



Molecular mechanisms involved into cancer development

Genetic alterations

- Oncogenes
- Tumor suppressor genes
- DNA repair genes
- Telomere and telomerase

Epigenetic alterations

- DNA-methylation
- Micro-RNA expression



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Genetic alterations

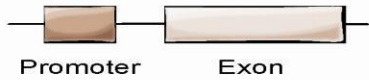
- **Oncogenes**
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- Micro-RNA expression

Mechanisms of protooncogene – oncogene transformation

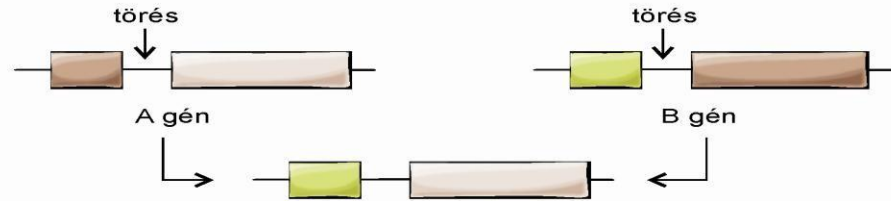
Protooncogén



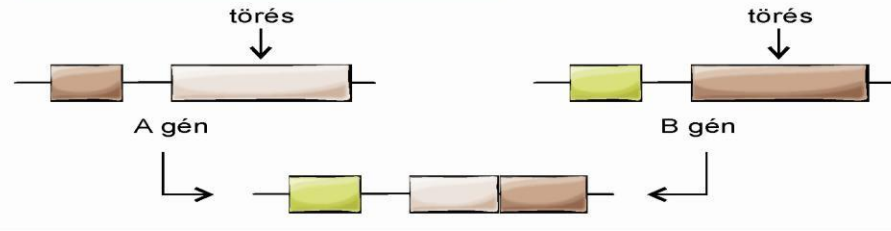
Pontmutáció / deléció / inszerció



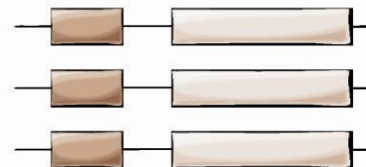
Transzlokáció (promotercsere)



Transzlokáció (fúziógén)

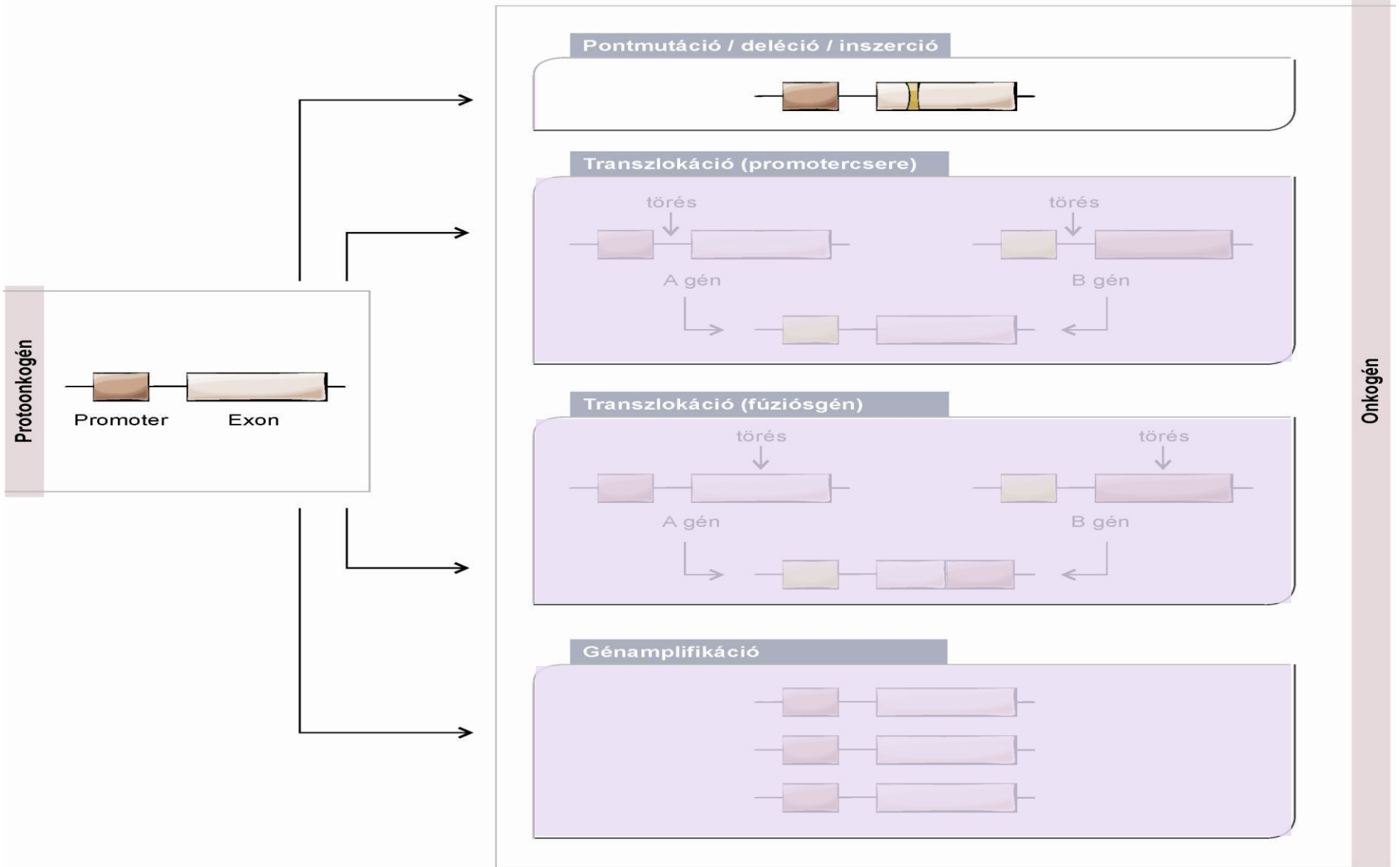


Génamplifikáció



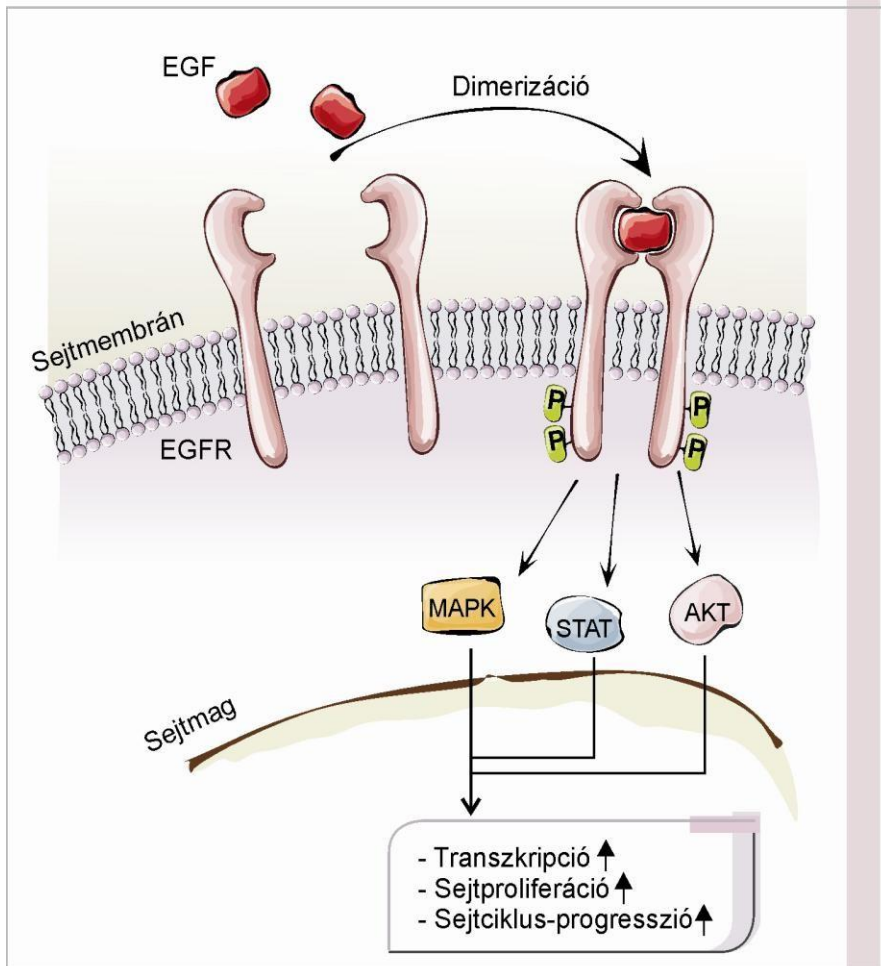
Onkogén

Mechanisms of protooncogene – oncogene transformation

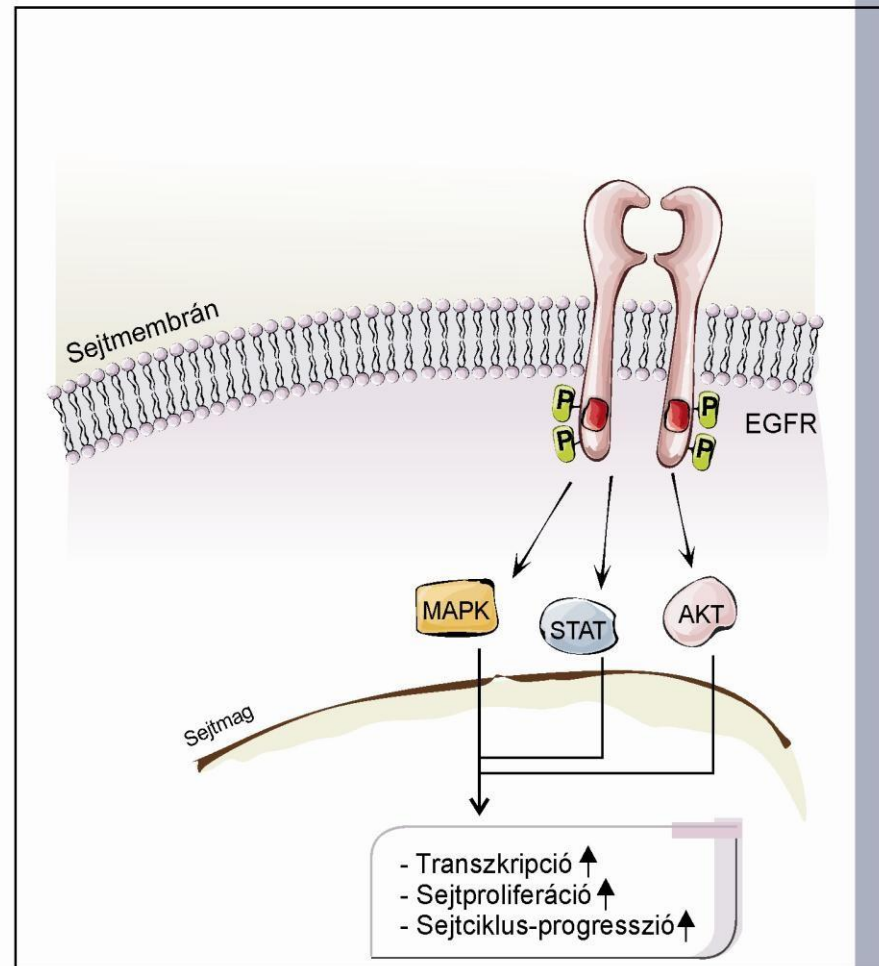


EGFR protooncogene – oncogene transformation (point mutation)

