6.4.5 Figure: Score vs log (number of entries).

Taken from Collins J.F., Coulson A.F.W., & Lyall A., CABIOS, 4 67-71, (1988).

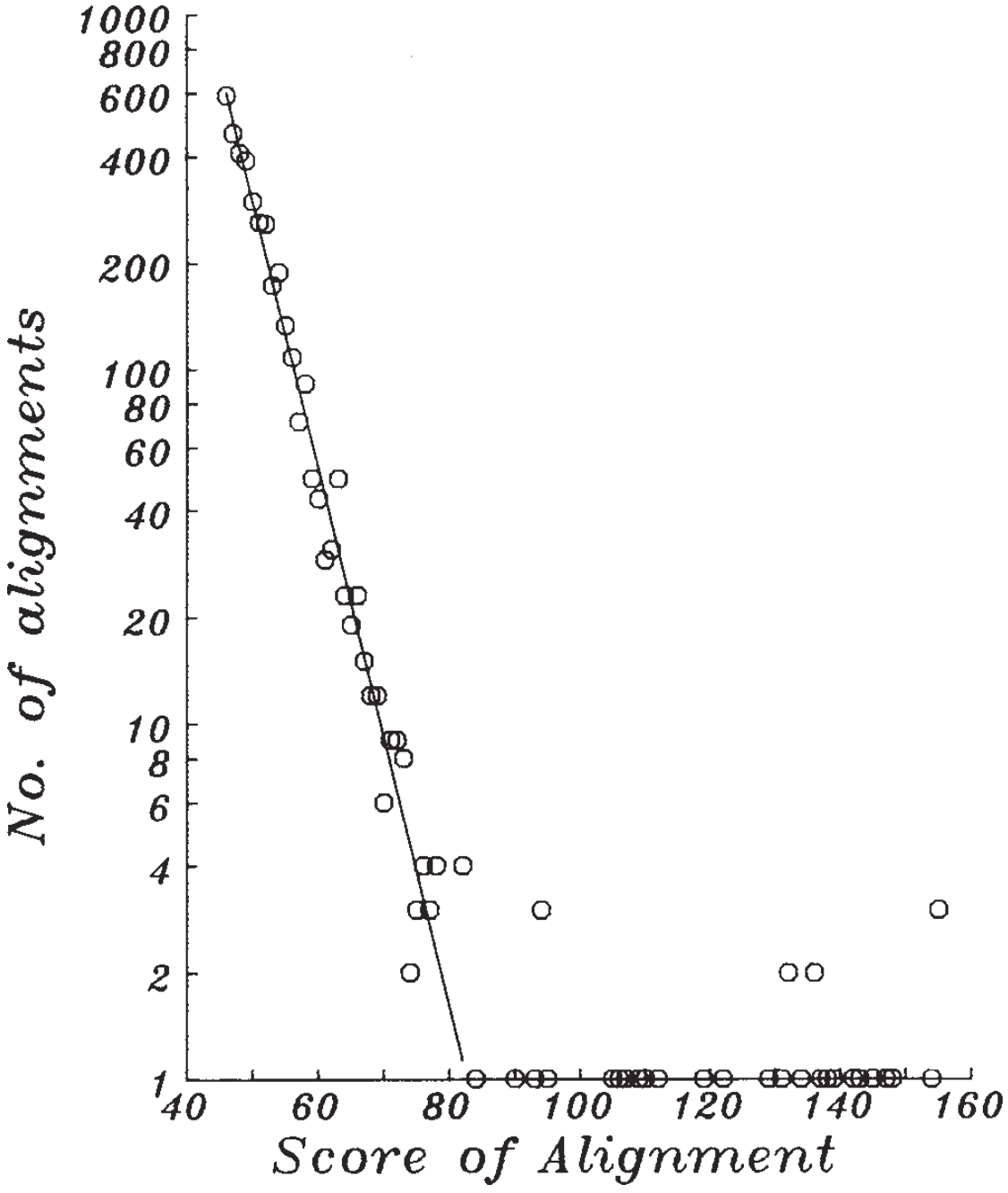


**Fig 1**. Plot of the number of reported alignments achieving a given score (logarithmic scale) against the score value, for a search of the protein database with bovine proglucagon as the query sequence, using the 100 PAM similarity table. The straight line is fitted to the lowest 97% of the reported part of the distribution. The top 49 results represent unequivocally related sequences.