

# Informatica Biomedica

## lezione16

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# Informatica Biomedica: Lezione 16

Bioinformatics Archives and Information Retrieval  
The archives

fonte essenziale: A- M- Lesk, [Introduction to Bioinformatics](#),  
Oxford University Press, 3rd Ed- (2008)

# Contents

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## Bioinformatics databases

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- ▶ **annotating** the compilation of biological information in a form most useful to support research-



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## Nucleic acid sequence databases

The nucleic acid sequence databases, as distributed, are collections of entries.

Each entry has the form of a text file containing data and annotations for a single contiguous sequence-

Many entries are assembled from several published papers reporting overlapping fragments of a complete sequence.

## Entries have a life history

Because of the desire on the part of the user community for rapid access to data, new entries are made available before completion of annotation and checking-

Entries mature through the classes:

Unannotated → Preliminary → Unreviewed → Standard

Rarely, an entry **dies**; a few have been removed when they are determined to be erroneous.

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- ▶ Surveys of molecular biology databases and servers

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