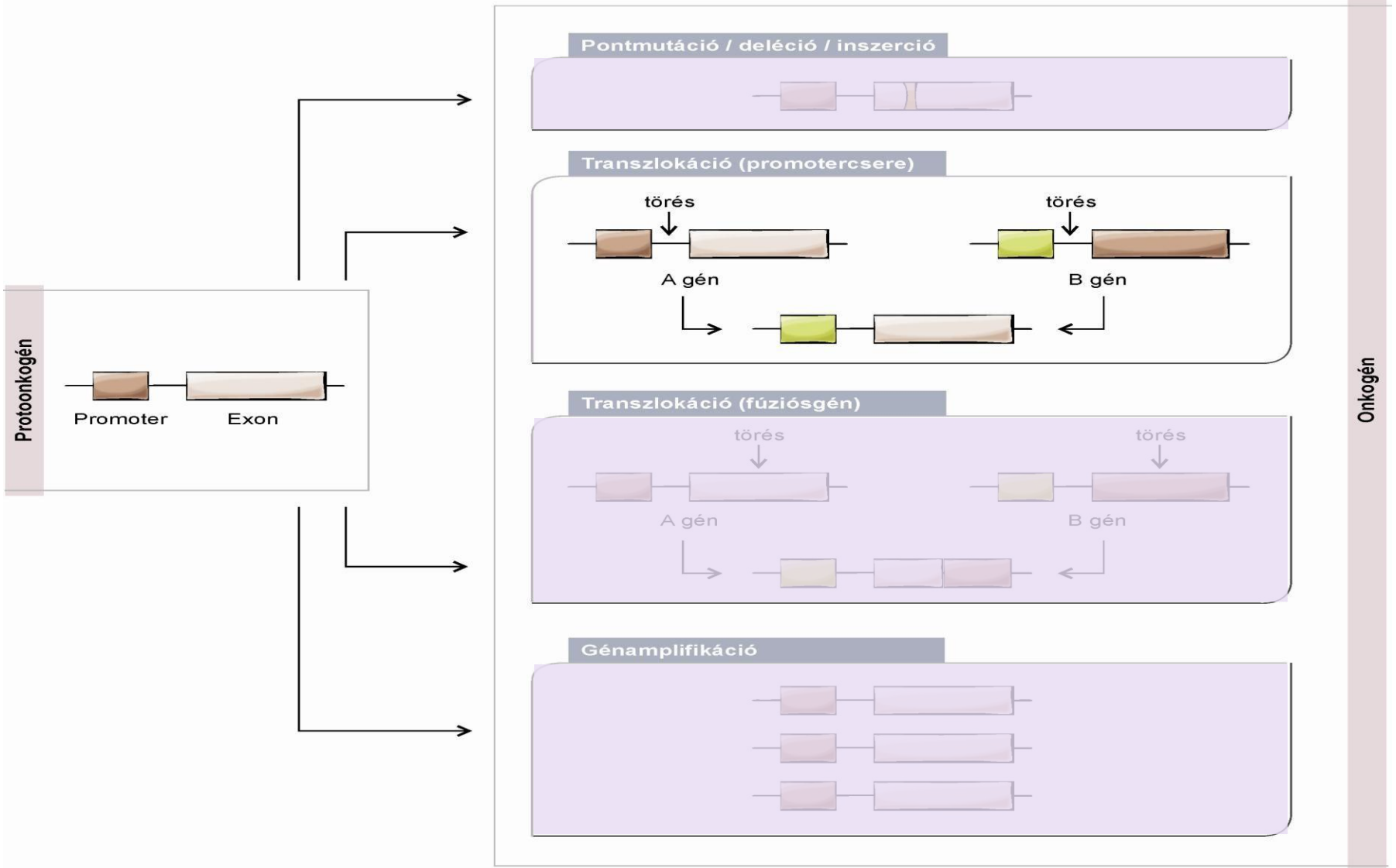
A. Protoonkogén EGFR



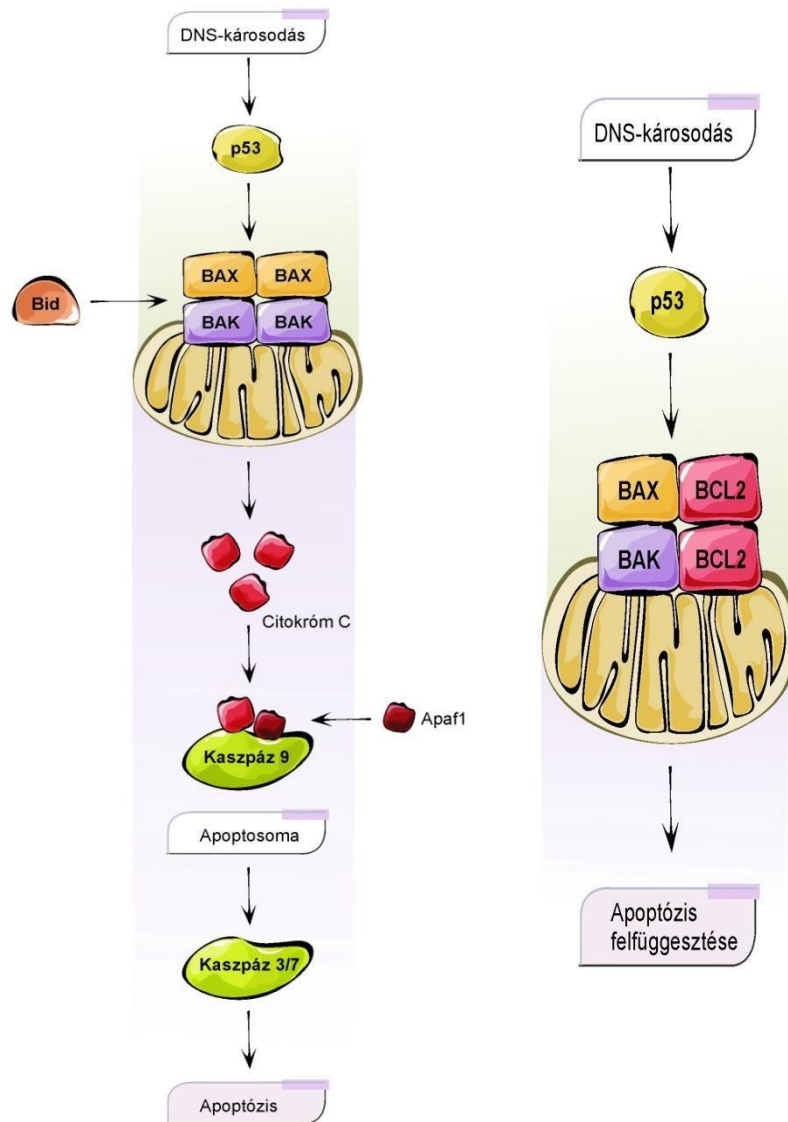
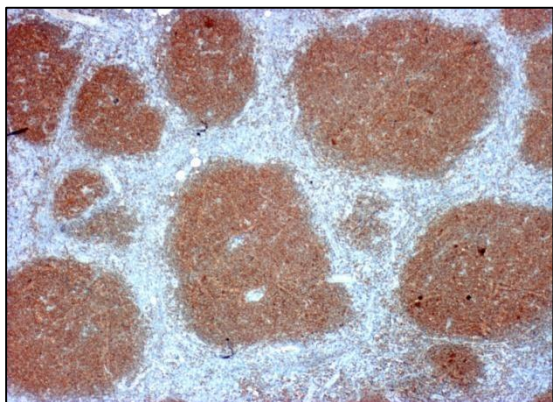
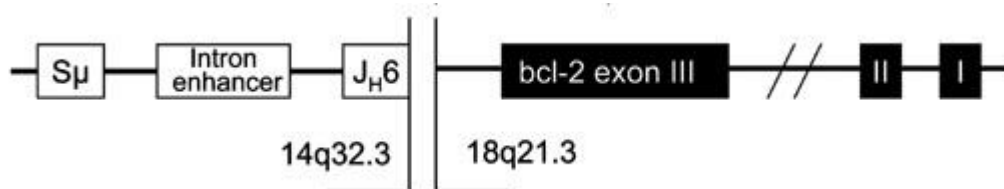
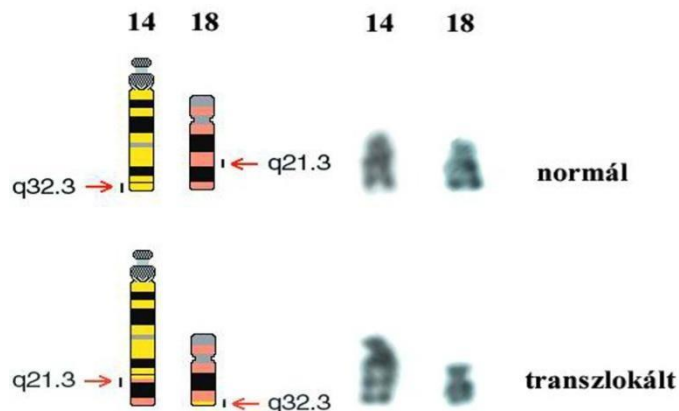
B. Onkogén EGFR



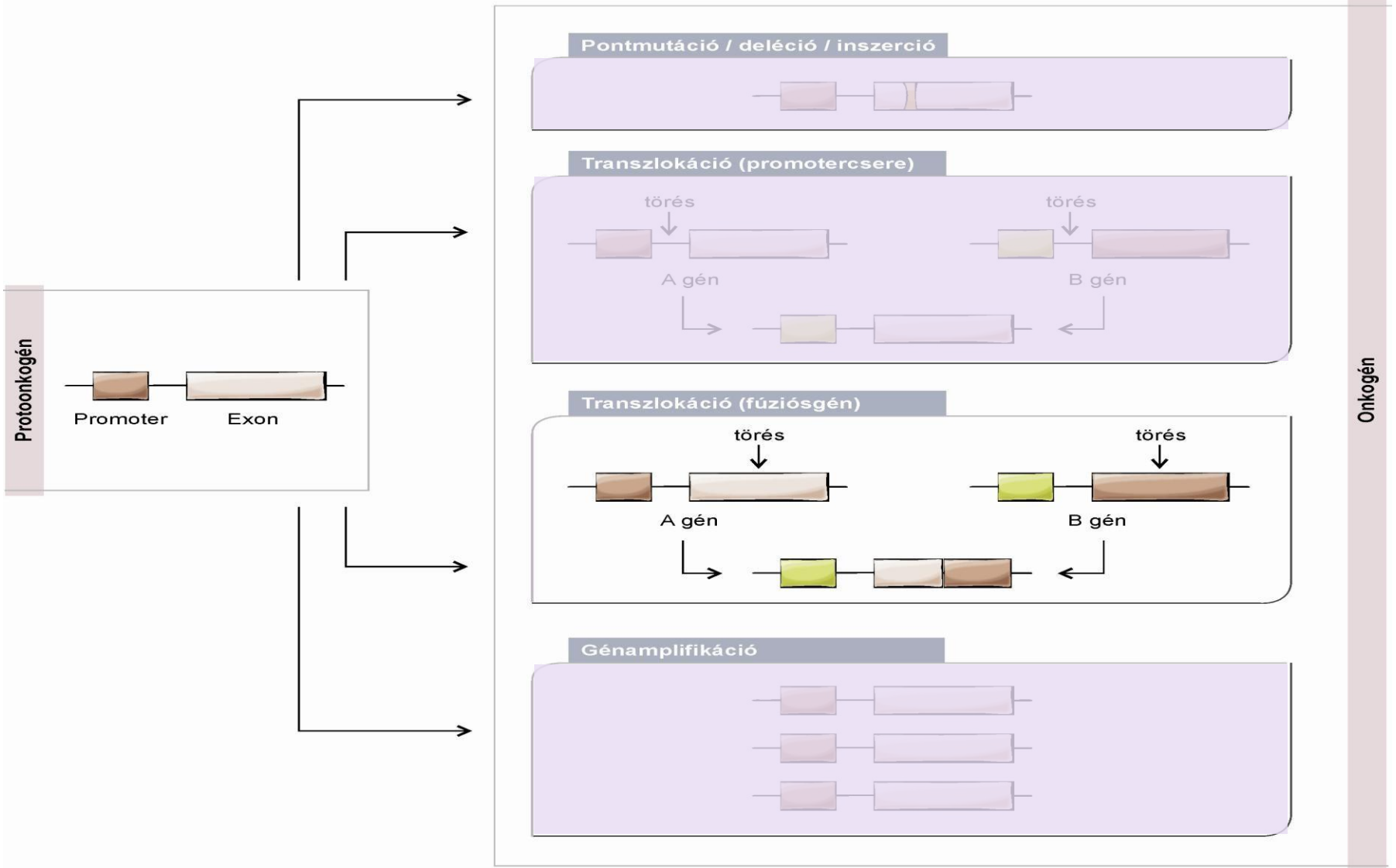
Mechanisms of protooncogene – oncogene transformation



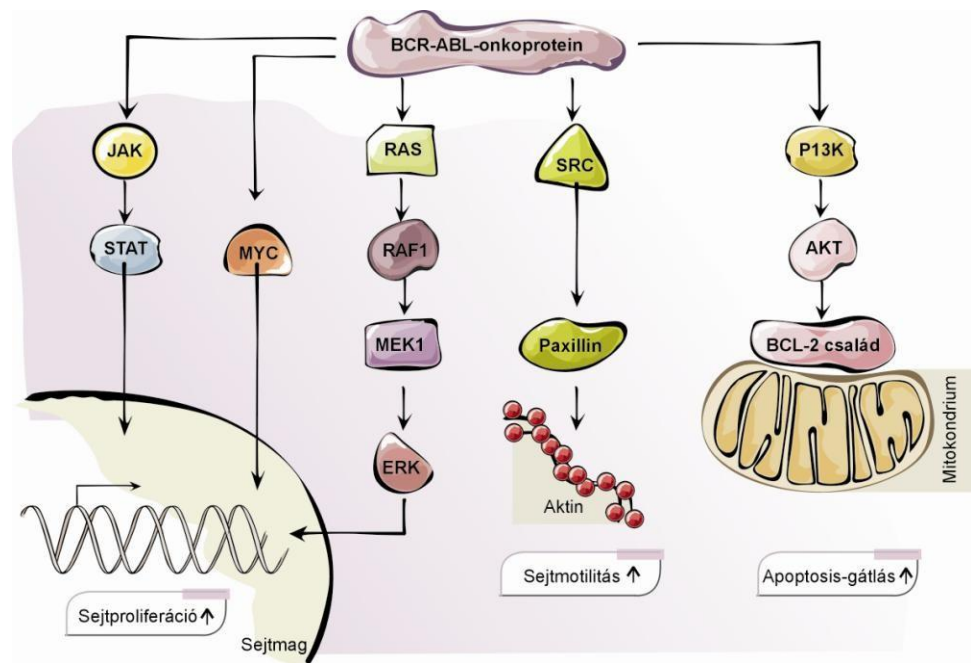
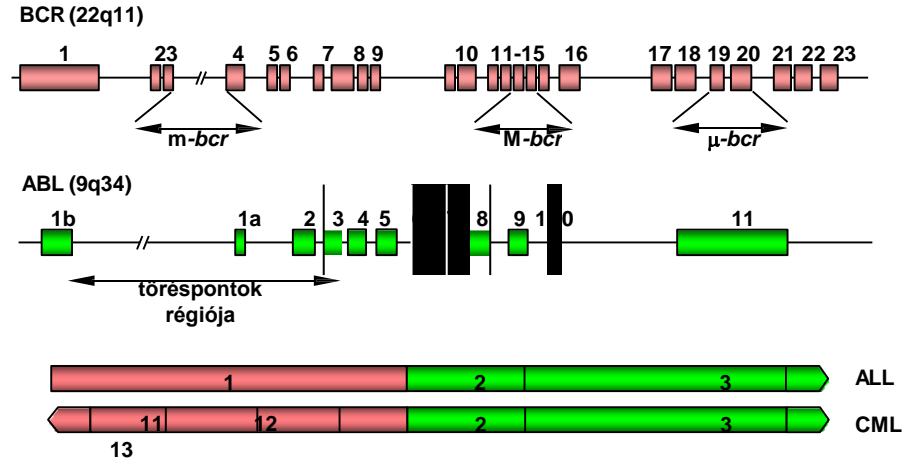
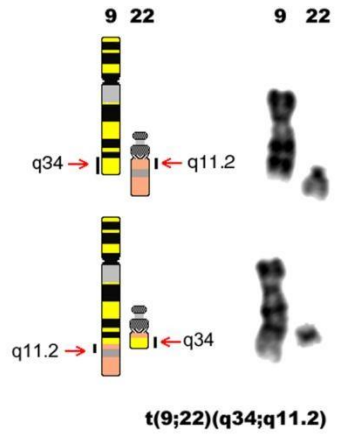
BCL2 protooncogene – oncogene transformation (translocation – promoter exchange)



