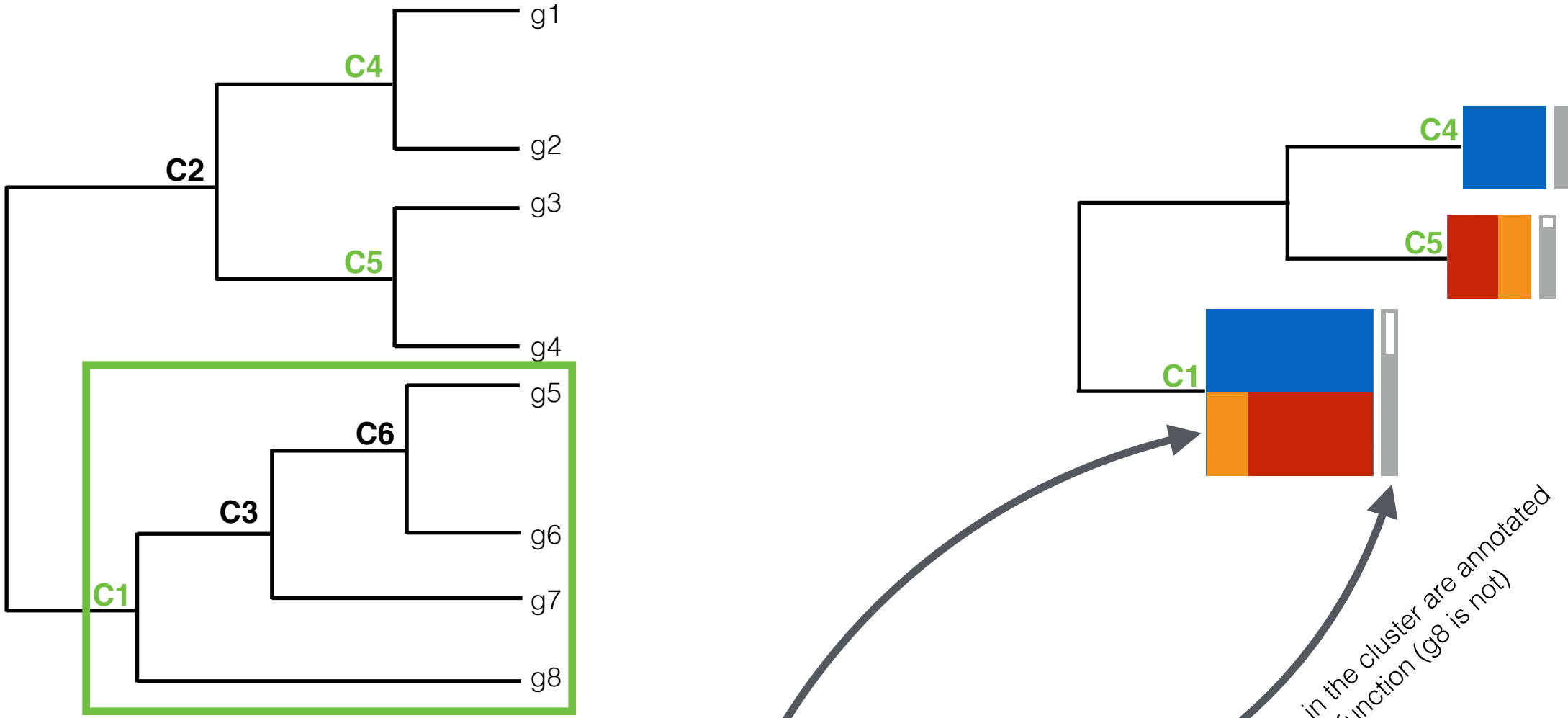


Dendrogram



Cluster C1

function	domain	P-value	Related genes
T1	BP	<0.05	g5, g6, g7
T2	BP	<0.05	g5, g6, g7
T3	CC	<0.05	g5, g6, g7
T4	MF	<0.05	g5, g6
T5	CC	<0.05	g6, g7
T6	CC	<0.05	g6, g7
T7	BP	<0.05	g6, g7
T8	BP	<0.05	g6, g7

75% of genes in the cluster are annotated to some function (g8 is not)