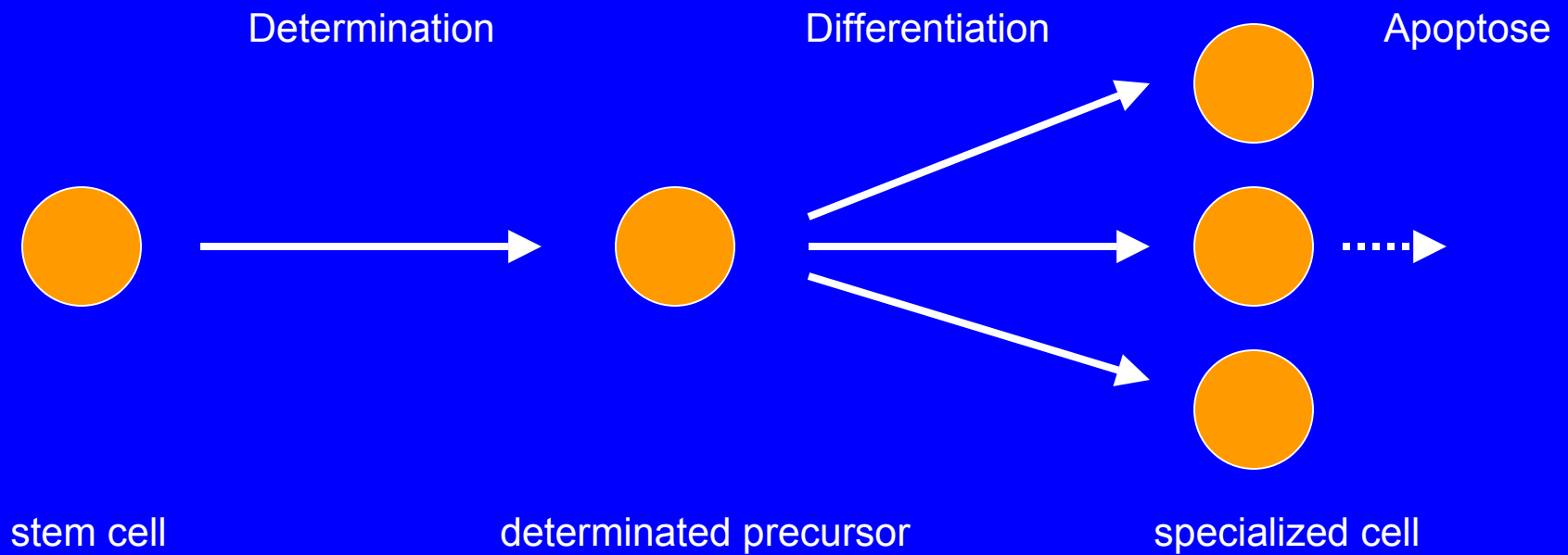


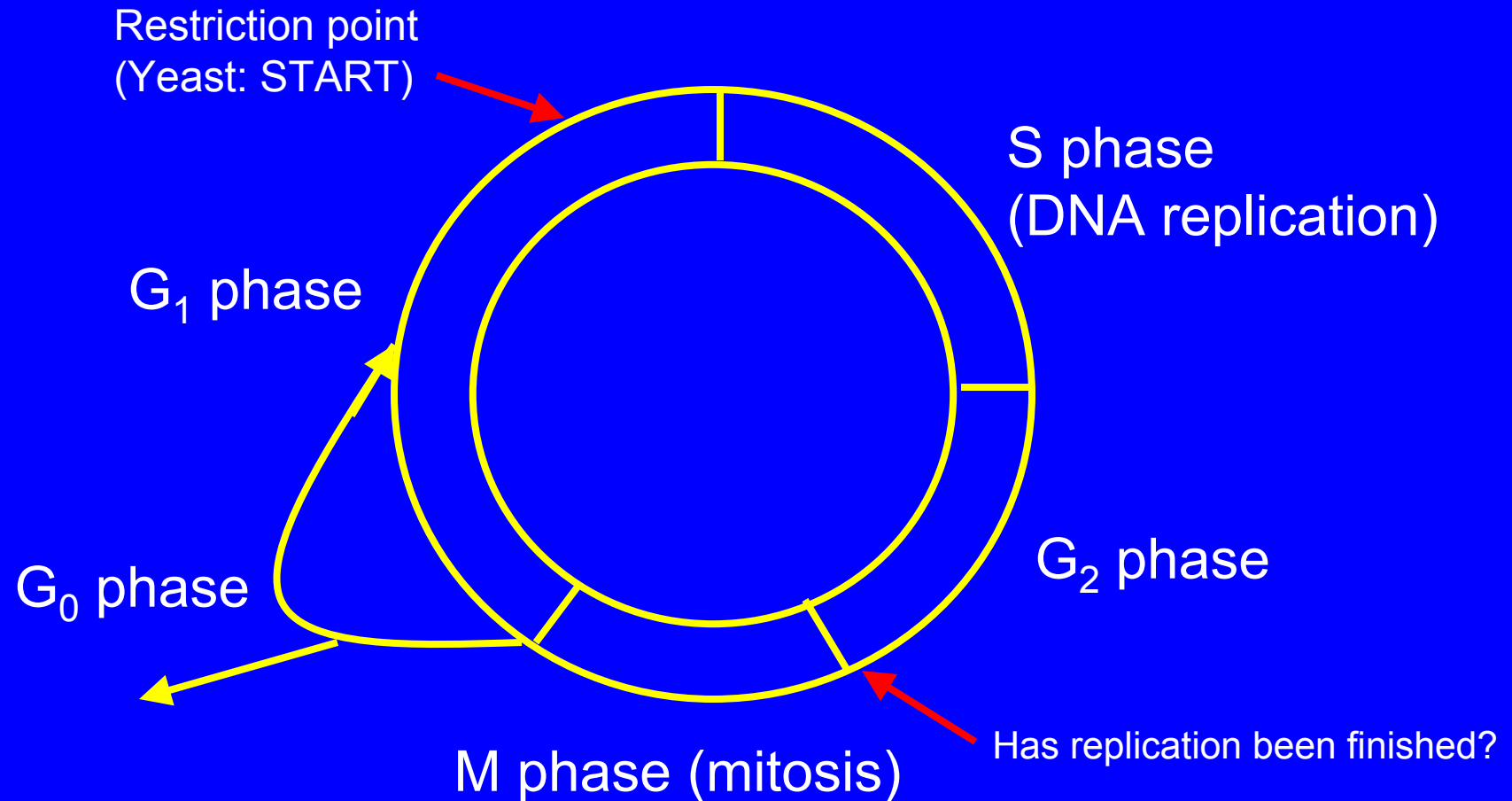
Biology of Cancer

- Developmental Biology: Determination and Differentiation
- Cell Cycle Regulation
- Tumor genes: Proto-Oncogenes, Tumor supressor genes
