

Basic Terminology

Genetics: The study of genes and their roles in inheritance

Genetic testing: Examines changes in an individual gene and/or set of genes that may account for the likelihood of a particular disease process

Genomics: The study of the complete set of a person's DNA/genes. Genomics also examines how genes interact with each other and how, or if, they are affected by environmental factors.

Genomic testing: Assesses an individual's genome (all of a person's genes, rather than a specific gene or set of genes)

Diploid: Containing two complete sets of chromosomes, one from each parent

Genome: 6 billion base pairs of DNA distributed unequally across 46 chromosomes (23 pairs) in a diploid genome

Exome: Part of the genome that includes just the DNA exons (about 1.5% of DNA in humans)

Exons: Sequences of DNA that encode proteins

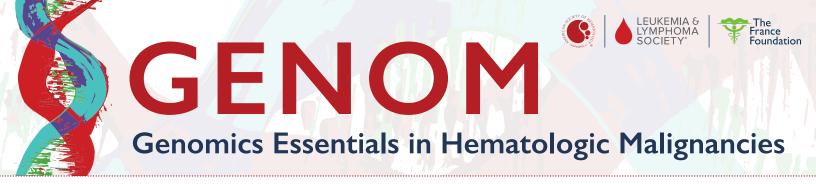
Transcriptome: Gene expression profile; all the mRNAs transcribed from the genes of a cell at a given time

Epigenetics: Modifications in gene expression due to environmental factors and/or behaviors that effect how proteins and cells function but not changes in DNA sequences

Somatic DNA: DNA in cells that are not involved in reproduction and have a full complement of chromosomes (23 pairs in humans)

Germline DNA: DNA in cells involved in reproduction (ie, sperm and eggs) that have half as many chromosomes as somatic cells (23 individual chromosomes in humans)

Germline susceptibility: Genes a person is born with that may confer increased risk of disease



Biomarkers

Biomarker: A biological molecule found in blood, other body fluids, or tissues that is a sign of a normal or abnormal process, or of a condition or disease. A biomarker may be used to see how well the body responds to a treatment for a disease or condition. Also called "molecular marker" and "signature molecule."

Biomarker testing: As applied to genomics, analysis of single or multiple somatic genetic events associated with disease outcome including response to therapy; could be at DNA, RNA, or molecular level

Diagnostic biomarker: Associated with the diagnosis of a particular disease

Prognostic biomarker: Associated with the overall disease outcome regardless of treatment

Predictive biomarker: Associated with either a favorable or unfavorable outcome from a particular treatment

Actionable biomarker: Used to make treatment decisions

Single nucleotide variant (SNV): Any difference in a single DNA base pair irrespective of how often it occurs

Single nucleotide polymorphism (SNP): A single base pair difference considered normal common genetic variation

DNA Alterations at the Chromosomal Level

Cytogenetics: Chromosomes in cells arrested in metaphase are stained through Giemsa banding; shows large gains, losses, and/or translocations of the entire chromosome complement

Karyotype: Image of normal human chromosomes lined up in pairs (include image)

Inversions: A portion of a chromosome which has broken off, turned upside down, and reattached

Trisomy: The presence of a third copy of a particular chromosome

Monosomy: The absence of one copy of a particular chromosome

Insertions or deletions: Changes the number of DNA bases in a gene by adding or removing a piece of DNA

Amplification: Small or large parts of chromosomes which have been duplicated or multiplied

Translocations: A piece of one chromosome which has broken off and attached to another chromosome



Variations/Mutations

Variation/mutation: A change in the DNA sequence

Point mutation: A change within a gene in which one base pair in the original DNA sequence is altered

Missense mutation: A point mutation (usually) that results in a change in amino acid

Deletion/insertion/duplication: A few bases may be deleted, inserted, or duplicated

Triplet repeats: Sequence of 3 nucleotides that is repeated multiple times in DNA. This is a mechanism of inherited genetic disease that often occurs with neurodegenerative diseases.

Frameshift mutation: Insertion or deletion of bases that results in a shift to the reading frame of the DNA resulting in changes in AAs and/or a premature stop codon

Somatic (acquired) mutations: Occur after conception, during a person's life, and are only present in certain cells

Germline (hereditary) mutations: Inherited from a parent. They occur before conception and are present throughout a person's life in almost every cell in the body.

