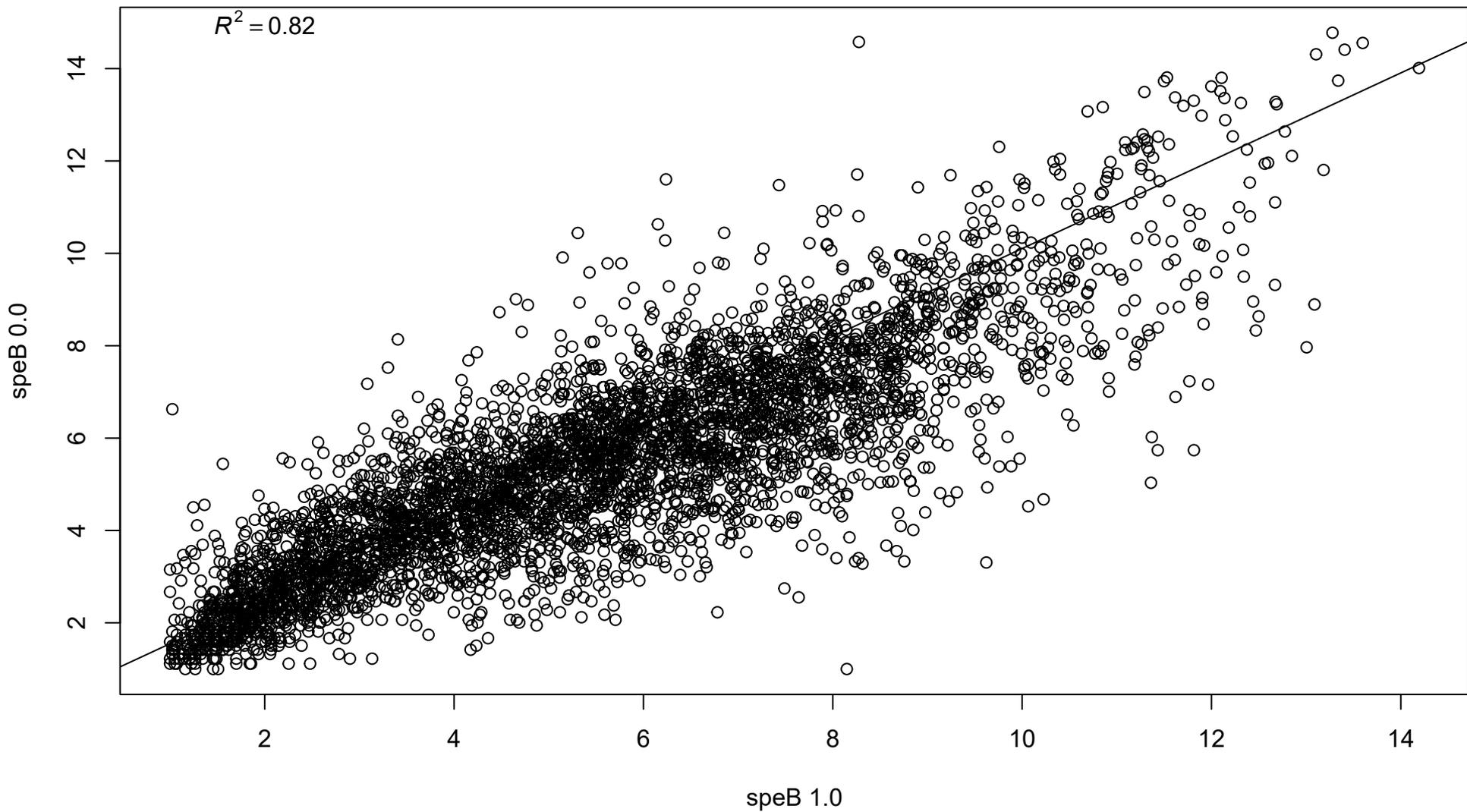
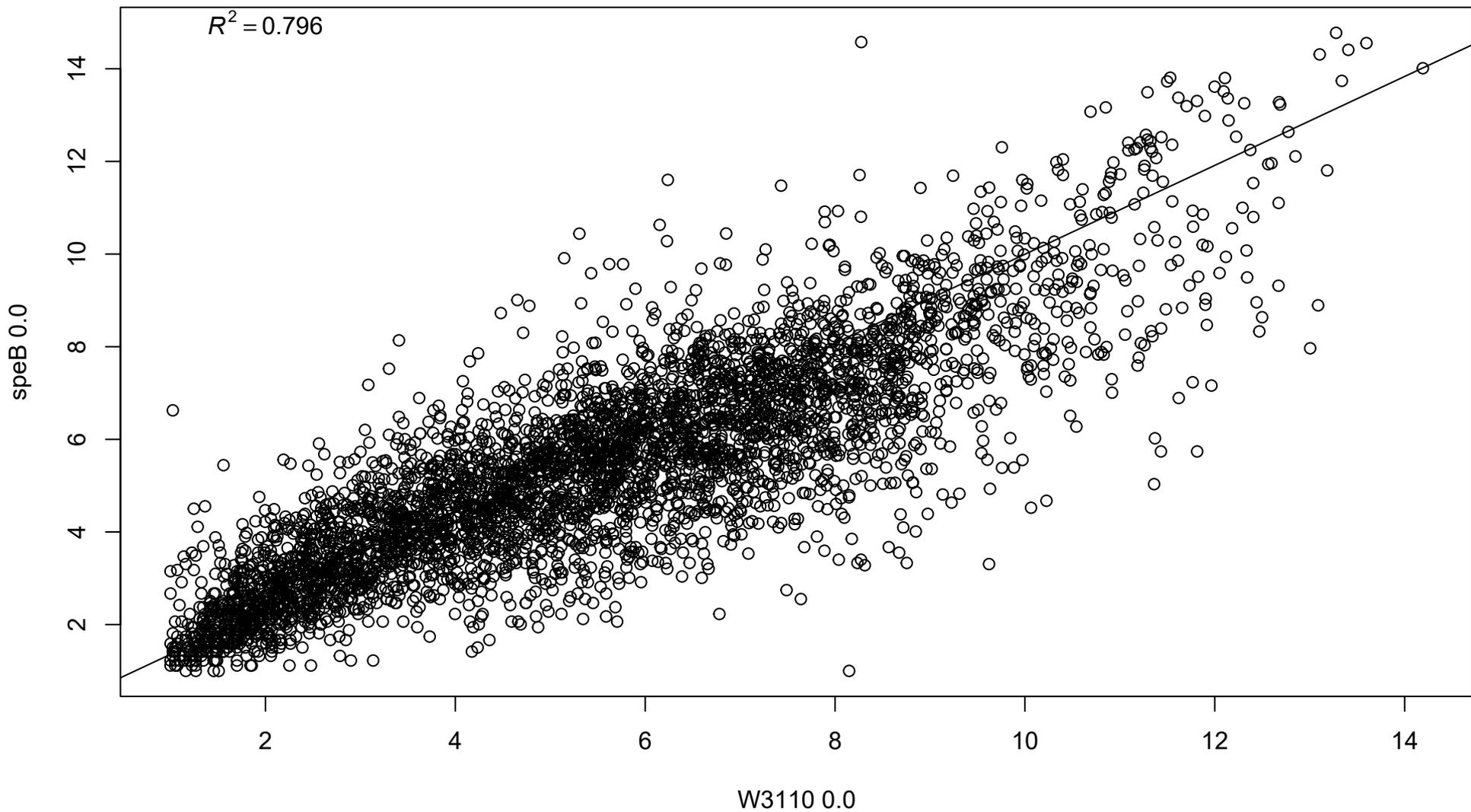


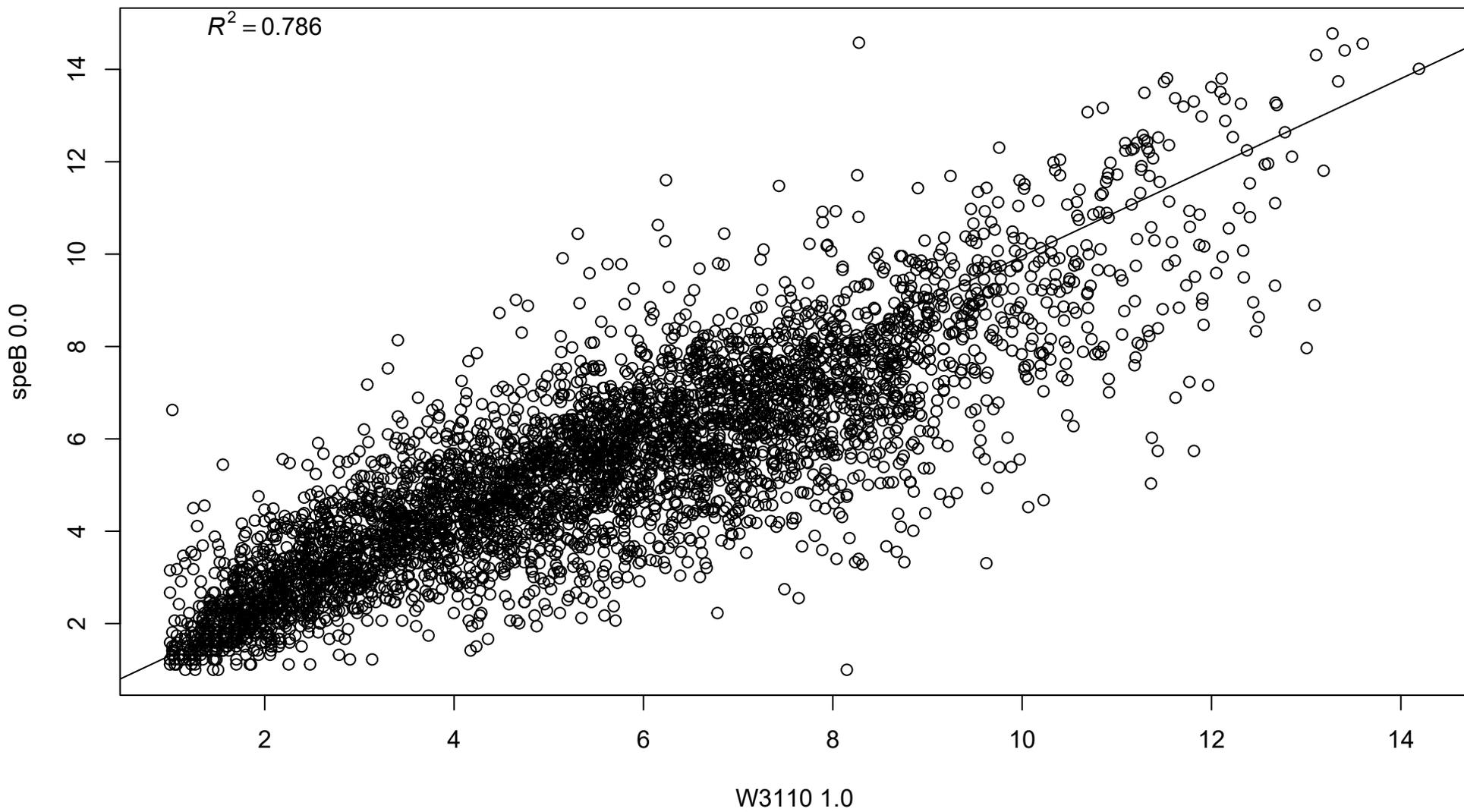
Comparison of the logCPM of the *speB* mutant with and without 1.0mM putrescine



