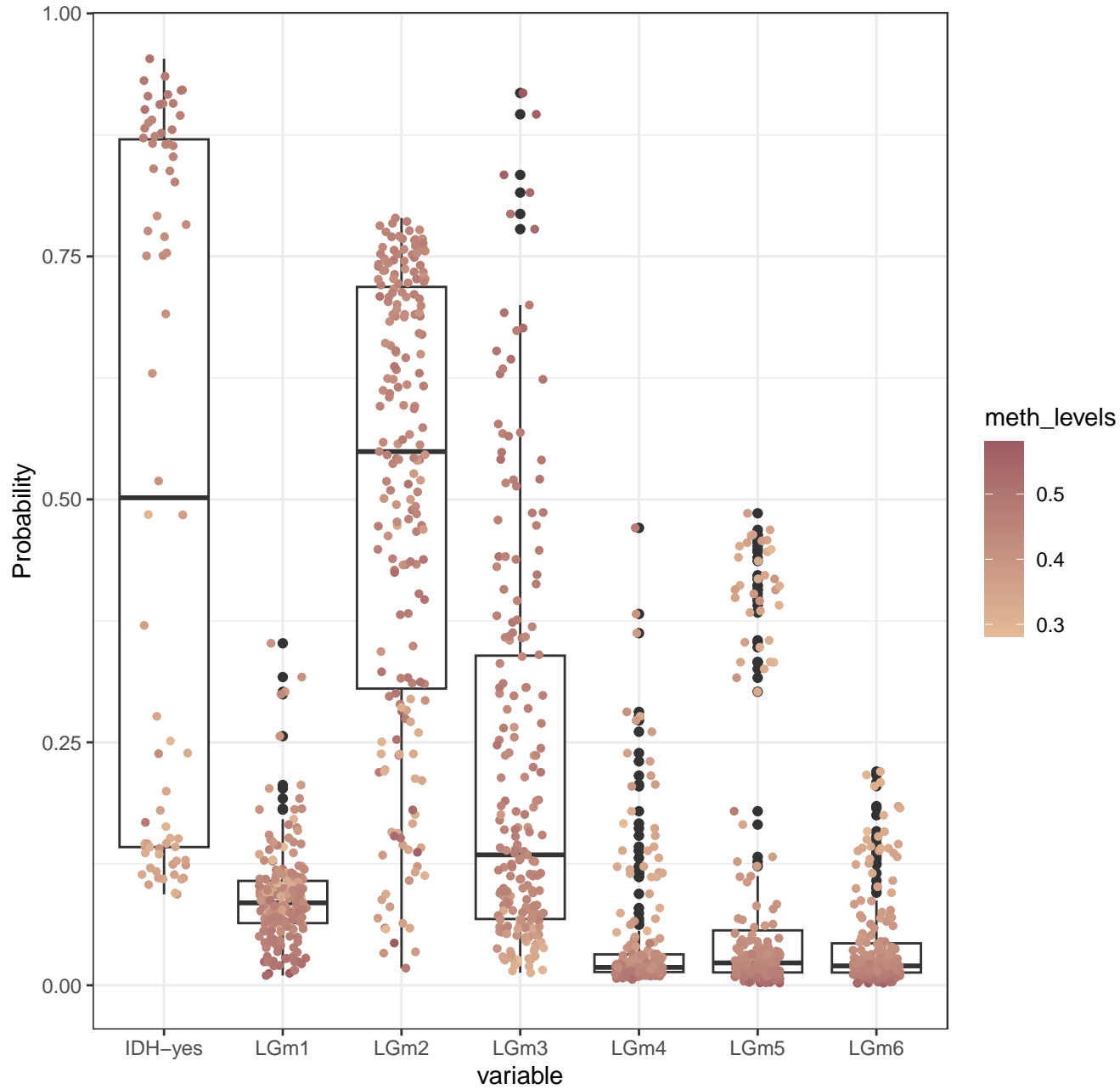
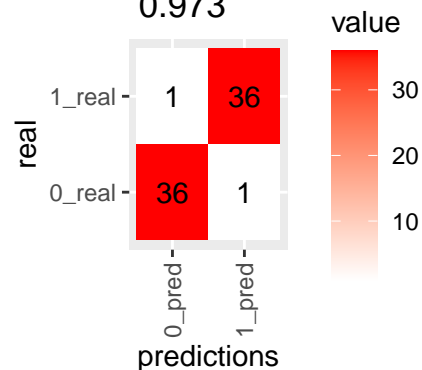


