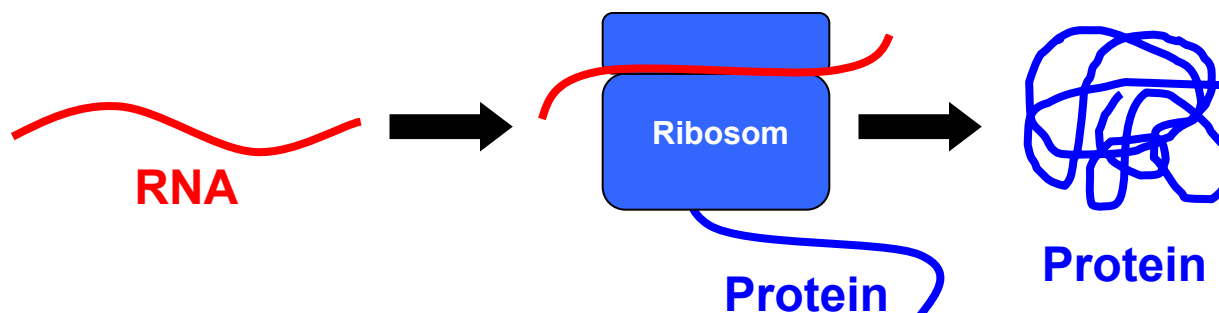


# Základní vlastnosti proteinů

## Teoretický úvod

Aplikovaná bioinformatika, Jaro 2013

# Proteiny

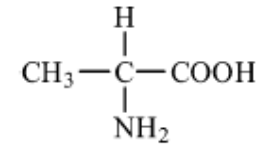


- Protein, polypeptid, bílkovina.
- Lineární polymer aminokyselin spojených peptidovými vazbami.
- Funkce: katalytická, regulační, transportní, zprostředkování pohybu, obranná, strukturální, zásobní.

# Proteinogenní aminokyseliny

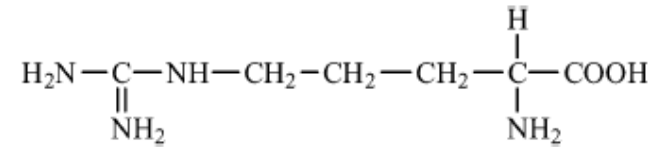
Alanine Ala

A



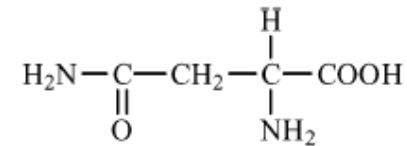
Arginine Arg

R



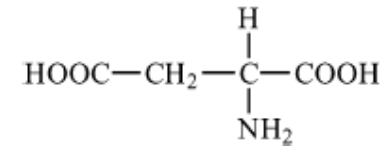
Asparagine Asn

N



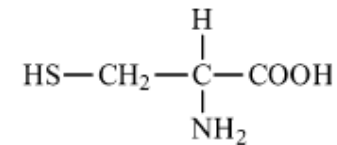
Aspartic acid Asp

D



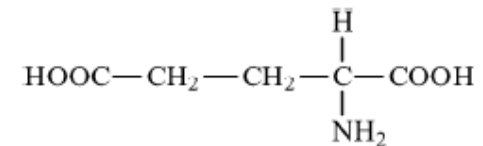
Cysteine Cys

C



Glutamic acid Glu

E

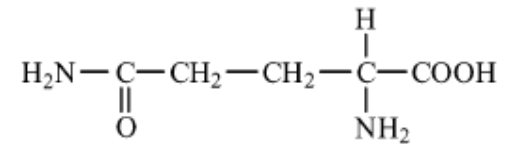


# Proteinogenní aminokyseliny

Glutamine

Gln

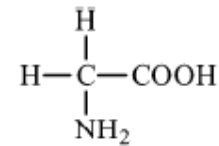
Q



Glycine

Gly

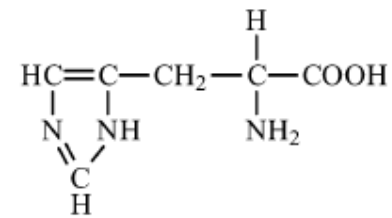
G



Histidine

His

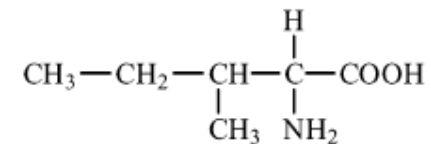
H



Isoleucine

Ile

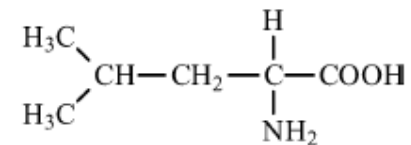
I



Leucine

Leu

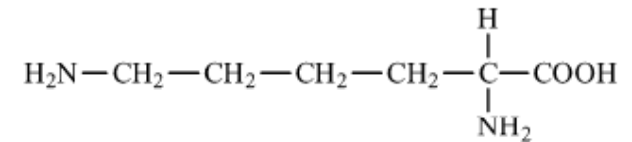
L



# Proteinogenní aminokyseliny

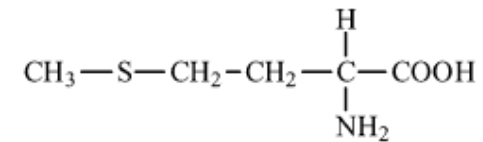
Lysine Lys

K



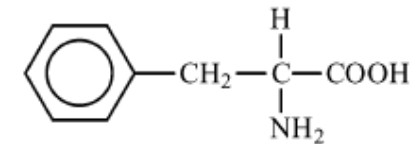
Methionine Met

M



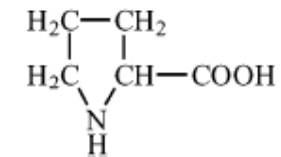
Phenylalanine Phe

F



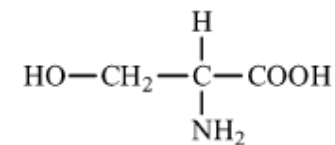
Proline Pro

P



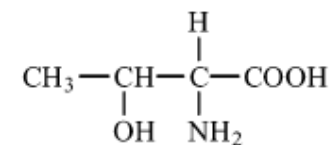
Serine Ser

S



Threonine Thr

T

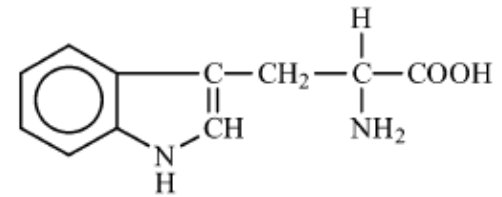


# Proteinogenní aminokyseliny

Tryptophan

Trp

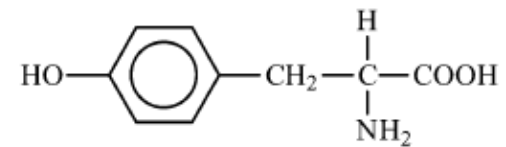
W



Tyrosine

Tyr

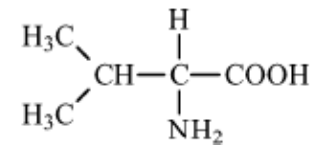
Y



Valine

Val

V



21. aminokyselina – selenocystein, Sec, U

22. aminokyselina – pyrrolysin, Pyl, O

## Part 2, Section B: THE ONE-LETTER SYSTEM (revision and updating of [11])

### 3AA-20. THE NEED FOR A CONCISE REPRESENTATION OF SEQUENCE

#### 3AA-20.1. General Considerations Regarding the One-Letter System

There are difficulties in using the three-letter system ([3AA-14](#) to [3AA-19](#)) in presenting long protein sequences. A one-letter code is much more concise, and is helpful in summarizing large amounts of data, in aligning and comparing homologous sequences, and in computer techniques for these processes. It may also be used to label residues in three-dimensional pictures of protein molecules.

