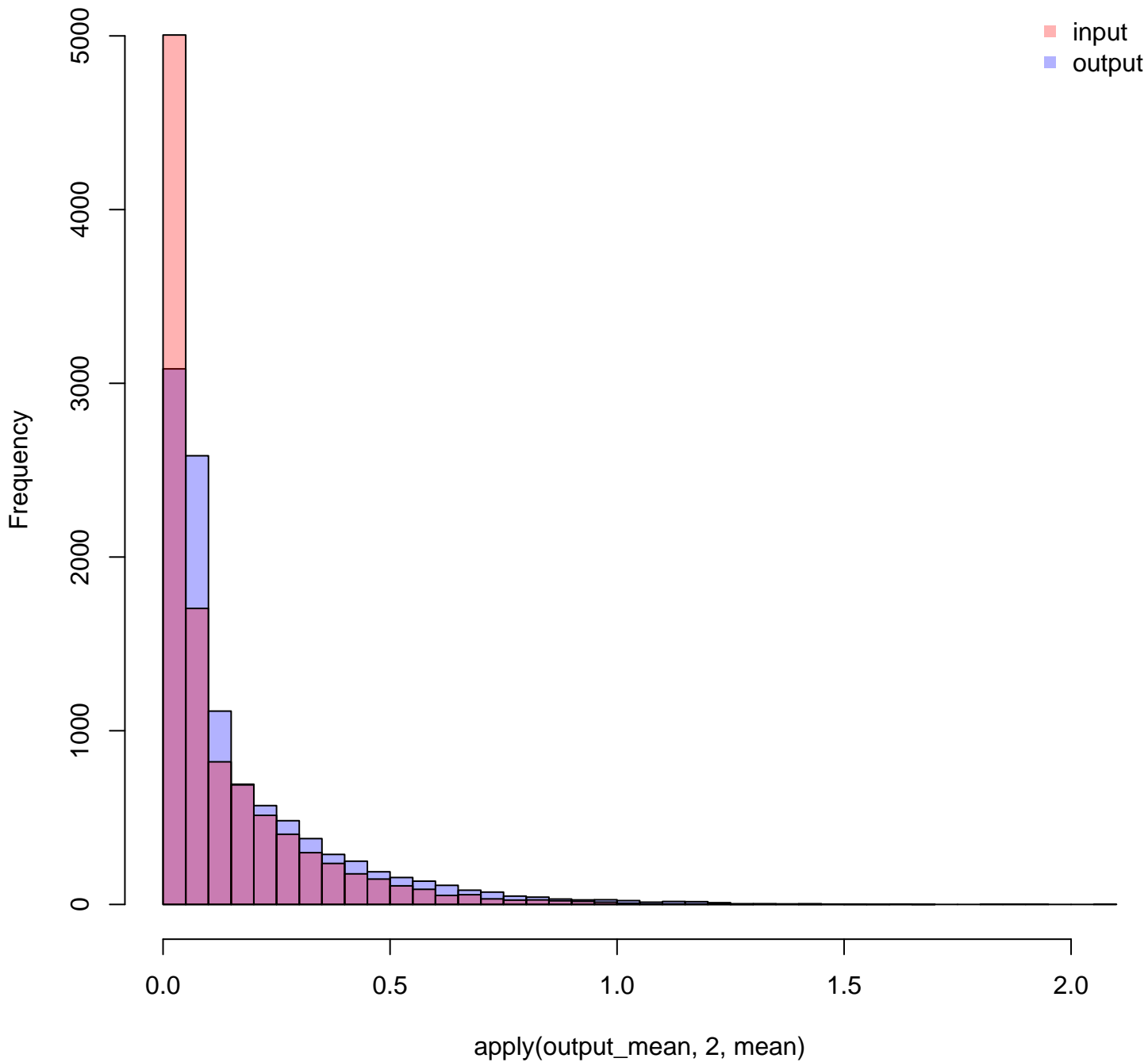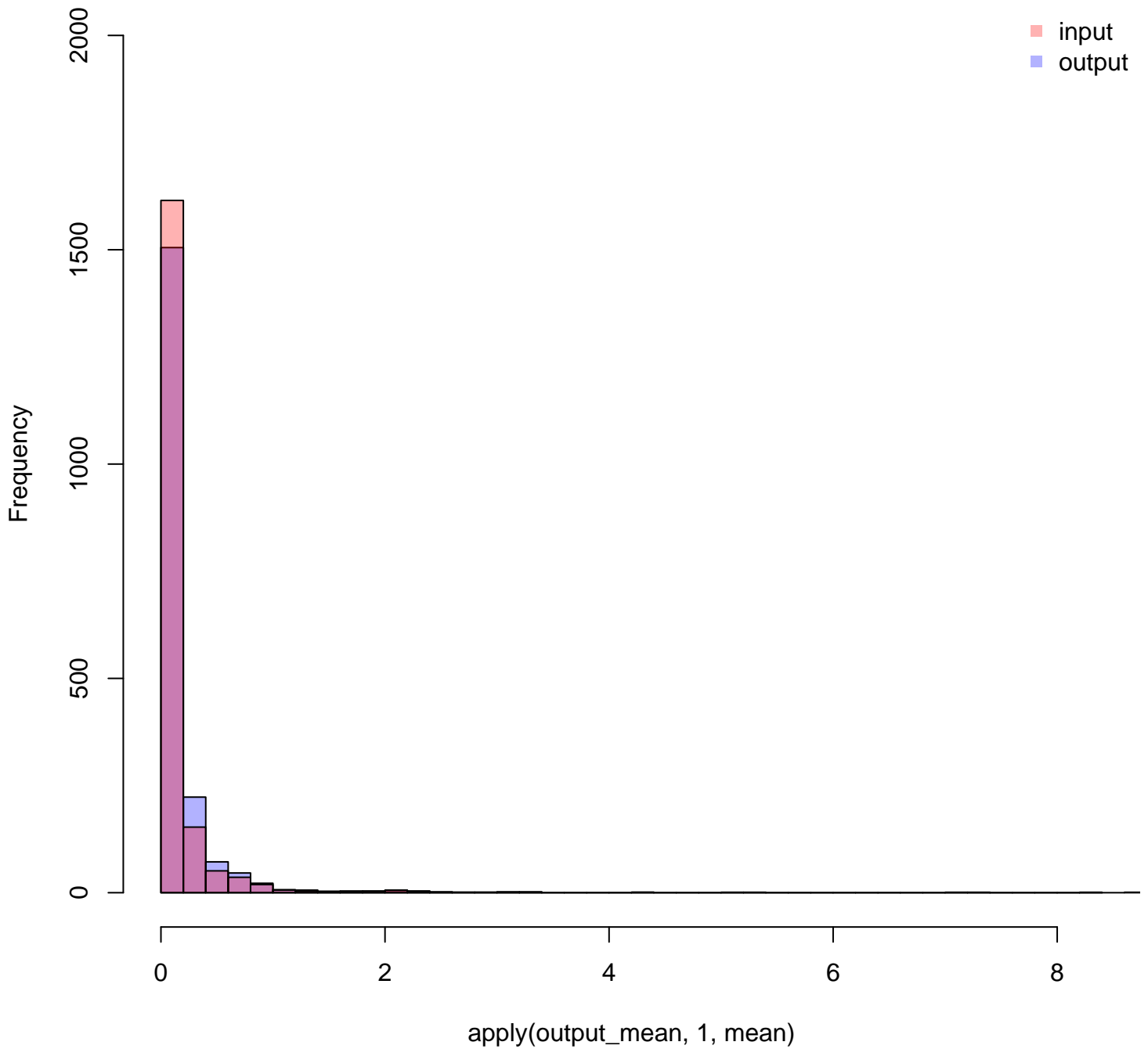