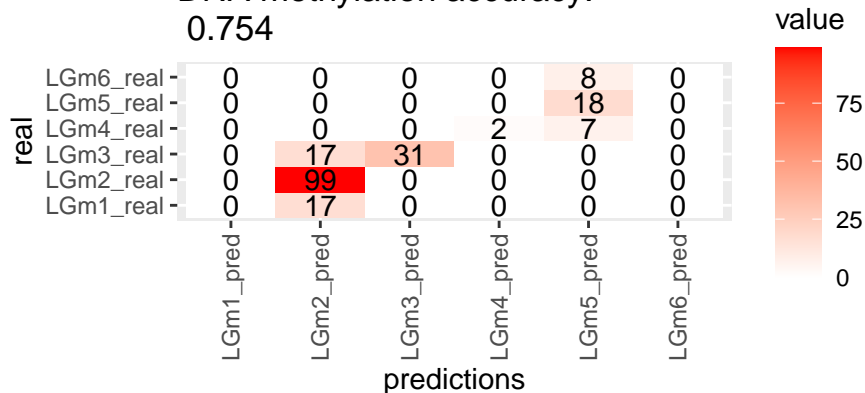
N.patients: 199-74



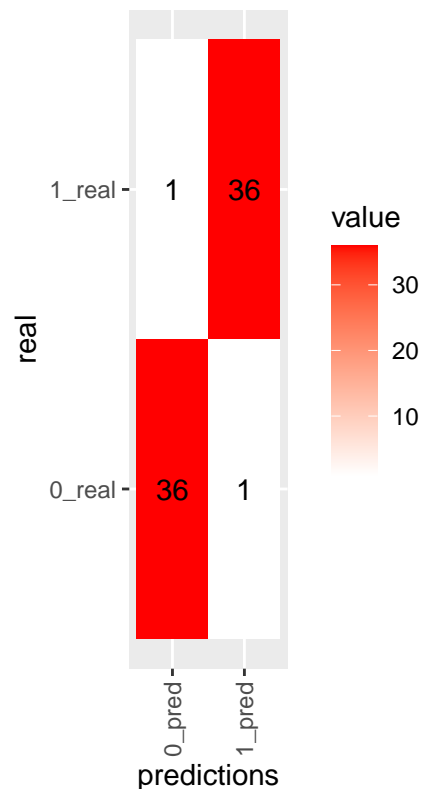
IDH model accuracy:
0.973



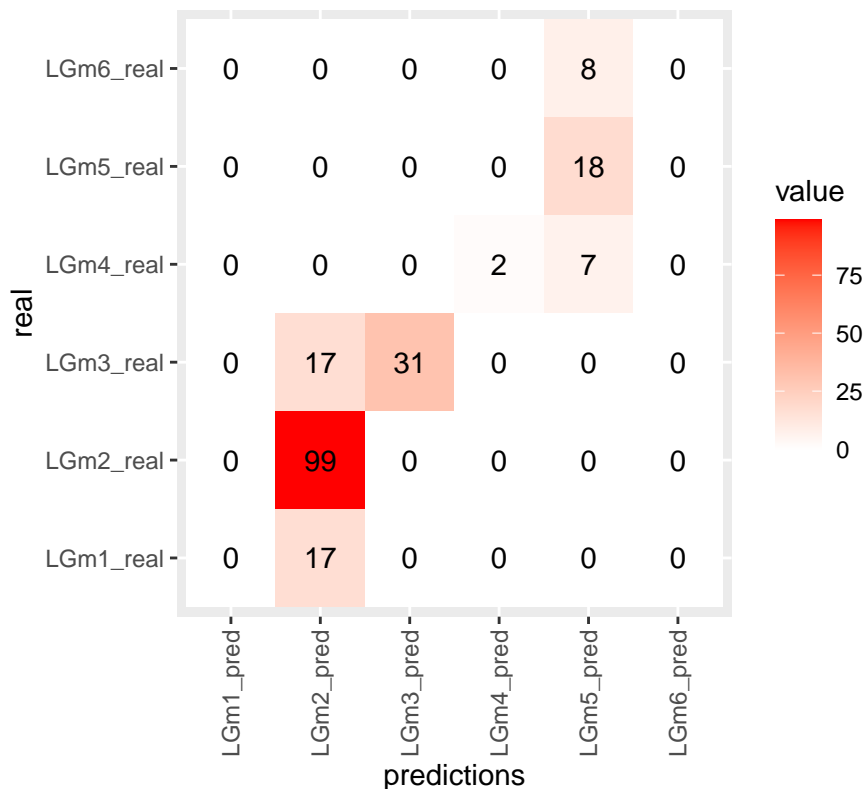
DNA methylation accuracy:
0.754

