NEOPLASIA



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2- Radiation and Physical Carcinogens:

- Sources:

-UV rays of sunlight, X-rays, radioactive isotopes, & nuclear fission (Bomb or reactors).

□U-V light:

- Effect depends on the intensity of exposure & quantity of melanin.
- At greatest risk are fair-skinned people who live in areas that receive a great deal of sunlight.

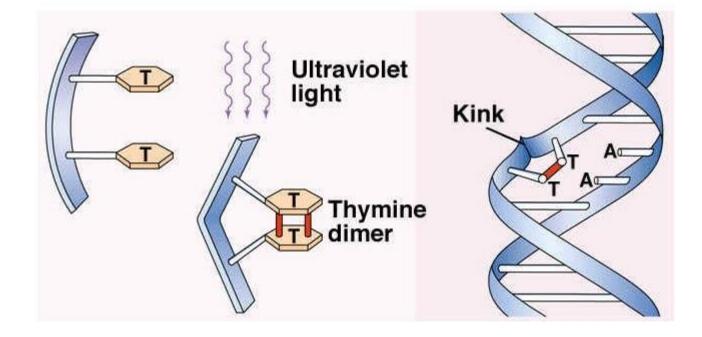
Skin Cancers, including:

- *Squamous Cell CA
- *Basal Cell CA
- *Melanoma

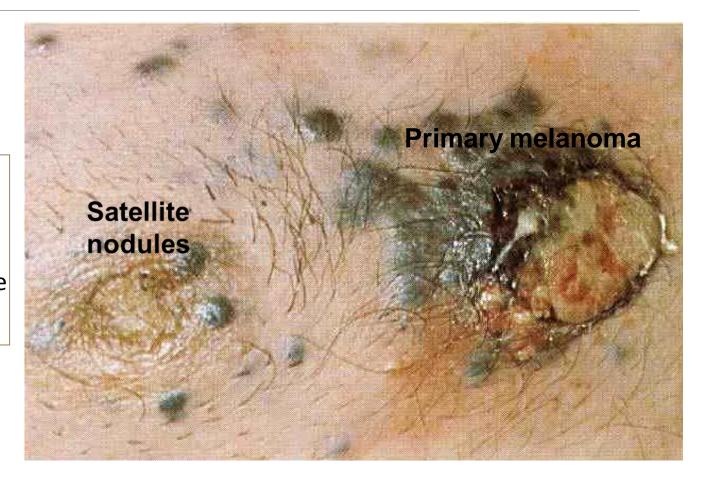
- Damage DNA by forming pyrimidine

dimers. This type of DNA damage is repaired by the nucleotide excision repair pathway. With extensive exposure to UV light, the repair systems may be overwhelmed, and skin cancer results.

Pyrimidine Dimer



Chest wall: Malignant melanoma with local spread: Many small metastatic satellite nodules have formed in the tissue around the pigmented primary melanoma.



□Ionizing Radiation:

- Explosions: 个 Leukemia, Breast, colon, thyroid, lung CA.
- Therapeutic radiation exposure of the head and neck $\rightarrow \uparrow$ Thyroid CA, Leukemia.
- Miners of radioactive elements (e. g uranium) have suffered a ten-fold increased incidence of lung cancer.
- Many of the pioneers in the development of X-rays develop skin cancers.

Mechanism: Free radical injury → chromosome breakage, translocations & less frequent point mutations.

□ Asbestos fiber inhalation: Mesothelioma & Lung CA

3- Microbial Carcinogens:

-Oncogenic DNA Viruses.

- 1. HPV
- 2. EBV
- 3. Hepatitis B. virus.
- 4. HHV-8 (Kaposi sarcoma herpes virus in AIDS)
- 5. Polyomavirus called Merkel cell virus

-Oncogenic RNA Viruses.

- HTLV-1
- H. pylori

1- HPV-Human Papilloma Virus:

- Several genetically distinct types:
 - **Types 1, 2, 4 & 7** → Benign squamous papilloma (wart)
 - Low-risk types (6, 11) → Genital Squamous Cell Papilloma (wart)
 - High-risk types (16, 18) \rightarrow Squamous Cell Carcinoma in cervix, vulva, perianal
 - Cervical severe dysplasia, SCCa in situ.
 - Oropharyngeal Carcinoma.