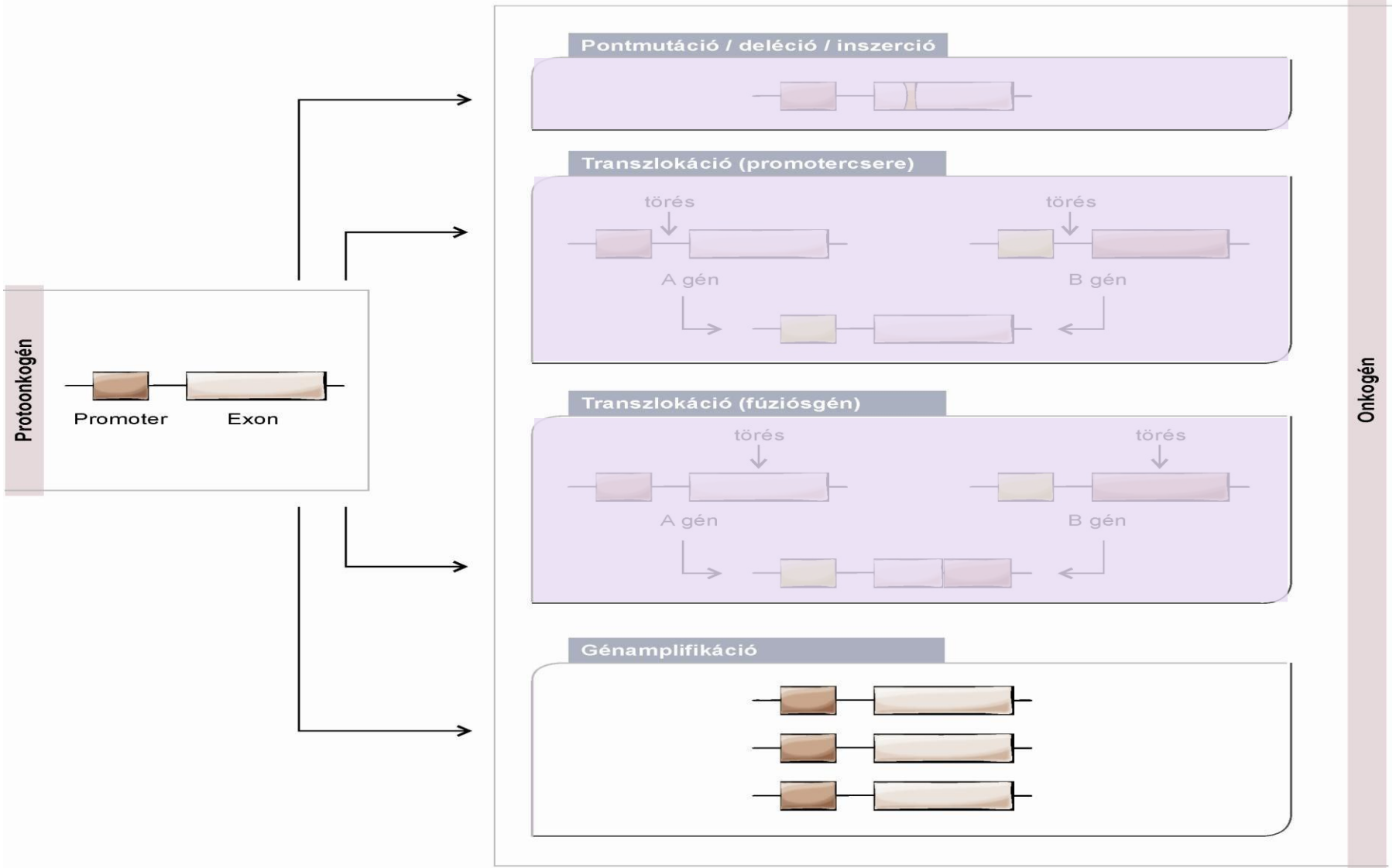
Mechanisms of protooncogene – oncogene transformation



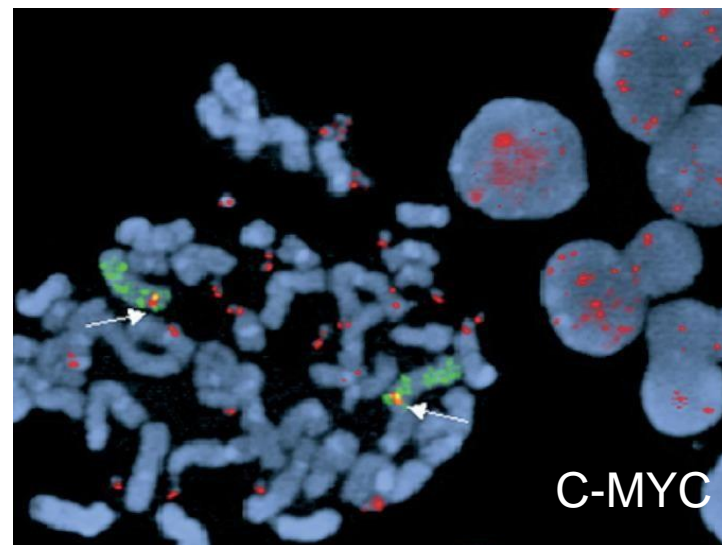
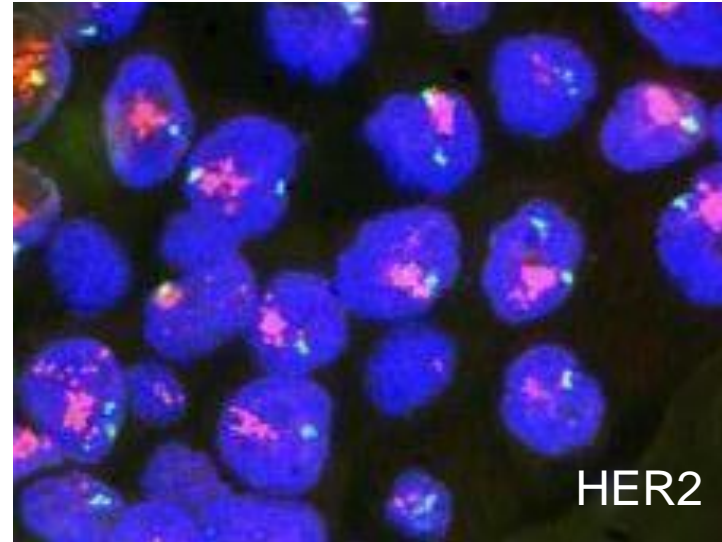
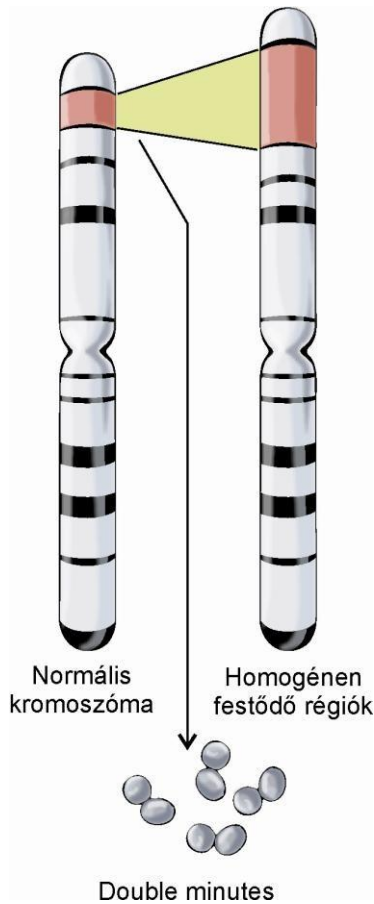
Protooncogene – oncogene transformation (translocation – gene fusion)



Mechanisms of protooncogene – oncogene transformation



Protooncogene – oncogene transformation (gene amplification)





Protooncogene activations involved in human cancer development

ONCOPROTEIN FUNCTION	PROTO-ONCOGENE	TYPE OF ACTIVATION	NEOPLASM
Growth factor	<i>SIS</i> <i>FGF-3</i> <i>HST</i>	Increased expression Amplification Increased expression	Astrocytoma, osteosarcoma Melanoma, breast-, bladder cancer Stomach cancer, Kaposi-sarcoma
Growth factor receptor	<i>EGFR</i> <i>FLT3</i> <i>RET</i> <i>KIT</i> <i>MET</i>	Increased expression Amplification Point mutation Point mutation Point mutation	Lung planocellular carcinoma, glioma Breast-, ovarium cancer MEN 2A, 2B, medullary carcinoma GIST Papillary renal cell carcinoma
GTP-binding proteins	<i>KRAS</i> <i>HRAS</i> <i>NRAS</i>	Point mutation Point mutation Point mutation	Lung-, colon-, pancreas cancer Kidney-, bladder cancer Melanoma, AML
Non-receptor thirosine kinase	<i>ABL</i> <i>JAK2</i>	Translocation Point mutation	CML, ALL PV, ET, MF
Nuclear transcription factor	<i>C-MYC</i> <i>N-MYC</i> <i>L-MYC</i> <i>C-JUN</i>	Translocation Amplification Amplification Increased expression	Burkitt lymphoma Neuroblastoma, SCLC SCLC Lung-, colon cancer
Ciklin and ciklin dependent kinase	<i>Ciklin D</i> <i>Ciklin E</i> <i>CDK4</i>	Translocation Amplification Increased expression Point mutation	Mantle cell lymphoma Breast-, oesophagus cancer Breast cancer Melanoma, glioblastoma
Antiapoptotic protein	<i>BCL2</i>	Translocation	Follicular lymphoma



