Bioinformatics Lab.

CHAPTER I TERMS 1

DNA (deoxyribonucleic acid)

RNA (ribo-)

Adenine, Cytosine, Guanine, Thymine

Gene Genome

Phosphodiester bond

5'3'

Upstream, Downstream

Template

Complement sequence

Antiparallel

Purine, Pyrimidine

Transition, Transversion

Central dogma
Transcription
Translation

Polymerase Ribosome

Promotor

Positive regulation Negative regulation

Gene code Triple code

Codon

Start codon: AUG

Stop codon: UAA, UGA, UAG

Reading frame Degeneracy

Amino acid structure

Polar vs. Nonpolar

Positively charged vs. Negatively charged

Open reading frame (ORF)

Intron and Exon

Splicing

GT-AG rule

Alternative splicing

Backbone and side chain

Amino group and Carboxylic acid group Amino terminus and Carboxy terminus

Restriction enzyme

EcoRI, Hind III

Blunt end

Cohensive/sticky end

Gel electrophoresis (garose, acrylamide)

Blotting

Hybridization Microarray

Cloning vector

cDNA

cDNA library

gDNA library

BAC library

PCR

Sequencing

Maxam-Gilbert method

Sanger method=Chain-termination method

Extra

NGS

Pyrosequencing

454 technology GS-FLX

Sorexa/Illumina HiSeq2000