Mode of action:

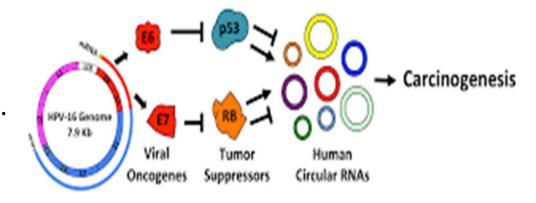
- HPV has transforming early genes (E6, E7), each of which has several activities that are pro-oncogenic and inactivate suppressor genes:

E6 protein binds & degrades **p53** → no apoptosis

E7 protein binds to **Rb** → releasing E2F transcription effect → activates cyclins & inhibit CDKIs--- promoting progression through the cell cycle.

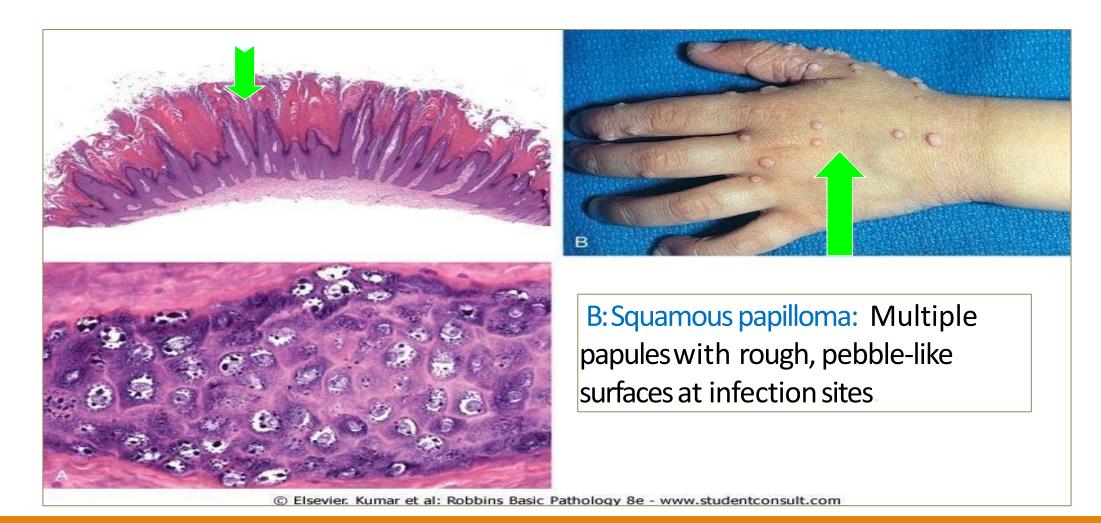
- E7 and E6 of high-risk types higher affinity for Rb and P53.
- In benign warts, the HPV genome is maintained in a nonintegrated form, while in cancers, the HPV genome is randomly integrated into the host genome.
- Integration interrupts a negative regulatory region in the viral DNA, resulting in overexpression of the E6 and E7 oncoproteins → Cell proliferation.

Regulation of Circular RNAs by HPV Oncogenes



A: Squamous papilloma:

Symmetrical papillary epidermal proliferation(top). Histology shows nuclear pallor, prominent keratohyalin granules

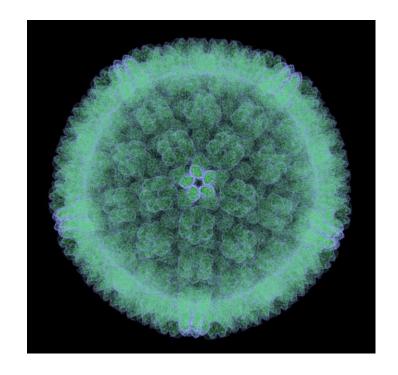


2- EBV - Epstein Barr Virus

- A member of the herpesvirus family.