The possibility of using one-letter symbols was mentioned by Gamow & Ycas [\[26\]](#) in 1958. Sorm *et al.* [\[27\]](#) systematized the idea in 1961 (see, for example, [\[28\]](#)), and Dayhoff and Eck used one-letter symbols derived partly from the code of Sorm *et al.* in their compilations of protein sequences ([\[29\]](#), latest edition [\[30\]](#)). IUB-IUPAC recommendations [\[11\]](#) were approved in 1968 on the basis of proposals of a subcommittee of W. E. Cohn, M. O. Dayhoff, R. V. Eck, and B. Keil, and these recommendations are given here with no substantial change.

PLLSASIVSAPVVTSETYVDIPGLYLDVAKAGIRDGKLQVILNVPTPY  
ATGNNFPGIYFAIATNQGVDGCFITYSSKVPESTGRMPFTLVATIDV  
GSGVTFVKGQWKSVRGSAMHIDSYASLSAIWGTAAPSSQSGNQGAET  
GGTGAGNIGGGGERDGTFNLPPIKFGVTALTHAANDQTIDIYIDDDP  
KPAATFKGAGAQDQNLGTKVLD SGNGRVRVIVMANGRPSRLGSRQVDI  
FKKSYFGIIGSEDGADDDYNDGIVFLNWPLG

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```
BclB -----ERDGI FTLPNIAFGV TALVNSSAPQTI EVFVDDNPKPAATFQ GAGTQ DANLNTQIVNSGK-G 62
BclC -----ERDGT FNLPPIKFGV TALHTAANDQTI DIYIDDDPKPAATFKGAGA QDQNLG TKVLD SGN-G 62
PA-IIL -----ATQGV FTLPANTRFGV TAFANSSGTQTVNVLVNN--ETAATFSGQSTNNAVIGTQVLNSGSSG 61
BclA ADSQTSSNRAGEFSI PPNTDFRAIFFANAAEQQH IKLFI GDSQEPAAAYHKL TTRDGPREAT--LNSGN-G 67
BclD -----DRNGN FSLPPNTAFKAI FYANAADRQDLKLF IDDAPEPAATFVGNSE DGVR LFT--LNSKG-G 60
Clustal Consensus      * *.:*.: * . .::: * :.: :.: :. ** . .: * ::* *
```