**mean expression per cell, DCA input vs output**
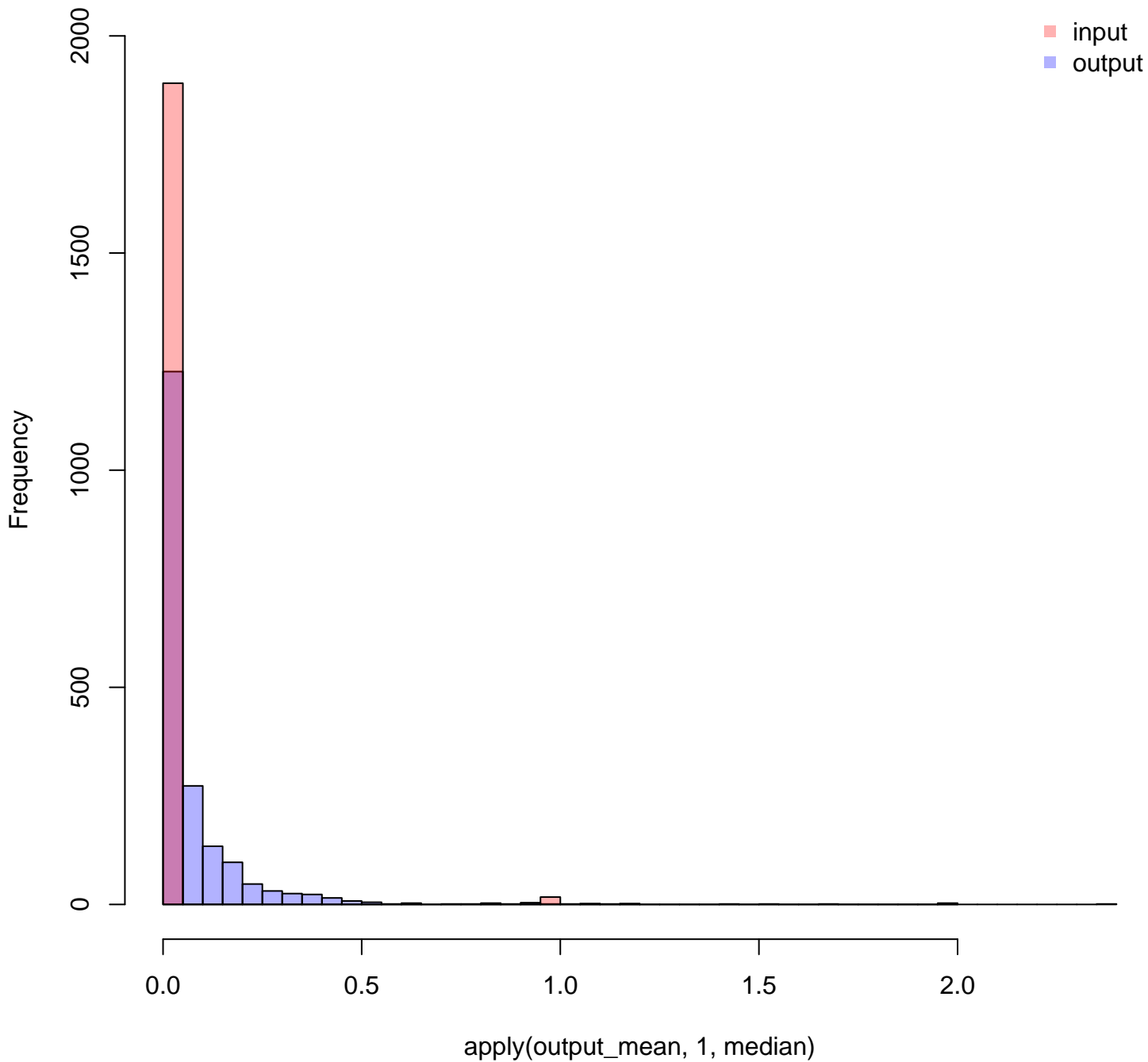
mean expression per gene, DCA