The version of BLAST that inputs a protein sequence and a regular expression pattern in it is called the
PHI-BLAST

In multiple alignment, alignment is scored by one of the following options
Scoring each column of the alignment and adding up the column scores

Data sequence library was established in 1980 by
European Molecular Biology Laboratory

Atlas of Protein Sequence and Structure provide useful information in the field of
Computational biology

One of the following statements is incorrect about EST databases
They are generated by double-barrel gun sequencing method

Analysis of nucleic acid composition,secondary structure, transcriptional factors and promoter sites, are components of
Genomic analysis

National Biomedical Research Foundation helps the scientists in
identifying and interpreting the information of protein sequences

A sequence-based database program that performs, one-by-one, a heuristic local alignment of the query sequence is the
BLAST

Macromolecular sequences was published under Atlas of Protein Sequence and Structure in the year
1965-1978

Which among the following options is Japanese sibling of GenBank?
DDBJ