

## lidský genom

jádro:

3.2 miliardy páru bází DNA  
(2x)

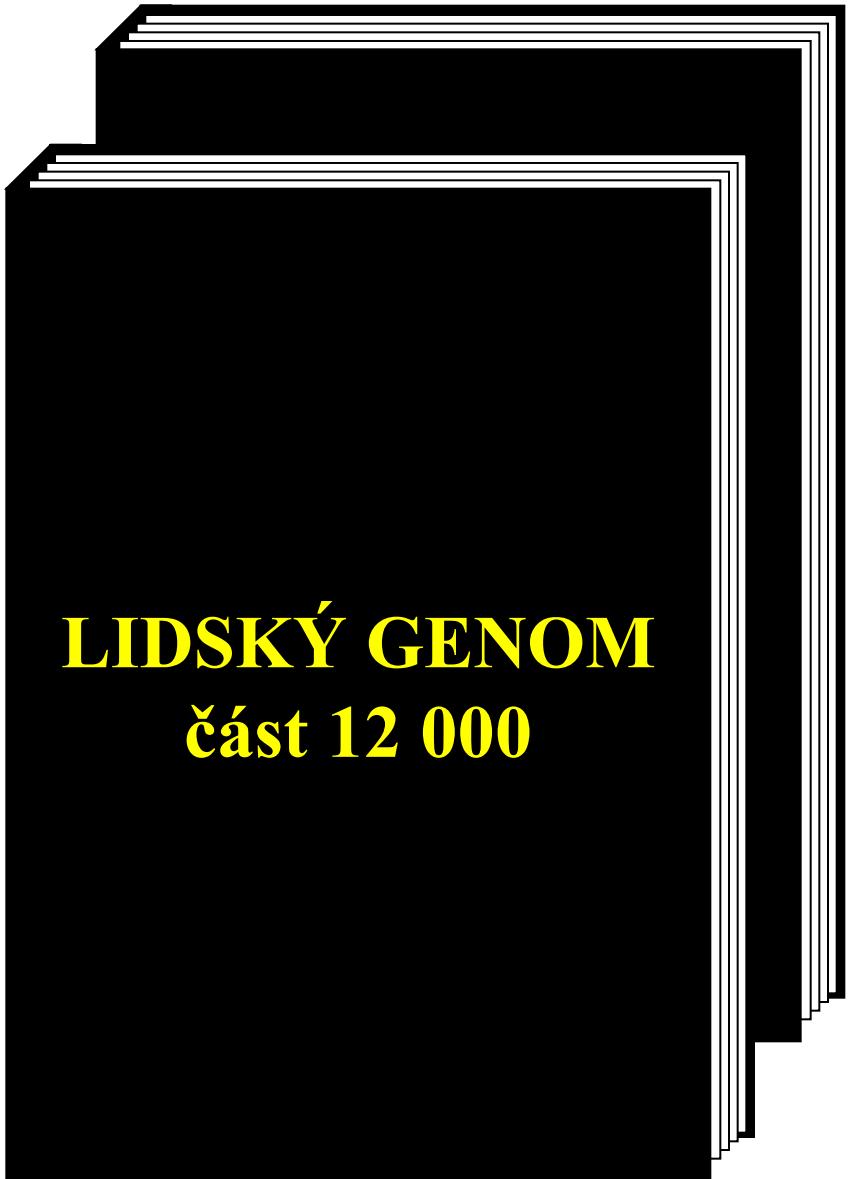
mitochondrie:

16.5 kb kruhová DNA  
(několik g/mt, stovky mt/b)  
geny pro 24 RNA + 13 proteinů

TGCTTGATGGATGGCCCTTAGCTAGCTCCTTTAGATCTAGTCGGCTAGCTATGCTTG  
GATCGATGGCCCTAGCTAGCTCCTTTAGATCTAGTCGGCTAGCTATGCTTGATCGA  
TGCCCTTAGCTAGCTCCTTTAGATCTAGTCGGCTAGCTATGCTTGATCGATGCC  
TTAGCTAGCTCCTTTAGATCTAGTCGGCTAGCTATGCTTGATCGATGCCCTAGCT  
AGCTCCTTTAGATCTAGTCGGCTAGCTATGCTTGATCGATGCCCTAGCTAGCTC  
TTTATAGATCTAGTCGGCTAGCTATGCTTGATCGATGCCCTAGCTAGCTCCTTTAG  
ATCTAGTCGGCTAGCTATGCTTGATCGATGCCCTAGCTAGCTCCTTTAGATCTAG  
TCGGCTAGCTATGCTTGATCGATGCCCTAGCTAGCTCCTTTAGATCTAGCTCCTTTAG  
AGCTAGTCGGCTAGCTATGCTTGATCGATGCCCTAGCTAGCTCCTTTAGATCTAG  
TCGGCTAGCTATGCTTGATCGATGCCCTAGCTAGCTCCTTTAGATCTAGCTCCTTTAG  
TAGCTATGCTTGATCGATGCCCTAGCTAGCTCCTTTAGATCTAGTCGGCTAGCTA  
TCCTGGATCGATGCCCTAGCTAGCTCCTTTAGATCTAGTCGGCTAGCTATGCTTG  
TCCTGGATCGATGCCCTAGCTAGCTCCTTTAGATCTAGTCGGCTAGCTATGCTTG  
GATCGATGCCCTAGCTAGCTAGCTCCTTTAGATCTAGTCGGCTAGCTATGCTTGATCGA  
TGCCCTTAGCTAGCTCCTTTAGATCTAGTCGGCTAGCTATGCTTGATCGATGCC  
TTAGCTAGCTCCTTTAGATCTAGTCGGCTAGCTATGCTTGATCGATGCCCTAGCT  
AGCTCCTTTAGATCTAGTCGGCTAGCTATGCTTGATCGATGCCCTAGCTAGCTC  
TTTATAGATCTAGTCGGCTAGCTATGCTTGATCGATGCCCTAGCTAGCTCCTTTAG  
ATCTAGTCGGCTAGCTATGCTTGATCGATGCCCTAGCTAGCTCCTTTAGATCTAG  
TCGGCTAGCTATGCTTGATCGATGCCCTAGCTAGCTCCTTTAGATCTAGTCGGC  
TAGCTATGCTTGATCGATGCCCTAGCTAGCTCCTTTAGATCTAGTCGGCTAGCTA  
TGCTTGATCGATGCCCTAGCTAGCTCCTTTAGATCTAGTCGGCTAGCTATGCTTG

1 strana

1 800 znaků



## LIDSKÝ GENOM

### část 12 000

1 kniha

150 stran

270 000 znaků

tloušťka 2 cm

---

celý genom

3.2 miliardy znaků

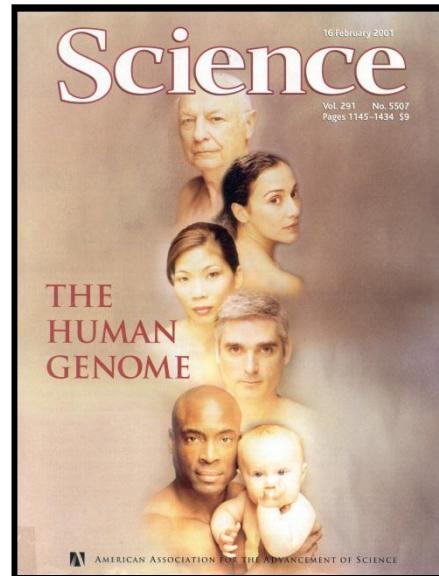
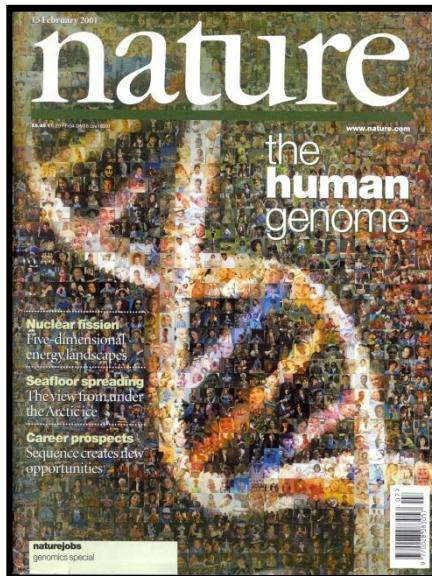
1.8 milionu stran

12 000 knih



[http://rene.spika.cz/prague\\_january/petrin\\_tower\\_1.html](http://rene.spika.cz/prague_january/petrin_tower_1.html)

NCBI: <http://www.ncbi.nlm.nih.gov/mapview/>  
Sanger/EBI: <http://www.ensembl.org/>  
UCSC: <http://genome.ucsc.edu>





PubMed

Entrez

BLAST

OMIM

Taxonomy

Structure

Search

Find in This View

Find

Advanced Search

MapViewer Home

Map Viewer Help

Human Maps Help

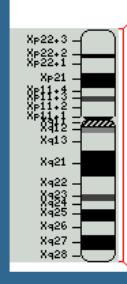
FTP

Chr. X Resource

Data As Table View

**Maps&Options**Compress Map 

Region Shown:


**Homo sapiens Map View build 31** **BLAST the Human Genome**Chromosome: [1](#) [2](#) [3](#) [4](#) [5](#) [6](#) [7](#) [8](#) [9](#) [10](#) [11](#) [12](#) [13](#) [14](#) [15](#) [16](#) [17](#) [18](#) [19](#) [20](#) [21](#)22 [X] [Y](#)

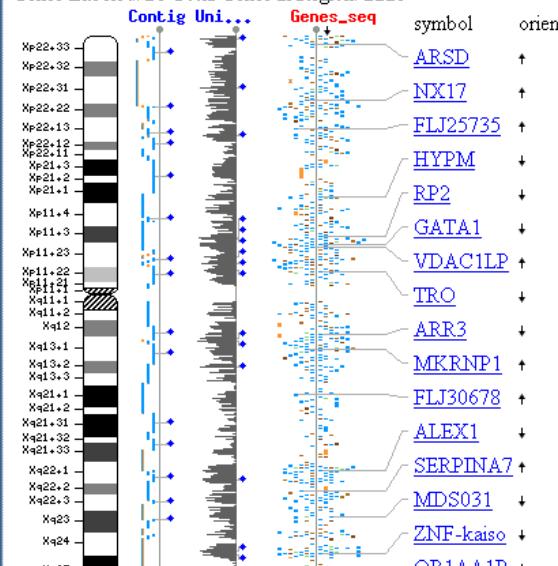
Master Map: Genes On Sequence

**Maps & Options**

Total Genes On Chromosome: 1110

Region Displayed: 0-151M bp [Download/View Sequence/Evidence](#)