- Tumor-Progression
- Example for Tumor-Progression: Colon cancer
- Chromosme Translocations in Leukemia

Determination and Differentiation



Cell Cycle

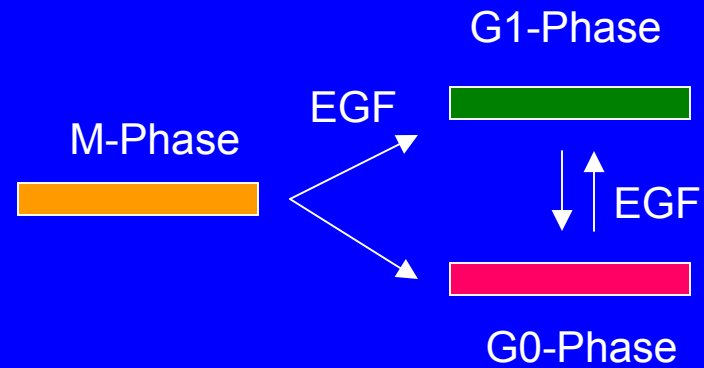
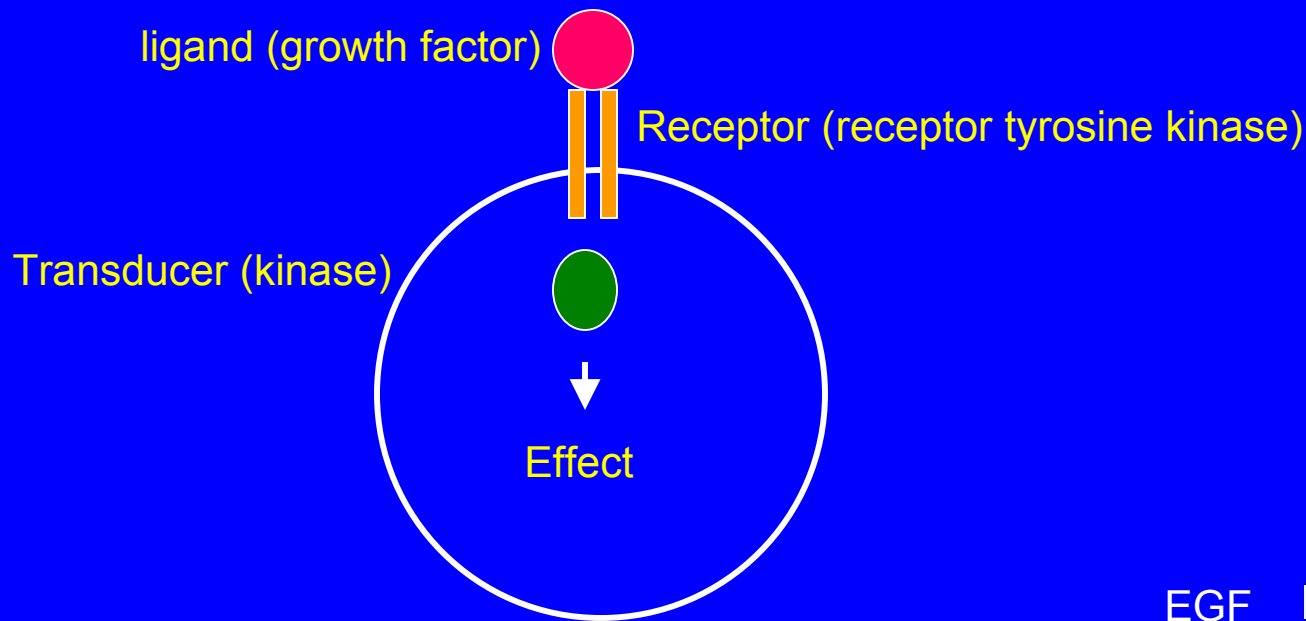