input vs output
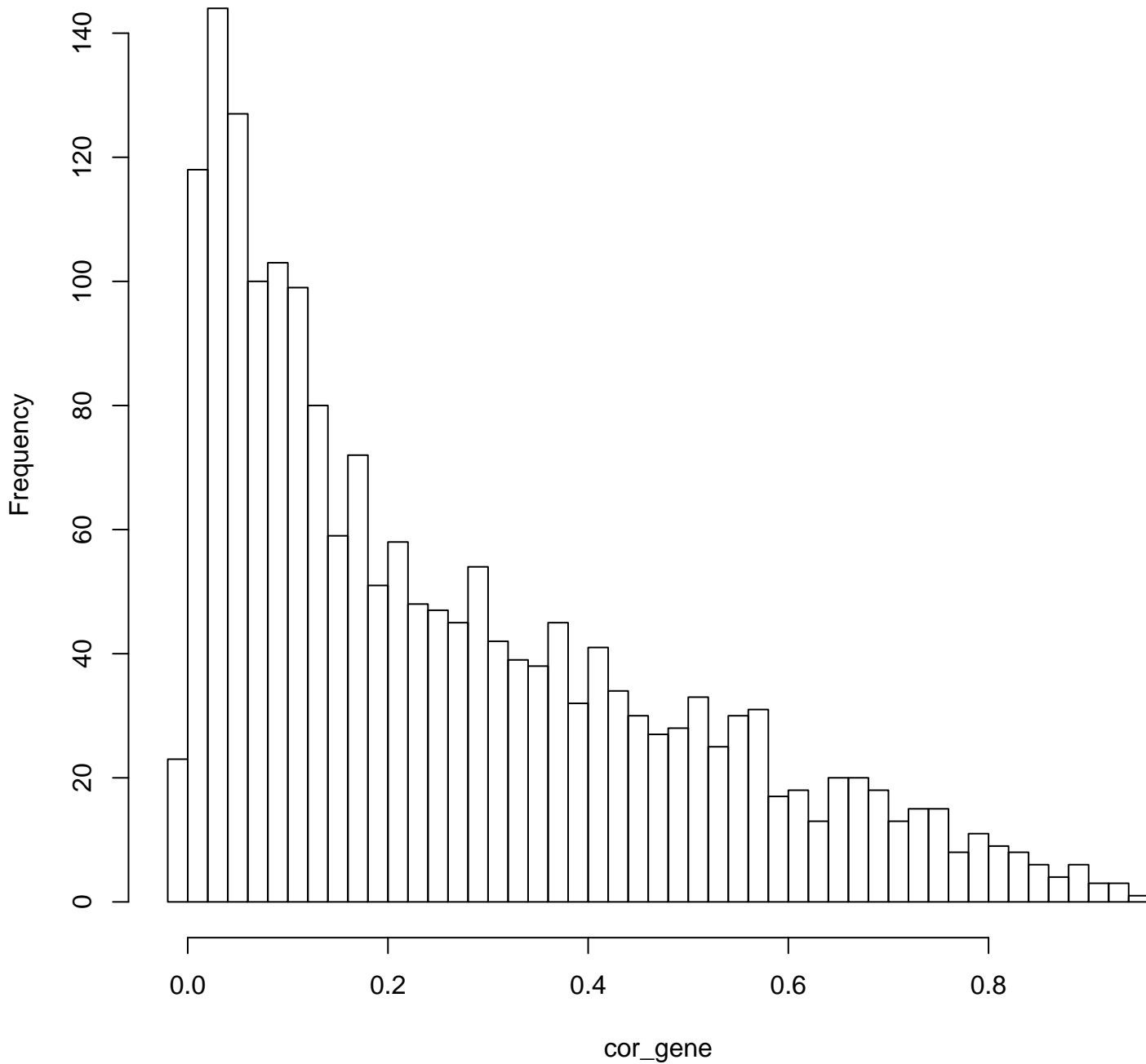
input
output