Genes Labeled: 20 Total Genes In Region: 1110

**FEATURES****source**

1..1251 /organism="Mus musculus" /mol_type="mRNA" /db_xref="taxon:10090" /clone="MGC:58456 IMAGE:6770406" /tissue_type="Testis, mouse" /clone_lib="NIH_MGC_169" /lab_host="DH10B" /note="Vector: pDNR-LIB"	1..1251 /gene="DOH6S2654E" /note="synonyms: X5L, XAP-5-like" /db_xref="LocusID: <a href="#">108161</a> " /db_xref="MGI: <a href="#">1351640</a> " 149..1153 /codon_start=1 /product="DOH6S2654E protein" /protein_id=" <a href="#">AAH49659.1</a> " /db_xref="GI:29436863" /db_xref="LocusID: <a href="#">108161</a> " /translation="MAQYKGTMREAGRAMHLIKKREKQKEQMEVLKQRIAEETIMKSKVDDKKFSAHYDAVEAELKSSTVGLVTLDNMKAKQEALLREREMQLAKREQUEQRRIQLEMLREKERRRERKRKISNLSFTLDEEEGQEDSRQAESAEVHSAGAKKNLKGKPNPDVDTSFLLPDREREEEENRLREELRQEWEAKREKVKGEEVEITFSYUDGSGRRTVRMSKGSTVQQFLKRALQGLRRDFRELRAAGVEQLMYVKEDLILPHYHTFYDFIVAKARGKSGPLSFDVHDDVRLLSDATMEKDESHAGKVVLRSWYEKNKHIFPASRWEPYDPEKKWDRTIR"
--	---

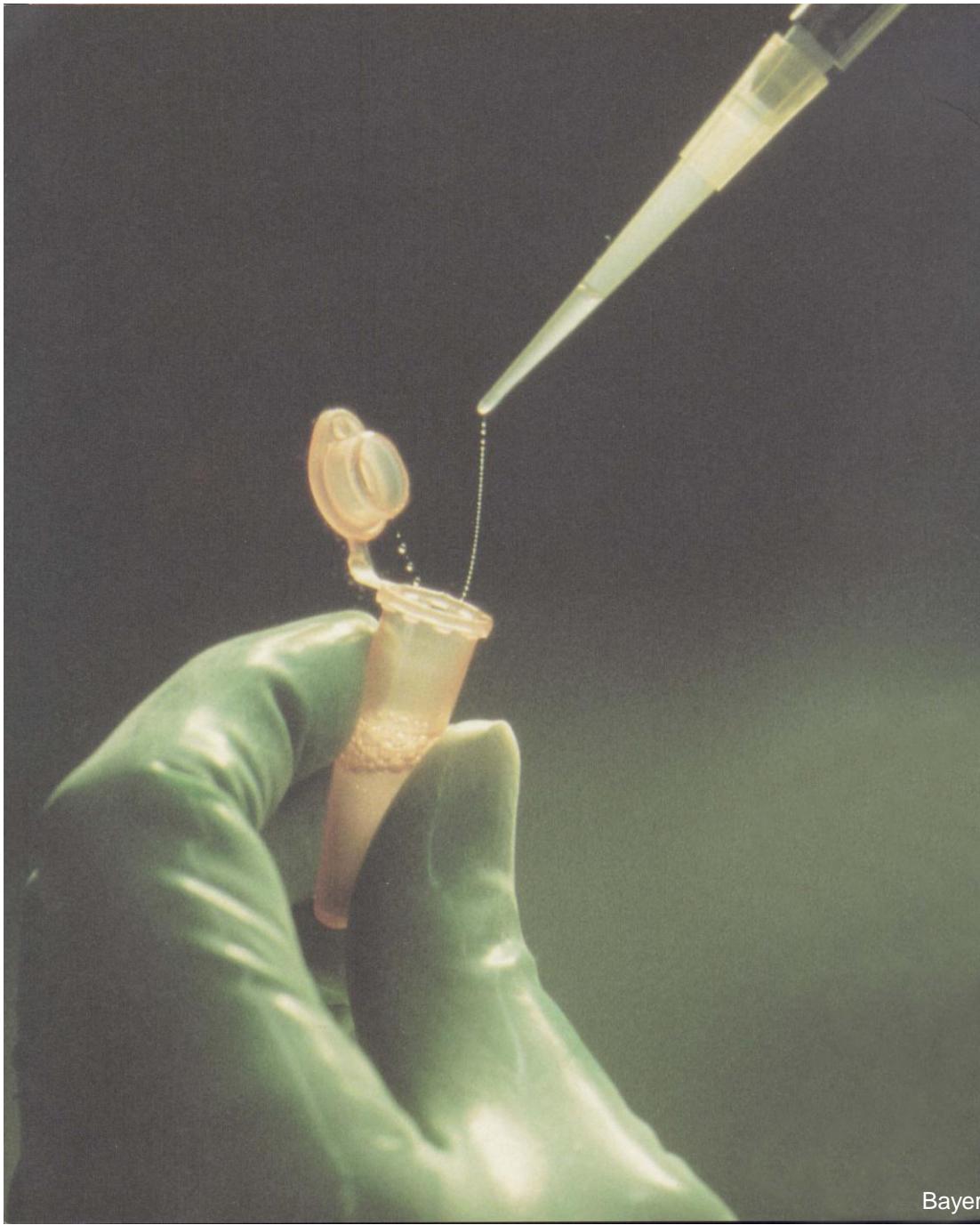
**CDS****gene****ORIGIN**

```

1 agaacaaaca atattcgcaa aagcaccaga aggaagagtc ttggctcata catcaaaaac
61 tgcagaatct gtgaactgac atcagacccca gaaggctacc agaaacaggg actggggcagg
121 ccaaaaagcc ttgcgtgaa ctgcaggcat ggccgcgtac aaaggccacca tgccccaaac
181 tggccggccc atgcacctga tcaagaagcg tgagaagcg aaggagcgat tggaggtgtct
241 gaaggcgcgc atgcaggagg agaccatcat gaagtcaaaa gtggacaaga agttctcgcc
301 acatctggac gccgtggagg ccggactgaa gtcccgatcg gtggggctgg tgaccctgaa
361 tgatcatgaa gcacaggagg agggccctgtct gaggagcgat gagatgcgc tgcccaaggag
421 ggaggcgtcg gagcaacgcgc ggatcacgtct ggagatgtcg cgccgagaagg agcgaaggcg
481 agaggcgtcg cgccaaatct ccacacgttc ttccacgttg gacgaggaaag aagggtacca
541 agaggacaggc cgccaaaggccg agatgtcccgaa ggtccacatgt gctggagccaa agaagaactt
601 gggcaagaat cccgtgtgg acacgagctt cctggccacac cgccgaggccg aggaggaggg

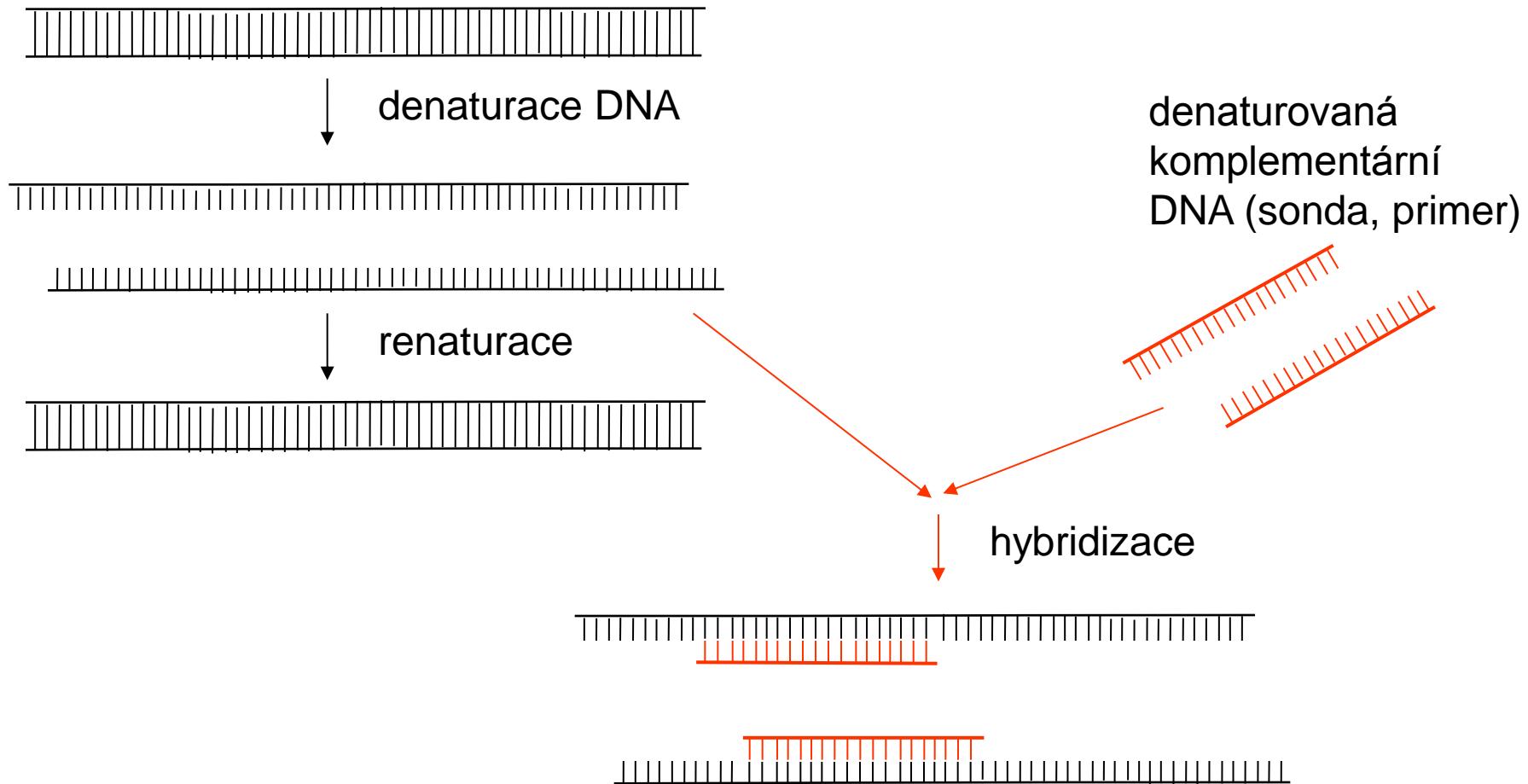
```

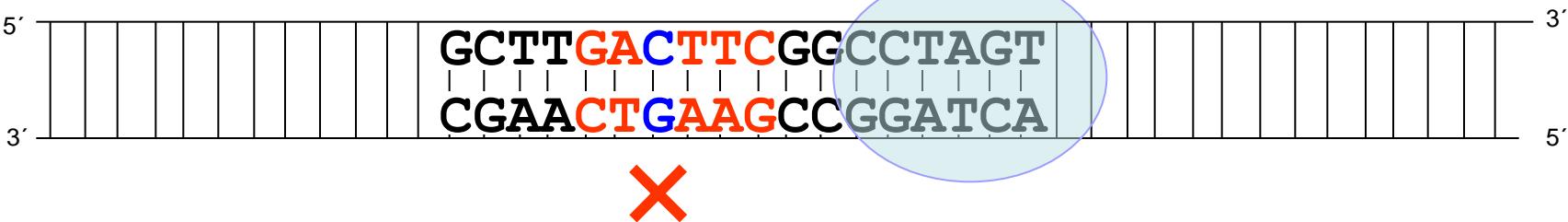
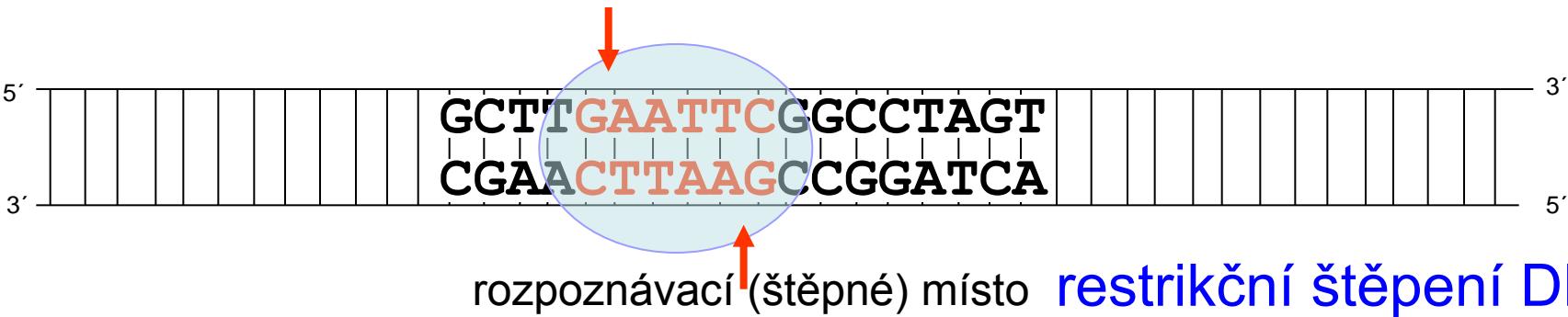
izolace  
DNA



