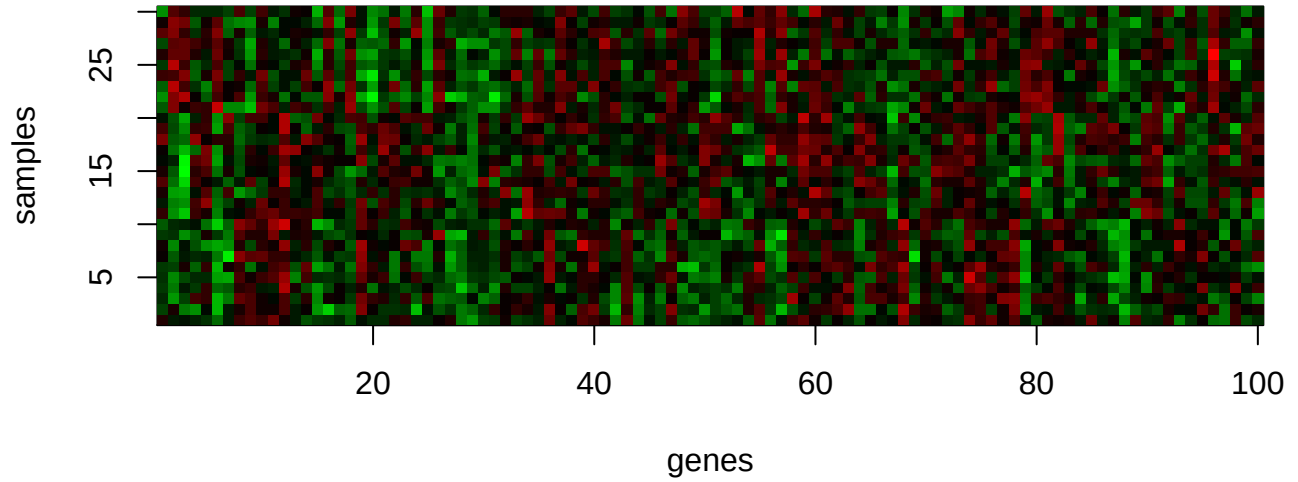
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80
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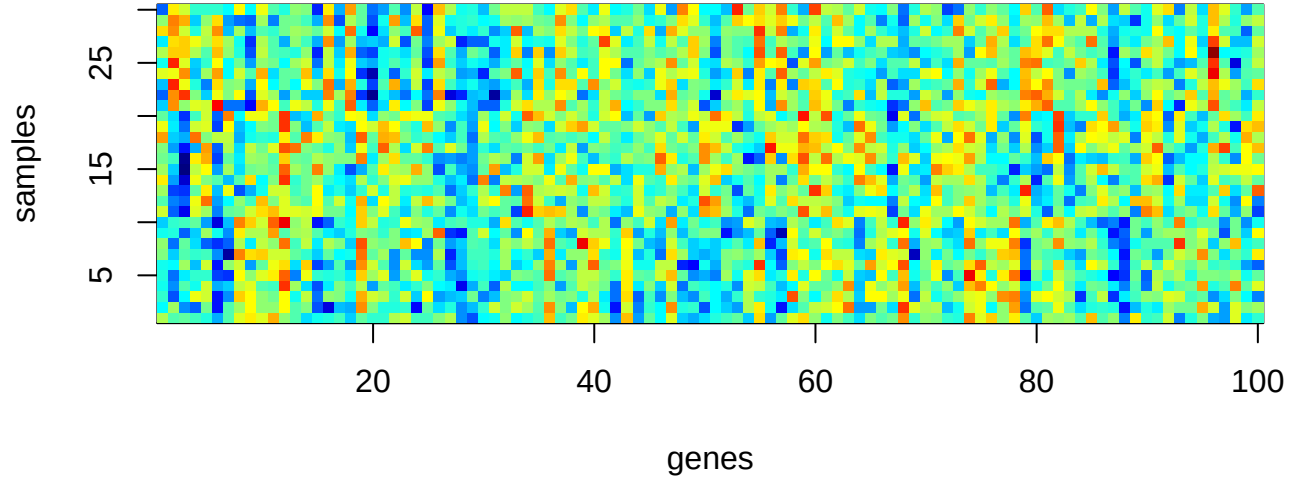
genes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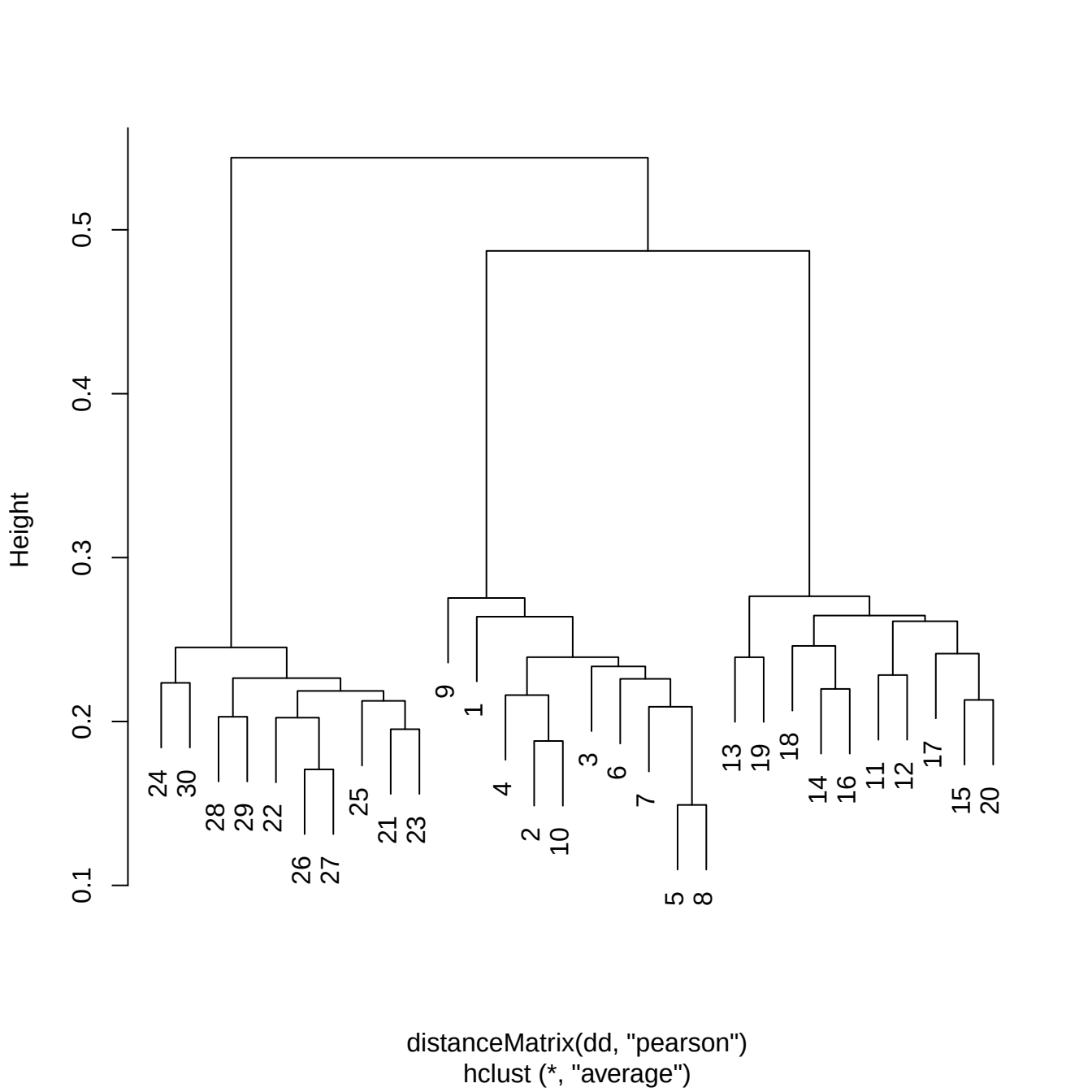