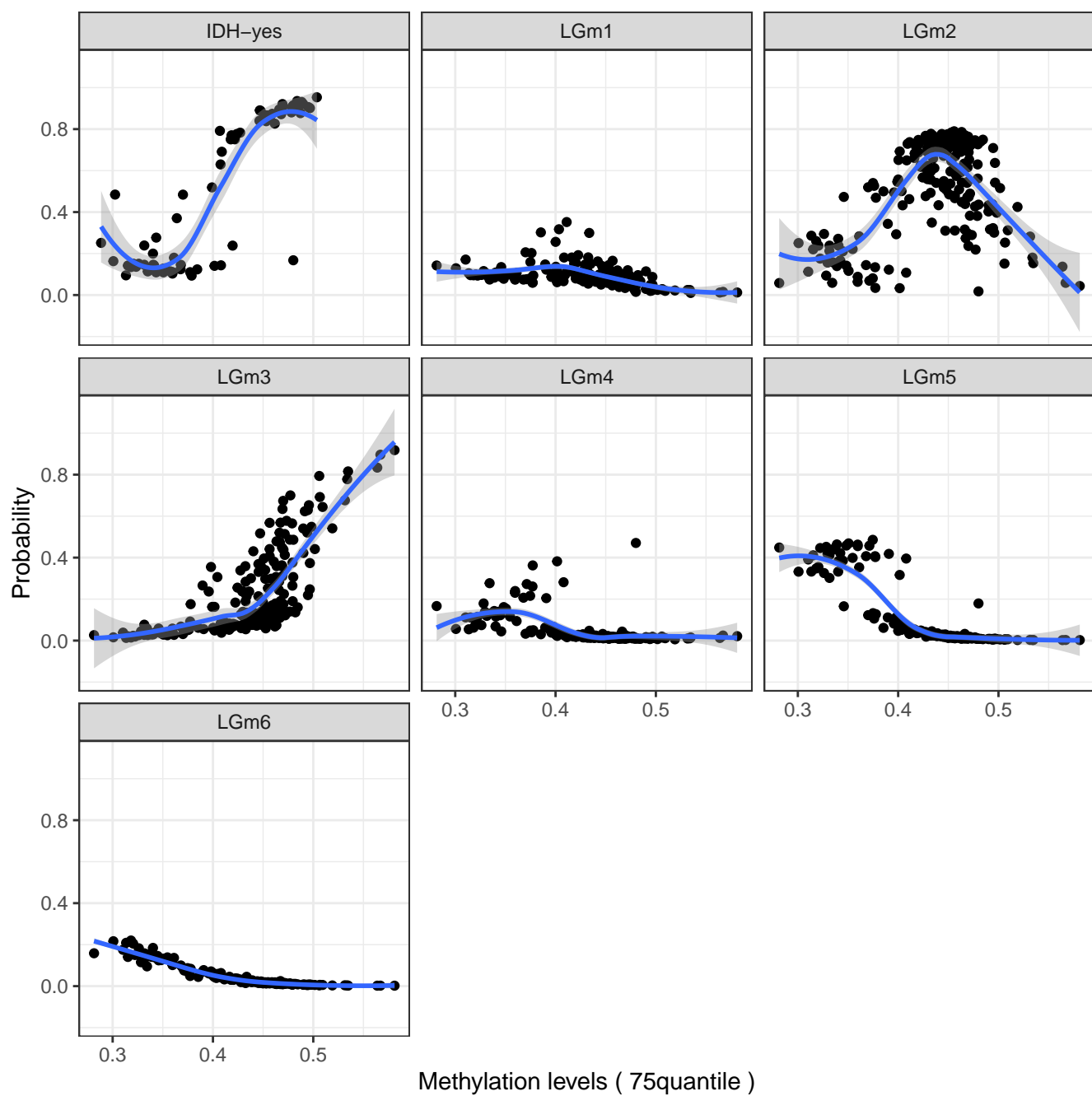


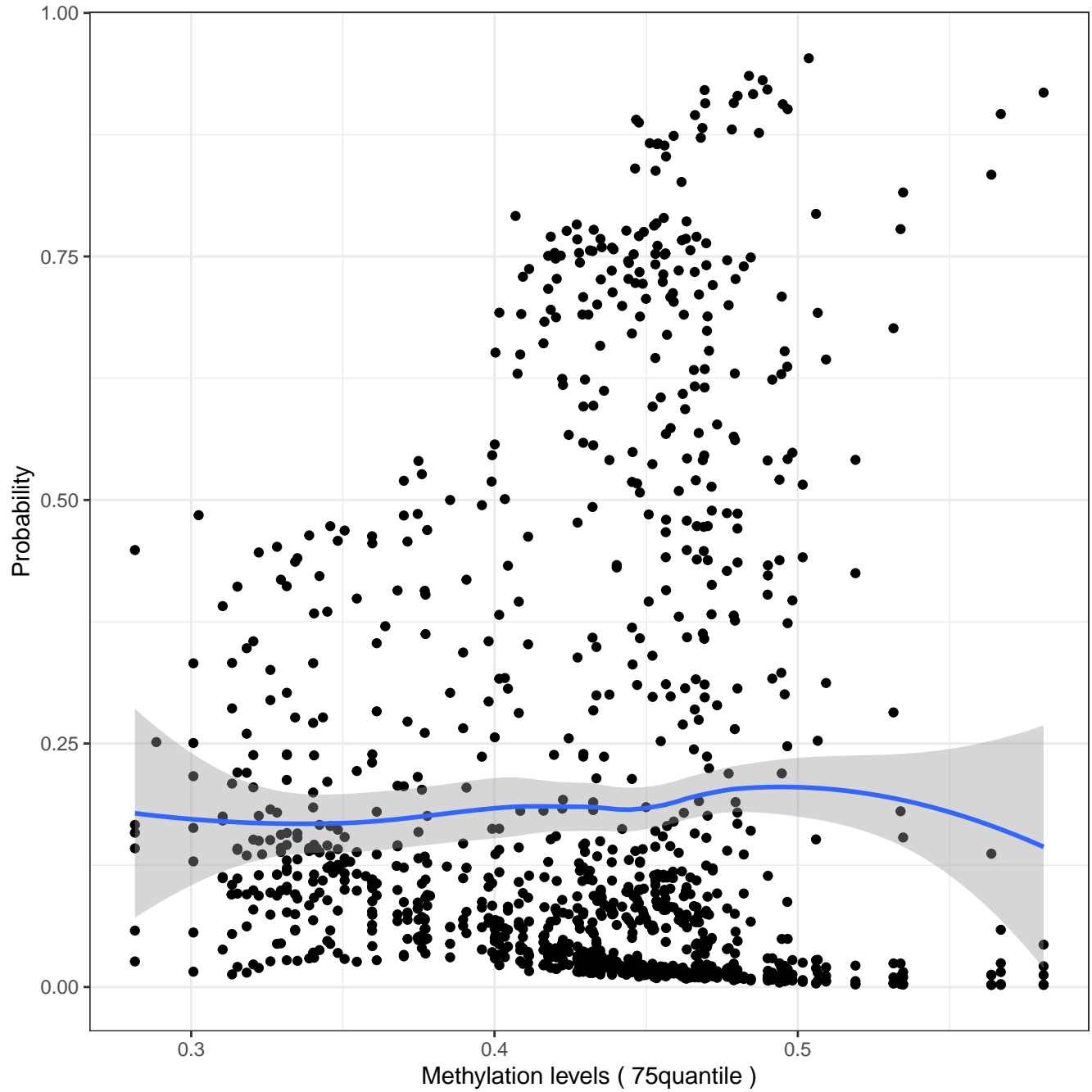
IDH model accuracy:
0.973

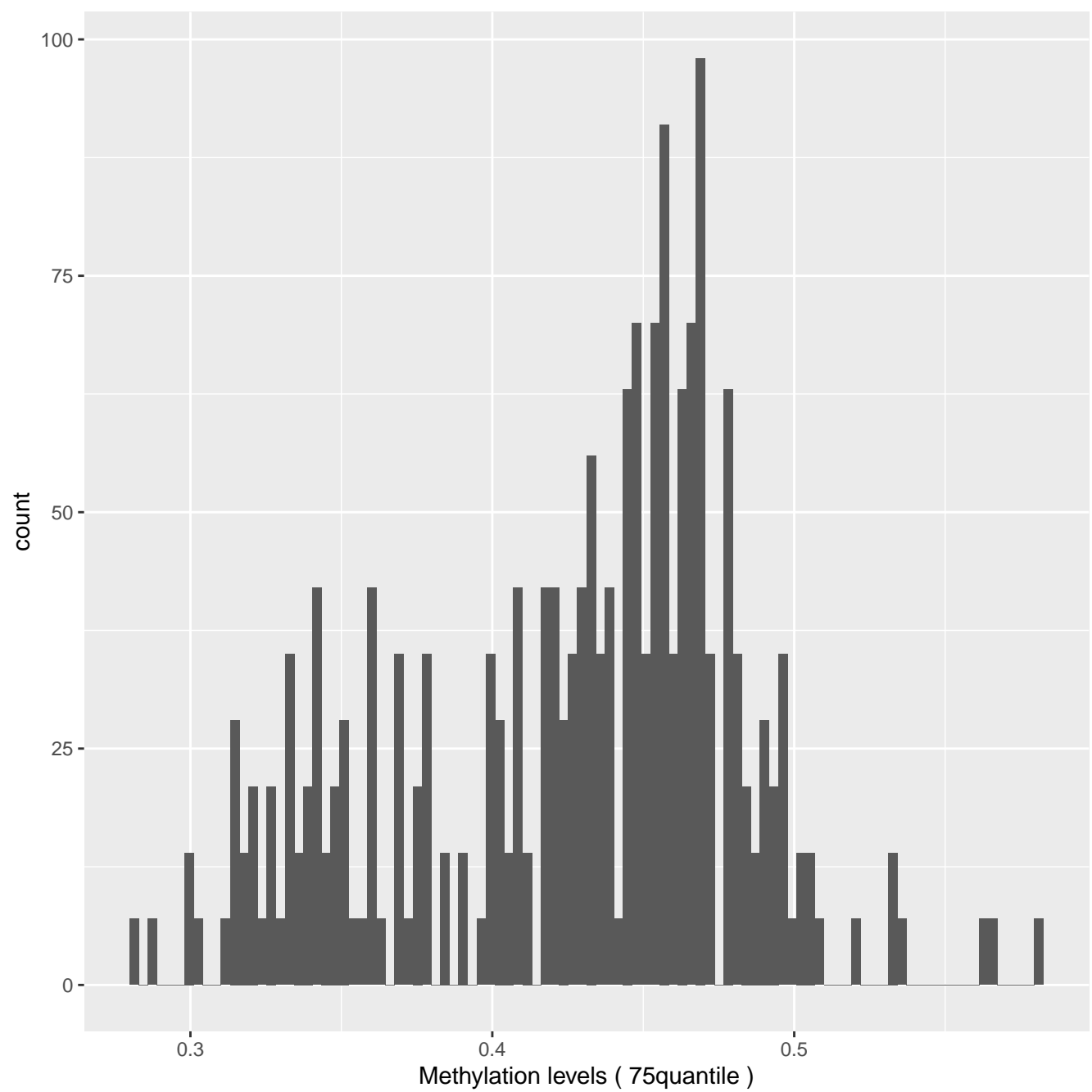


DNA methylation accuracy:
0.754

