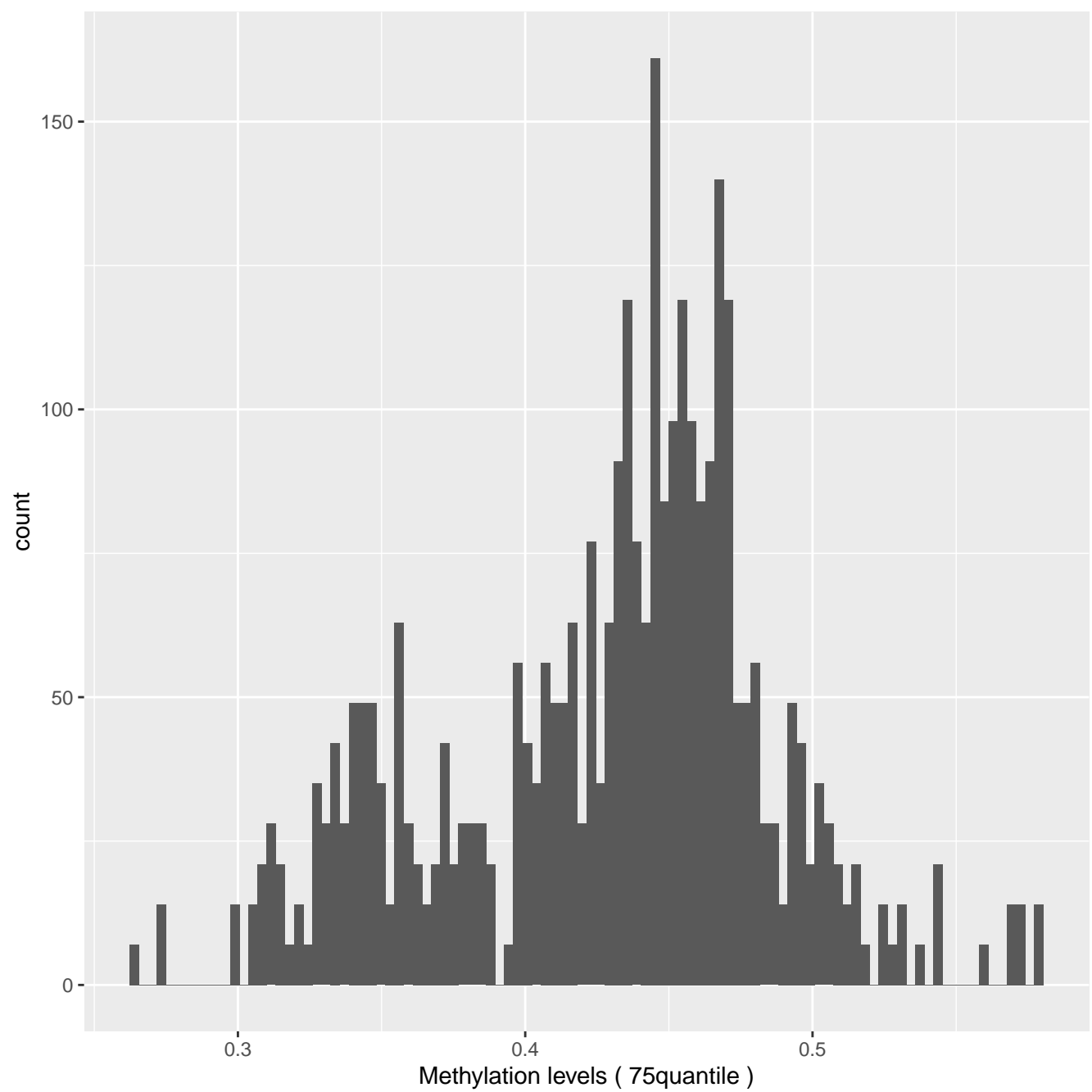


DNA methylation accuracy:  
0.806

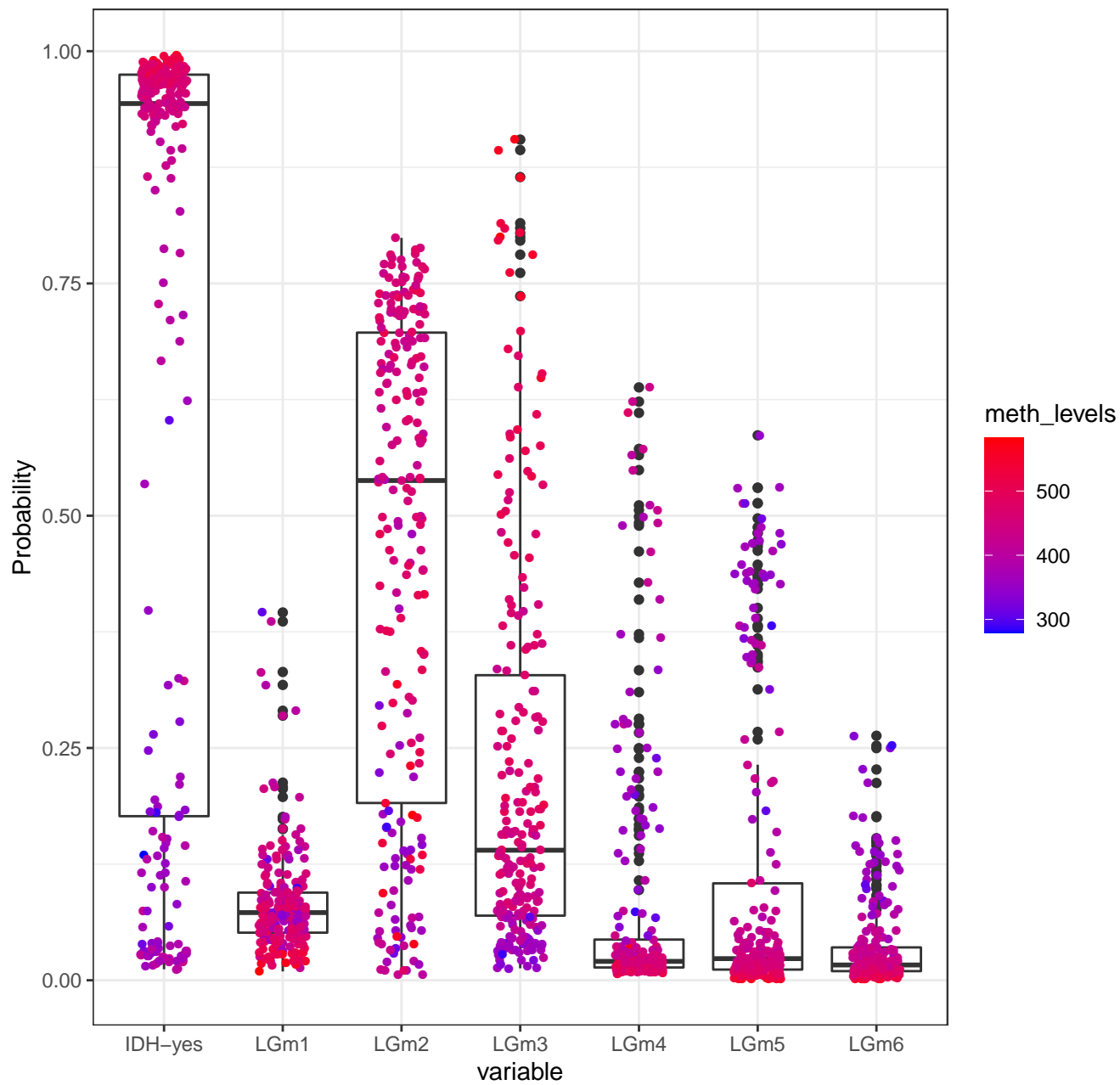




