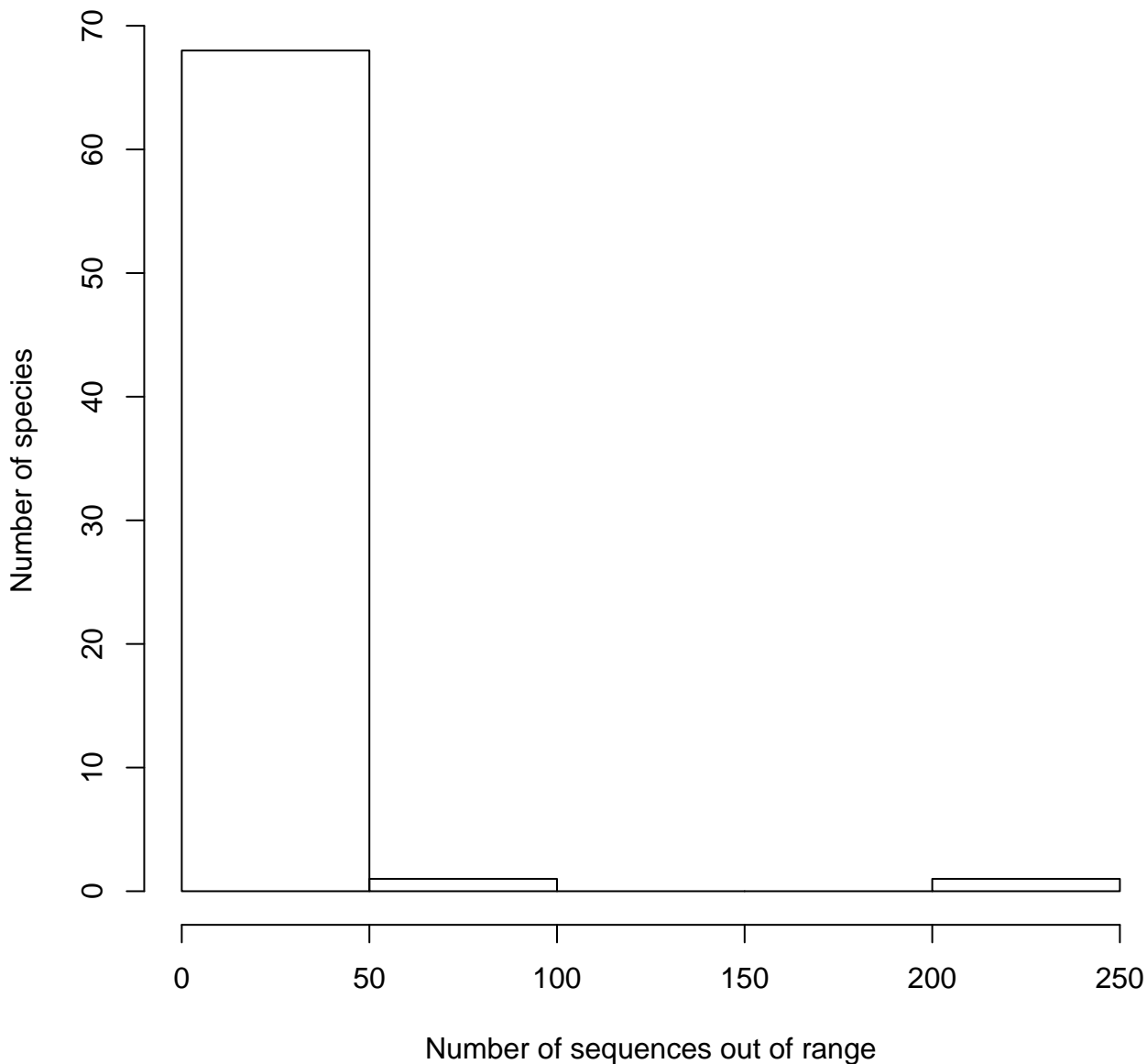