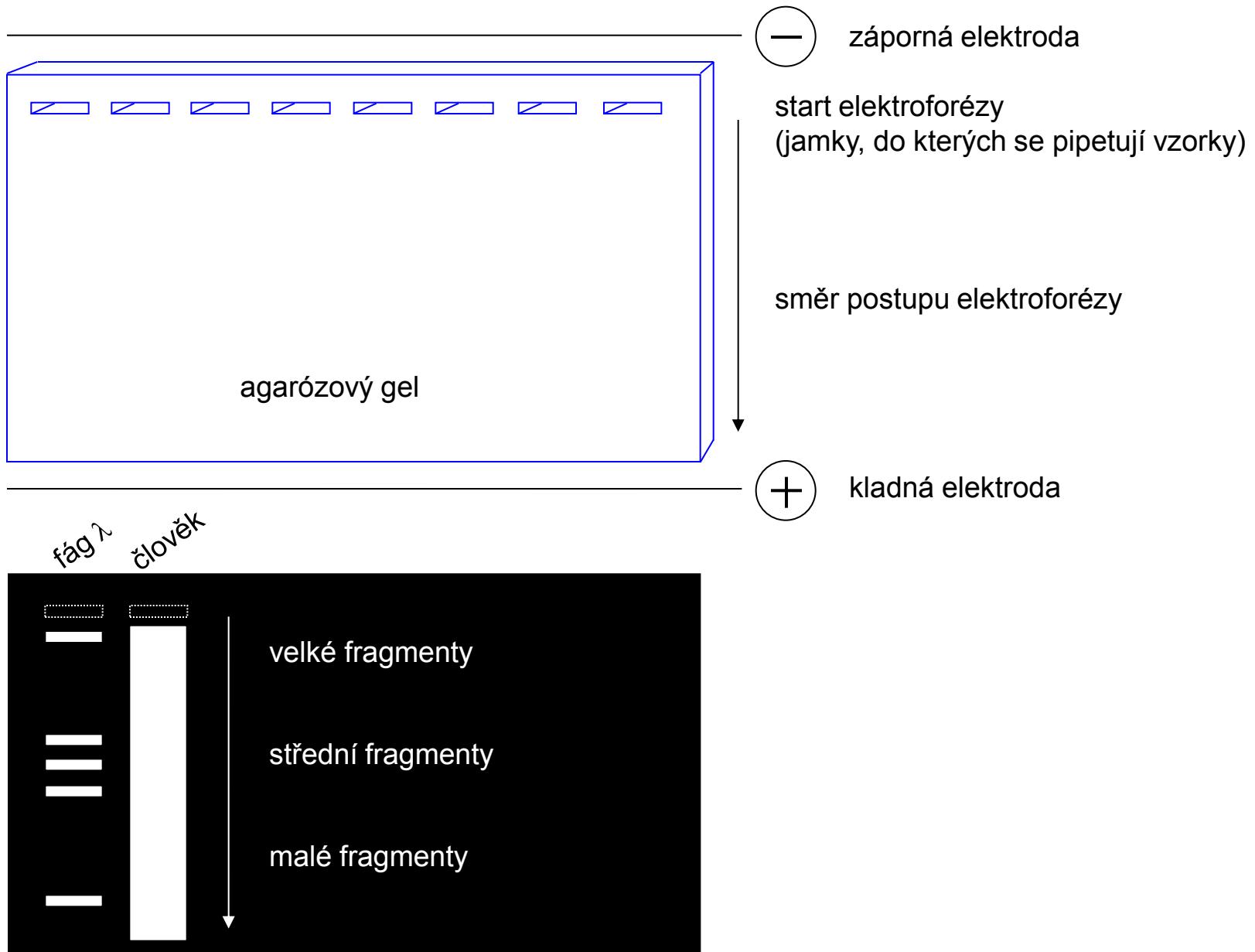
Bayer

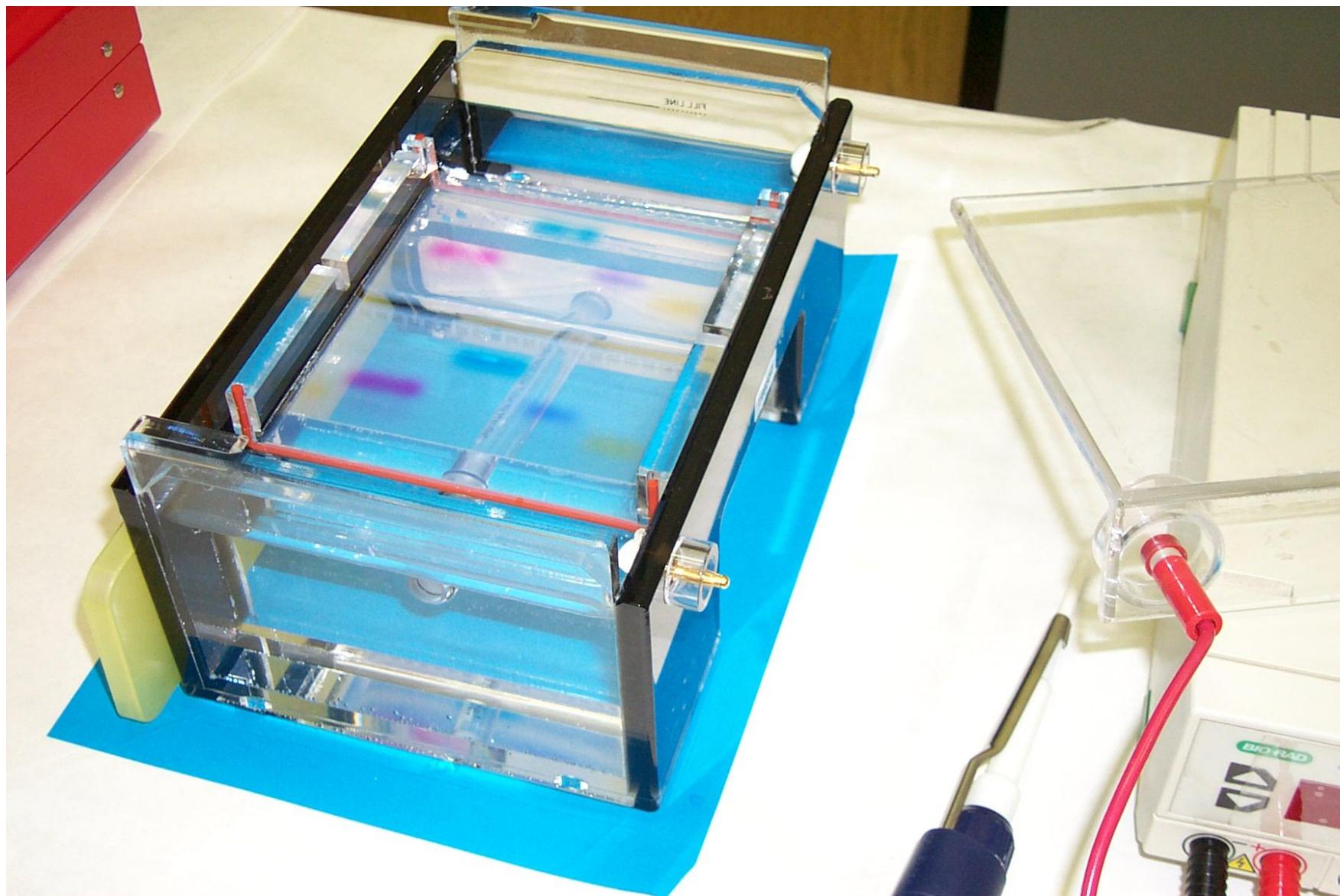
# hybridizace DNA

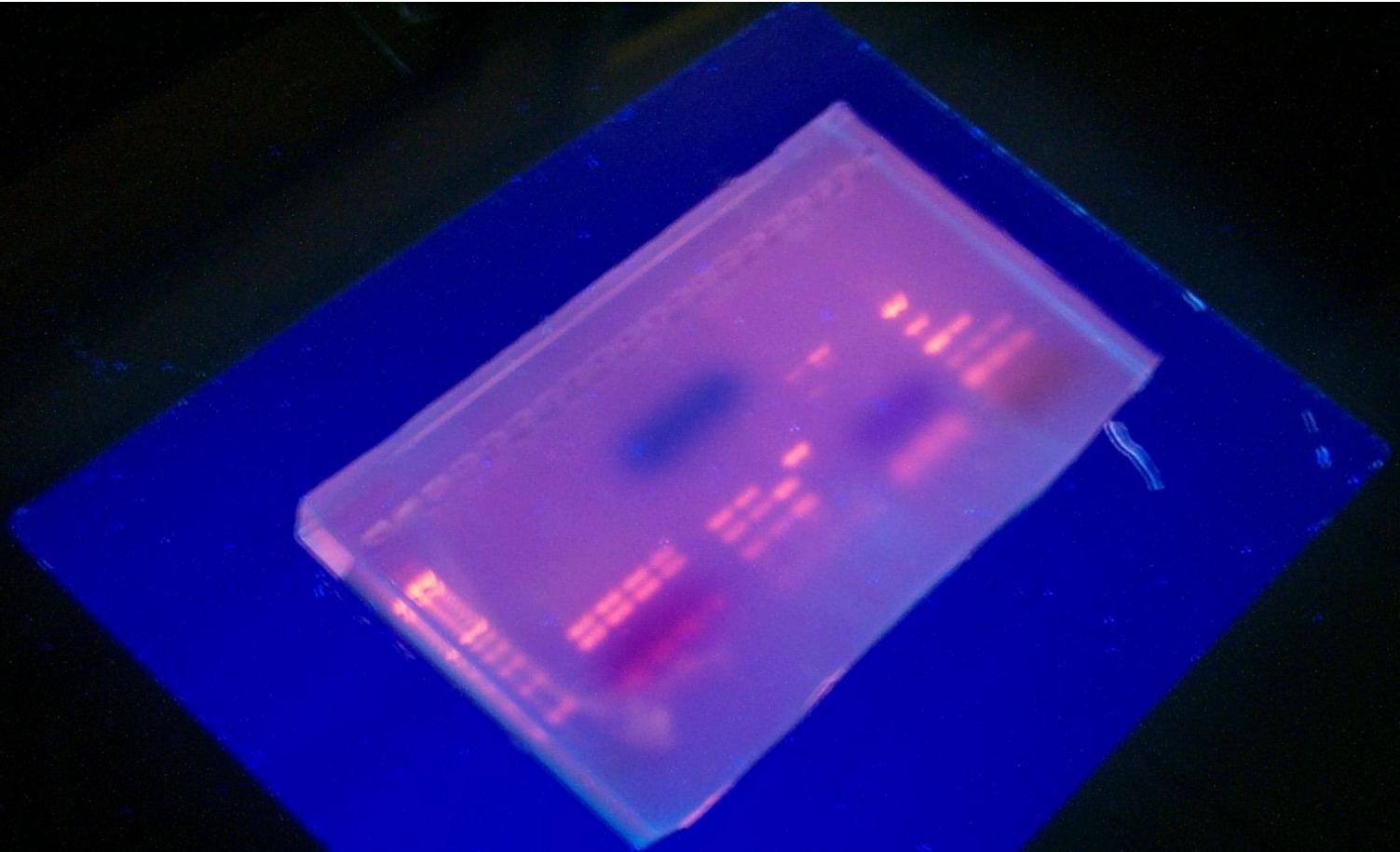


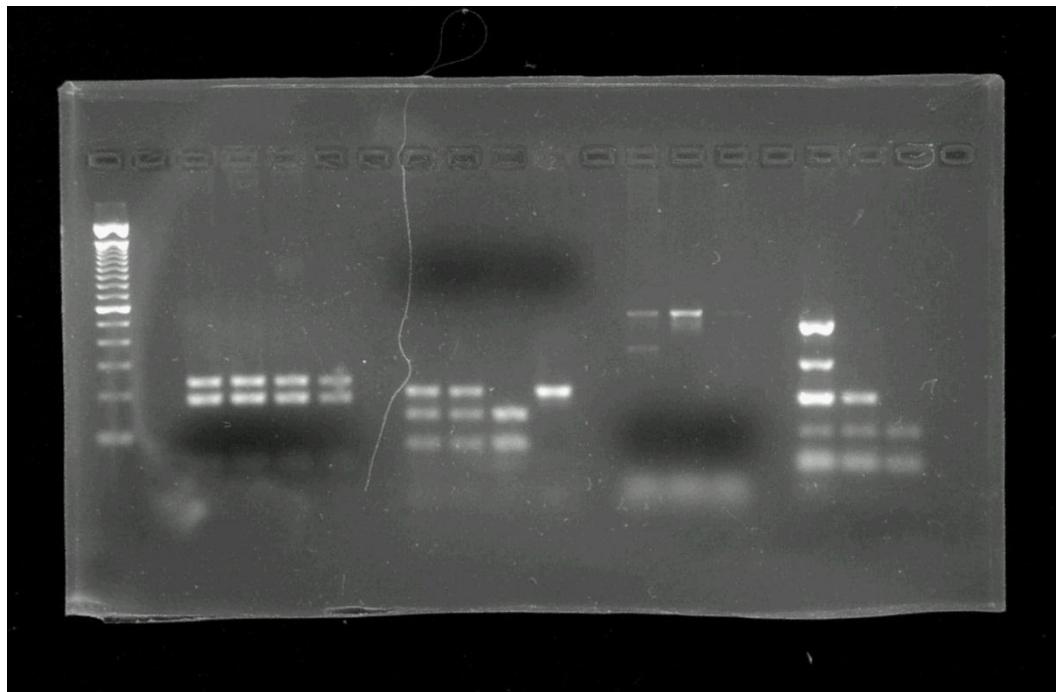


# gelová elektroforéza DNA

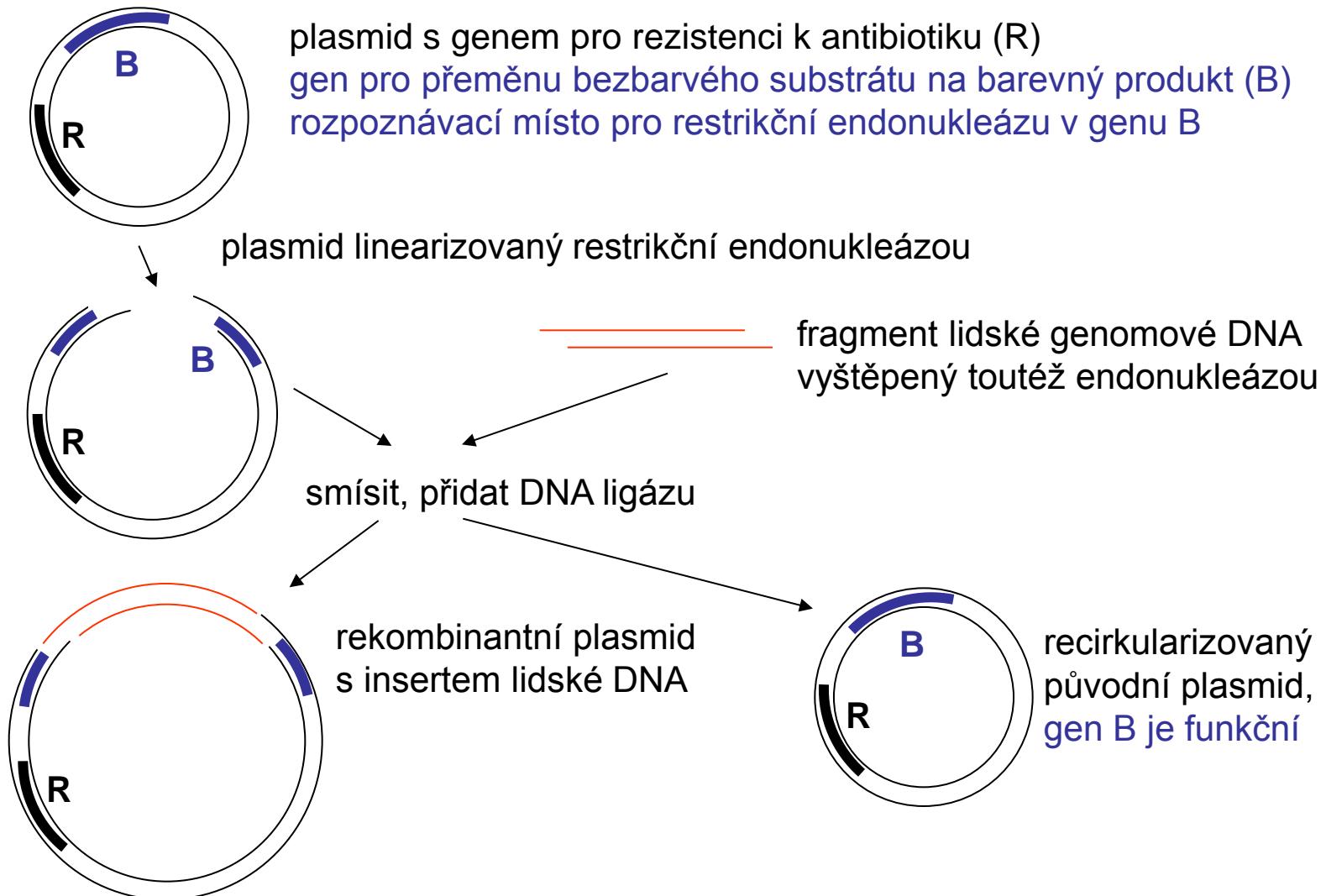


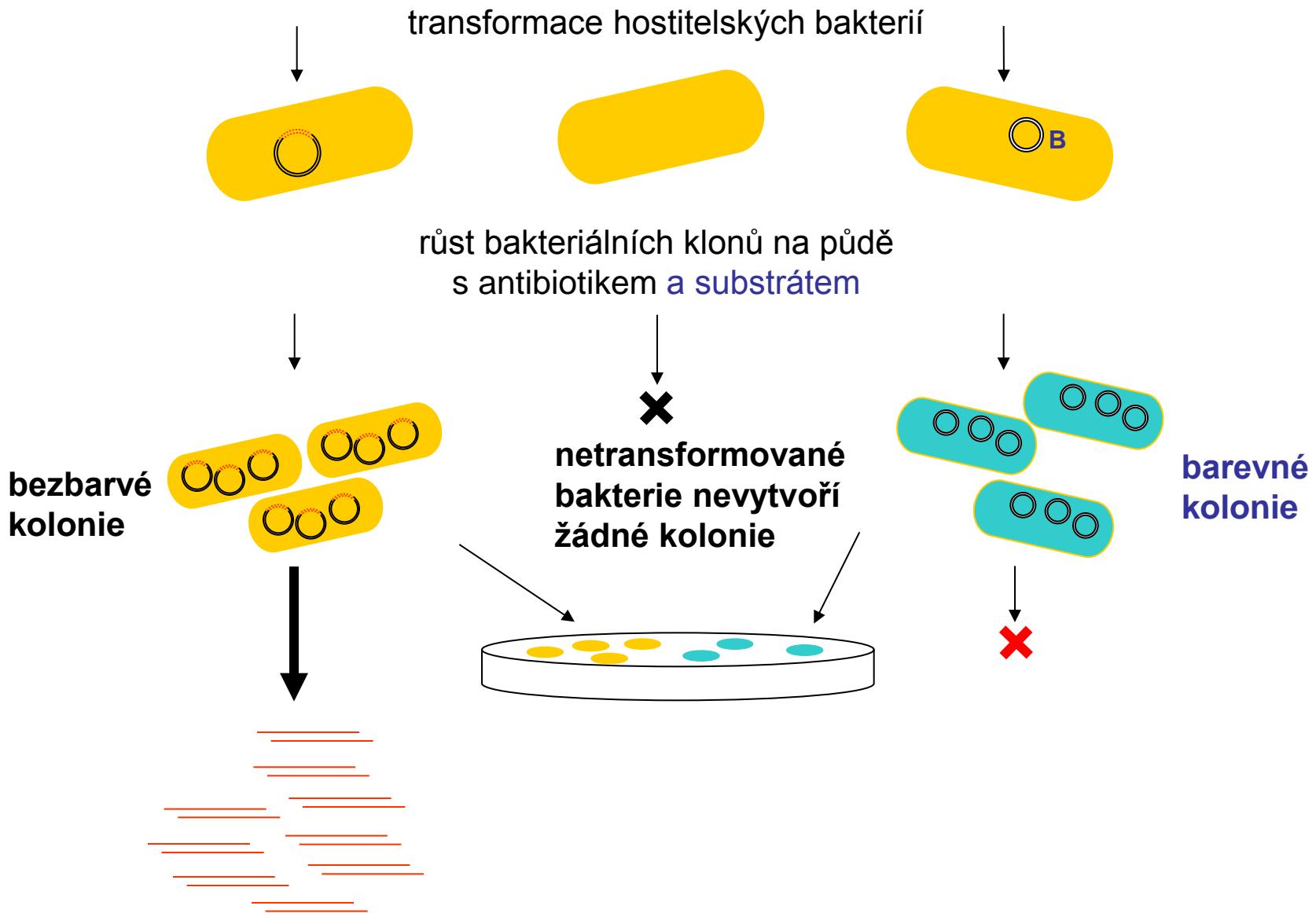