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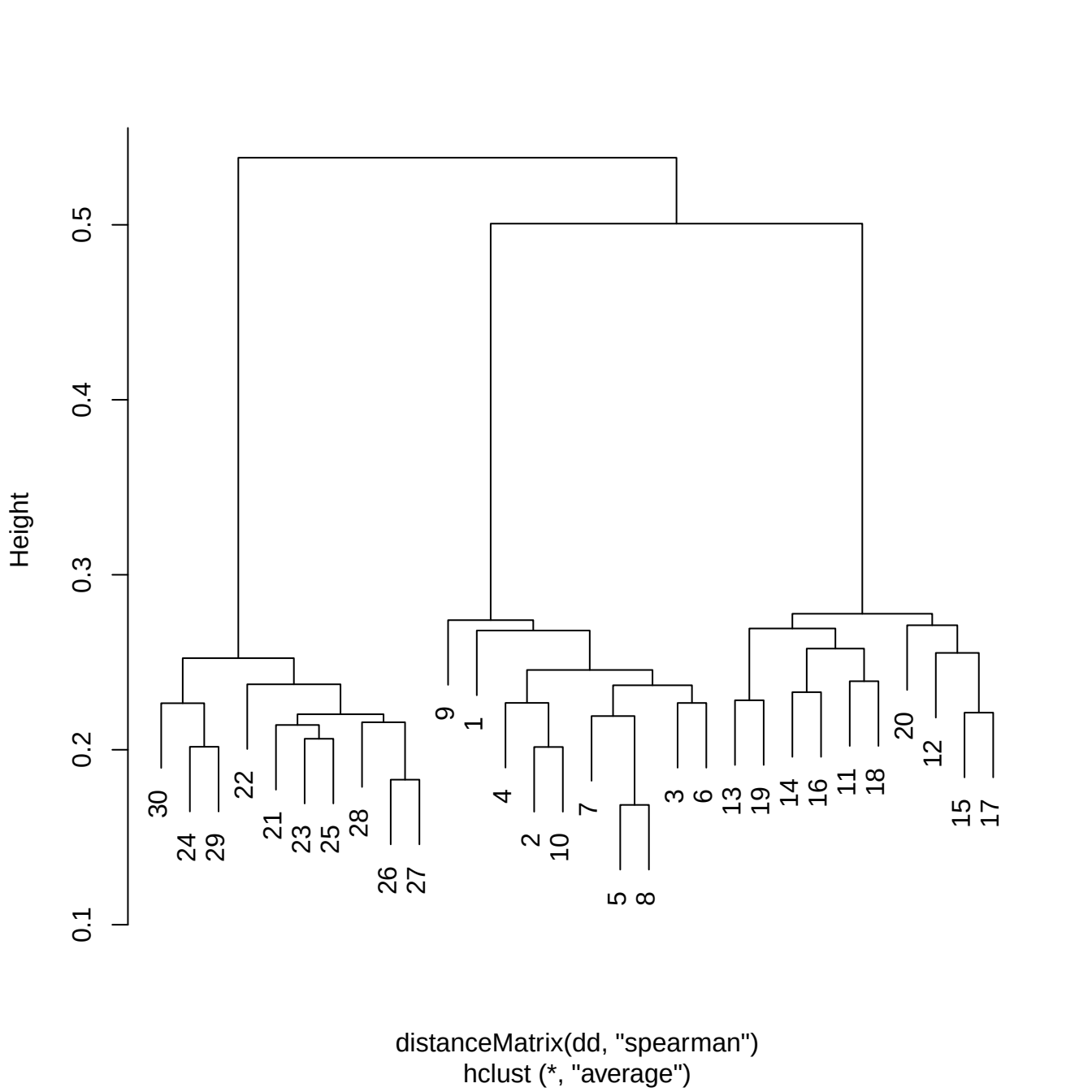
hclust (*, "average")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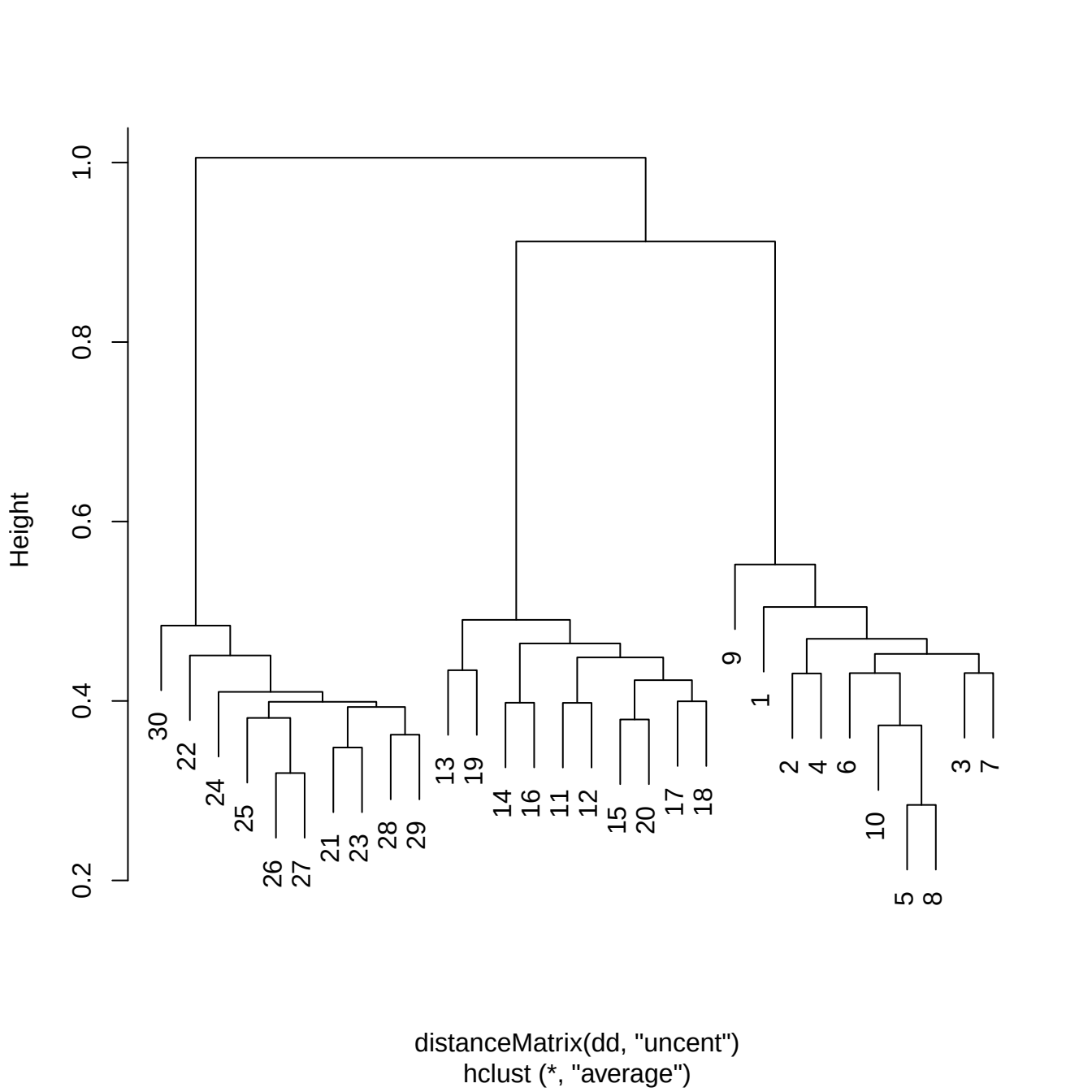
hclust (*, "average")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