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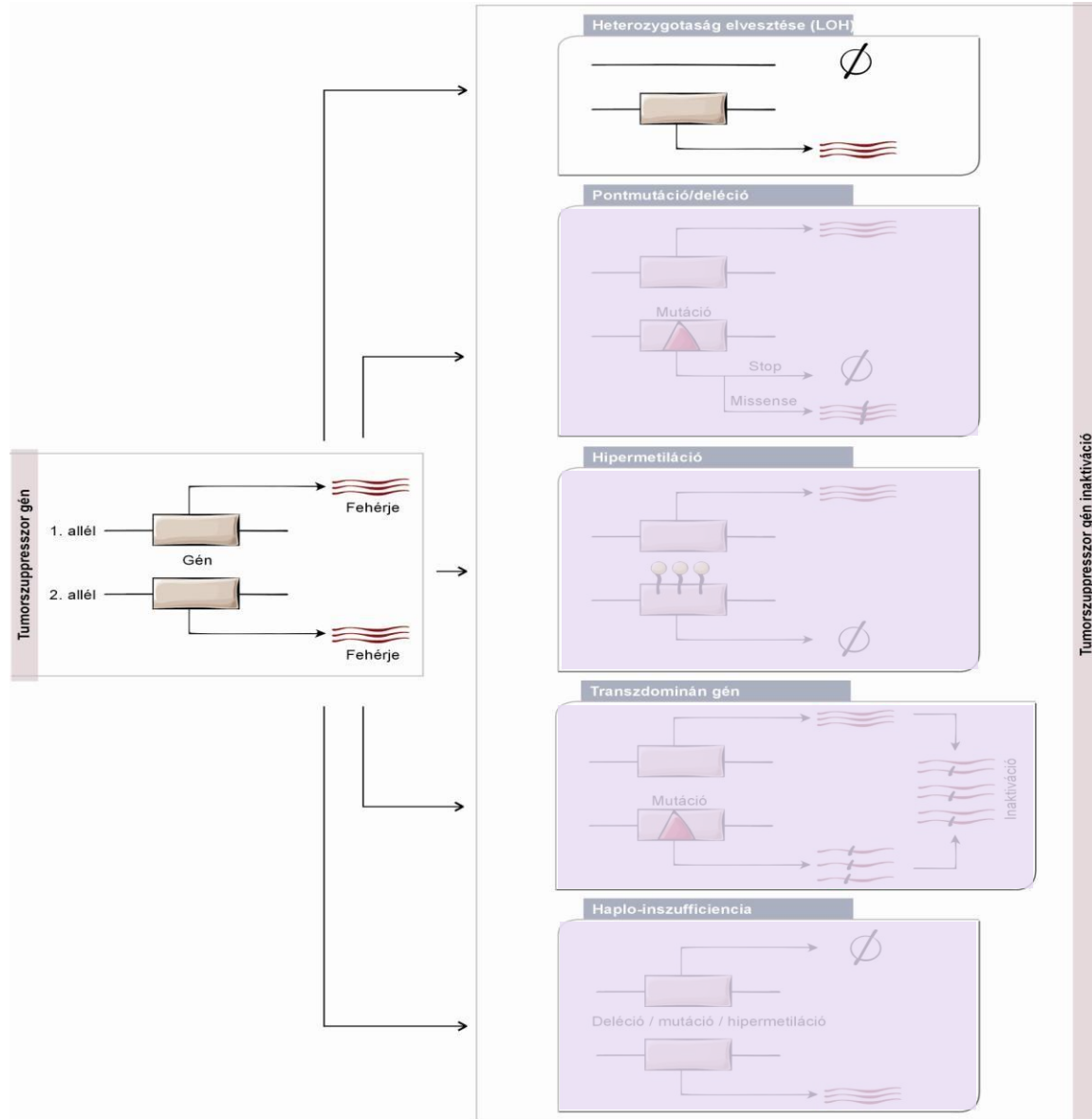
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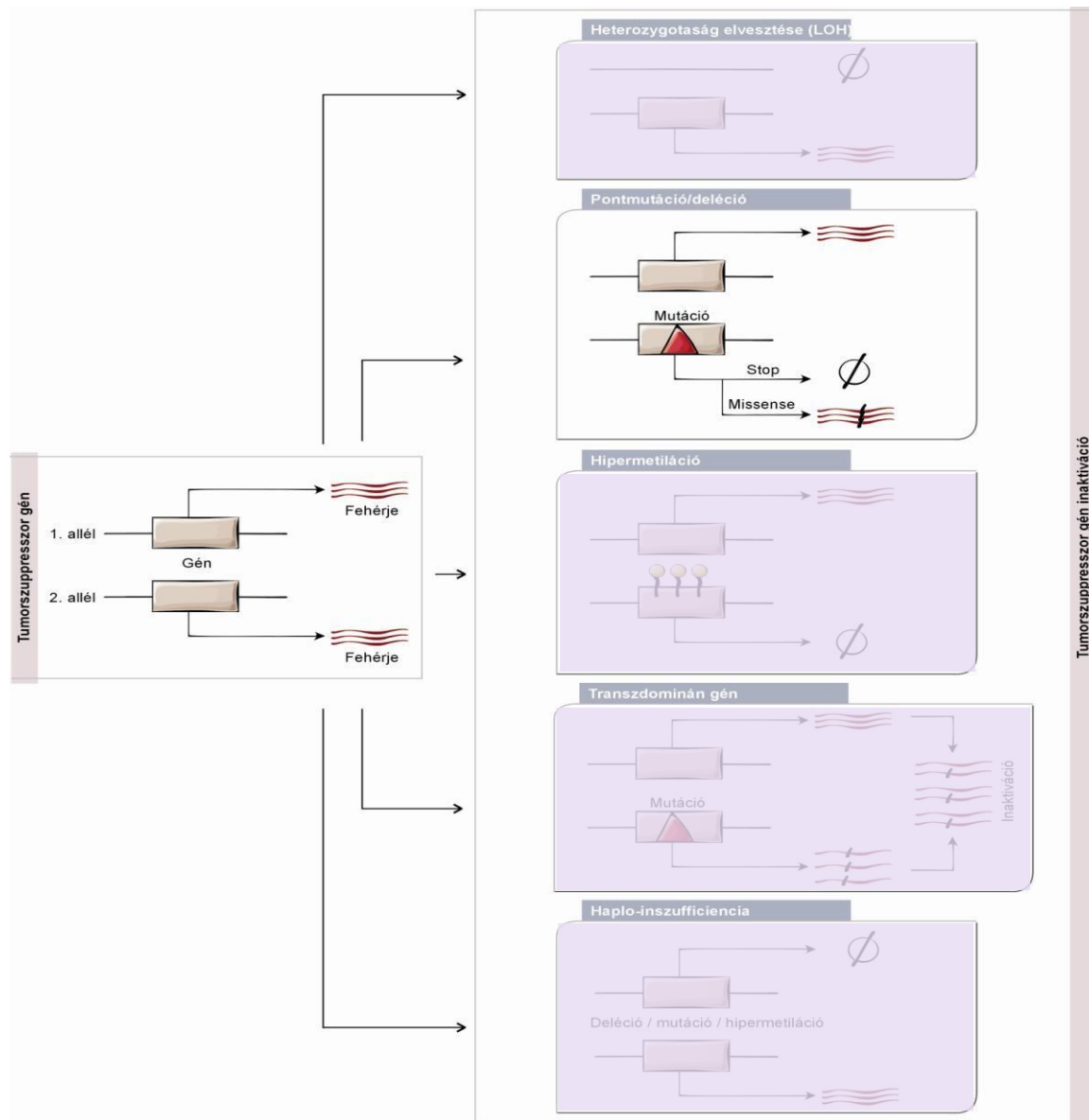
Epigenetic alterations

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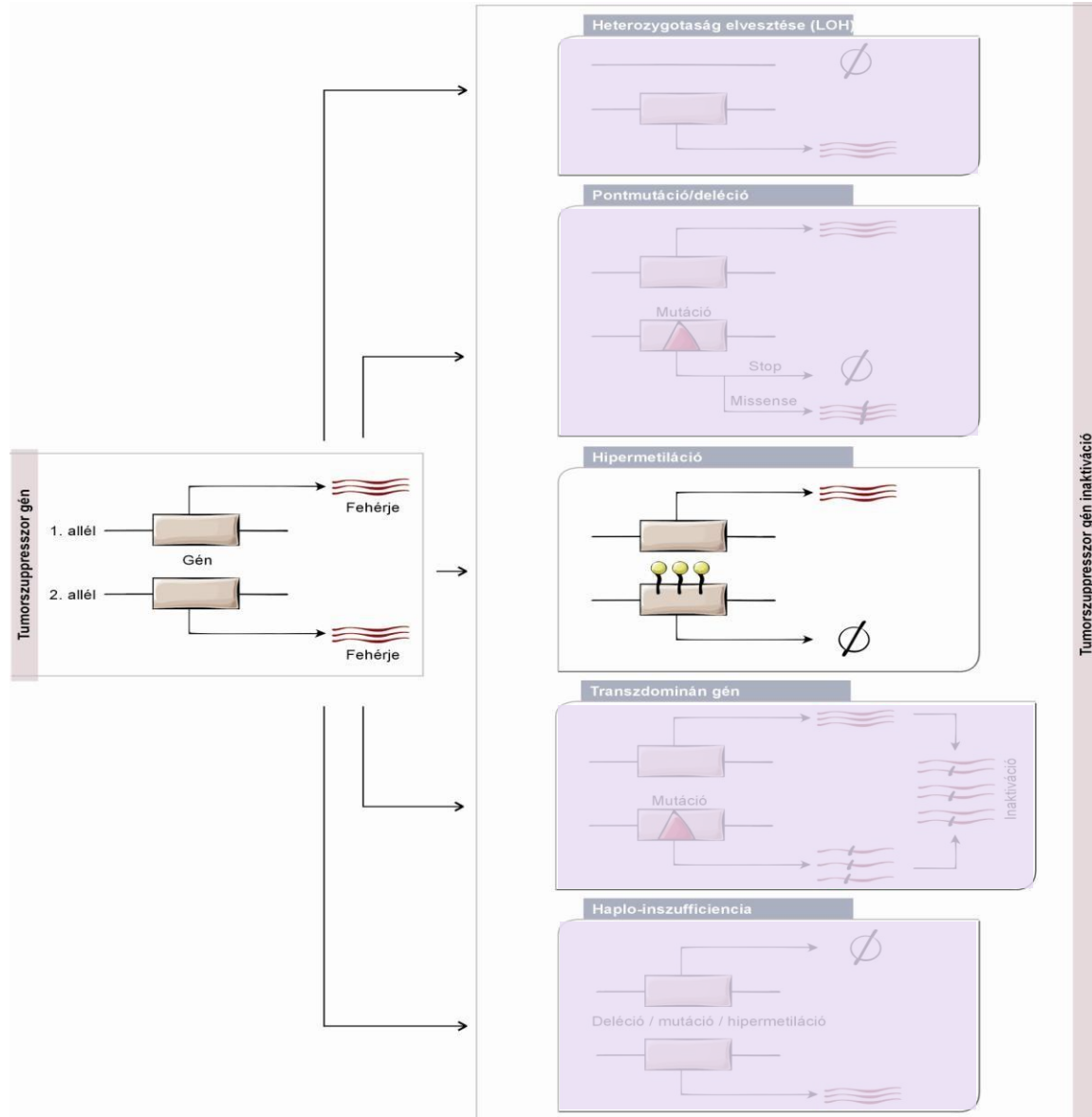
Mechanisms of tumor suppressor gene inactivations