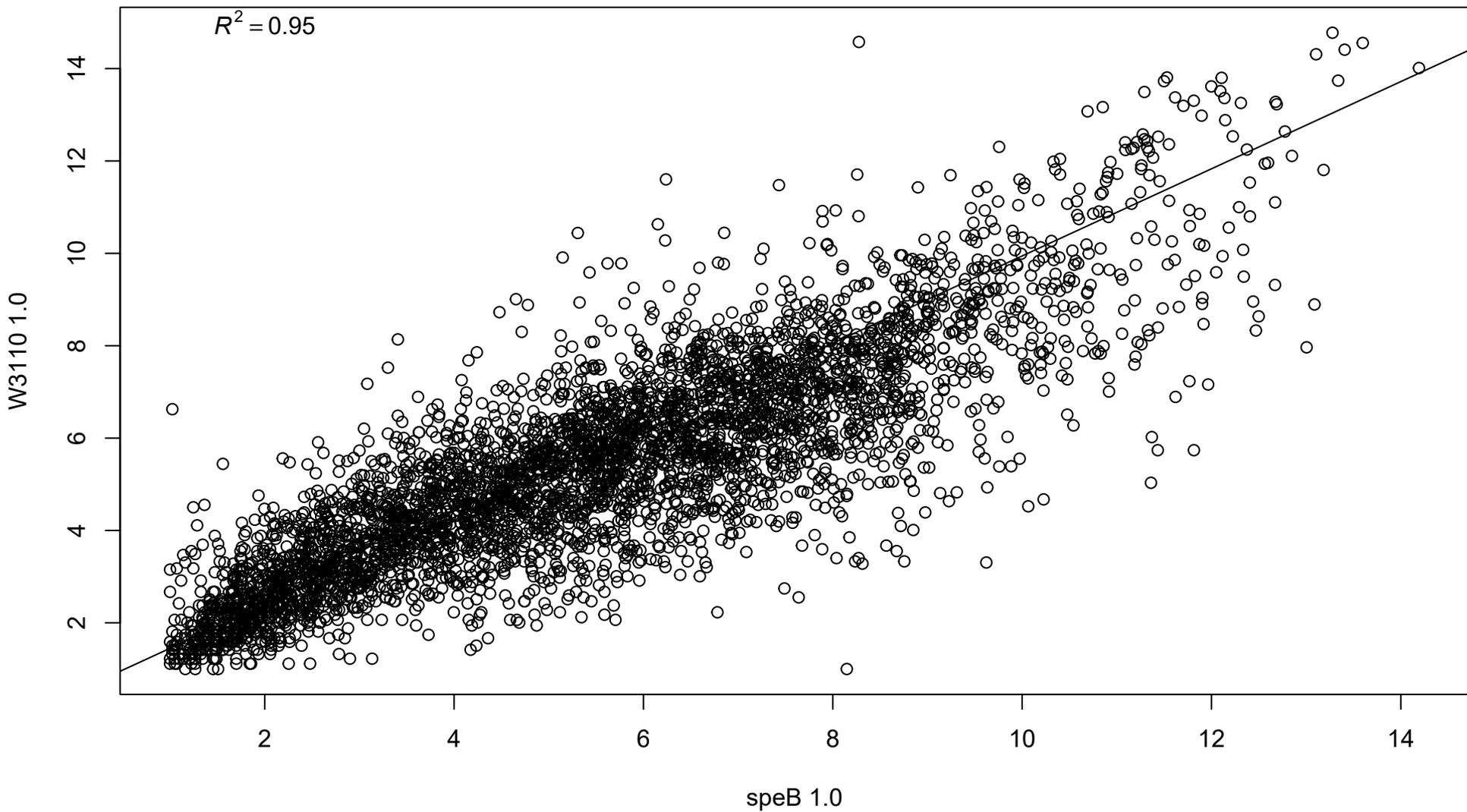
Comparison of the logCPM of W3110 and a speB mutant without putrescine



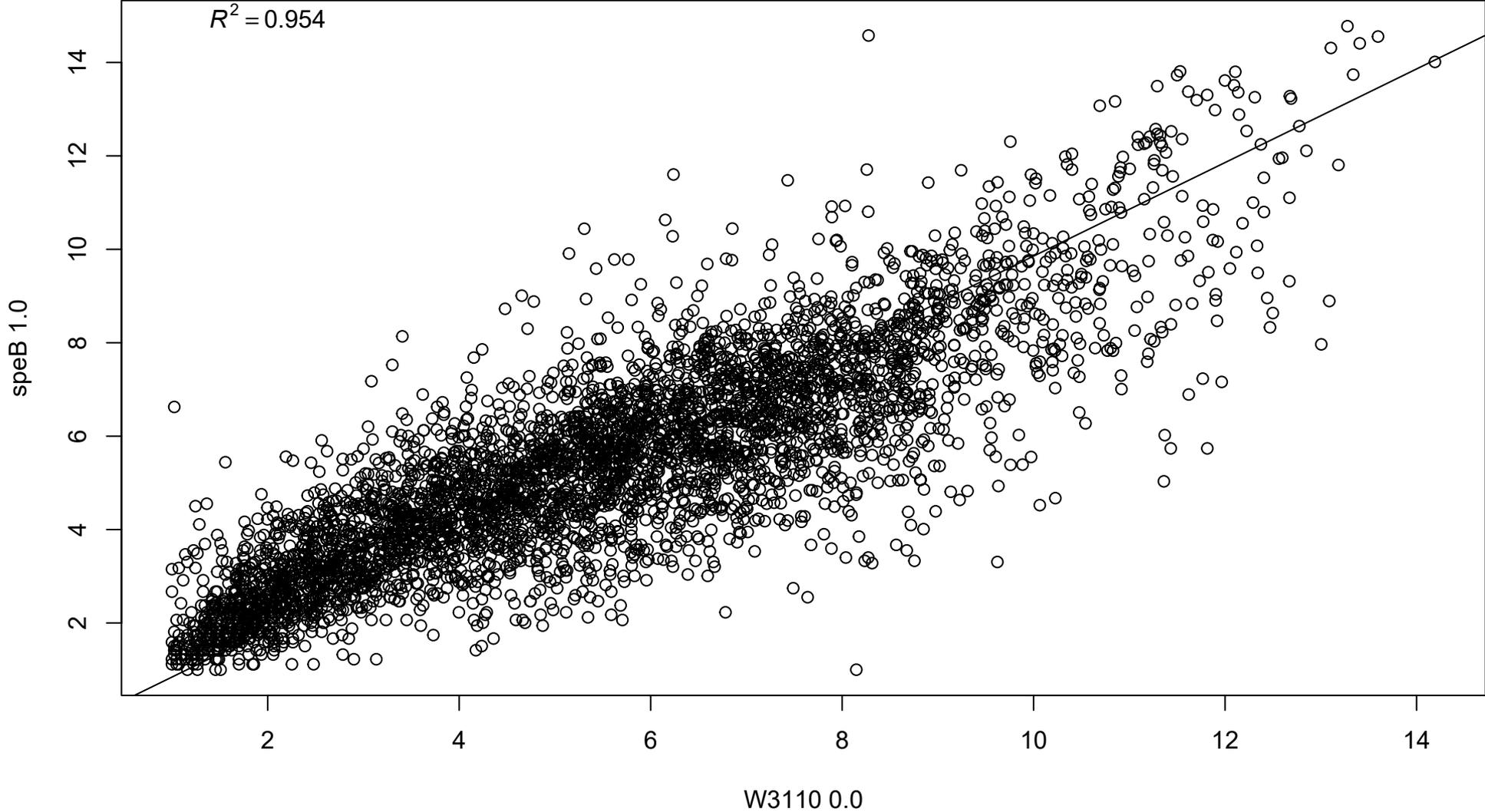