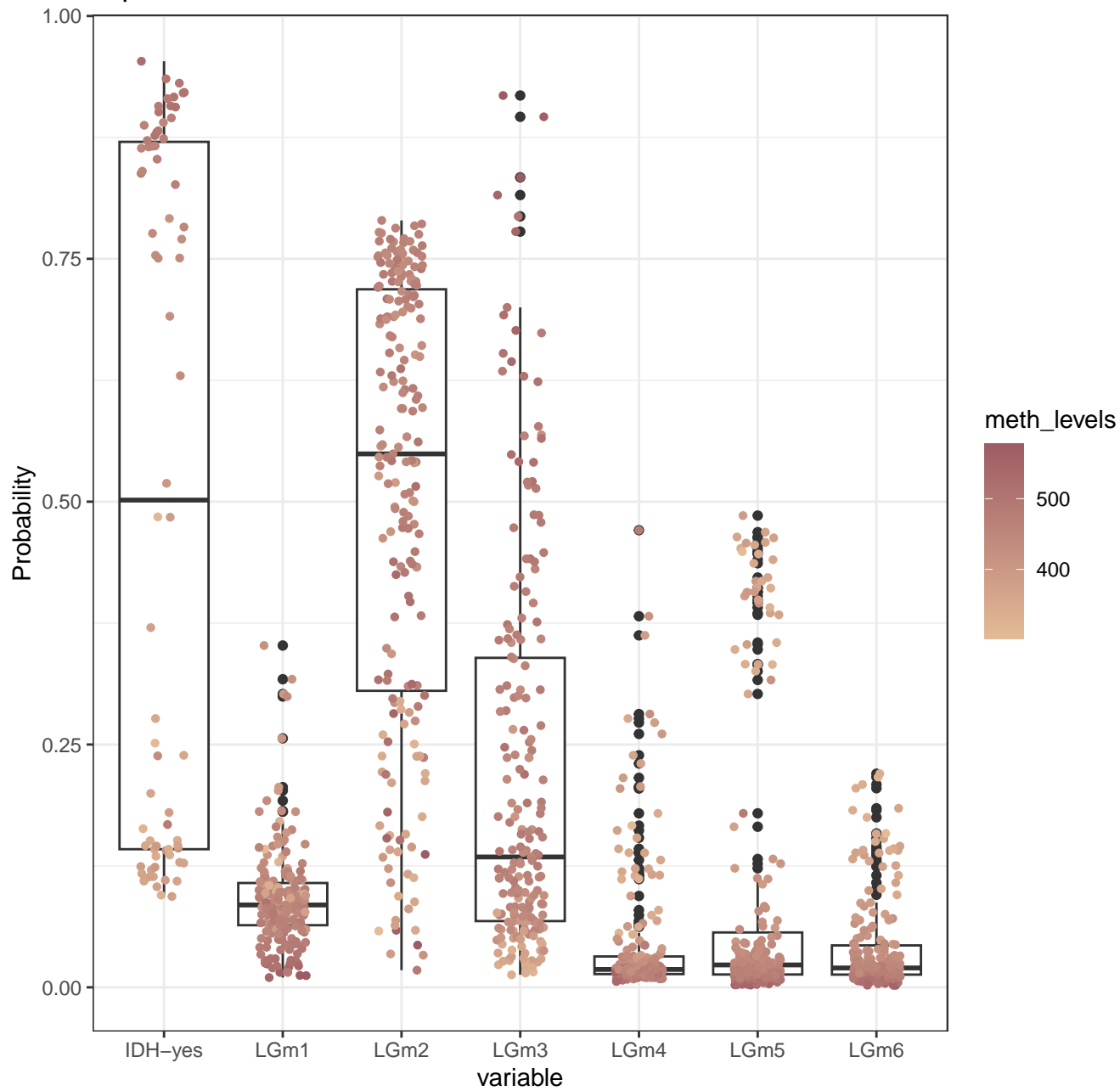




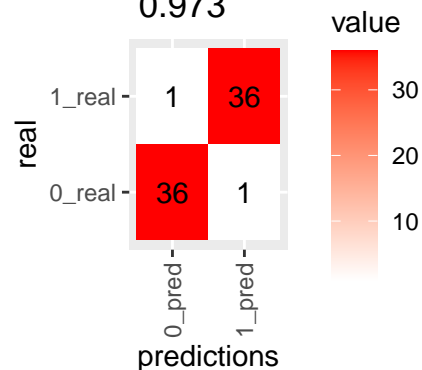




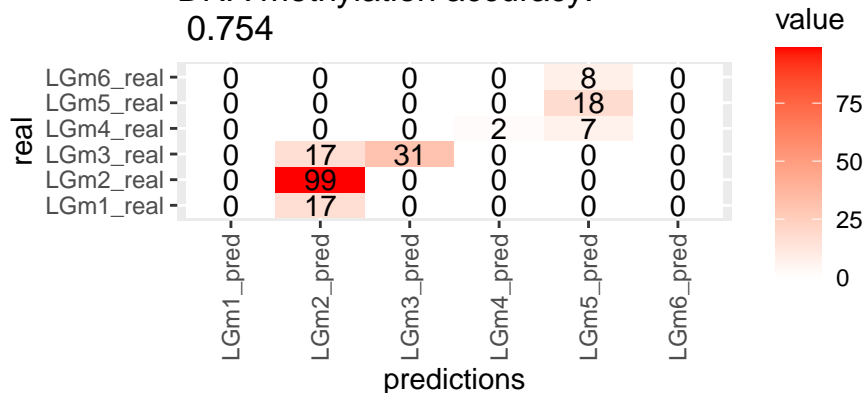
N.patients: 199-74



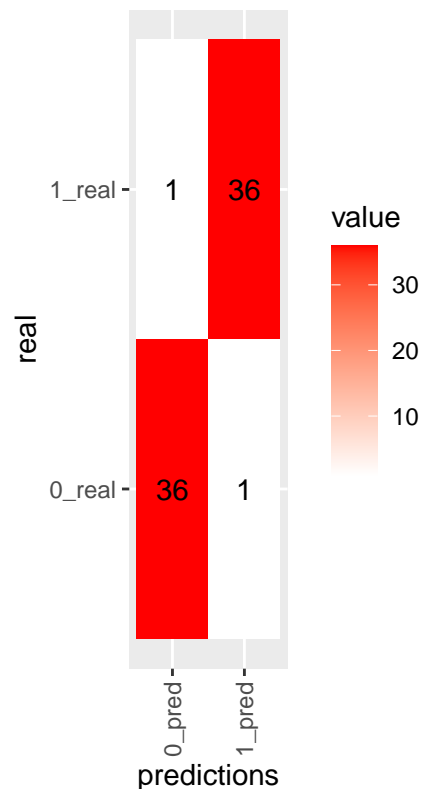
IDH model accuracy:
