

DNA Identification Sequences

Specimen: Deer Antler

Description: Nucleotide collection (nt)

Program: BLAST 2.8.1+

Molecule Type: Nucleic acid



DNA barcode test is a molecular biology technique that uses a recognized and relatively short DNA fragment in the genome to identify species. It is fast, accurate, and free from the morphological characteristics of the sample. It is suitable for medicinal materials and bases. The identification of the original species is an effective supplement to the traditional Chinese medicine identification method. This study is based on the ITS sequence to identify the source of wild Deer Antler.

XYSY1

GAATAAATGTTGATATAGGATGGGATCCCCGCCTCTGCTGGGTCGAAGAAAGTTGTATTTAAATTC-
GGTCTGTTAGTAGTATTGTAATTCGGCTGCTAGTACAGGAAGTGAGAGAAGTAATAATACAGCAGT-
GATTAAGACAGATCATACAATAAAGGGGTTTGATACTGTGATATAGCAGGAGGTTTTATATTAATA-
ATTGTTGTGATAAAGTTAATTGCTCCTAAAATTGAGGAGACACCTGCTAAGTGTAAAGAGAAAAATGTTA-
AGTCTACTGAAGCTCCTGCGTGAGCTAGGTTACCAGCTAAAGGAGGGTAAACAGTTCAACCTGTTCTG-
CCCCAGCTTCAATTATGGATGATGCTAGAAGAAGTAGAAAAGAGGGAGGGAGAAGTCAGAAGCTTATAT-
TATTTATCCGAGGGAATGCTATATCTGGGGACCAATTATTAGAGGGACAAGTCAATTACCAATCCTCCAAT-
TATAATTGGCATTACTATAAAGAAAATTATTACGAATGCATGTGCGGTTACAATTACATTATAAAATTTGATC-
GTCTCCGAGTAGGGTCCAGGTTGGCCAGTTGAGCAGCGATTAGTAAGCTTAGGGCAGTTCCTACTATG-
CCTGCTCAAGCACCATAATAGTAAAT



XYSY2

GAATAAATGTTGATATAGGATGGGATCCCCGCCTCTGCTGGGTCGAAGAAAGTTGTATTTAAATTC-
GGTCTGTTAGTAGTATTGTAATTCGGCTGCTAGTACAGGAAGTGAGAGAAGTAATAATACAGCAGT-
GATTAAGACAGATCATACAATAAAGGGGTTTGATACTGTGATATAGCAGGAGGTTTTATATTAATA-
ATTGTTGTGATAAAGTTAATTGCTCCTAAAATTGAGGAGACACCTGCTAAGTGTAAAGAGAAAAATGTTA-
AGTCTACTGAAGCTCCTGCGTGAGCTAGGTTACCAGCTAAAGGAGGGTAAACAGTTCAACCTGTTCTG-
CCCCAGCTTCAATTATGGATGATGCTAGAAGAAGTAGAAAAGAGGGAGGGAGAAGTCAGAAGCTTATAT-
TATTTATCCGAGGGAATGCTATATCTGGGGACCAATTATTAGAGGGACAAGTCAATTACCAATCCTCCAAT-
TATAATTGGCATTACTATAAAGAAAATTATTACGAATGCATGTGCGGTTACAATTACATTATAAAATTTGATC-
GTCTCCGAGTAGGGTCCAGGTTGGCCAGTTGAGCAGCGATTAGTAAGCTTAGGGCAGTTCCTACTATG-
CCTGCTCAAGCACCATAATAGTAAATACAAAGGTTG



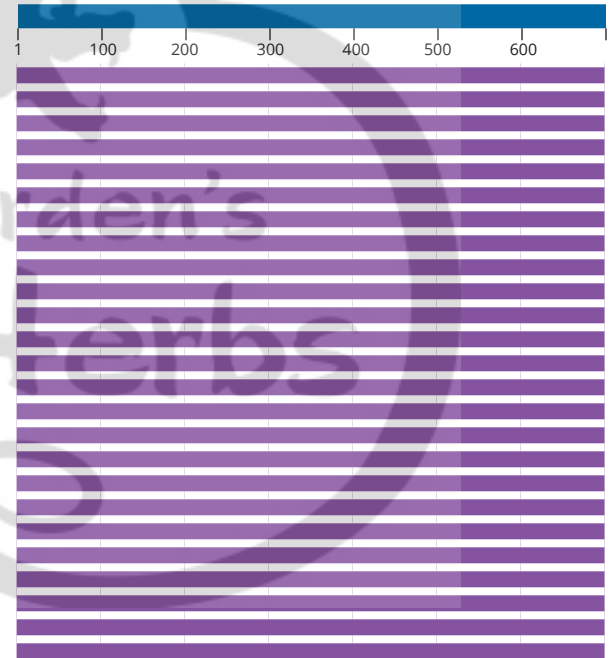
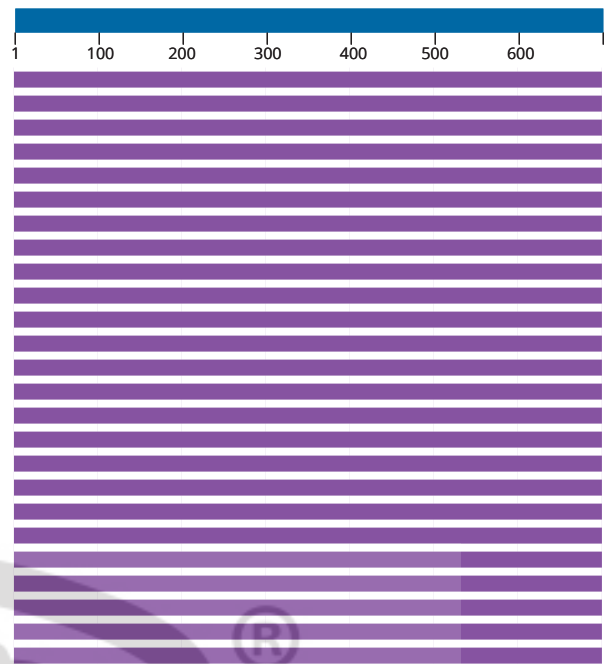
Conclusion: Through the processing and analysis of the sample and data, we identified the closest species of the sample to the Latin name ***Cervus Nippon***, which identified the highest similarity species as Deer Antler with a maximum similarity of 99.67%.



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Alignment Scores

<40 40-50 50-80 80-200 ≥200



99.7%

99.67% similarity to species:

Cervus Nippon