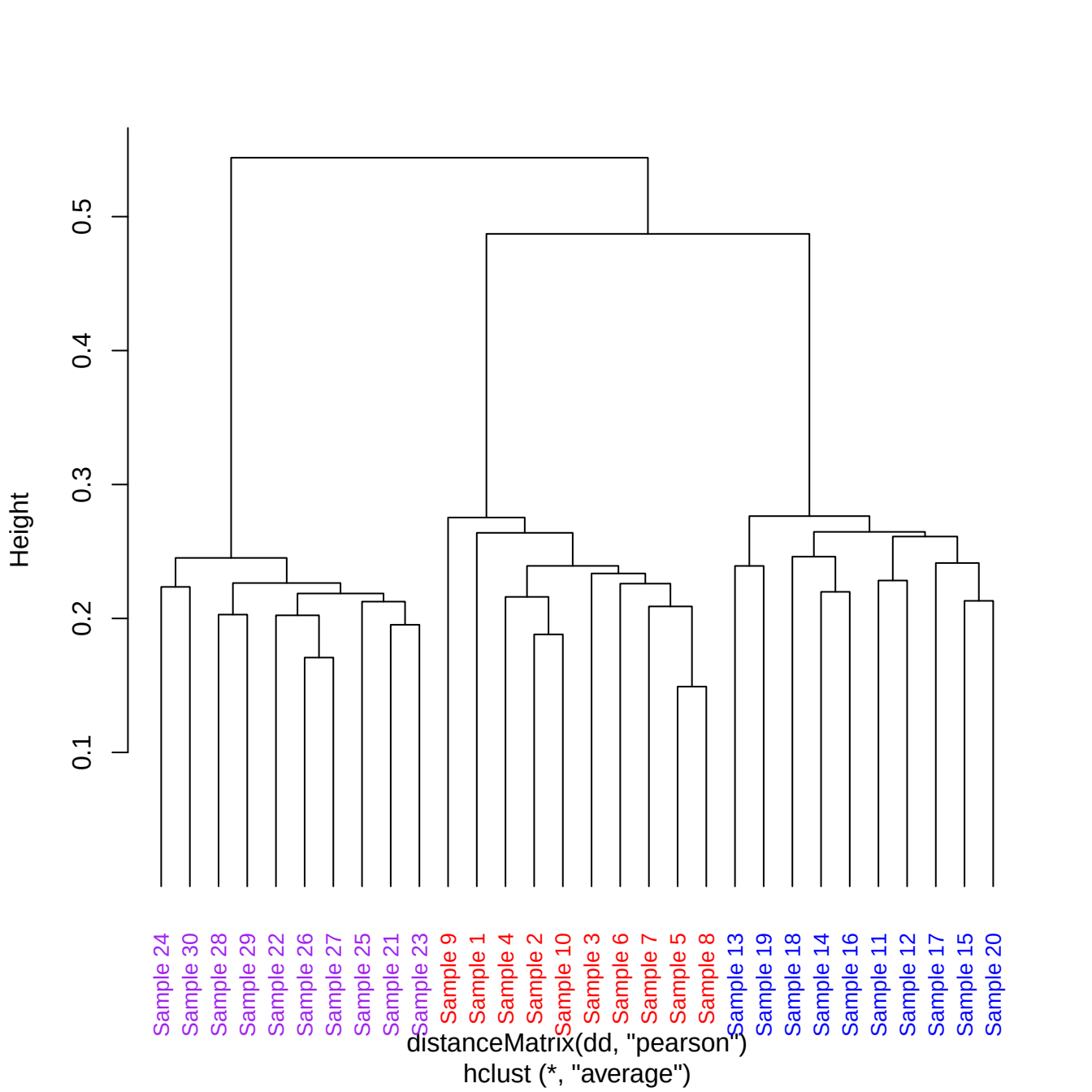
7
distanceMatrix(dd, "uncent")
hclust (*, "average")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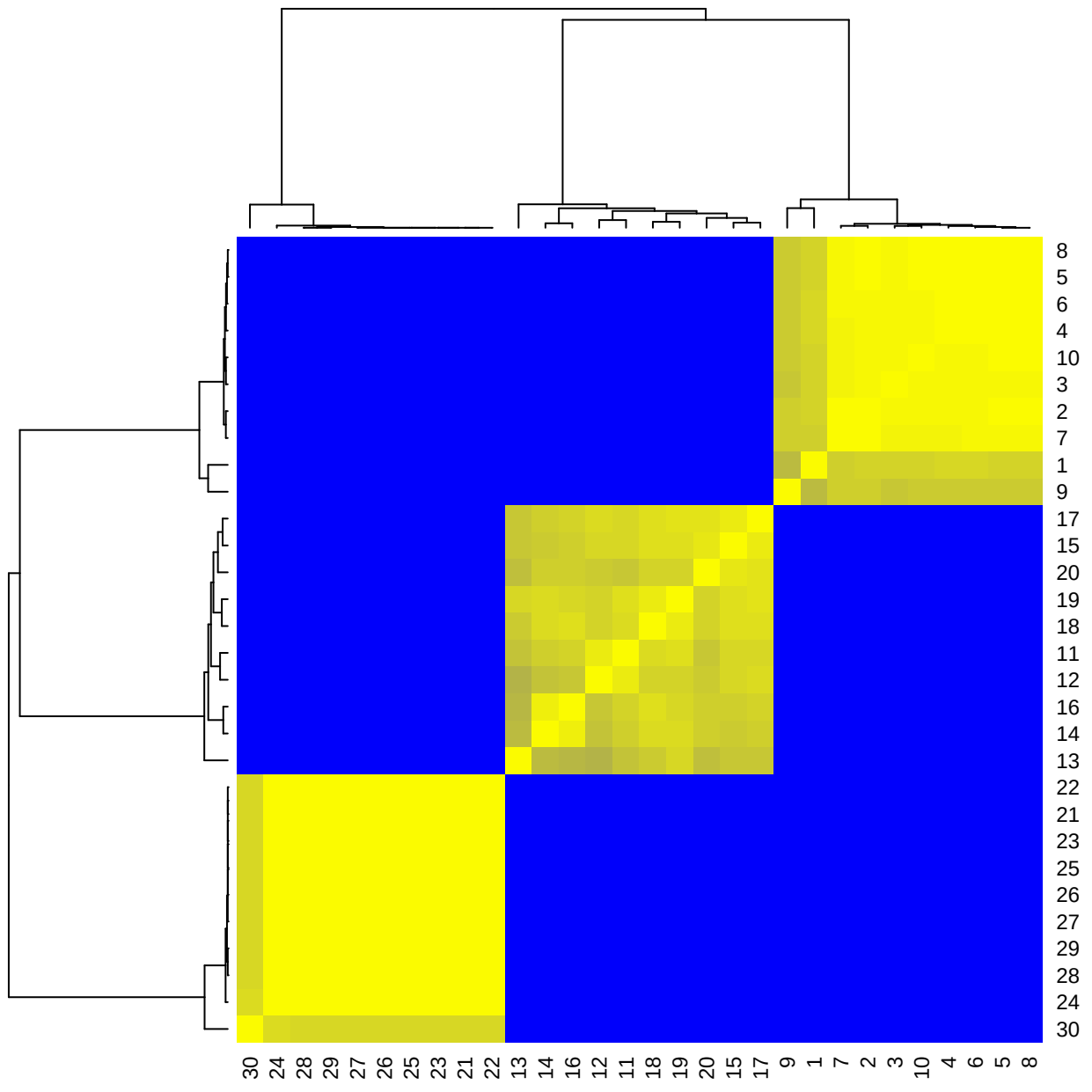
Sample 23
Sample 9
Sample 1
Sample 4
Sample 2
Sample 10
Sample 3
