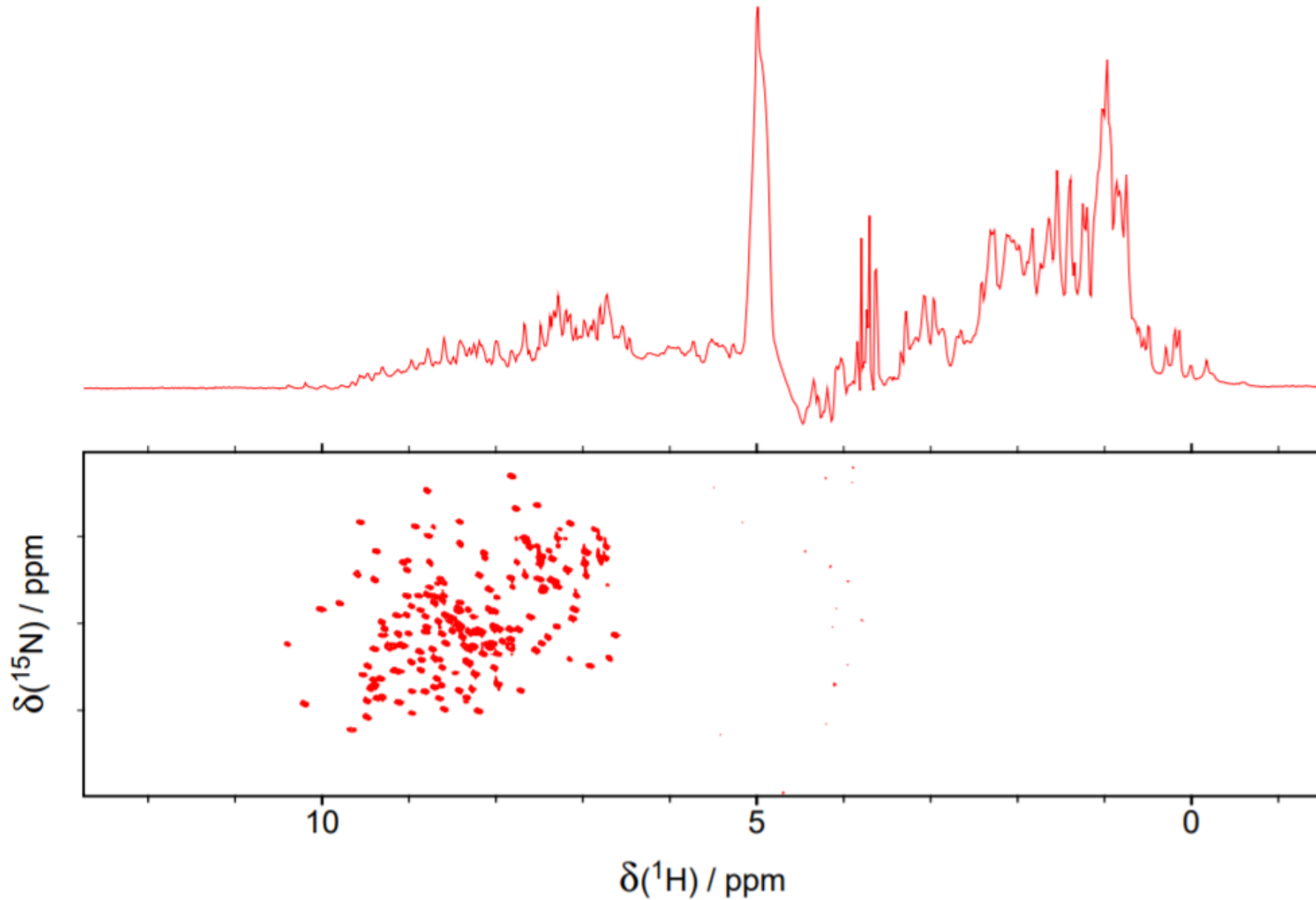


Nukleární magnetická rezonance

Viktor Bartošík

10. 12. 2023

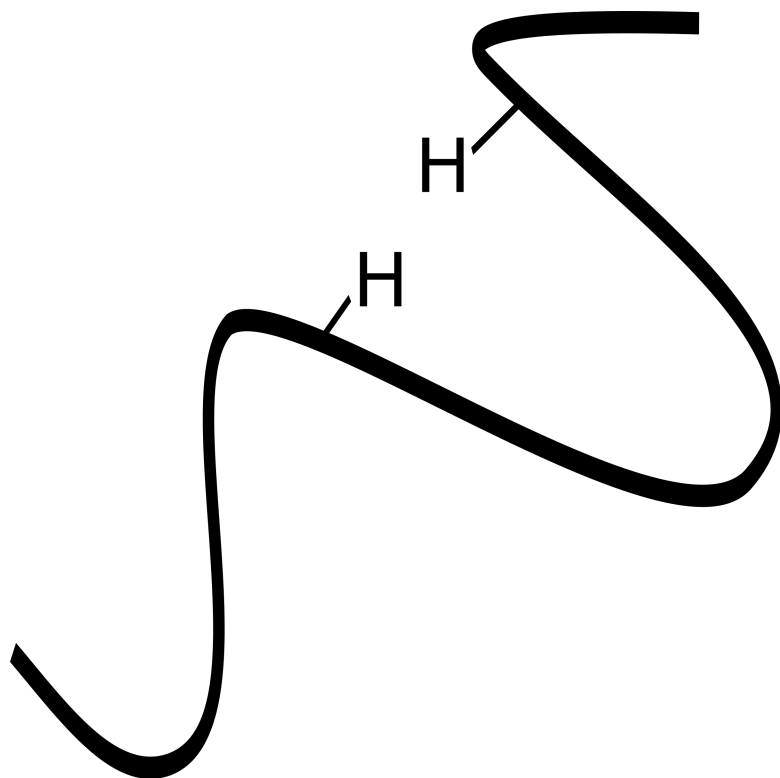
NMR proteinů



Experimenty

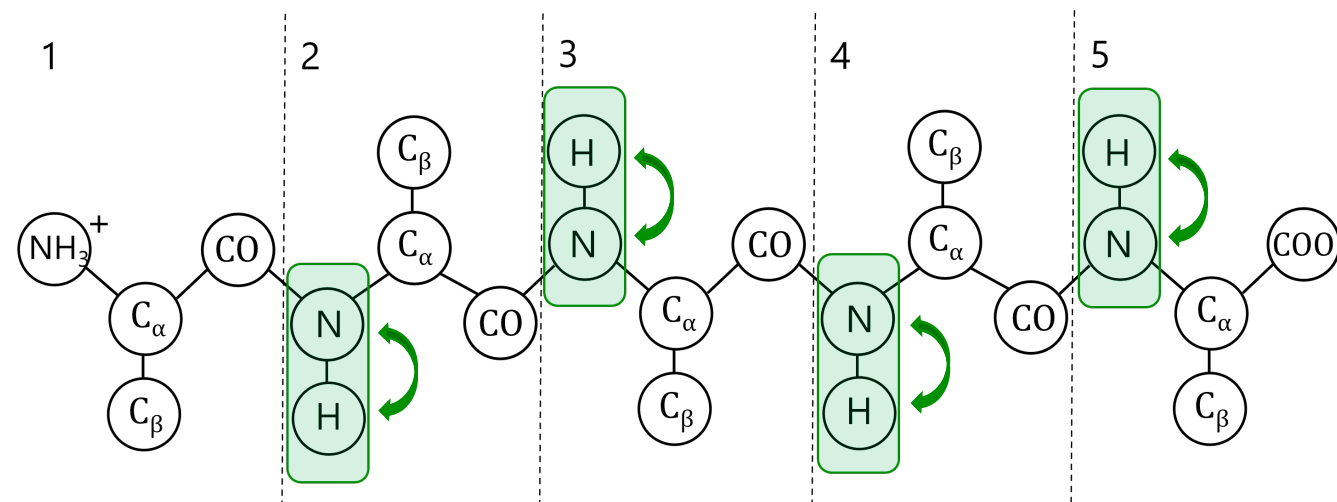
korelace přes prostor

- NOESY

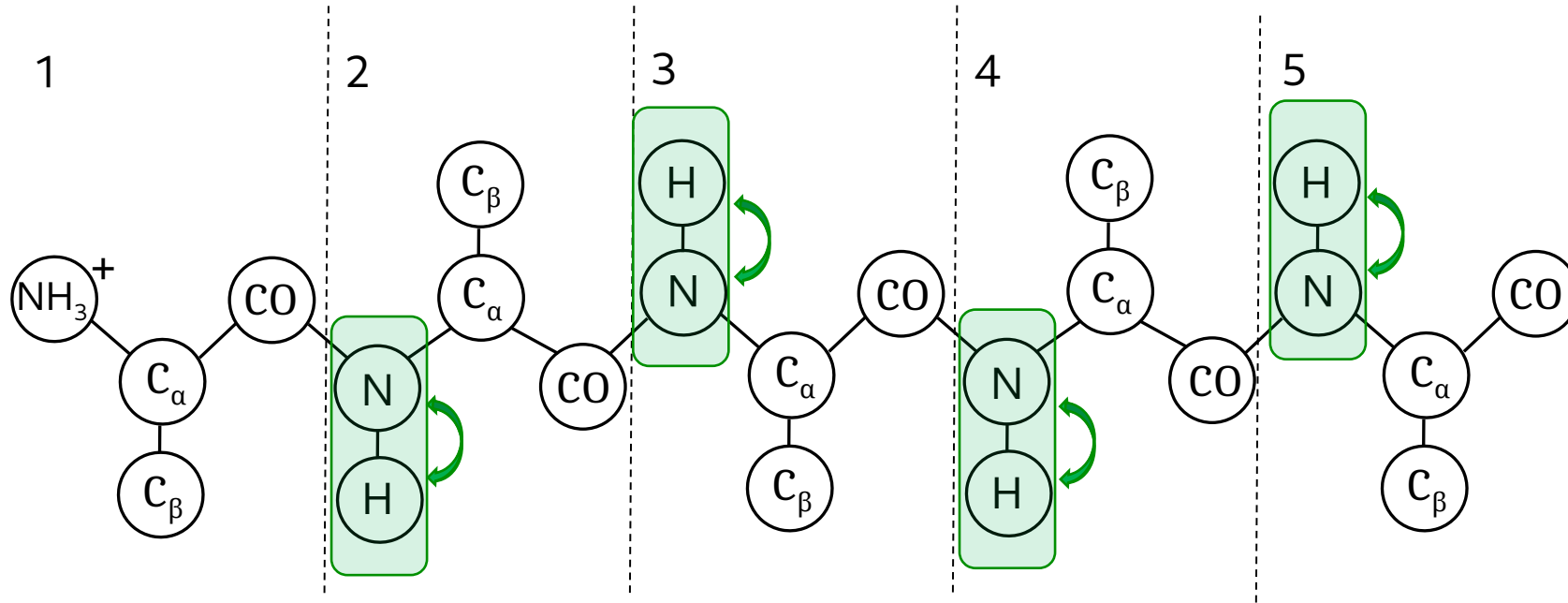


korelace přes vazby

- HSQC
- HNCACB
- HN(CO)CACB

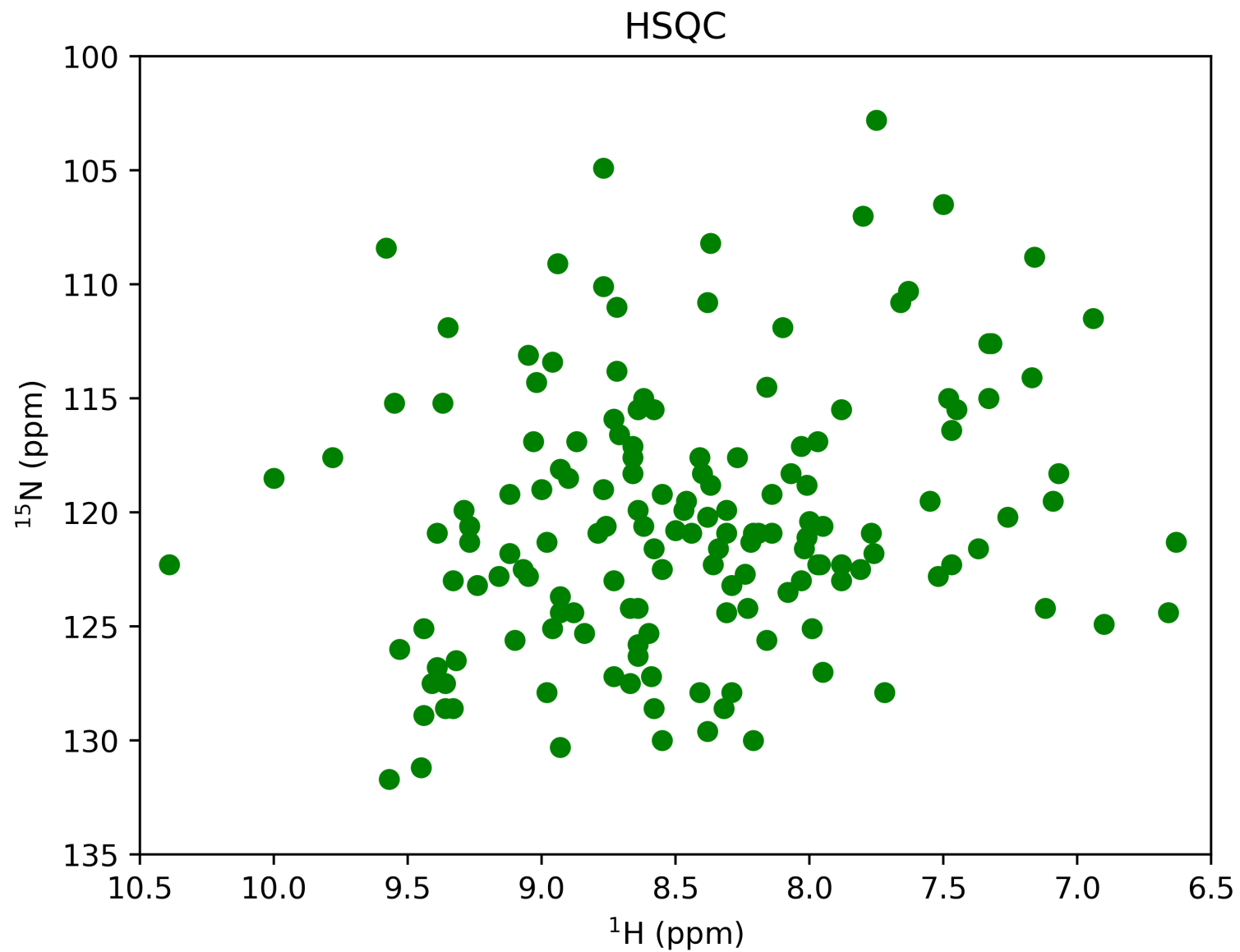


Sekvenční Přiřazení HSQC

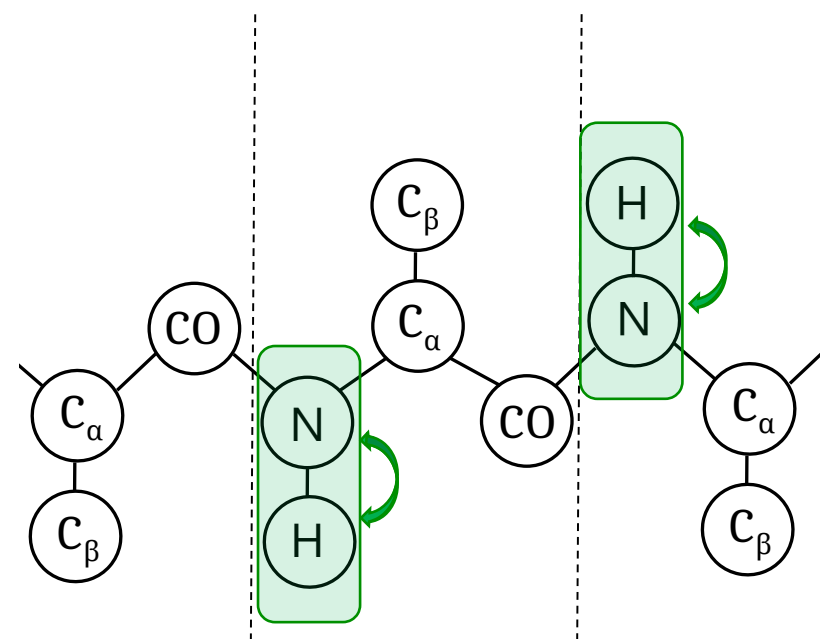


HSQC: korelace H-N vazby

2D HSQC

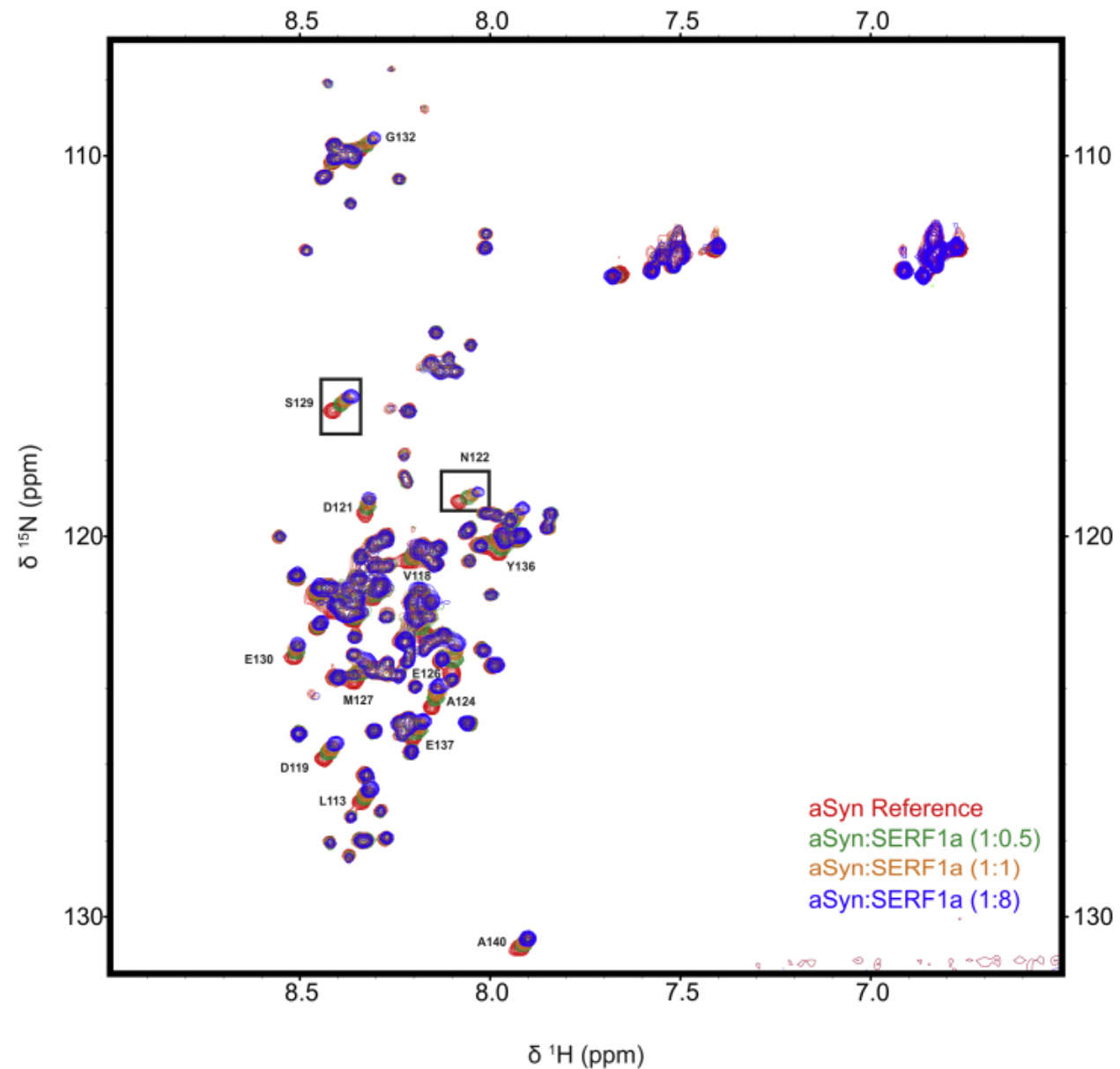
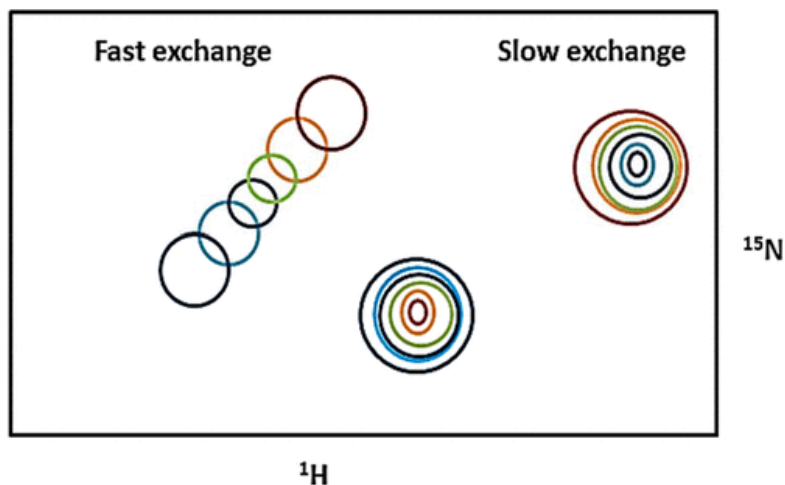


HSQC: korelace H-N vazby

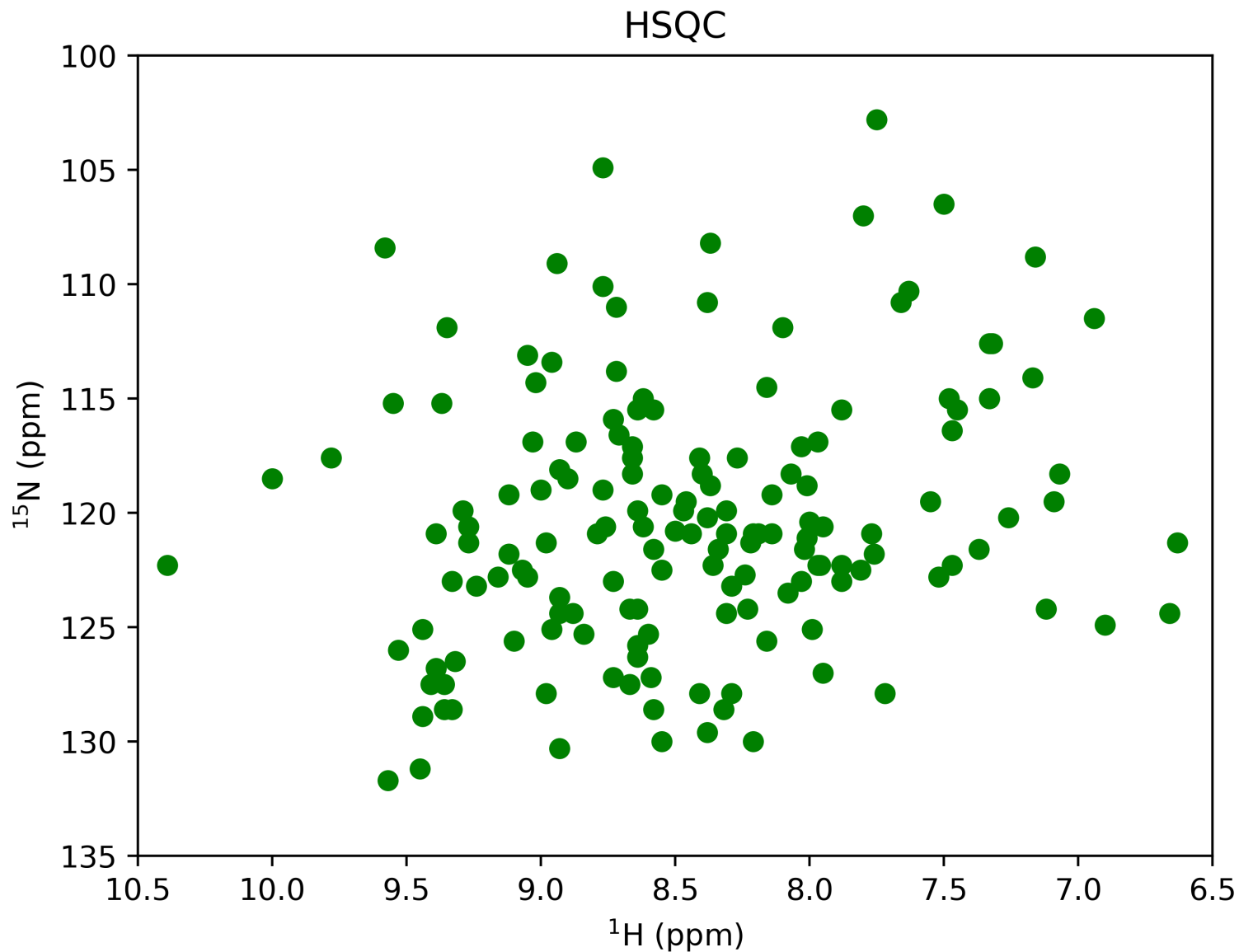


Titrační experimenty

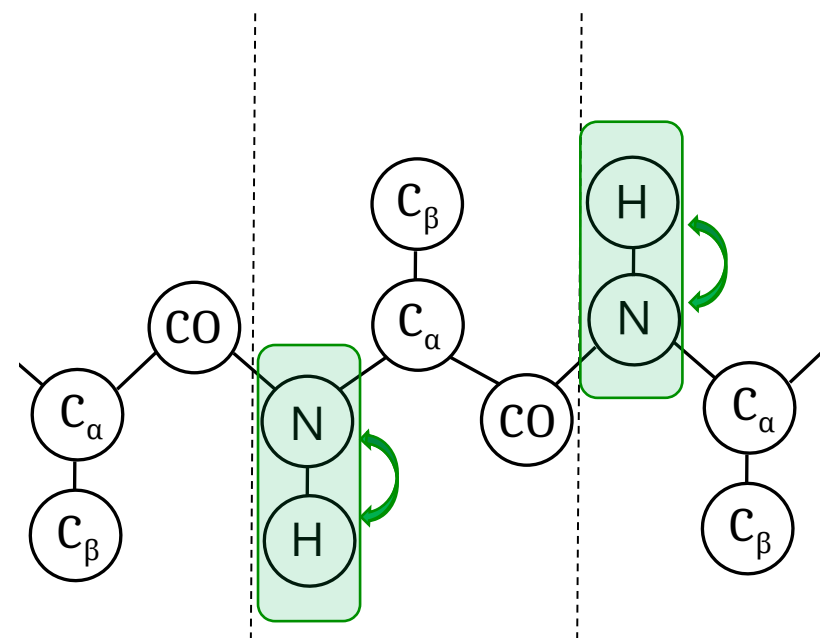
- isotopicky značený protein titrovaný neznačeným ligandem
- posun peaků v okolí **vazebných míst** nebo **při konformačních změnách**
- určení afinity – K_d



2D HSQC



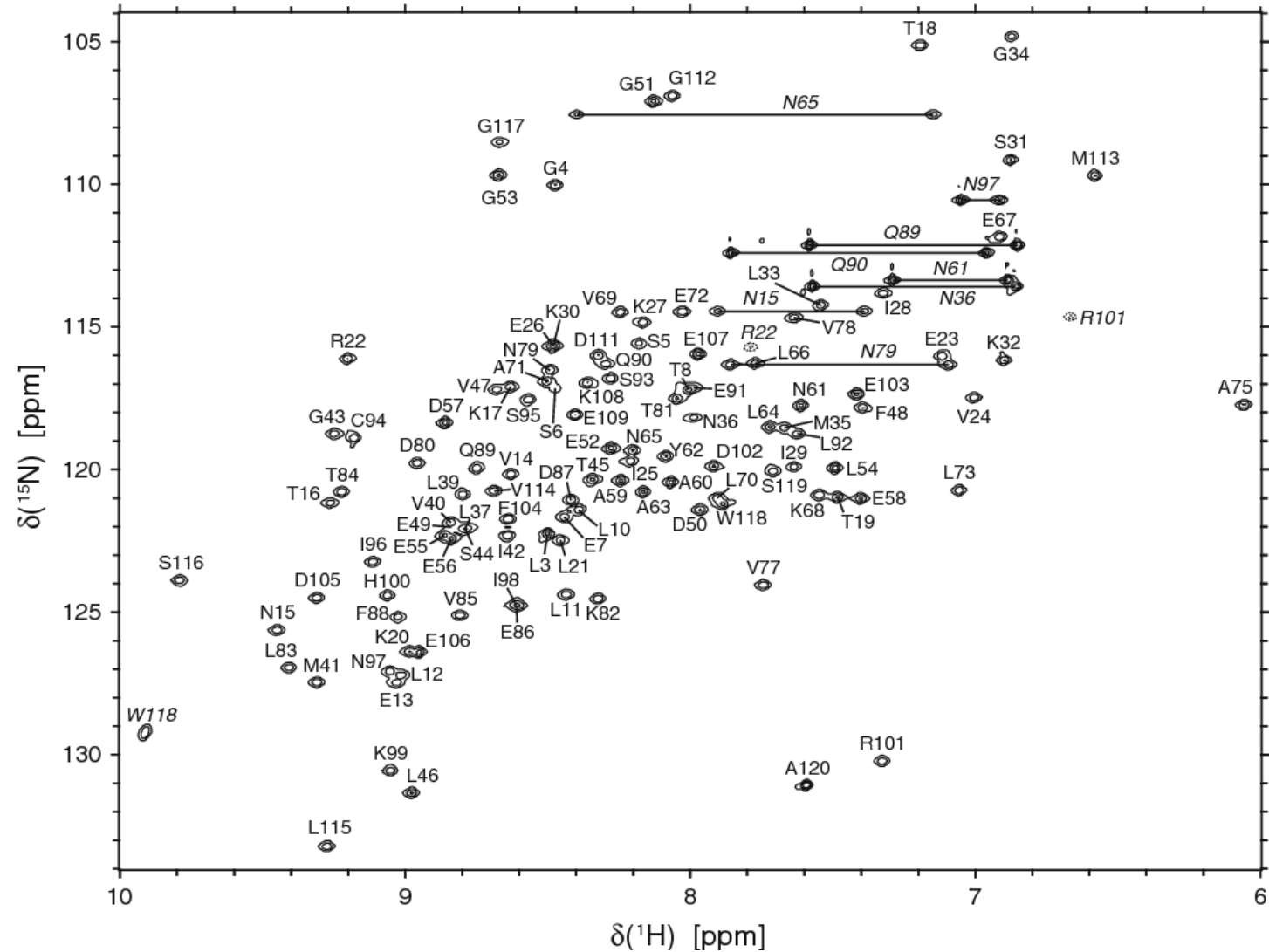
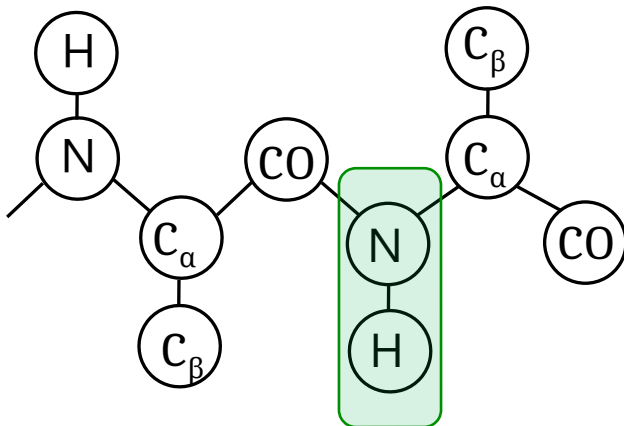
HSQC: korelace H-N vazby