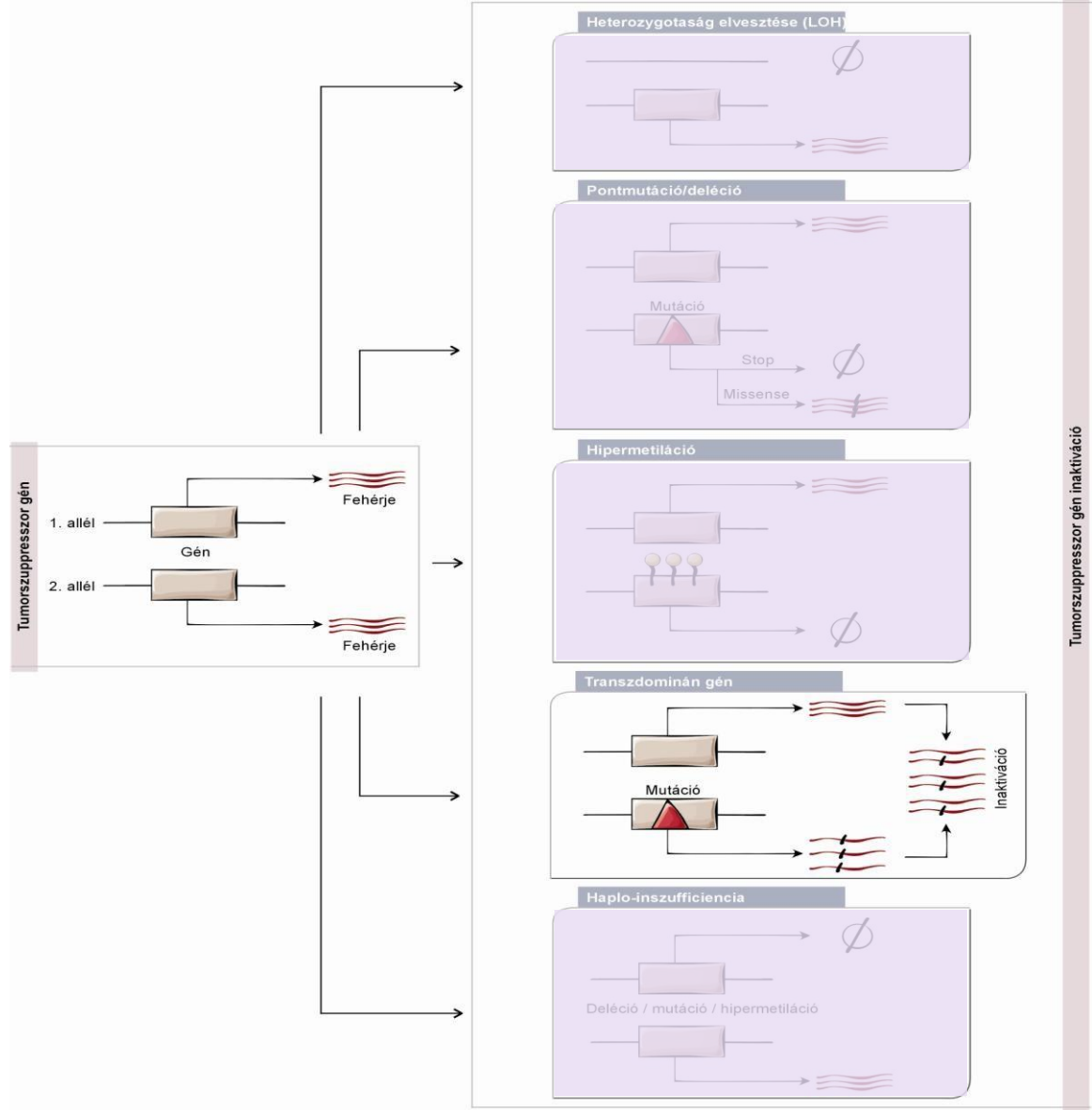
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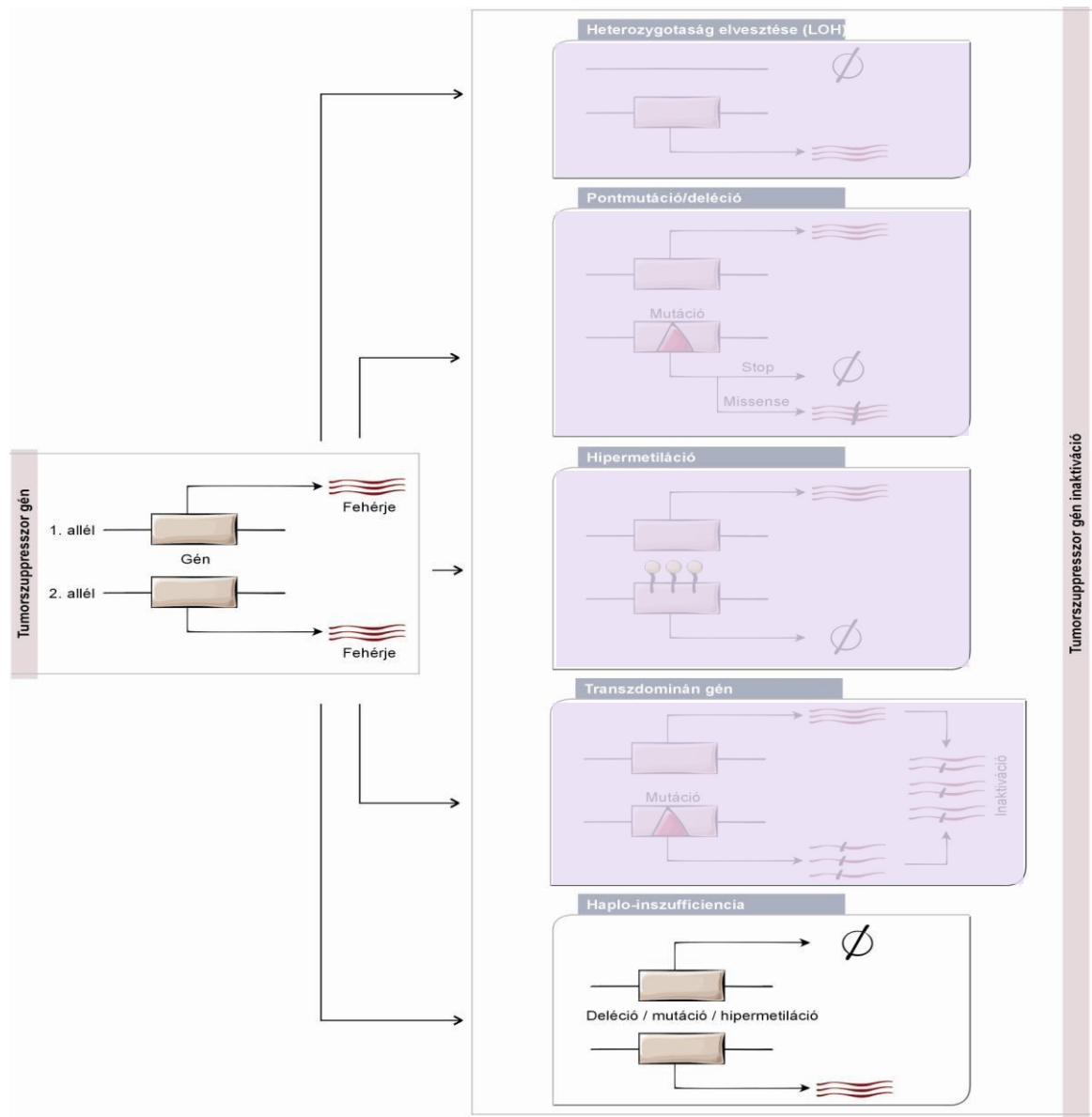
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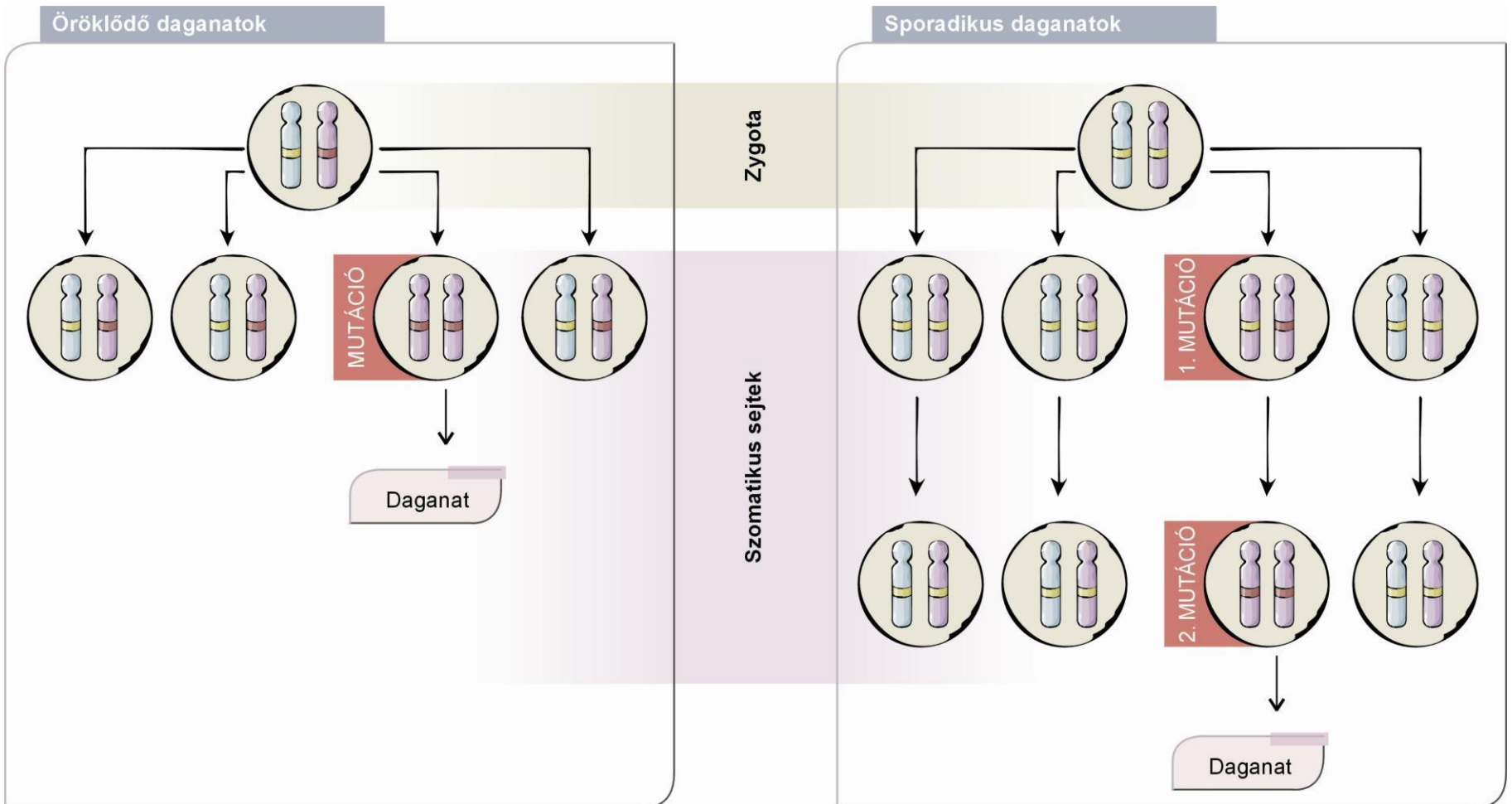
Mechanisms of tumor suppressor gene inactivations



Mechanisms of tumor suppressor gene inactivations



Tumorsuppresszor gene inaktivation in hereditary and sporadic cancers

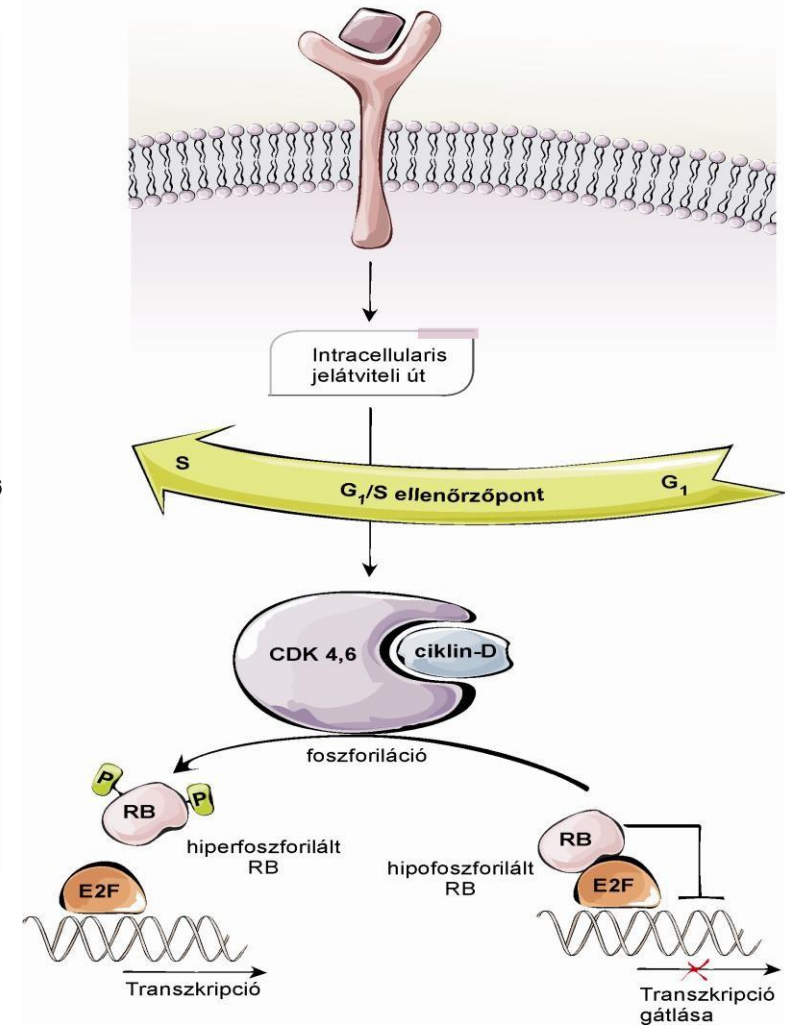
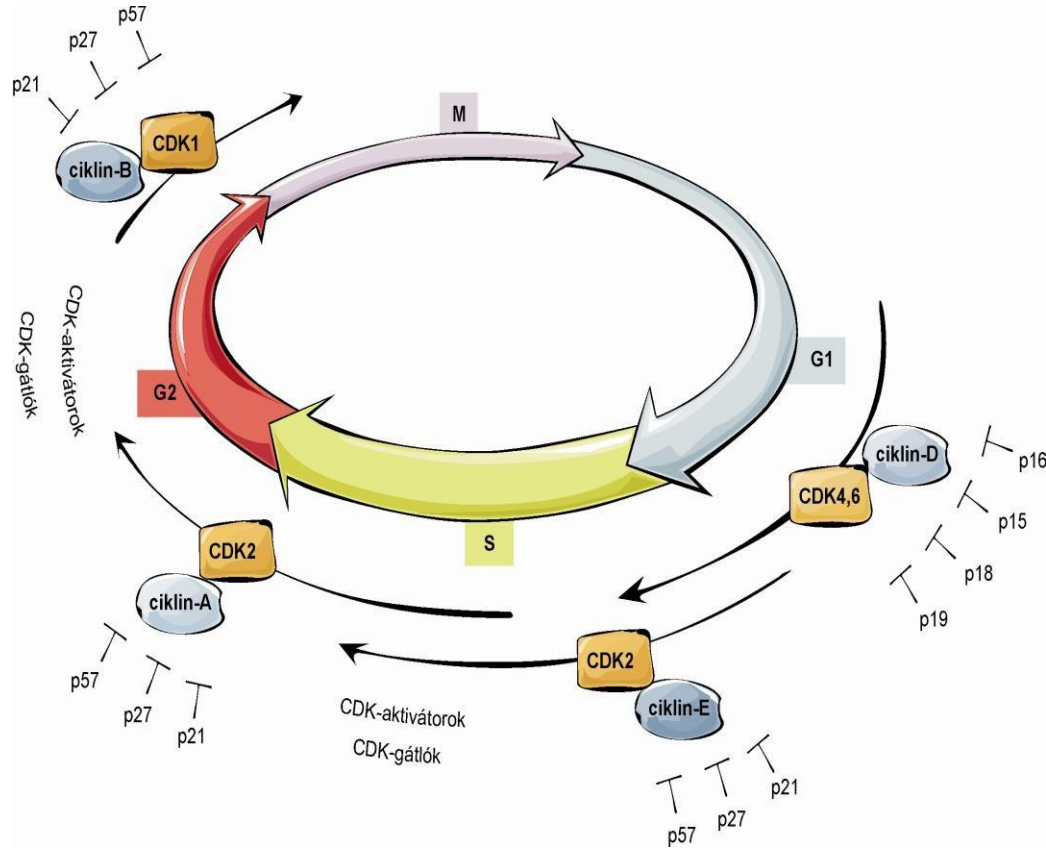




Tumorsuppressor genes involved in human cancer development

GENE	FUNCTION	HEREDITARY NEOPLASMS	SPORADIC NEOPLASMS
<i>RB</i>	Cell cycle regulation	Retinoblastoma, osteosarcoma	Retinoblastoma, osteosarcoma, emlő-, colon-, tüdőcarcinoma
<i>p53</i>	Cell cycle block, apoptosis, cell aging	Li-Fraumeni-syndroma	Most human cancer
<i>APC</i>	β -katenin block	Familial adenomatous polyposis, melanoma	Colon-, stomach-, pancreas cancer
<i>WT1</i>	Nuclear transcription	Wilms-tumor	Wilms-tumor
<i>p16/p16</i>	Ciklin dependent kinase inhibitor	Melanoma	Breast-, eosophagus-, prostata-, pancreas cancer
<i>BRCA1</i>	DNA-repair	Breast-, ovarium cancer	Breast-, ovarium cancer
<i>BRCA2</i>	DNA-repair	Breast cancer	Stomach cancer
<i>VHL</i>	Elongin inhibitor	vonHippel-Lindau-syndrome	Renal cell carcinoma, haemangioma
<i>NF1</i>	RAS, p21 inhibitor	Neurofibromatosis 1	Neuroblastoma, melanoma
<i>NF2</i>	Citoskeleton stability	Neurofibromatosis 2	Schwannoma, meningeoma
<i>PTEN</i>	Tyrosine kinase inhibitor	Cowden-syndrome	Endometrium-, prostate cancer