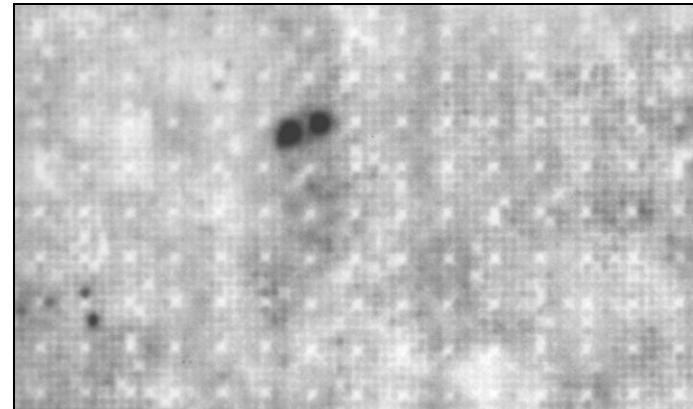
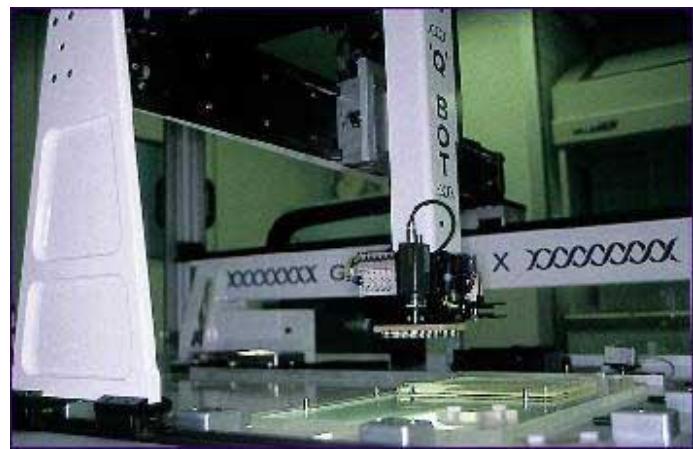
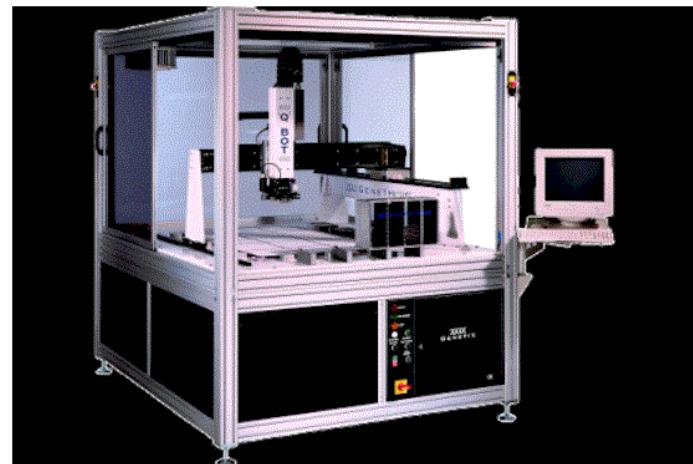




# klonování DNA

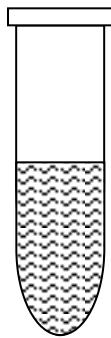
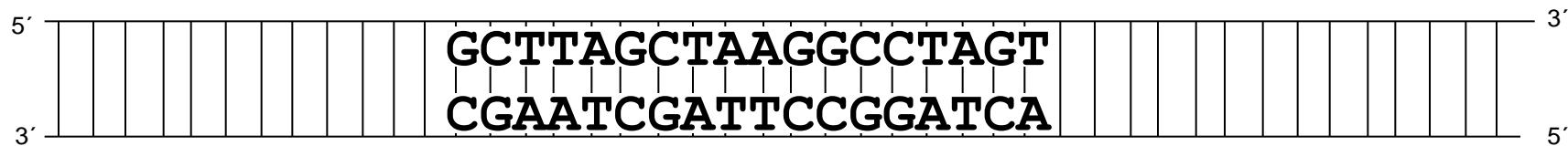




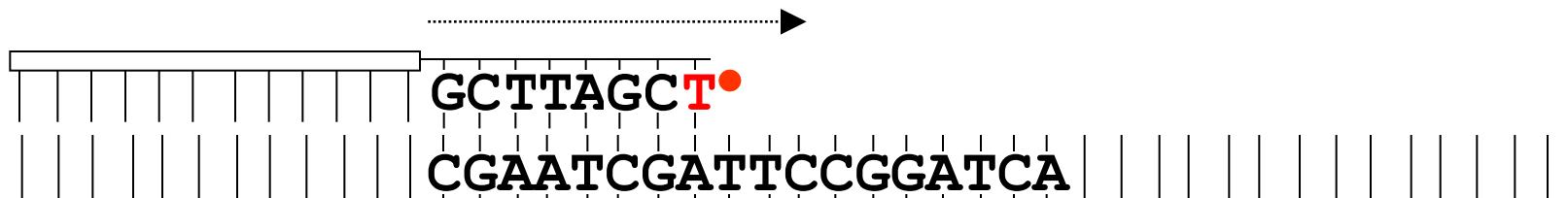


knihovny DNA