Cell Cycle

Important barriers in the Cell Cycle:

- G1 → S: transcription has to be stopped
- G1 → S: DNA-repair has to be finished
- G2 → M: replication has to be finished
- G0 ↔ G1: is there any need for proliferation

Signal Transduction



EGF: Epidermal Growth Factor

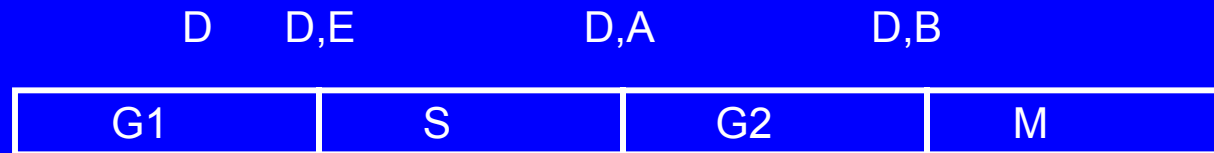
Cell Cycle: Regulation

Cyclins: regulatory proteins, active at specific stages in the cell cycle

CDK: cycline dependent kinases

CDI: CDK inhibitor

Cycline:



CDI (e.g. p27, p21)

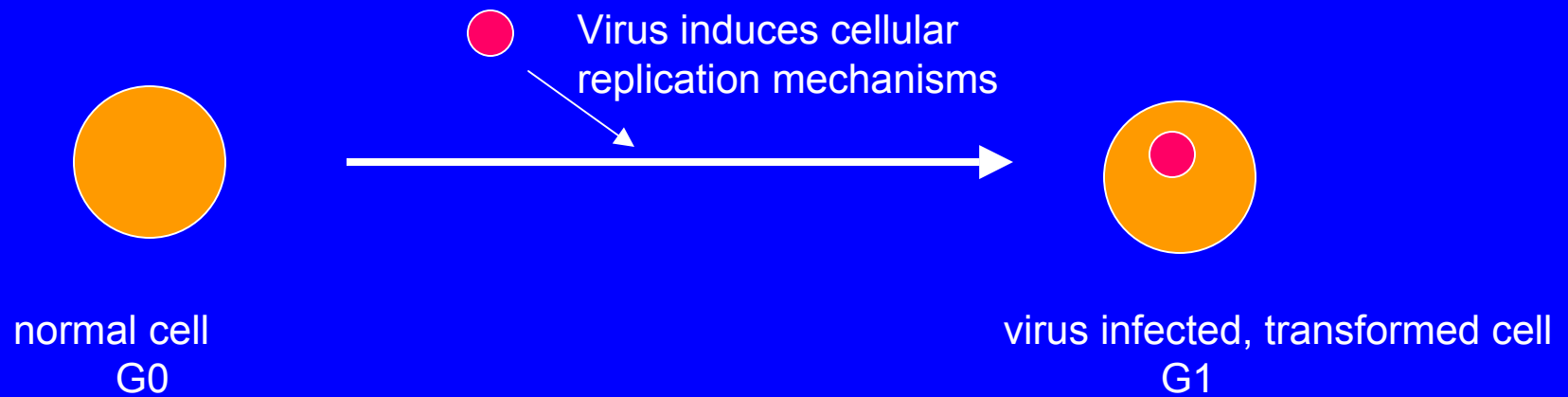
Tumor genes: Knowledge from Viruses

- Viruses need mechanisms for replication
- many use hosts replication machinery
- Consequence: Viruses have to disable the hosts strict regulation mechanisms

Possible mechanisms:

- active interaction of a viral protein in the cells regulatory pathways (e.g. v-ras)
- inactivation of a hosts regulatory factor by binding of a viral protein (e.g. T-Antigen from SV40)

Tumor genes: Knowledge from Viruses



Oncogenes

Oncogenes:

viral proteins which interact with the cellular control mechanisms to overcome the strict regulation of proliferation (v-ras, v-myc, v-abl, ...)