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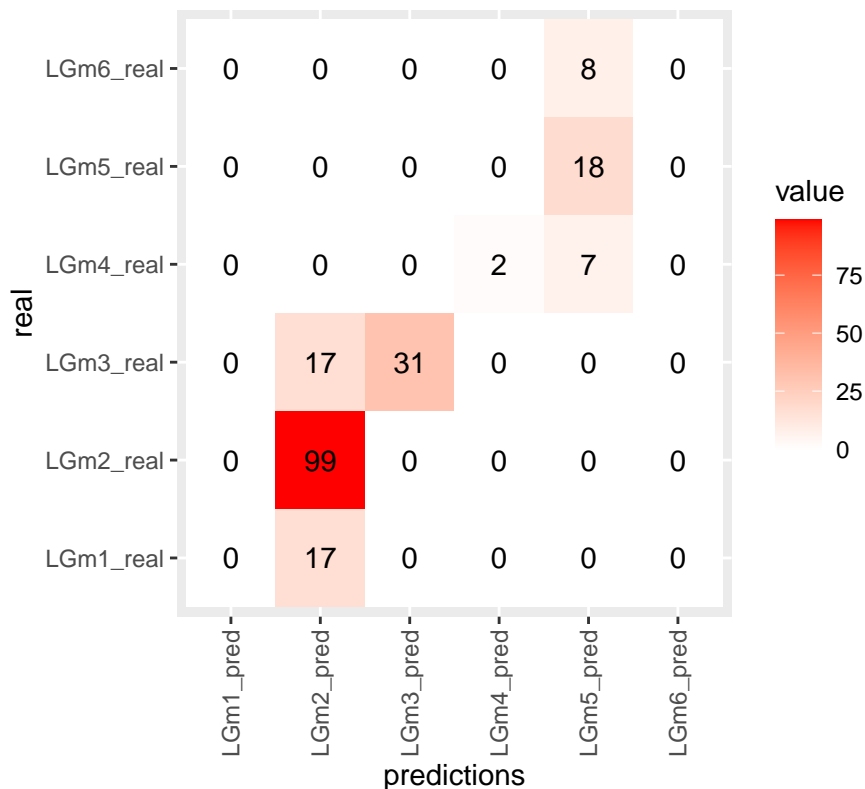
DNA methylation accuracy:
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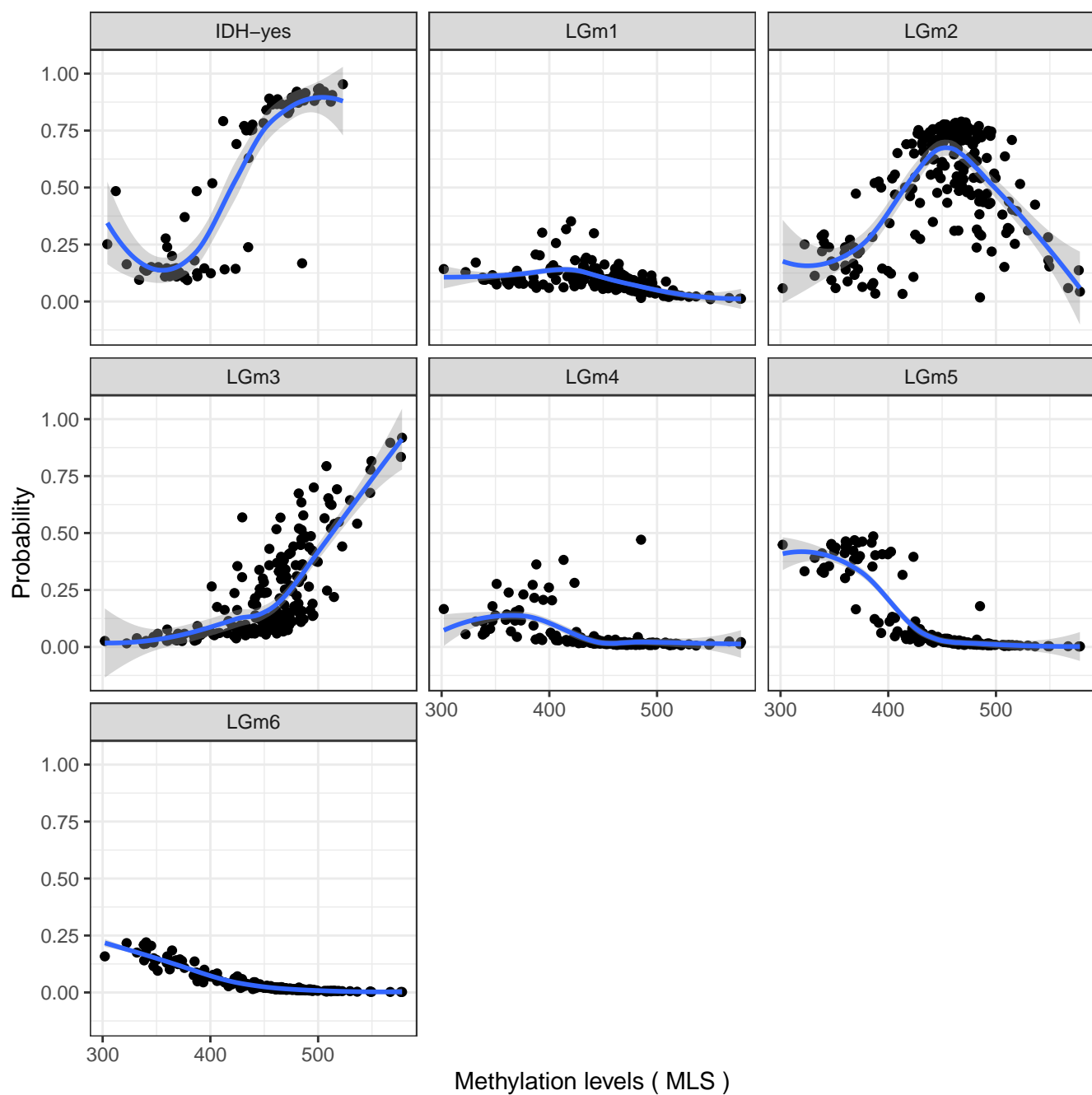


