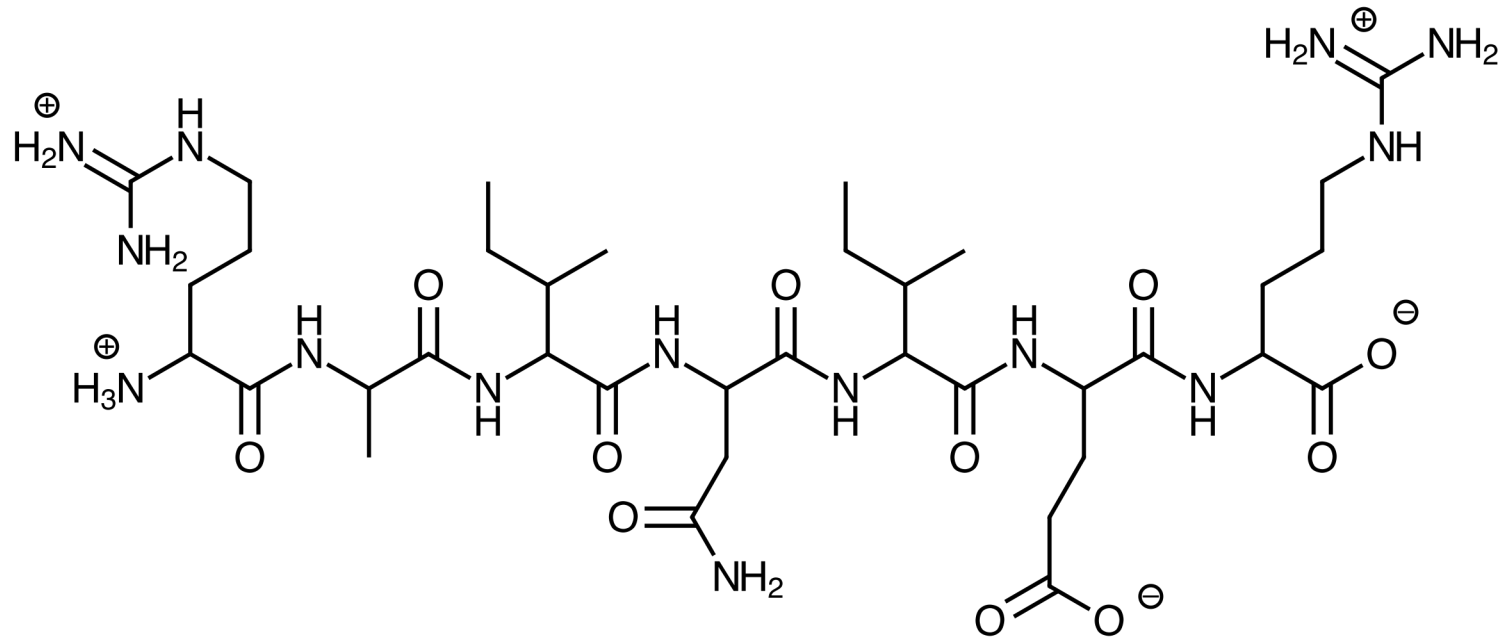
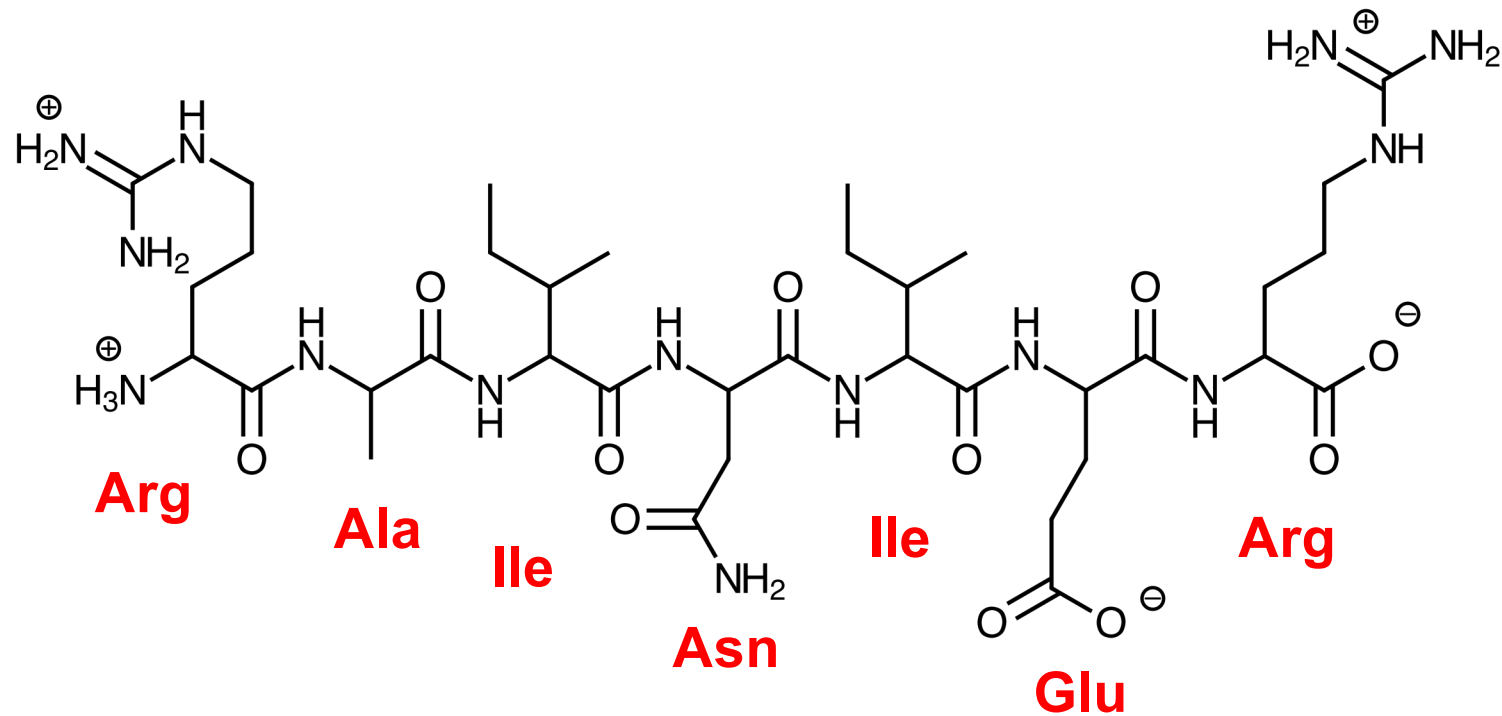


Who Am I?