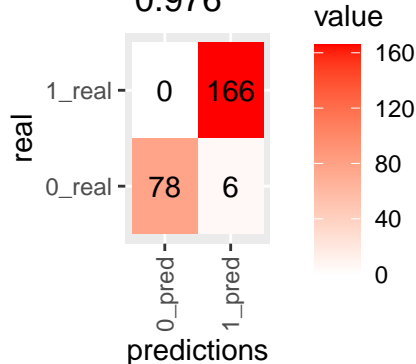




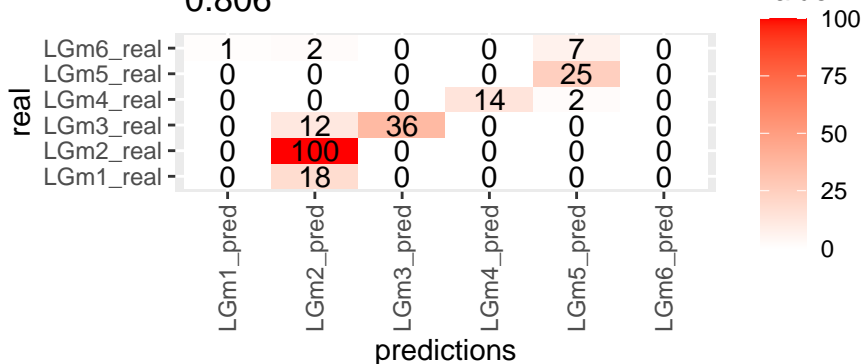
N.patients: 217–250



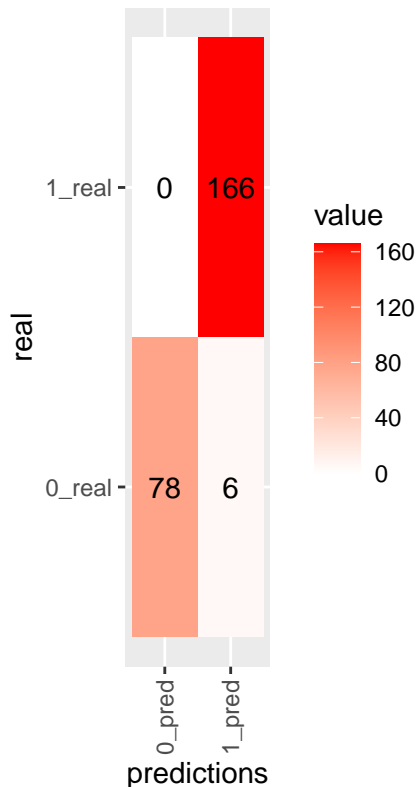