```
BclB KVRVVVTANGKPSKIGSRQVDIFK-----KTYFGLVGS EDGGDGDYNDGIA IILNWPLG 115
BclC RVRVIVMANGRPSRLGSRQVDIFK-----KSYFGIIGSE DGADDDYNDGIVFLNWPLG 115
PA-IIL KVQVQVSVNGRPSDLVSAQVILT N-----ELNFALVGS EDGTDNDYNDAVVVINWPLG 114
BclA KIRFEVSVNGKPSATDARLAPINGKKS DGS PFTVNF GIVVSE DGHDSYNDGIVVLQWPIG 128
BclD KIRIEASANGRQSATDARLAPL----SAGD--TVWL GWLGA EDGADADYNDGIVILQWPIT 115
Clustal Consensus  :::. . .**:* * : . : :. : :*** * ****. :. :. :** : 49
```



One-letter symbol	Three-letter symbol	Amino acid
A	Ala	alanine
B	Asx	aspartic acid or asparagine
C	Cys	cysteine
D	Asp	aspartic acid
E	Glu	glutamic acid
F	Phe	phenylalanine
G	Gly	glycine
H	His	histidine
I	Ile	isoleucine
K	Lys	lysine
L	Leu	leucine
M	Met	methionine
N	Asn	asparagine
P	Pro	proline
Q	Gln	glutamine
R	Arg	arginine
S	Ser	serine
T	Thr	threonine
V	Val	valine
W	Trp	tryptophan
X**	Xaa	unknown or 'other' amino acid
Y	Tyr	tyrosine
Z	Glx	glutamic acid or glutamine (or substances such as 4-carboxyglutamic acid and 5-oxoproline that yield glutamic acid on acid hydrolysis of peptides)

### *Note on the Choice of Symbols*

Initial letters of the names of the amino acids were chosen where there was no ambiguity. There are six such cases: cysteine, histidine, isoleucine, methionine, serine and valine. All the other amino acids share the initial letters A, G, L, P or T, so arbitrary assignments were made. These letters were assigned to the most frequently occurring and structurally most simple of the amino acids with these initials, alanine (A), glycine (G), leucine (L), proline (P) and threonine (T).

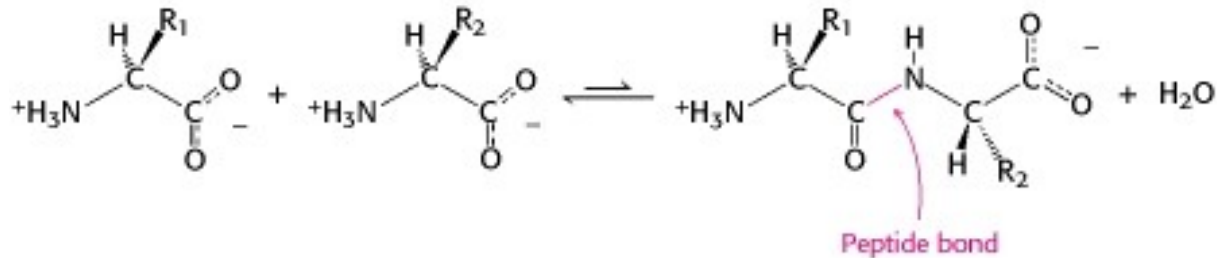
Other assignments were made on the basis of associations that might be helpful in remembering the code, e.g. the phonetic associations of F for phenylalanine and R for arginine. For tryptophan the double ring of the molecule is associated with the bulky letter W. The letters N and Q were assigned to asparagine and glutamine respectively; D and E to aspartic and glutamic acids respectively. K and Y were chosen for the two remaining amino acids, lysine and tyrosine, because, of the few remaining letters, they were close alphabetically to the initial letters of the names. U and O were avoided because U is easily confused with V in handwritten material, and O with G, Q, C and D in imperfect computer print-outs, and also with zero. J was avoided because it is absent from several languages.