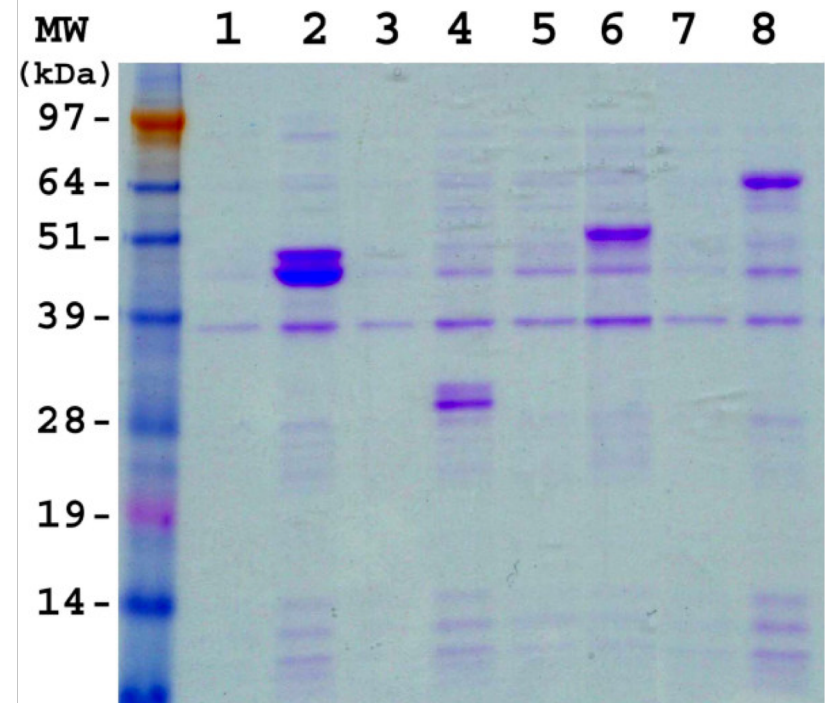
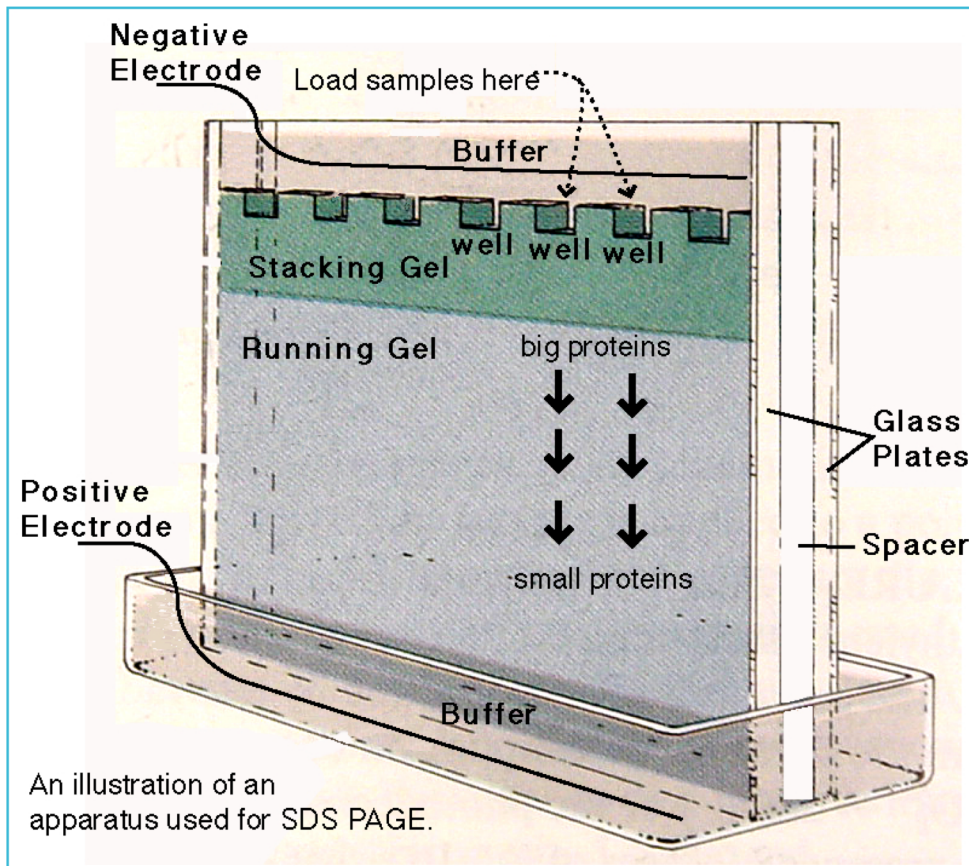
...and who geeked out this week?

Who Am I?



...and who geeked out this week?

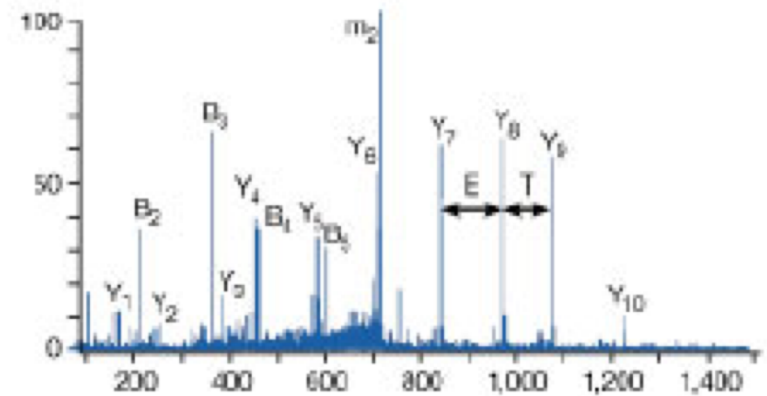
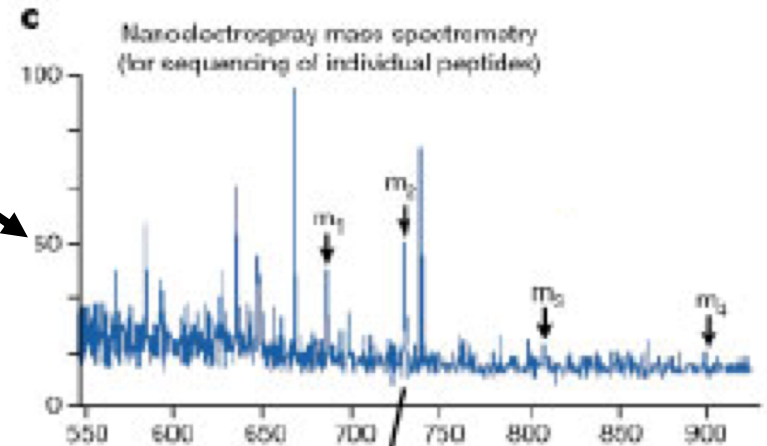
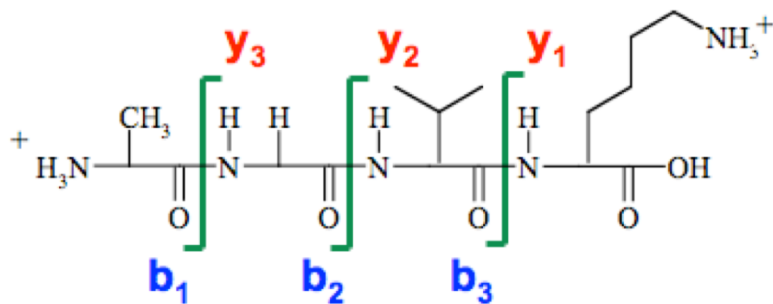
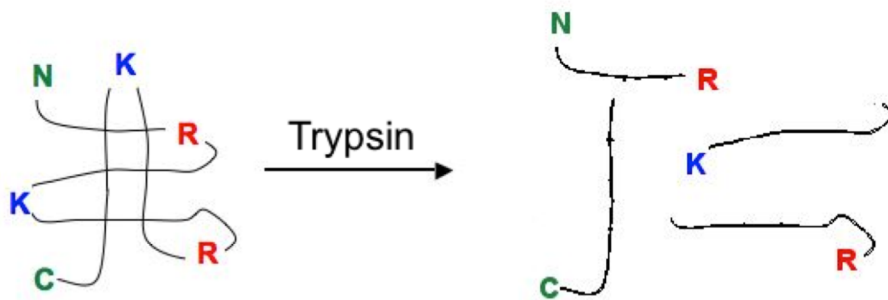
Denaturing Polyacrylamide Gel Electrophoresis



https://ww2.chemistry.gatech.edu/~lw26/bCourse_Information/4581/techniques/gel_elect/page_protein.html