The role of inactivation of *RB* tumoursuppressor gene in cancer development

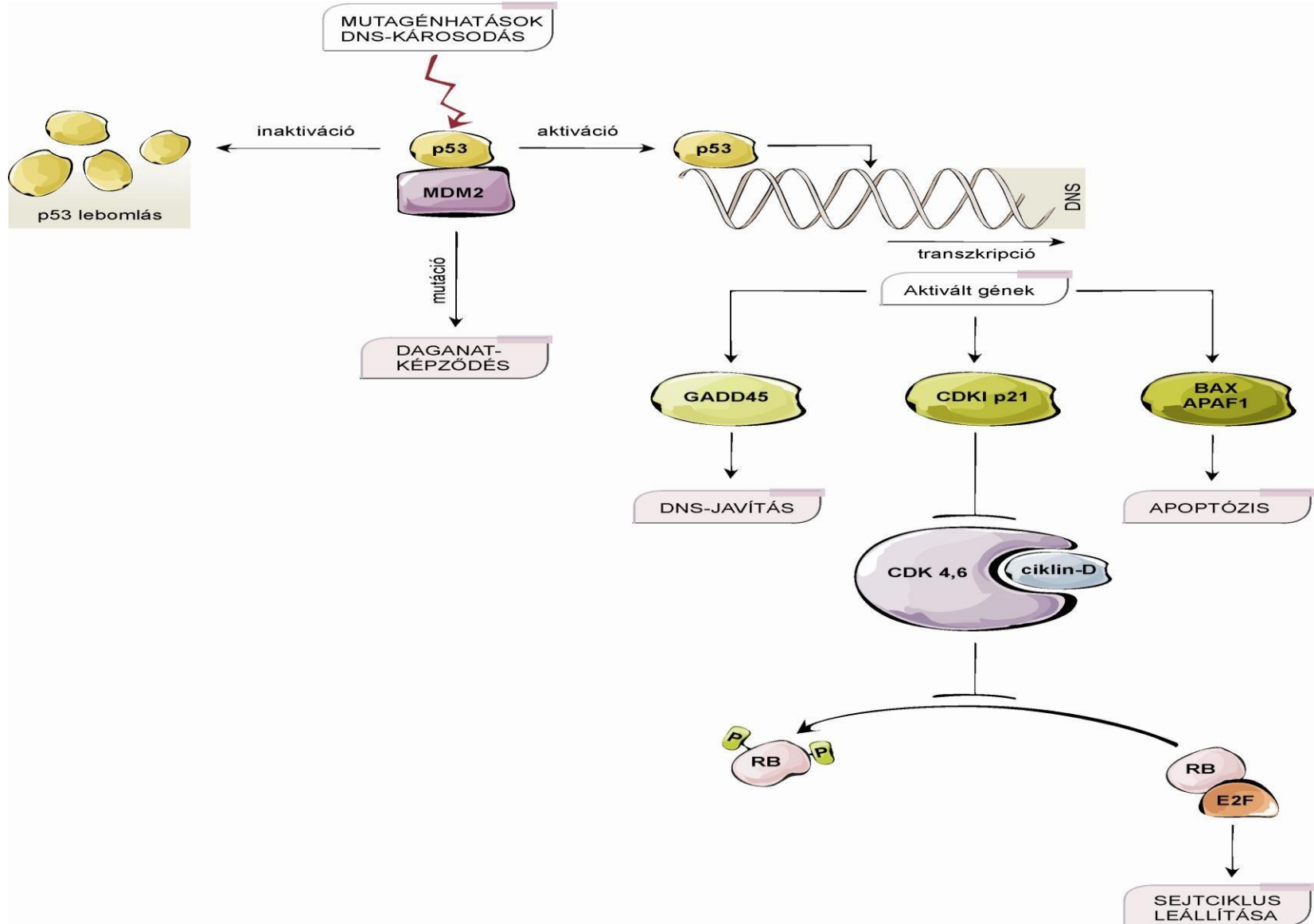




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The role of inactivation of p53 tumoursuppressor gene in cancer development



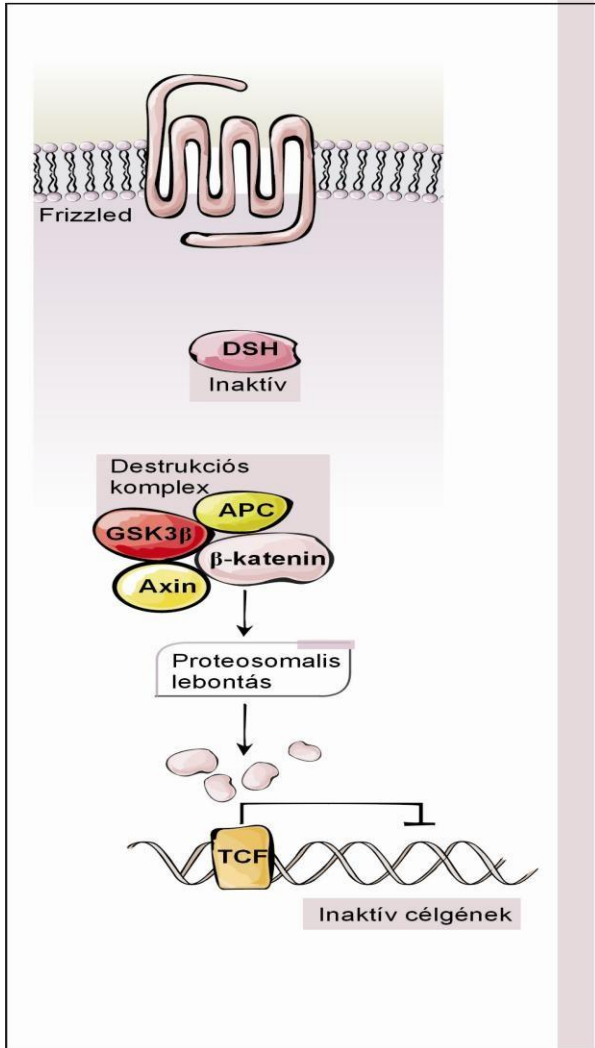


Tumorsuppressor genes involved in human cancer development

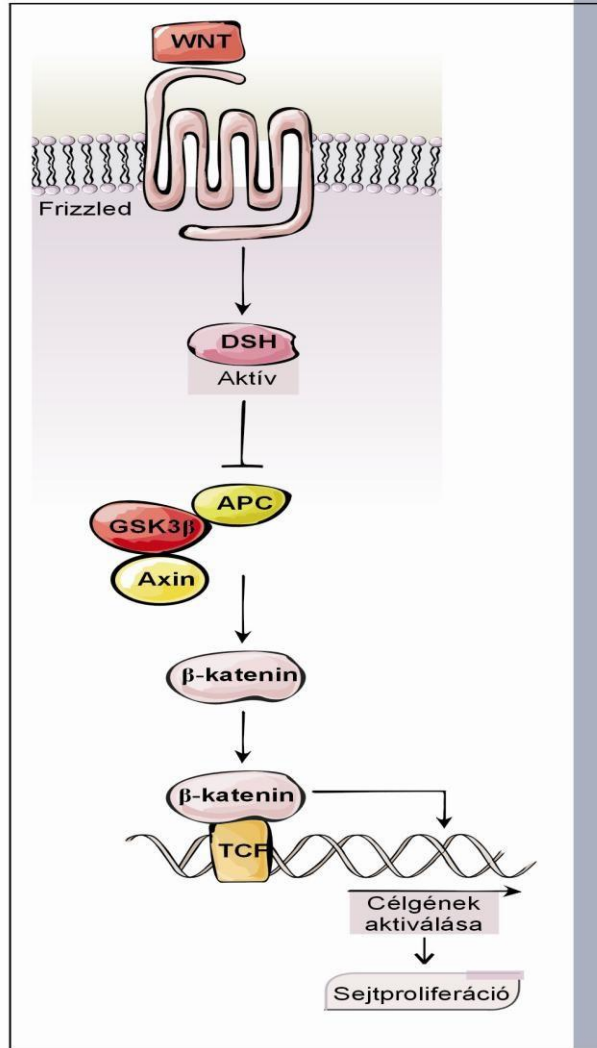
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The role of APC in the development of adenomatous polyposis coli

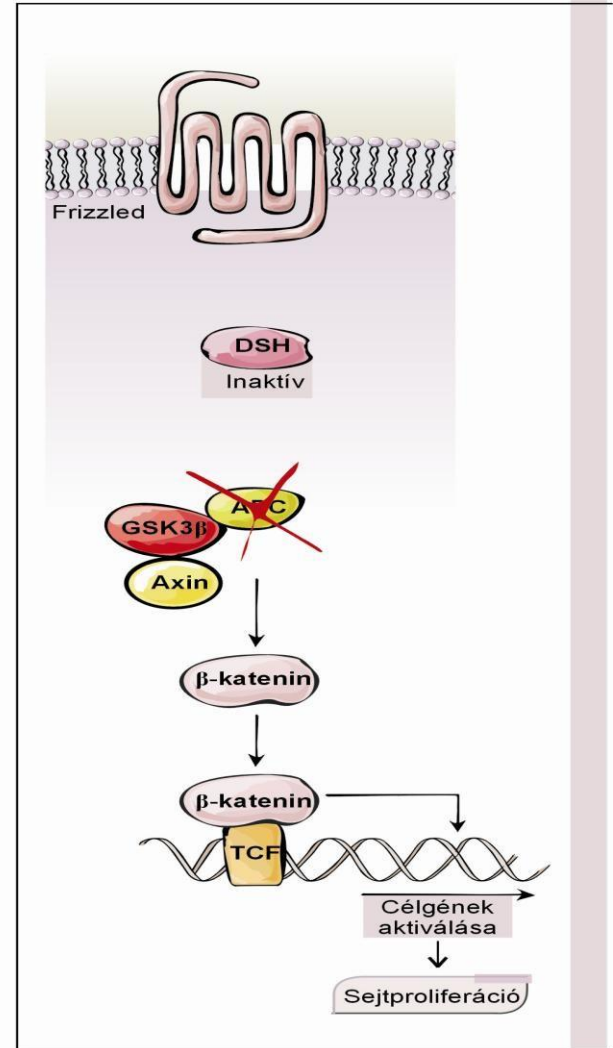
A. Inaktív WNT jelátviteli út



B. Aktív WNT jelátviteli út



C. WNT jelátviteli út APC inaktiváció esetén





Molecular mechanisms involved into cancer development

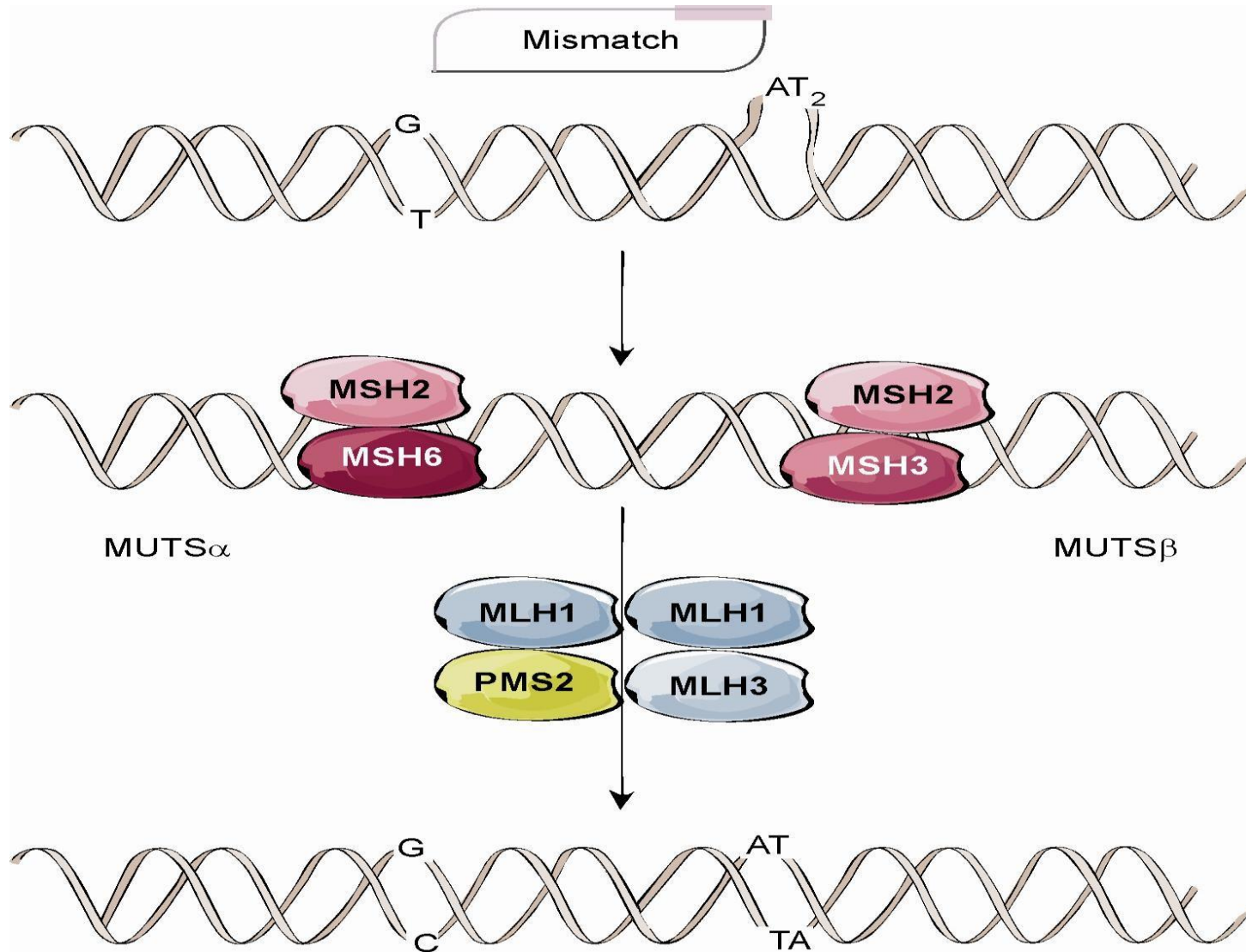
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- **DNS repair genes**
- Telomer and telomerase

