Genetics Engineering (Zoo-455)

Human Genome Project Lecture-7

Introduction to the human genome project:

- The Human Genome Project (HGP) was the international, collaborative research program whose goal was the complete mapping and understanding of all the genes of human beings.
- □ HGP has three major goals:
 - 1. Determine the nucleotide sequences of all DNA in the human genome.
 - 2. Identify the location and sequence of every human gene.
 - 3. Producing linkage maps, through which inherited traits (such as those for genetic disease) can be tracked over generations.
 - 4. comparisons of genomes within and across species.



Time line of the human genome project:



□ The Human Genome Project officially begins in 1990 and completes in 2003.

Results of the HGP:

- □ HGP has sequenced all of the DNA base pairs of human chromosomes.
- □ HGP has sequenced 3 billion base pairs (ATGC).
- □ HGP covers about 99% of the human genome's gene-containing regions.
- □ The identification of approximately 20.000 protein-coding genes in human.
- Many of protein-coding genes produce more than one protein product by alternative splicing of the primary transcript of the gene. Each ORFs produce more than 2 to 3 different proteins.
- □ The identification of more than 3 million human genetic variations, called single nucleotide polymorphisms (SNPs).
- □ The generation of full-length complementary DNAs (cDNAs) for more than 70% of known human and mouse genes.
- □ The identification of the mouse genome sequence, published in December 2002.
- The identification of the rat genome sequence, produced in November 2002.

Exploring the human genome:



2003



Identification of common genetic variation in 270 individuals from 4 populations (Europe, China, Japan, and Nigeria)

2008

1000 Genomes

A Deep Catalog of Human Genetic Variation



Whole genome sequencing and complete description of human genetic diversity in >1000 individuals from multiple world populations.

What is a genome?

- Genome: All the DNA for a human.
- Gene: Specific sequences on DNA that can be expressed into proteins.
- **Human genome contains:**
 - Nucleus: 3 billion base pairs (ATGC) packaged in 46 chromosome.
 - **Mitochondrion**: 16600 base pairs (ATGC).
 - 20.000 protein-coding genes.
 - 4 millions DNA variants/individual.
 - 50% is high copy number repeats.
 - About 10% is transcribed (made into RNA).
 - Only 1.5% codes for protein (98.5% non-coding DNA).
- SNP = Single nucleotide polymorphism: DNA sequence variation in which a single nucleotide A, T, C or G differs between members of the same species.
- □ SNP occurs commonly within a population (> 1%).



How to read the genome?

- 1. Genotyping: Process of determining genetic differences between individuals by using a set of markers
- 2. Sequencing: Process of determining the full nucleotide order of a DNA sequence
- Technology allows the study and comparison of both genes and proteins.
- Bioinformatics: is the use of computer databases to organize and analyze biological data.
- **Proteomics**: is the study and comparison of proteins



