

## MCBG2039A - CELLS AND ORGANISMS

INTRODUCTORY MICROBIOLOGY



## THE MICROBIAL WORLD

CHAPTER I



# MOLECULAR BIOLOGY AND THE UNITY AND DIVERSITY OF LIFE

#### 1.14 - Molecular Basis of Life

1.15 - Woese and the Tree of Life



#### MOLECULAR BASIS OF LIFE

- Ability to grow bacteria rapidly under controlled conditions makes them excellent models for fundamental nature of life
- Led to foundations of molecular biology, genetics, and biochemistry
- Metabolic model chemistry: Certain macromolecules and reactions are universal

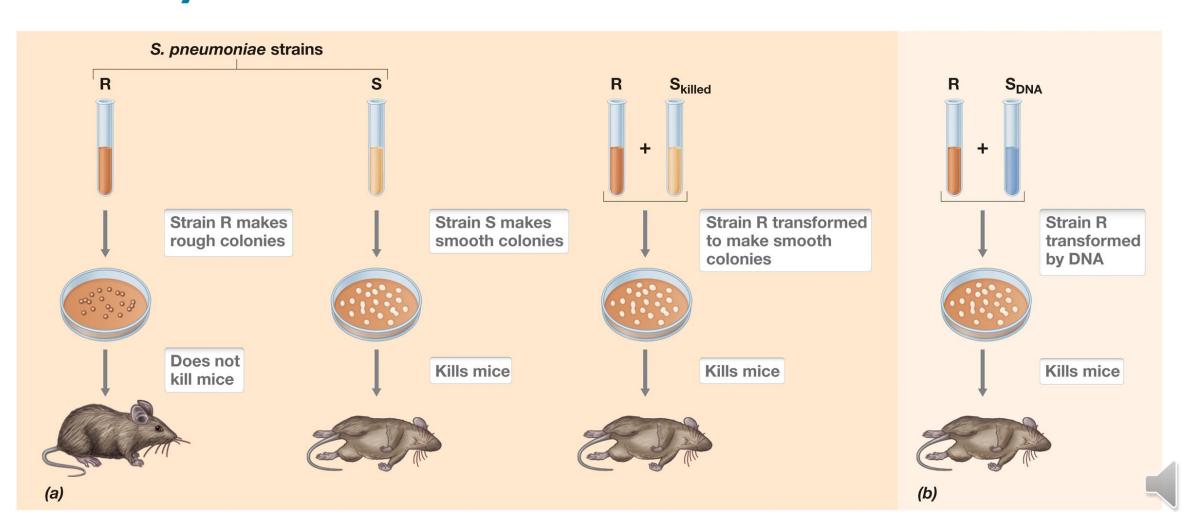


#### MOLECULAR BASIS OF LIFE

- Cracking the Code of Life
  - Bacteria can transfer DNA amongst each other horizontal gene transfer
    - Frederick Griffith, Streptococcus pneumoniae and transformation
    - Avery-MacLeod-McCarty experiment (1944)
    - James Watson, Francis Crick, Rosalind Franklin: structure of DNA
    - Emile Zuckerkandl and Linus Pauling: molecular sequences and evolutionary relationships



# Early Evidence That DNA Is the Molecular Basis of Heredity



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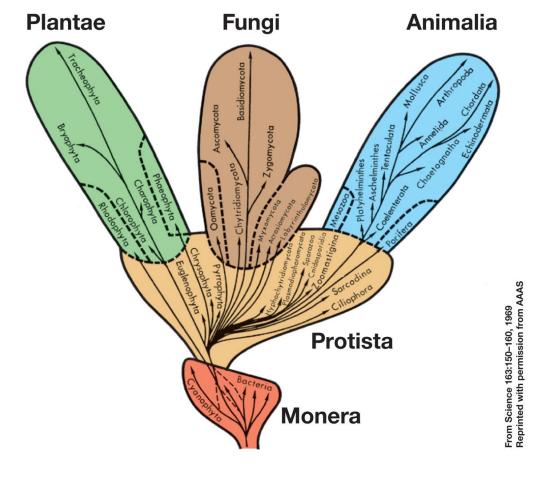
# MOLECULAR BIOLOGY AND THE UNITY AND DIVERSITY OF LIFE