Sample 6
Sample 7
Sample 5
Sample 8
Sample 13
Sample 19
Sample 18
Sample 14
Sample 16
Sample 11
Sample 12
Sample 17
Sample 15
Sample 20
distanceMatrix(dd, "pearson")
hclust (*, "average")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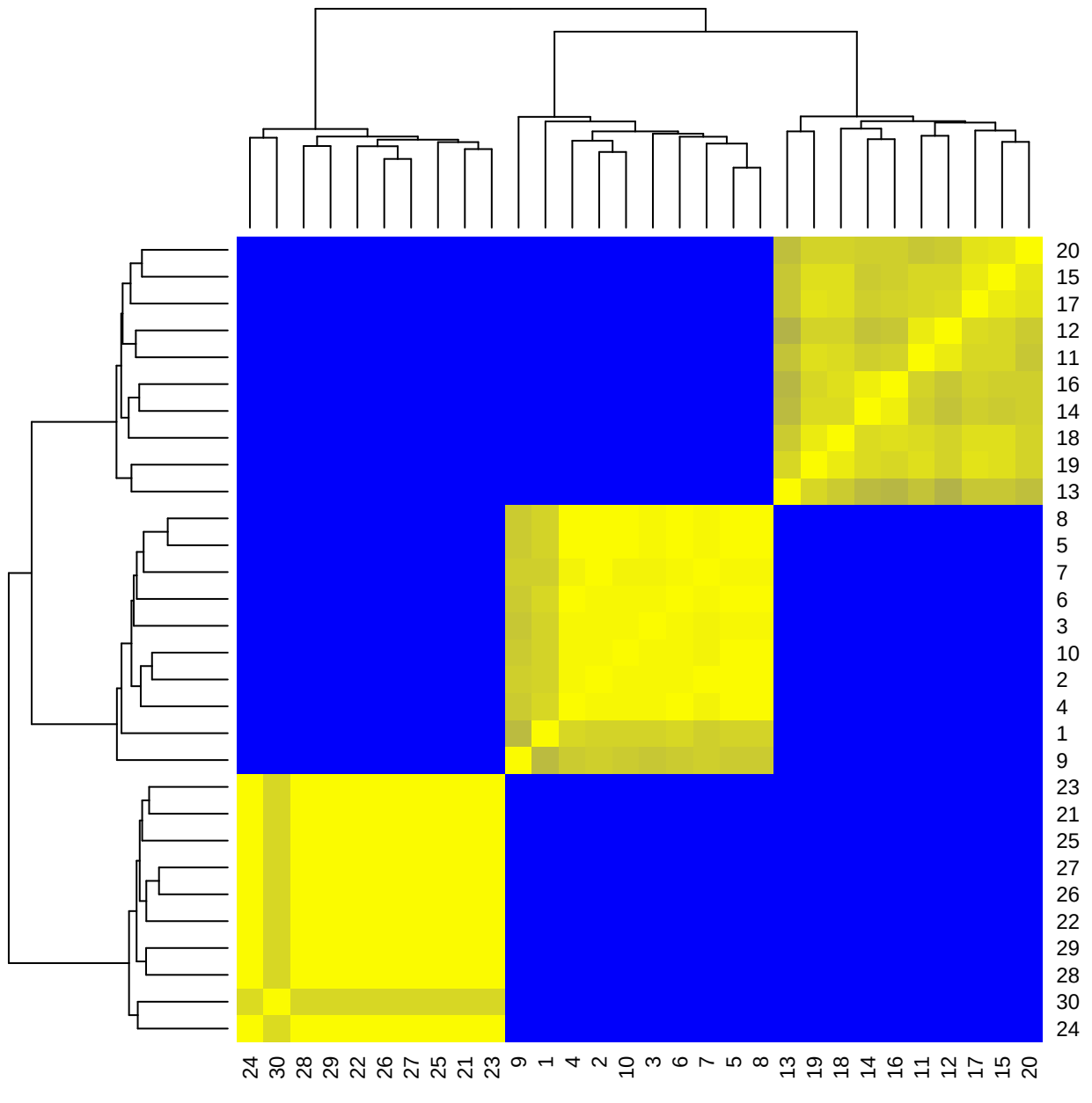