Frequency

apply(output_mean, 1, mean)

median expression per cell, DCA input vs output

median expression per gene, DCA input vs output

input
output

Frequency

apply(output_mean, 1, median)
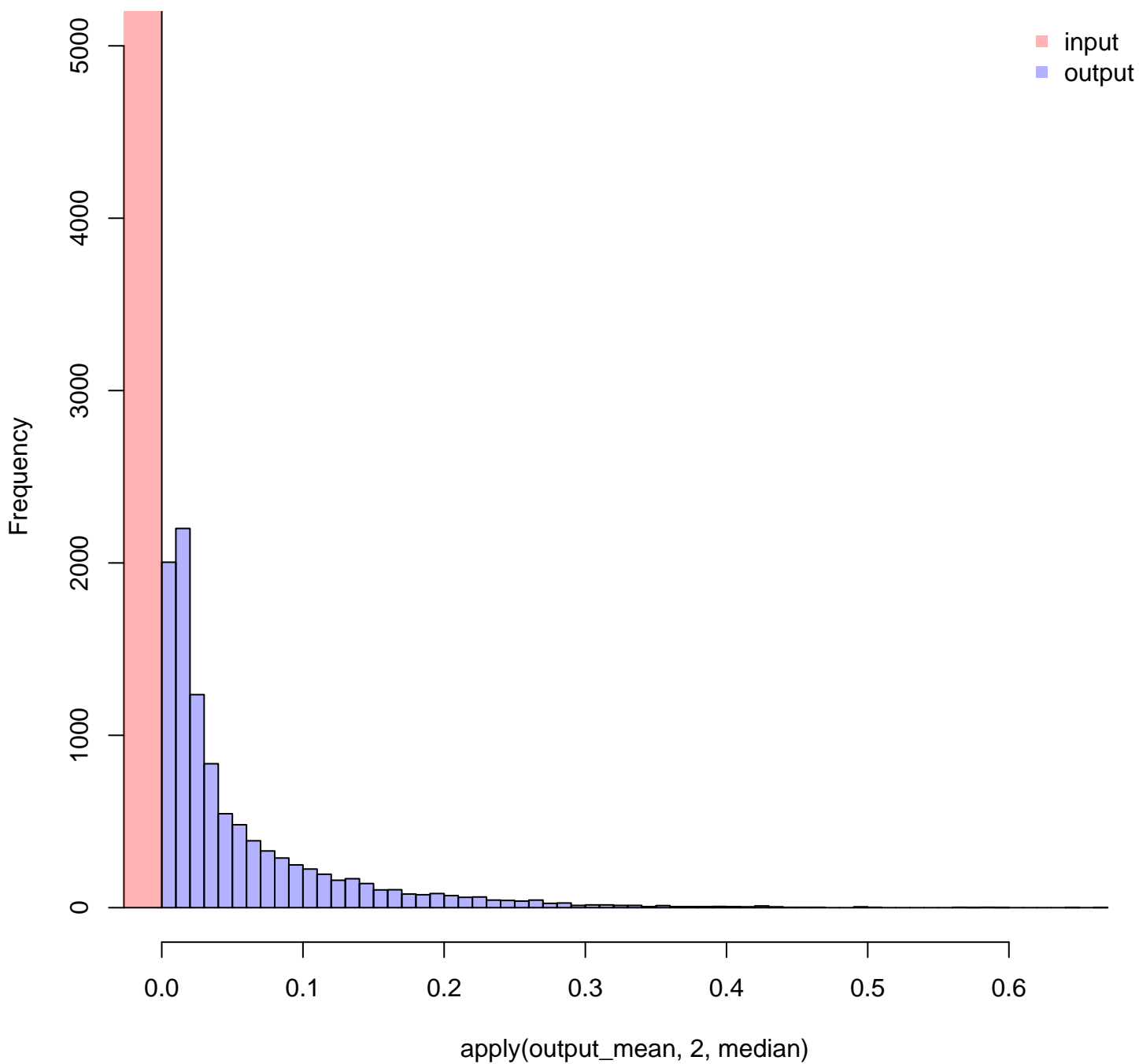
correlation of mean expression, input vs output, per gene

Frequency

cor_gene

mean vs variance