Associated with:

- Burkitt Lymphoma
- Other B-cell Lymphoma
- Hodgkin lymphoma (Subset)
- Nasopharyngeal Carcinoma



Mode of action in Burkitt lymphoma:

- In endemic cases, EBV is identified in tumor cells.

EBV (has LMP1 oncogene) enters B cells ———
 Induces B cell proliferation

Prevents apoptosis by activating BCL2.

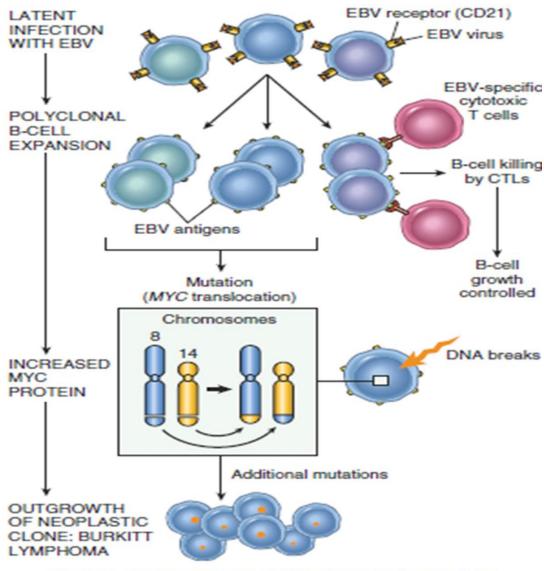


Fig. 6.34 Possible evolution of EBV-induced Burkitt lymphoma.

But...

-Endemic infection & Malnutrition may play a role in \downarrow immunity (Lost T cell control)---So a small number of the EBV-infected B cells survive and with the acquisition of specific mutations, most notably \Longrightarrow Dysregulation of c-MYC by translocation t(8;14) \Longrightarrow BURKITT Lymphoma

In nasopharyngeal carcinoma:

- In contrast to Burkitt lymphoma, 100% of nasopharyngeal carcinomas obtained from all parts of the world contain EBV.
- The uniform association of EBV with nasopharyngeal carcinoma suggests that EBV has a central role in the genesis of the tumor, but the restricted geographic distribution indicates that genetic or environmental cofactors, or both, also contribute to tumor development.
- LMP1 is expressed on epithelial cells activating cell proliferation

3- Hepatitis B & Hepatitis C Viruses

- About 70% to 85% of hepatocellular carcinomas are caused by HBV or HCV.
- The HBV and HCV genomes do not encode any viral oncoproteins
- -Multifactorial oncogenic effect but mainly Immunologically mediated chronic inflammation with hepatocyte death, leading to regeneration and genomic damage.

The oncogenic effect of HBV seems to be:

- (1) First, by causing chronic liver cell injury & accompanying regeneration, HBV predisposes the cells to mutations, caused possibly by environmental agents
- (2) virus-induced gene damage in regenerating liver cells may set the stage for multistep carcinogenesis.

In addition:

The HBV contains HBx gene, may more directly promote the development of cancer:

Acts as growth-promoting gene

HBx inactivates suppressor functions, such as TP53.

The HCV (RNA virus) has HCV core protein which may induce proliferation

4- Oncogenic RNA Viruses:

HTLV-1 (Human T-Cell Leukemia Virus Type 1):

- HTLV-1 has a tropism for CD4+ T cells, and T cells are the major target for neoplastic

transformation.

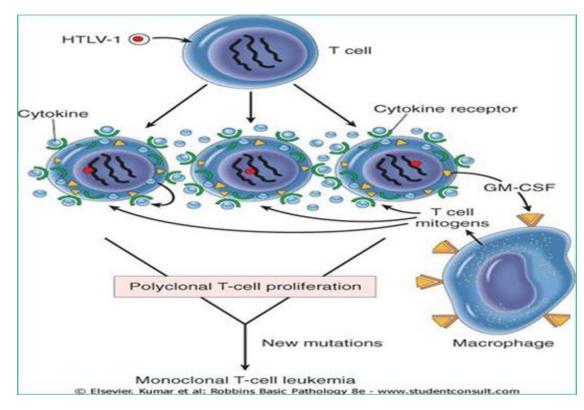
-Induces adult T-cell leukemia/lymphoma (ATLL)

- Transmitted sexually, blood or breast milk.

