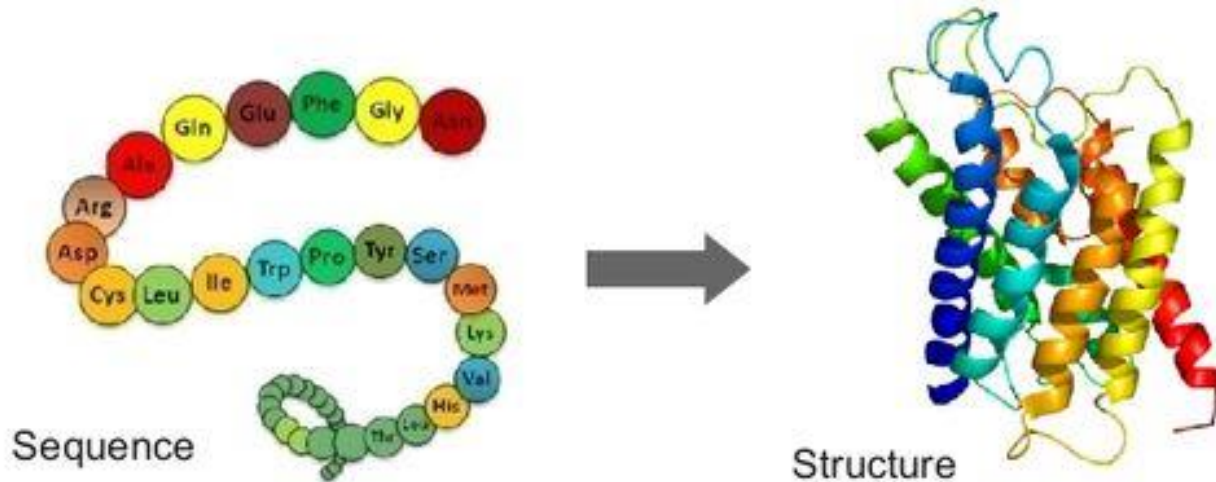


# Prerequisite Knowledge

The same day I do this module, I cover:

- 1) Amino acid structures and groupings
- 2) Peptide bond formation
- 3) N- and C-termini