**Kterou aminokyselinu v tomto HSQC spektru neuvидíme?
Které další NH korelace naopak vidět můžeme?**

2D spektrum proteinu - HSQC

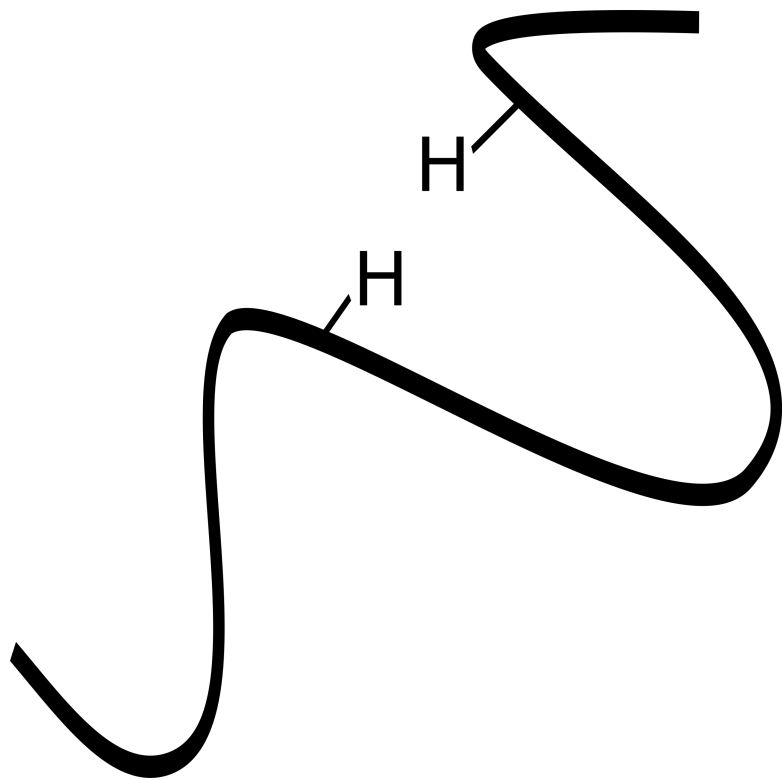
- korelace ^{15}N a ^1H peptidové vazby v proteinové páteři
(který vodík se váže na který dusík)
- 1 peak \approx 1 aminokyselina
- **otisk prstu proteinu**



Experimenty

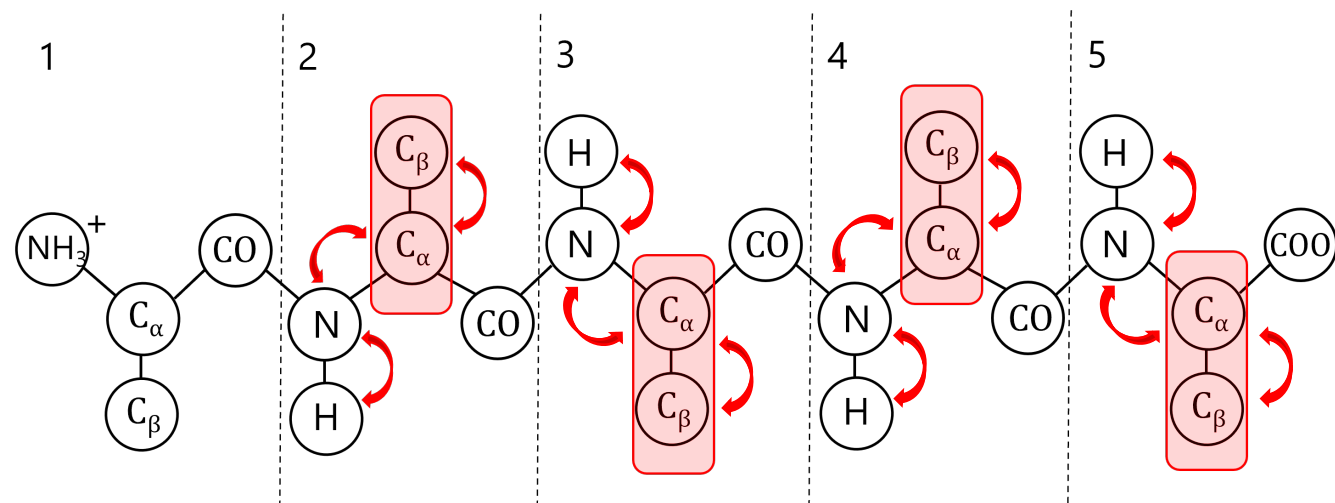
korelace přes prostor

- NOESY



korelace přes vazby

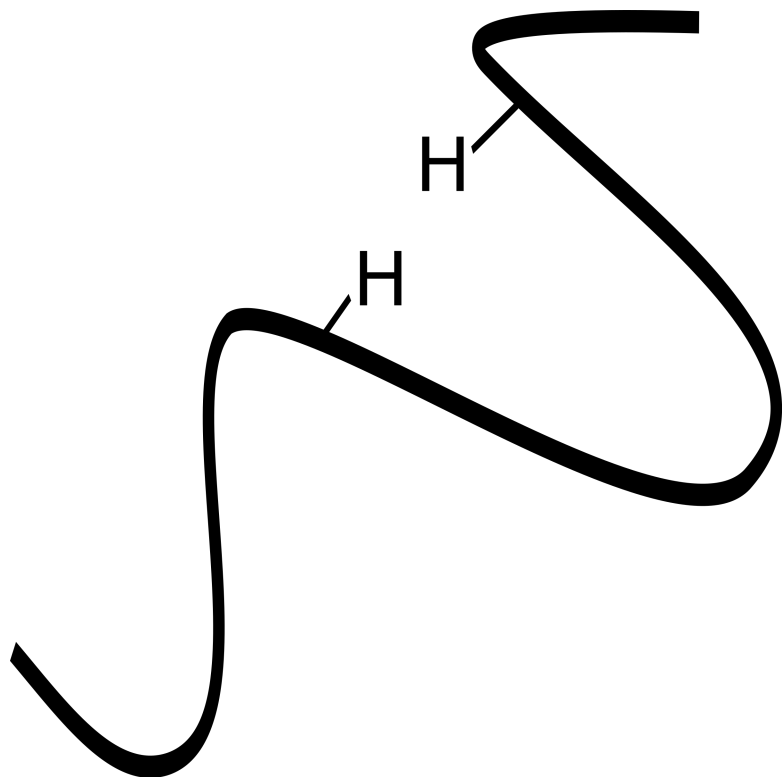
- HSQC
- **HNCACB**
- HN(CO)CACB



Experimenty

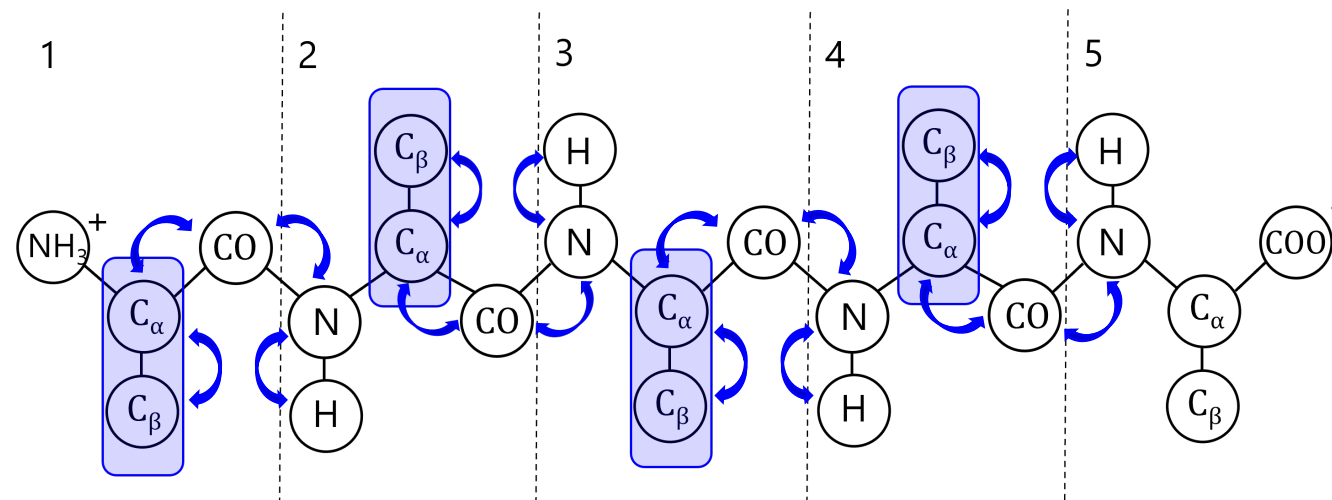
korelace přes prostor

- NOESY



korelace přes vazby

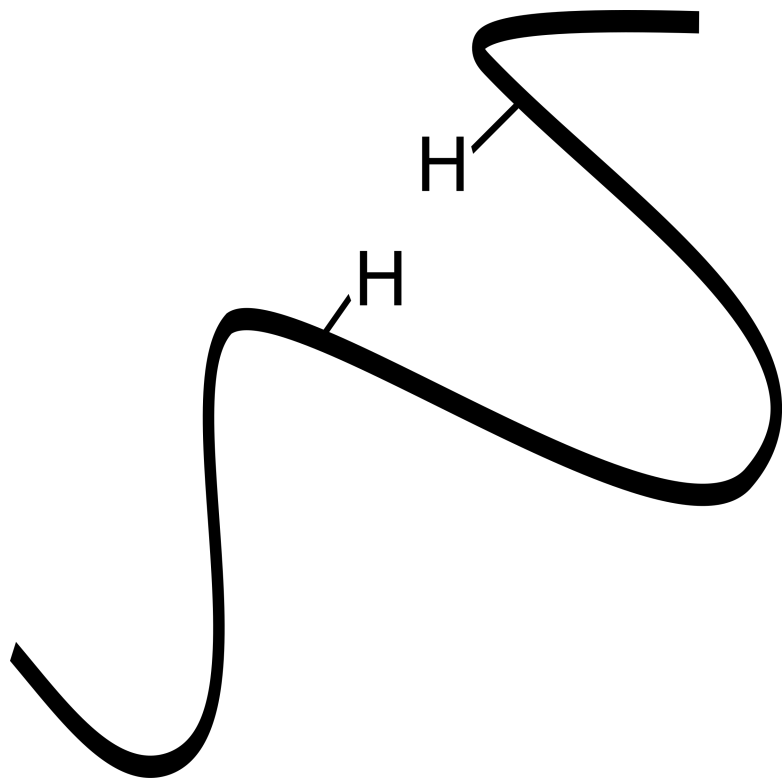
- HSQC
- HNCACB
- **HN(CO)CACB**



Experimenty

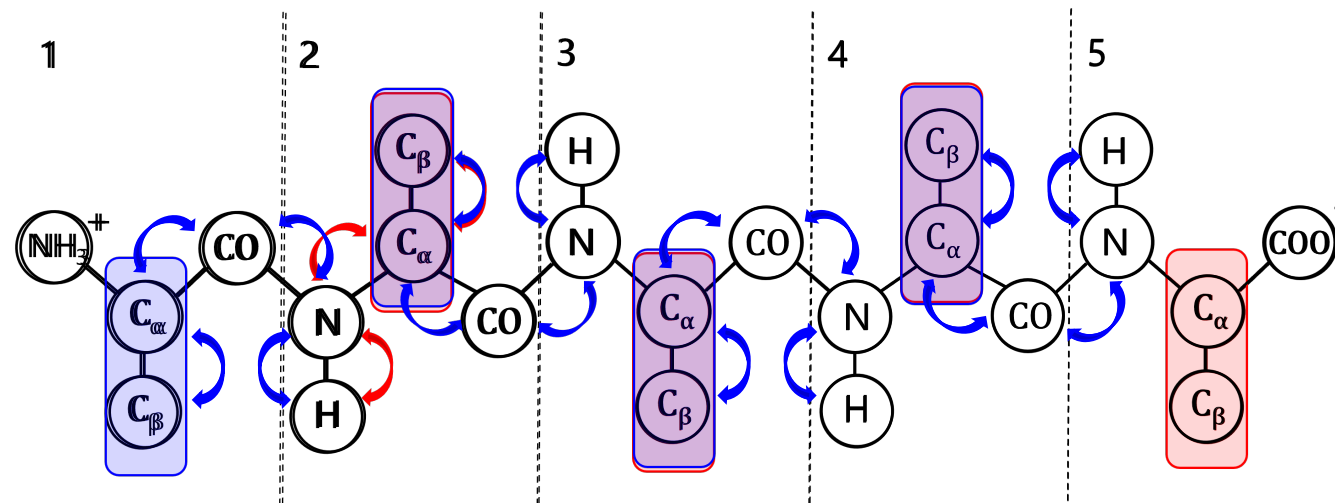
korelace přes prostor

- NOESY



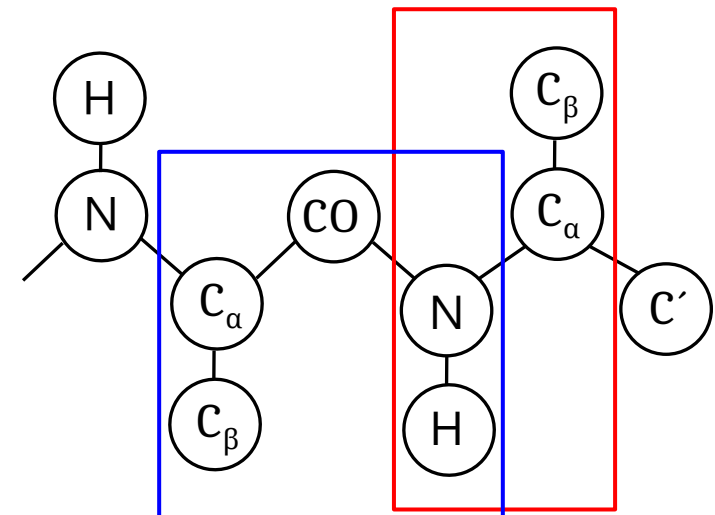
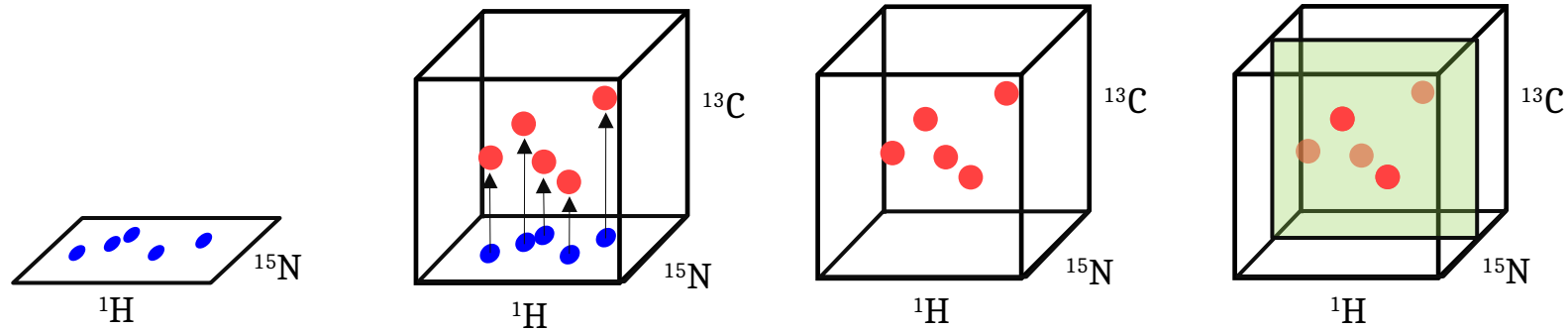
korelace přes vazby

- HSQC
- HNCACB
- **HN(CO)CACB**



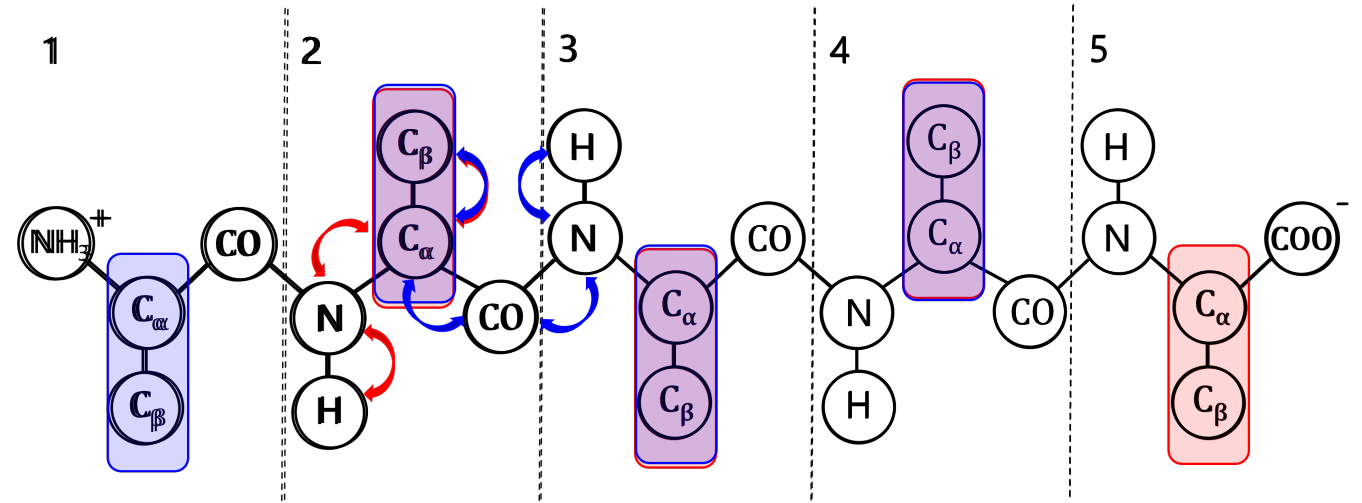
Přiřazení HSQC

- 3D experimenty
- další zvýšení rozlišení
- přiřazení peaků aminokyselinám
- (assignment)
- série komplementárních spekter
HNCACB, **HN(CO)CACB**

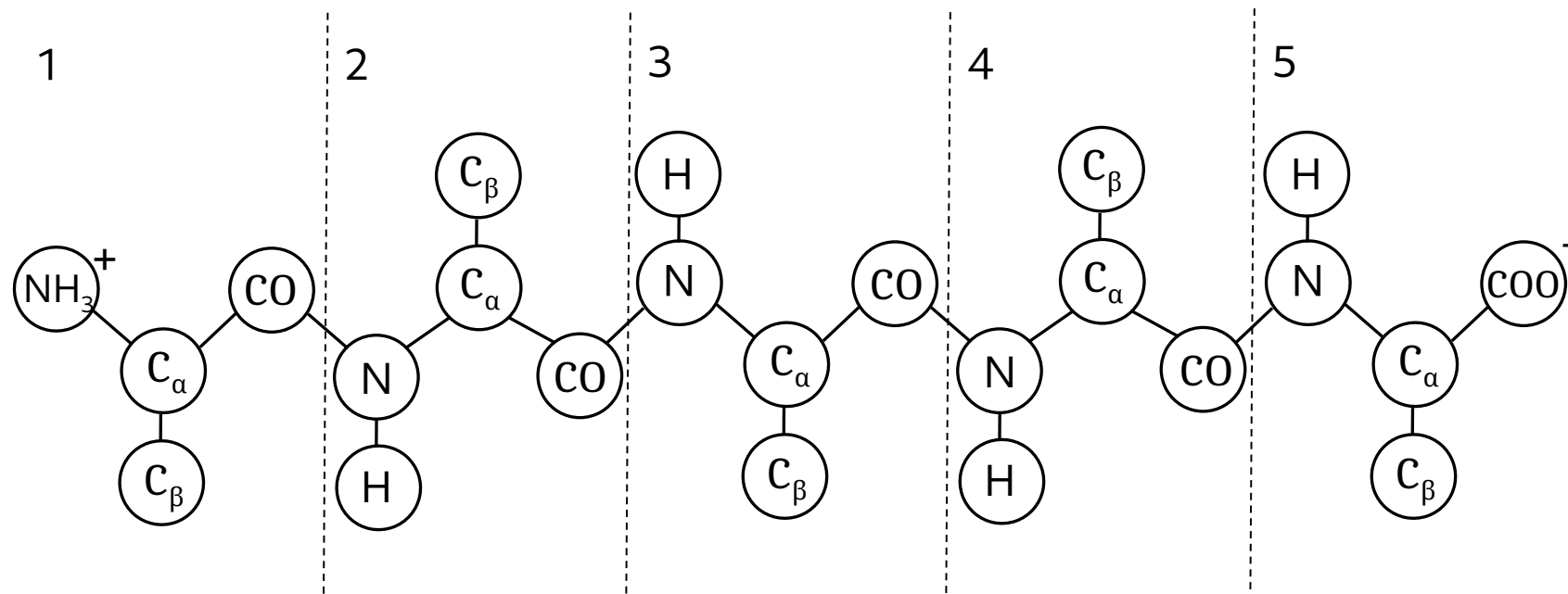


Přiřazení HSQC proteinu

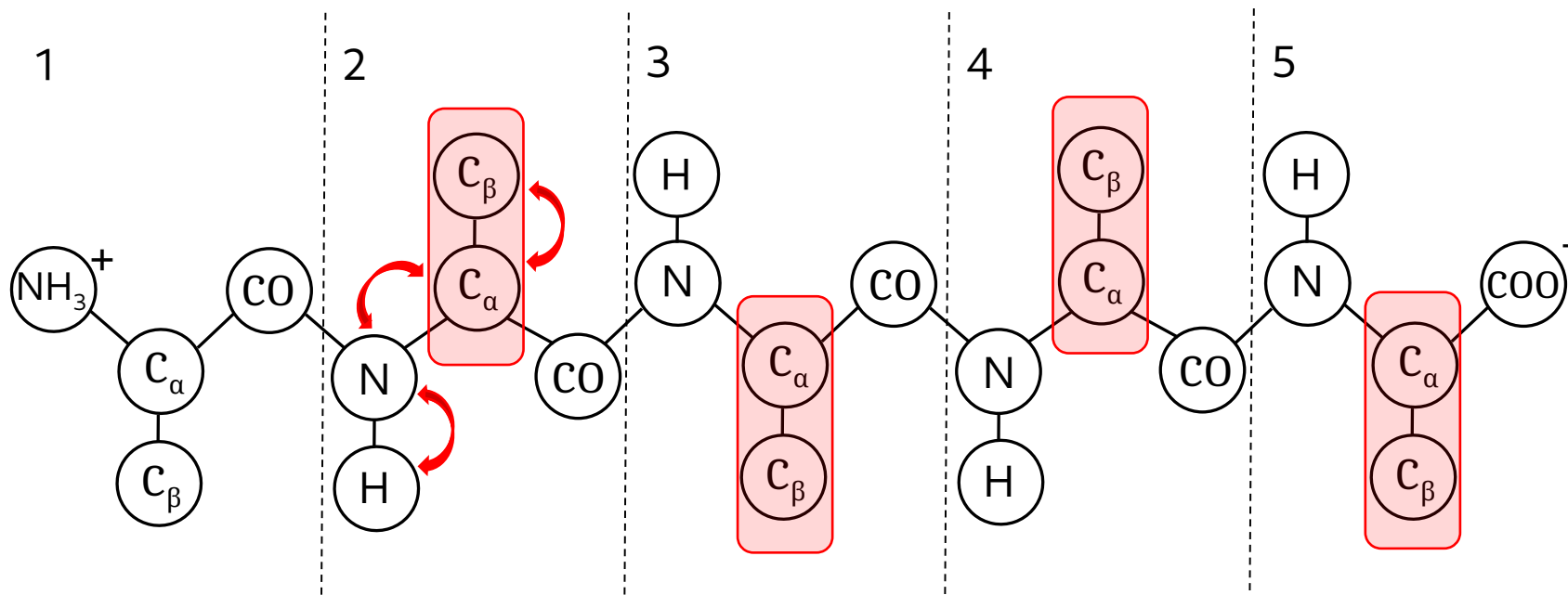
- 3D experimenty
- další zvýšení rozlišení
- přiřazení peaků aminokyselinám
- série komplementárních spekter
HNCA (současné AMK)
HN(CO)CA (předchozí AMK)



Sekvenční Přiřazení HSQC

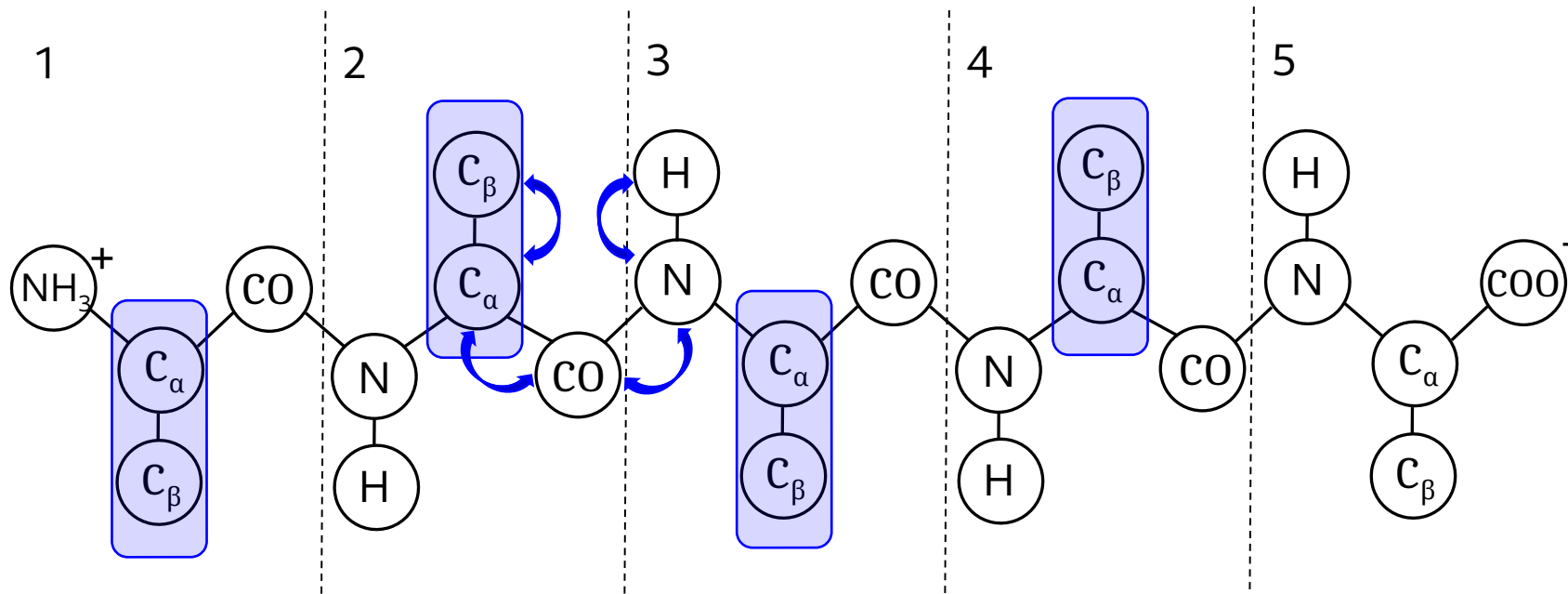


Sekvenční Přiřazení HSQC



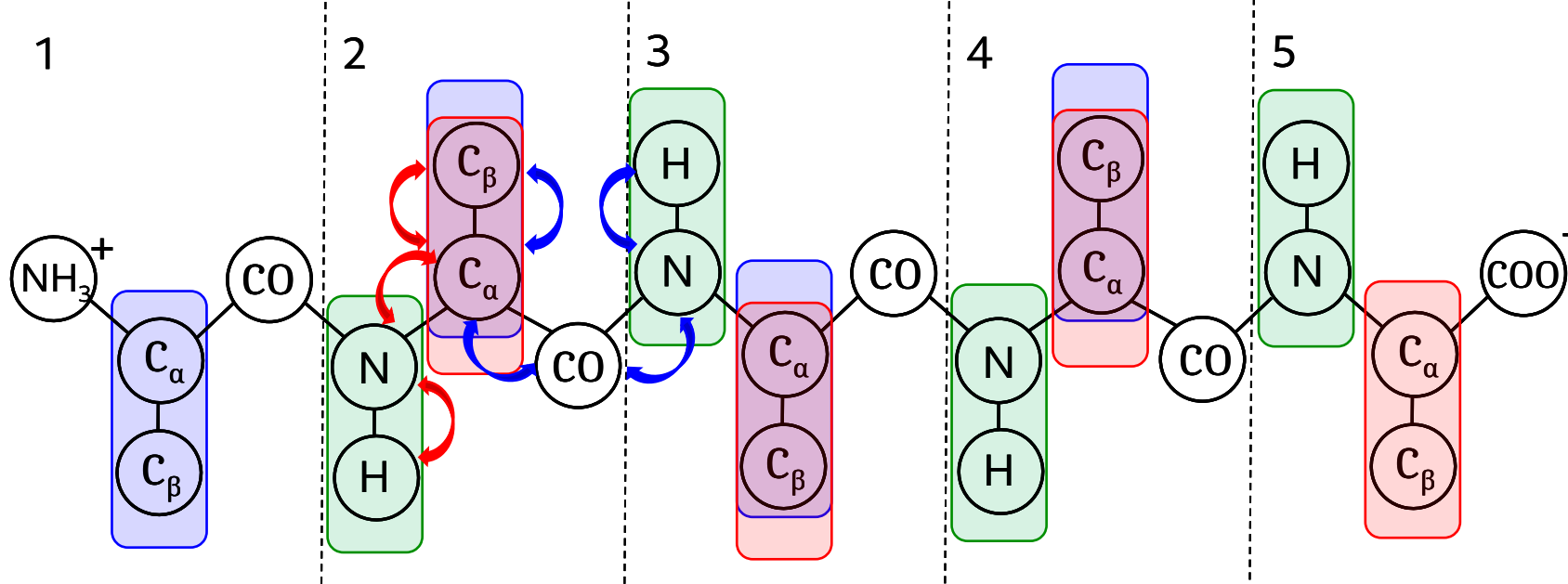
HNCACB: C_{α} a C_{β} současné AMK

Sekvenční Přiřazení HSQC



HN(CO)CACB: C_α a C_β předchází AMK

Sekvenční Přiřazení HSQC

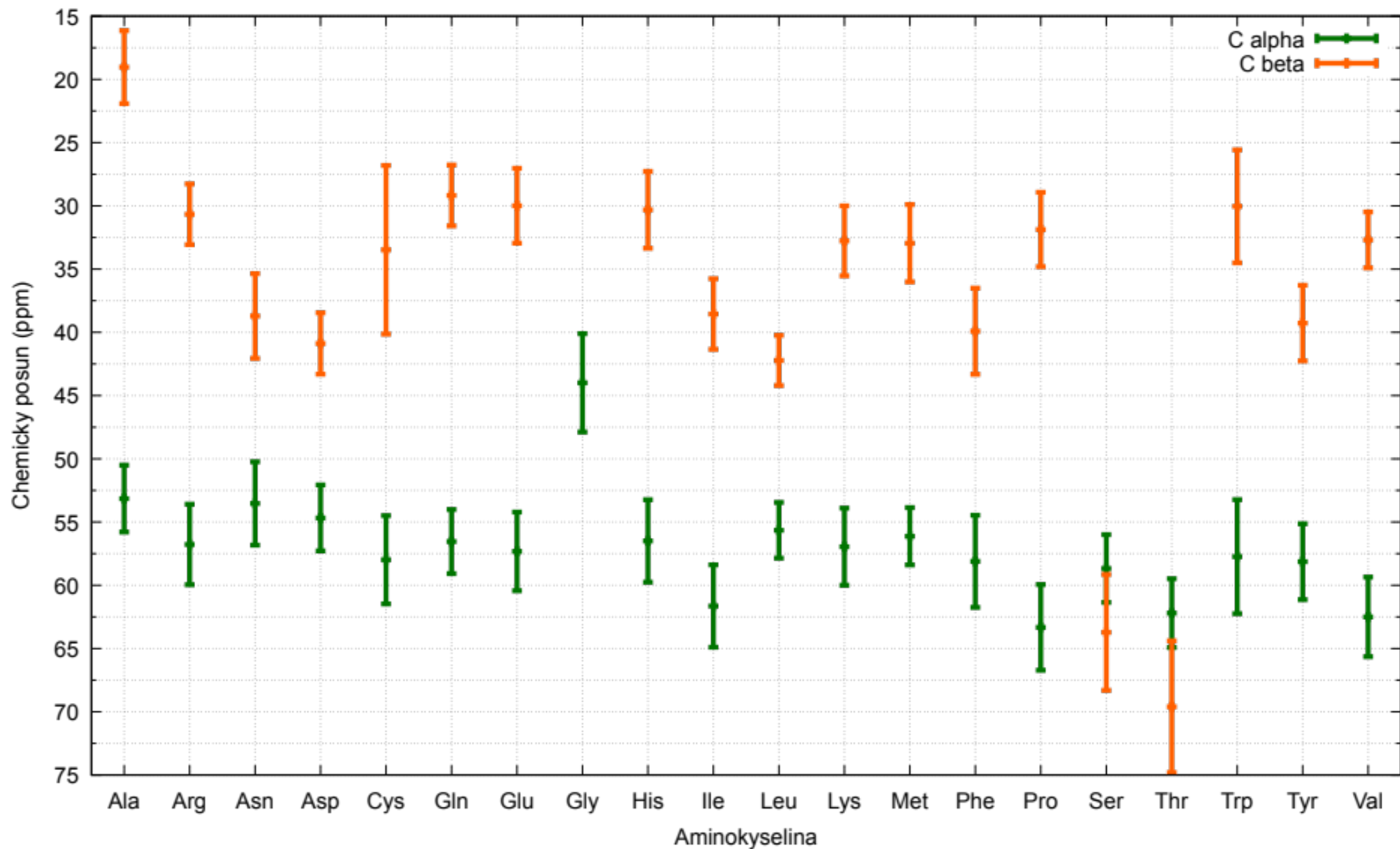


HSQC: korelace H-N vazby

HNCACB: C α a C β současné AMK

HN(CO)CACB: C α a C β předchozí AMK

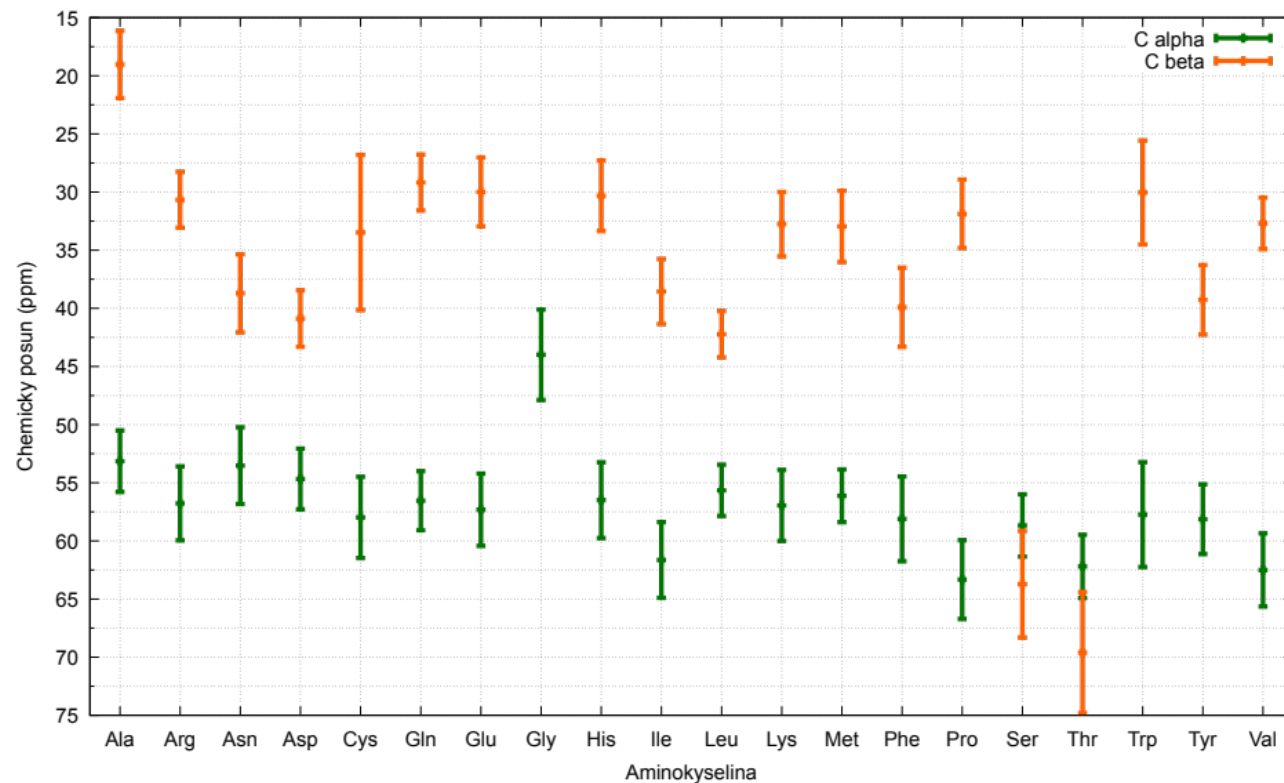
Typické chemické posuny C α a C β aminokyselin