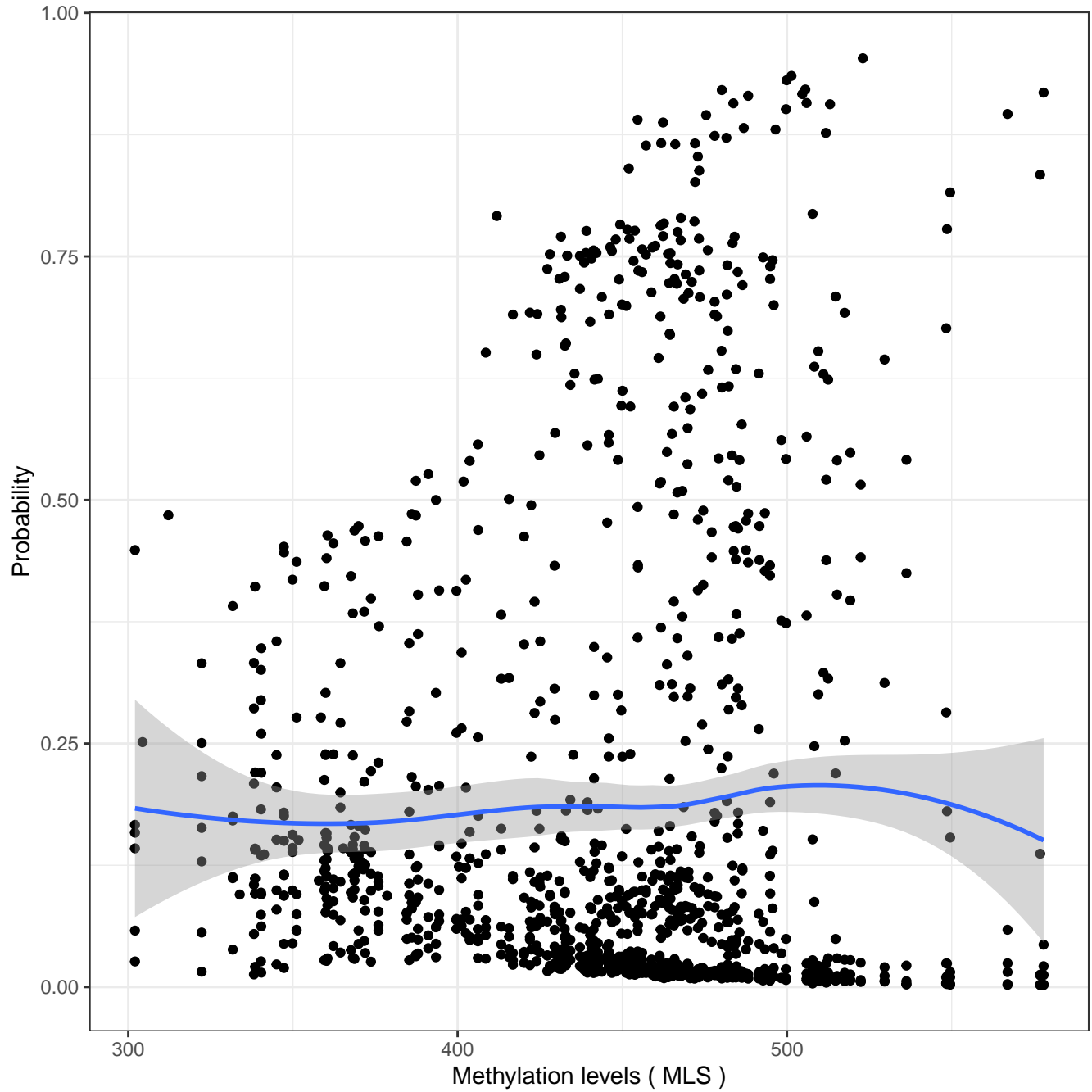
IDH model accuracy:
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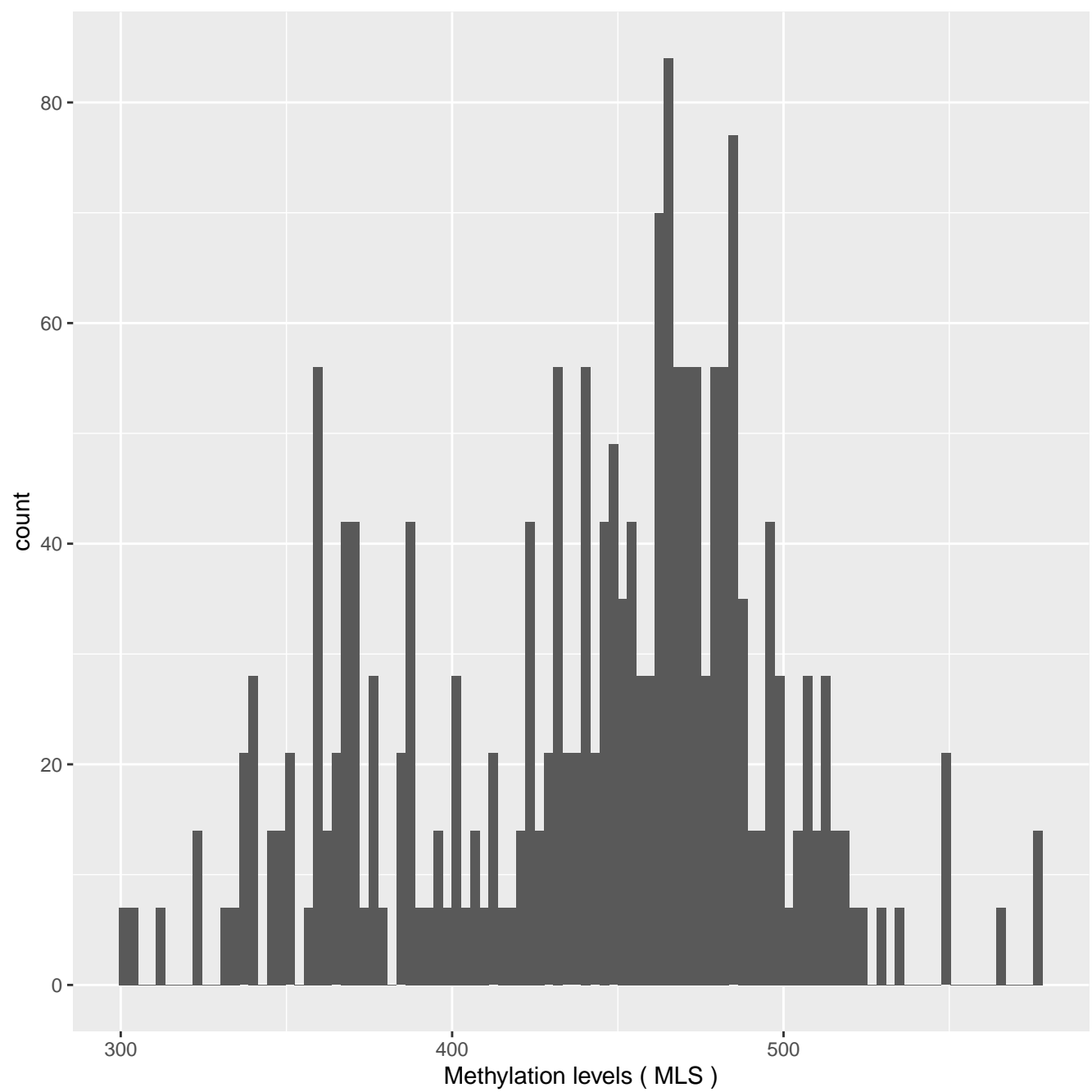