Pathologic Variants

Activating mutations: Turns on the function of a gene that may normally be carefully regulated

Fusions: Portions of DNA that are attached to each other that should not be

Clonal mutations: In every cell of the cancer

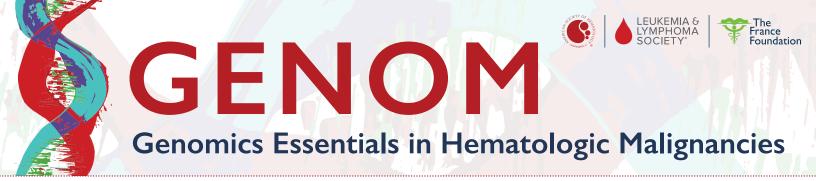
Subclonal mutations: Only in a subset of cells of the cancer

Driver mutations: Thought to be key in promoting the initiation and/or survival of the cancer

Passenger mutations: Random mutations found within the cell that do not contribute directly to promoting the initiation and/or survival of the cancer; may be acquired prior to driver mutation

Variant of undetermined significance (VUS): A genetic change in the germline of unknown impact on the function of the gene

Pathogenic mutations: Directly contributes to disease development



Pathologic Variants (cont.)

Actionable mutations: There is a specific therapy indicated

Clonal hematopoiesis of indeterminate potential (CHIP): Somatic mutation that occurs with aging in the normal blood cells; associated with increased subsequent risk of blood cancer and cardiovascular disease

T cell clonality: V(d)J rearrangements; unique T cell receptor used to diagnose a T cell malignancy

B cell clonality: IGH Clonality; unique B cell receptor sequence for a given B cell cancer that can be used to determine measurable residual disease (MRD, see Testing for definition) or for prognostic purposes in CLL

Testing

FISH (fluorescent in situ hybridization): Uses a fluorescent probe to detect very small gains, losses, or translocations within a cell; FISH testing is able to detect changes that are too small to be seen on basic cytogenetic tests

Immunohistochemistry (IHC): Uses an antibody that is tagged with a dye or fluorescence against a cellular protein to study over- or under-expression of certain proteins

Flow cytometry: Cells in suspension with fluorescent-tagged antibodies are passed through a laser beam that activates fluorescence when the antibody target is present; used to detect whether multiple proteins are co-expressed on the same cell and to determine cell phenotype, which in turn strongly contributes to determining the diagnosis

Sequencing: Laboratory tests that examine the exact order of the four bases of DNA

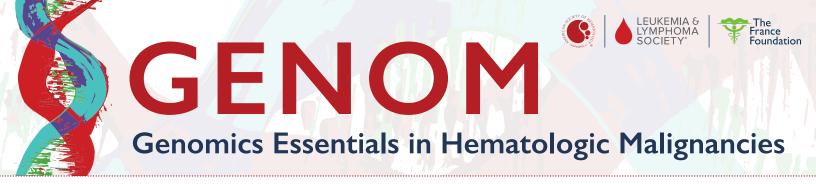
Sanger sequencing: Low throughput sequencing often used to characterize one gene at a time

Next generation sequencing (NGS): High throughput sequencing used to characterize many genes or an entire genome at once; higher volume and more rapid than traditional Sanger sequencing

Single cell sequencing: Sequencing of RNA or DNA at single cell resolution

Cell-free sequencing (liquid biopsy): Sequencing of DNA in plasma

Polymerase chain reaction (PCR): Specific DNA probes are used to amplify a sequence to detect the presence or absence of a sequence of DNA; can detect very small amounts of DNA



Testing (cont.)

Genome editing (e.g., CRISPR/Cas9): Approach to modifying a gene of interest by introducing a specific mutation into the endogenous gene

Precision medicine: A form of medicine that uses information about a person's own genes or proteins to prevent, diagnose, or treat disease. In cancer, precision medicine uses specific information about a person's tumor to help make a diagnosis, plan treatment, determine treatment effectiveness, or make a prognosis.

Measurable (previously minimal) residual disease (MRD): Describes detection of low levels of cancer cells after therapy; can be used to test for specific mutations. Definition has changed to measurable because minimal is dependent on what is measurable given the current state of technology.