Proto-Oncogenes:

Cellular proteins which correspond to the viral Oncogenes but which are strictly regulated. Mutations in these genes could transform a cell into a tumor cell (c-ras, c-myc, c-abl, ...).

Oncogenes: ras

- ras is a small protein involved in Signal Transduction from cell surface receptors (receptor tyrosine kinases) into the nucleus, where Transcription Factors are activated
- active form: ras-GTP/ inactive form: ras-GDP
- is activated through the so called GDP/GTP-exchange factor(GEF,GRF)
- helps to activate a Serine/Threonine Kinase (raf)
- negatively controlled by an intrinsic phosphatase activity which hydrolyses GTP to GDP (induced by a protein called GAP)

Oncogenes: ras

v-ras (viral form): constitutive activity
=> Oncogene

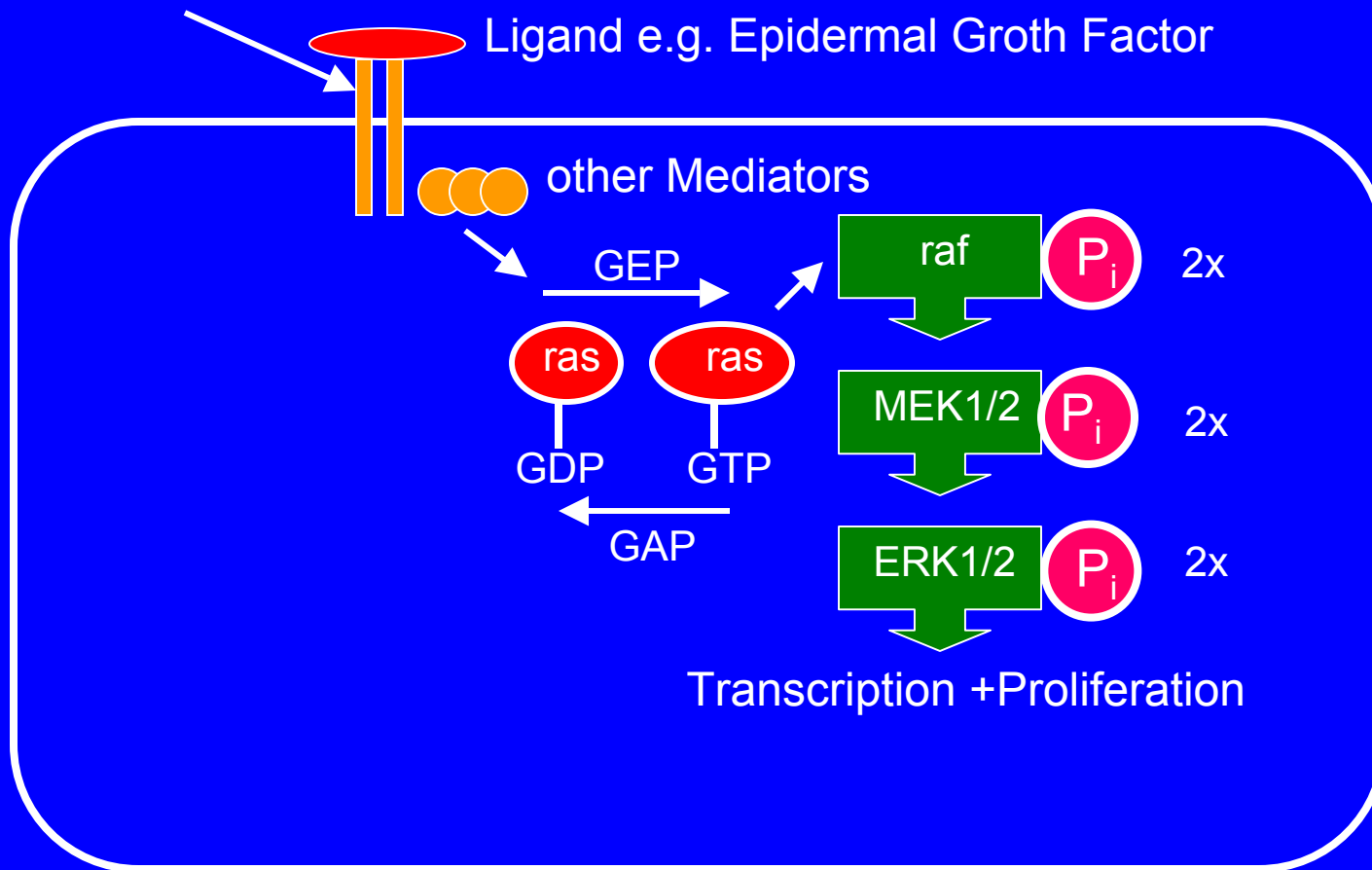


c-ras (cellular form): activity is regulated
=> Proto-Oncogene

Proto-Oncogenes: c-ras

Receptor Tyrosine Kinase

Ligand e.g. Epidermal Groth Factor



Tumor suppressor genes

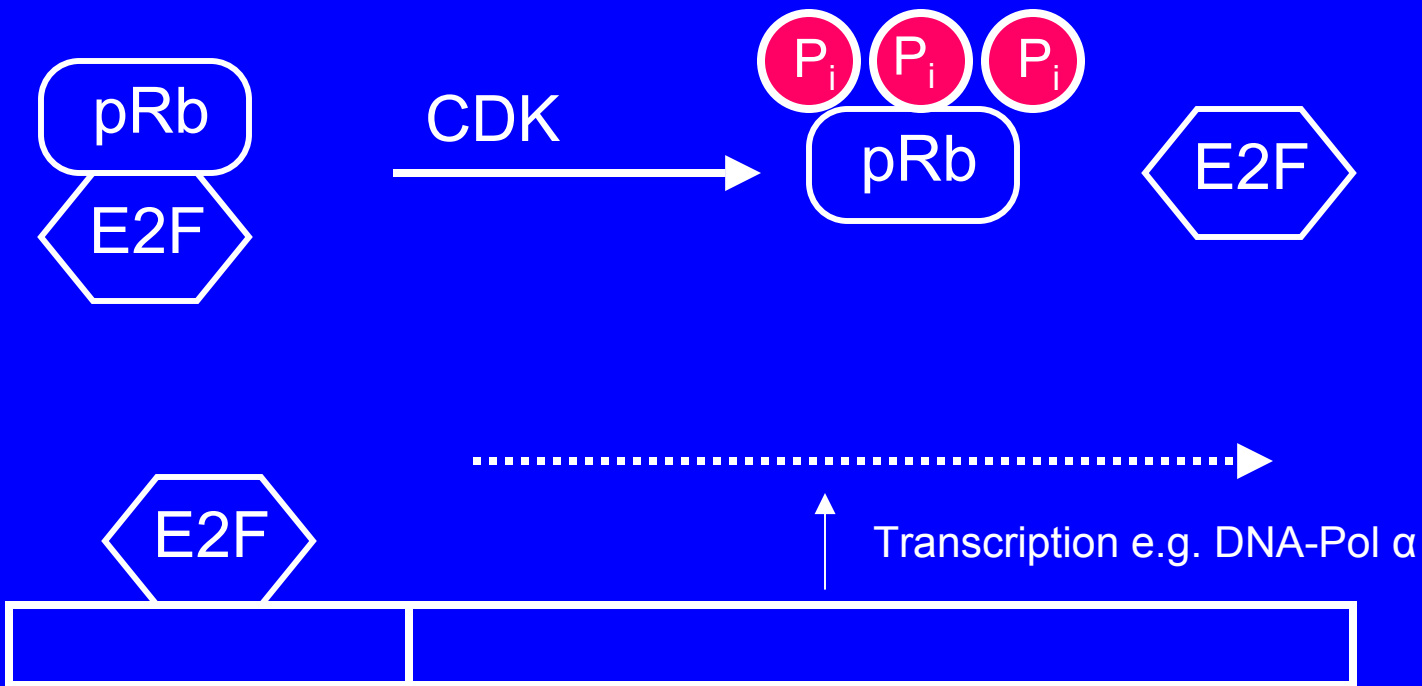
Gene products which are normally responsible for negative control of transcription and proliferation

Examples:

pRb inhibits transcription factors of the E2F-family, which are needed to get into the S-Phase of the cell cycle (Restriction Point)