Two other symbols are often necessary in partly determined sequences, so B was assigned to aspartic acid or asparagine when these have not been distinguished; Z was similarly assigned to glutamic acid or glutamine. X means that the identity of an amino acid is undetermined, or that the amino acid is atypical. See the [Addendum](#) for an alternative use of X.

21. aminokyselina – selenocystein, Sec, U

22. aminokyselina – pyrrolysin, Pyl, O

# Peptidová vazba



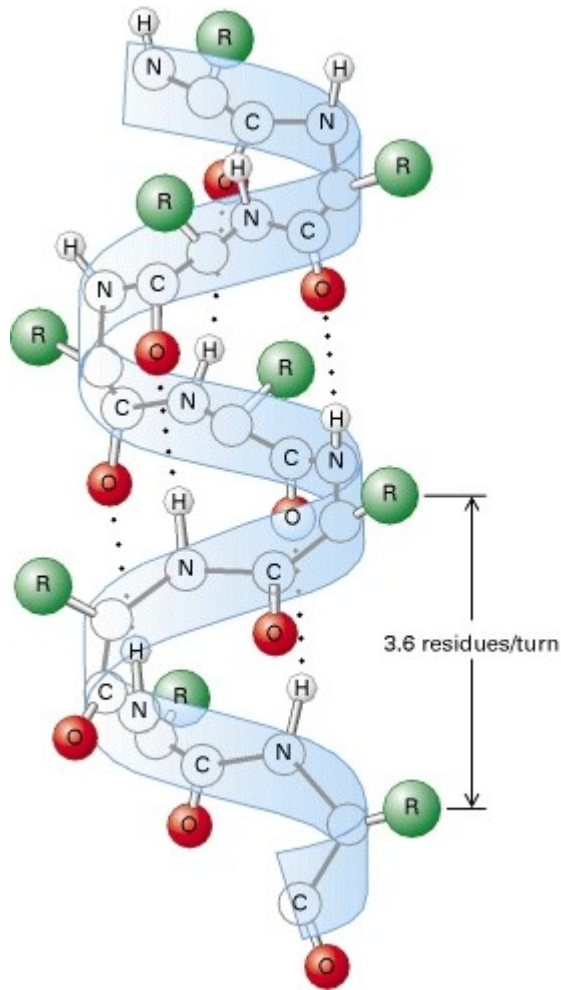
**N** MTWCKTMIDQGRSWPHCYYGMAA  
DTYYKKLTPGHTQVGITILMGAC  
GCCCGTGCRNMSDETGCWWCGTA  
HSPGCTDEQLRCGLVCGT **C**

**N** → **C**

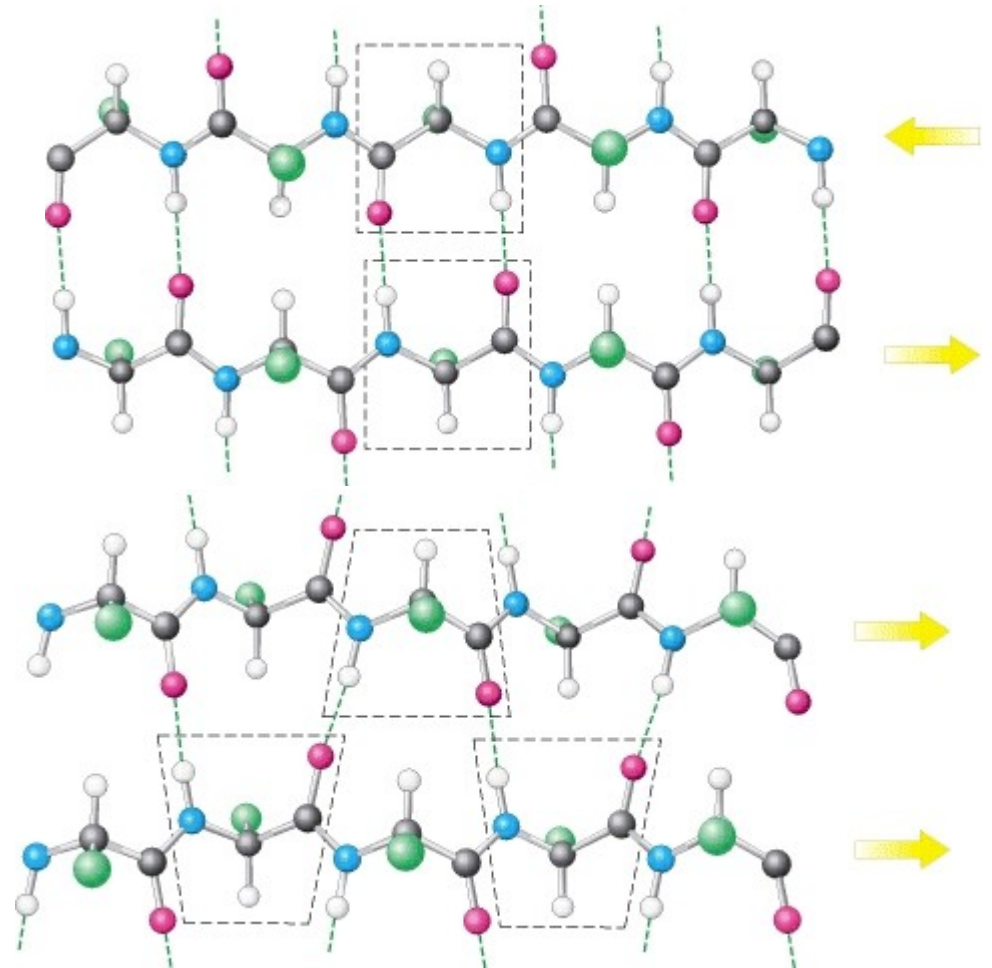
# Proteinogenní aminokyseliny

- Stavební jednotky proteinů:  $\alpha$ -L-aminokyseliny.
- 20 standardních proteinogenních aminokyselin.
- Alifatické (Gly, Ala, Val, Leu, Ile).
- **Sírné** (Cys, Met).
- **S OH** skupinou (Ser, Thr).
- **Kyselá a z nich odvozená** (Glu, Gln, Asp, Asn).
- **Bazické** (Lys, Arg).

# Sekundární struktura

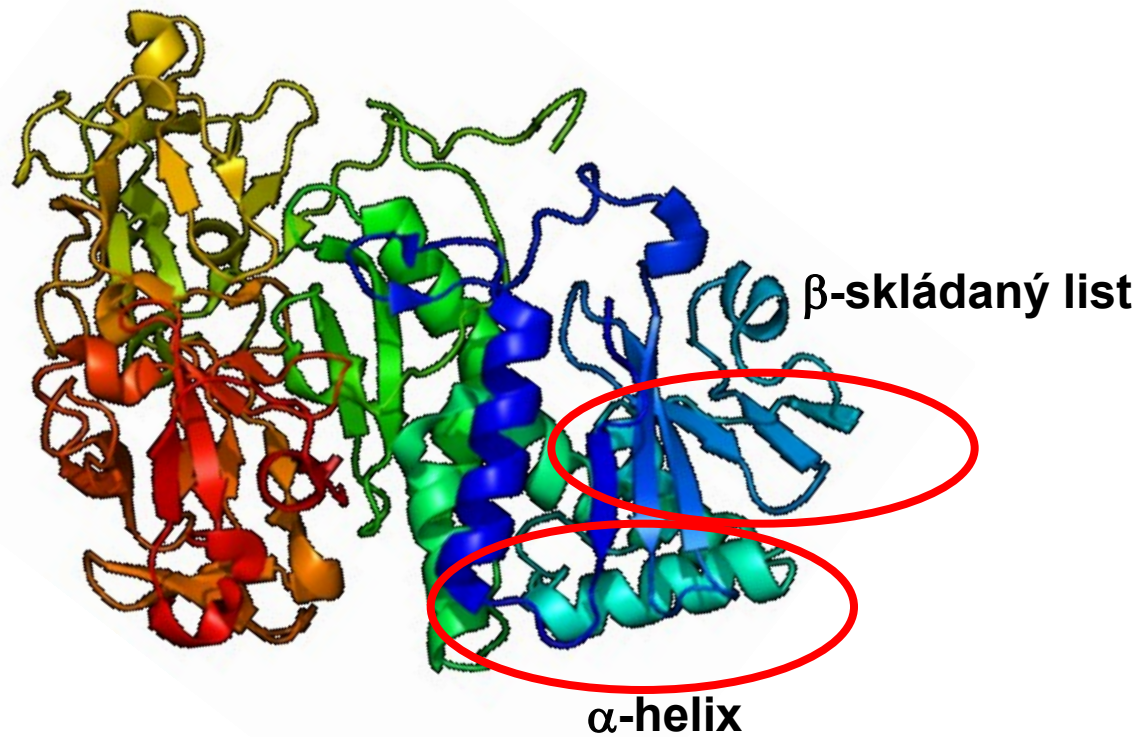


$\alpha$ -helix



$\beta$ -skládání list

# Terčiární struktura



# **Jak se chovají proteiny? (v laboratoři)**

- **Limitované množství proteinu (cena, dostupnost).**
- ***In vitro* mohou rychle ztrácet aktivitu (nutná správná sekundární, terciární a někdy i kvarterní struktura).**
- **Mohou být nestabilní (některé velmi nestabilní) mimo své optimální prostředí v buňce (organismu).**
- **K výrazné destabilizaci a denaturaci může docházet již za laboratorní teploty (25 °C).**

**...zlobí.**

# Práce s proteiny

- **Většina savčích proteinů začíná denaturovat již při teplotách nad 40 °C. Při teplotě 95 °C dochází k úplné denaturaci téměř všech proteinů během několika minut. K výrazné destabilizaci a denaturaci může docházet již za laboratorní teploty (25 °C).**

**S proteiny pracujeme „na ledu“.**





# Práce s proteiny

- Proteiny jsou štěpeny proteasami a peptidasami. Optimum těchto enzymů je 37 °C, za nižší teploty se jejich aktivita snižuje (ale jsou aktivní i při 4 °C). Proteasy se do vzorku dostanou neopatrnou manipulací a jsou také produkovány mikroorganismy.