# Amino Acids – an introduction to Proteins

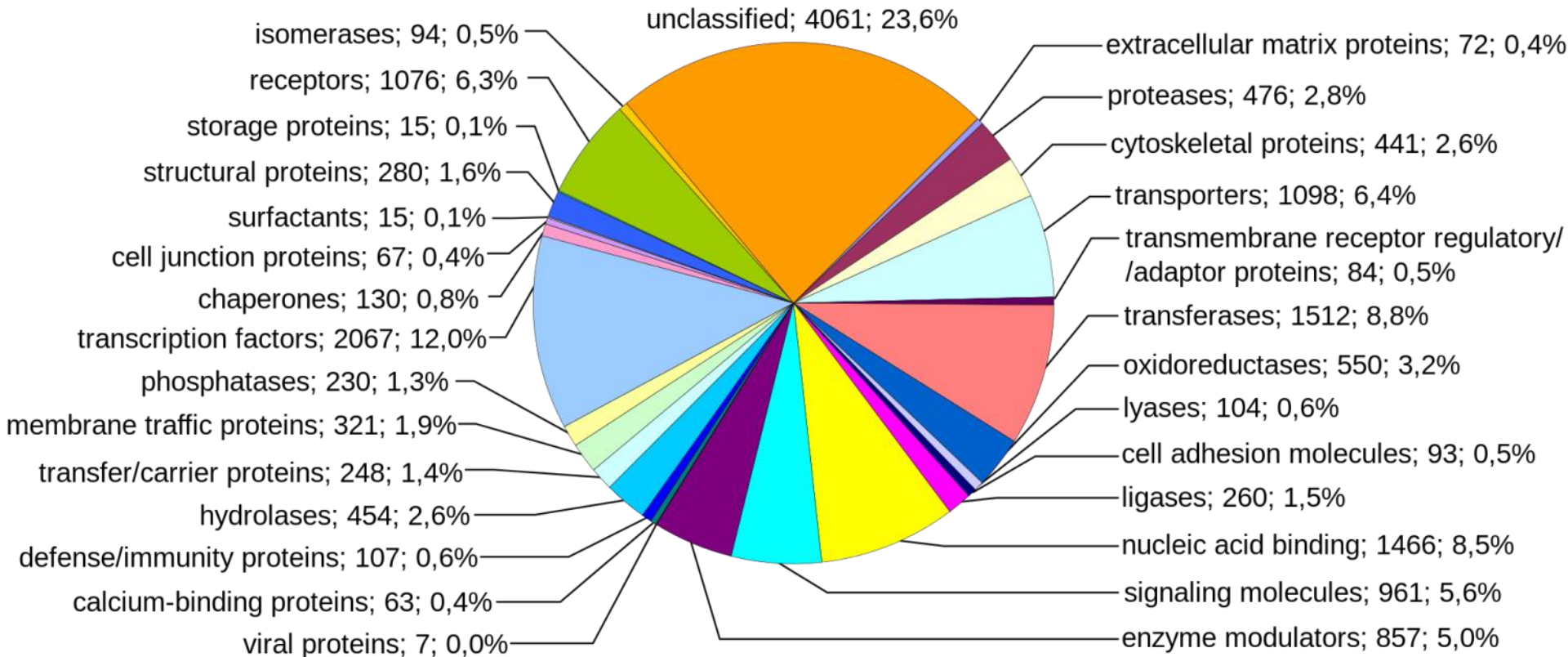


Over the next couple of weeks we will look at the how proteins are made:

- Primary structure (string of amino acids)
- Secondary structure (common, stabilized 3D features)