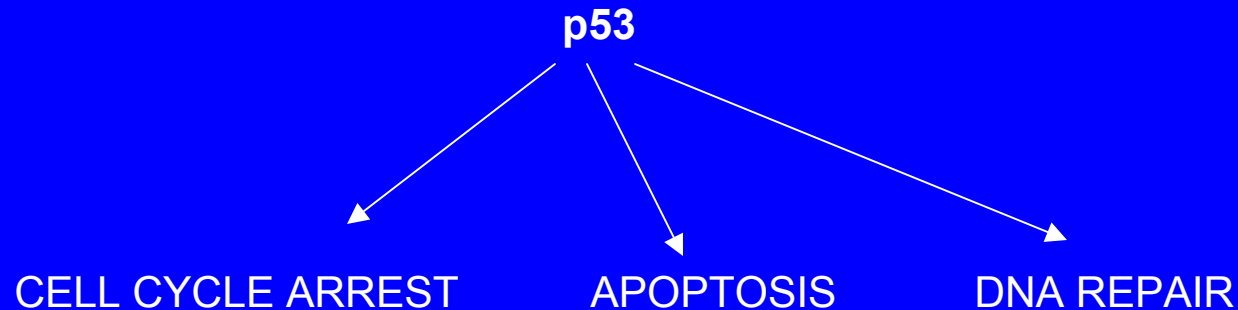
p53 induces transcription of the CDK-inhibitor (CDI) p21 which causes a cell cycle arrest (one function)
p53 is found upregulated in cells with a high level of DNA-damage

Tumor suppressor genes: rb



Tumor suppressor genes: p53

DNA damage
hypoxia
nucleotide depletion
heat shock
viral oncoprotein



Neoplastic transformation

- a tumor has a monoclonal origin
- a single mutation of a proto-Oncogene or Tumor suppressor gene is not enough for a Neoplastic Transformation
- mutations of a proto-Oncogene have a dominant effect
- mutations of a Tumor suppressor gene are recessive
- only a chain of multiple genomic changes leads to a Neoplastic Transformation (Tumor Progression)