<http://www.clinicalmolecularallergy.com/content/4/1/12/figure/F1>

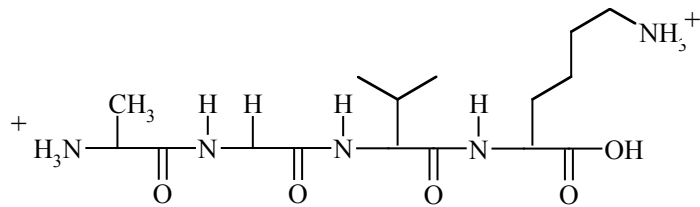
MS/MS Sequencing



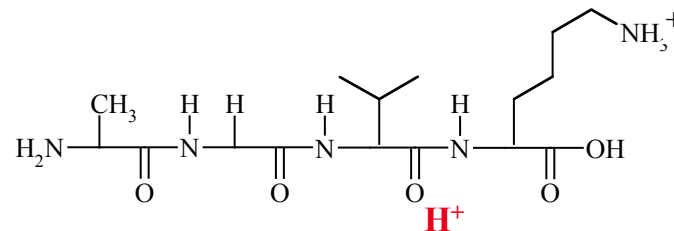
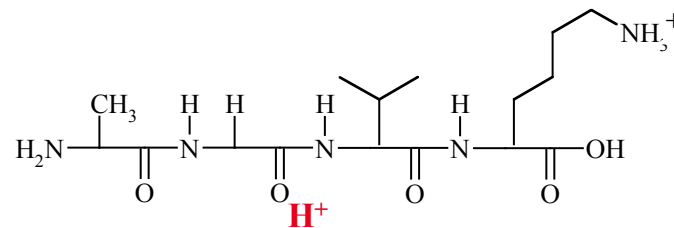
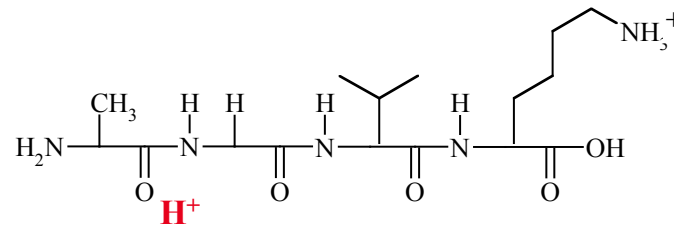
A peptide sequence tag derived from the above data is used to search nucleotide and protein sequence databases

Migration of N-terminal Proton

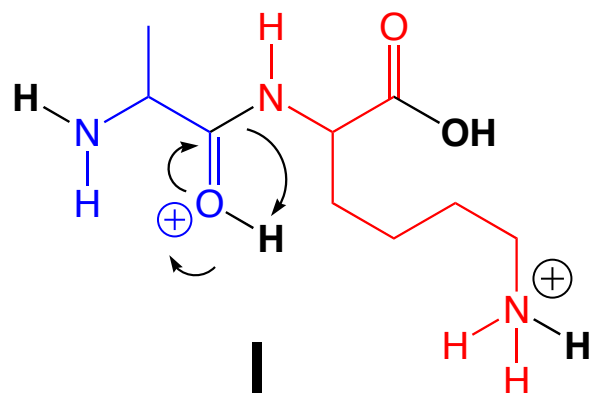
| | | |
|--------|------------------------|--------------|
| Mass = | Ala | 71.0 |
| | Gly | 57.0 |
| | Val | 99.1 |
| | Lys | 128.1 |
| | OH | 17.0 |
| | <u>3 H⁺</u> | <u>3.0</u> |
| | | 375.2 |



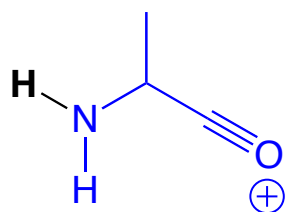
$$m/z = 375.2/2 = 187.6$$



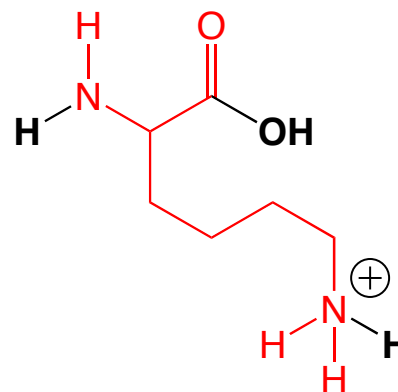
Collisionally Induced Dissociation



| | | |
|---------|--------------|------|
| b_1 : | Ala | 71.1 |
| | H^+ | 1.0 |
| | <hr/> | |
| | | 72.1 |

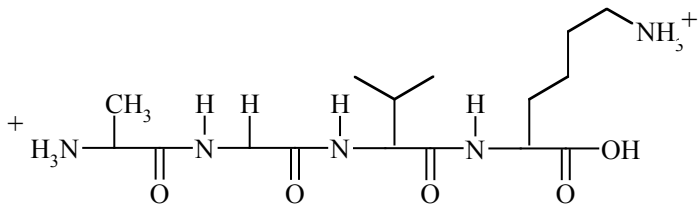
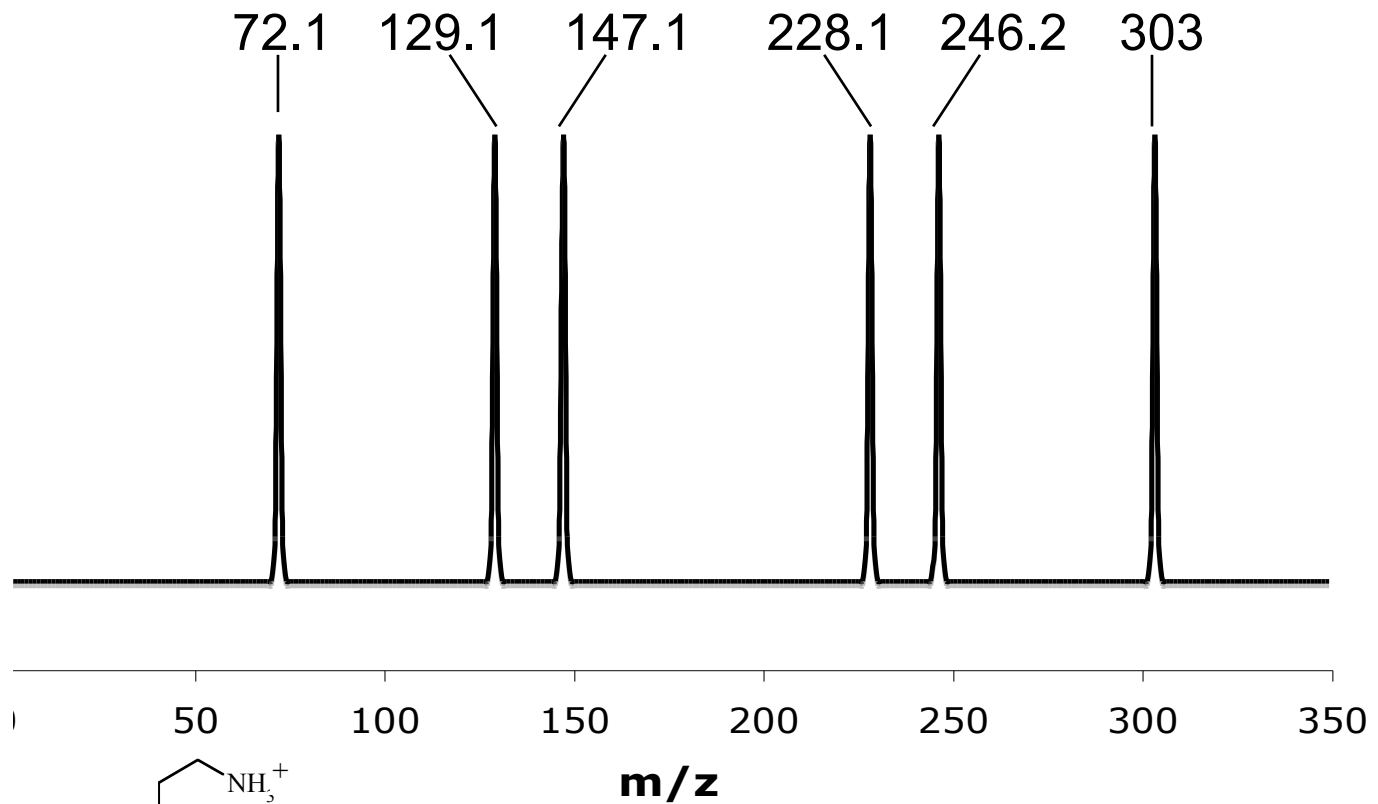


| | | |
|---------|---------------|-------|
| y_1 : | Lys | 128.1 |
| | 2H^+ | 2.0 |
| | OH | 17.0 |
| | <hr/> | |
| | | 147.1 |