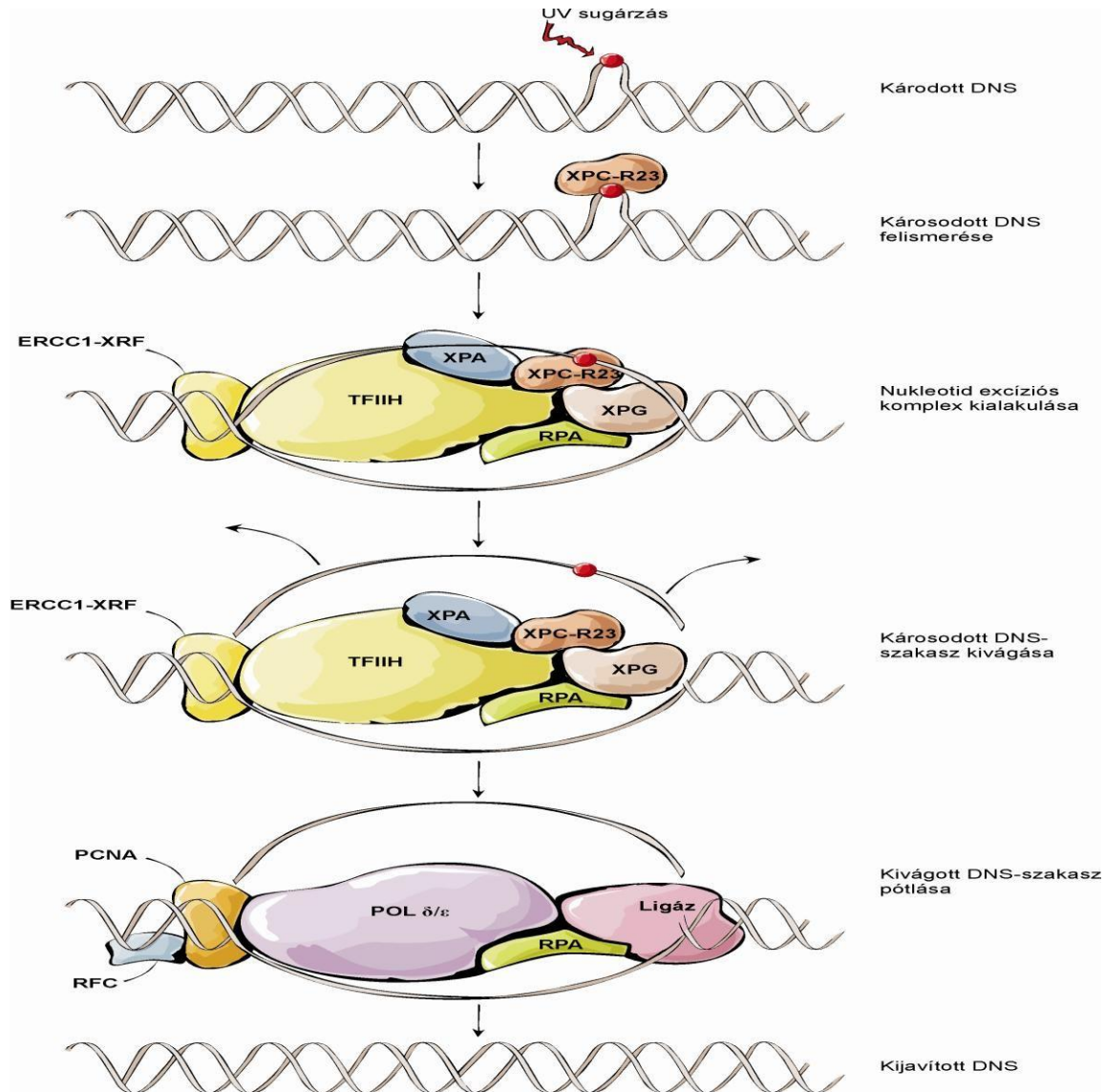
Epigenetic alterations

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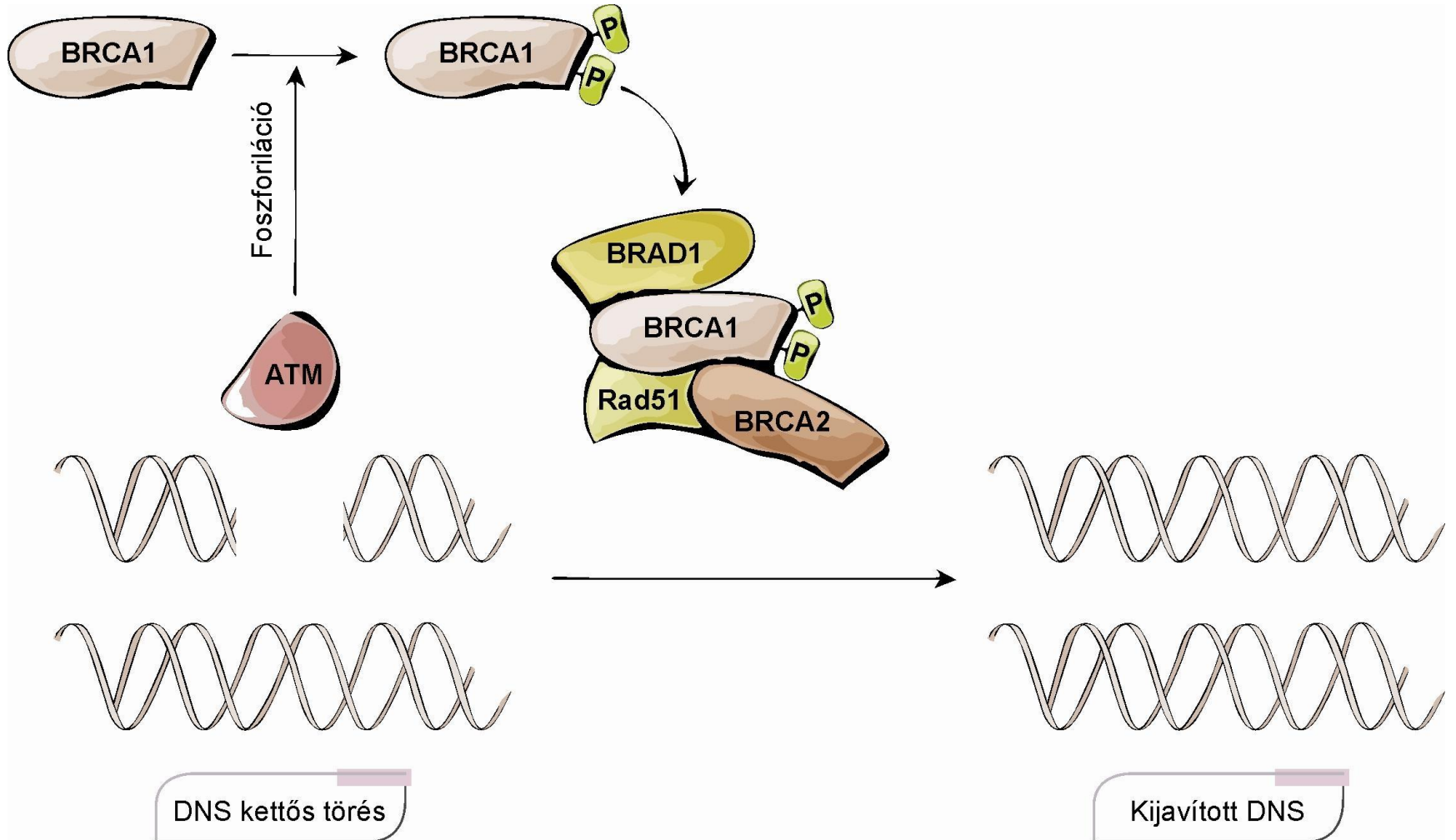
The role of DNA mismatch repair in cancer development



The role of nucleotide excision repair in cancer development



The role of homologous recombination repair in cancer development





The role of DNA repair gene mutations in cancer development

RPAIR SYSTEM	GENES	NEOPLASM
Mismatch repair	<i>MSH2</i> <i>MLH1</i> <i>MSH6</i>	Colorectal-, ovarium-, endometrium-, stomach cancer
Nucleotide excision repair	<i>XPA</i> <i>XPC</i> <i>XPG</i>	Basocellular-, planocellular carcinoma, melanoma
Homolog recombination repair	<i>BRCA1</i> <i>BRCA2</i> <i>ATM</i>	Breast-, ovarium-, prostata carcinoma Breast-, ovarium-, pancreas, stomach carcinoma, melanoma Ataxia teleangiectasia, breast carcinoma, different sporadic neoplasms



Molecular mechanisms involved into cancer development

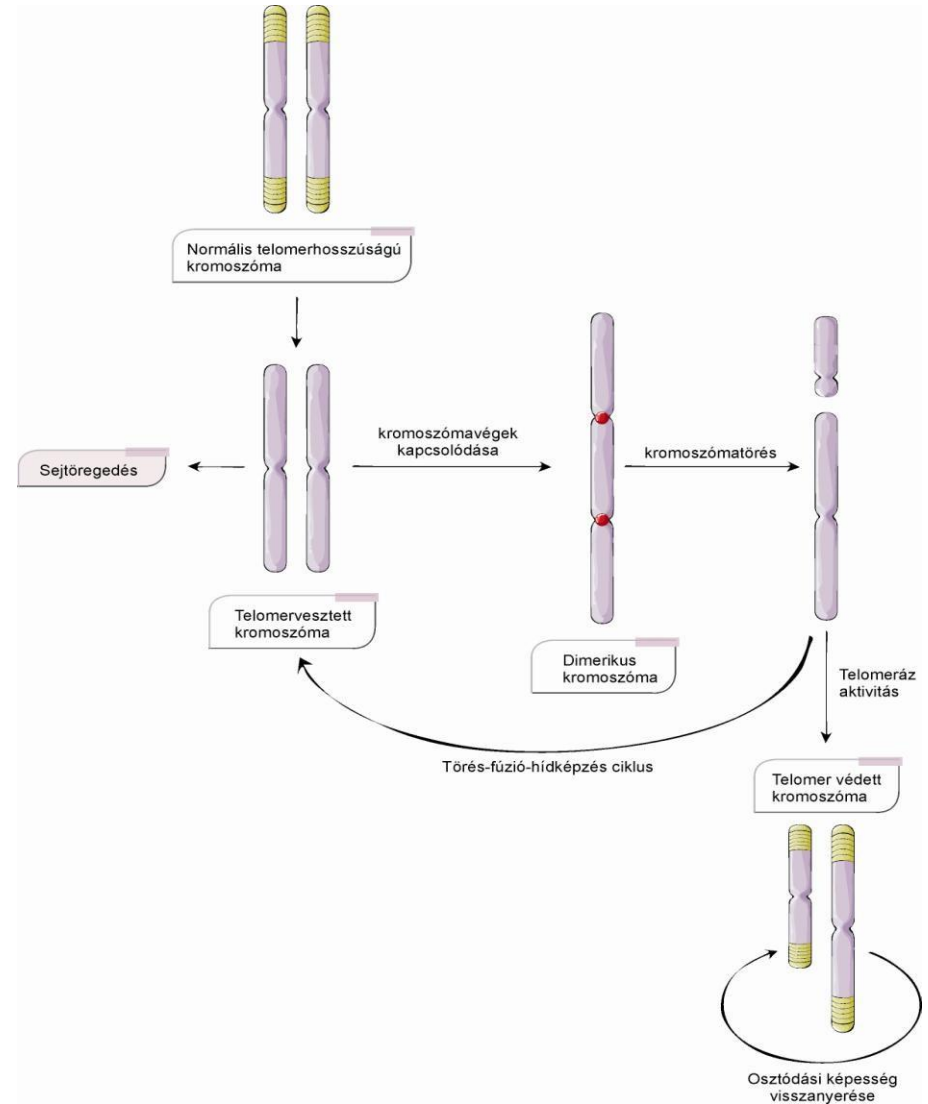
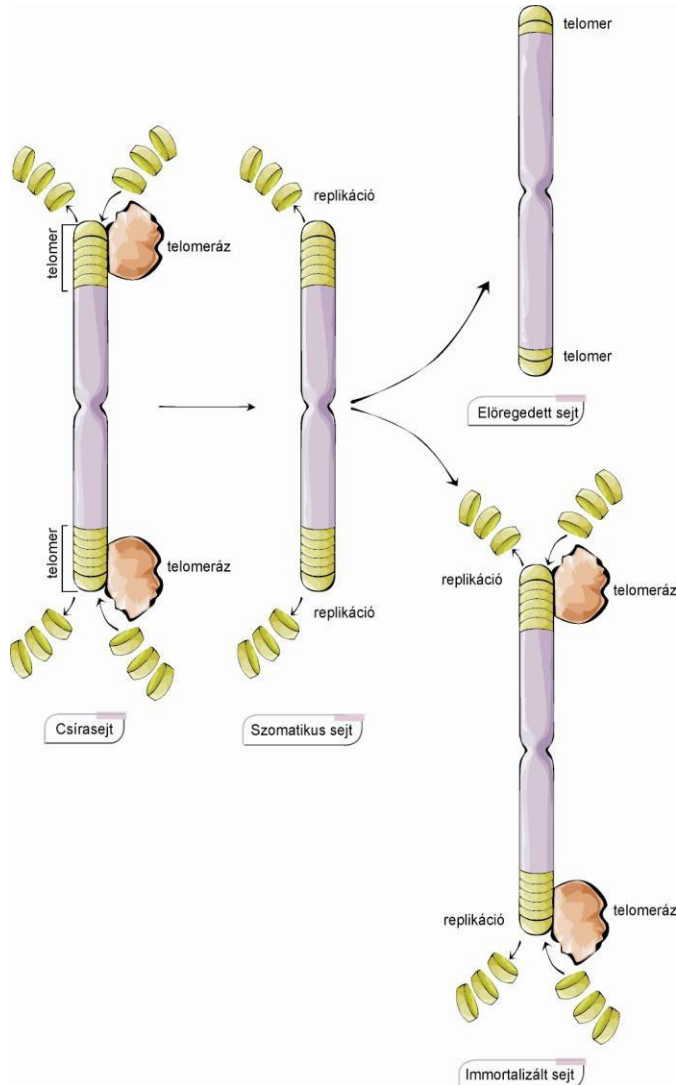
Genetic alterations

