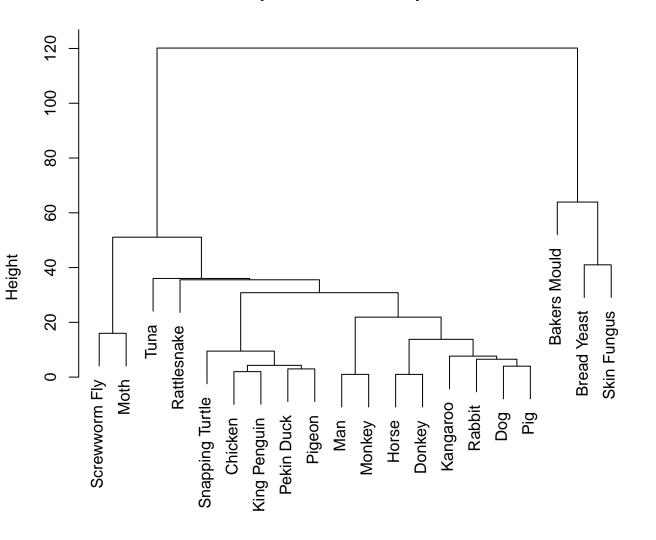
Classification hiérachique ascendante a partir du dataset Mutations



m hclust (*, "ward.D2")