IDH model accuracy:  
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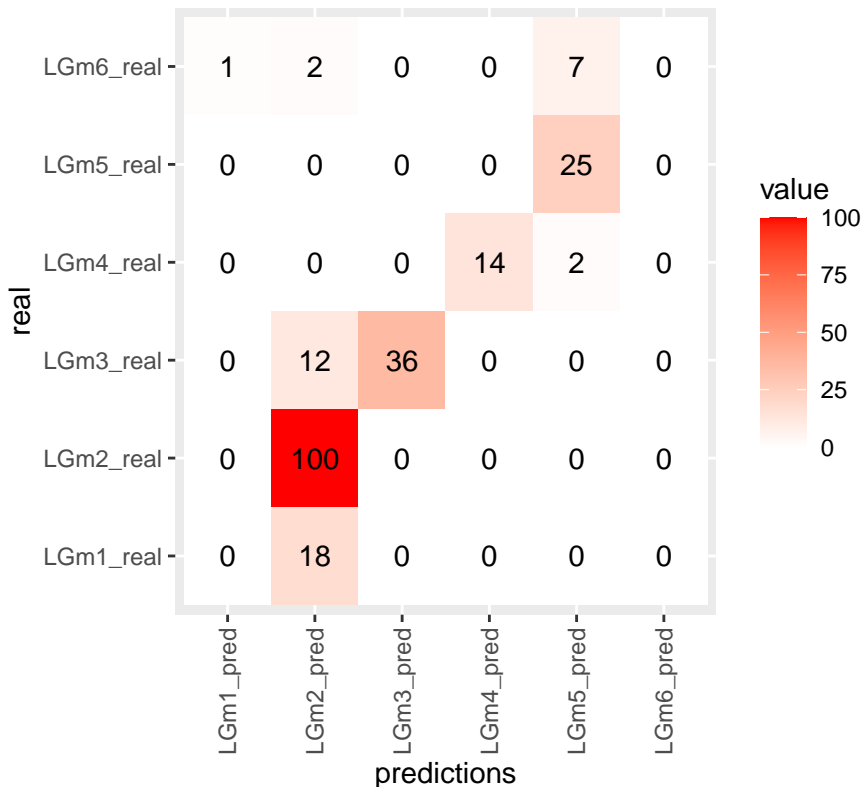