**Minimalizace kontaminace vzorku (ochranné pomůcky) a použití inhibitorů proteas.**



# Práce s proteiny

- Všechny vzorky jsou dříve či později kontaminovány bakteriemi. Mikroorganismy si na proteinech pochutnají i při 4 °C a produkují proteasy.

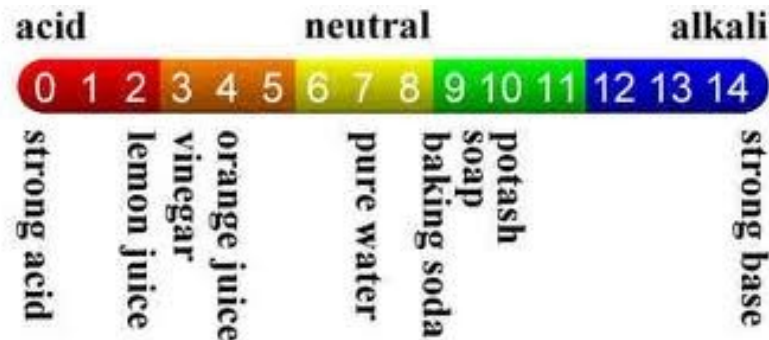
**Přídavek antibakteriálních látek (0,02-0,05 % azid sodný)**



# Práce s proteiny

- Proteiny jsou aktivní (a stabilní) v určitém rozmezí pH. A to může být pro některé proteiny velmi úzké... Fyziologické pH pro většinu proteinů je cca 7,2-7,4. Silně kyselé nebo zásadité prostředí proteiny denaturuje.

**Nutné kontrolované prostředí – pufrý o vhodném pH.**



# Práce s proteiny

- Proteiny jsou aktivní (a stabilní) v určitém rozmezí pH. A to může být pro některé proteiny velmi úzké... Fyziologické pH pro většinu proteinů je cca 7,2-7,4. Silně kyselé nebo zásadité prostředí proteiny denaturuje.

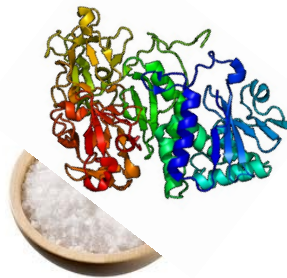
## Nutné kontrolované prostředí – pufrы o vhodném pH.

- Pufr, tlumivý roztok, ústojný roztok, ústoj: látka (směs látek) schopná udržovat stabilní pH po přidavku silné kyseliny nebo zásady do systému.  
Příklad: slabá kyselina/její sůl, HA/A<sup>-</sup>.  
„Přirodní“ x syntetické pufrы. Pufrы nesmí interagovat s proteiny nebo interferovat s jejich funkcí!

# Práce s proteiny

- Proteiny vyžadují pro svou aktivitu (a stabilitu) určitou koncentraci solí (iontů). Vysoká i nízká koncentrace solí může způsobovat agregaci a precipitaci. Proteiny většinou nejsou stabilní v čisté vodě.

**Nutná optimalizovaná koncentrace soli v roztoku.**



# Práce s proteiny

- Při práci s nízkými koncentracemi proteinů ( $< 1 \text{ mg/ml}$ ) se může výrazně projevit ztráta způsobená vazbou na stěny použité nádoby (zkumavky).

Pokud je to možné, lze použít inertní proteiny (BSA, cca  $2 \text{ mg/ml}$ ), které vazbě zabrání.



# Práce s proteiny

## Protein LoBind Tubes

Při použití těchto zkumavek dochází k minimálním ztrátám proteinů. V těchto zkumavkách nedochází k redukci aktivity enzymů ani k denaturačním efektům způsobeným vazbou na povrch běžných zkumavek. Příště již žádné trápení se saturací povrchu BSA nebo silikonizací běžných plastových zkumavek pro práci s malými množstvími proteinů. Zkumavky jsou vyrobeny speciální technologií z nejčistšího polypropylenu.