- Tertiary structure (entire 3D folding pattern)
- Quaternary structure (more than one amino acid chain)

# What are proteins good for?

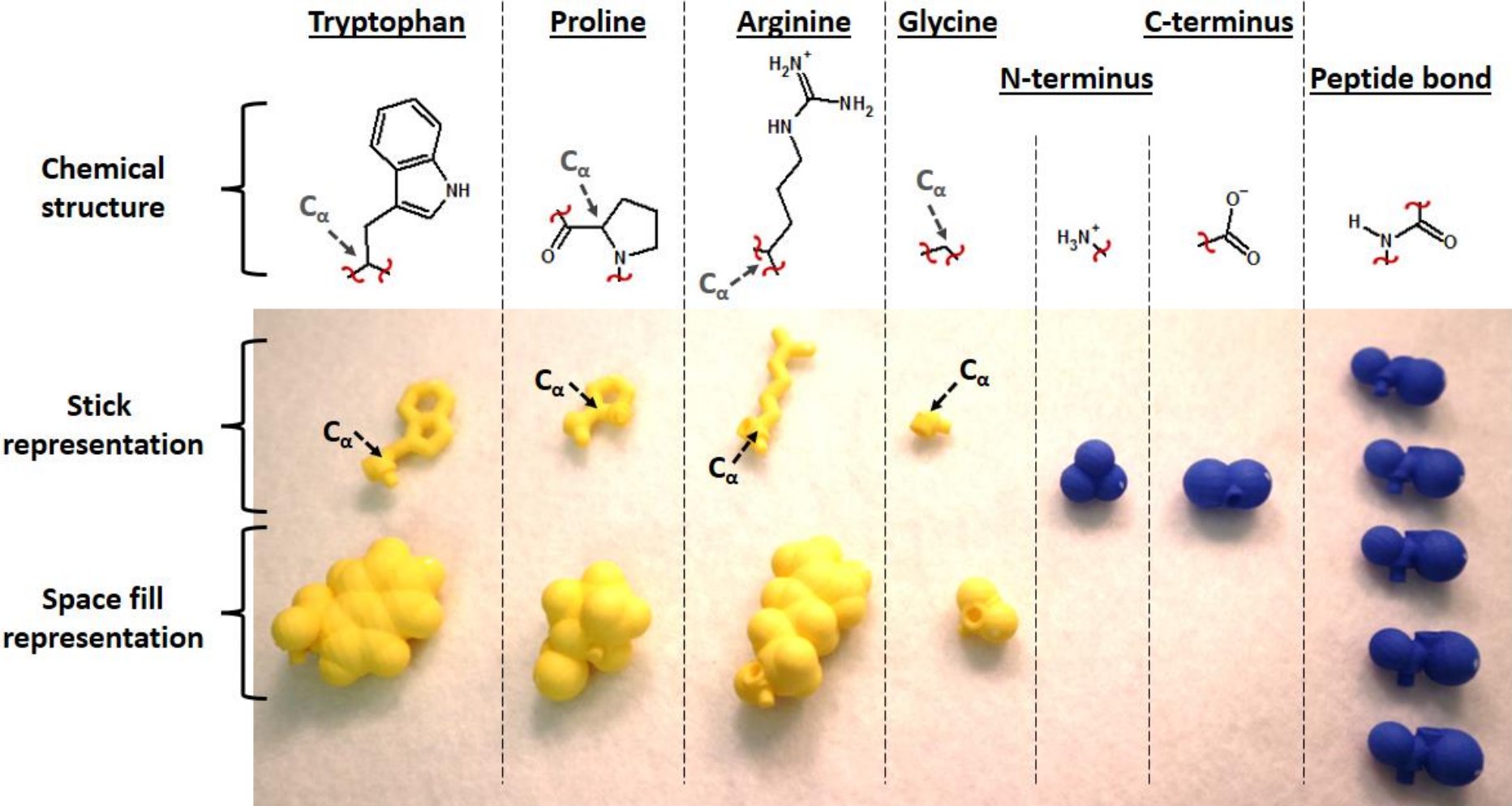
Proteins grouped by function from the Human Genome