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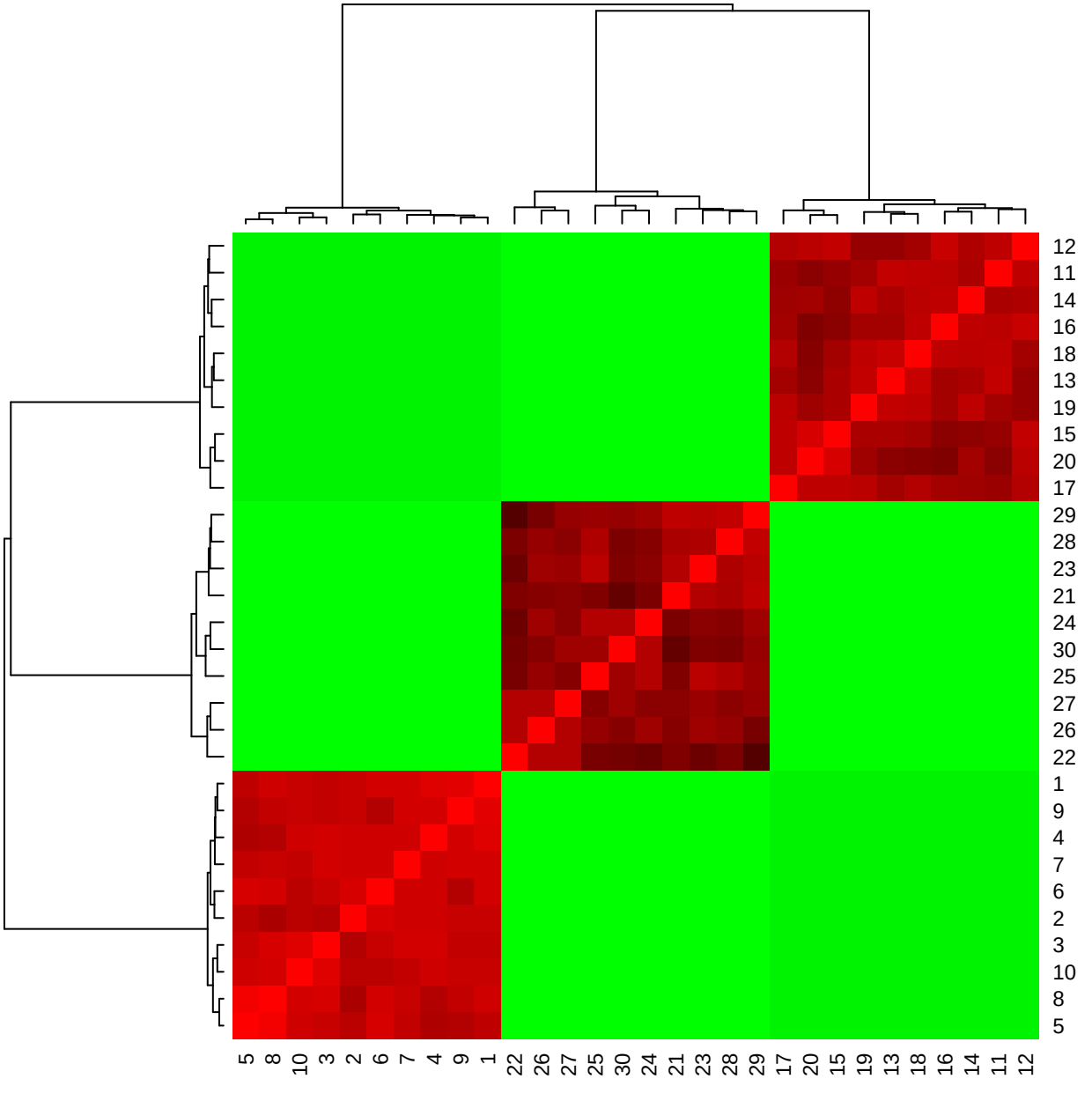
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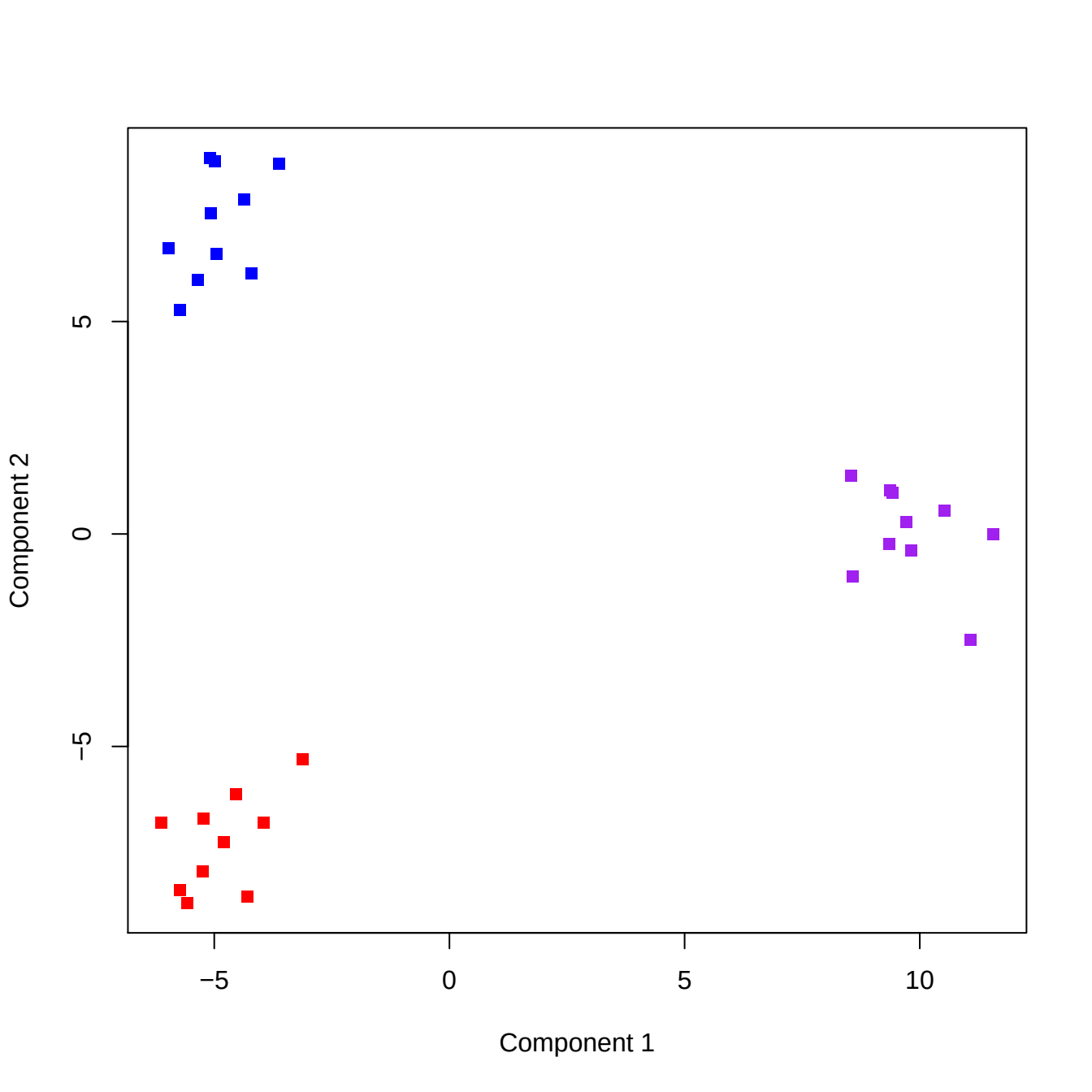
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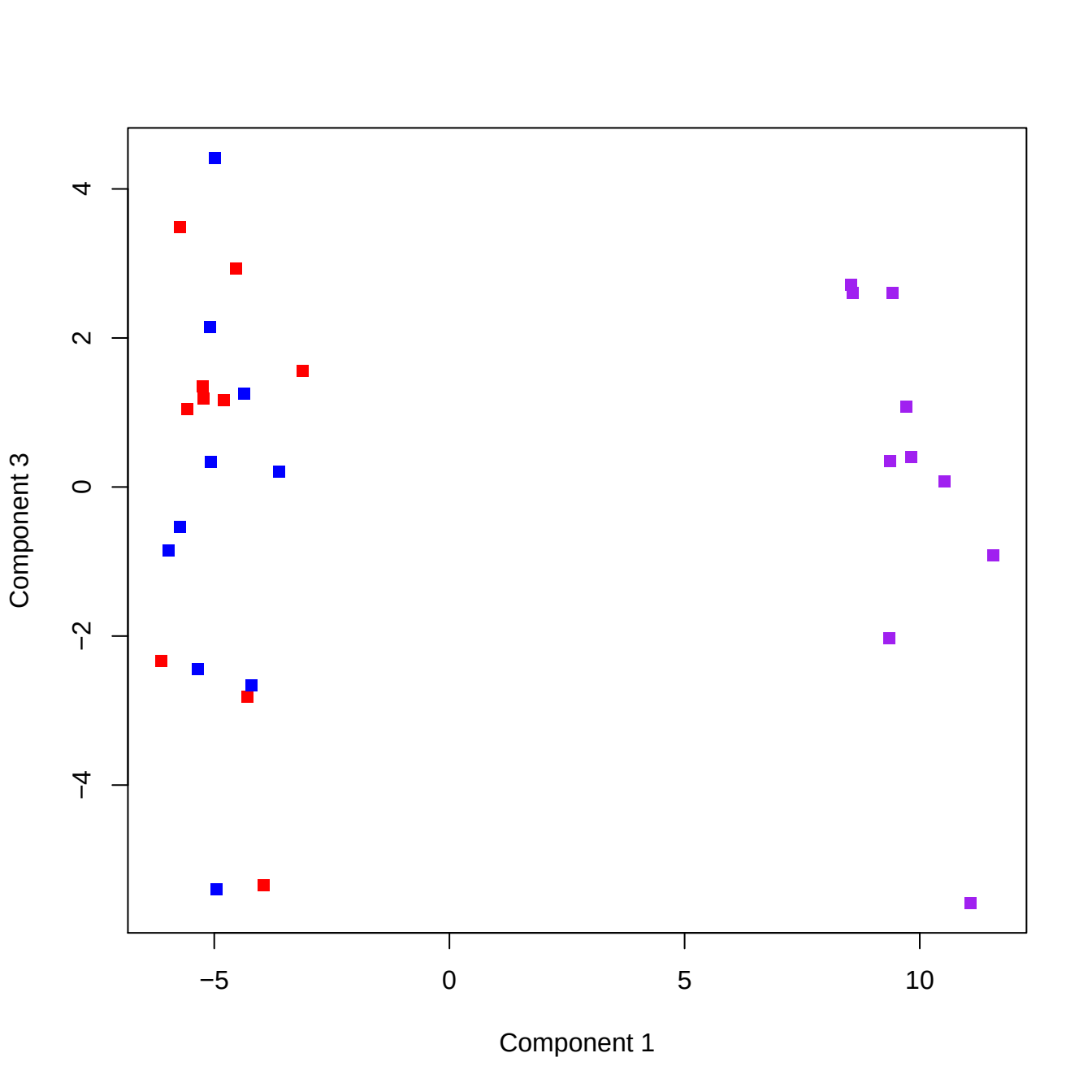