### Specifikace:

- ▶ minimální ztráty proteinů (méně než 3%, BSA 1 µg/ml)
- ▶ vnitřní povrch zkumavek nemá žádný potah (např. silikonem)
- ▶ bez DNA, DNAs, RNAs a PCR inhybitorů (PCR clean)
- ▶ vyráběné jako Safe-Lock zkumavky ve velikostech 0,5 ml, 1,5 ml a 2 ml
- ▶ vhodné i pro centrifugaci při vysokých otáčkách

# Práce s proteiny

- Proteiny mohou být rovněž poškozeny mechanicky při příliš energickém míchaní nebo třepání!

Nutná opatrná a jemná manipulace s proteiny.



Vortex



Míchačka



# Skladování proteinů

Characteristic	Storage Condition			
	Solution at 4°C	Solution in 25-50% glycerol or ethylene glycol at -20°C	Frozen at -20° to -80°C or in liquid nitrogen	Lyophilized (usually also frozen)
Typical shelf life	1 month	1 year	Years	Years
Requires sterile conditions or addition of antibacterial agent	Yes	Usually	No	No
Number of times a sample may be removed for use	Many	Many	Once; repeated freeze-thaw cycles generally degrade proteins	Once; it is impractical to lyophilize a sample multiple times

# Skladování proteinů

- Lyofilizace – mrazová sublimace.
- Odpařování vody ze zmraženého vzorku za sníženého tlaku.
- Nedostatek vody zabraňuje růstu mikroorganismů a inhibuje enzymy (proteasy).
- Nepoškozuje vzorek v takovém rozsahu jako jiné způsoby dehydratace (vysoká teplota, vysoušedla).
- Jednoduchá rehydratace.

---

---

Lyophilized  
(usually also frozen)

---

Years

---

No

---

Once; it is  
impractical to  
lyophilize a sample  
multiple times

---


# Skladování proteinů

Characteristic	Storage Condition			
	Solution at 4°C	Solution in 25-50% glycerol or ethylene glycol at -20°C	Frozen at -20° to -80°C or in liquid nitrogen	Lyophilized (usually also frozen)
Typical shelf life	1 month	1 year	Years	Years
Requires sterile conditions or addition of antibacterial agent	Yes	Usually	No	No
Number of times a sample may be removed for use	Many	Many	Once; repeated freeze-thaw cycles generally degrade proteins	Once; it is impractical to lyophilize a sample multiple times

**Sterilní zkumavky, sterilizace filtrací.  
Inhibitory proteas.**

# Skladování proteinů

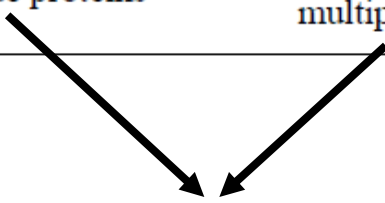
Kryoprotektanty zabraňují tvorbě krystalků ledu a poškození proteinu.



	Storage Condition			