DNA methylation accuracy:  
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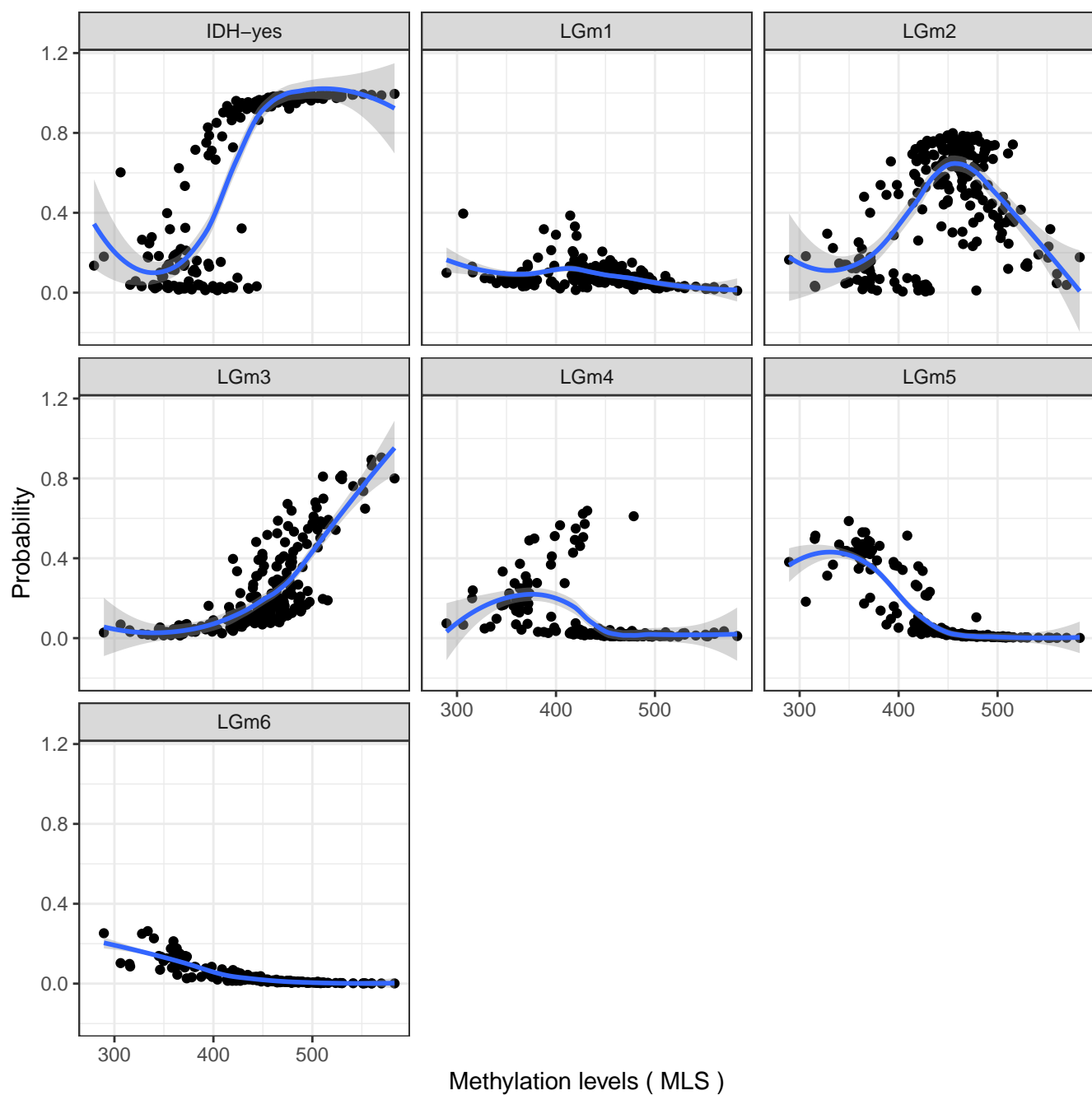