Just about everything!

Proteins control almost every action and reaction—including thought!

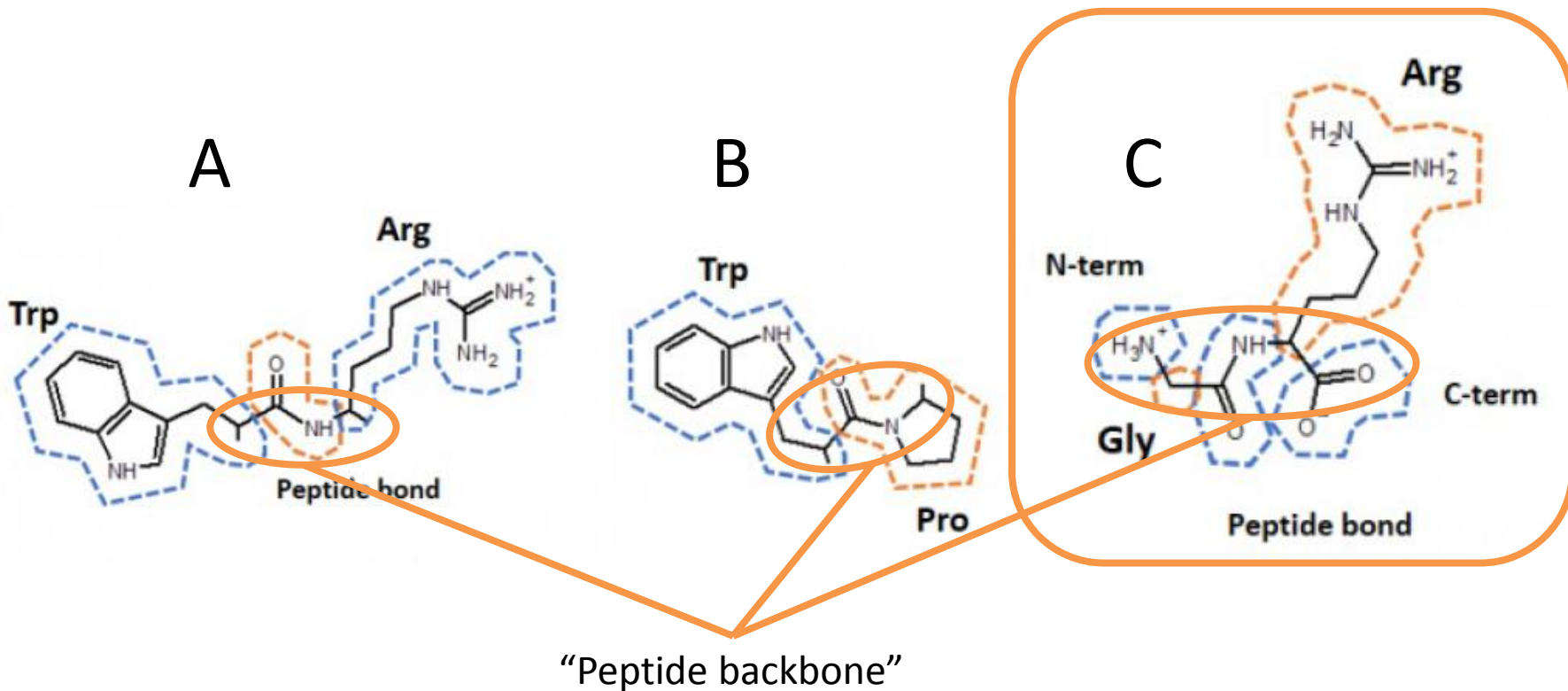
# Connecting Amino Acids Activity



If doing an electronic version of the activity, provide link here

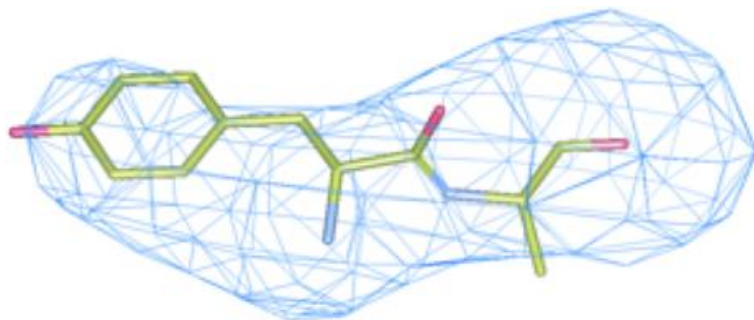
# Amino Acid Clicker Question:

Which one of these dipeptides will have the greatest rotation around the  $C_{\alpha}$ s?