Mode of action:

Virus **TAX** gene--- TAX protein which:

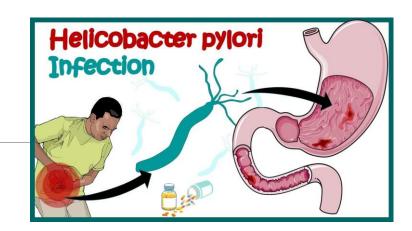
- Increased survival and growth of infected cells.
- Suppresses action of TP53 & CDKI
- Increased genomic instability.



POLYCLONAL \rightarrow MONOCLONAL \rightarrow LEUKEMIA (3-5% of cases - latent period 40-60 yrs).

5- Helicobacter pylori:

- First described as a cause for peptic ulcer.
- Multifactorial etiology in gastric carcinoma & gastric lymphoma



Carcinoma:

Chronic gastritis \rightarrow atrophy \rightarrow intestinal metaplasia \rightarrow dysplasia \rightarrow Gastric Carcinoma

- This sequence occurs in only 3% after a long latent period
- In adenocarcinoma, H. pylori contains Cytotoxic Associated gene A ($Cag\ A$) \rightarrow Cell proliferation

<u>Lymphoma:</u>

Chronic gastritis \rightarrow mucosal lymphoid follicles \rightarrow reactive polyclonal B cells \rightarrow monoclonal B cells \rightarrow MALT lymphoma

Early in the course of the disease, eradicating *H. pylori* with antibiotics causes regression of the lymphoma by removing the antigenic stimulus for T cells.

Carcinogenesis Molecular basis of cancer

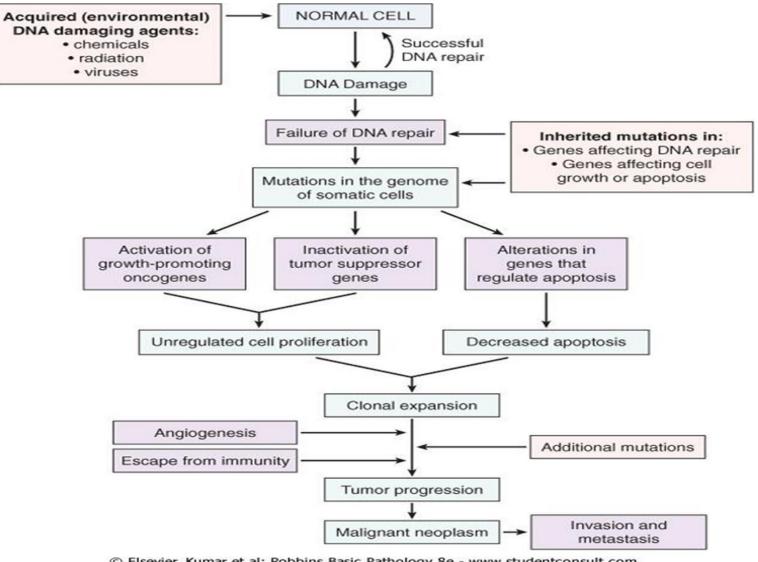
Neoplasms arising from a single clone of cells

Monoclonal proliferation

Cells that are genetically identical to the unit from which they were derived

- Non-lethal (non-killing) genetic damage (or mutation) lies at the heart of carcinogenesis

- Tumors arise from clonal growth of transformed cells that have developed mutations in several classes. of genes:
 - Growth promoting protooncogenes
 - Growth inhibiting tumor suppressor genes
 - Genes regulating apoptosis
 - Genes involved in DNA repair



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Genetic Lesions in Tumors

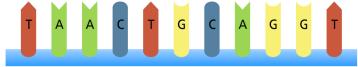
1. Point mutation:

-Change in a single base in a nucleotide sequence (altering amino acid residues) may activate an oncogene or inactivate a tumor suppressor.