IDH model accuracy:  
0.976

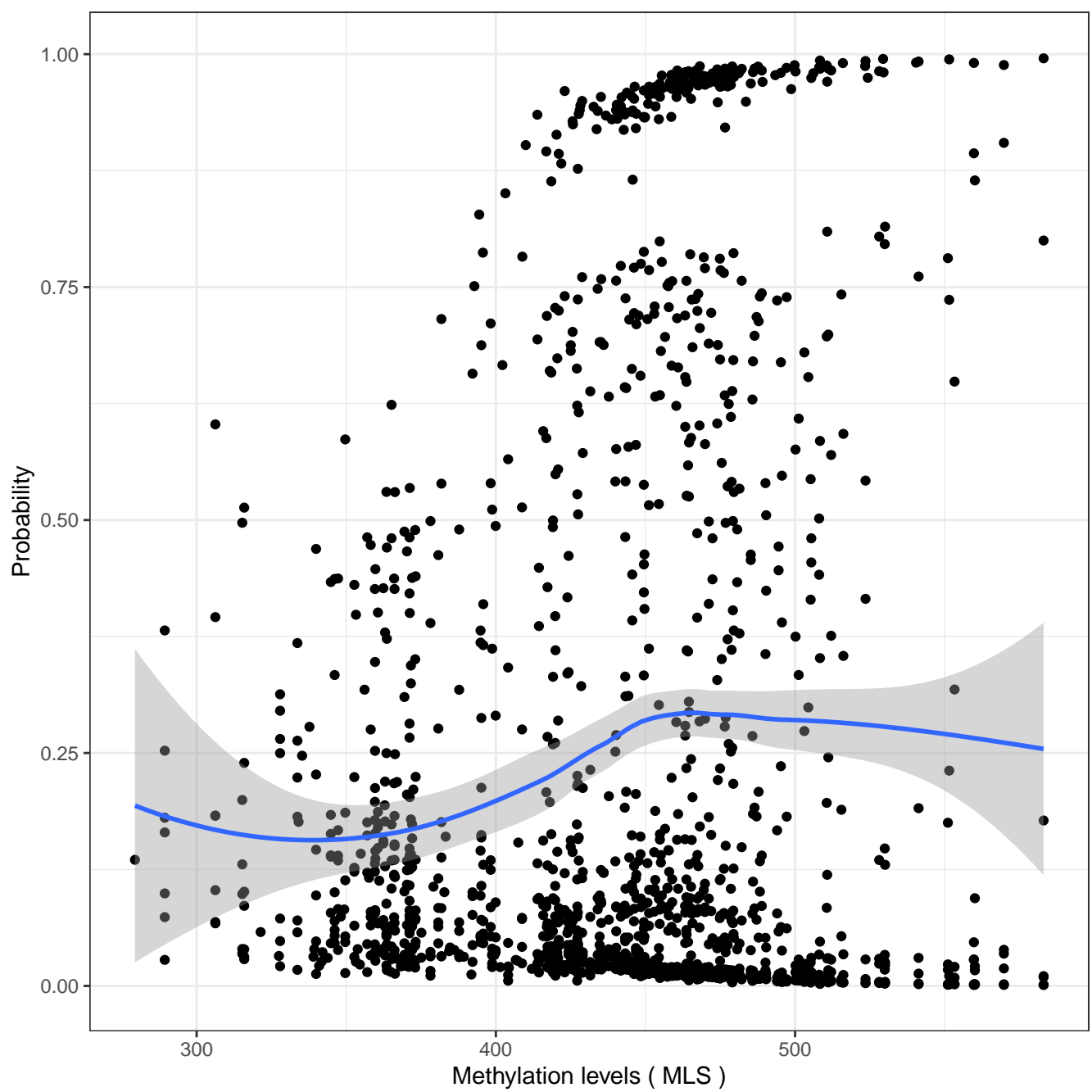


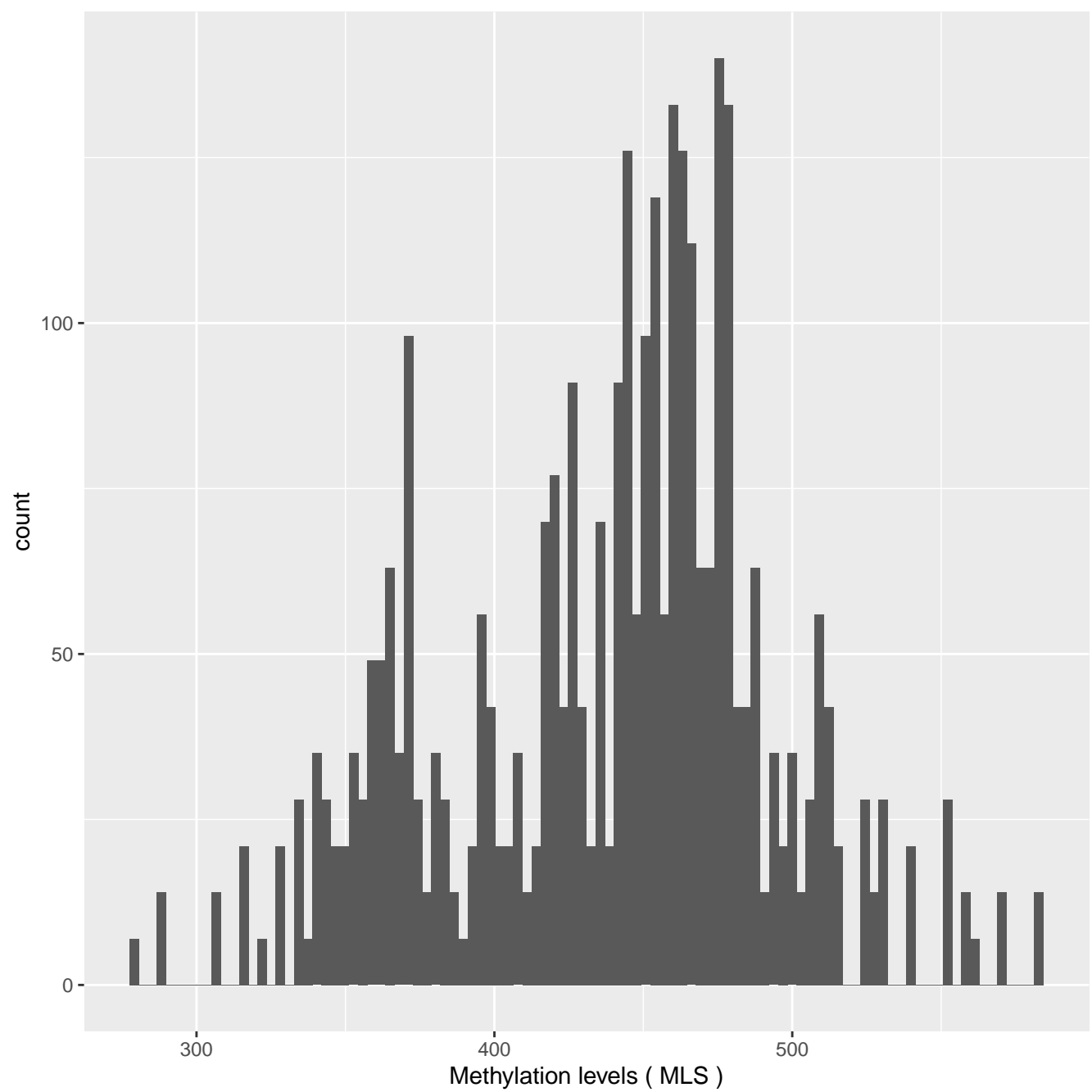