Characteristic	Solution at 4°C	Solution in 25-50% glycerol or ethylene glycol at -20°C	Frozen at -20° to -80°C or in liquid nitrogen	Lyophilized (usually also frozen)
Typical shelf life	1 month	1 year	Years	Years
Requires sterile conditions or addition of antibacterial agent	Yes	Usually	No	No
Number of times a sample may be removed for use	Many	Many	Once; repeated freeze-thaw cycles generally degrade proteins	Once; it is impractical to lyophilize a sample multiple times

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	Storage Condition			
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Number of times a sample may be removed for use	Many	Many	Once; repeated freeze-thaw cycles generally degrade proteins	Once; it is impractical to lyophilize a sample multiple times



**Nutné připravit několik alikvotů (částí zásobního roztoku) proteinu.**

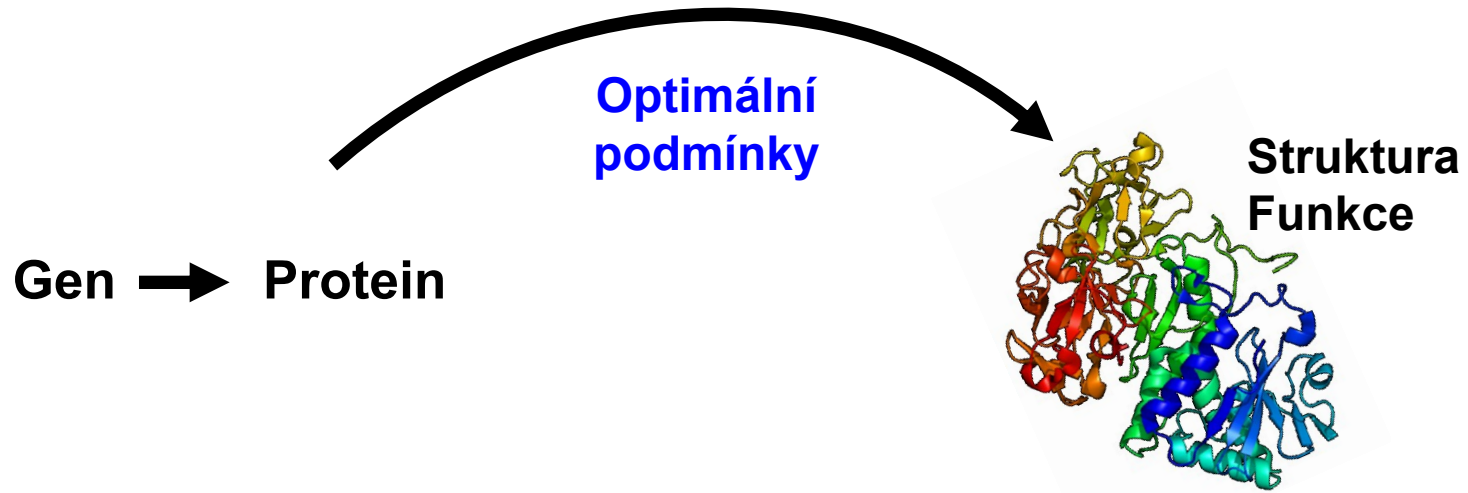
# Skladování proteinů

	Storage Condition			
Characteristic	Solution at 4°C	Solution in 25-50% glycerol or ethylene glycol at -20°C	Frozen at -20° to -80°C or in liquid nitrogen	Lyophilized (usually also frozen)
Typical shelf life	1 month	1 year	Years	Years
Requires sterile conditions or addition of antibacterial agent	Yes	Usually	No	No
Number of times a sample may be removed for use	Many	Many	Once; repeated freeze-thaw cycles generally degrade proteins	Once; it is impractical to lyophilize a sample multiple times

**Proteiny mohou být lyofilizací nebo zamražením nevratně poškozeny!**

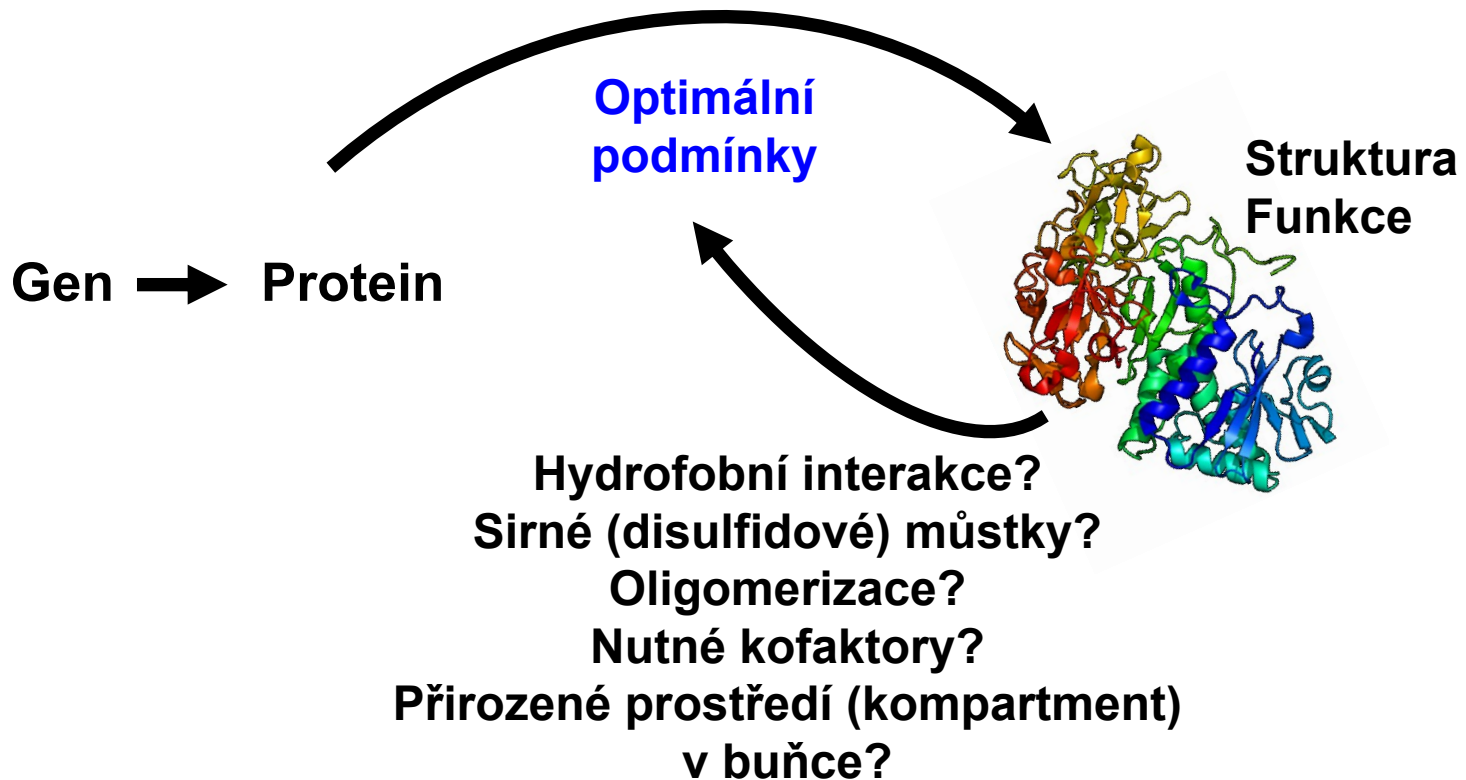
# Práce s proteiny

Protein je správně sbalený, aktivní, v dostatečném množství a koncentraci



# Práce s proteiny

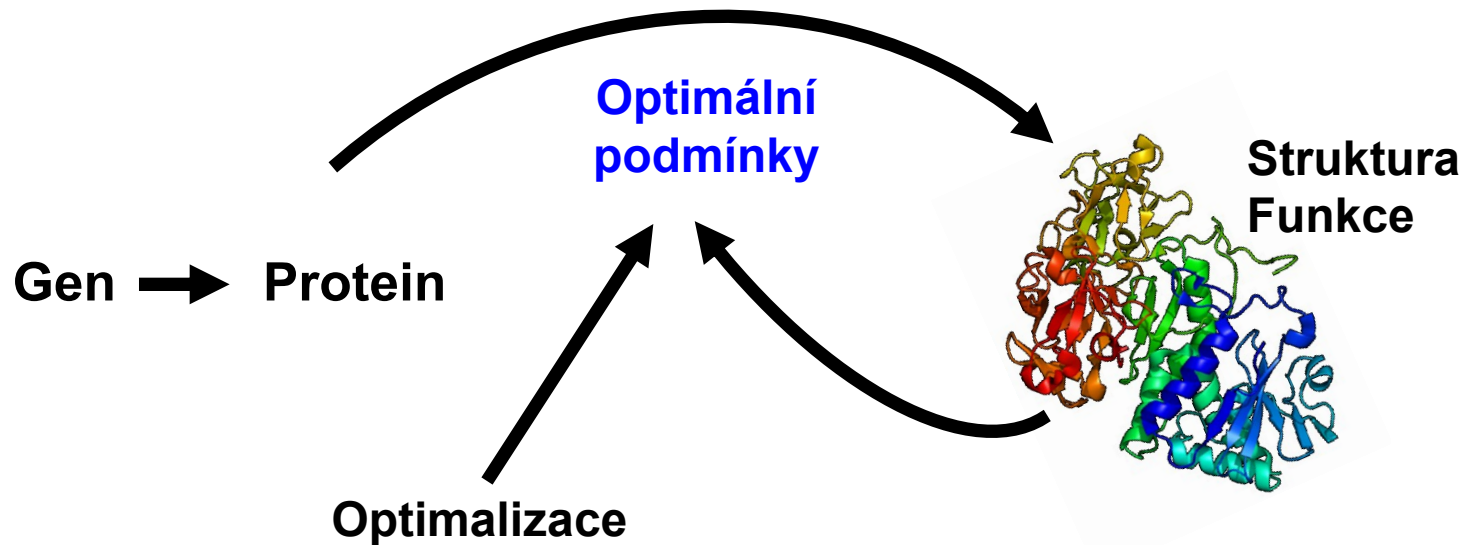
Protein je správně sbalený, aktivní, v dostatečném množství a koncentraci





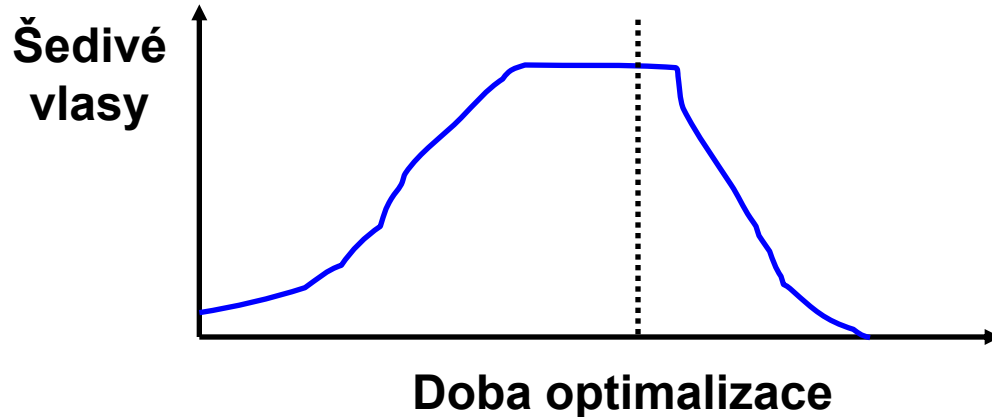
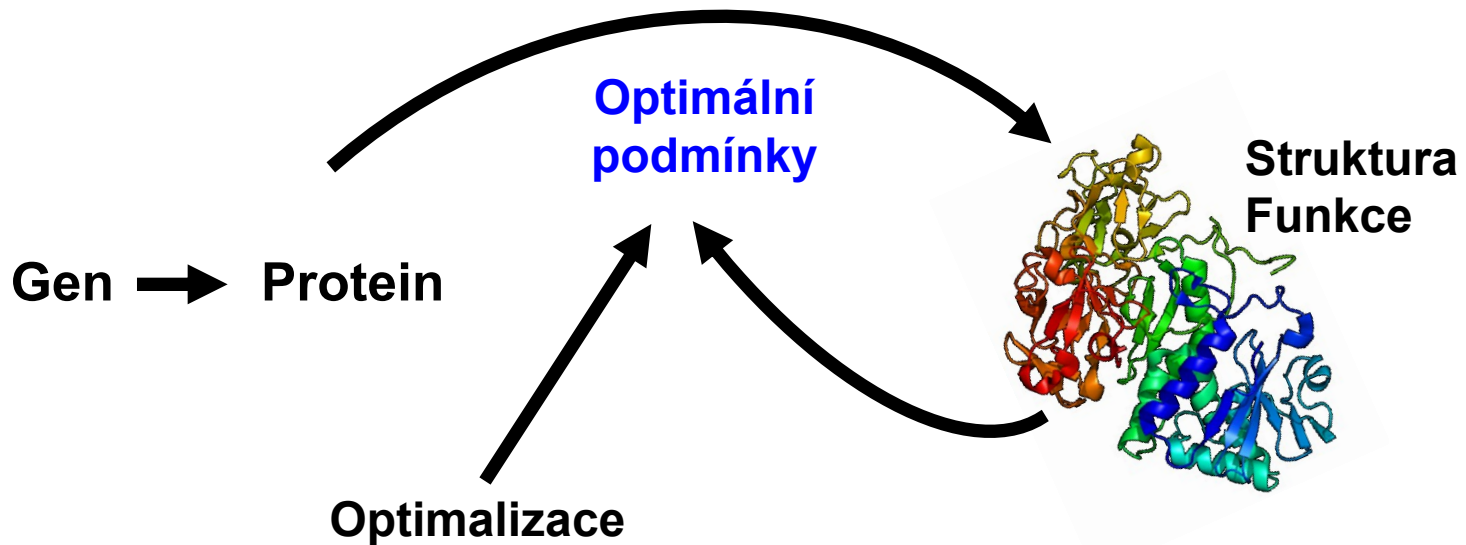
# Práce s proteiny

Protein je správně sbalený, aktivní, v dostatečném množství a koncentraci



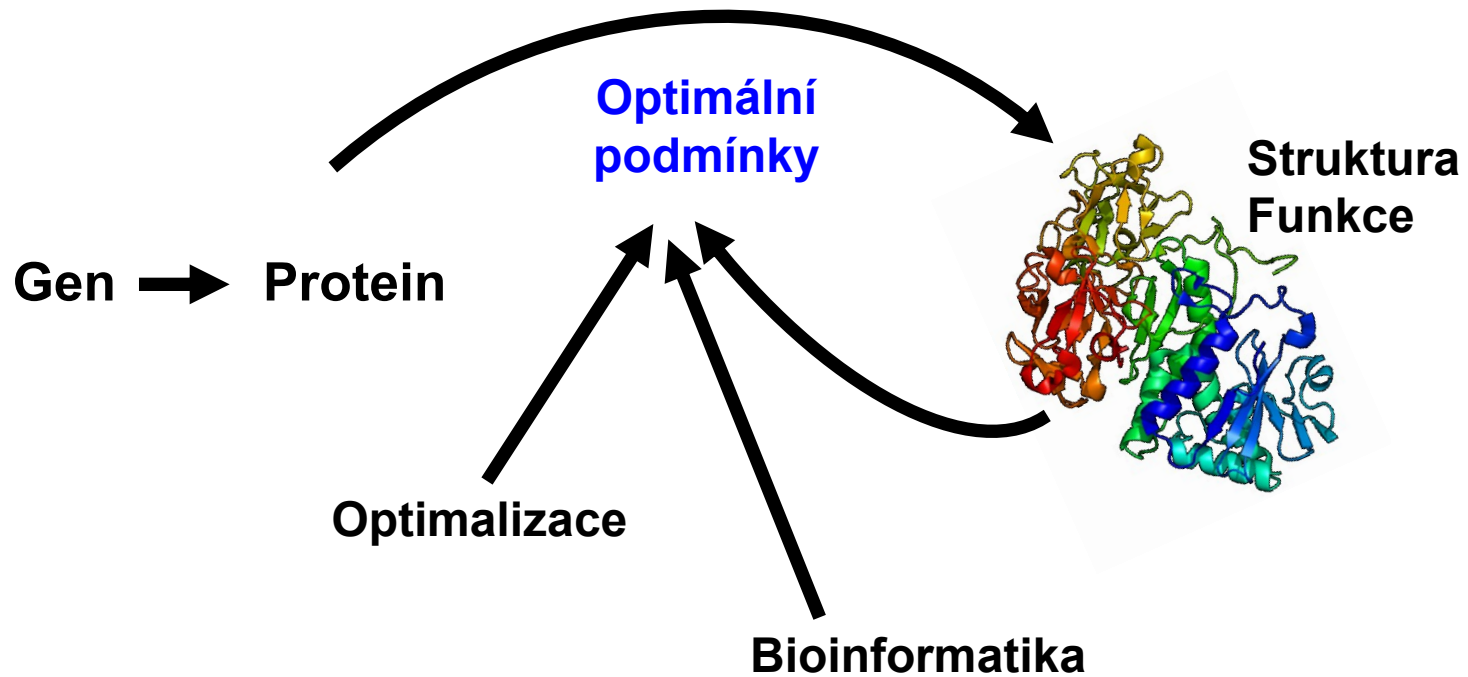
# Práce s proteiny

Protein je správně sbalený, aktivní, v dostatečném množství a koncentraci



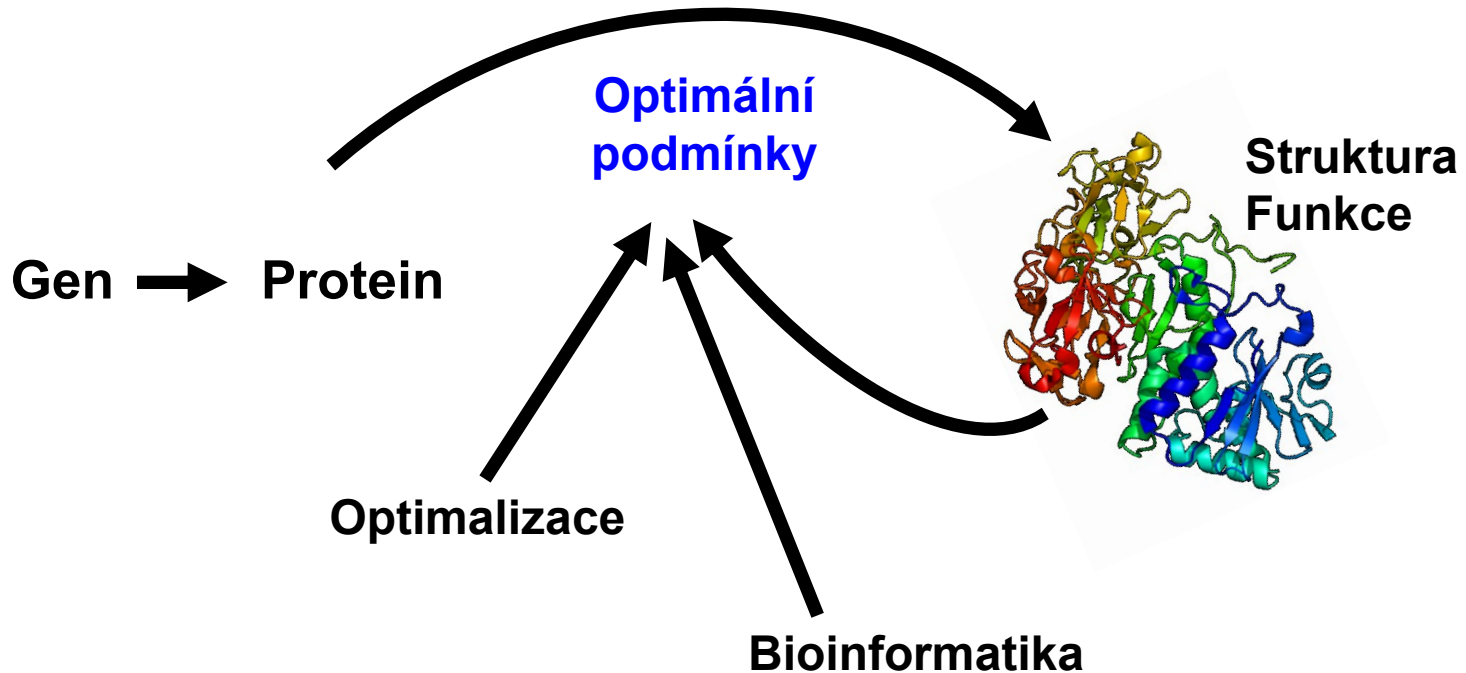
# Práce s proteiny

Protein je správně sbalený, aktivní, v dostatečném množství a koncentraci



# Práce s proteiny

Protein je správně sbalený, aktivní, v dostatečném množství a koncentraci



