

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)