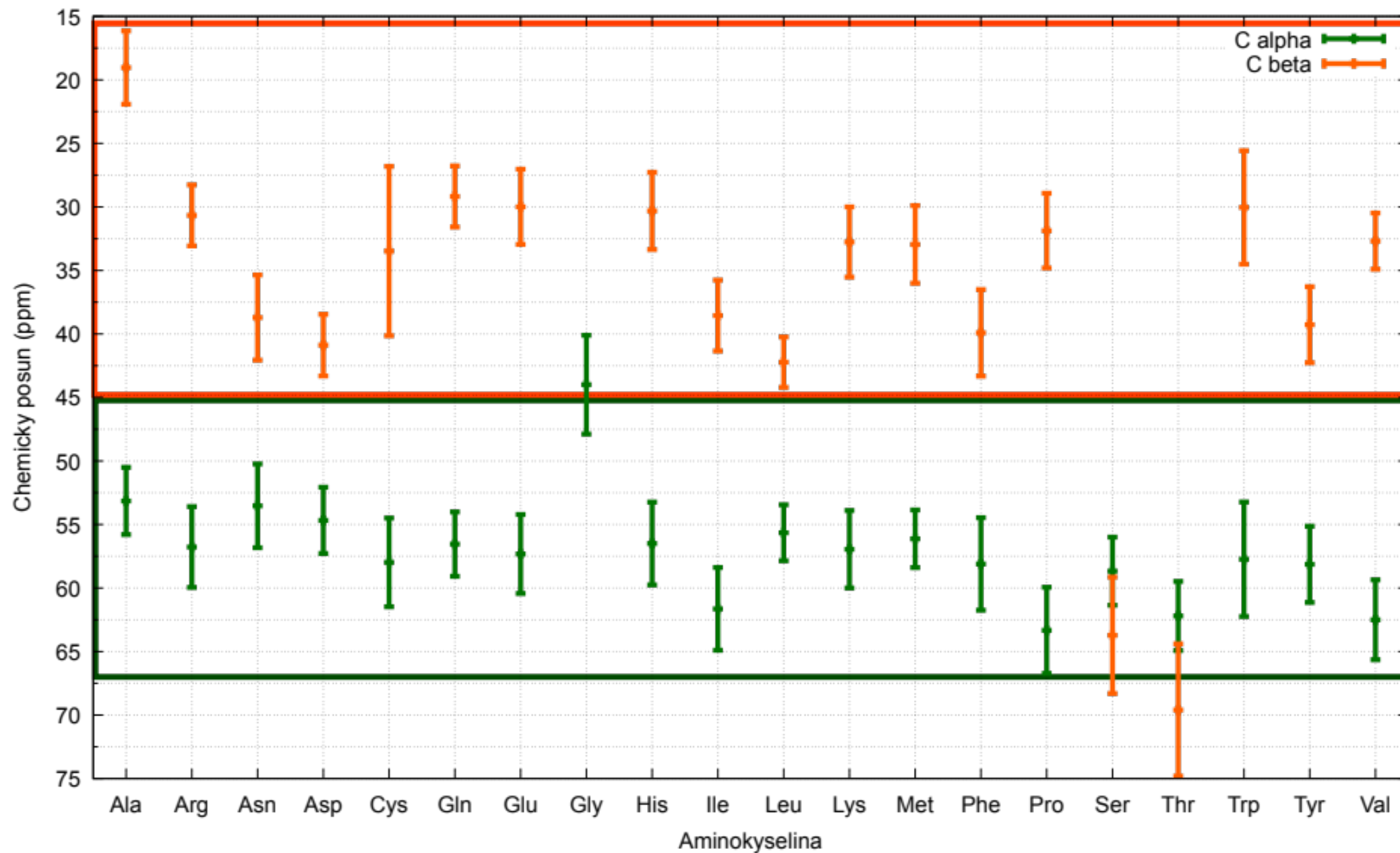
Typické chemické posuny C α a C β aminokyselin

průměrné hodnoty z BMRB
(Biological Magnetic Resonance Databank)

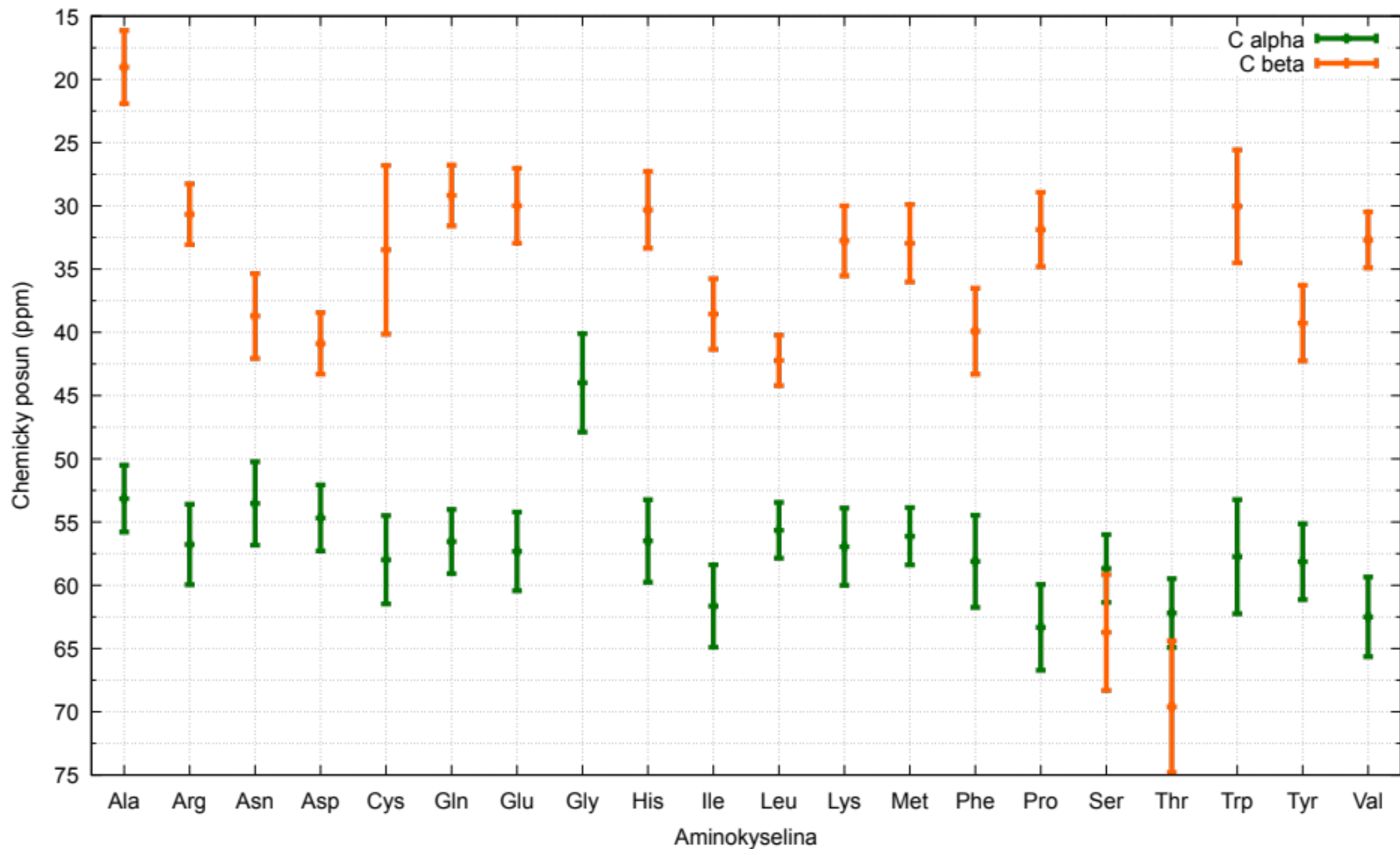
<https://bmr.io/>

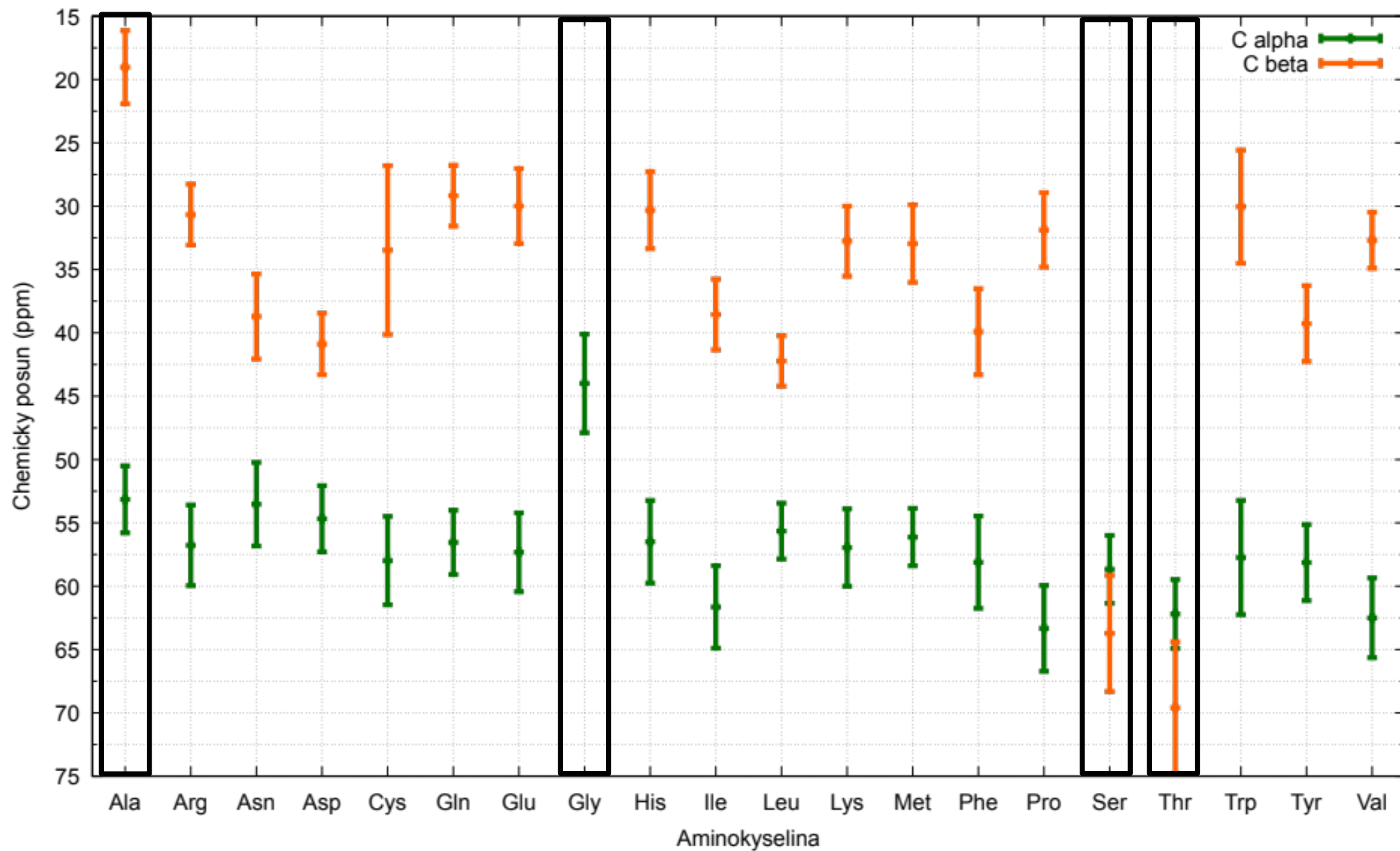


Typické chemické posuny C α a C β aminokyselin

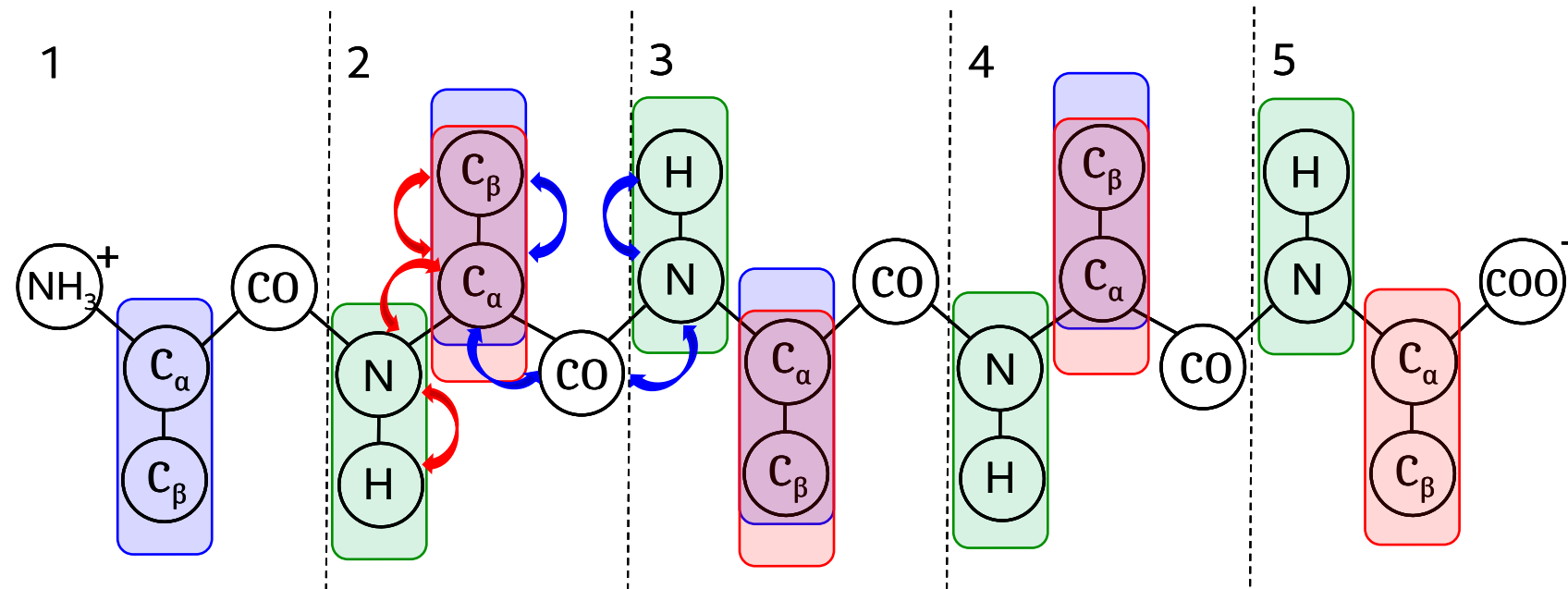


Které aminokyseliny mají specifickou pozici C α a C β ?





Sekvenční Přiřazení HSQC

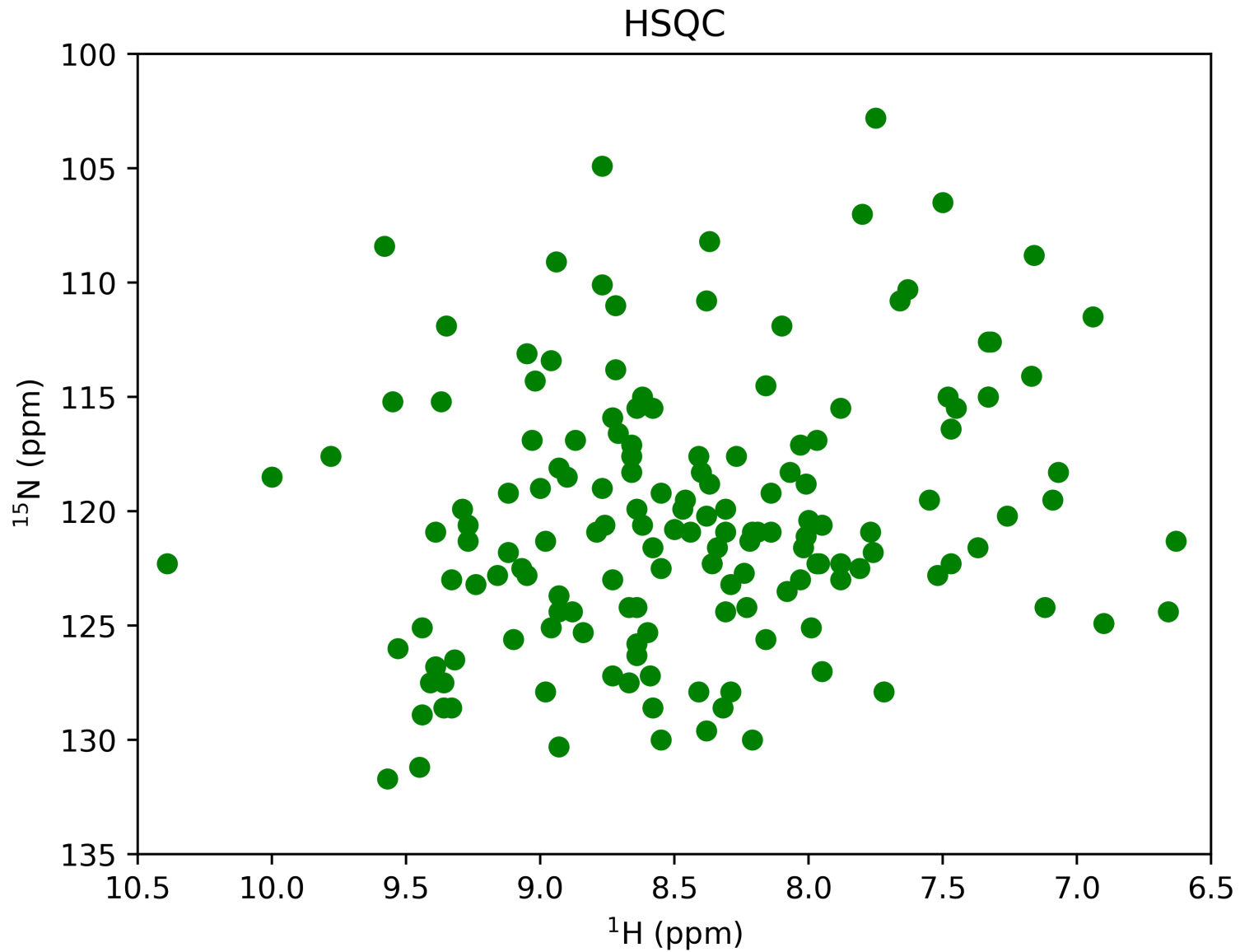


HSQC: korelace H-N vazby

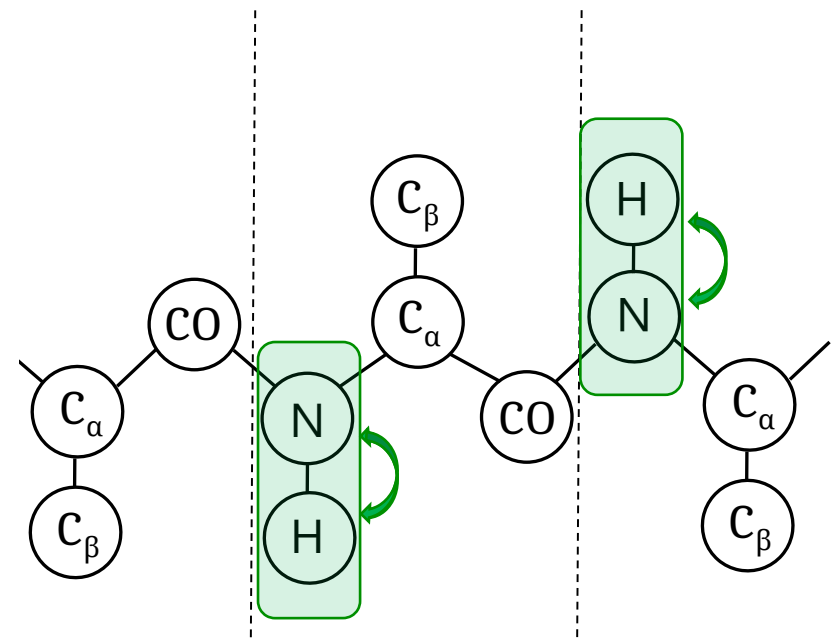
HNCACB: C α a C β současné AMK

CACB(CO)NH: C α a C β předchozí AMK

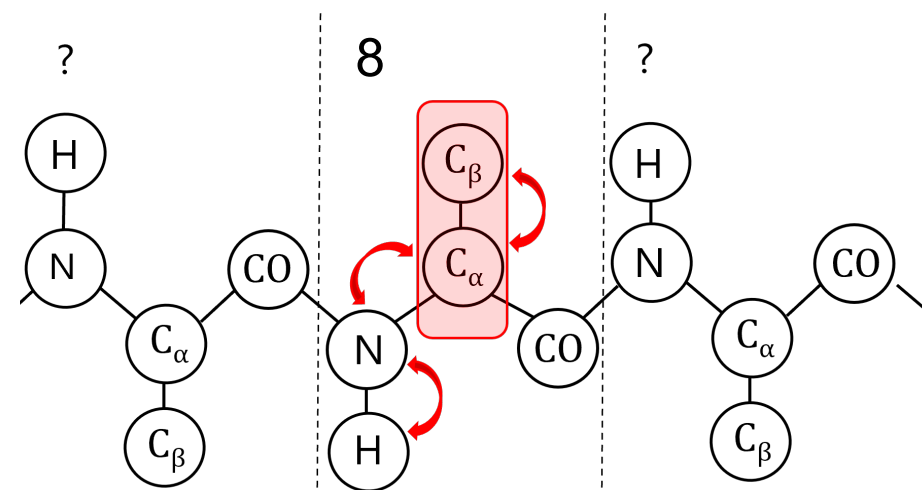
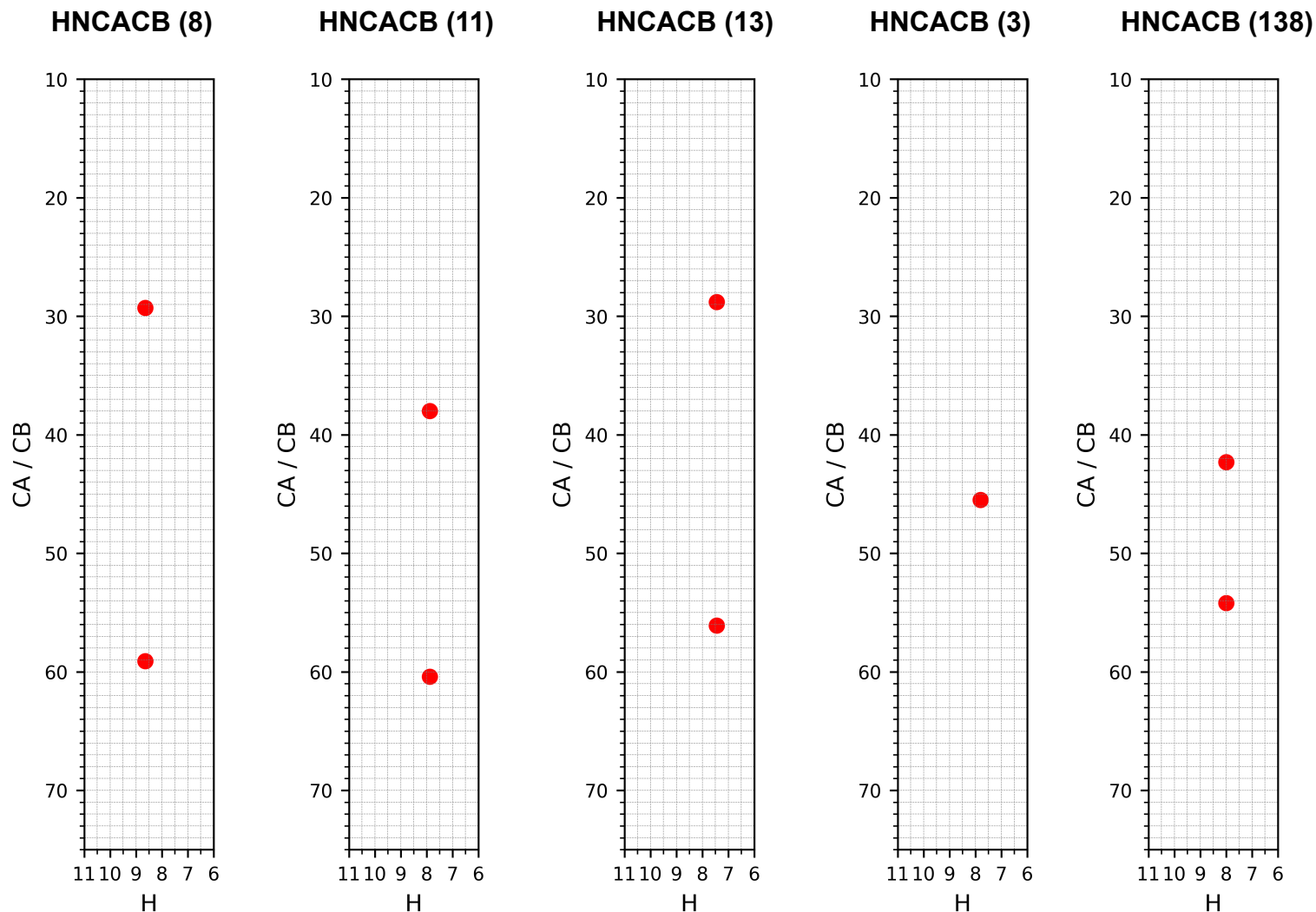
2D HSQC



HSQC: korelace H-N vazby



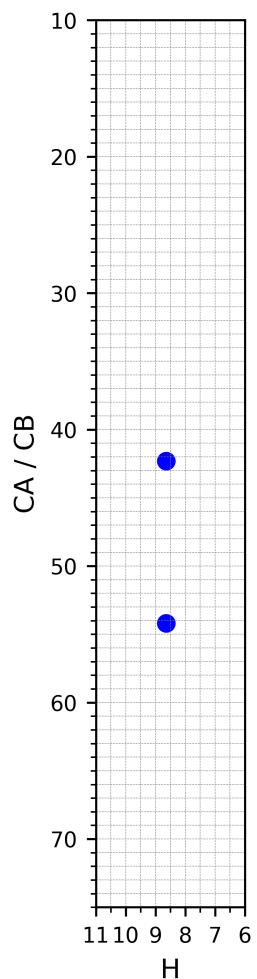
Postup při přiřazování - HNCACB



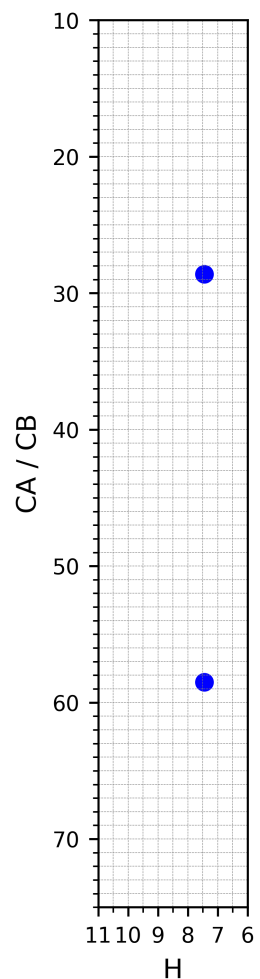
HNCACB: C_α a C_β současné AMK

Postup při přiřazování – HN(CO)CACB

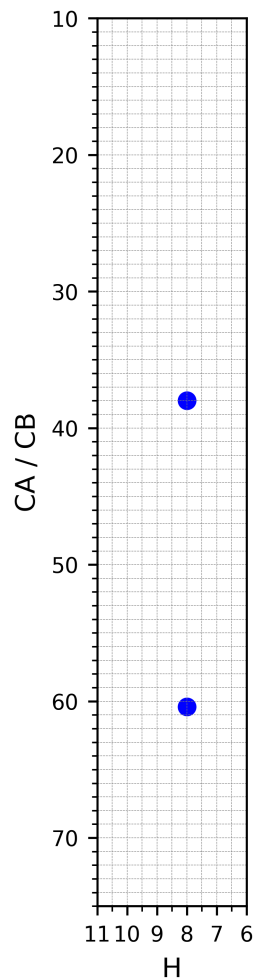
HN(CO)CACB (8)



