

# Etiology of chromosomal aberrations

## Chromosomal aberrations

1. Abnormalities in chromosome number
  1. aneuploidy
    - monosomy
    - trisomy (or tetrasomy, pentasomy...)
  2. polyploidy
    - triploidy
    - tetraploidy,...
2. Abnormalities in chromosome structure
  1. balanced
    - translocation
    - inversion
    - insertion
  2. unbalanced
    - deletion (incl. ring chromosome)
    - duplication
    - isochromosome

## Etiology of congenital chromosomal aberrations

- Origin of aneuploidies and polyploidies (see question No. 34 – Abnormalities in chromosome number, their causes and clinical presentations in man)
- Origin of structural aberrations:
  - chromosome breaks and rearrangements during gamete formation (in meiosis) or in pre-gametic mitotic divisions of gonadal cells – resulting in stable products having one centromere and two telomeres
    - one centromere is necessary for regular segregation in mitosis (unstable dicentric chromosomes undergo secondary rearrangements)
    - telomeres maintain the integrity of the ends of linear chromosome structure (in deleted chromosomes new telomeres are added by \*\*telomere synthesis or by mechanism of telomere capture)
  - causes of spontaneous breaks – see below (external effects)

## Etiology of acquired chromosomal aberrations

(= chromosome breaks and rearrangements during mitotic divisions of somatic cells)

External effects (physical, chemical, biological)

- random environmental factors – spontaneous breaks (UV light, ionizing radiation – cosmic rays, medical radiation (X-rays), drugs, viral infections)
- professional exposition (mutagens: chemicals – alkylating agents, intercalation substances...; radiation)
- oncological treatment (chemotherapy, radiotherapy)

## Hereditary syndromes of chromosome instability

- congenital defects of repair mechanisms - mostly double-strand DNA breaks repair
- rare genetic disorders with AR inheritance, higher predisposition to cancer development
- higher level of chromosome breaks and rearrangements detected in cytogenetic analysis
  - ataxia teleangiectasia (defect of ATM gene - important for double-strand DNA breaks repair)
  - xeroderma pigmentosum (defect of nucleotide excision repair)
  - Bloom syndrome (extreme genome instability, high level of sister chromatid exchanges - SCEs, high frequency of mutations)
  - Fanconi anemia
  - Nijmegen breakage syndrome

Methods of analysis of acquired chromosomal aberrations (see question No. 31 – Methods of chromosomal examination)