Fragment Spectrum of AGVK

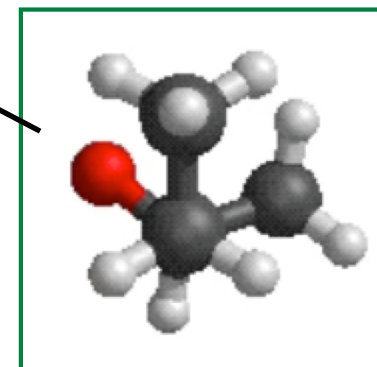
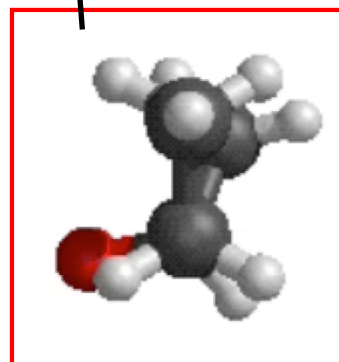
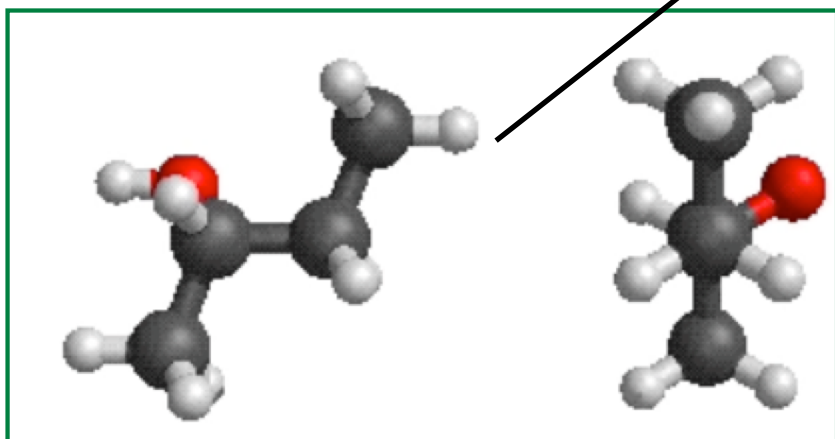
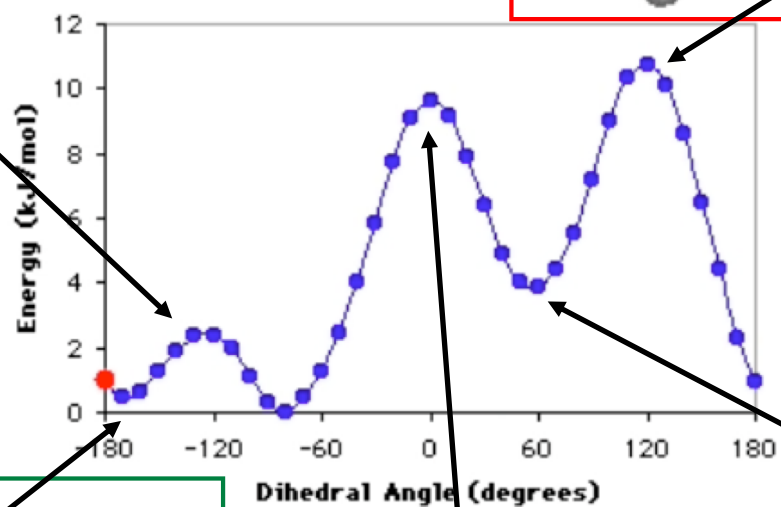
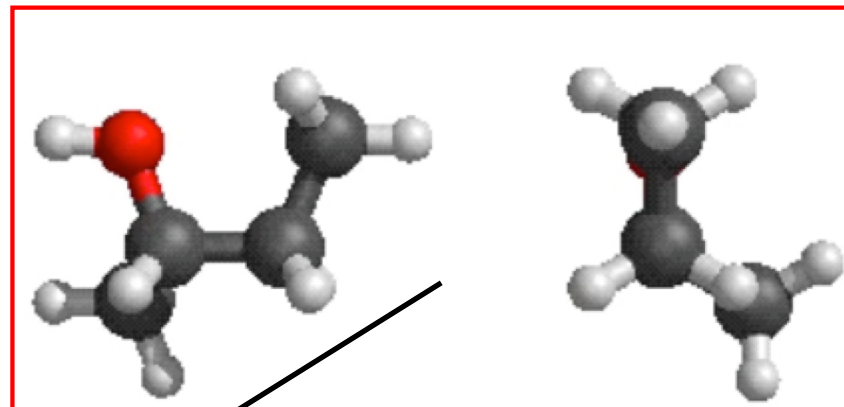
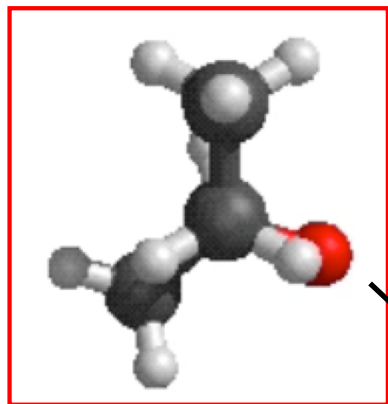
$$m/z = 375.2/2 = 187.6$$