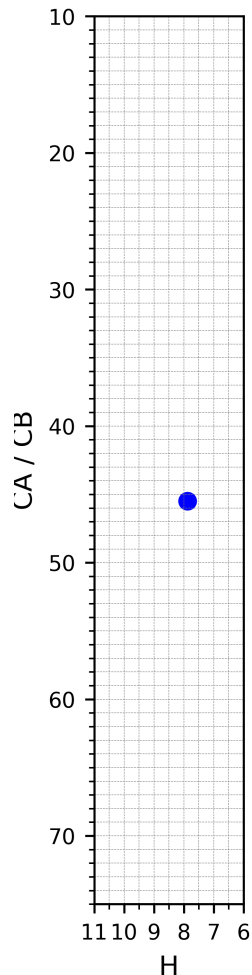
HN(CO)CACB (13)



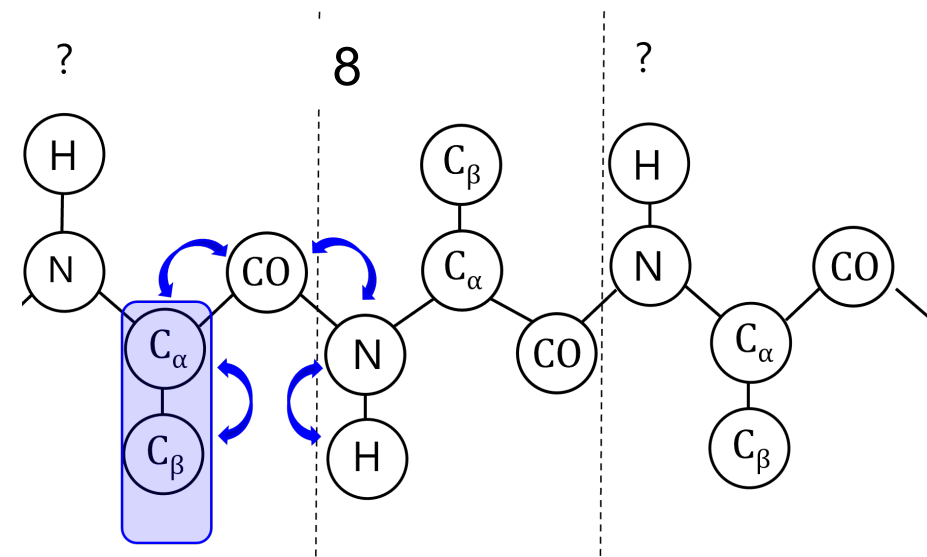
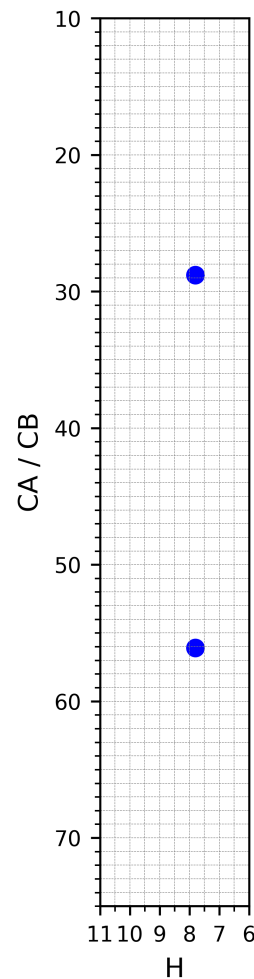
HN(CO)CACB (138)



HN(CO)CACB (11)



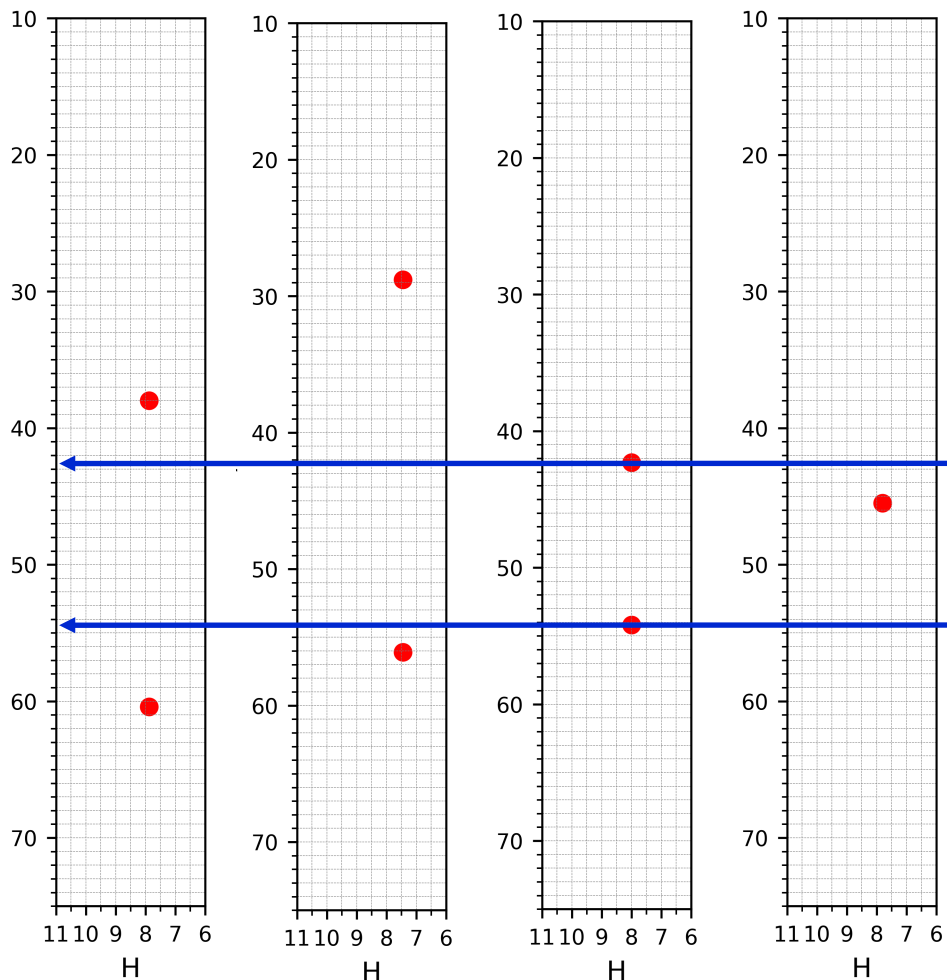
HN(CO)CACB (3)



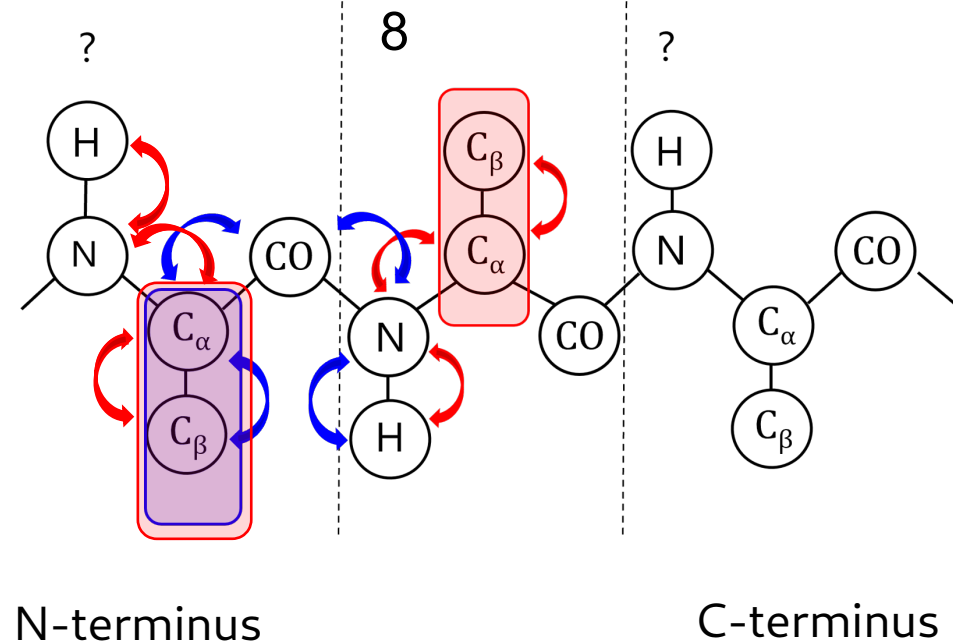
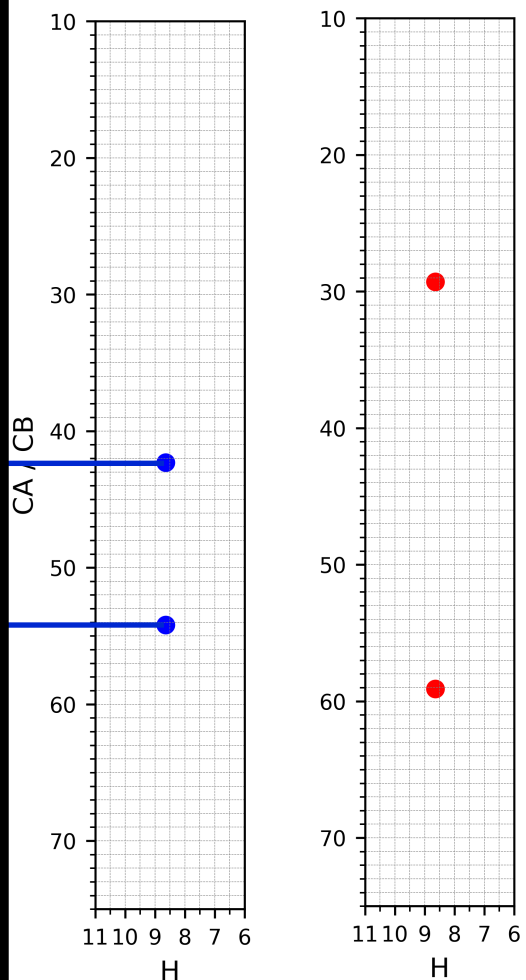
HN(CO)CACB: C α a C β předchozí AMK

Postup při přiřazování -138-8

HNCACB (11) HNCACB (13) HNCACB (138) HNCACB (3)



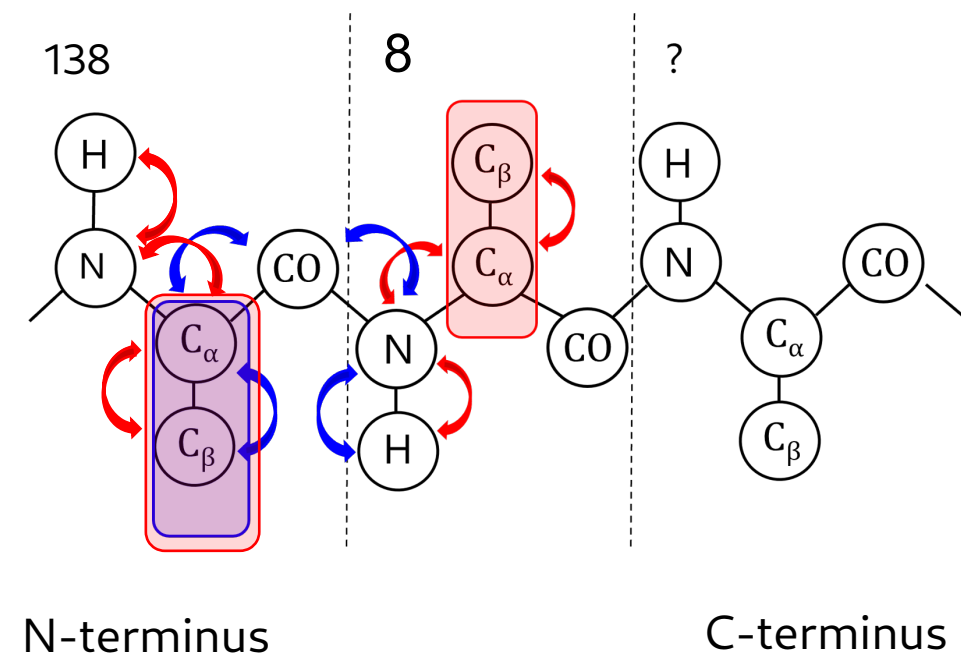
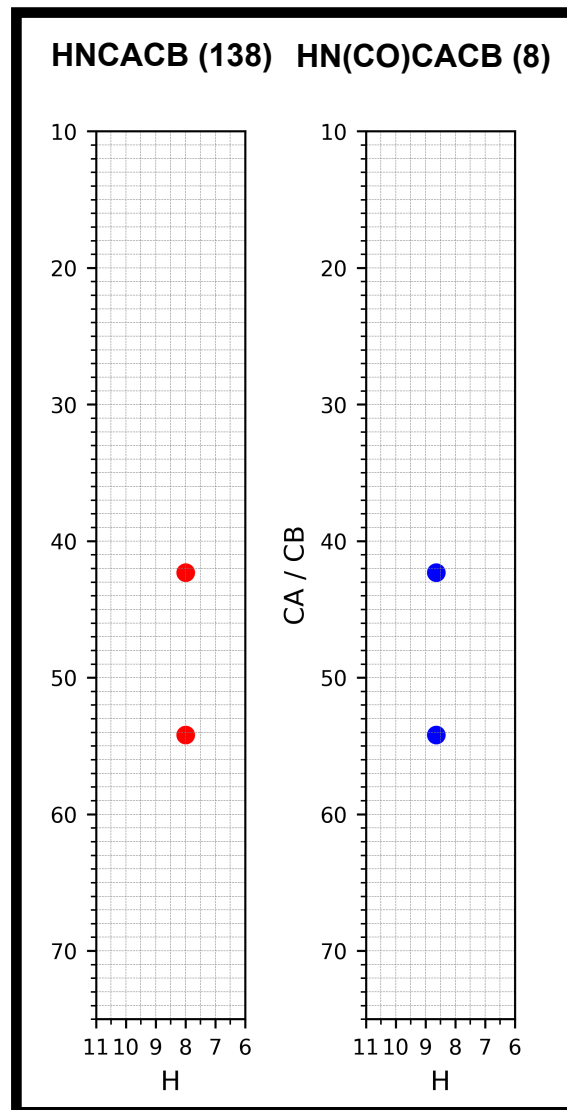
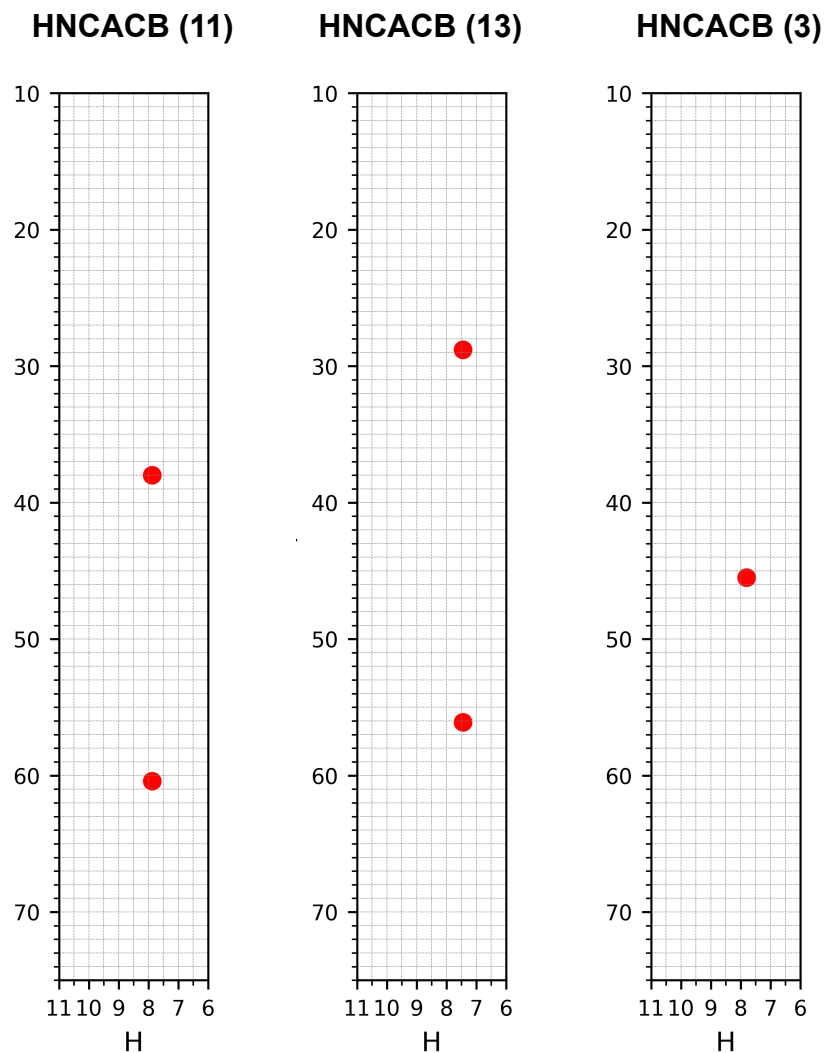
HN(CO)CACB (8) HNCACB (8)



HNCACB: C_α a C_β současné AMK

HN(CO)CACB: C_α a C_β předchozí AMK

Postup při přiřazování - 138-8

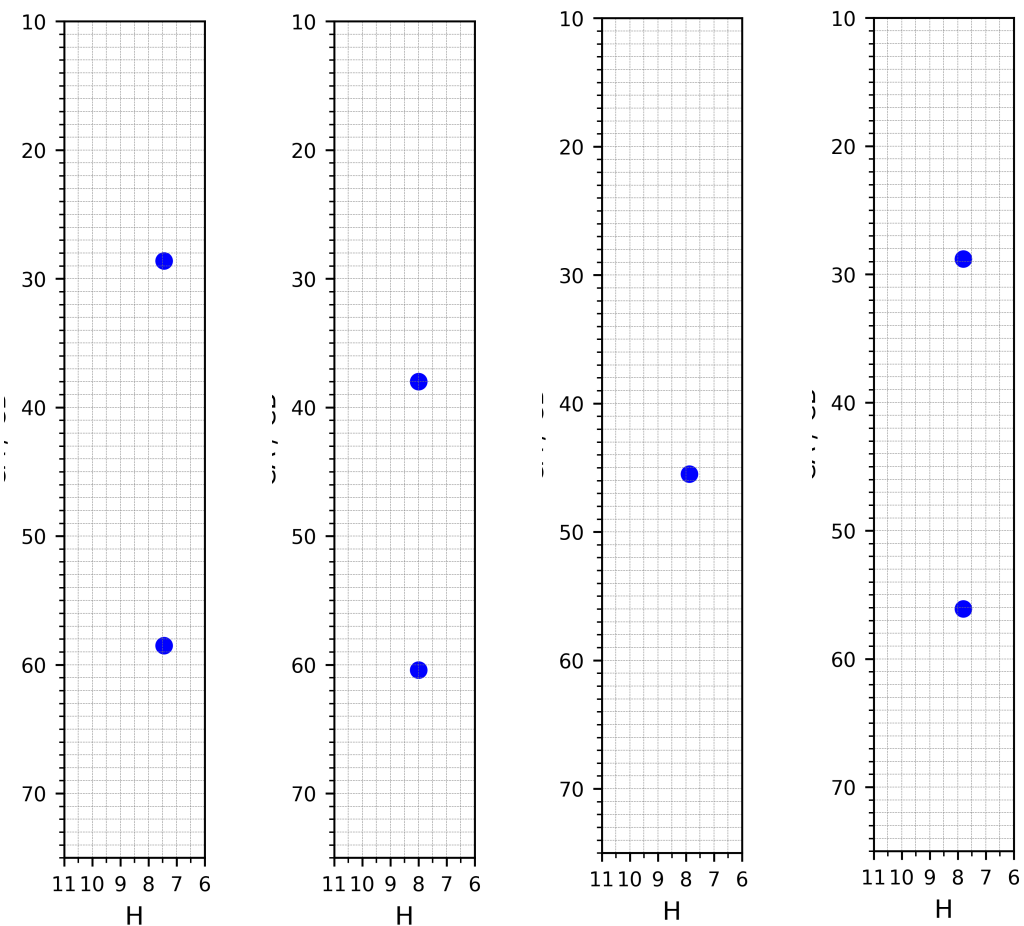


HNCACB: C_α a C_β současné AMK

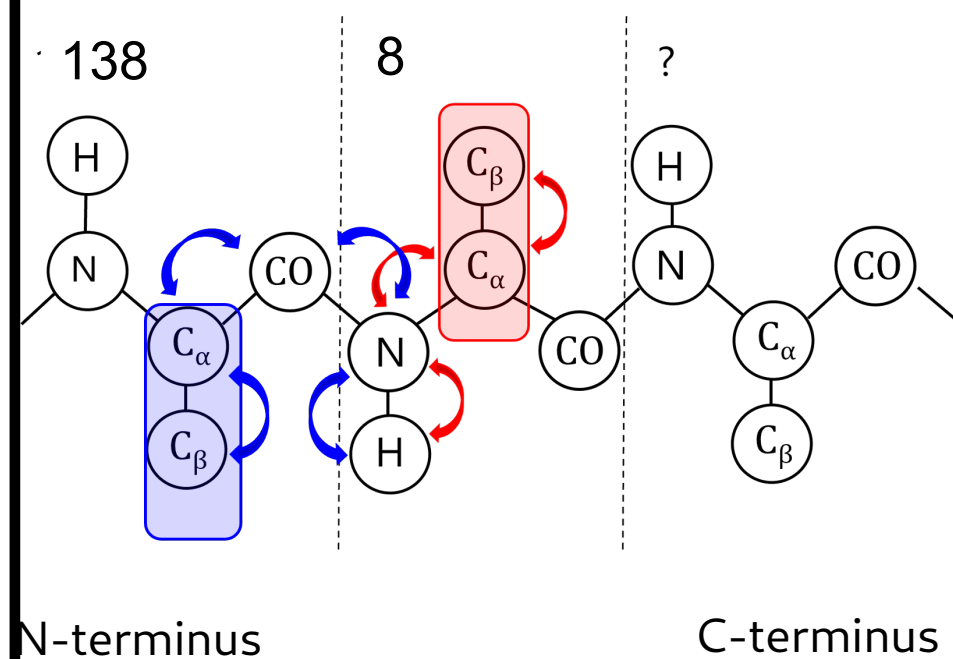
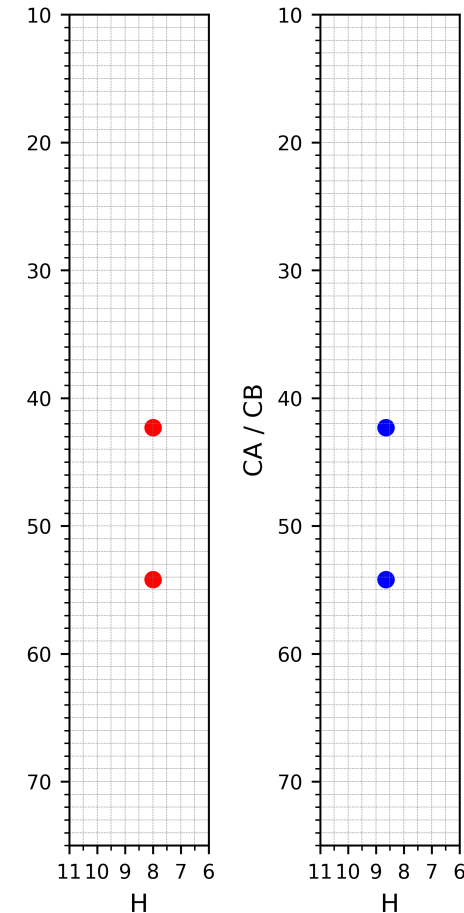
HN(CO)CACB: C_α a C_β předchozí AMK

Postup při přiřazování - 138-8

HN(CO)CACB (13) HN(CO)CACB (138) HN(CO)CACB (11) HN(CO)CACB (3)



HNCACB (138) HN(CO)CACB (8)

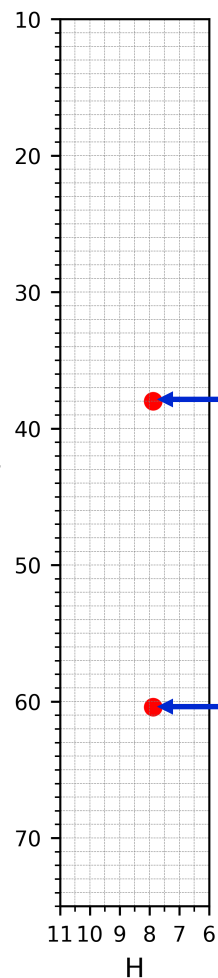


HNCACB: C_α a C_β současné AMK

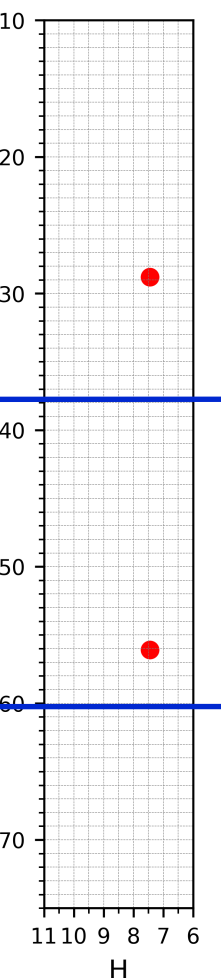
HN(CO)CACB: C_α a C_β předchozí AMK

Postup při přiřazování -11-138-8

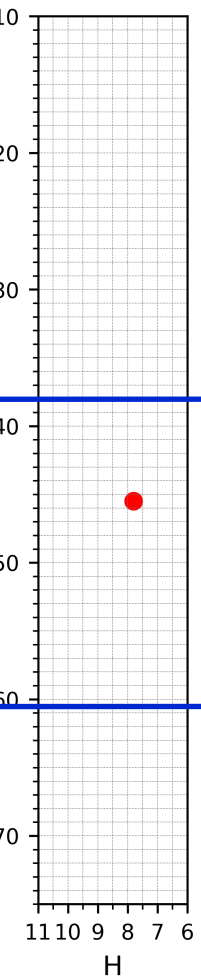
HNCACB (11)



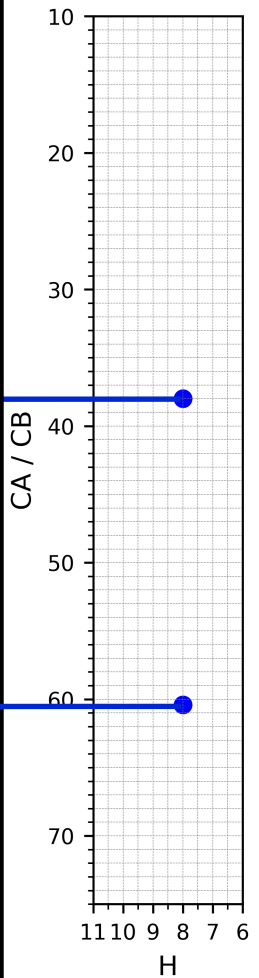
HNCACB (13)



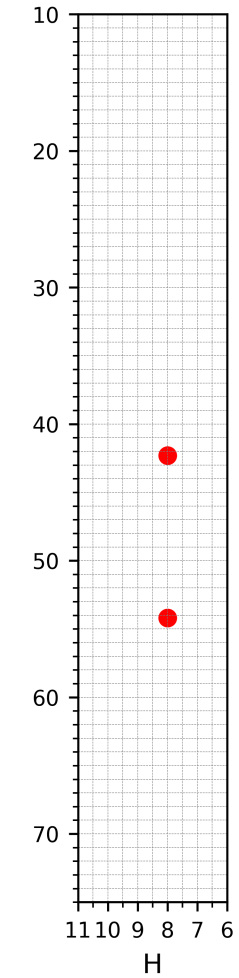
HNCACB (3)



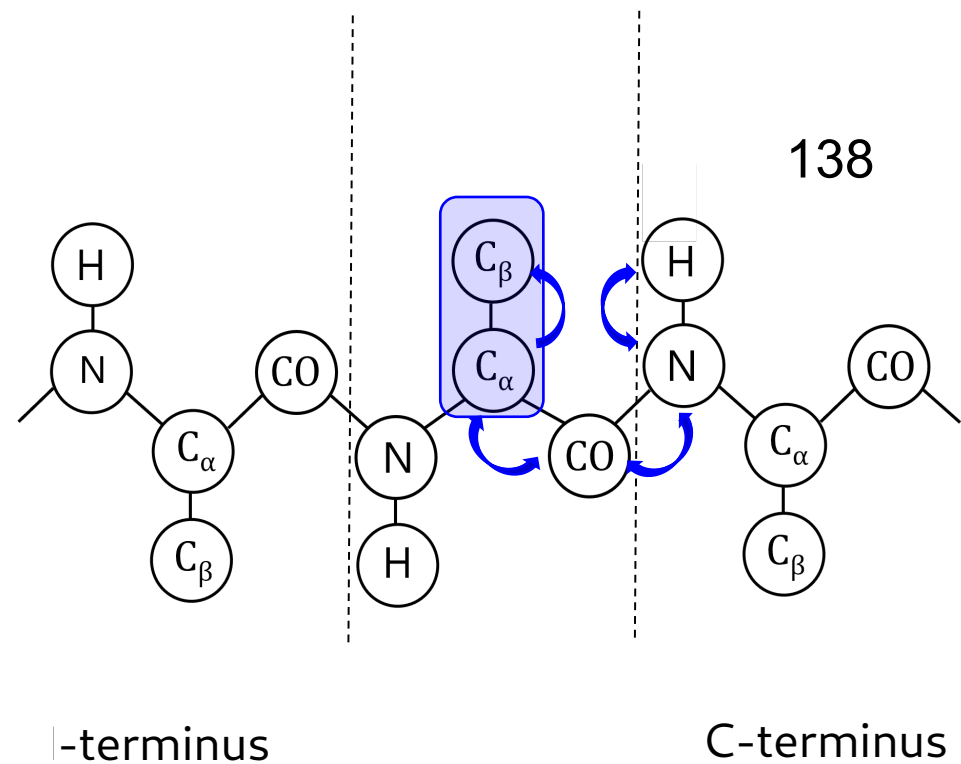
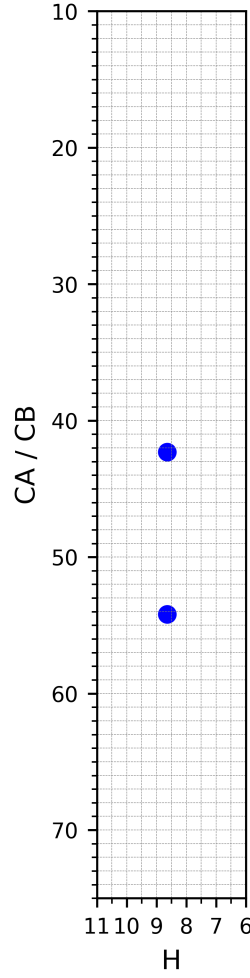
HN(CO)CACB (138)



HNCACB (138)