1.14 - Molecular Basis of Life

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- Ribosomal RNA (rNA) present in all cells made it possible to build the first tree of life
- First attempt by Ernst Haeckel (1866)
- Robert Whitaker (1969) Five kingdom classification scheme

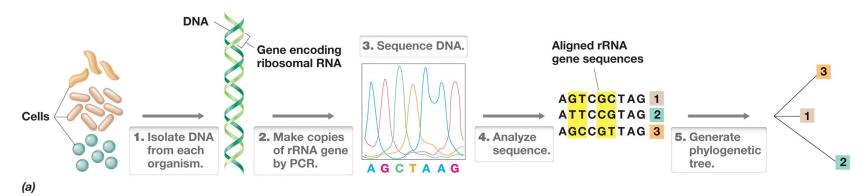


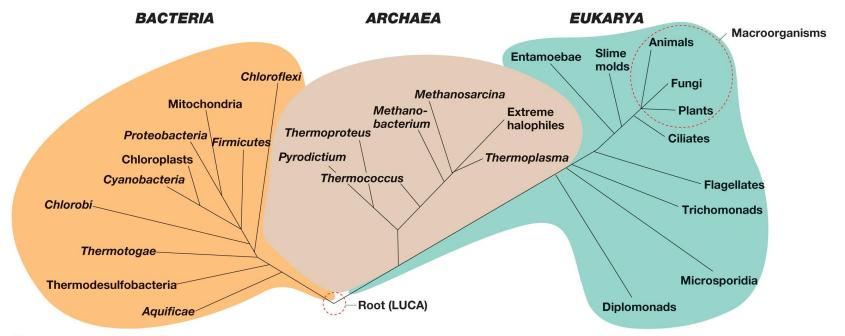


- Carl Woese (1928–2012) realized rRNA sequences could be used to infer evolutionary relationships
  - discovered rRNA from methanogens distinct from Bacteria and Eukarya
  - named new group Archaea
  - found evolutionary relationships between all cells could be revealed by rRNA analysis



# **Evolutionary Relationships and the Phylogenetic Tree of Life**



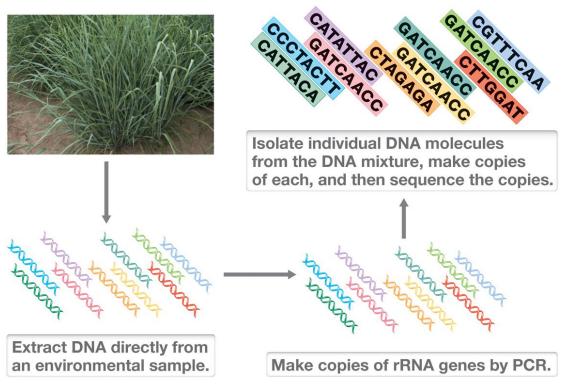




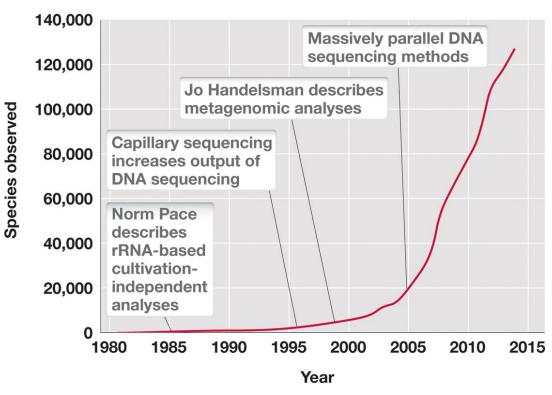
- Phylogenetic tree: depicts phylogeny (evolutionary history) of all cells
  - clearly shows three domains
  - Root is LUCA (last universal common ancestor)
  - evolution along two paths to form Bacteria and Archaea
  - Archaea later diverged to distinguish Eukarya from Archaea
- Cultivation-independent methods show most microbes have not been cultured yet



# Analysis of Environmental rRNA Genes Leads to Discovery of New Microbial Species



(a) Cultivation-independent analysis of rRNA genes



(b) Revealing the extent of microbial diversity



- DNA sequencing technology improvements have improved ability to study Bacteria and Archaea
  - Can sequence entire genomes
  - Metagenomics: microbial genomes/fragments can be recovered from environmental DNA samples