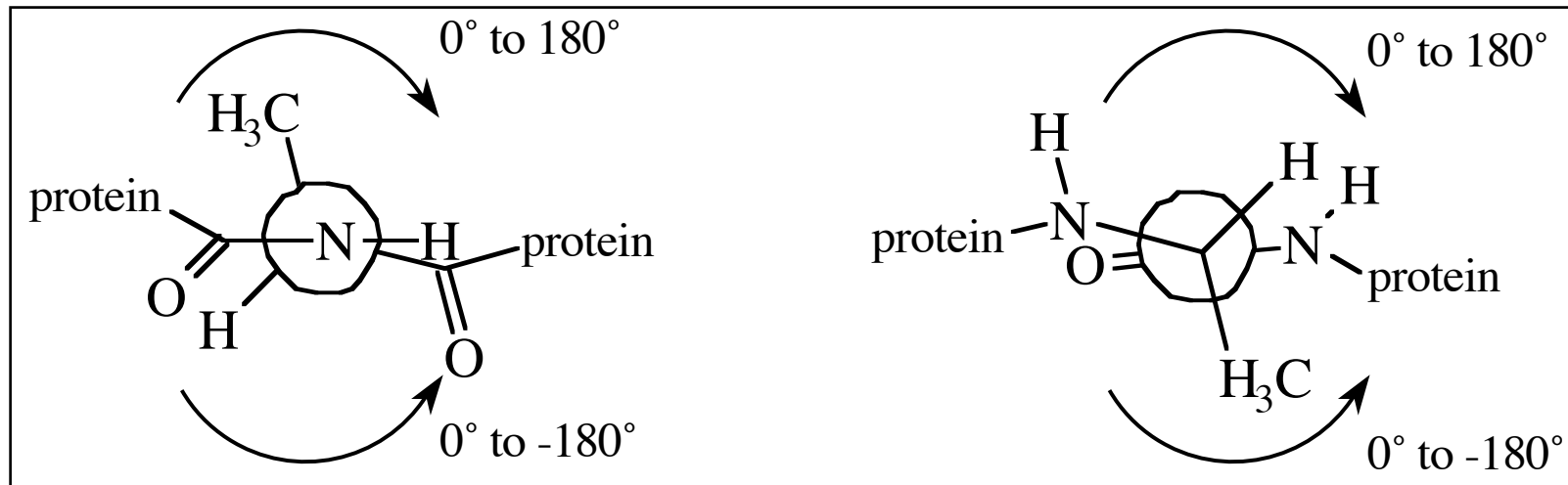
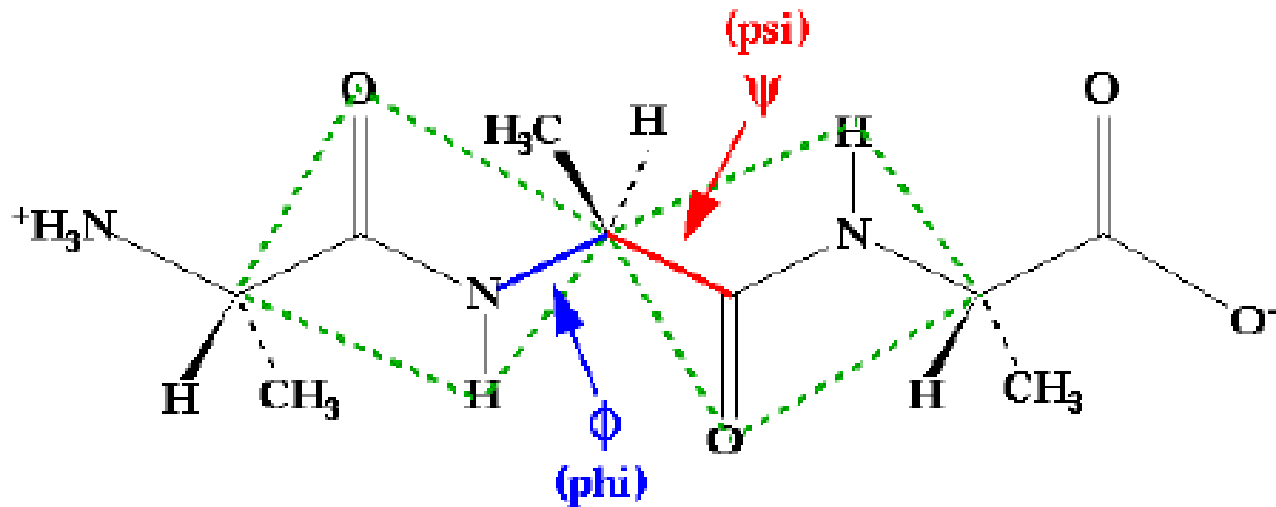
Properties of the Twenty

| Amino Acid | Abbreviations | Residue Mass (Da) | Side Chain pK _a | $\Delta\Delta G^\circ$ transfer (kcal/mol) |
|---------------|---------------|-------------------|----------------------------|--|
| Alanine | Ala, A | 71.09 | - | -0.87 |
| Arginine | Arg, R | 156.19 | 12 | 15.93 |
| Asparagine | Asn, N | 114.11 | - | 5.22 |
| Aspartic Acid | Asp, D | 115.09 | 4 | 9.71 |
| Cysteine | Cys, C | 103.15 | 8 | -0.34 |
| Glutamine | Gln, Q | 128.14 | - | 6.51 |
| Glutamic Acid | Glu, E | 129.12 | 4 | 7.78 |
| Glycine | Gly, G | 57.05 | - | 0.00 |
| Histidine | His, H | 137.14 | 7 | 5.63 |
| Isoleucine | Ile, I | 113.16 | - | -4.00 |
| Leucine | Leu, L | 113.16 | - | -4.00 |
| Lysine | Lys, K | 128.17 | 10 | 6.52 |
| Methionine | Met, M | 131.19 | - | -1.42 |
| Phenylalanine | Phe, F | 147.18 | - | -2.05 |
| Proline | Pro, P | 97.12 | - | <0 |
| Serine | Ser, S | 87.08 | - | 4.36 |
| Threonine | Thr, T | 101.11 | - | 3.53 |
| Tryptophan | Trp, W | 186.21 | - | -1.40 |
| Tyrosine | Tyr, Y | 163.18 | 10 | 1.09 |
| Valine | Val, V | 99.14 | - | -3.11 |