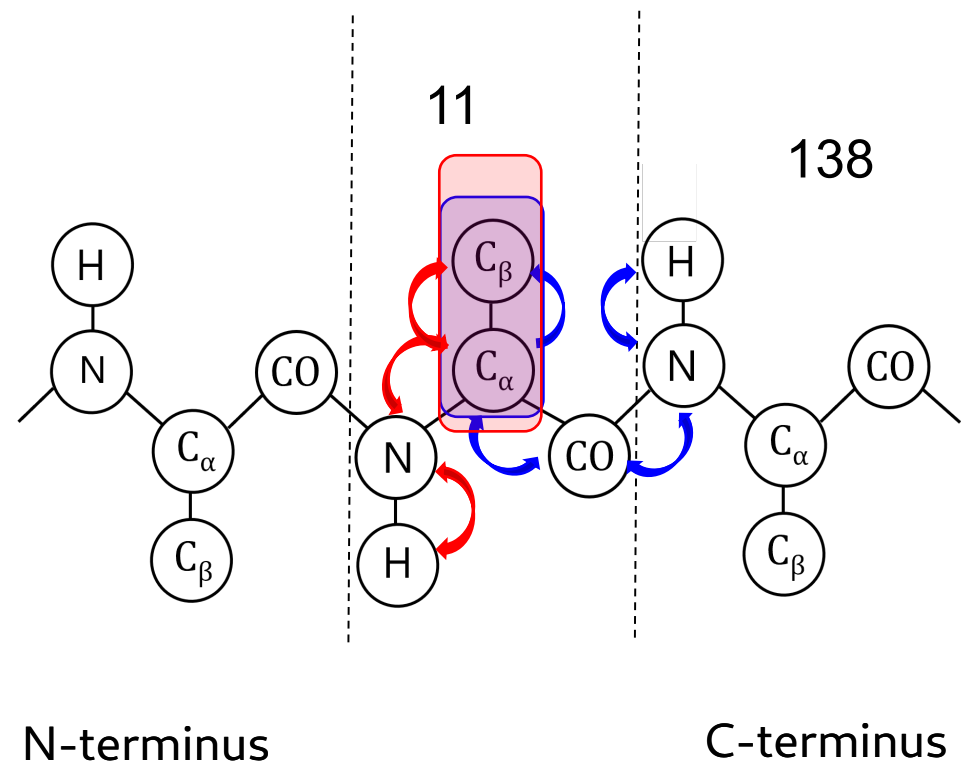
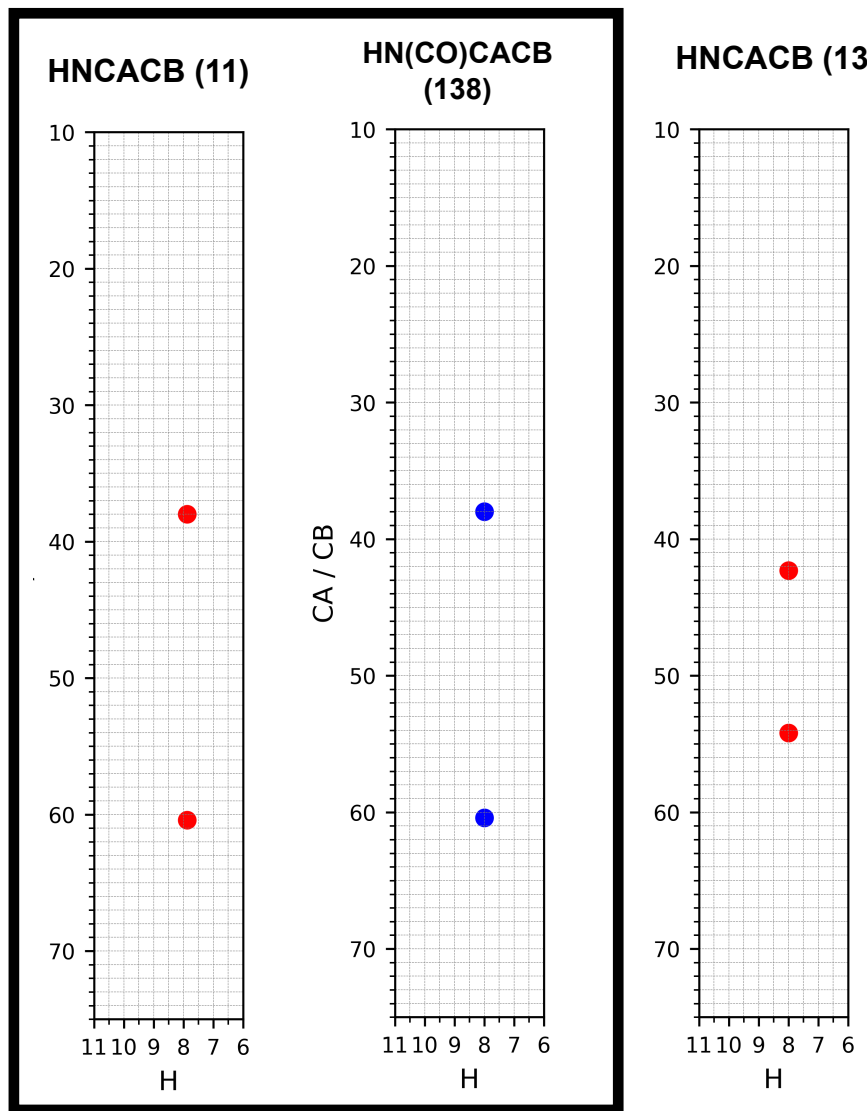
HN(CO)CACB (8)



HNCACB: $C\alpha$ a $C\beta$ současné AMK

HN(CO)CACB: $C\alpha$ a $C\beta$ předchozí AMK

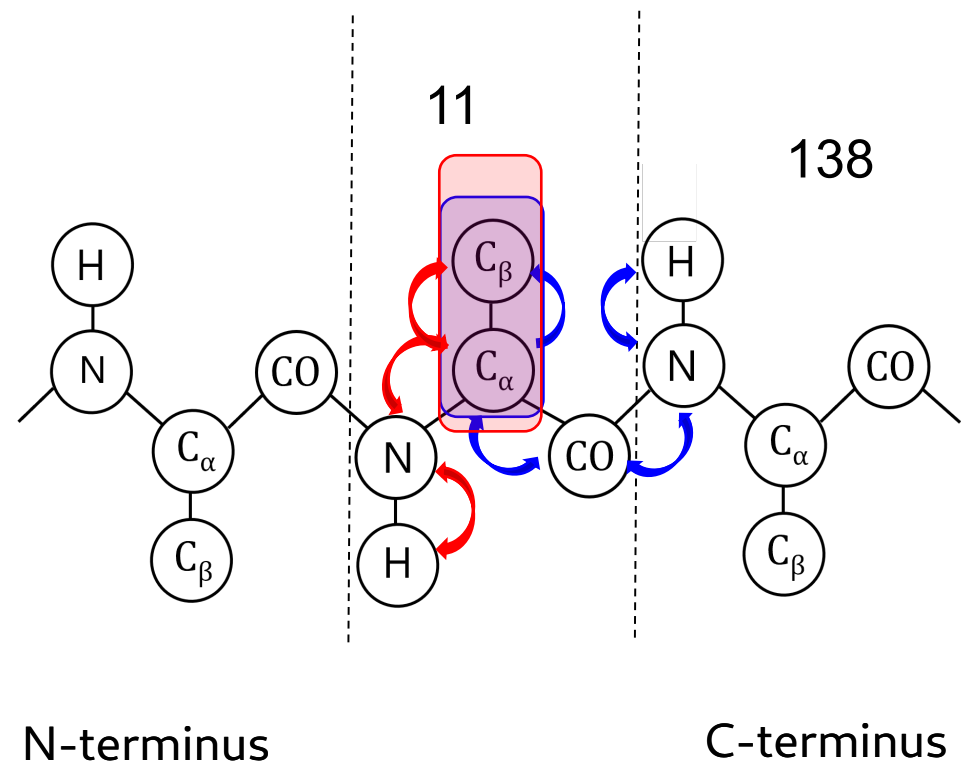
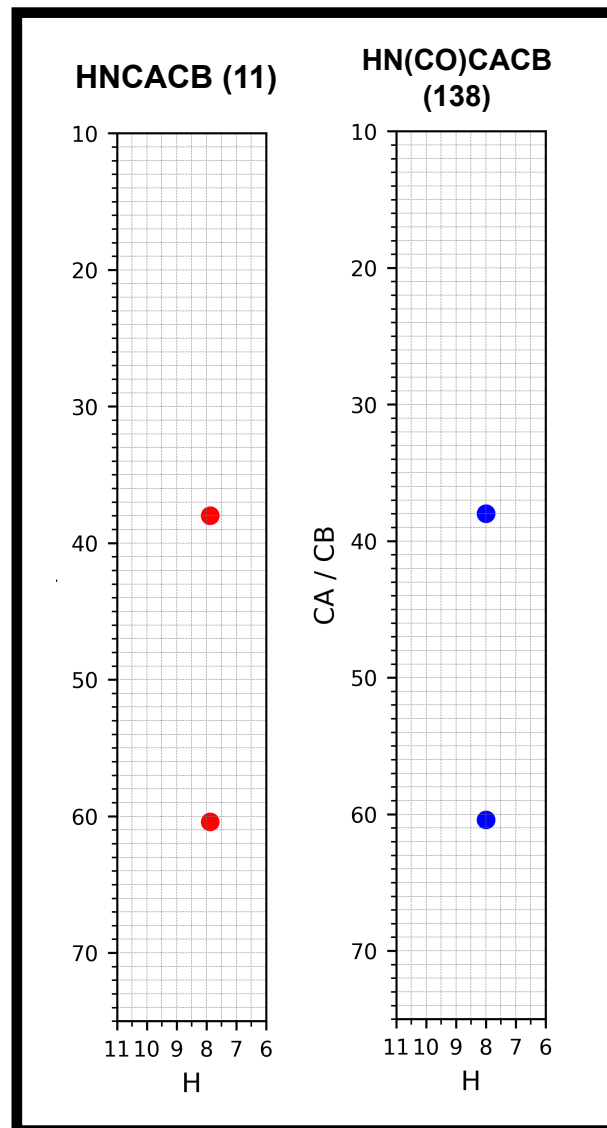
Postup při přiřazování - 11-138-8



HNCACB: C α a C β současné AMK

HN(CO)CACB: C α a C β předchozí AMK

Postup při přiřazování - 11-138-8

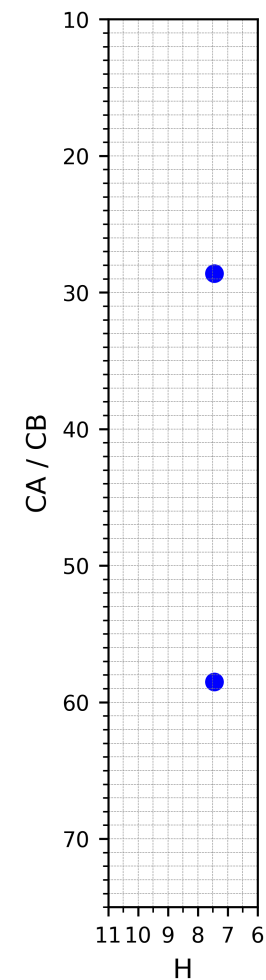


HNCACB: C α a C β současné AMK

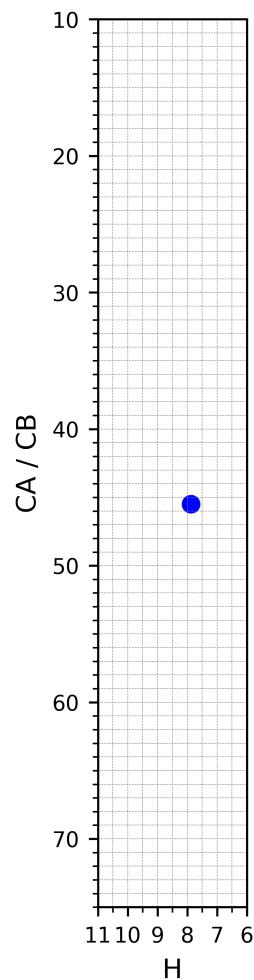
HN(CO)CACB: C α a C β předchozí AMK

Postup při přiřazování - 11-138-8

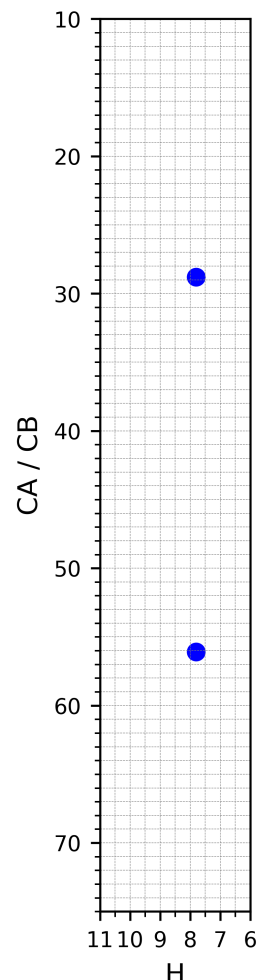
HN(CO)CACB (13)



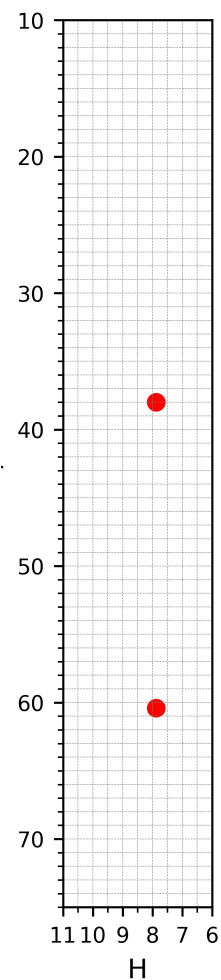
HN(CO)CACB (11)



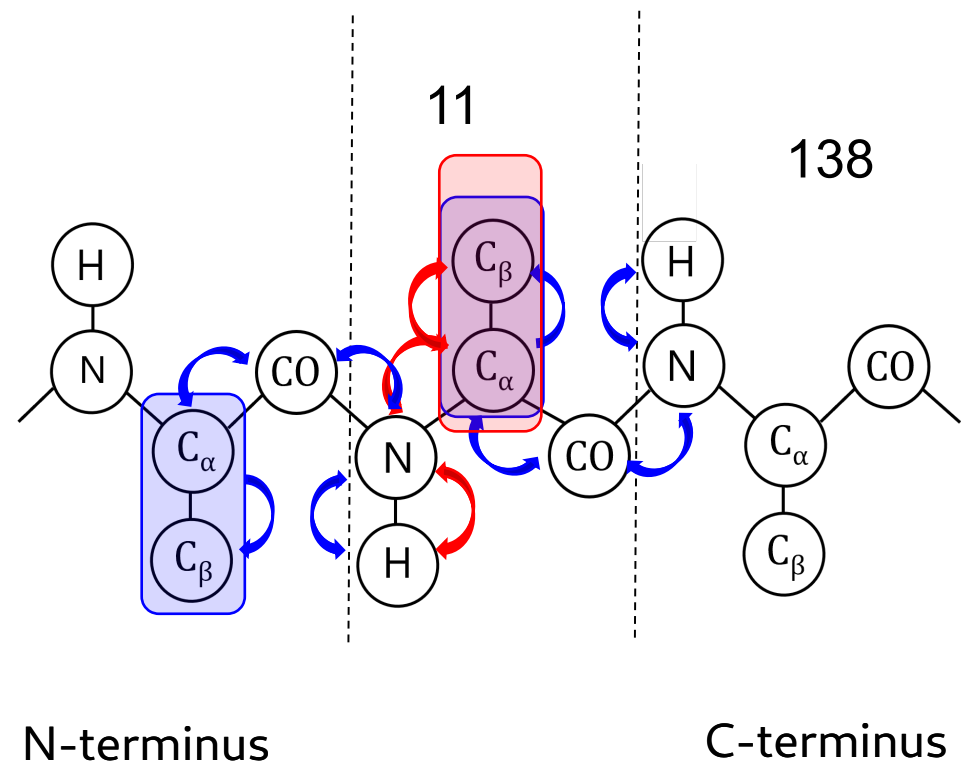
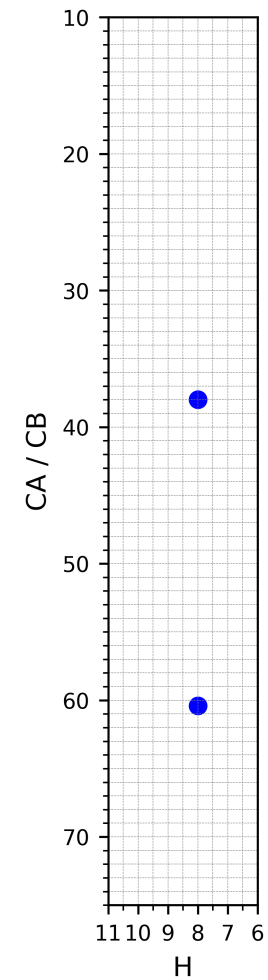
HN(CO)CACB (3)



HNCACB (11)



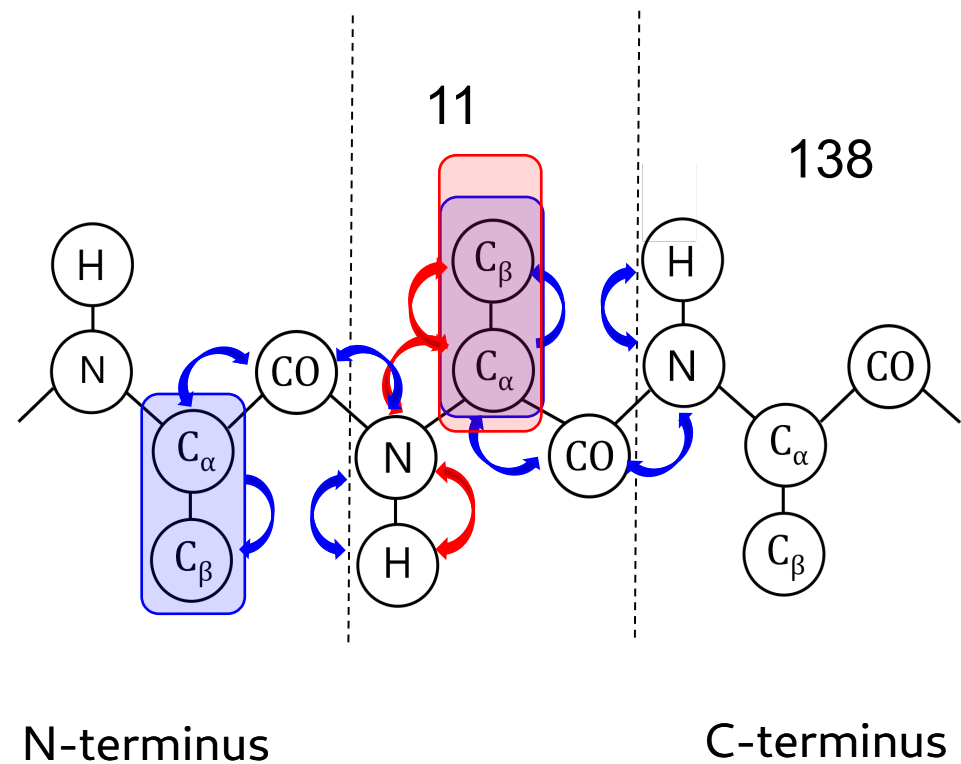
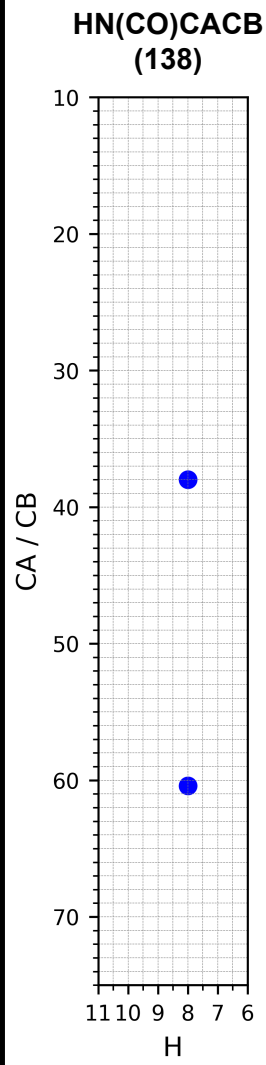
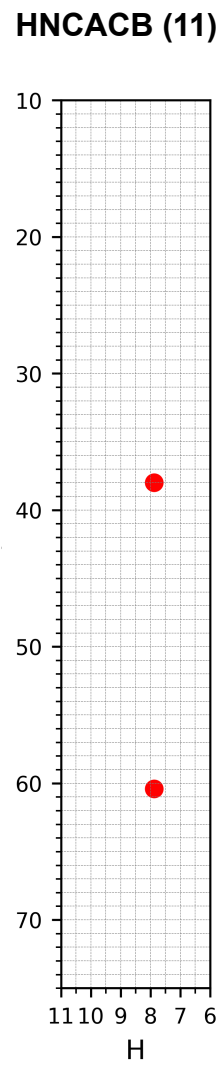
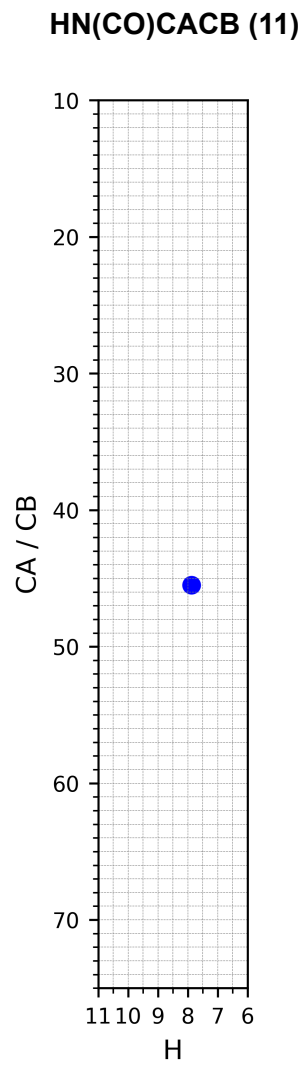
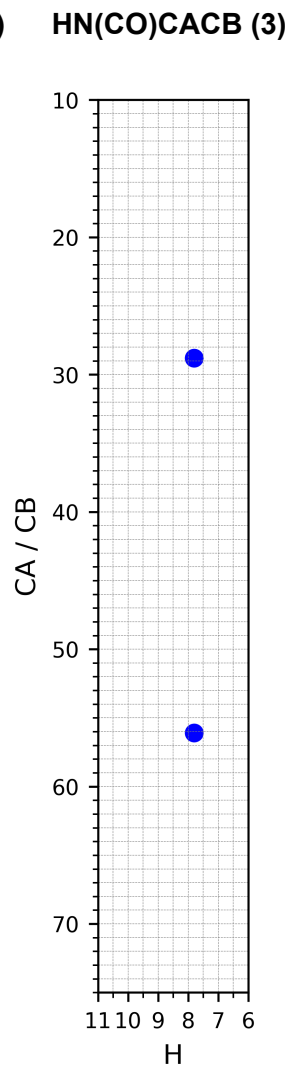
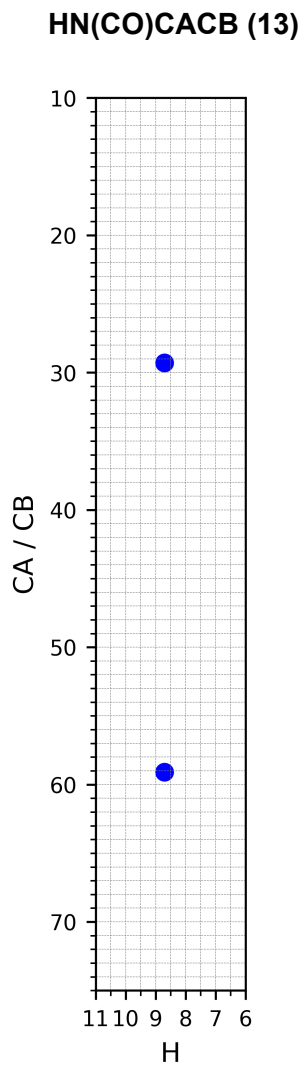
HN(CO)CACB (138)



HNCACB: C α a C β současné AMK

HN(CO)CACB: C α a C β předchozí AMK

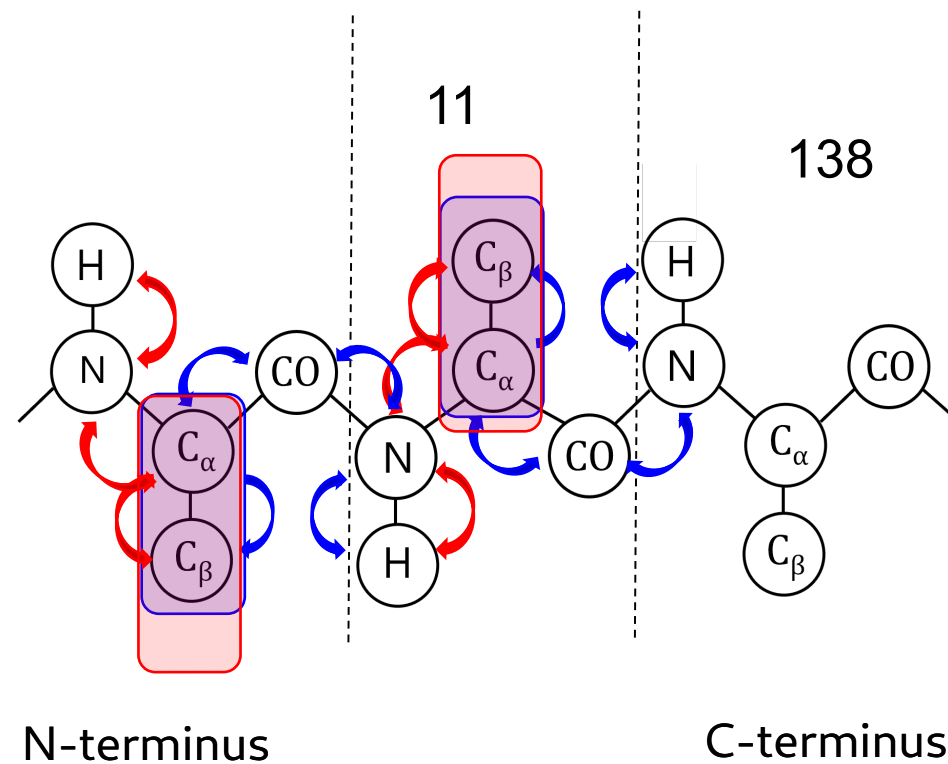
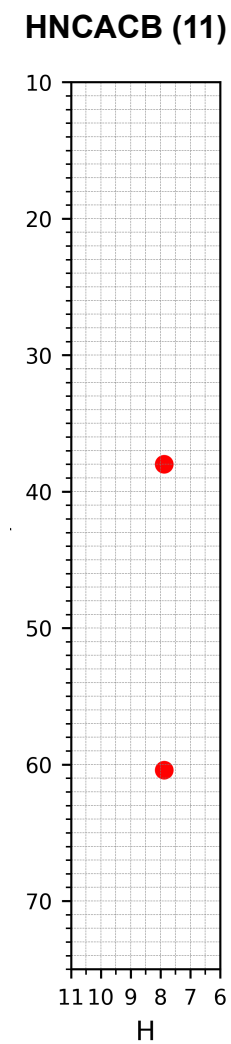
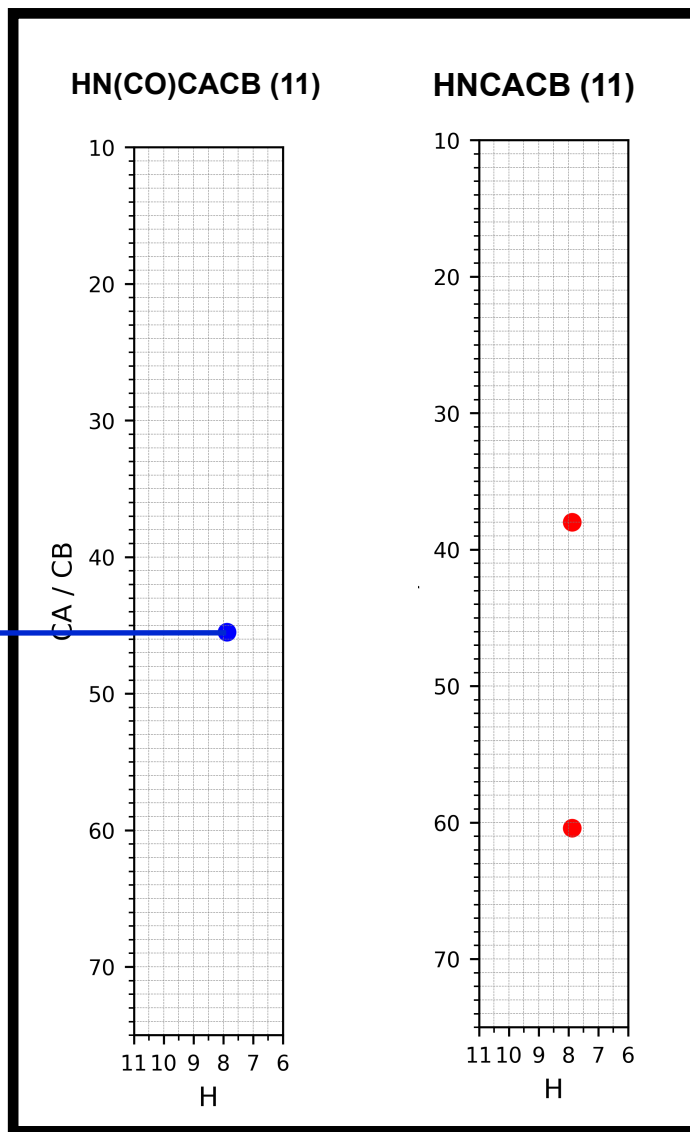
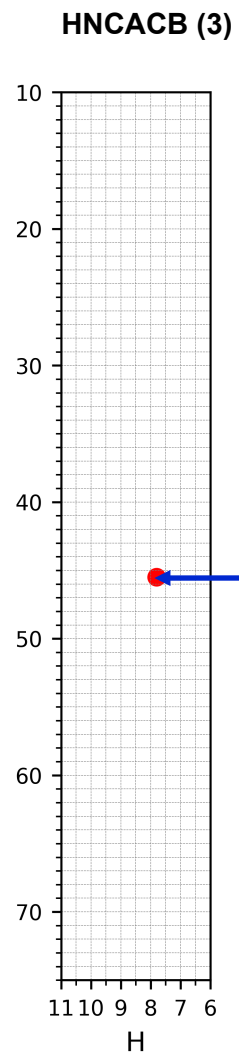
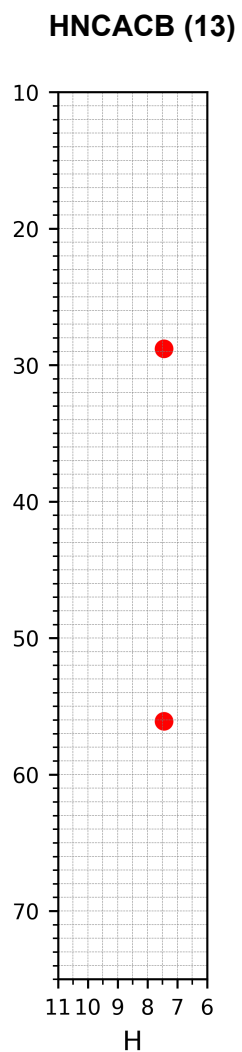
Postup při přiřazování - 11-138-8



