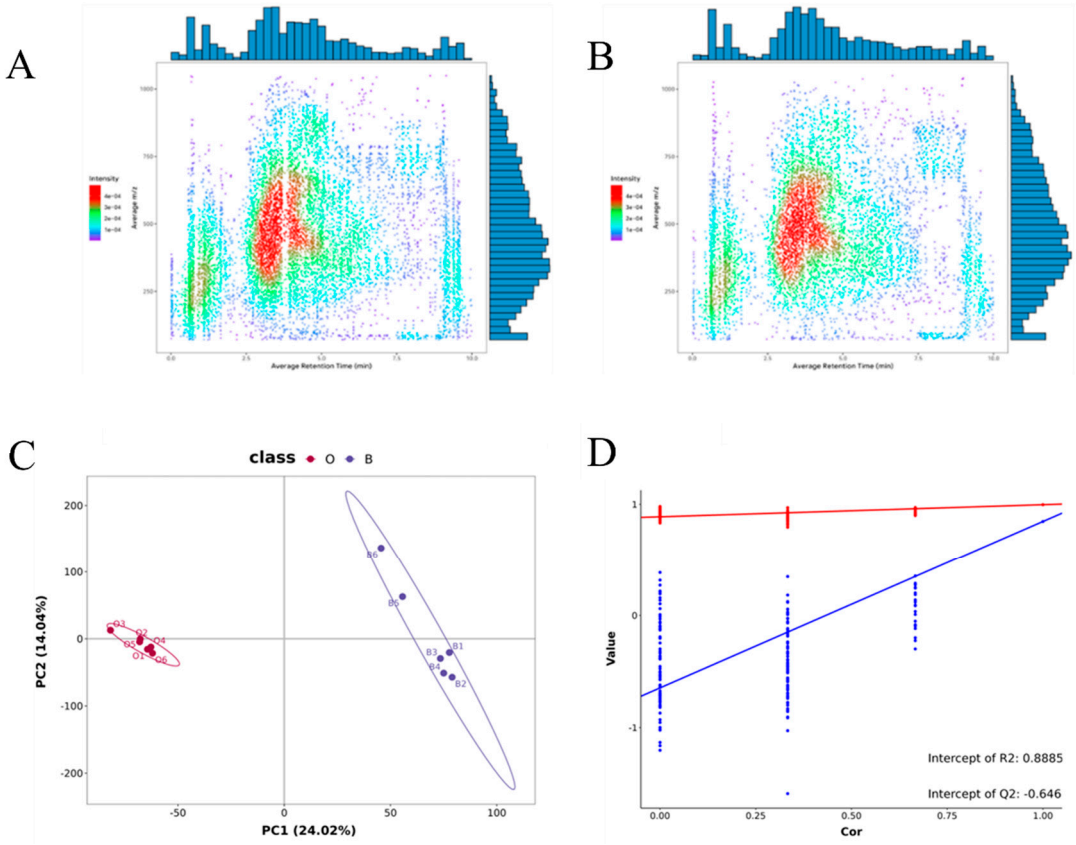


Figure S3 Validation of the Accuracy of Metabolome Sequencing Data from *ZmbHLH30* Overexpression Materials

(A) Negative ion mode,

(B) Positive ion mode,

(C)The explanation rate of the model in the X-axis direction,

(D)The prediction rate of the model.