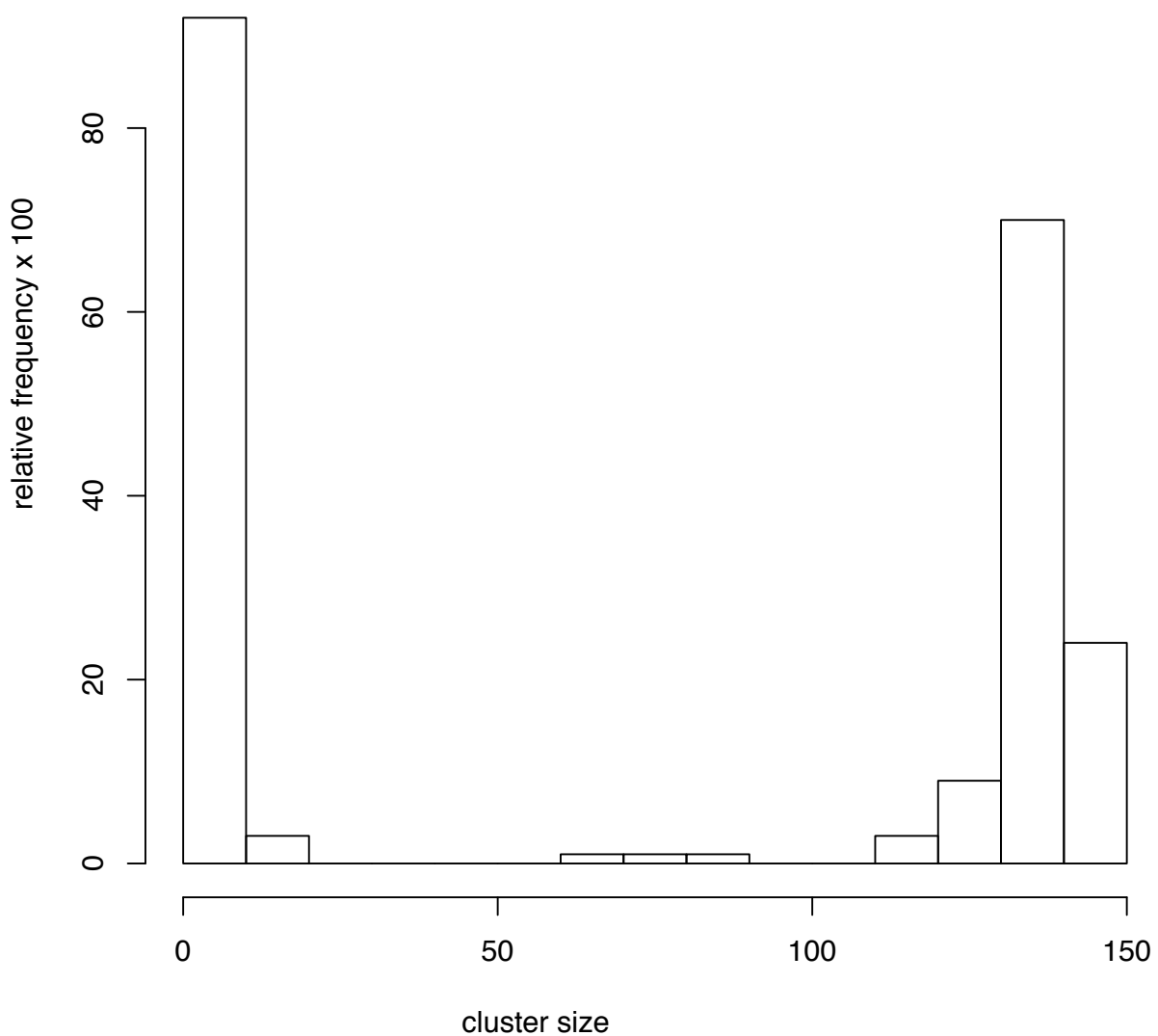




Distribution of the number of control sequences clustering together with the patient's own sequences in those patients, whose sequences failed to form a monophyletic cluster.