DNA methylation accuracy:
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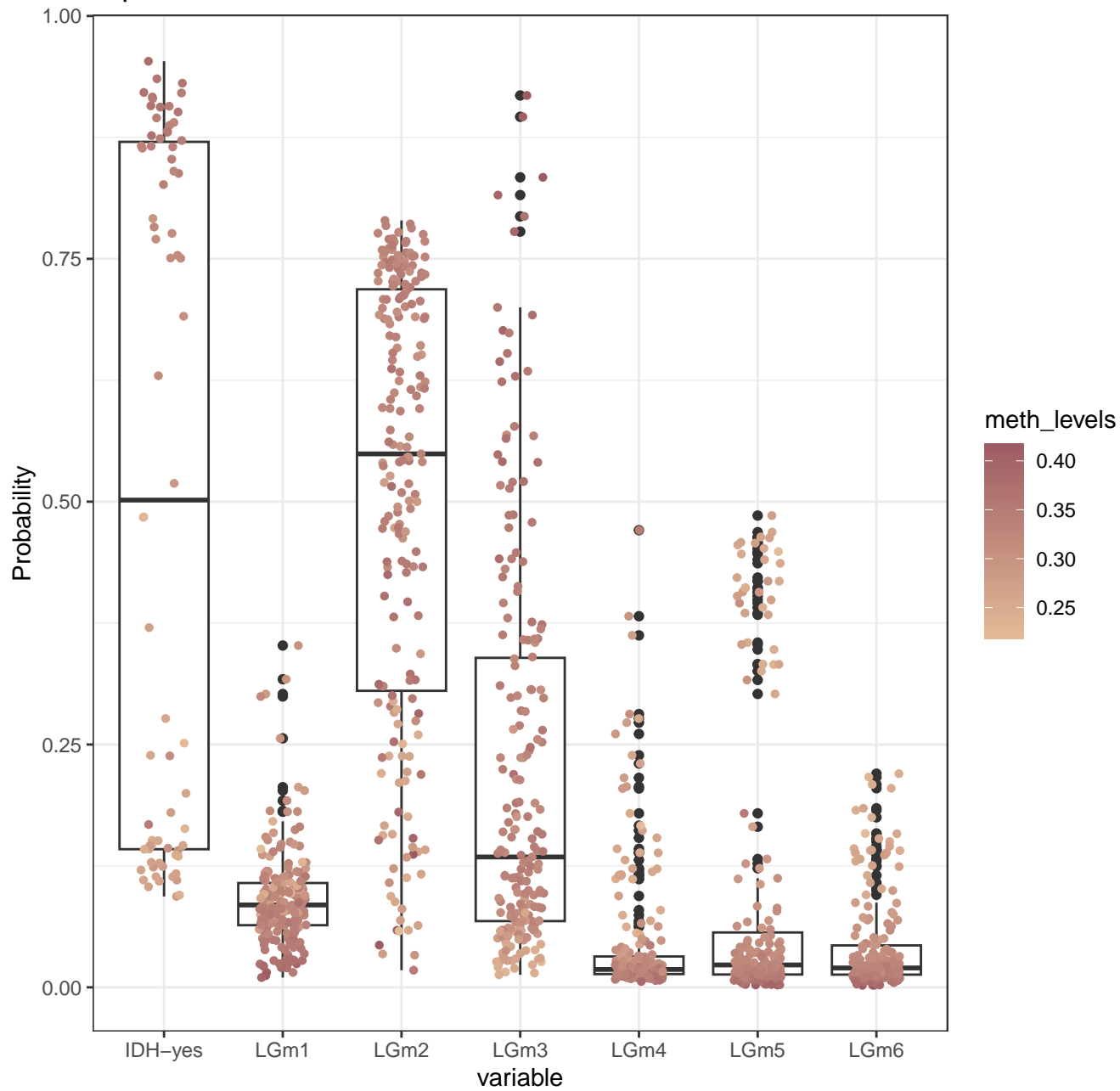