```
PLLSASIVSAPVVTSETYVD  
IPGLYLDVAKAGIRDGKLQV  
ILNVPTY
```

**Predikce  
vlastností**

# Predikce vlastností proteinů

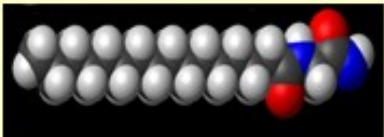


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**Molecular Biology of the Cell, 4<sup>th</sup> edition**

**Bruce Alberts, Alexander Johnson, Julian Lewis, Martin Raff, Keith Roberts, and Peter Walter**

**New York: Garland Science; 2002.**

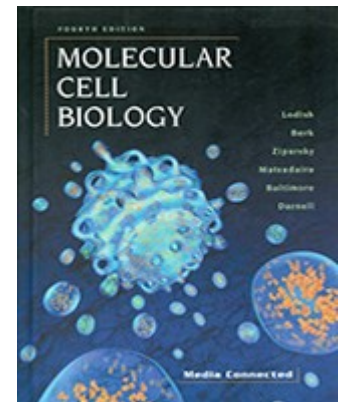
**ISBN-10: 0-8153-3218-1 ISBN-10: 0-8153-4072-9**

**Molecular Cell Biology, 4<sup>th</sup> edition**

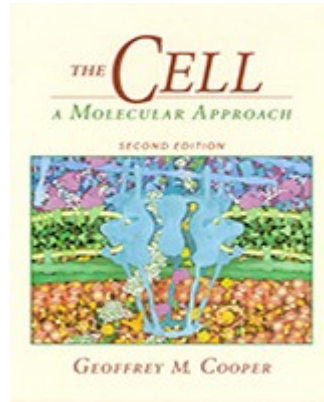
**Harvey Lodish, Arnold Berk, S Lawrence Zipursky, Paul Matsudaira, David Baltimore, and James Darnell**

**New York: W. H. Freeman; 2000.**

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**The Cell, 2<sup>nd</sup> edition, A Molecular Approach**

**Geoffrey M Cooper**

**Boston University**

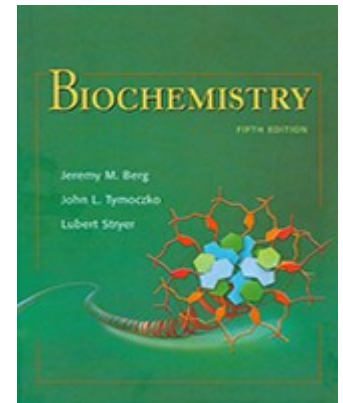
**Sunderland (MA): Sinauer Associates; 2000.**

**ISBN-10: 0-87893-106-6**

**Biochemistry, 5<sup>th</sup> edition**  
**Jeremy M Berg, John L Tymoczko, and**  
**Lubert Stryer**

**New York: W H Freeman; 2002.**

**ISBN-10: 0-7167-3051-0**



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McCrorry DC, Coeytaux RR, Yancy WS Jr, et al.  
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Report | Comparative Effectiveness Research, Health Care
-  [Outpatient Case Management for Adults With Medical Illness and Complex Care Needs \[Internet\]](#)  
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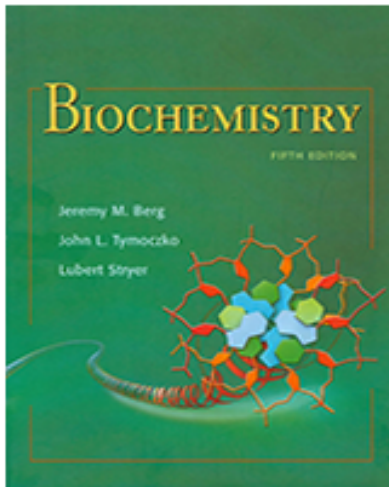
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**Brian W. Matthews.** Hydrophobic Interactions in Proteins, in Encyclopedia of Life Sciences (ELS), John Wiley & Sons, Ltd: Chichester, 2001.

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Tech Tip #43 Protein stability and storage: [www.piercenet.com](http://www.piercenet.com)