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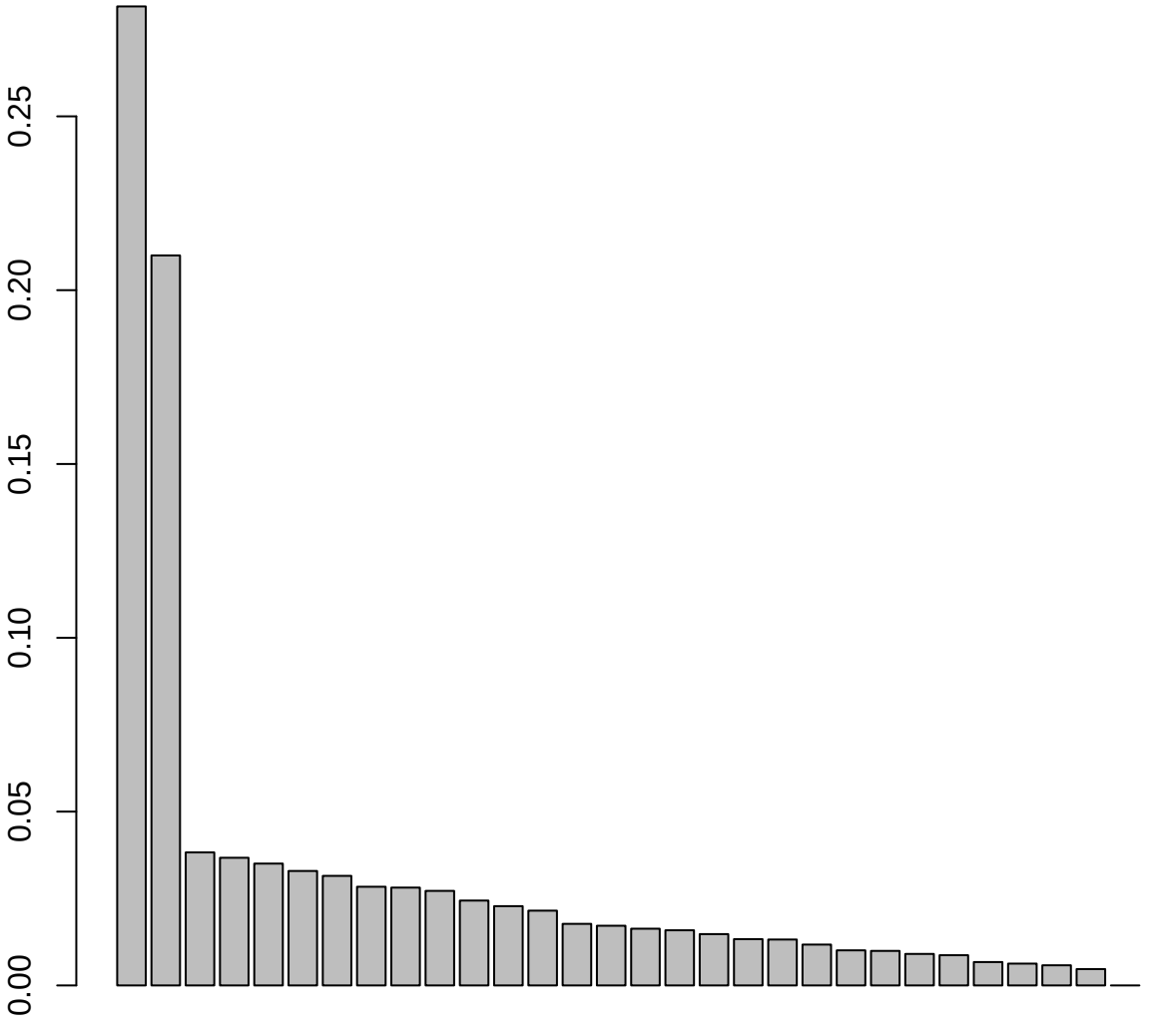
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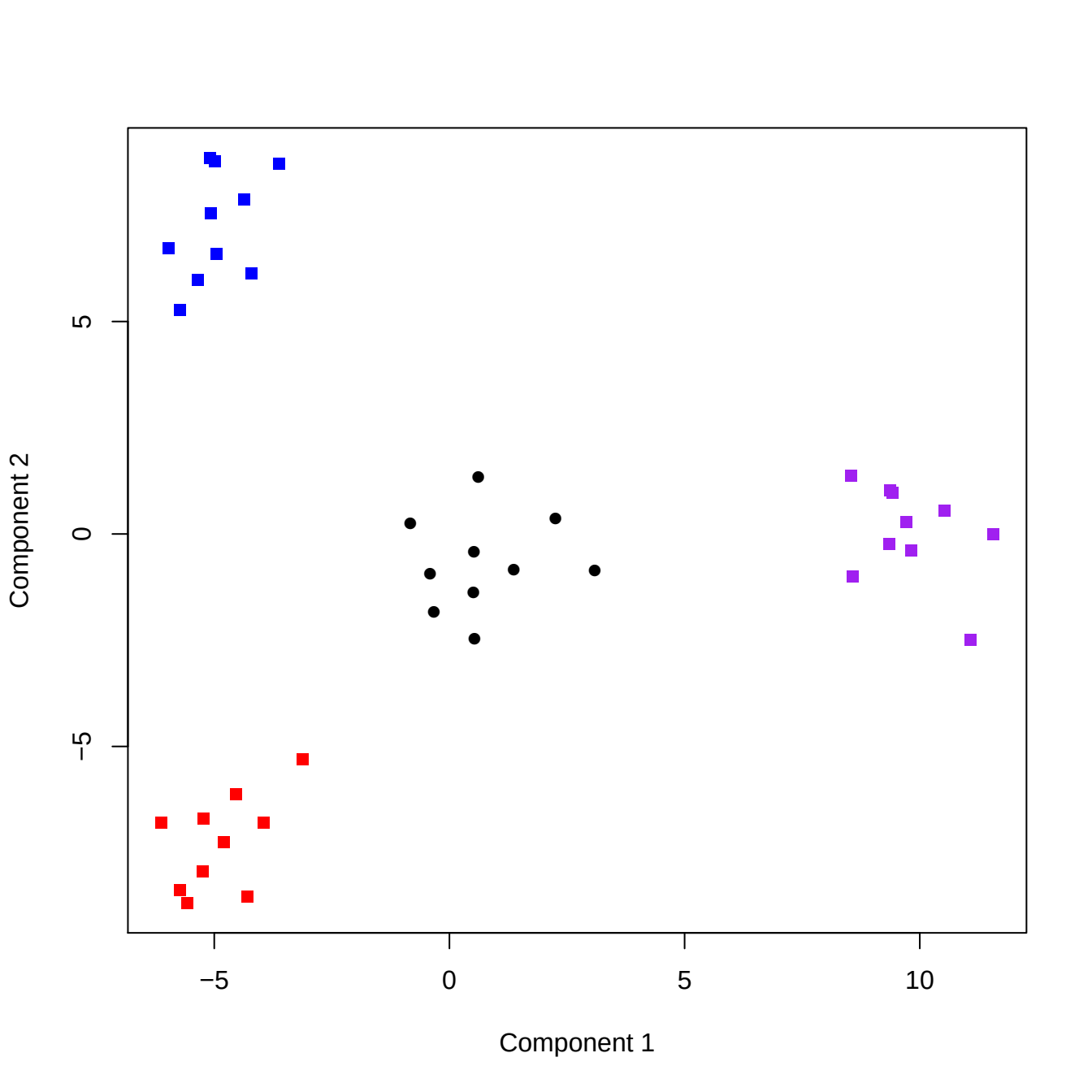
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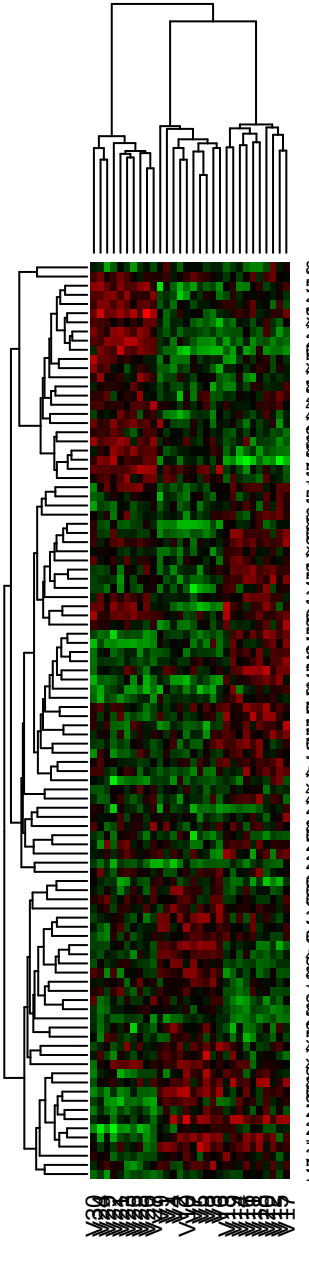