HNCACB: C α a C β současné AMK

HN(CO)CACB: C α a C β předchozí AMK

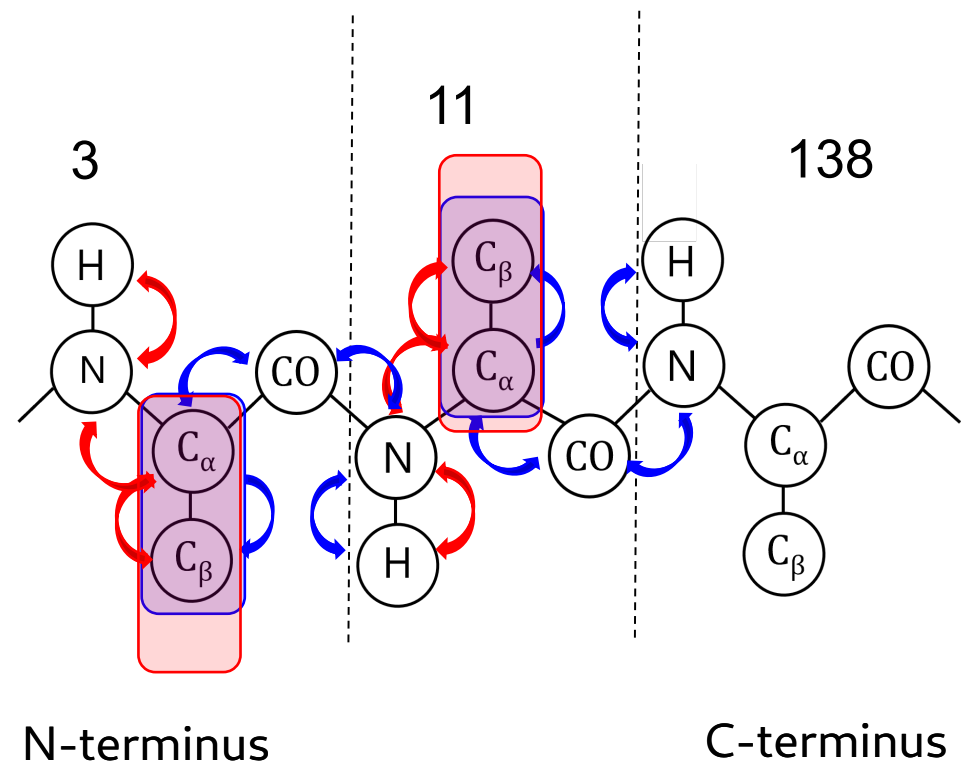
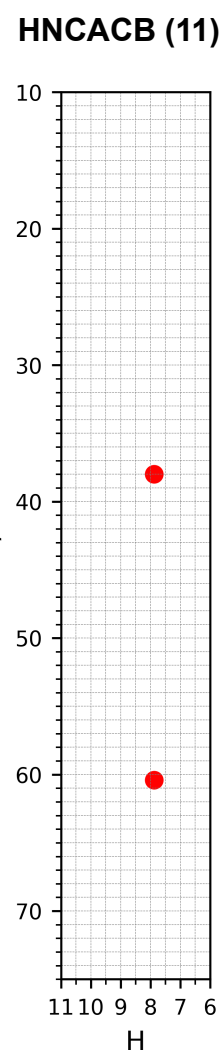
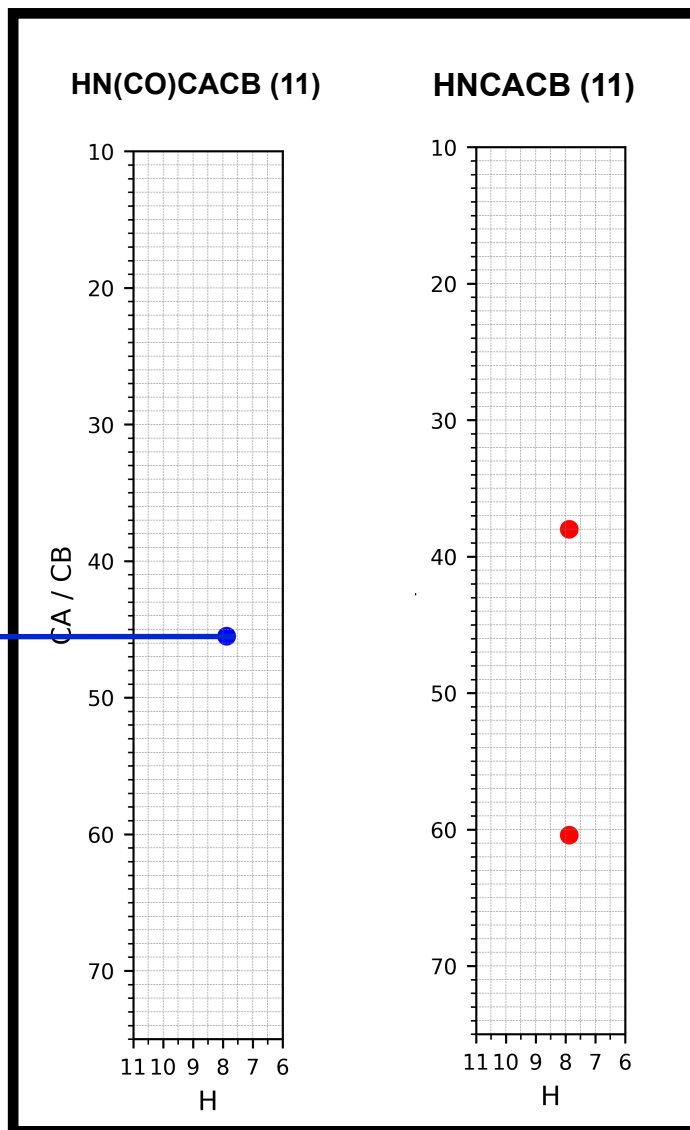
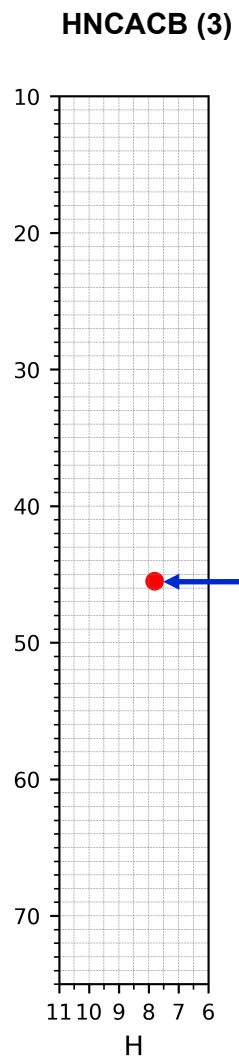
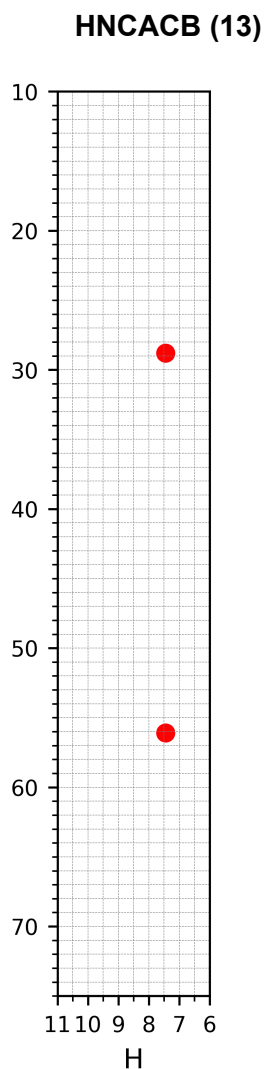
Postup při přiřazování – 3-11-138-8



HNCACB: C α a C β současné AMK

HN(CO)CACB: C α a C β předchozí AMK

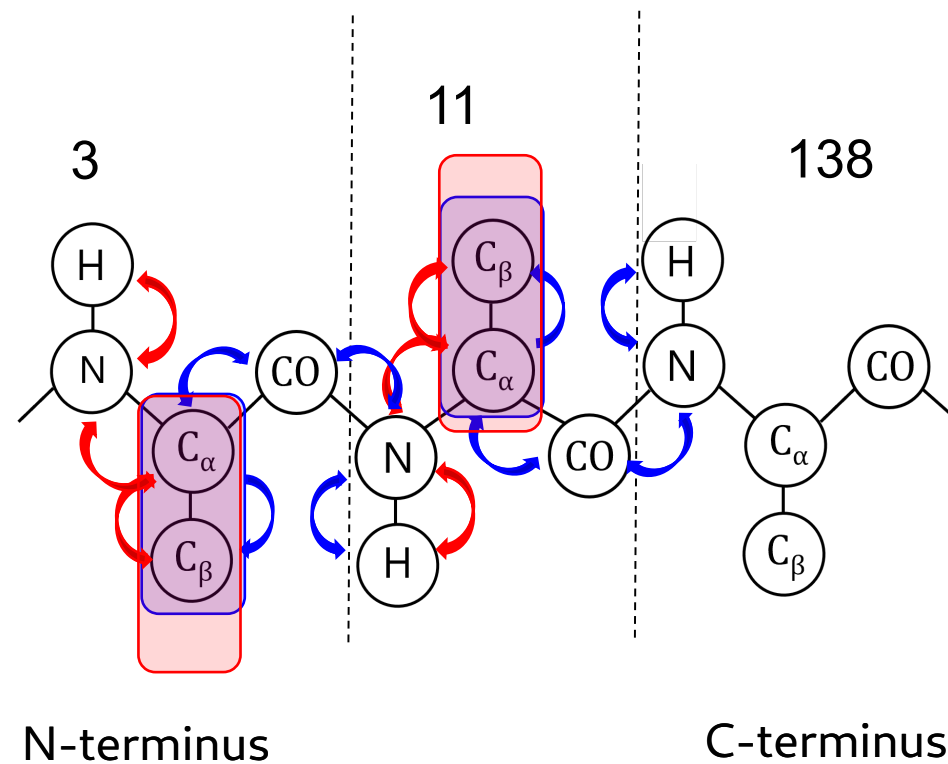
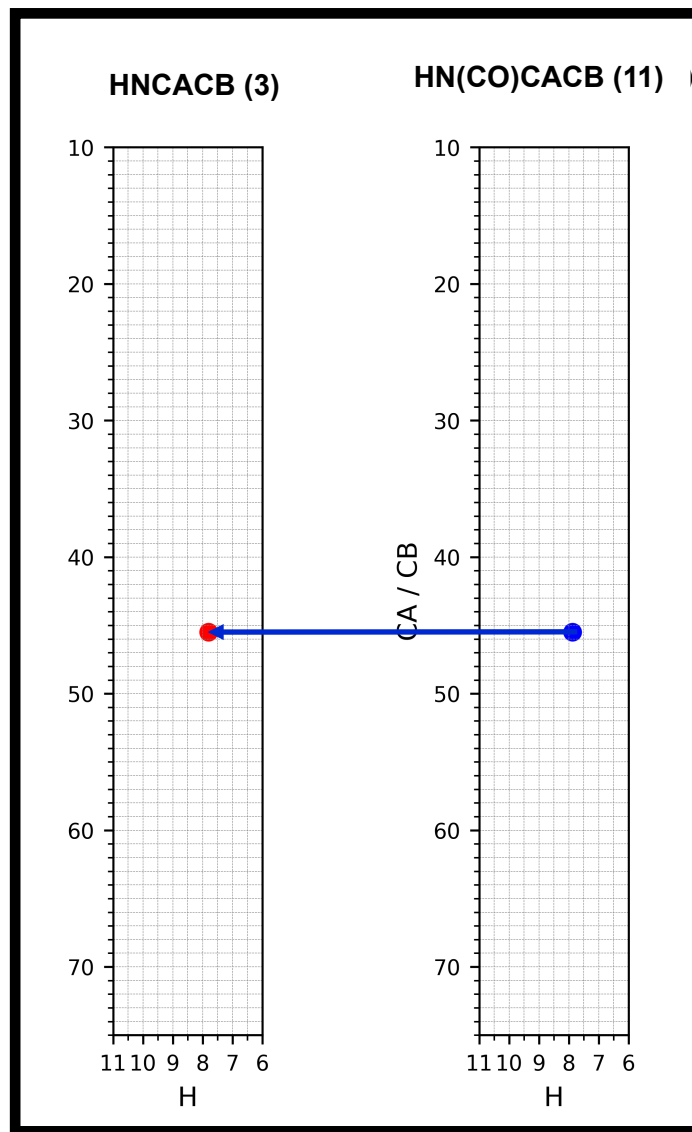
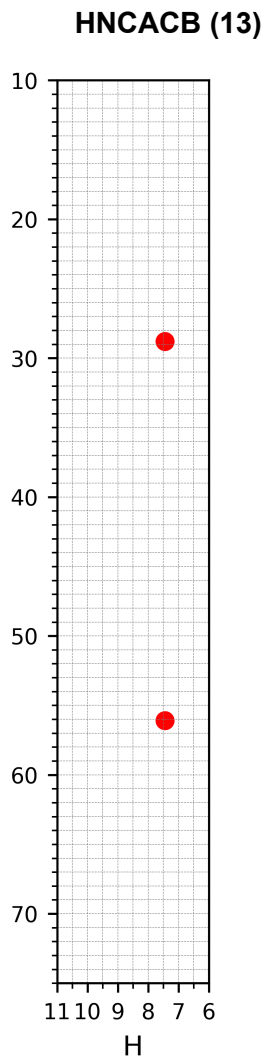
Postup při přiřazování - 3-11-138-8



HNCACB: C_{α} a C_{β} současné AMK

HN(CO)CACB: C_{α} a C_{β} předchozí AMK

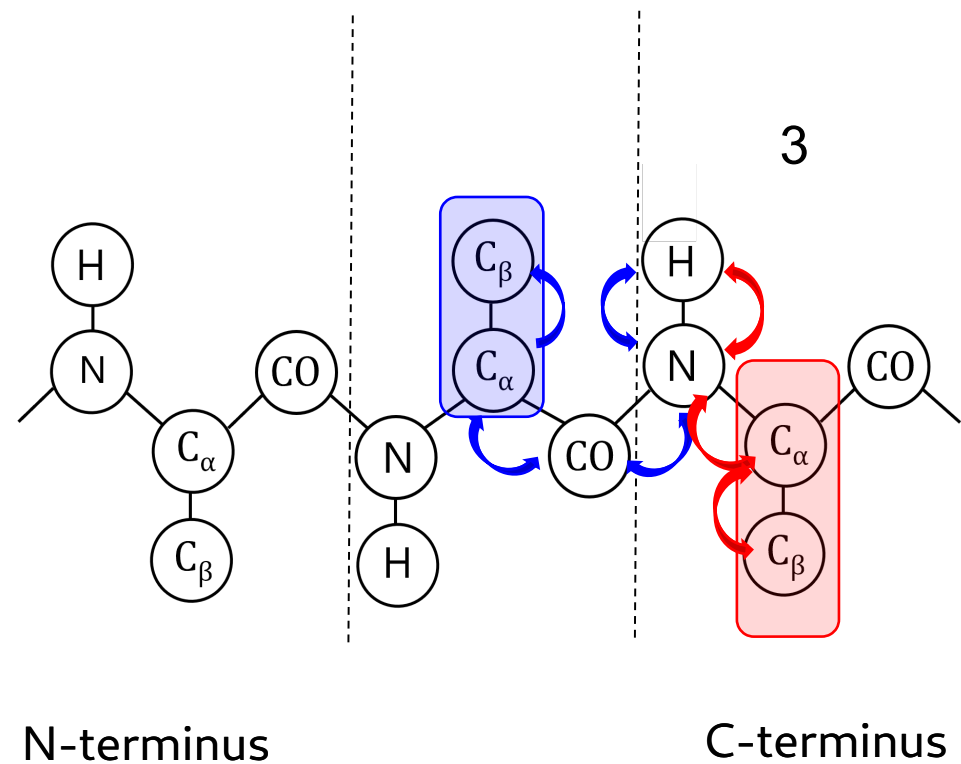
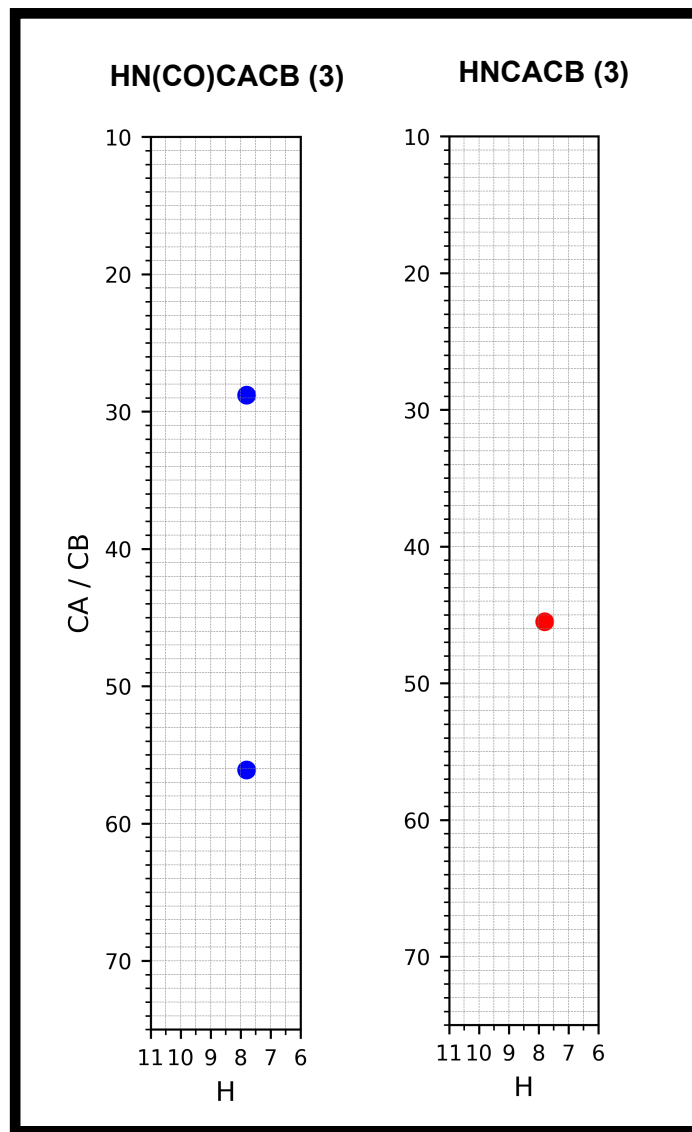
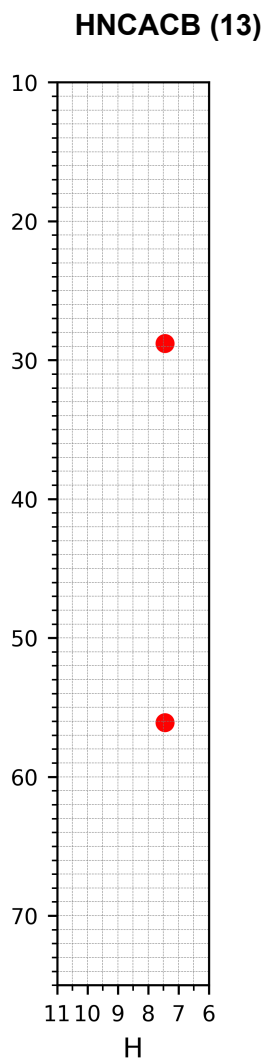
Postup při přiřazování - 3-11-138-8



HNCACB: C α a C β současné AMK

HN(CO)CACB: C α a C β předchozí AMK

Postup při přiřazování - 3-11-138-8

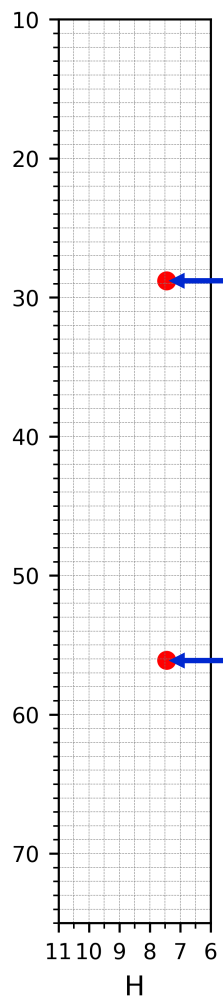


HNCACB: C α a C β současné AMK

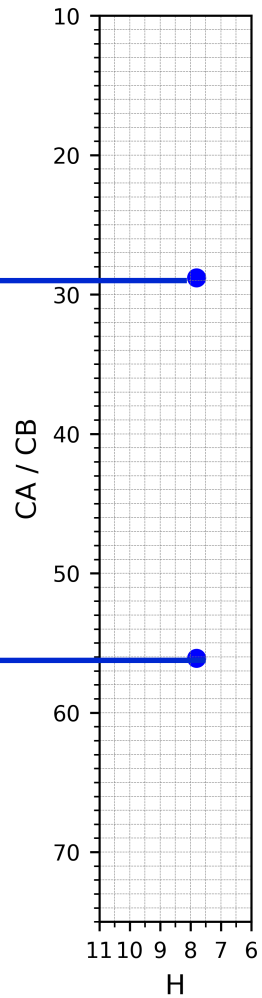
HN(CO)CACB: C α a C β předchozí AMK

Postup při přiřazování – 13-3-11-138-8

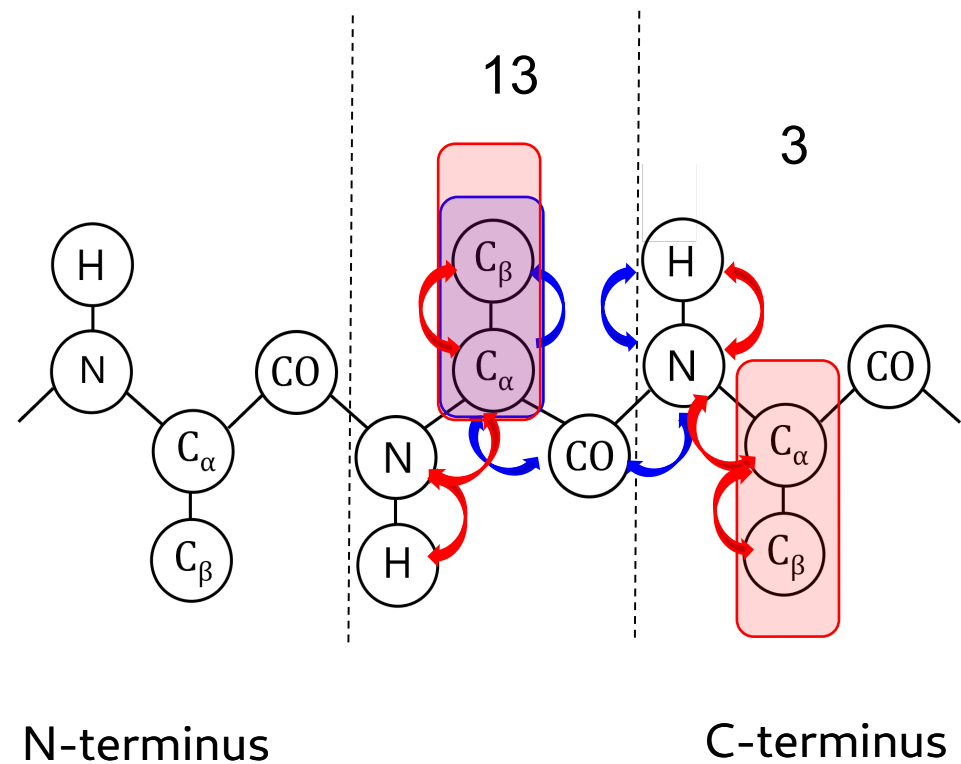
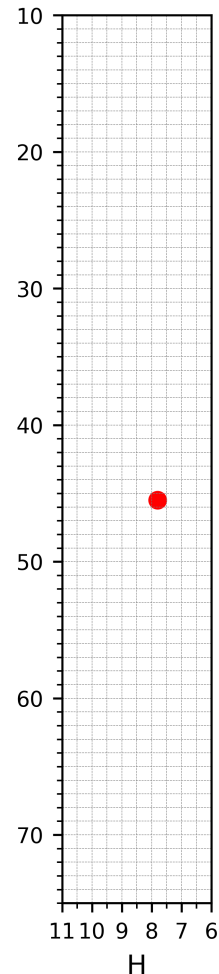
HNCACB (13)



HN(CO)CACB (3)



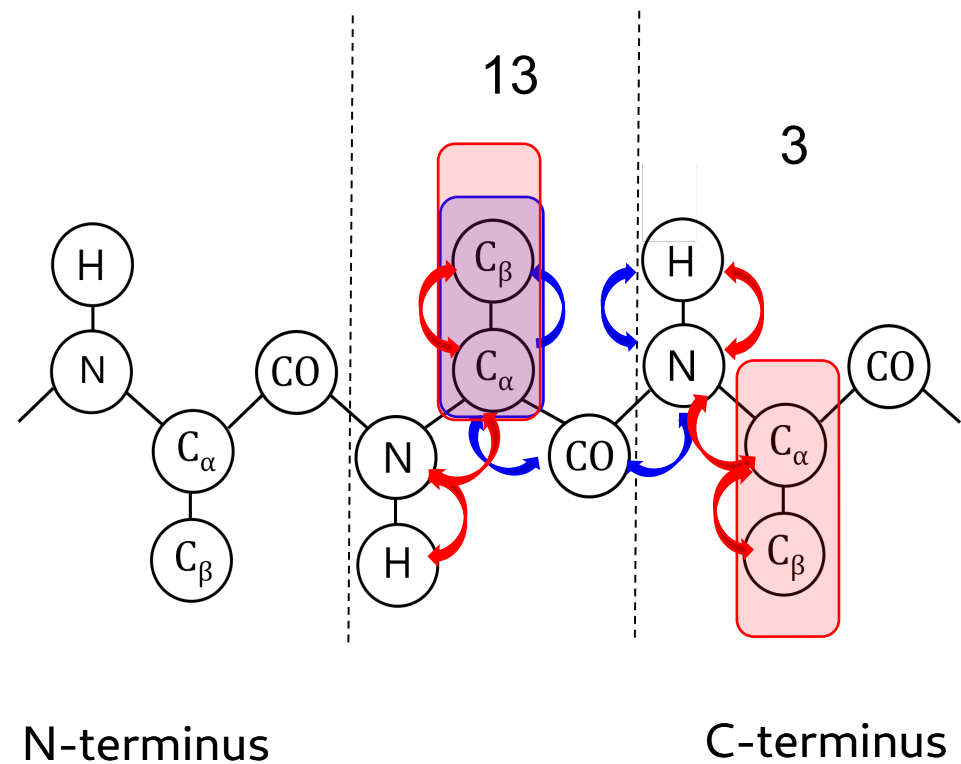
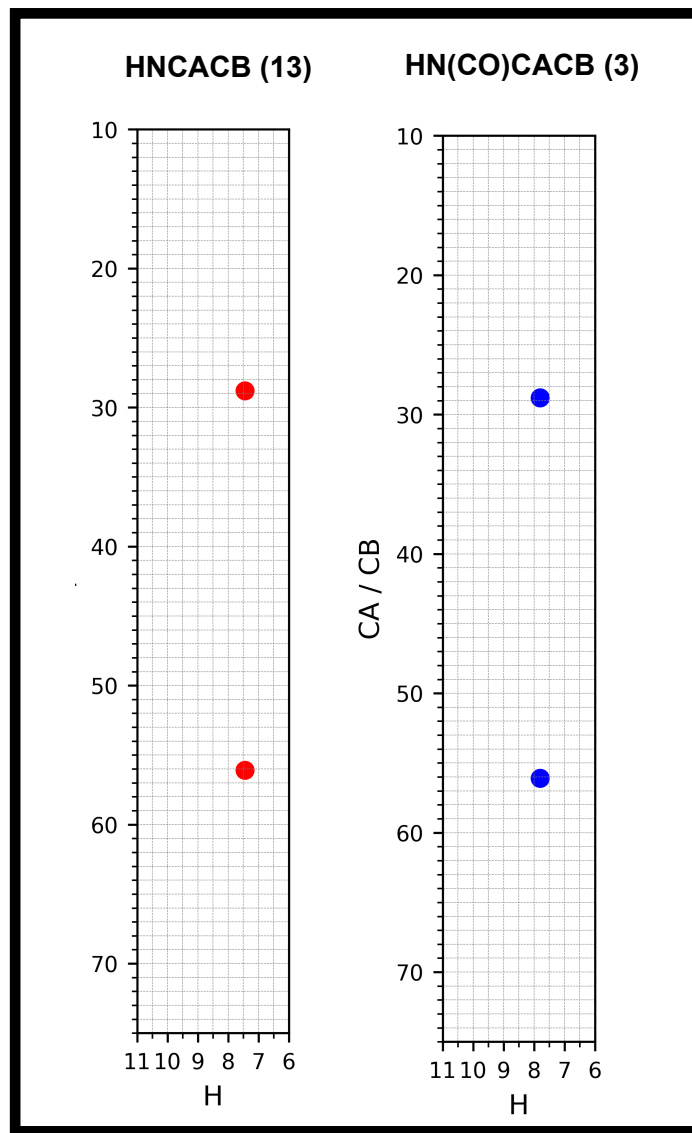
HNCACB (3)



HNCACB: C_α a C_β současné AMK

HN(CO)CACB: C_α a C_β předchozí AMK

Postup při přiřazování - 13-3-11-138-8

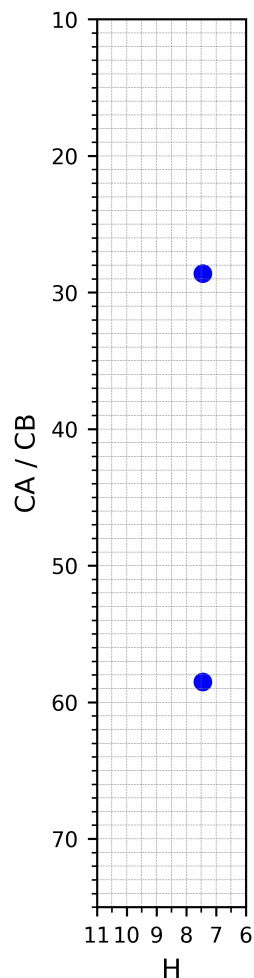


HNCACB: C α a C β současné AMK

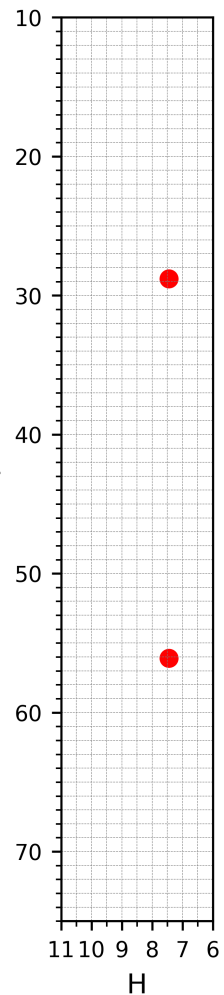
HN(CO)CACB: C α a C β předchozí AMK

Postup při přiřazování - 13-3-11-138-8

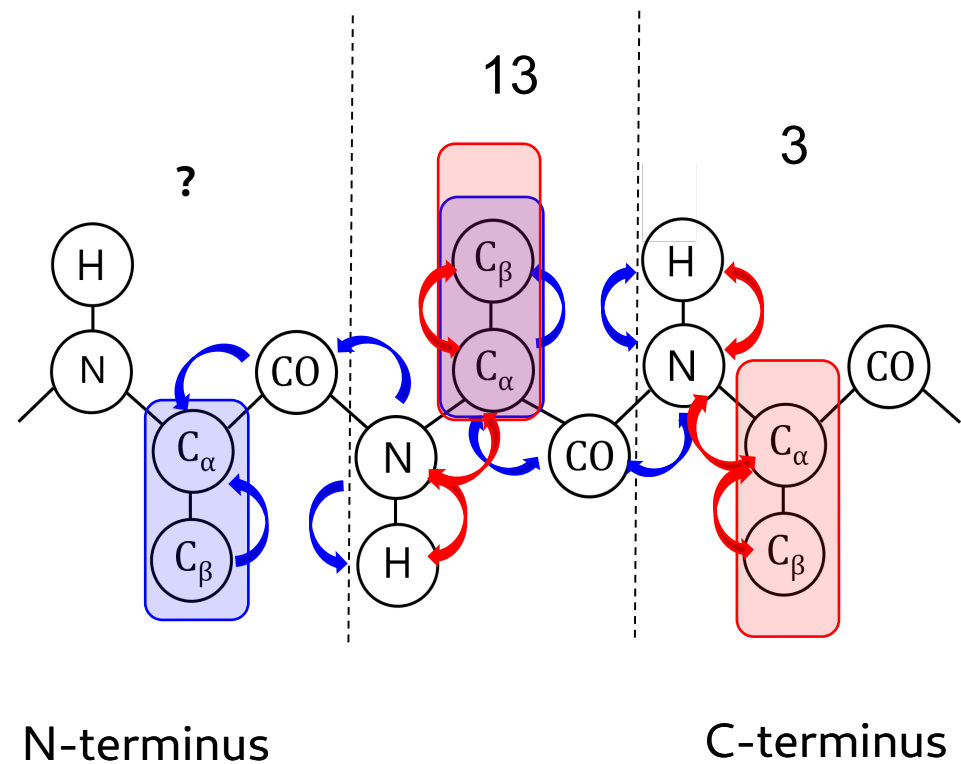
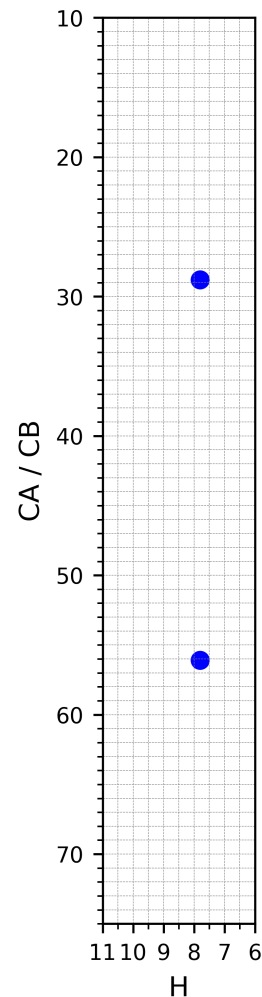
HN(CO)CACB (13)