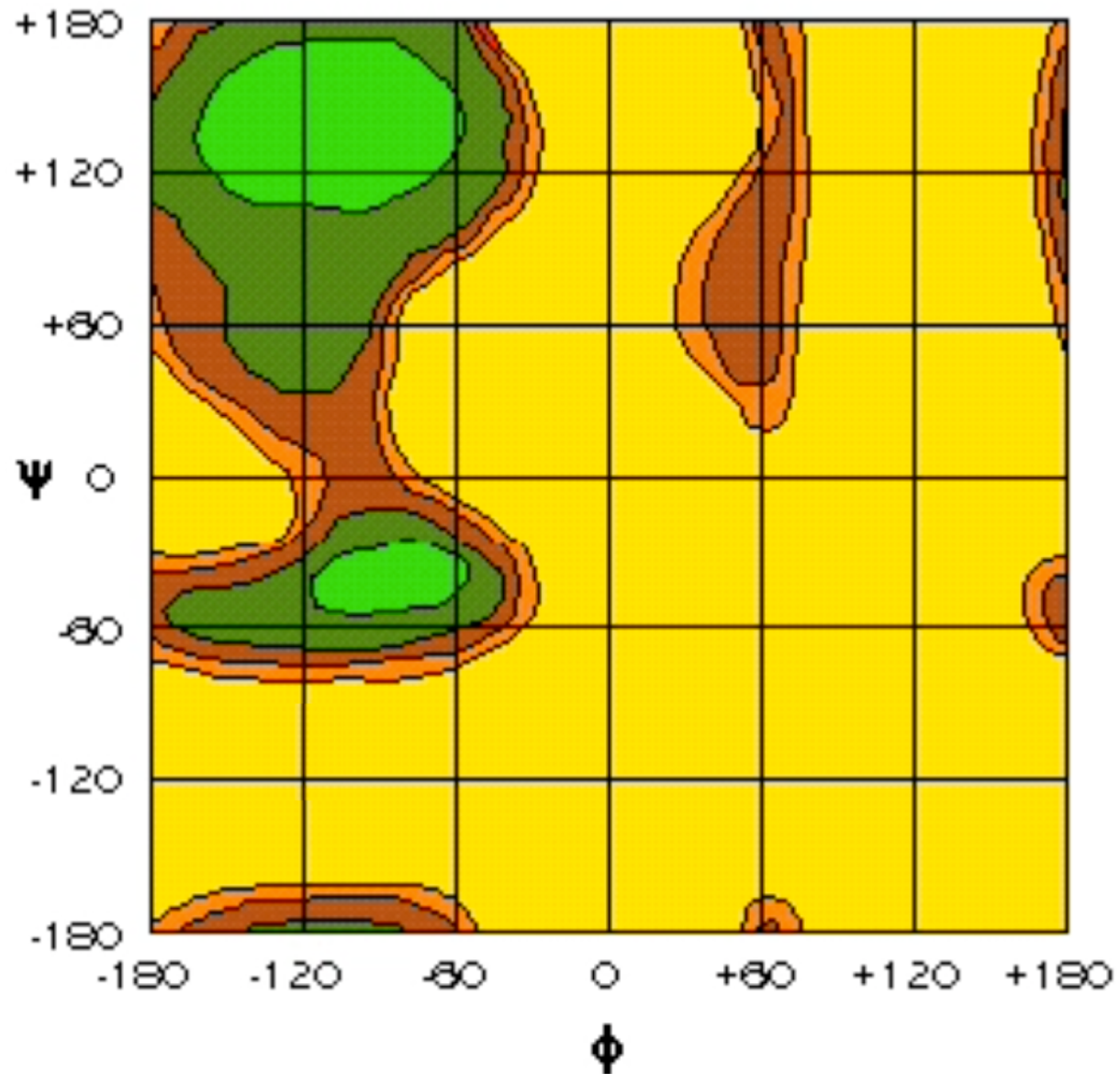
S-2-Butanol Conformers

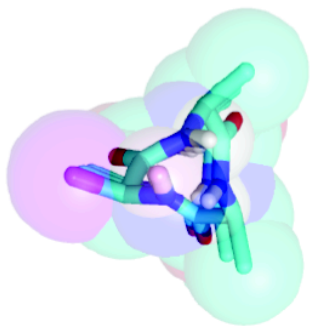


Phi (ϕ) and Psi (ψ) Dihedrals

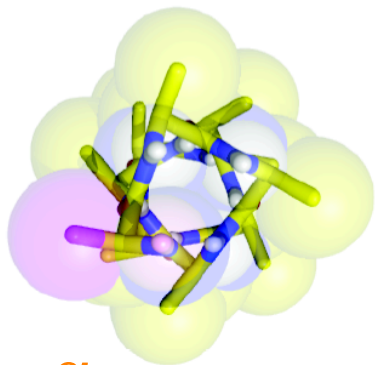


Ramachandran plot

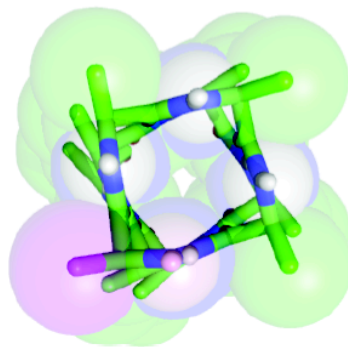




3_{10}

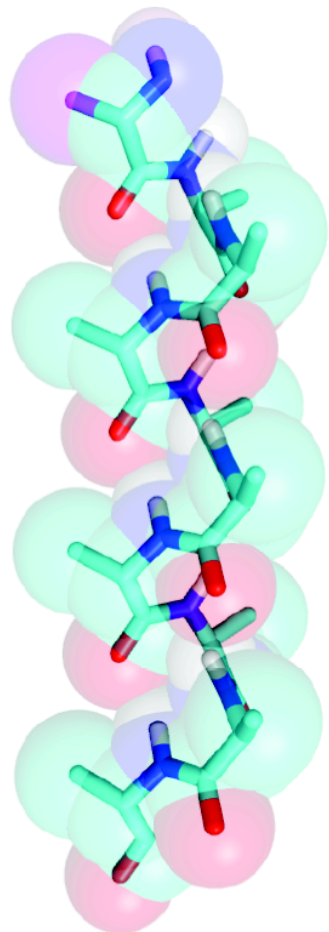


α

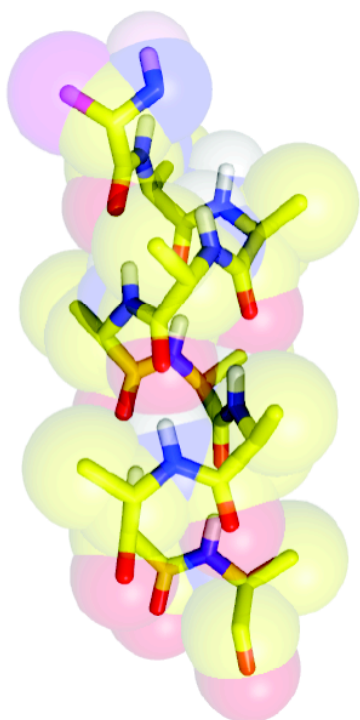


π

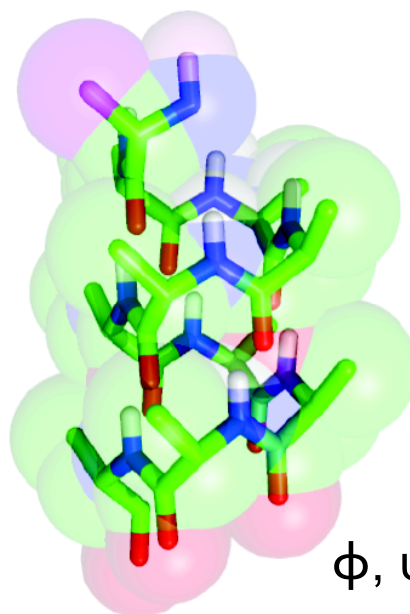
Three Possible
Right-handed
Helices



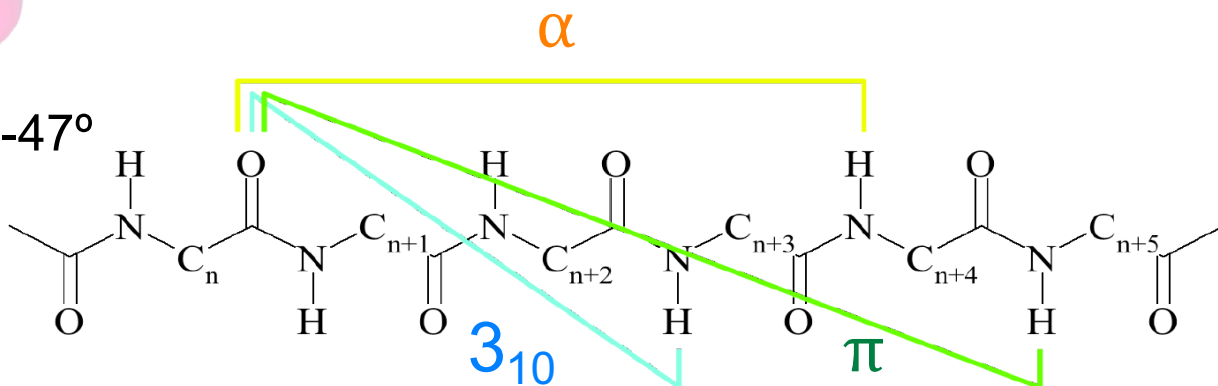
$\phi, \psi = -74^\circ, -4^\circ$



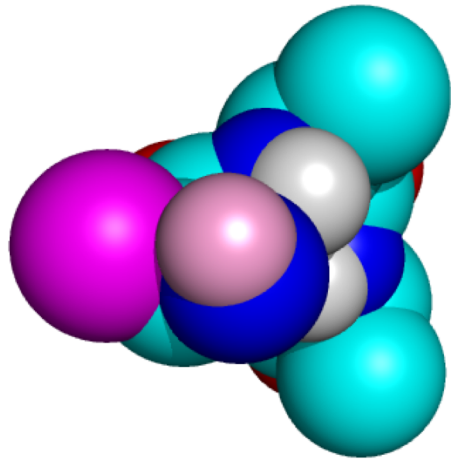
$\phi, \psi = -56^\circ, -47^\circ$