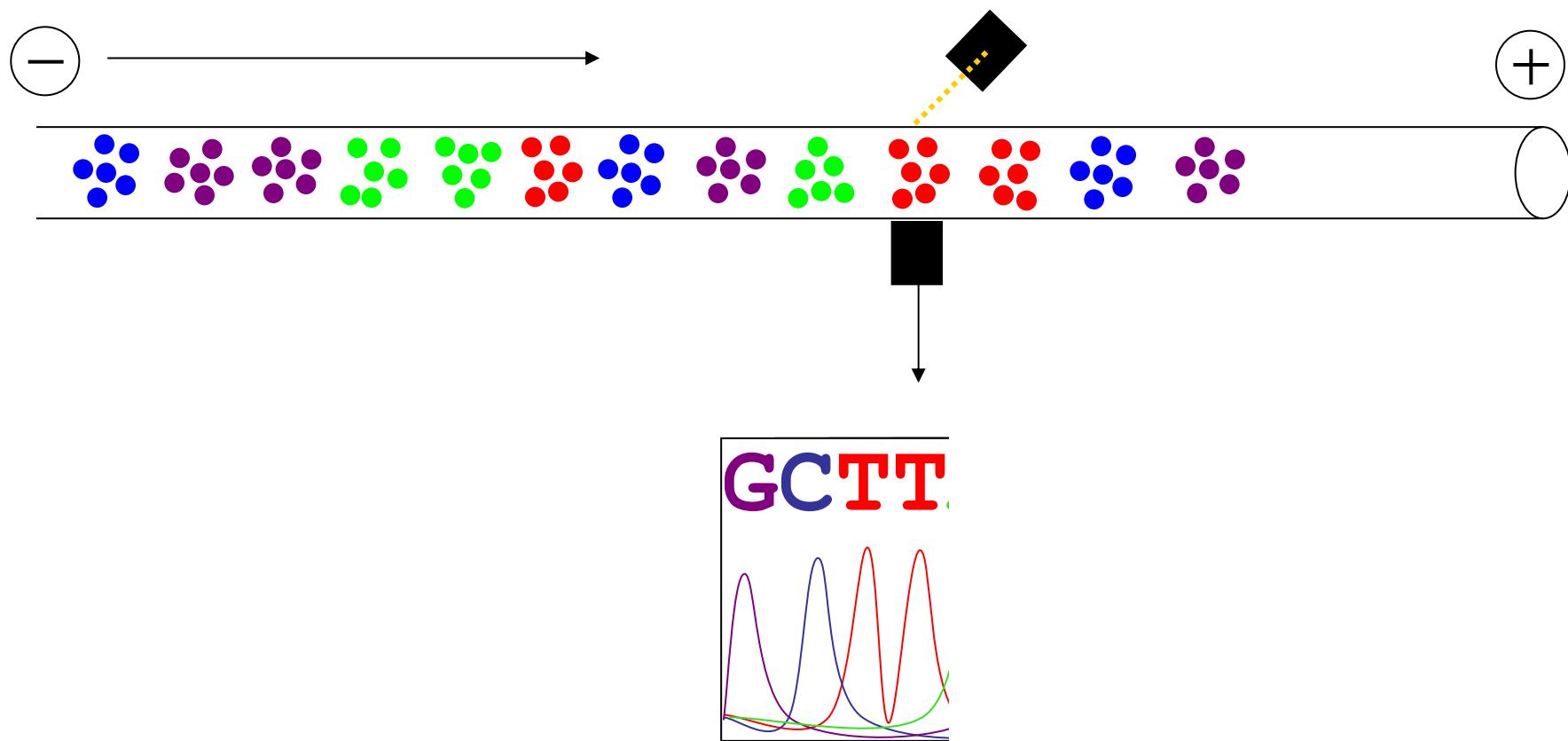
# sekvenování DNA



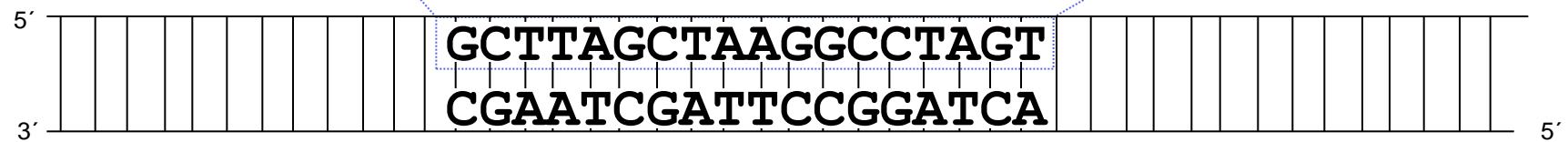
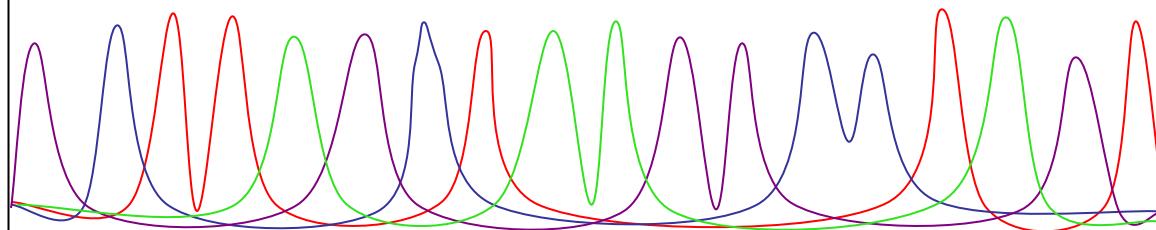
denatuovaná DNA  
+ primer + polymeráza  
+ deoxynukleotidy G, A, T, C  
+ dideoxynukleotidy **G<sup>•</sup>**, **A<sup>•</sup>**, **T<sup>•</sup>**, **C<sup>•</sup>**