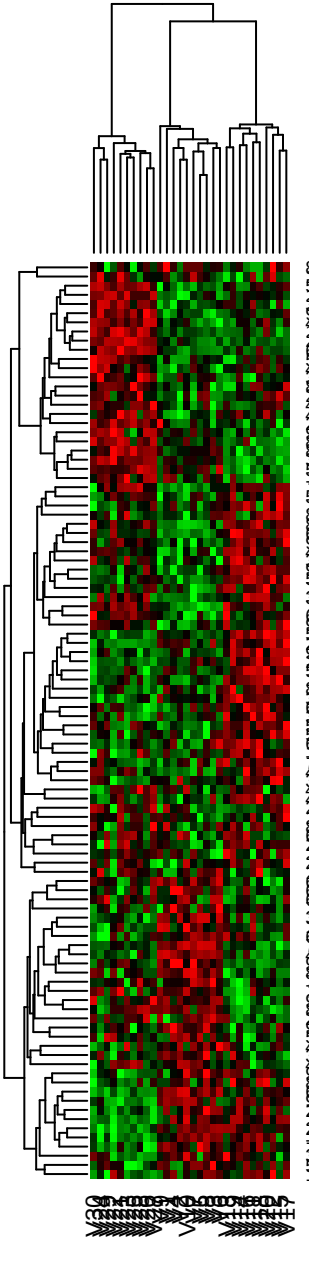
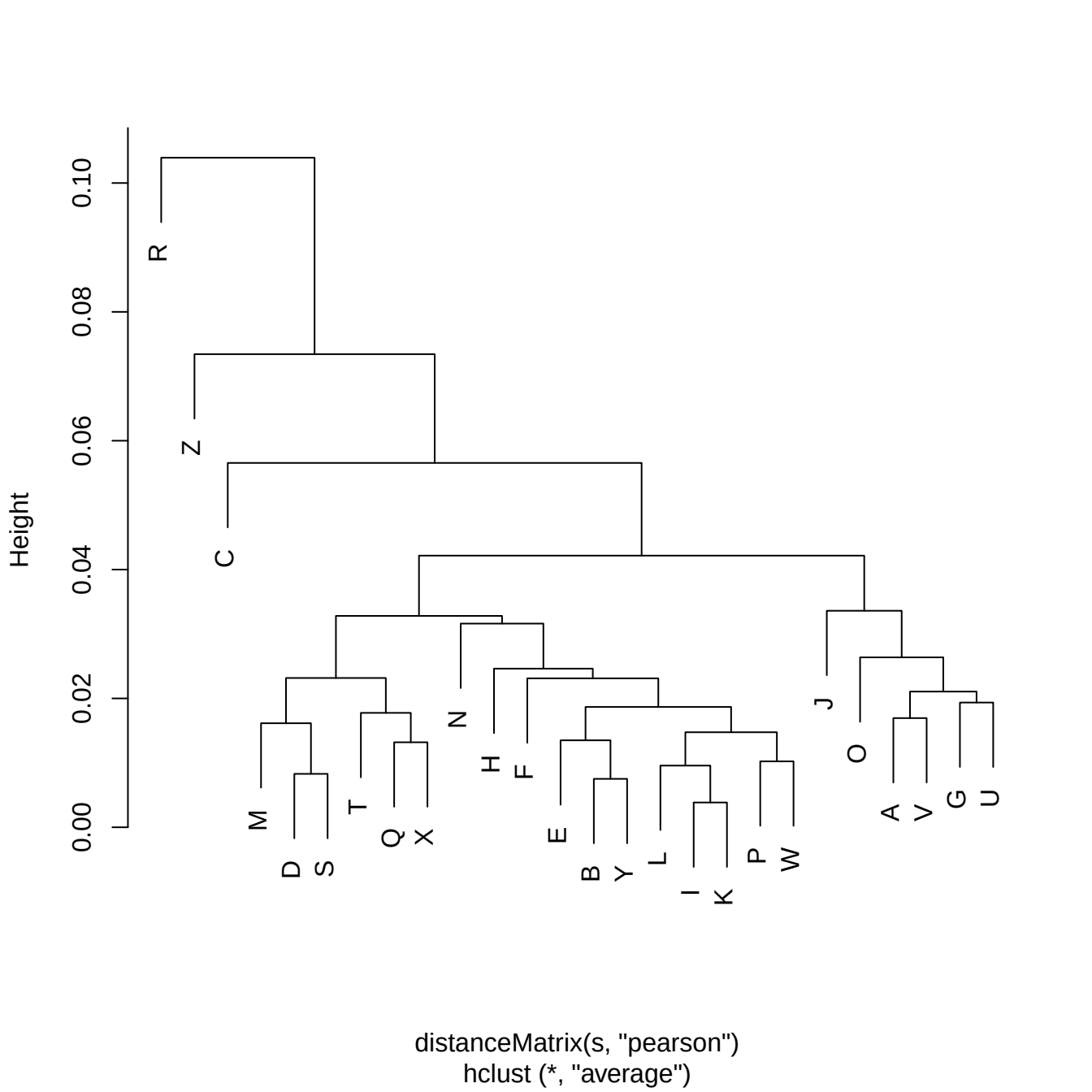
Component 1



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K
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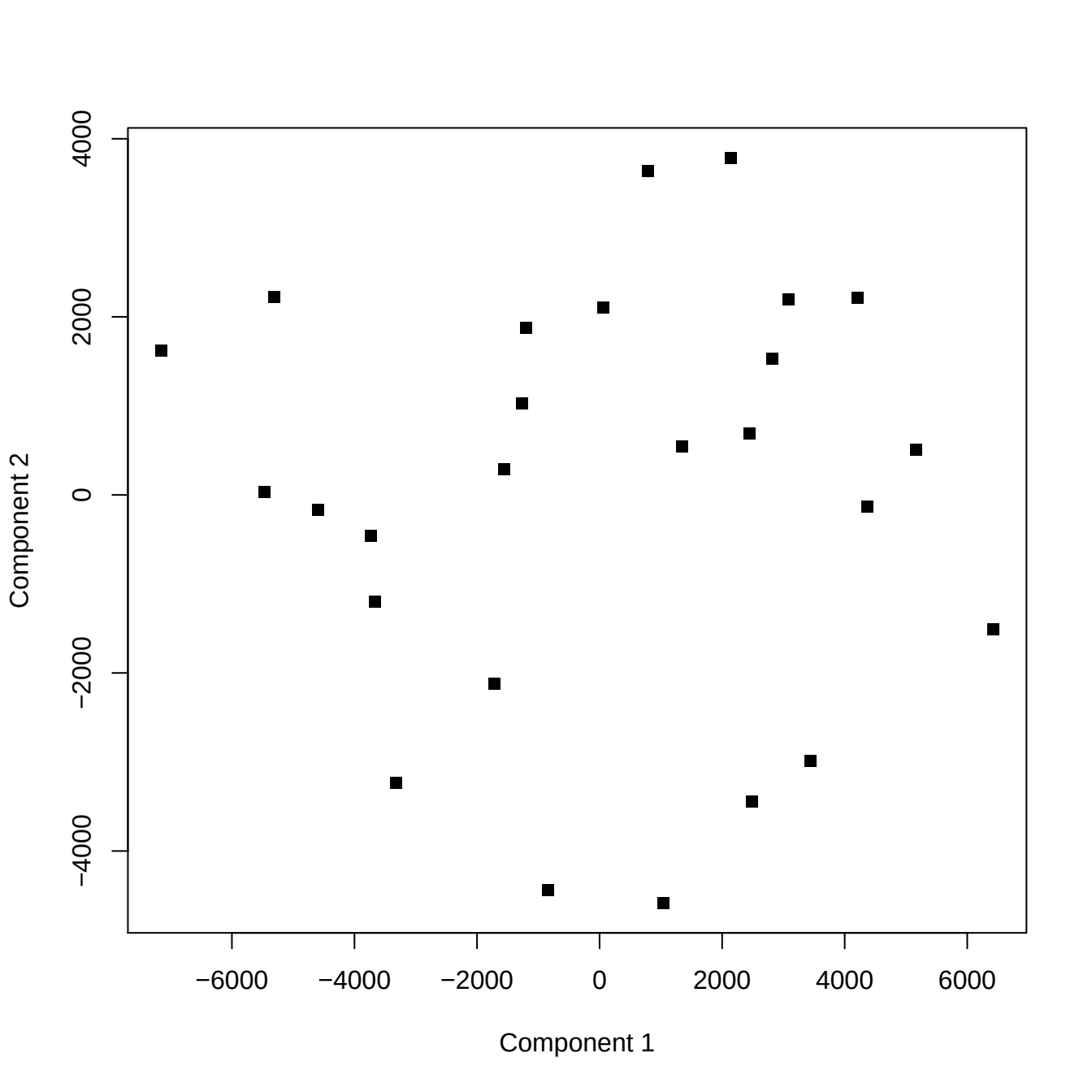
J
O