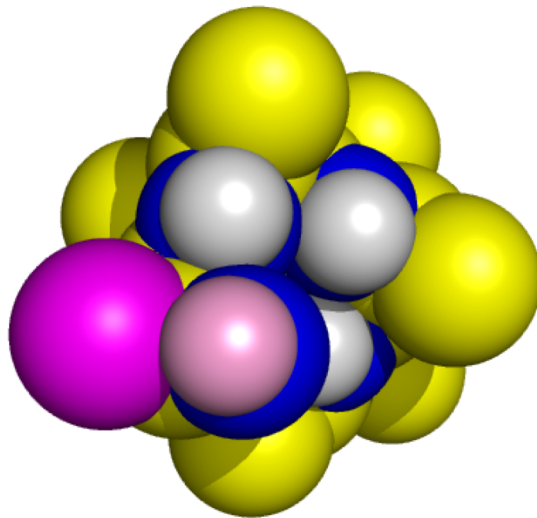
$\phi, \psi = -59^\circ, -70^\circ$



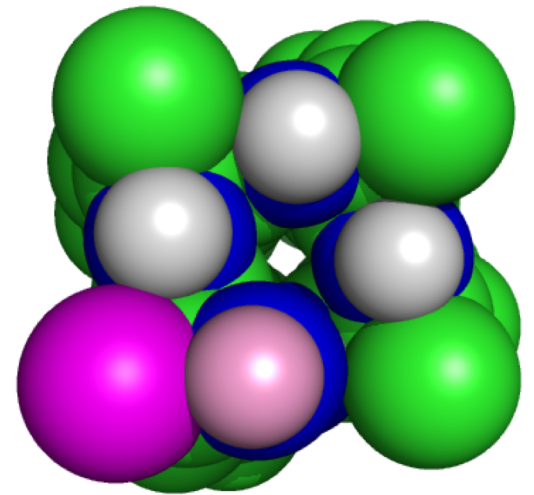
Three Helices



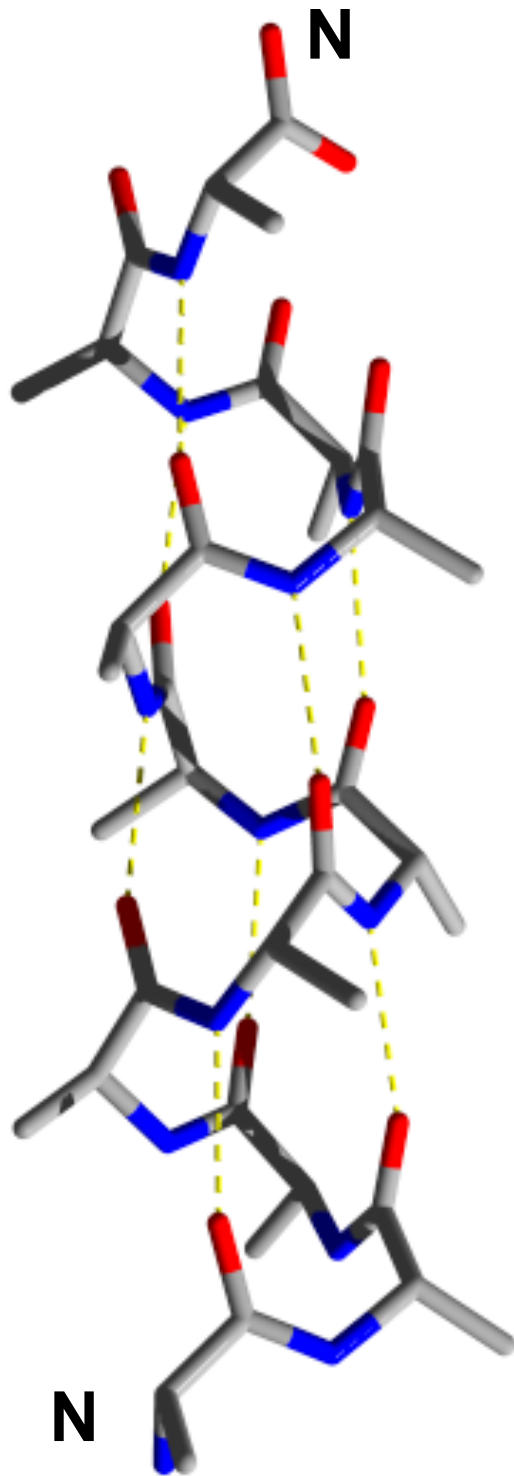
3_{10}



α



π



α -Helix Statistics

Right-handed twist

$\phi, \psi = -47^\circ, -57^\circ$

3.6 residues/turn

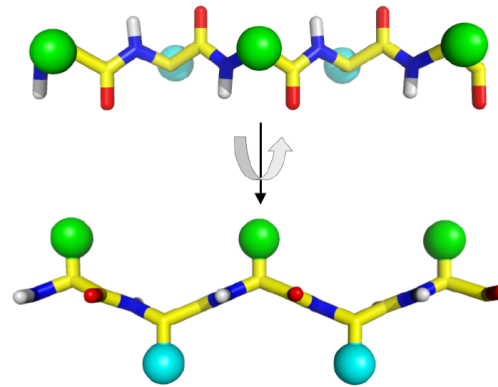
Rise/AA along axis = 1.5 Å

Rise/turn along axis = 5.4 Å

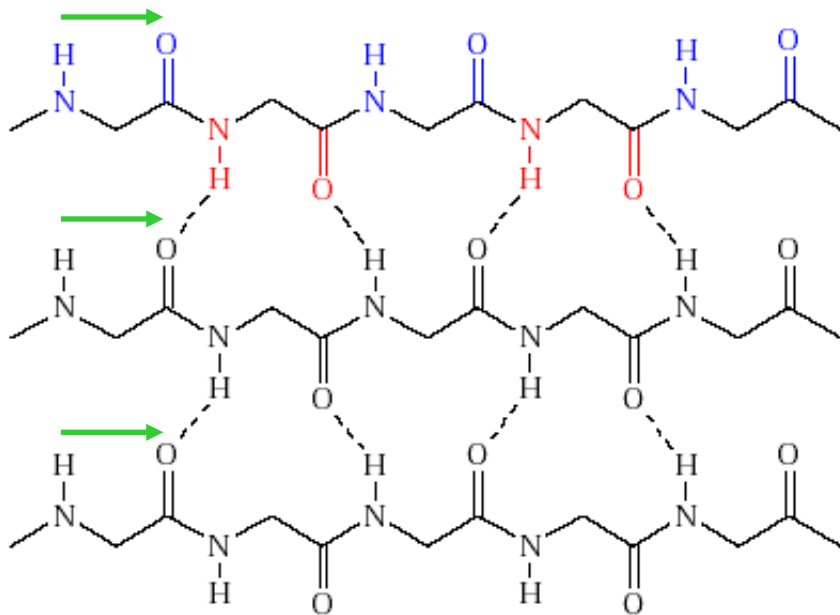
Helix diameter = 4.6 Å

Beta Strands

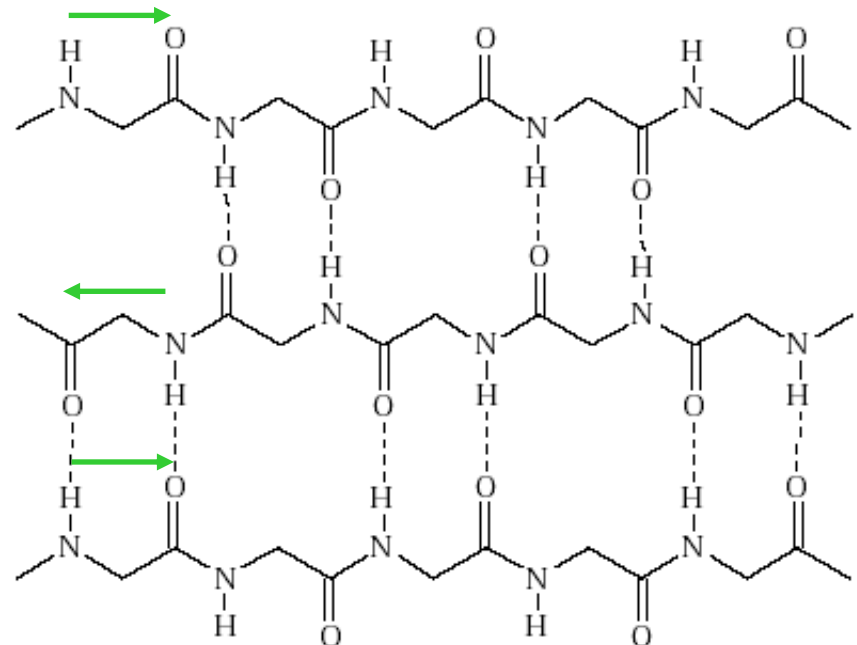
$\phi \approx -120^\circ$ to -140° , $\psi = 110^\circ$ to 130°