Neoplastic transformation

spontaneous or induced DNA damage



primary molecular lesion



genomic instability

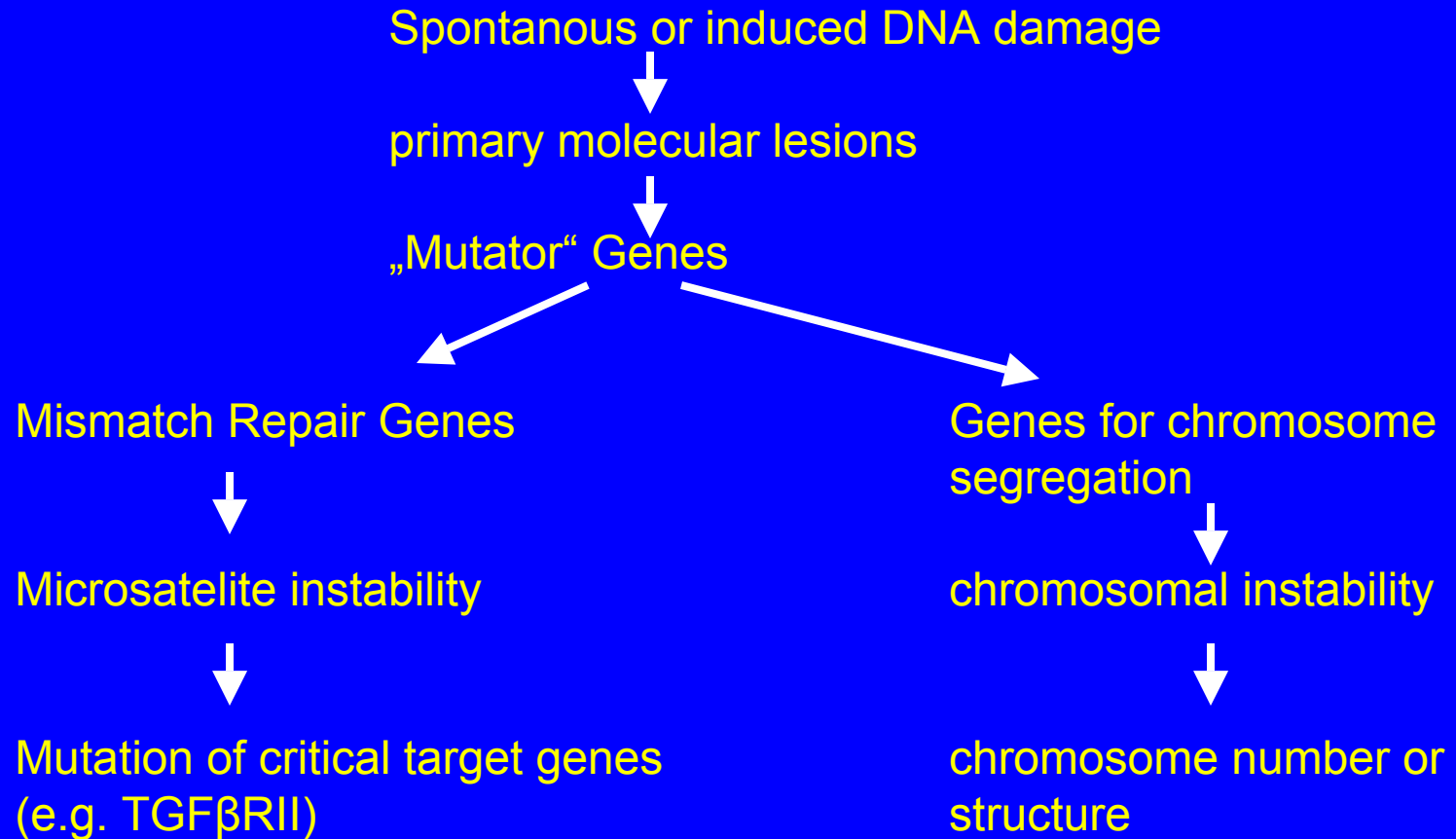


transforming molecular lesion



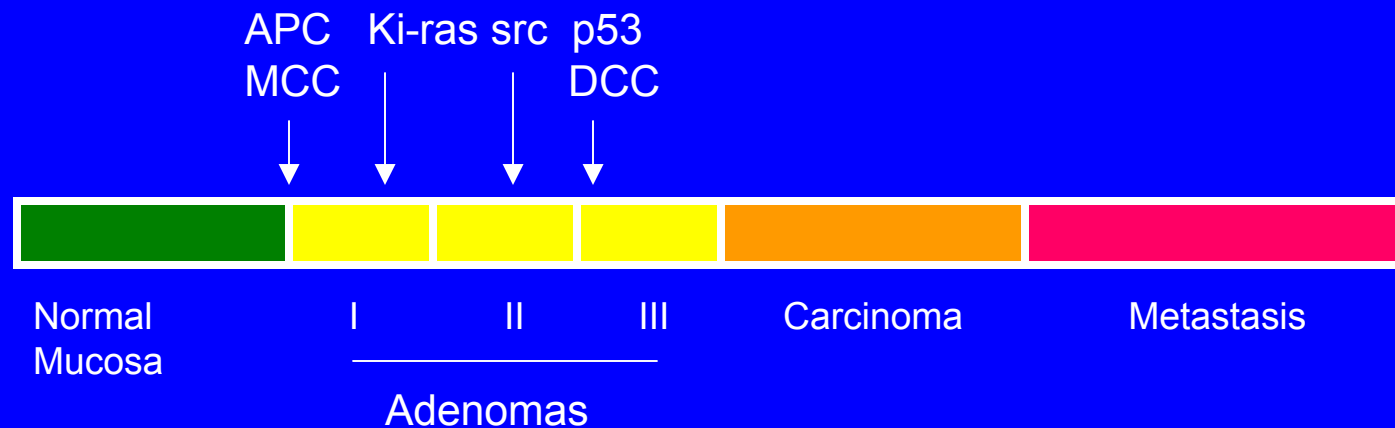
Neoplastic Transformation

Neoplastic transformation



Neoplastic transformation

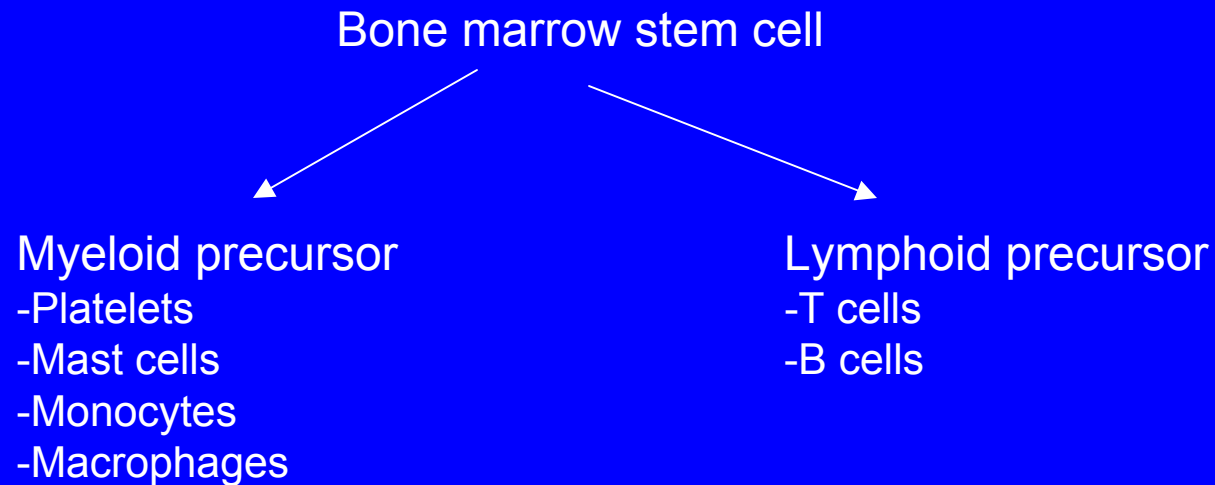
Genetic alterations during the Familial Adenomatous Polyposis Syndrome (FAP)