A
V
G
U
distanceMatrix(s, "pearson")
hclust (*, "average")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2000
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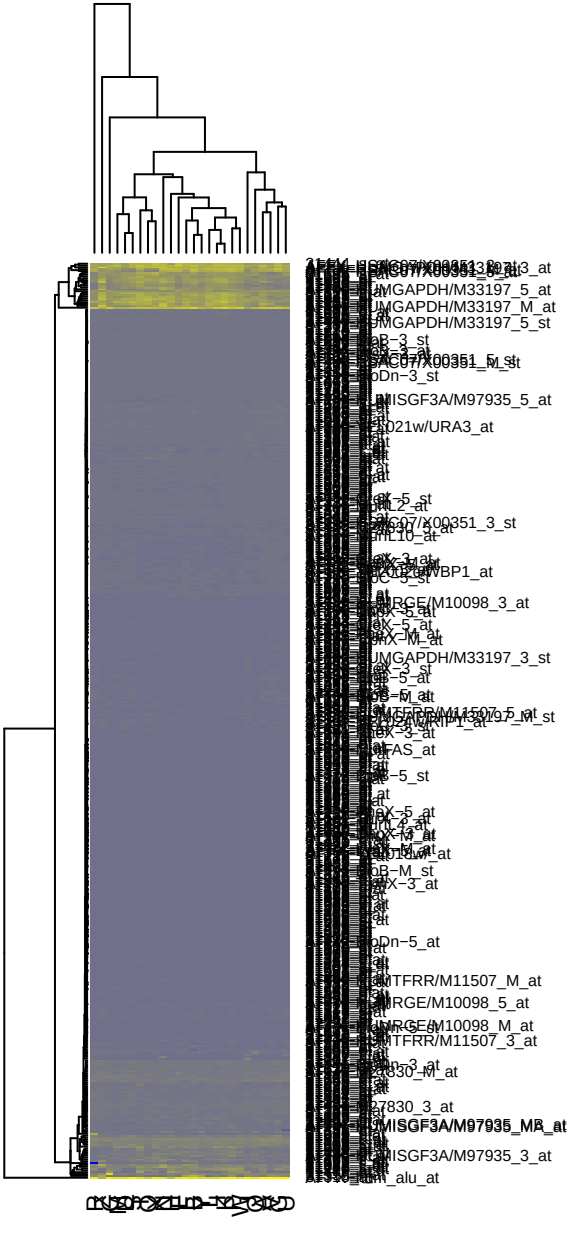
Component 1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