

Comparing Prokaryotic and Eukaryotic Cells

Basic chemical components/elements of a cell

CHOPKNS CaFe (its) Mg (ood)

TABLE 2.2 Chemical comp	osition of a prokaryotic cell*	Rem: 70-85% Water	
Molecule	Percent of dry weight*	Molecules per cell	Different kinds
Total macromolecules	96	24,610,000	~2500
Protein	55	2,350,000	~1850
Polysaccharide	5	4.300	2' 🗭
Lipid	9.1	22,000,000	44
Lipopolysaccharide	3.4	1,430,000	1
DNA	3.1	2.1	1
8 RNA	20.5	255,500	€660> €
Total monomers	3.0		~350
Amino acids and precursors	0.5		~100
Sugars and precursors	2		~50
Nucleotides and precursors	0.5		~200
Inorganic ions	1		18
Total	100%		
Total a Data from Neidhardt, F. C., et al. (eds. American Society for Microbiology, Was b Dry weight of an actively growing cei c Assuming peptidoglycan and glycog d There are several classes of phosphol between species and because of differe), 1996. Escherichis coli and Salmonella typ hingtor, DC. I of E. coli $\approx 2.8 \times 10^{-13}$ g; total weight m to be the major polysaccharides press pids, each of which exists in many kind	(70% water) = 9.5 × 10 ⁻¹³ g. ent.	
L	50% 10%	Wall 10–20%	•

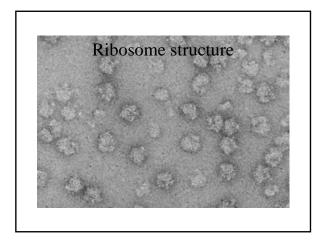
Take Home Message:

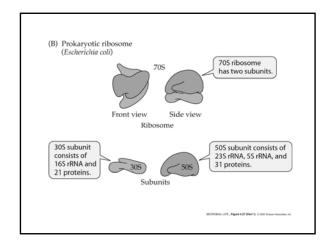
Proteins are #1 by weight

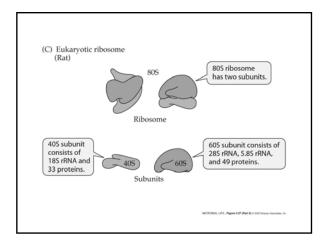
Lipids are #1 by number

Peptidoglycan is 1 jumbo molecule

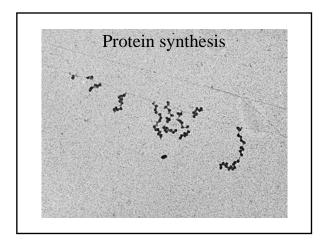
Comparing Prokaryotic and Eukaryotic Cells Classification of prokaryotic cellular features: Invariant (or common to all) • Ribosomes: Sites for protein synthesis – aka the grand translators • Cell Membranes: The barrier between order and chaos • Nucleoid Region: Curator of the Information







Property	Prokaryote	Eukaryote
Overall size	70S	805
Small subunit	305	40S
Number of proteins	~21	~30
RNA size (number of bases)	16S (1500)	18S (2300)
Large subunit	50S	60S
Number of proteins	~34	~50
RNA size	235 (2900)	285 (4200)
(number of bases)	5S (120)	5.8S (160)
		5S (120)



Importance of a Molecular Biological Approach

- **Traditional culturing** techniques isolate ~1% of the total bacteria in marine ecosystems, thereby severely underestimating diversity and community structure.
- Because nutrient-rich **culture media** have been historically used during enrichment procedures, bacteria which may be dominant in natural communities are selected against in favor of copiotrophic (weedy) bacteria.
- SSU rRNAs and their respective genes are excellent descriptors of microbial taxa based on phylogeny.