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Epigenetic alterations

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The role of telomere and telomerase in cancer development





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- Micro-RNA expression



Molecular mechanisms involved into cancer development

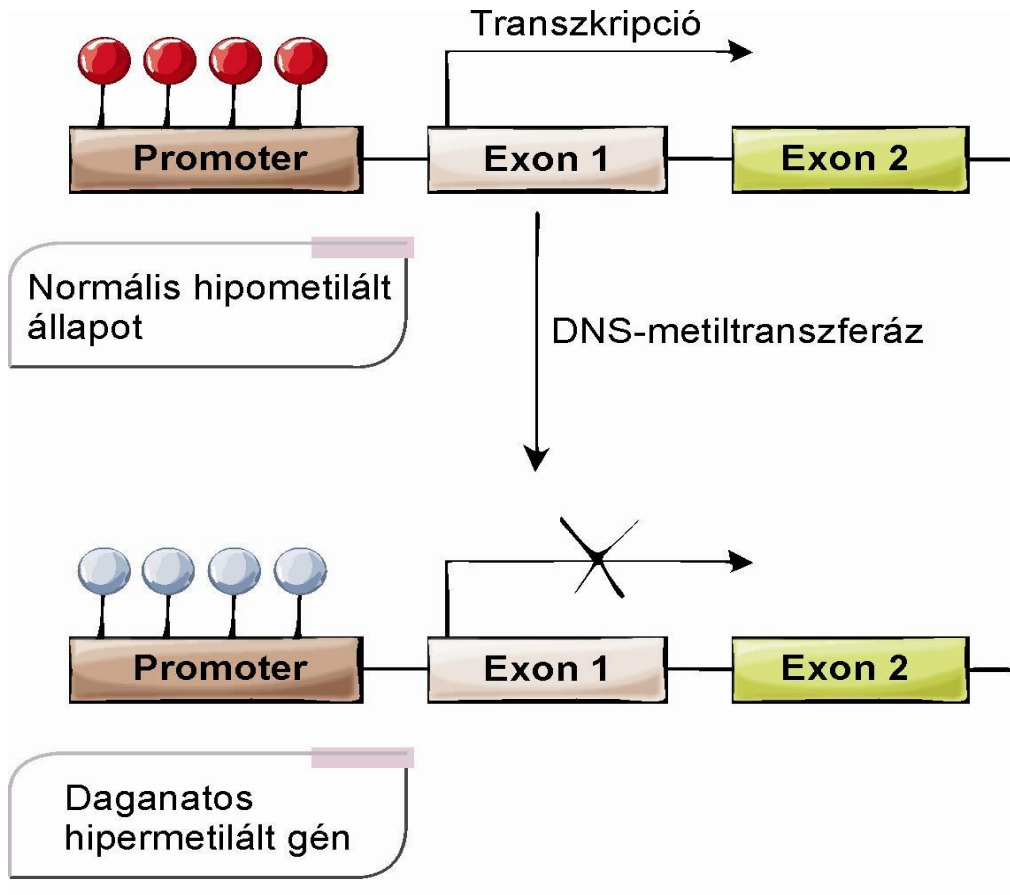
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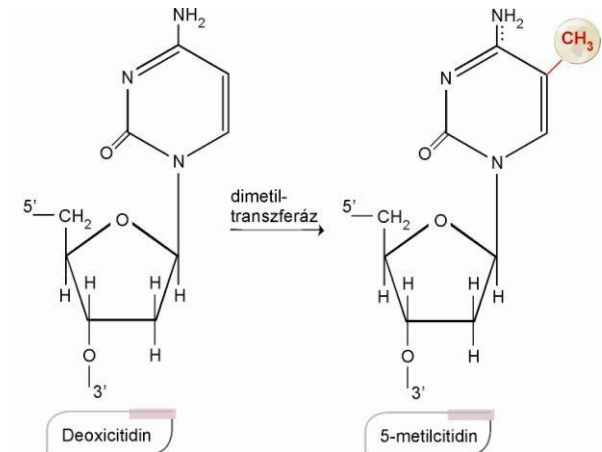
Epigenetic alterations

- DNA-methylation
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Hipermetiláció szerepe a daganatok kialakulásában



-  Hipometilált CpG
-  Hipermetilált CpG





Frequently hypermethylated genes in different cancers

GENE	NEOPLASM
<i>hMLH1</i>	Stomach-, colorectali-, ovarium carcinoma
<i>APC</i>	Colorectal carcinoma
<i>ARF</i>	Colorectal carcinoma
<i>RB</i>	Retinoblastoma
<i>CDKN2B</i>	Leukaemia
<i>CDKN2A</i>	Different neoplasmsípusok
<i>VHL</i>	Renal cell carcinoma
<i>p16INK4A</i>	Colorectal-, urinary bladder-, oesophagus-, lung carcinoma, lymphoma
<i>BRCA1</i>	Breast-, ovarium carcinoma



Molecular mechanisms involved into cancer development

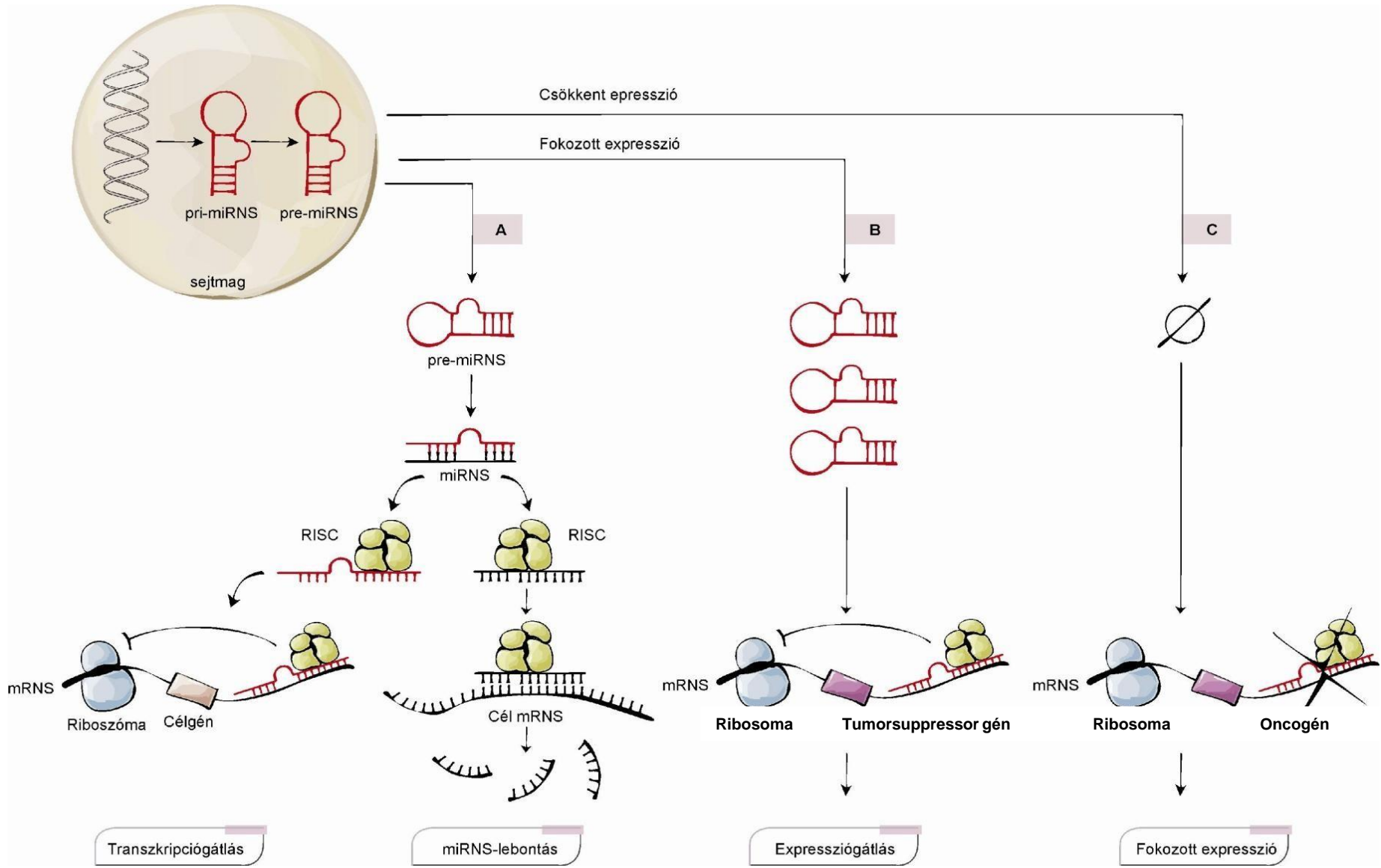
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The role of miRNS in the development of cancer



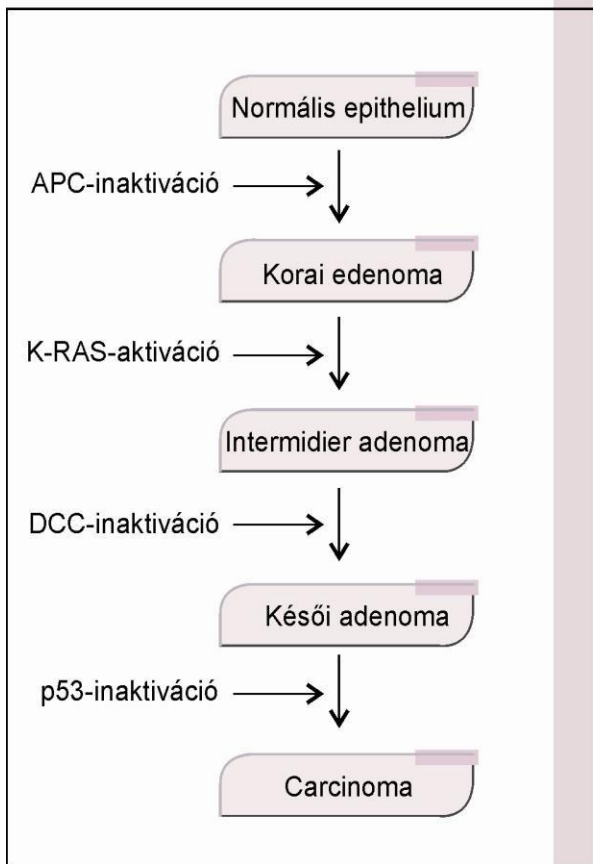


The role of miRNAs in the development of cancer

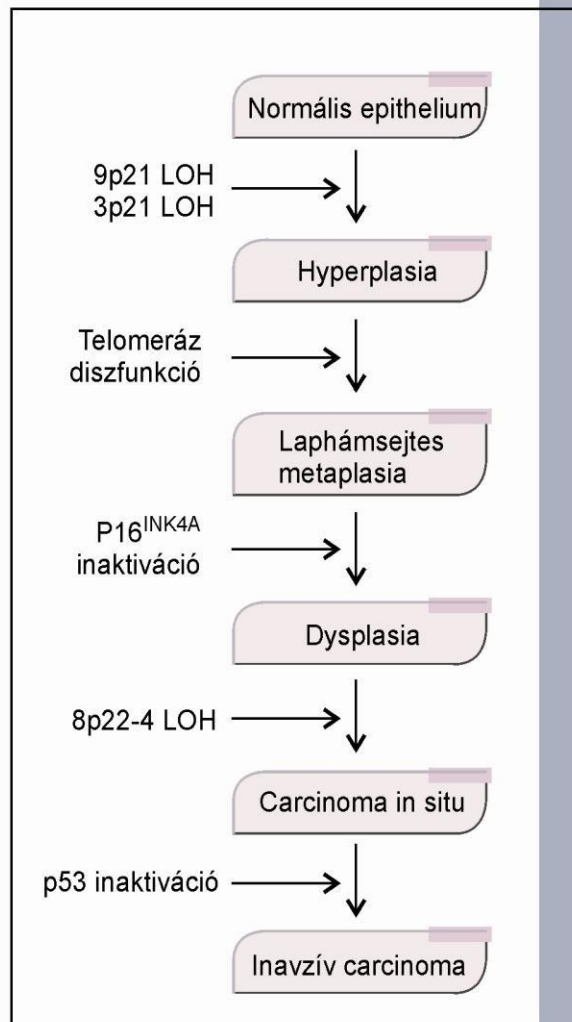
miRNA type	miRNA	Neoplasm
Onkogene miRNA	miR-17-92	Lung cancer
	miR-372-373	Testis germinal tumors
	miR-21	Glioblastoma, breast cancer
	miR-155	Breast-, colon-, lung cancer, B-cell lymphoma
	miR-146	Breast-, pancreas-, prostate cancer
	miR-121,122	Papillary thyroid cancer
Tumor suppressor miRNA	miR-127	Urinary bladder-, prostate cancer
	miR-15a	CLL
	miR-145	Colon-, breast cancer
	Let-7	Related to prognosis of lung cancer

Multistep genetic alteration in cancer development

A. Colonicarcinoma



B. Tüdő laphámsejtes carcinoma



C. Köpenysejtes lymphoma