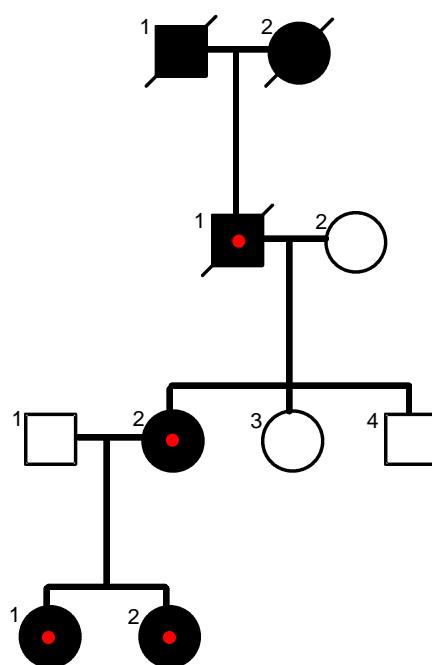


GCTTAGCTAAGGCCTAG<sup>T</sup>•  
GCTTAGCTAAGGCCTAG<sup>G</sup>•  
GCTTAGCTAAGGCCTA<sup>A</sup>•  
GCTTAGCTAAGGCC<sup>T</sup>•  
GCTTAGCTAAGGCC<sup>C</sup>•  
GCTTAGCTAAGGC<sup>C</sup>•  
GCTTAGCTAAGG<sup>G</sup>•  
GCTTAGCTAA<sup>G</sup>•  
GCTTAGCTAA<sup>A</sup>•  
GCTTAGCTA<sup>A</sup>•  
GCTTAGCT<sup>T</sup>•  
GCTTAGC<sup>C</sup>•  
GCTTAG<sup>G</sup>•  
GCTTA<sup>A</sup>•  
GCT<sup>T</sup>•  
GCT<sup>T</sup>•  
GC<sup>C</sup>•  
G<sup>G</sup>•

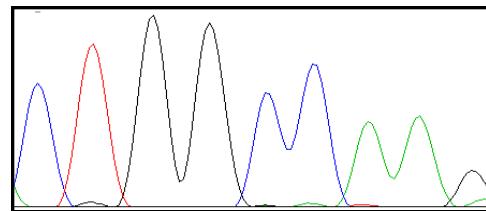


**GCTTAGCTAAGGCCTAGT**



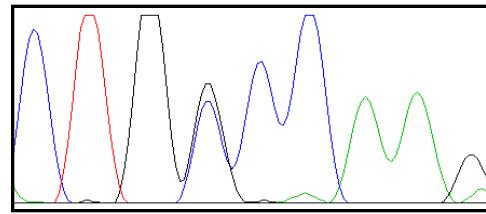


normální sekvence



C	T	G	G	C	C	A	A	G
137		138		139				
Leu		Ala		Lys				

heterozygot

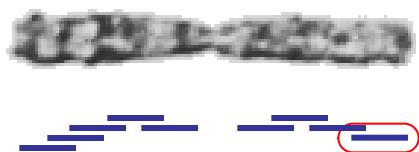


C	T	G	G	C	C	A	A	G
137		138		139				
Leu		Ala		Lys				
<b>Pro</b>								

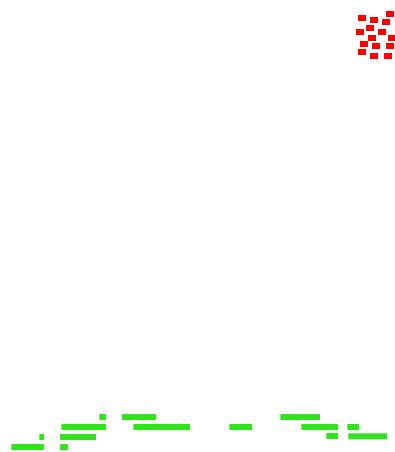


mezinárodní konsorcium

DNA 12 anonymních dárců

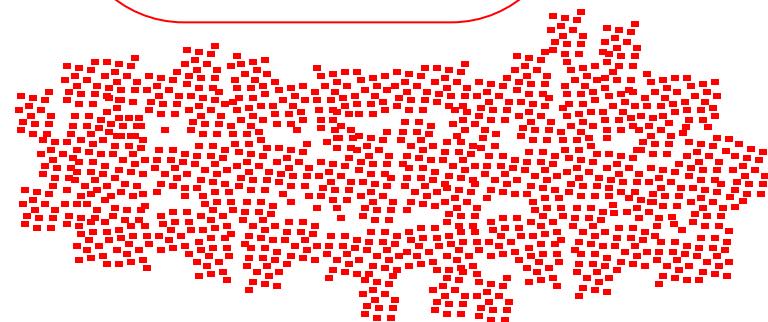


mapování



Celera Genomics

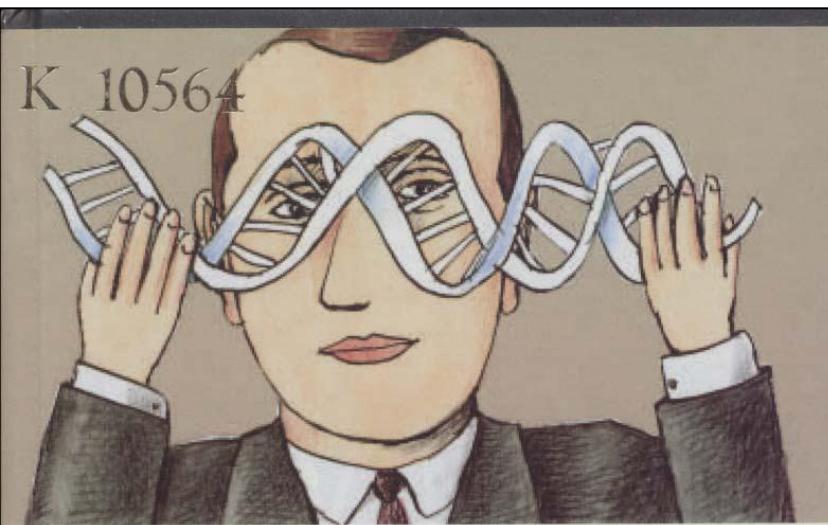
5 dárců různé etnicity



whole-genome shotgun



paired-end sequencing, scaffolds



KEVIN  
DAVIES

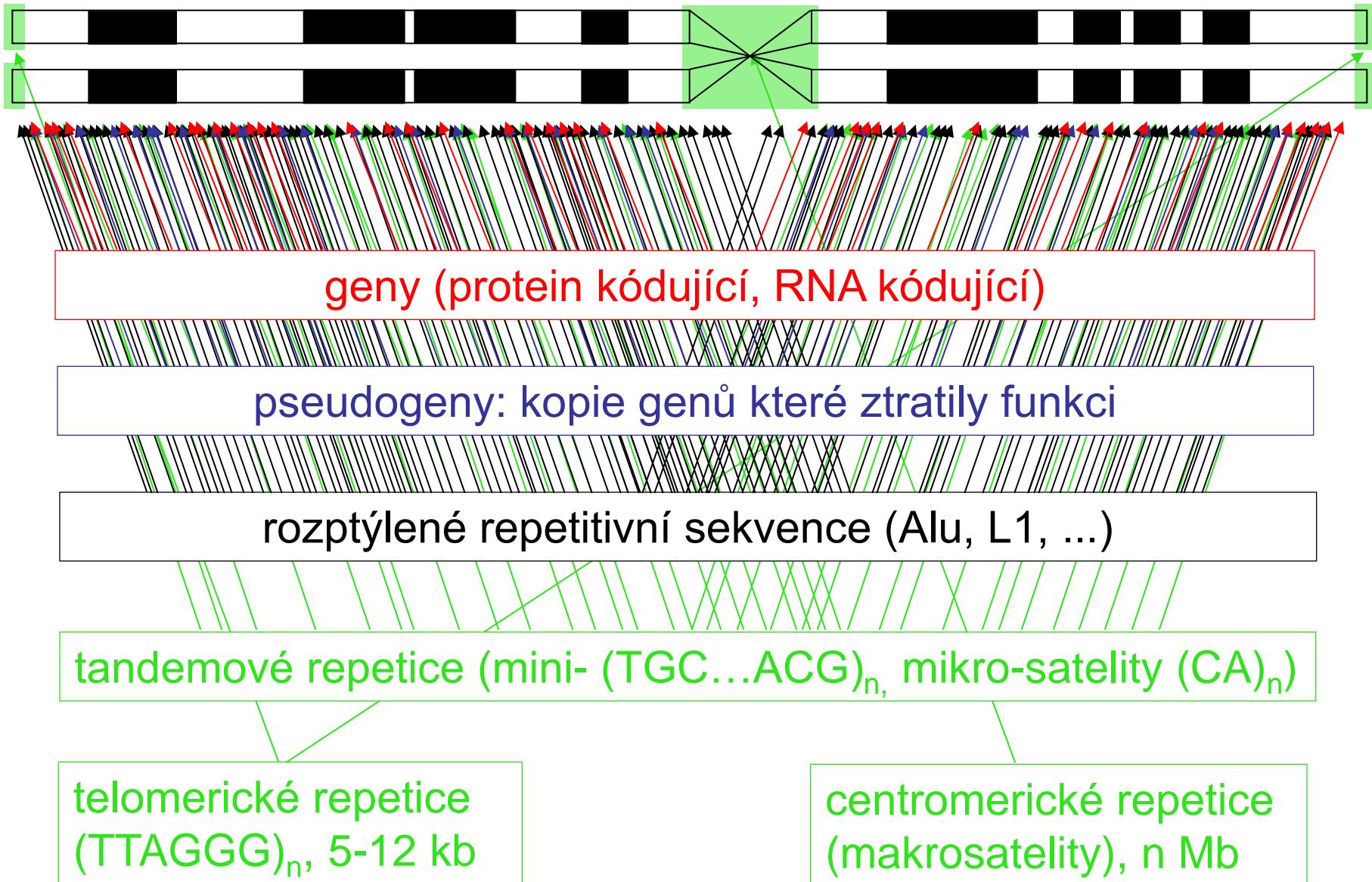
Rozluštěný  
genom

Příběh  
největšího  
vědeckého  
objevu  
naší doby

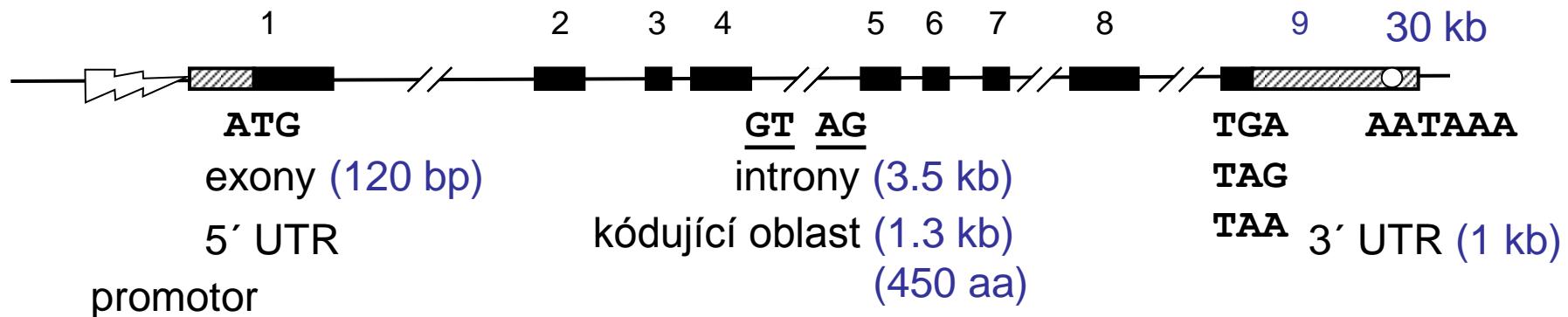
K A

FÉNI

# typy sekvencí v lidském genomu



## geny kódující proteiny



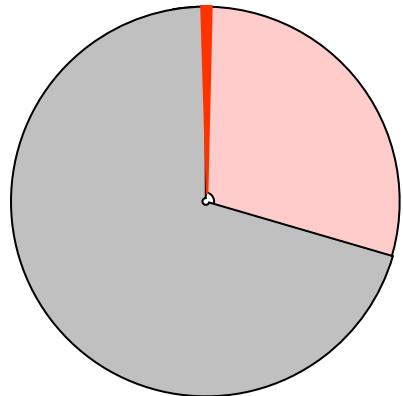
nejdelší gen: dystrofin (2.5 Mb, 89 exonů)  
nejvíce exonů: titin (363 exonů)

## genové rodiny

počet genů: méně než 25 tisíc

## geny kódující nekódující RNA (ncRNA)

## pseudogeny



geny (exony + introny) cca. 1/4

exony protein kódujících genů cca. 1.5%

mezigenová DNA



‘junk DNA’ 98.5%

podstata, původ a funkce ‘junk DNA’ ?