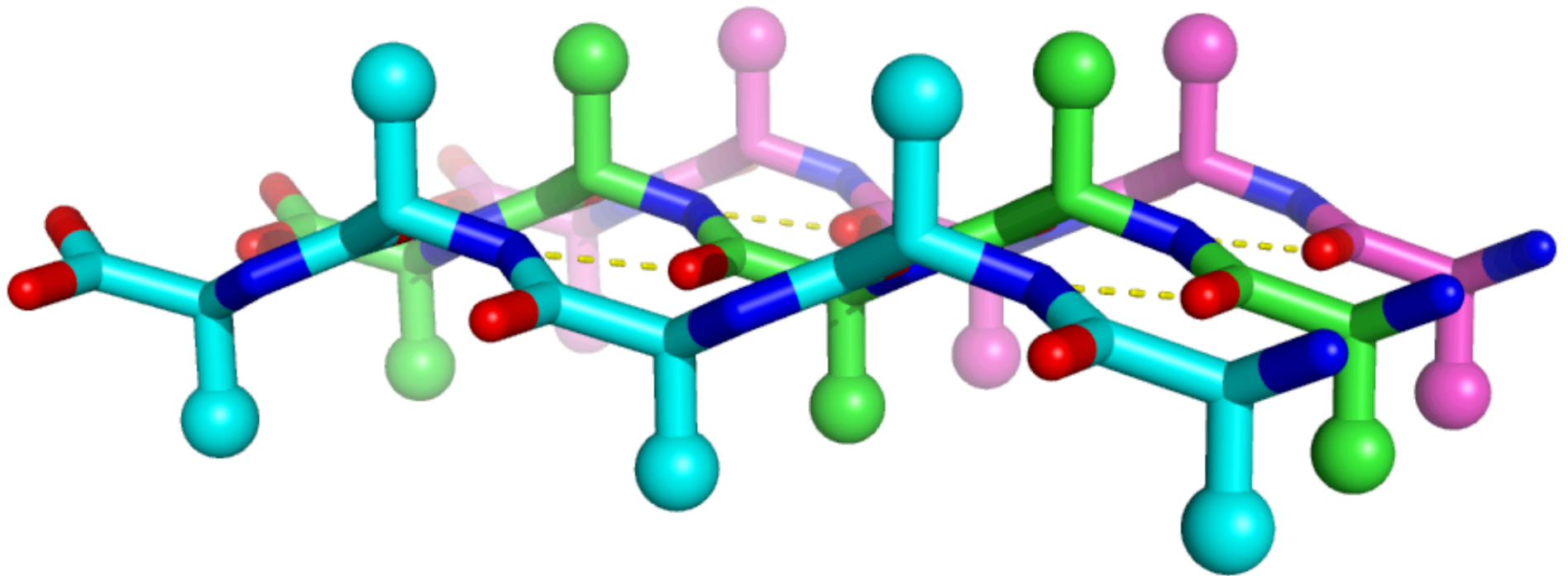
Parallel β -Sheet



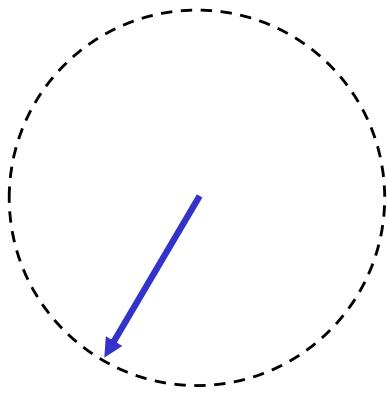
Antiparallel β -Sheet



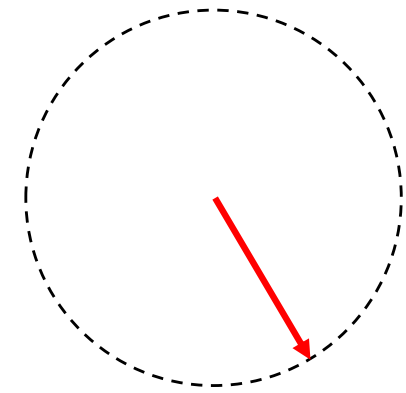
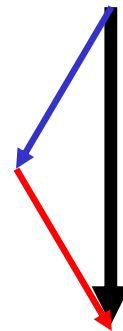
Beta Pleated Sheet



Plane-Polarized Light



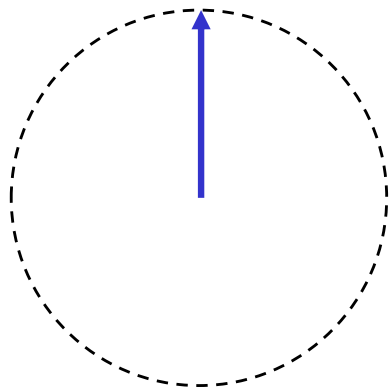
Left Circularly
Polarized Light



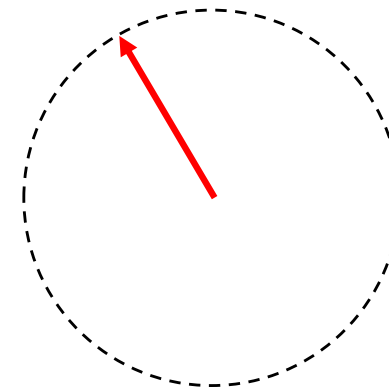
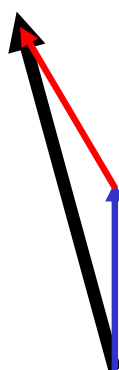
Right Circularly
Polarized Light

Equal populations of both
Forms of polarized light create
That oscillate in a plane

Optical Rotation

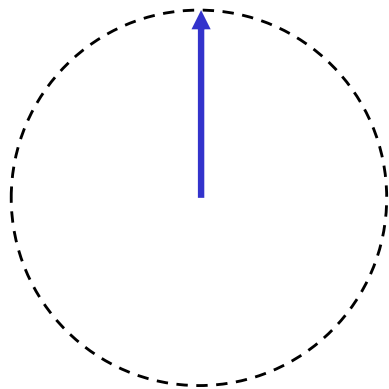


Left Circularly
Polarized Light

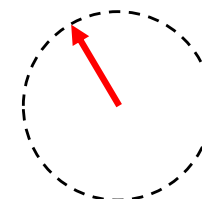
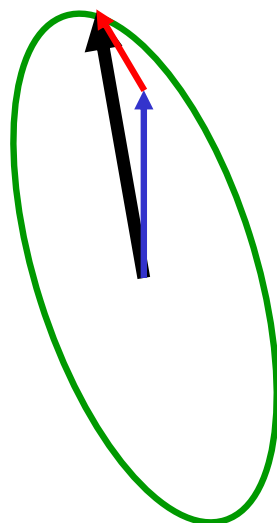


Right Circularly
Polarized Light –
Retarded by Chiral
compound, fields now
sum to give rotated
plane of polarization

Circular Dichroism



Left Circularly Polarized Light



Right Circularly Polarized Light –
Retarded and absorbed
by Chiral
Compound

CD Spectra of 2° Structure

θ = ellipticity

$[\theta]$ = molar ellipticity

$[\theta] = 3298(\epsilon_L - \epsilon_R)$