HNCACB (13)



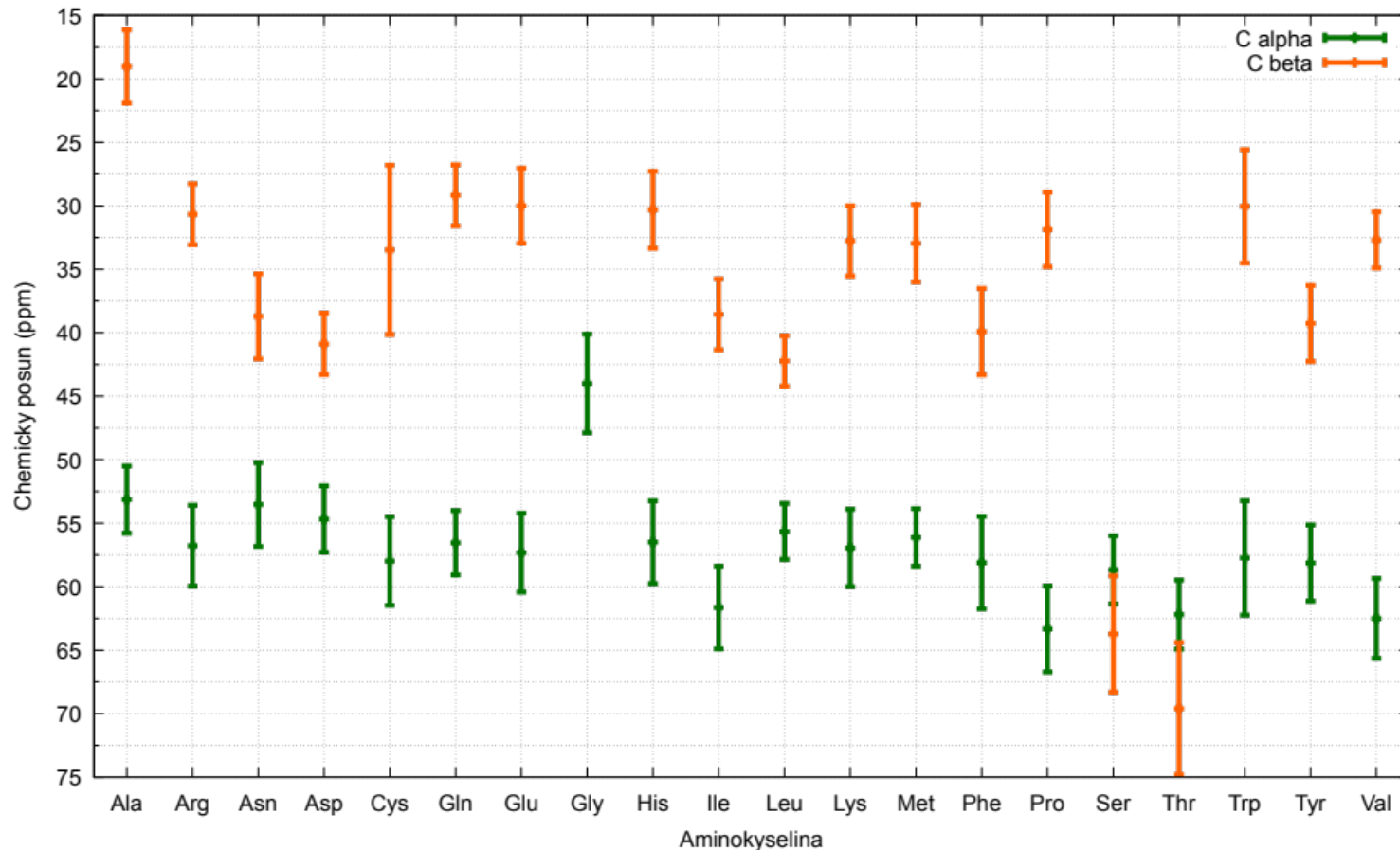
HN(CO)CACB (3)



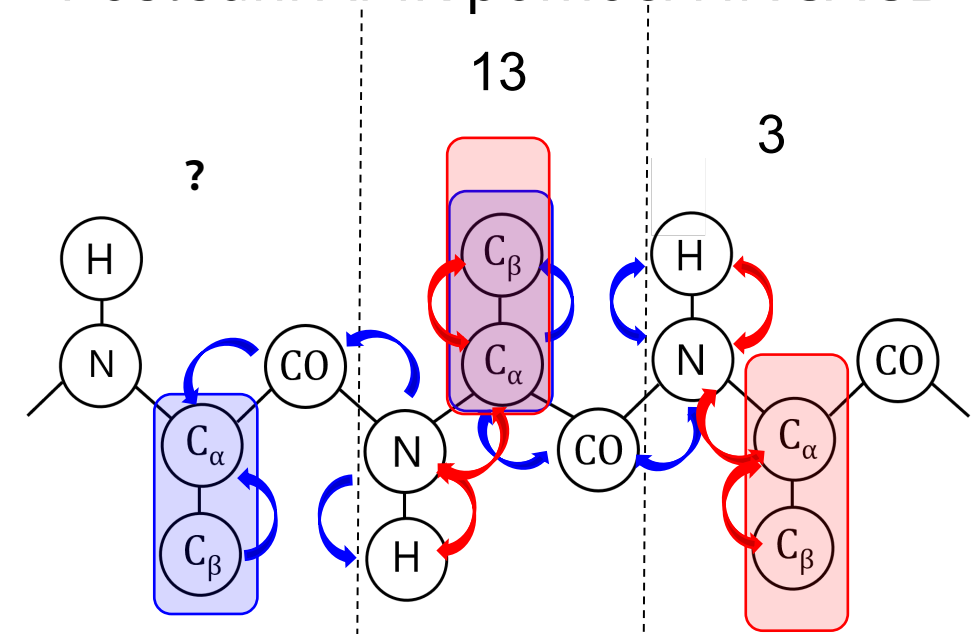
HNCACB: C α a C β současné AMK

HN(CO)CACB: C α a C β předchozí AMK

Úkol č. 2 – určit o jaké aminokyseliny se jedná na základě chemického posunu CA a CB

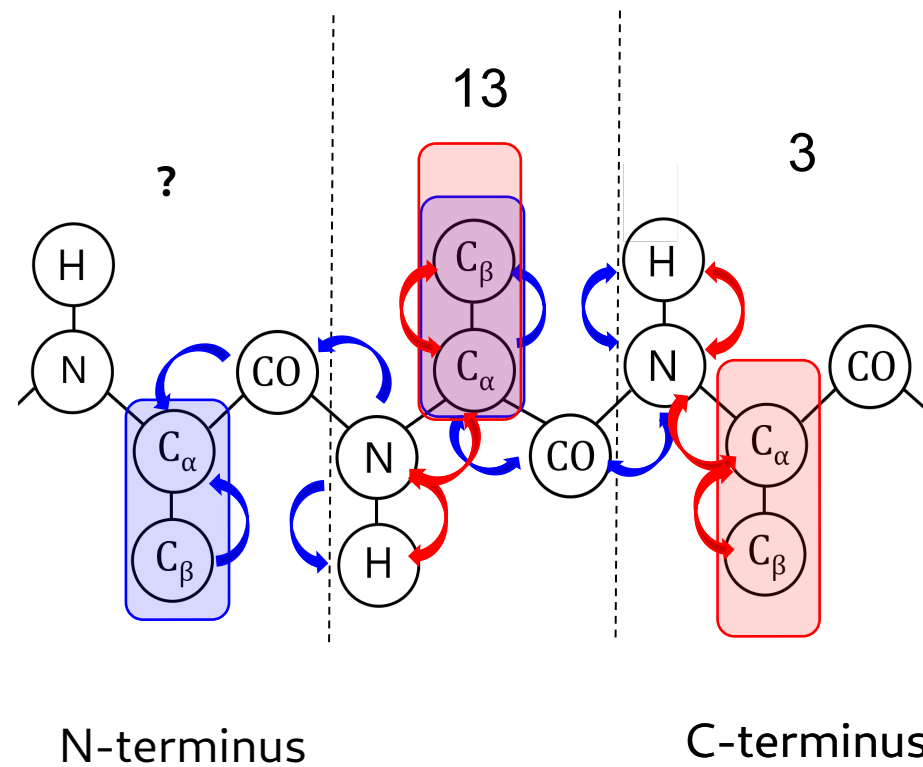
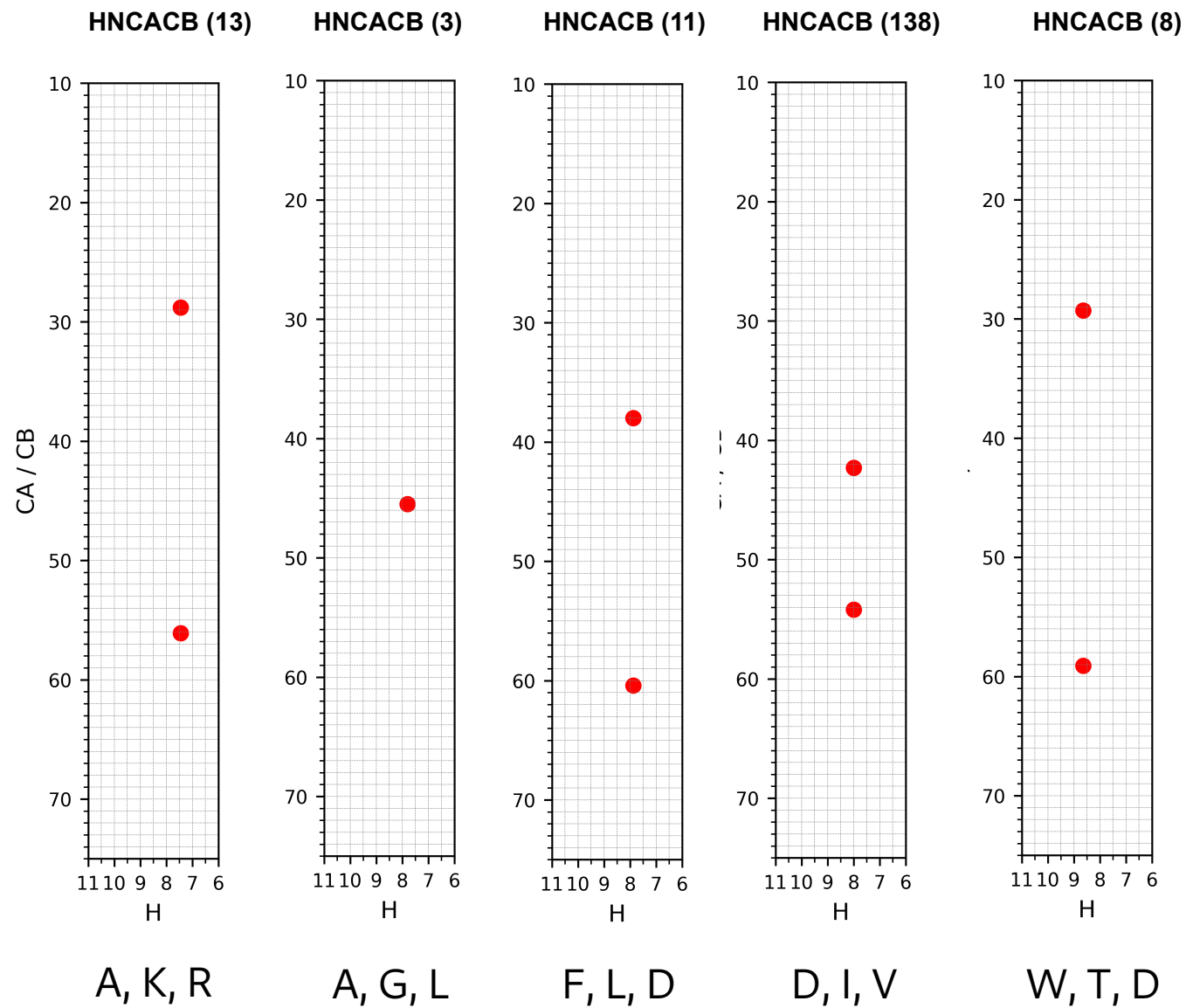


- První AMK na N-terminu – pomocí HN(CO)CACB
- Poslední AMK pomocí HNCACB

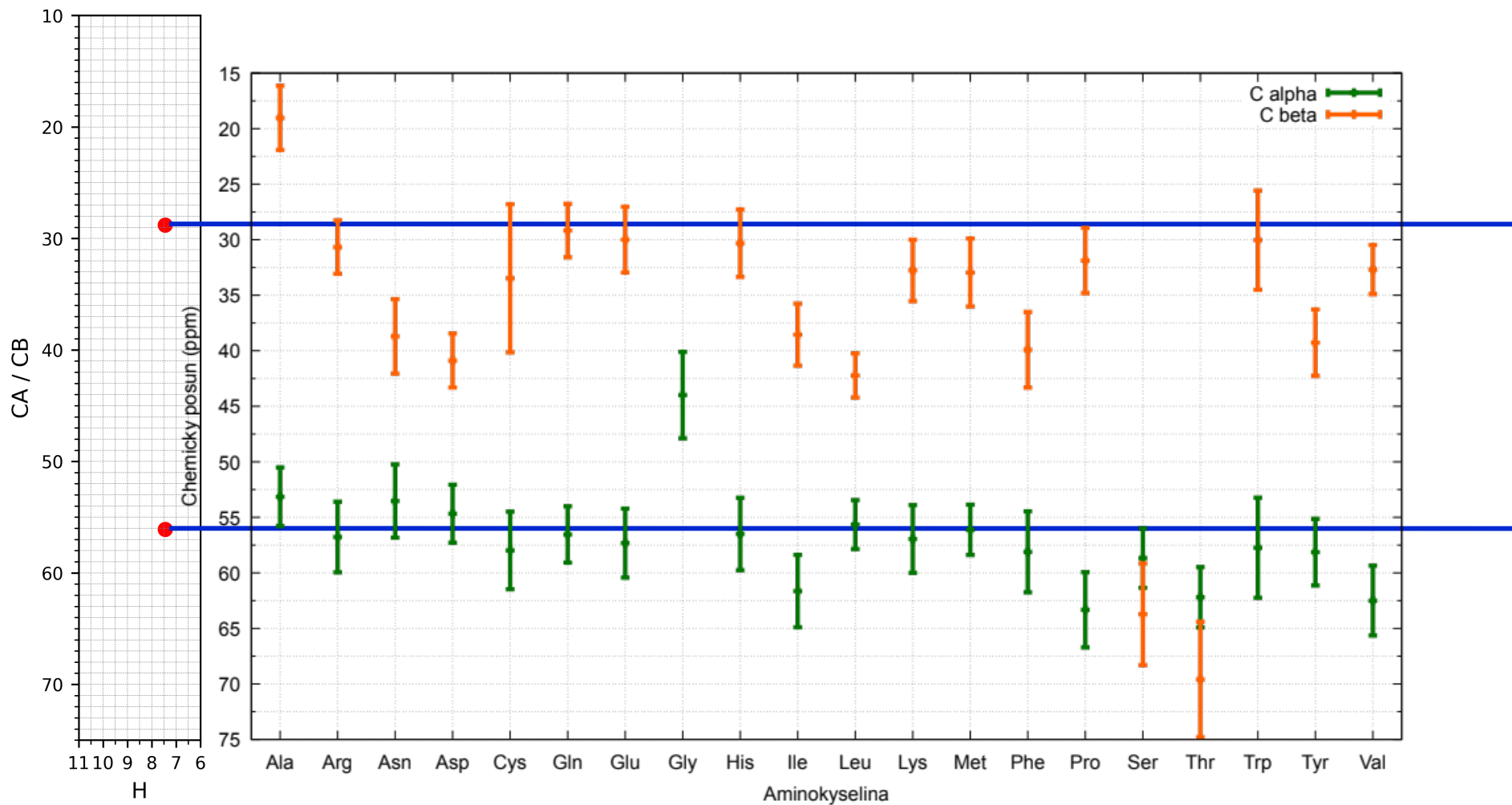


N-terminus

C-terminus

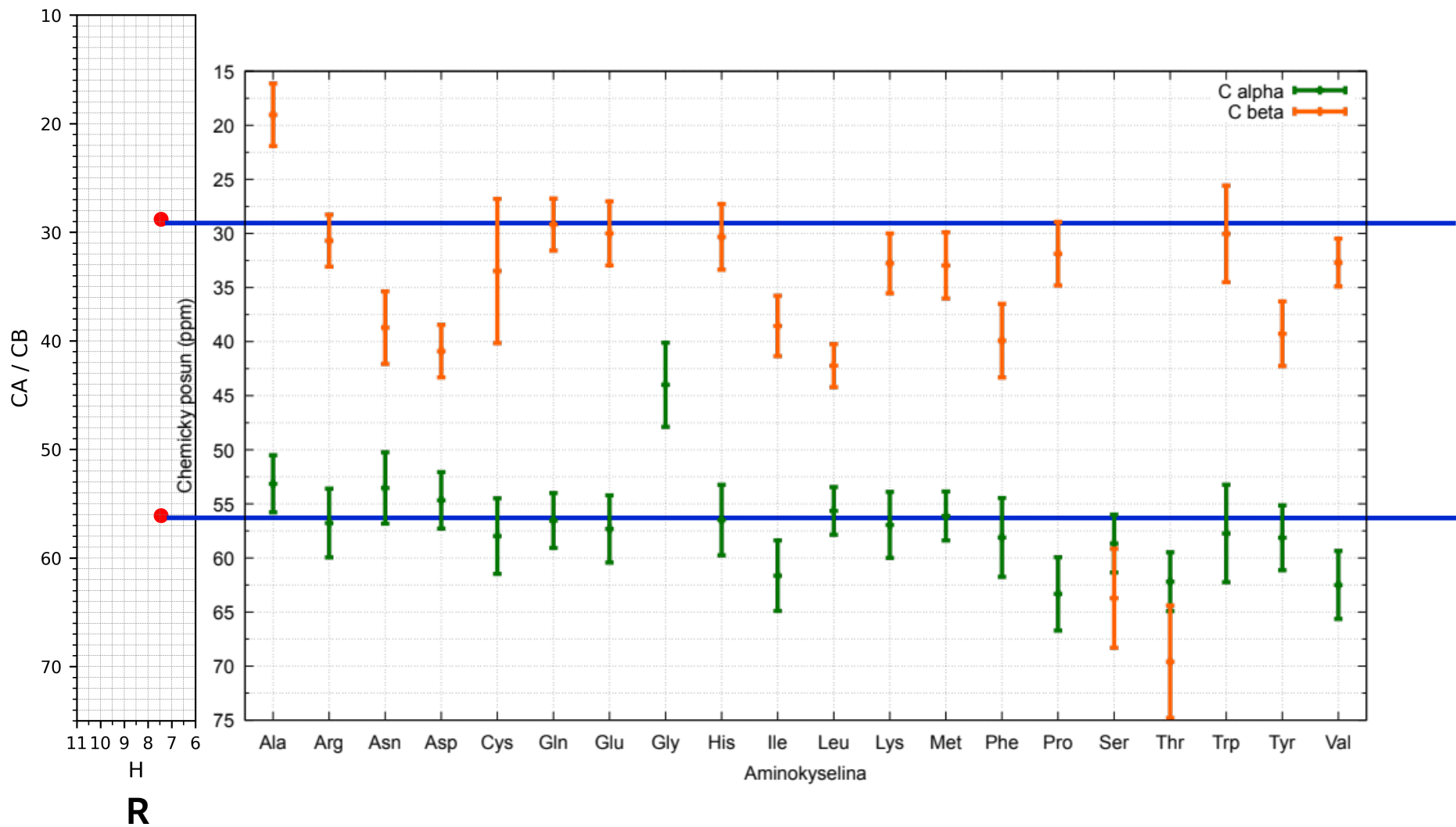


HNCACB (13)

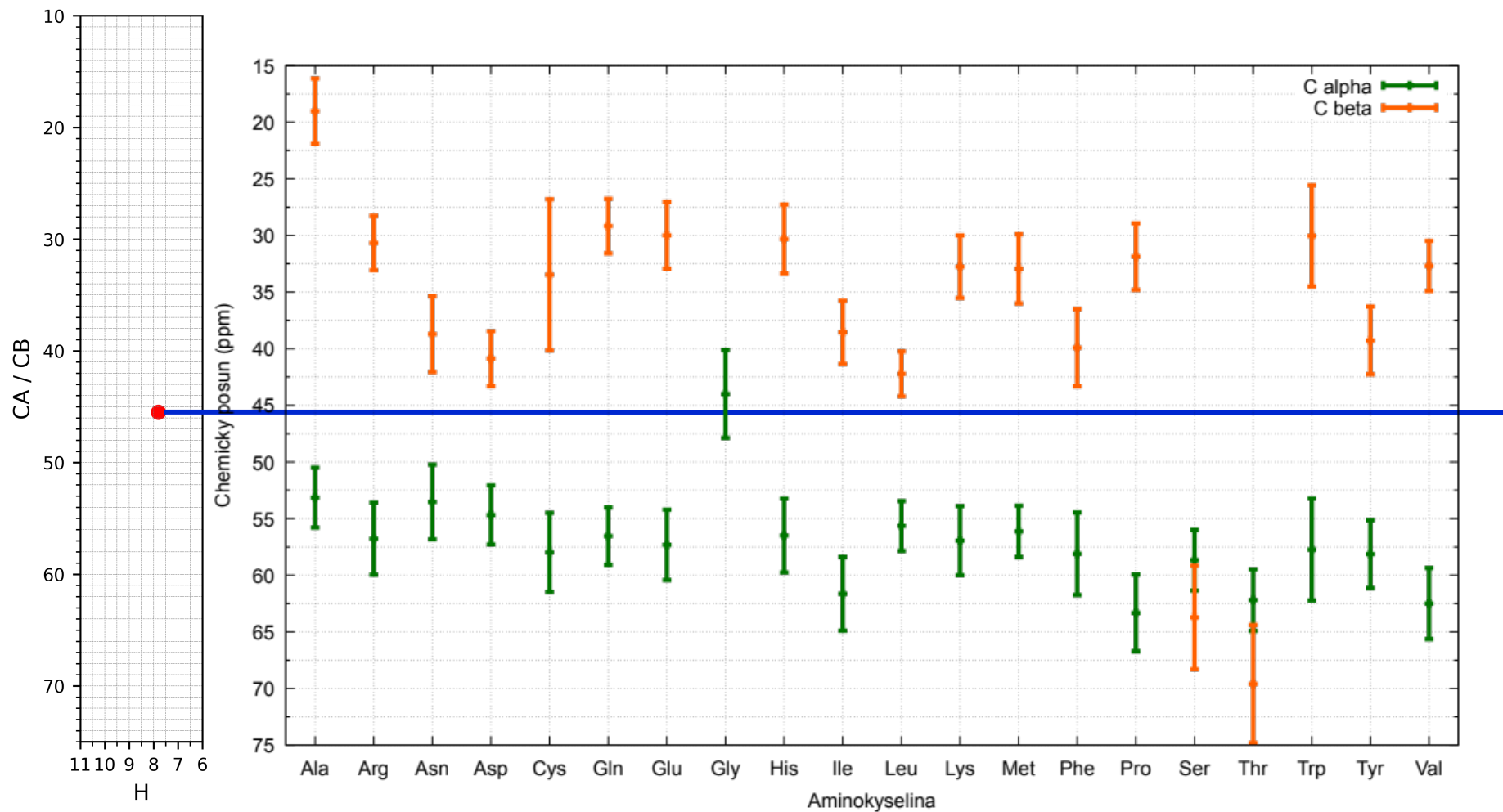


A, K, R

HNCACB (13)

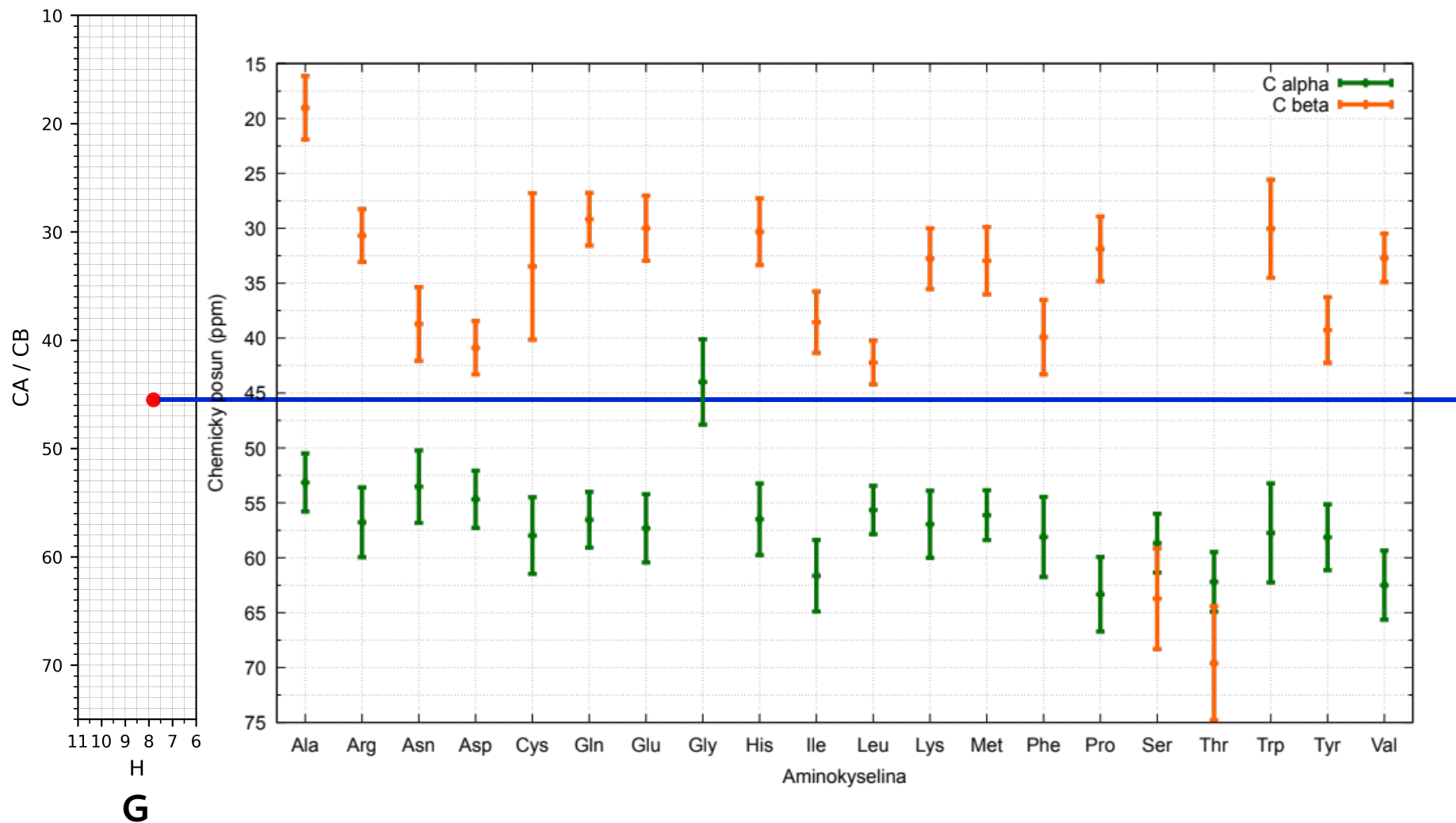


HNCACB (3)

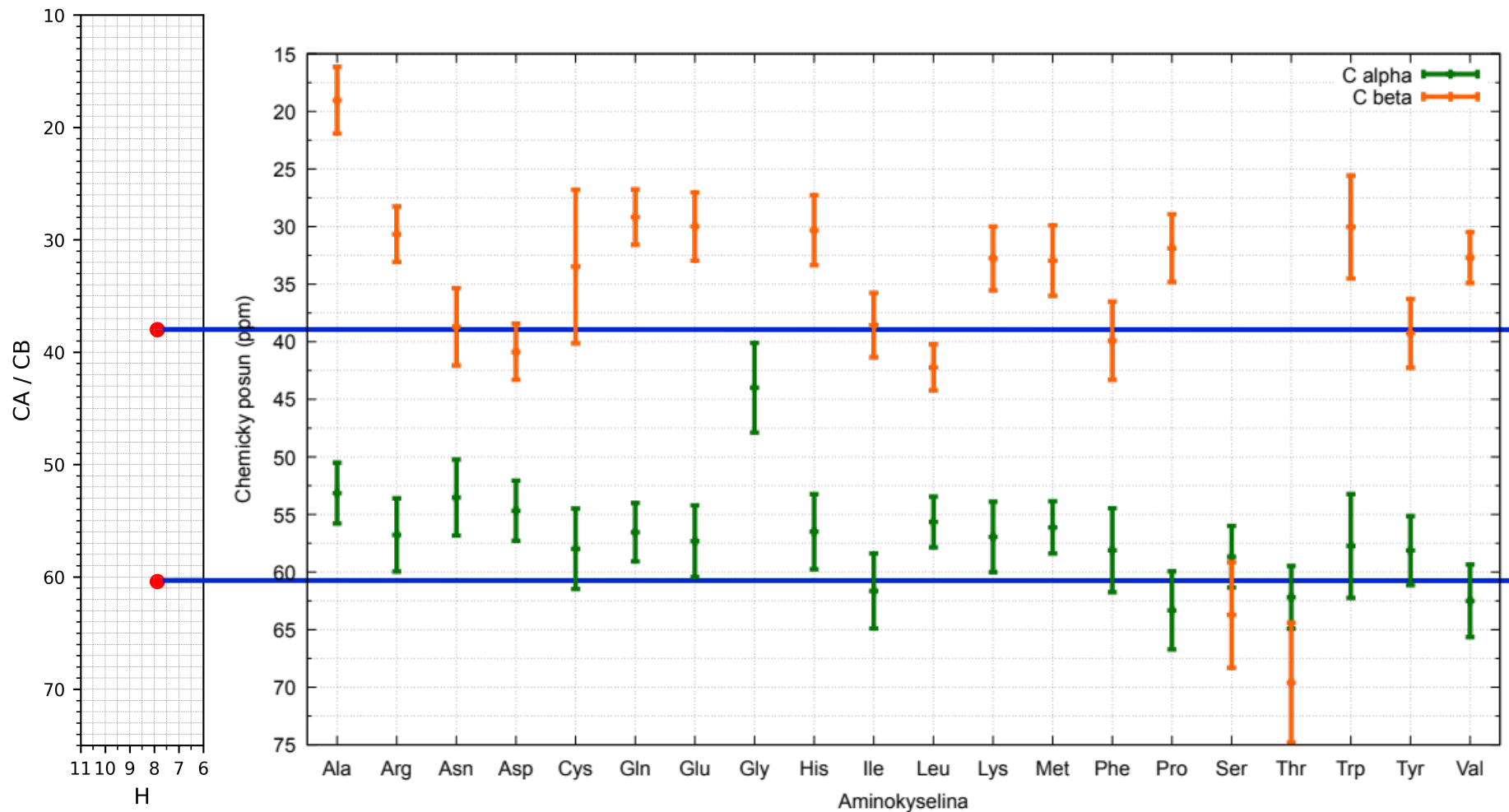


A, G, L

HNCACB (3)