**2500**



**1**

nepravidelná distribuce genů v genomu

**600**



**13**

**1500**



**19**

**300**



**21**

**700**



**22**

**400**



**18**



původ genů

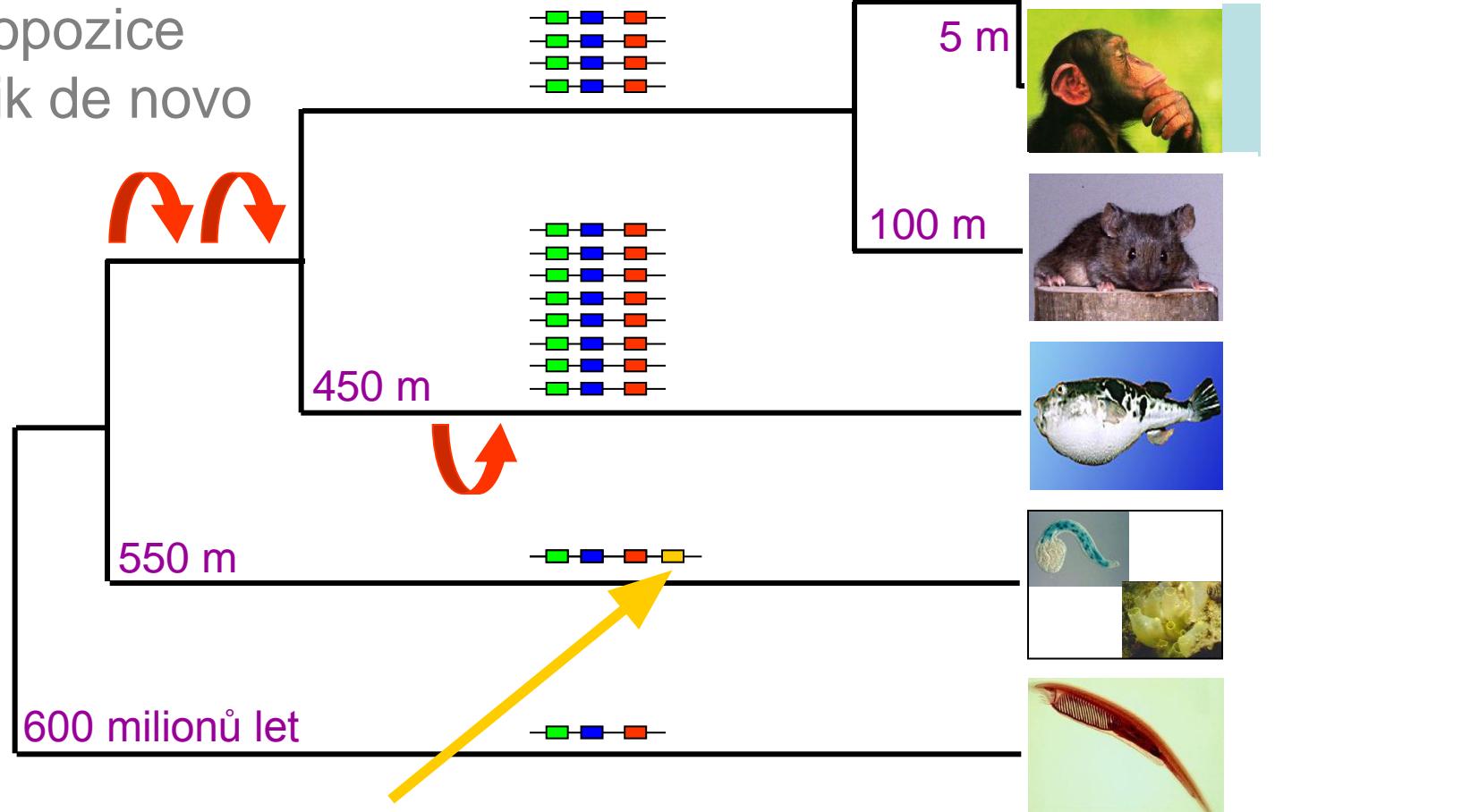
celogenomové duplikace

duplikace segmentů

tandemové duplikace

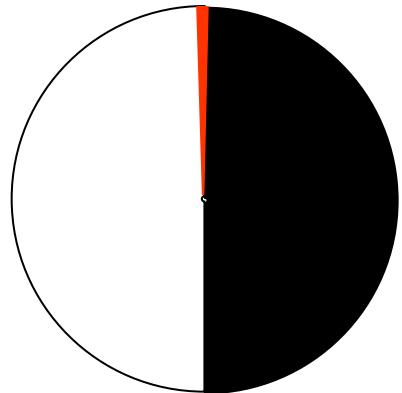
retropozice

vznik de novo

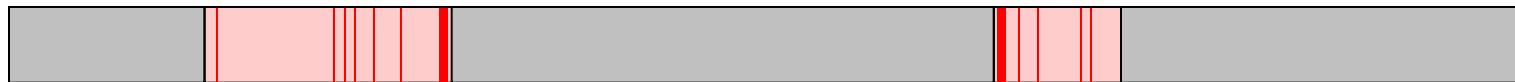


horizontální transfer genů  
(také transfer z genomu organel)

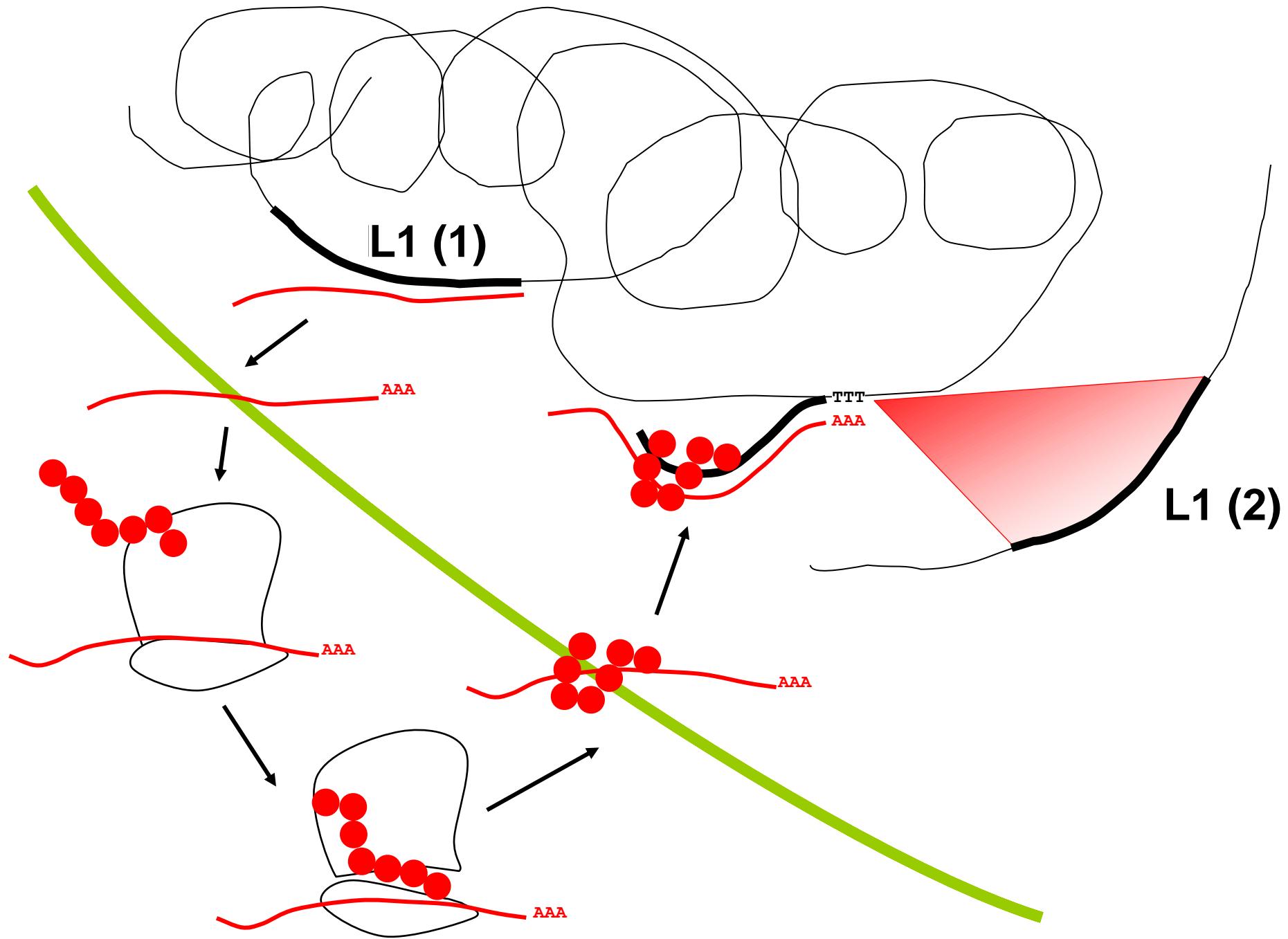
## repetitivní DNA



exony cca. 1.5%  
repetitivní DNA cca. 50%  
nerepetitivní DNA



## mobilní (soběcká) DNA



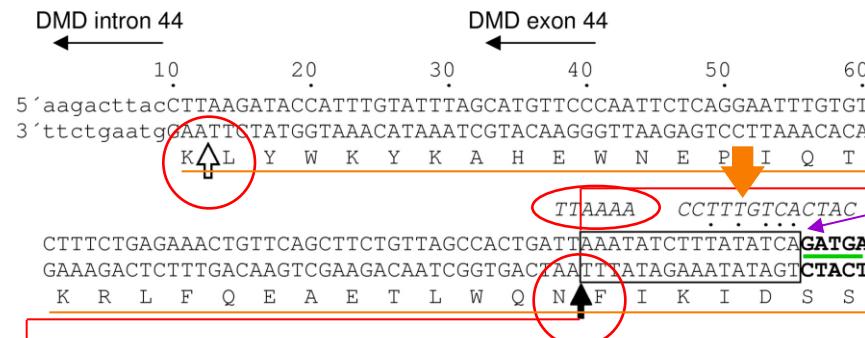
# DMD - Duchennova muskulární dystrofie

Xp21  
gen DMD

2.5 Mb

~89 exonů

inzerční mutace



DMD intron 44                            DMD exon 44

5' aagacttac CTTAAGATAACCATTGTATTAGCATGTTCCAATTCTCAGGAATTGTGT  
3' ttctgaatg **AATT**STATGGTAAACATAAATCGTACAAGGGTTAACAGTCCTAAACACA  
K A L Y W K Y K A H E W N E P I Q T