Comparison of the logCPM of W3110 with 1.0mM putrescine and a speB mutant without putrescine



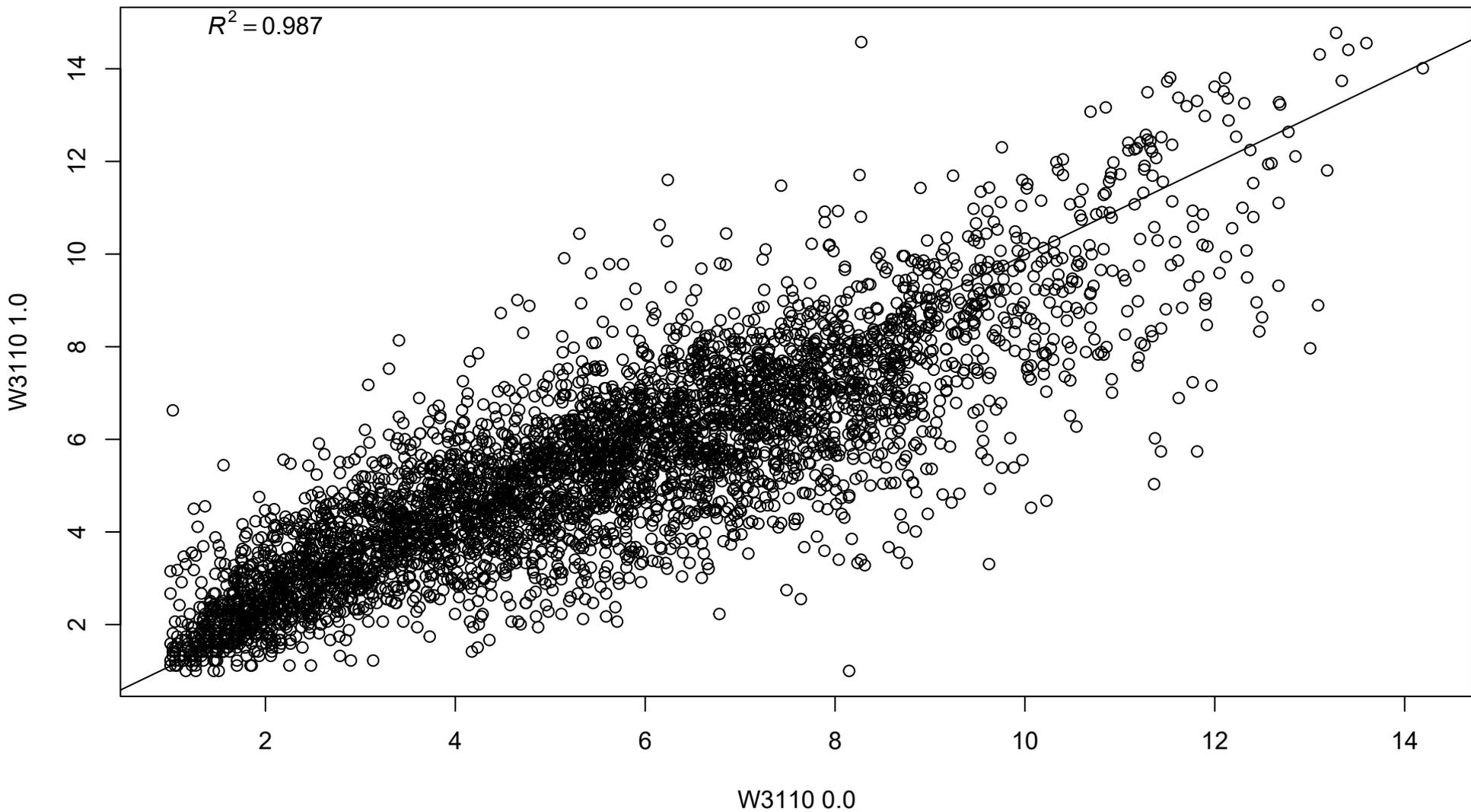
Comparison of the logCPM of W3110 and a speB mutant with 1.0mM putrescine



Comparison of the logCPM of W3110 without putrescine and a speB mutant with 1.0mM putrescine



Comparison of the logCPM of W3110 with and without 1.0mM putrescine



Supplementary File 1: Expression graphs comparing the logCPM of the transcriptomes of the average of the three replicates of the W3110 with and without 1 mM putrescine and the *speB* mutant with and without 1 mM putrescine. A regression line was calculated and the correlation between each set of transcriptomes noted on the graph. Higher R^2 values indicate greater similarity between the transcriptomes.