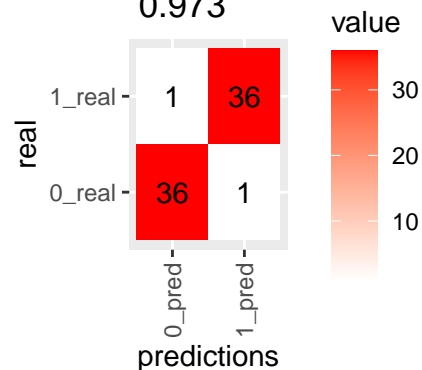




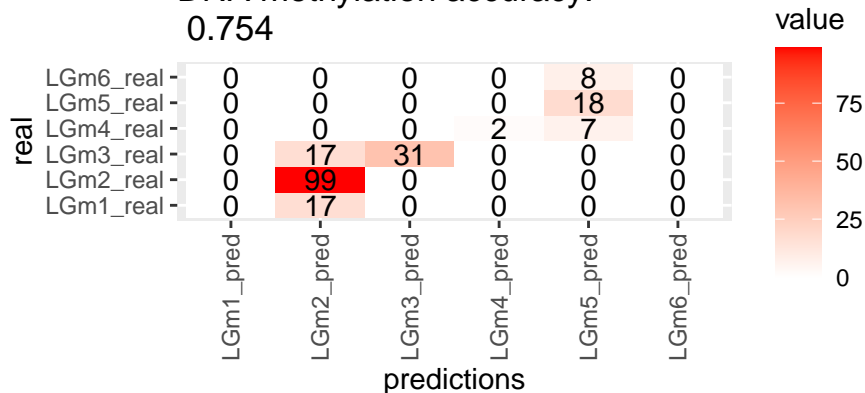
N.patients: 199-74



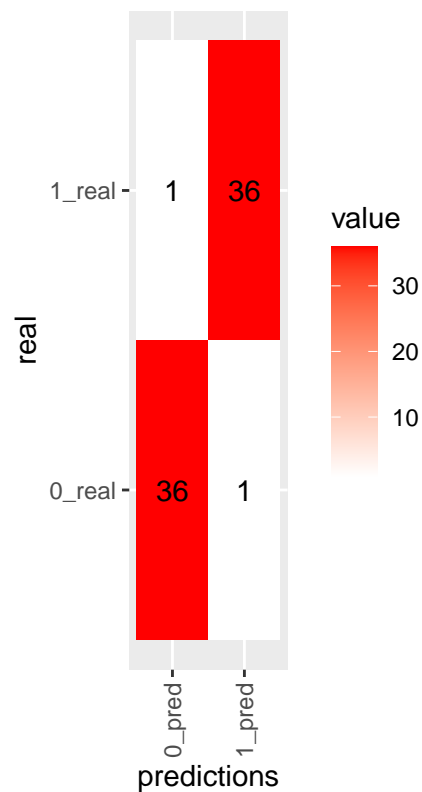
IDH model accuracy:
0.973



DNA methylation accuracy:
0.754



IDH model accuracy:
0.973



DNA methylation accuracy:
0.754