APC: aden polyp. Coli
MCC: mutated in colon cancer
DCC: deleted in colon cancer

src: gene for a protein tyrosine kinase

Leukemia



Lymphoblastic Leukemia: Transformation of a lymphoid determined stem cell (acute or chronic, ALL, CLL)

Myeloid Leukemia: Transformation of a myeloid determined stem cell (acute or chronic, AML, CML)

Leukemia

Different Leukemias are mostly characterized through simple Translocations between different chromosomes (=> fusion genes)

best known example: Philadelphia Chromosome t(9;22)(q34;q11) (CML,ALL)

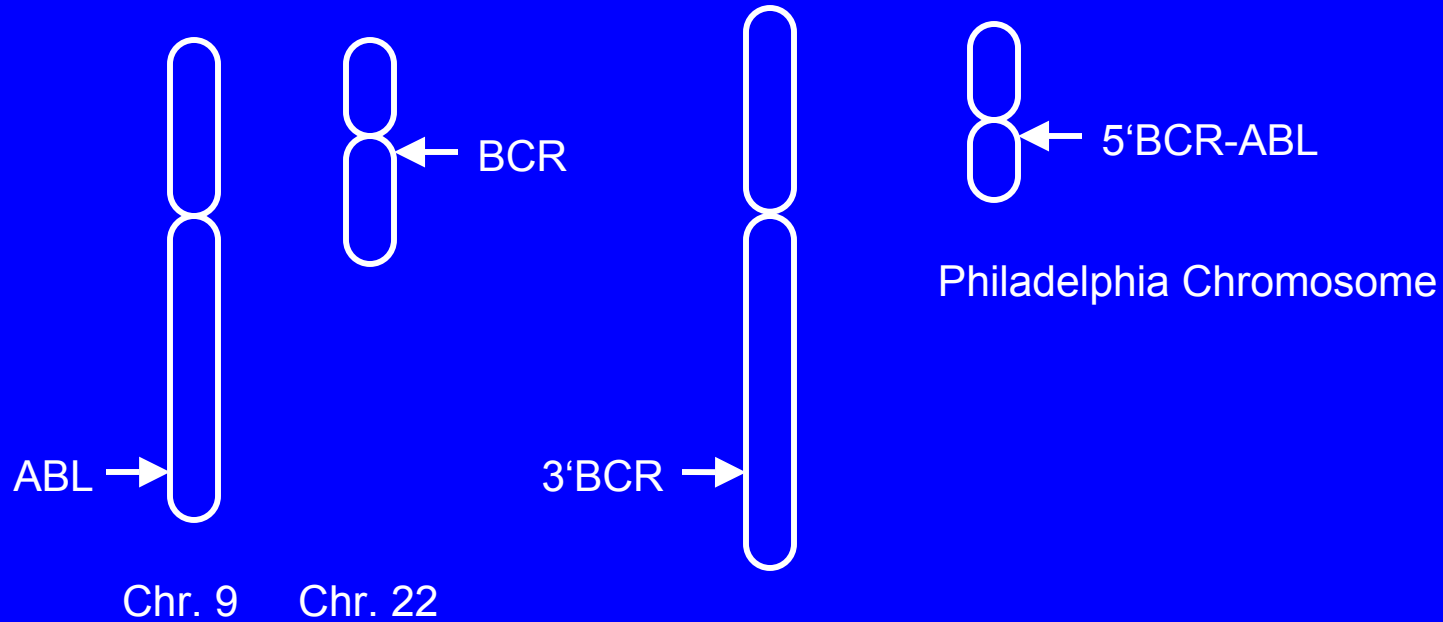
⇒ **Fusion gene: BCR-ABL**

c-abl: tyrosine kinase which probably is involved in cell cycle regulation

bcr(breakpoint cluster region): maybe involved in signal transduction

Fusion gene: constitutive tyrosine kinase activity contributed by c-abl

Leukemia



Other examples: $t(8,14)$, $t(2,8)$, $t(8,22)$ \Rightarrow fusion of the proto-Oncogene c-myc on chromosome 8 with one immunoglobulin-locus \Rightarrow overexpression of c-myc

Summary

- Cancer arises from a wrong cell cycle regulation
- one monoclonal mutation leads to genomic instability
- Neoplastic Transformation is the consequence of the following multiple genetic alterations