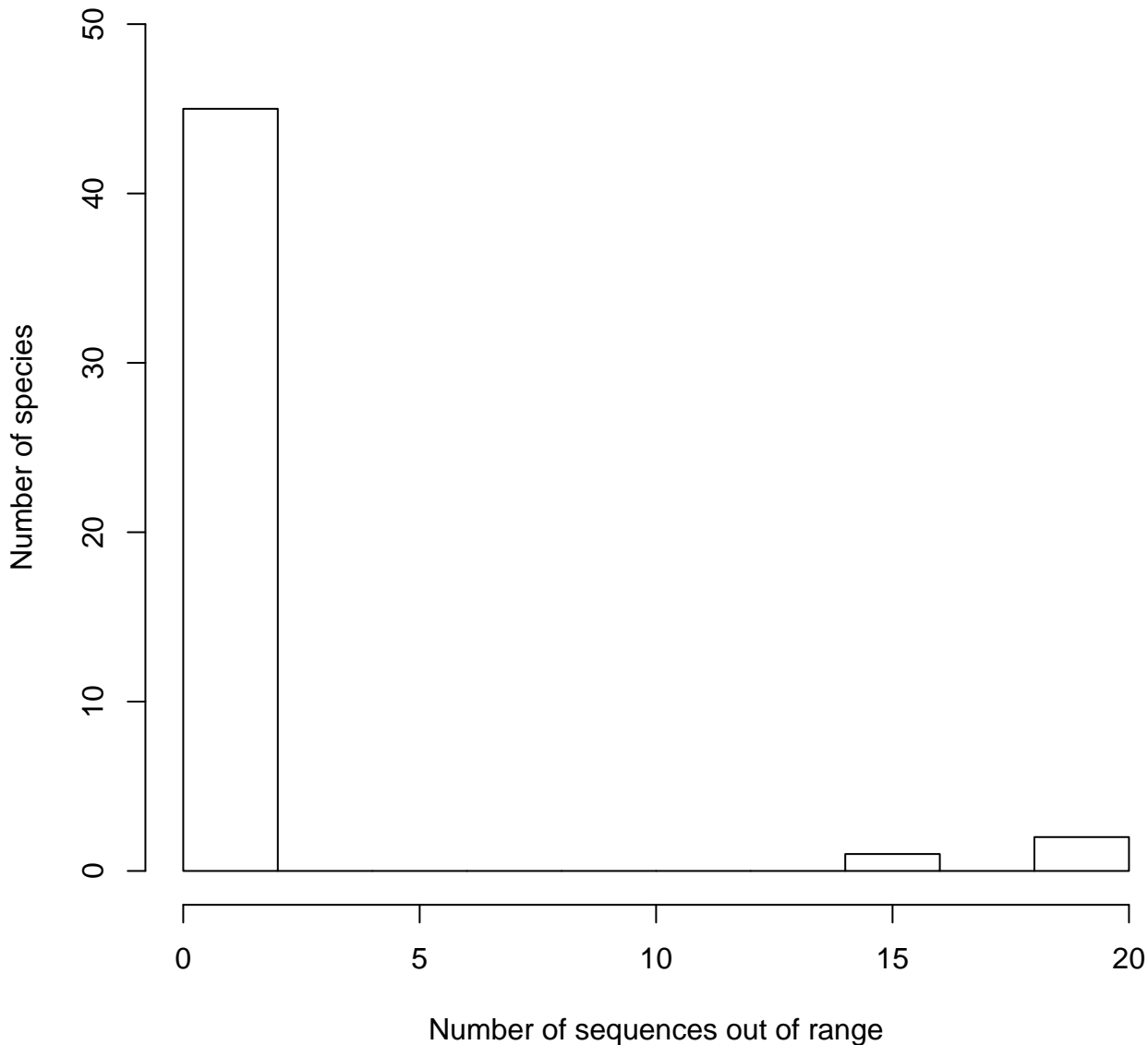