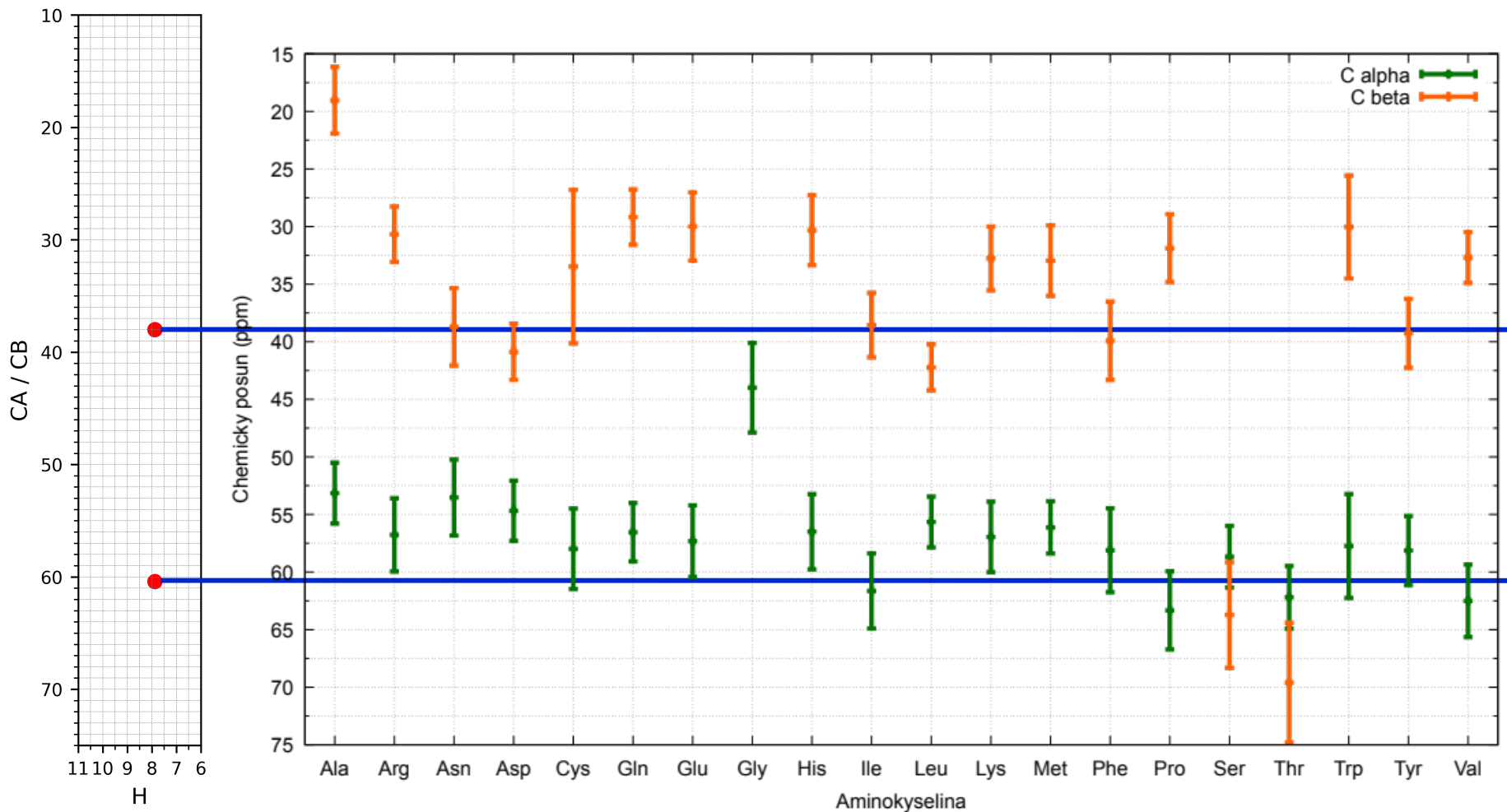


HNCACB (11)



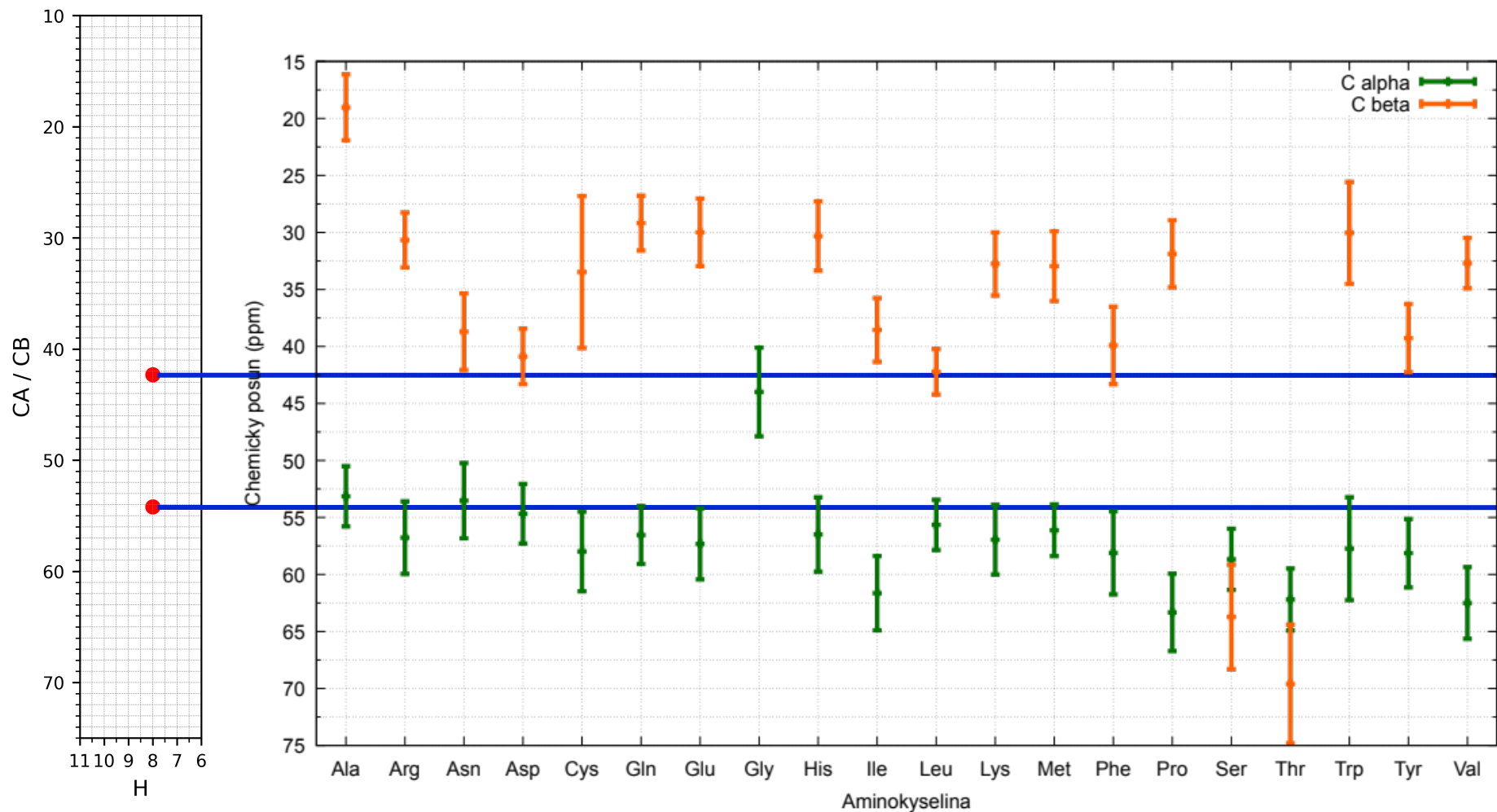
F, L, D

HNCACB (11)



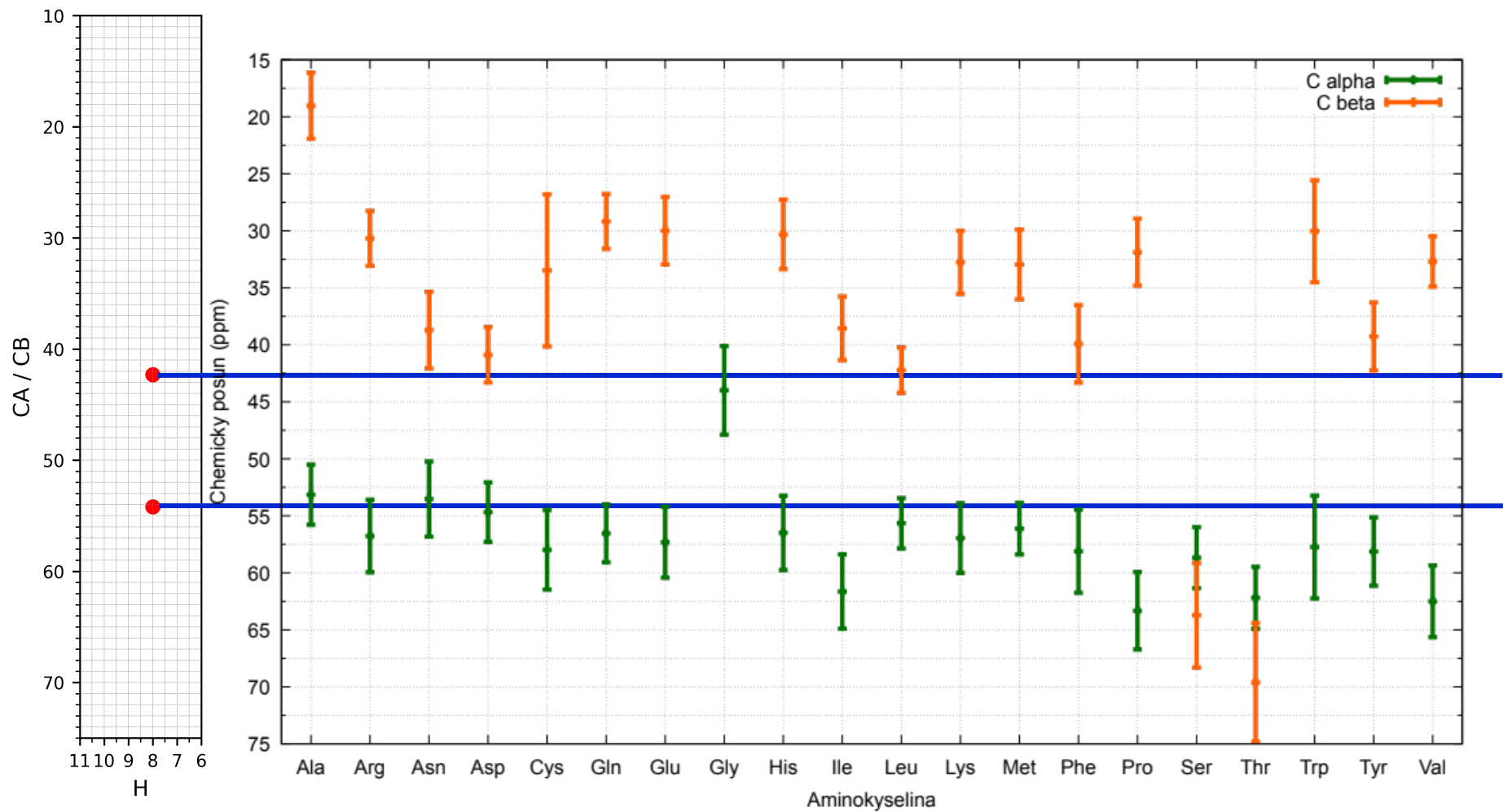
F

HNCACB (138)



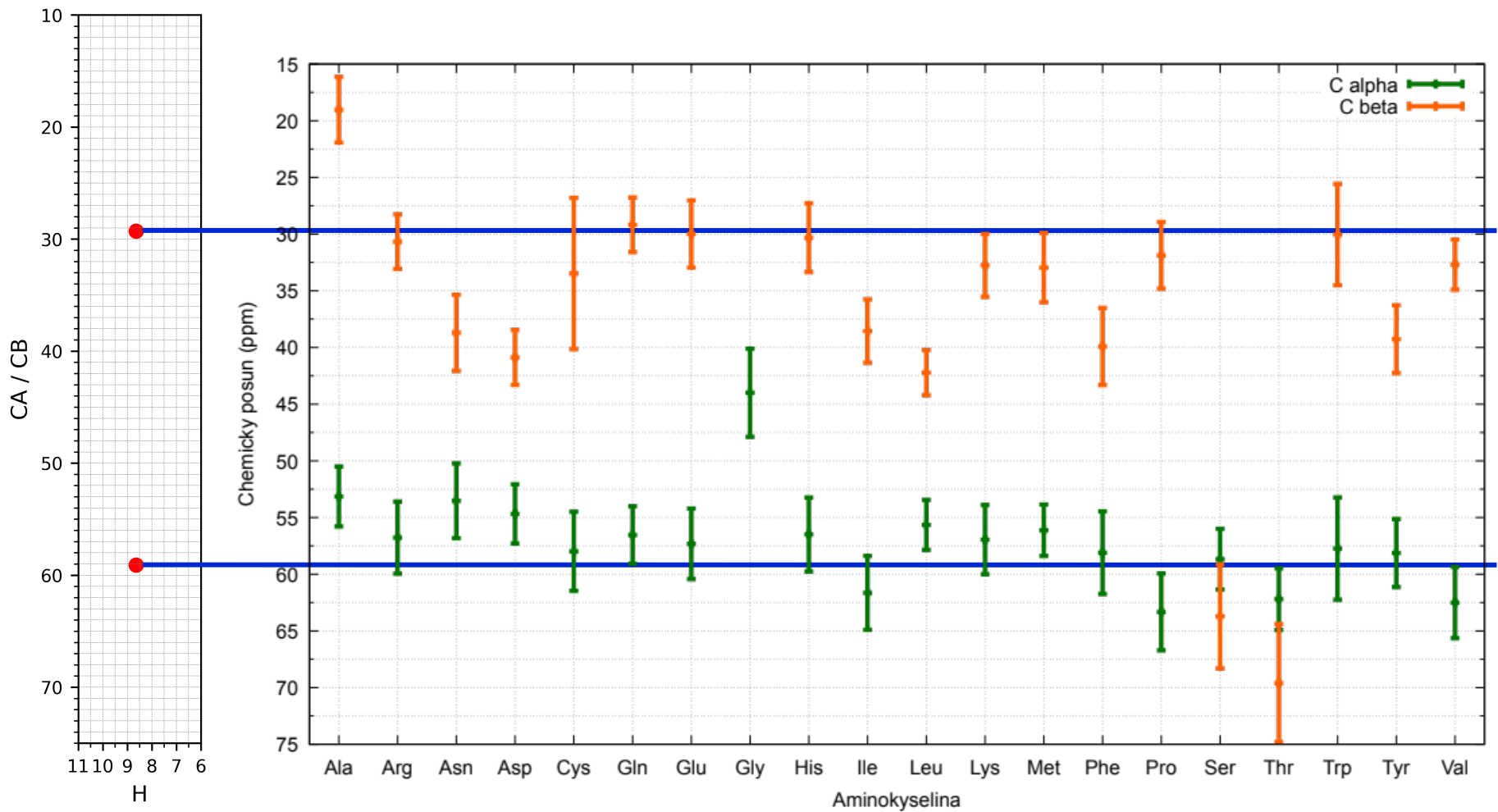
D, I, V

HNCACB (138)



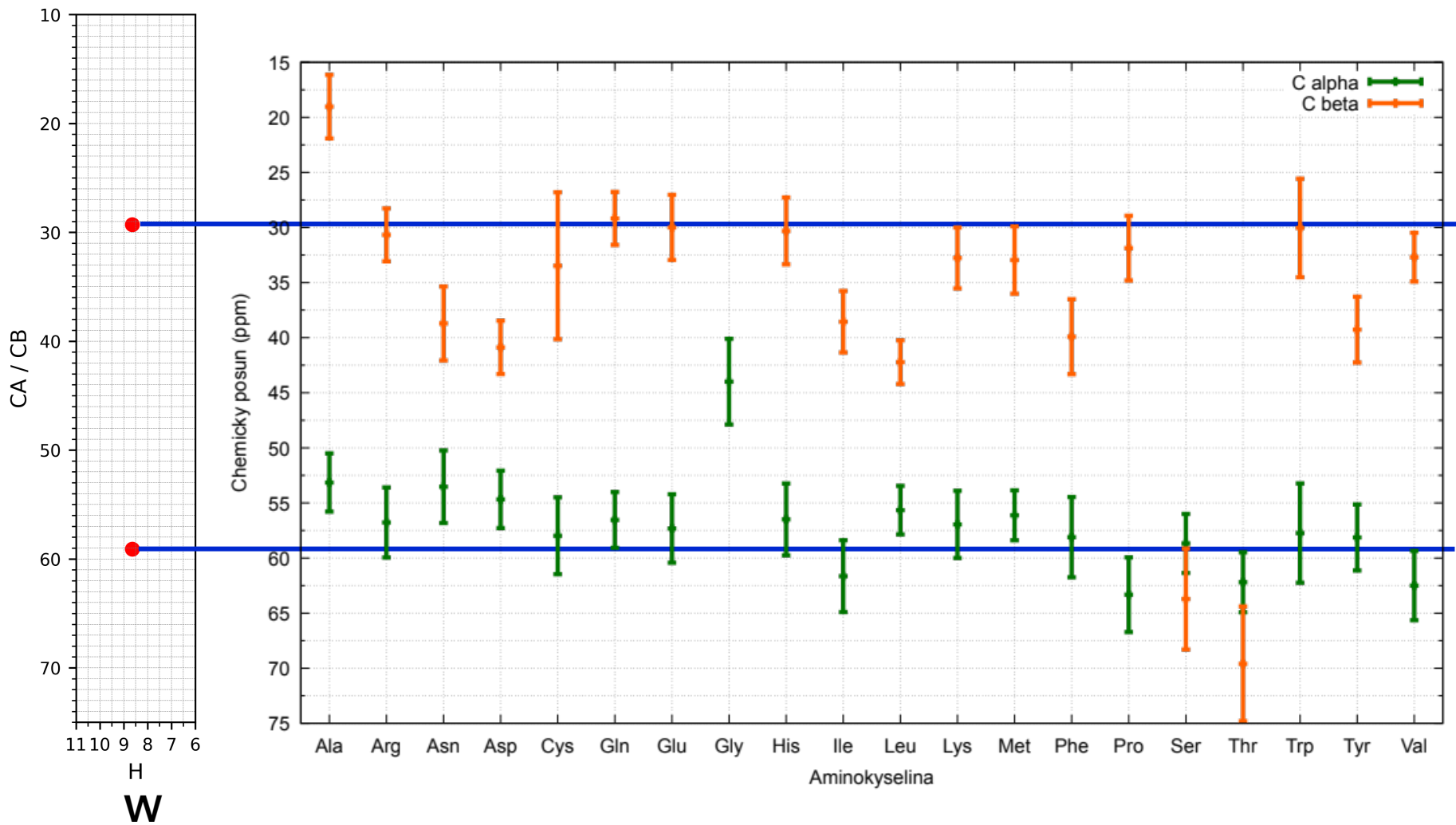
D

HNCACB (8)

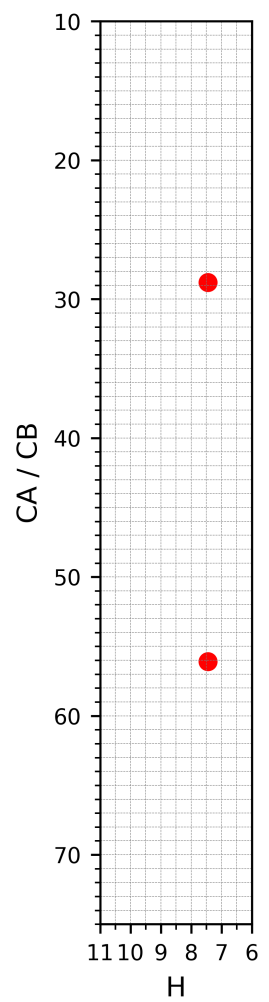


W, T, D

HNCACB (8)

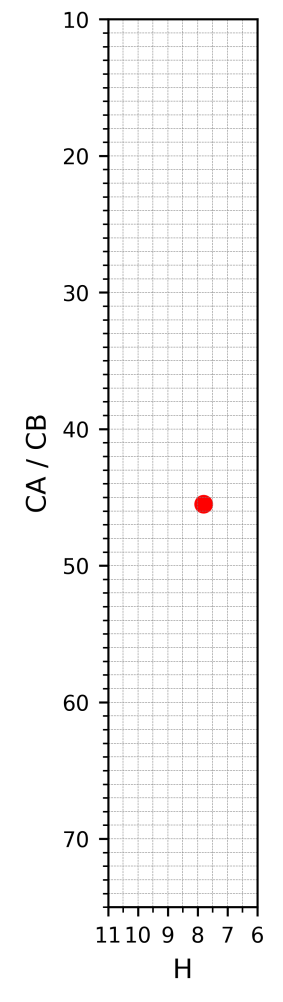


HNCACB (13)



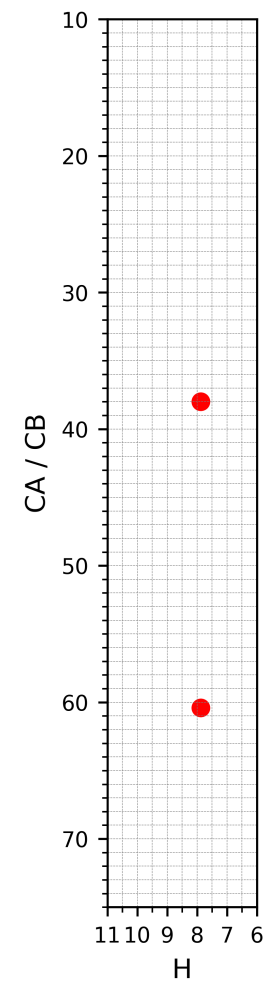
R

HNCACB (3)



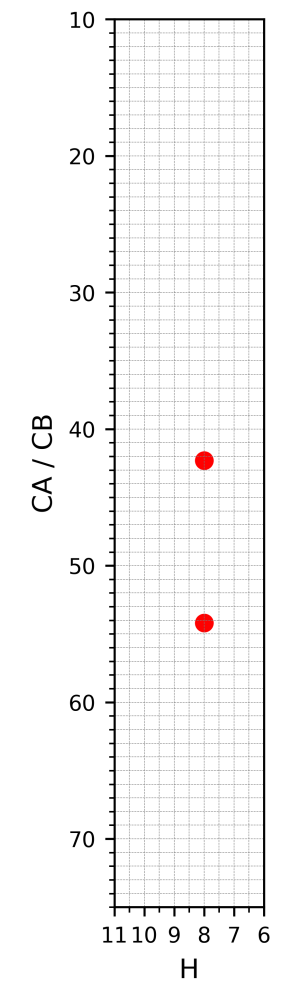
G

HNCACB (11)



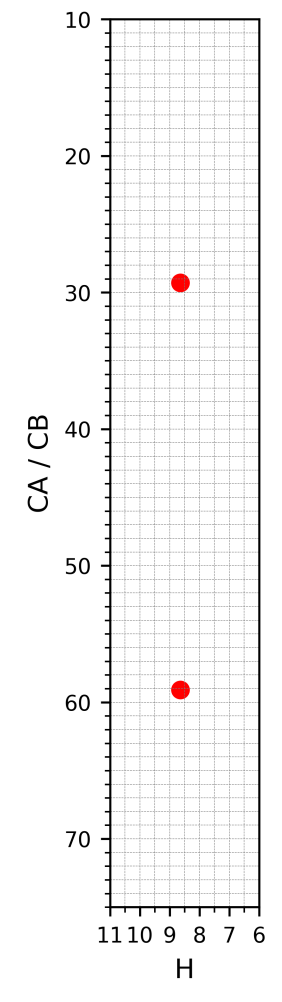
F

HNCACB (138)



D

HNCACB (8)



W

Úkol č. 3 – najdi přiřazení v sekvenci proteinu

10 20 30 40 50 60
MSHHWGYGKH NGPEHWHKDF PIAKGERQSP VDIDTHTAKY DPSLKPLSVS YDQATSLRIL

70 80 90 100 110 120
NNGHAFNVEF DDSQDKAVLK GGPLDGTYRL IQFHFHWGSL DGQGSEHTVD KKKYAAELHL

130 140 150 160 170 180
VHWNTKYGDF GKAVQQPDGL AVLGIFLKVG SAKPGLQKV DVLDSIKTKG KSADFTNFDP

190 200 210 220 230 240
RGLLPESLDY WTYPGSLTTP PLLECVTWIV LKEPISVSSE QVLKFRKLNF NGEGEPEELM

250 260
VDNWRPAQPC RGF~~D~~WLKNRQ IKASFK

R G F D W

Úkol č. 3 – najdi přiřazení v sekvenci proteinu

10 20 30 40 50 60
MSHHWGYGKH NGPEHWHKDF PIAKGERQSP VDIDTHTAKY DPSLKPLSVS YDQATSLRIL

70 80 90 100 110 120
NNGHAFNVEF DDSQDKAVLK GGPLDGTYRL IQFHFHWGSL DGQGSEHTVD KKKYAAELHL

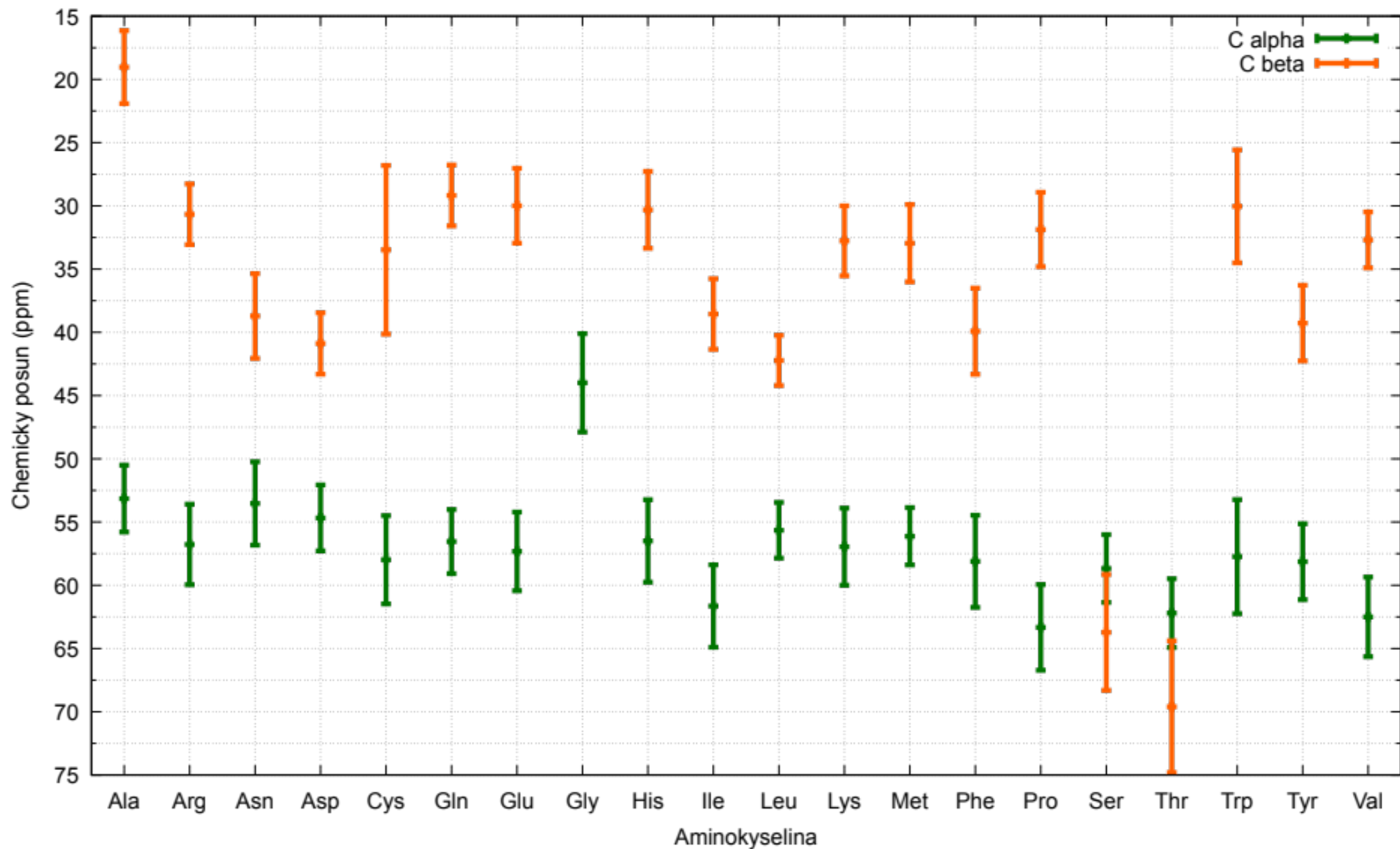
130 140 150 160 170 180
VHWNTKYGDF GKAVQQPDGL AVLGIFLKVG SAKPGLQKV DVLDSIKTKG KSADFTNFDP

190 200 210 220 230 240
RGLLPESLDY WTYPGSLTTP PLLECVTWIV LKEPISVSSE QVLKFRKLNK NGEGEPEELM

250 260
VDNWRPAQPC **RGFDW** LKNRQ IKASFK

R G F D W

Typické chemické posuny C α a C β aminokyselin

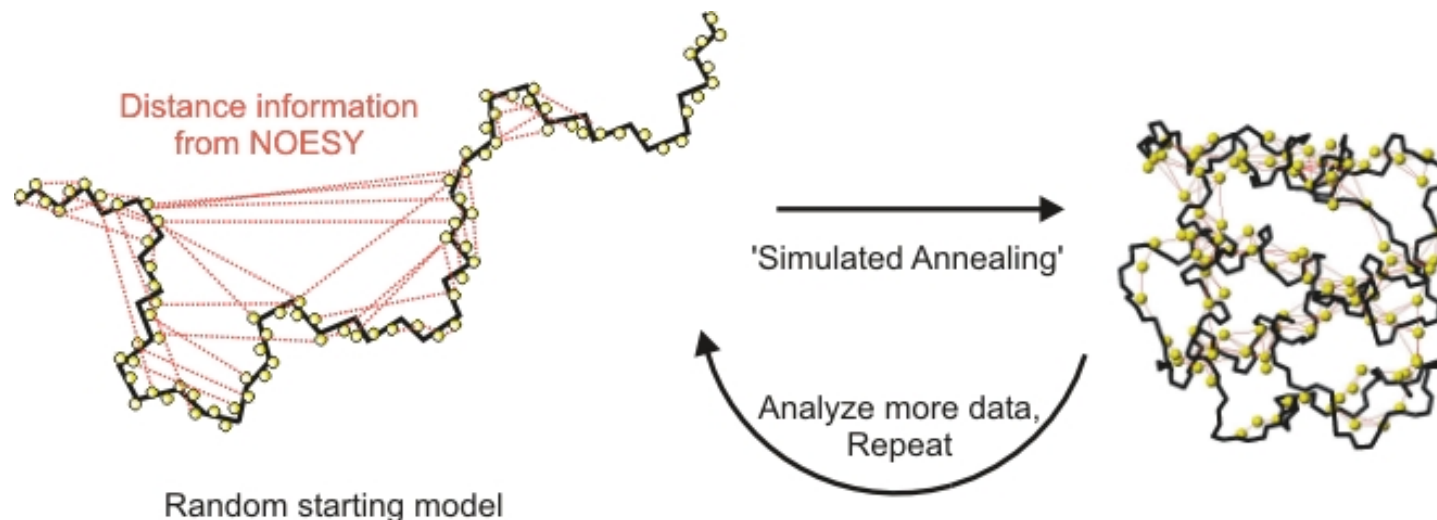


Struktura

- **sekvenční přiřazení** –interakce přes vazbu přes skalární J-coupling
- **určení vzdáleností mezi atomy ^1H** – interakce přes prostor NOESY
 - intenzita klesá s r^6 vzdálenosti
- při řešení struktury **molekulová dynamika** omezená experimentálními daty

Molekulární dynamika

- získané vzdálenosti z NOESY experimentu
- další experimentální data (torzní úhly z chemických posunů)
- simulované žíhání (zahřívání a ochlazení)
- opakujeme (např. 8x)
- vybereme struktury s nejnižší energií (20/100)



Výsledná struktura

- překryv několika (např. 10, 20) energeticky nejvýhodnějších struktur