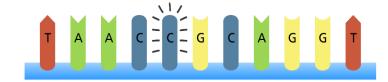
e.g. RAS oncogene.

TP53, tumor suppressor gene.





Point mutation

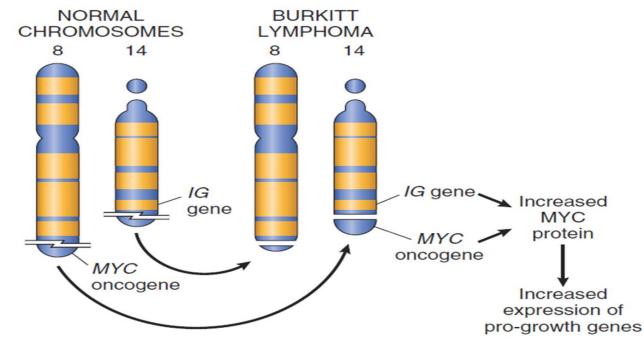


2. Translocation:

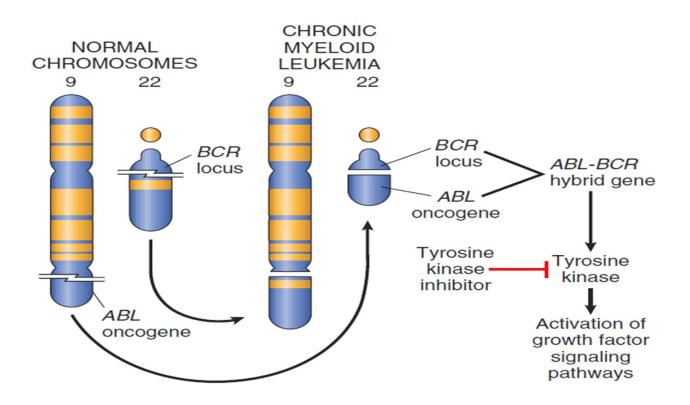
These rearrangements can activate proto-oncogenes in two ways:

1. Result in overexpression of proto-oncogenes by removing them from their normal regulatory elements and placing them under the control of an inappropriate, highly active promoter or enhancer

Burkitt Lymphoma : t (8;14)



- 2. Other oncogenic gene rearrangements create fusion genes encoding novel chimeric proteins.
- Chronic myeloid leukemia :t(9;22)(PHILADELPHIA Chromosome)→ Fusion Gene is produced: BCR-ABL (tyrosine kinase activity)



3. Chromosomal deletions:

-Deletion of specific regions of chromosomes may result in the loss of particular tumor suppressor genes.

e.g. Retinoblastoma, RB gene ch13

4. Gene amplification:

- Gene amplification, with consequent overexpression and hyperactivity of otherwise normal proteins.
- Such amplification may produce several hundred copies of the gene.

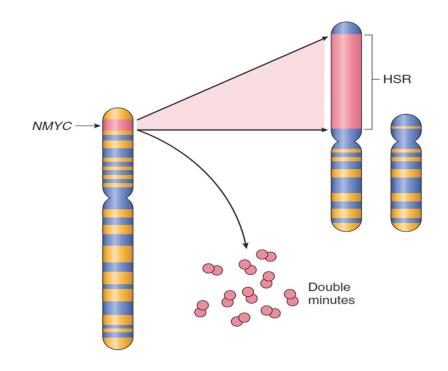
Two mutually exclusive patterns are seen:

- Double minutes: Small fragments of extrachromosomal DNA
- Homogenous staining regions produced by chromosomal segments with various lengths and uniform staining intensity.

Examples:-

- Neuroblastoma: N-MYC

- Breast carcinoma: HER2/Neu



5. Chromosomes loss or gain:

- Change from the normal multiples of 23 (Aneuploidy).

6. Epigenetic changes:

- -Reversible, heritable changes in gene expression that occur without mutation.
- Involves posttranslational modifications of histones and DNA methylation
- This may silence tumor suppressor genes & repair genes, leading to carcinogenesis