DNA methylation accuracy:  
0.806



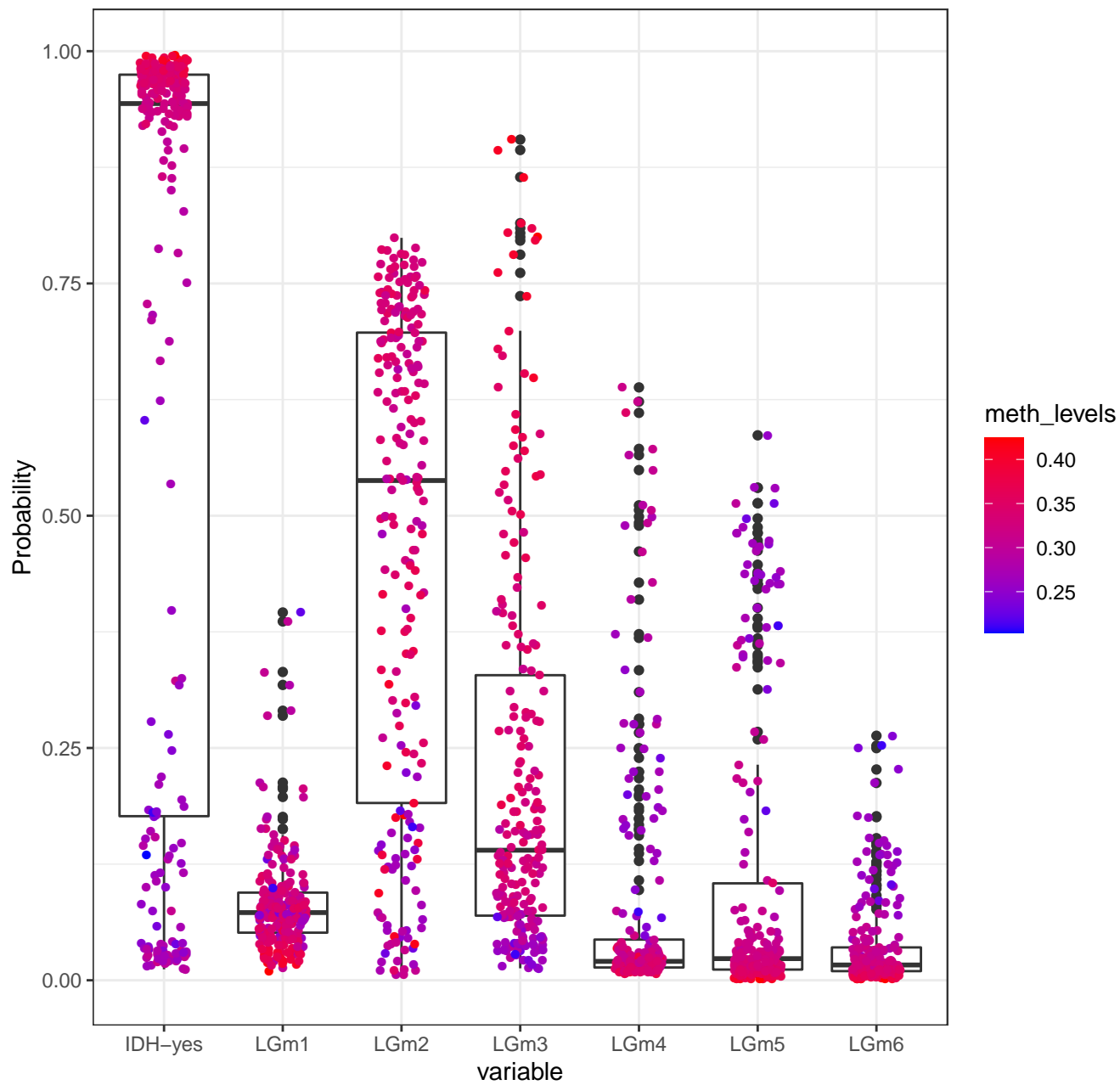




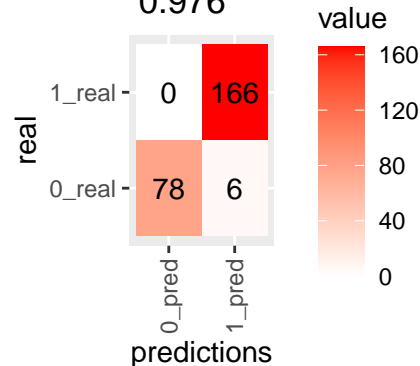