TTTCTGAGAAACTGTTCACTGATTTAGGTAAACATCGTACAAGGGTTAACAGTCCTAAACACA  
GAAAGACTCTTGACAAGTCGAAGACAATCGGTGACAAATTATAGAAATATAGI **CTACT**  
K R L F Q E A E T L W Q N F I K I D S S

**GTAGGGTTGC**AAAAAATTTCTCCATGTTGAGGTTGCCGTTCACTCTGATGGTAGTTTC  
**CATCCAACG**TTTAAAGAGGGTACAACATCCAACCGACAAGTGAGACTACCATCAAAG  
Y T A F I K E W T T P Q R N V R I T T E

**TTTGCTGTGC**AGAACGCTCTTAGTTAAATTAGATCCATTGTCAATTGTCTTTGT  
AAAACGACACGTCCTCGAGAAATCAAATTAACTAGGGTAAACAGTTAAACAGAAAACA  
K A T C F S K L K I L D W K D I K D K T

**TGCCATTGCT**TTGGTGTGGACATGAAGTCCTGCCACGCCATGTCTGAATGGT  
ACGGTAACGAAAACCACAAACCTGTACTTCAGGAACGGGTGCGGATAACAGGACTTACCA  
A M A K P T K S M F D K G V G I D Q I T

duplicace cílové sekvence

AATGCCTAGGTTTCTCTAGGGTTTATGGTTAGGTTAACGTTAAATCTTAAT  
TTACGGATCCAAAAGAAGATCCAAAATACAAAATCCAAATTGCAAATTAGAAATTA  
I G L N E E L T K I T K P K V N L D K I

CCATCTGAATTGATTTGTATAAGGTGAAGGAAGGGATCCAGTTCAGCTTCTACA  
GGTAGAACTTAACTAAAACATATTCCACATTCTCCCTAGGTCAAAGTCGAAGATGT  
W R S N I K T Y P T L F P D L K L K R C

TATGGCTAGCCAGTTTCCCAGCACCATTTATTAAATAGGGATCCTTCCCCATTGCTT  
ATACCGATGGTCAAAGGGTGTGTTAAATAATTATCCCTTAGGAAAGGGGTAAACGAA  
I A L W N E W C W K N F L S D G W Q K

GCGGCATTAT

GTTTTTCTCAGGTTGTCAAAGATCAGATAAGTGTAGATATGCCGCAAAAAAAAAA  
CAAAAGAGTCCAAACAGTTCTAGTCTATCAACATCTATACGCCGTTTTTTTTTT  
N K E P K D F I L Y N Y I H P F F F F F

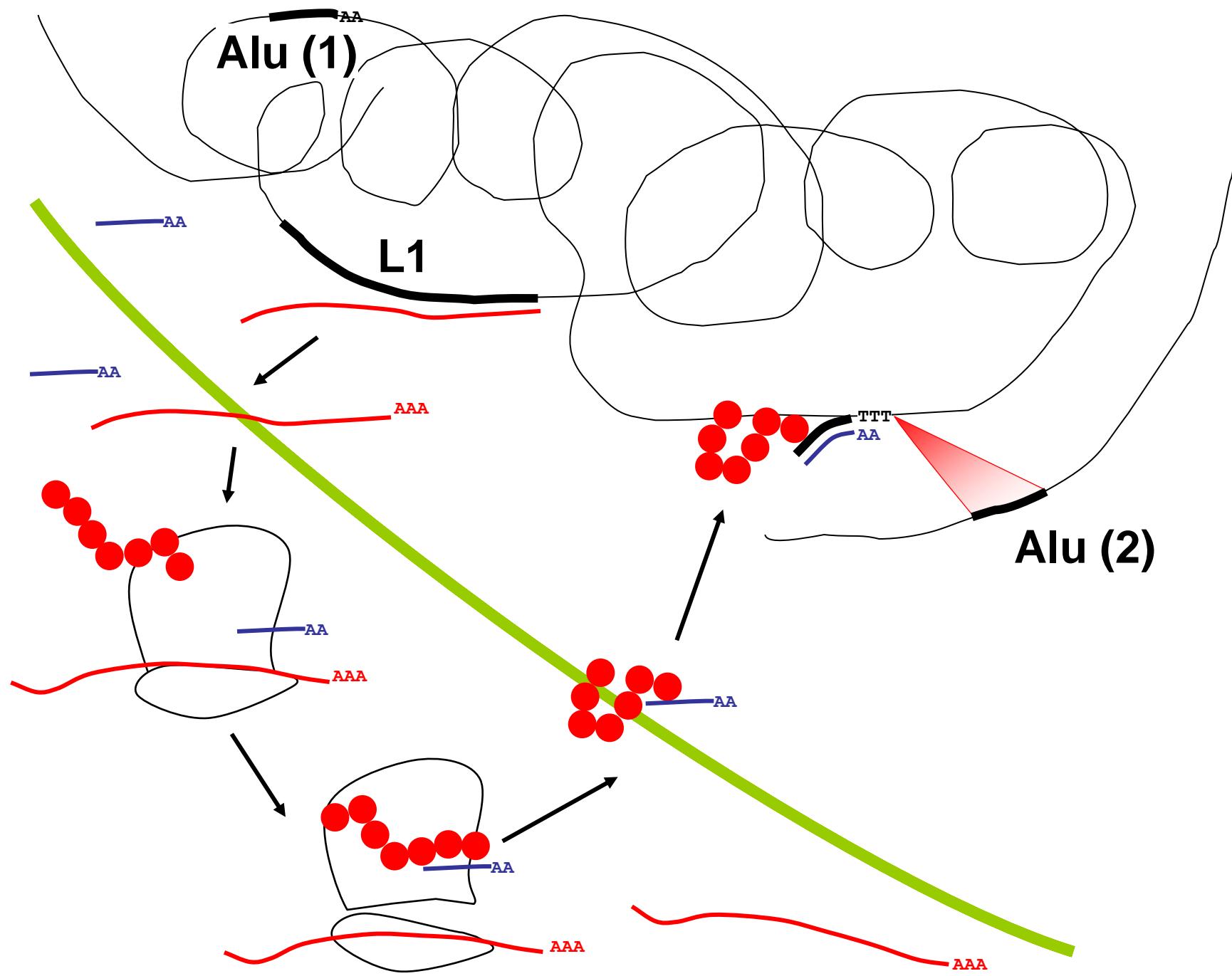
AAAAAAAAAAAAAAAAAAAAAA  
TT  
F F F F F F F F F F F F

TTTCTCAACAGATCTGTCAAATCGCtcgcaggtaaaagcatatggatcaagaaaaataga 3'  
AAAGAGTTGTCTAGACAGTTAGCGGacgtccatttcgtatacctagttcttttatct 5'  
K E V S R D F R

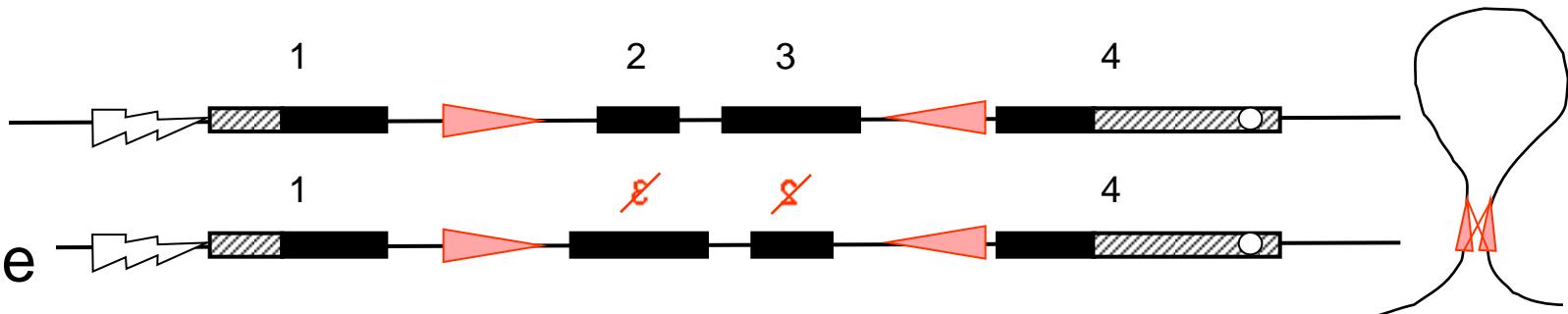
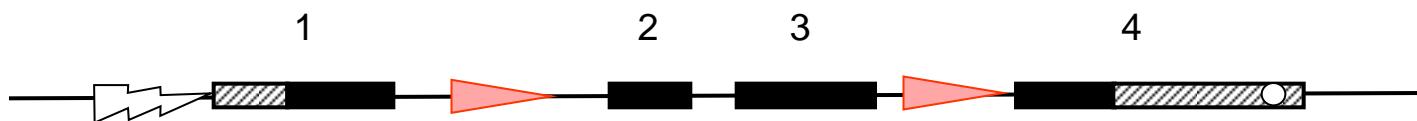
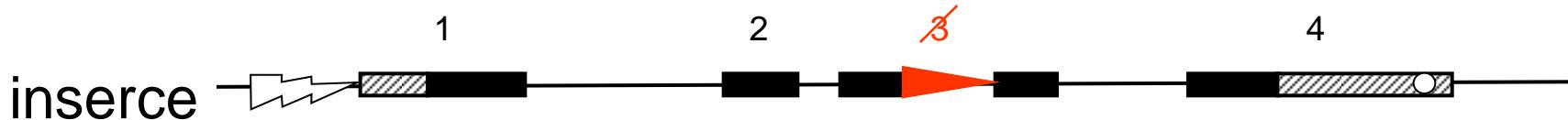
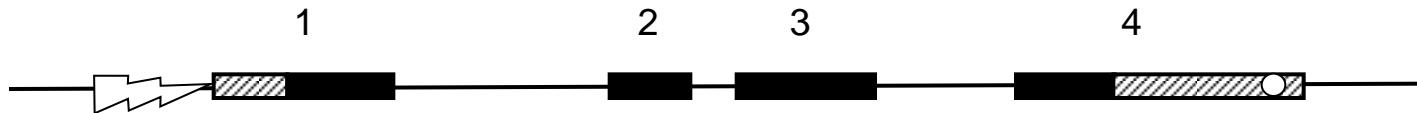
DMD exon 44

DMD intron 43

L1



# vliv transpozonů na genom



fluidita/plasticita genomu