Number of Genbank sequences out of range

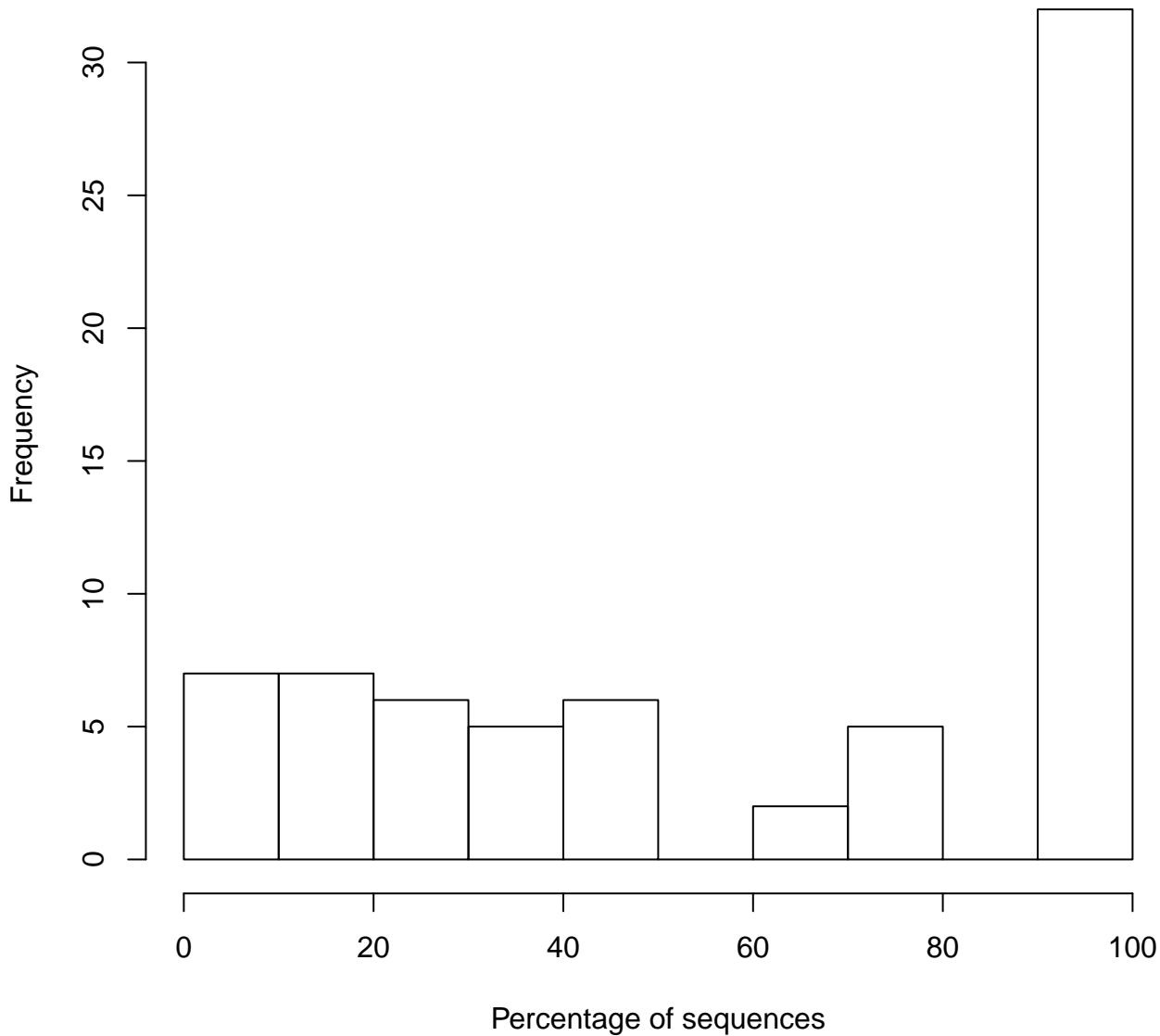


Number of Genbank sequences out of range

Interval between 0 and 20



Percentage of sequences out of range per species



Percentage of sequences out of range per species

Number of total sequences does not equal the number of sequences out of range