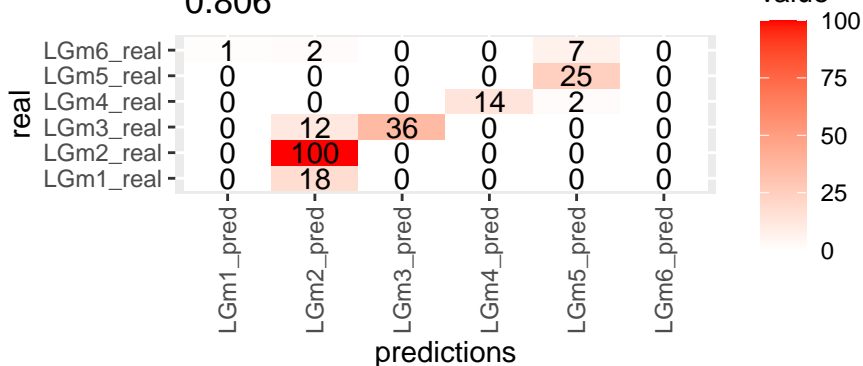
N.patients: 217-250



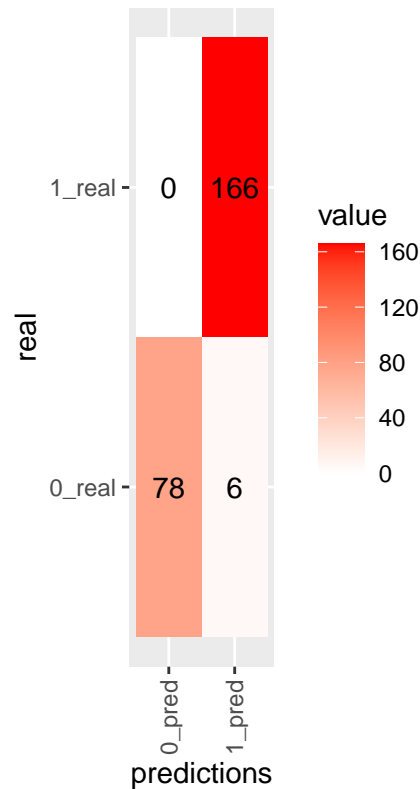
IDH model accuracy:  
0.976



DNA methylation accuracy:  
0.806



IDH model accuracy:  
0.976



DNA methylation accuracy:  
0.806