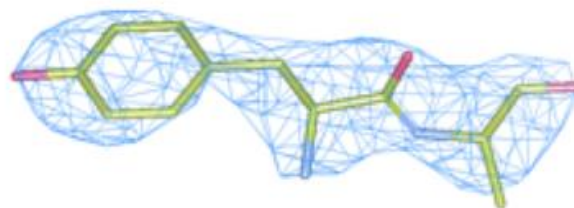




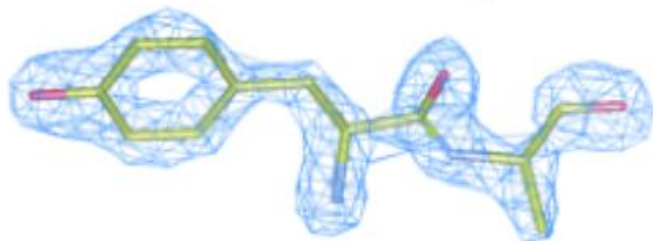
# X-ray crystallography



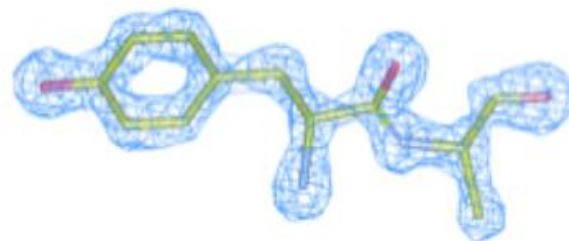
5.0 Å resolution,  $2.5\sigma$



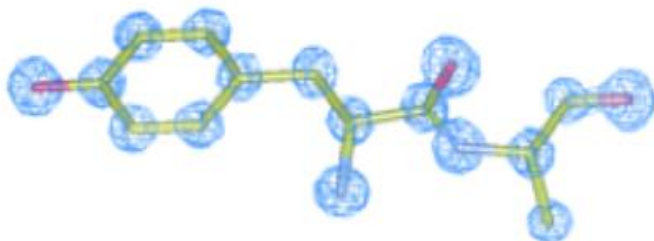
3.0 Å resolution,  $6.5\sigma$



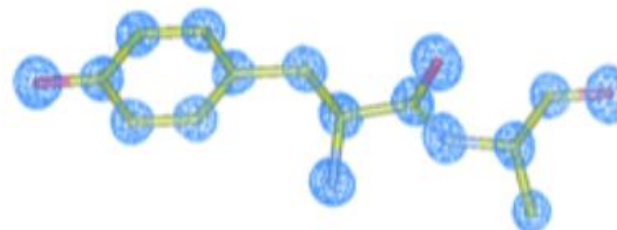
2.0 Å resolution,  $8.5\sigma$



1.5 Å resolution,  $10\sigma$

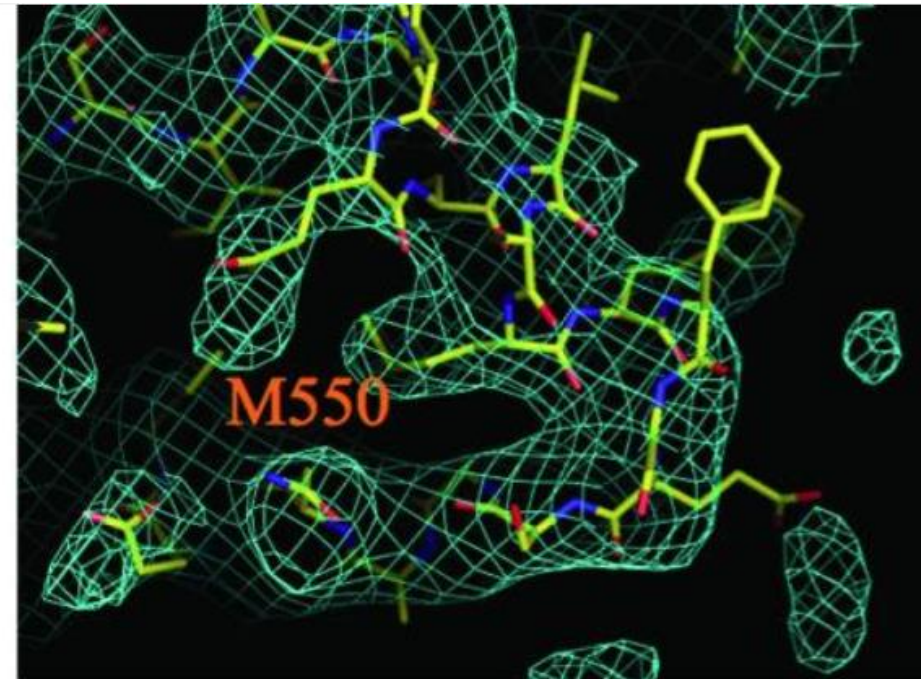
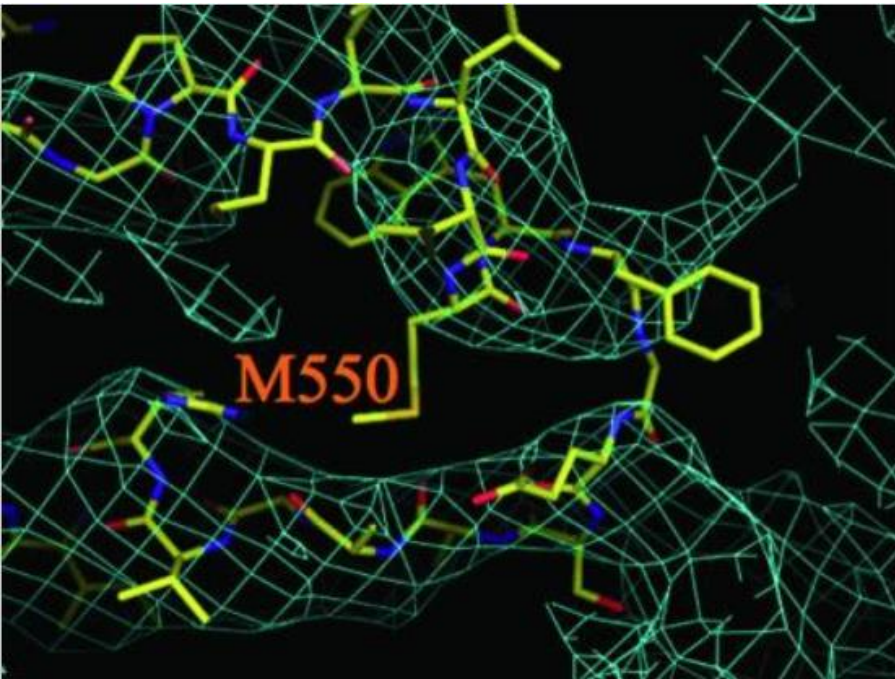


1.0 Å resolution,  $14\sigma$



0.6 Å resolution,  $12\sigma$

# Evaluating our trust of crystal structures



## 1308B (Code to continue)

Brunger AT, DeLaBarre B, Davies JM, Weis WI. X-ray structure determination at low resolution. *Acta Crystallographica Section D: Biological Crystallography*. 2009;65(Pt 2):128-133. doi:10.1107/S0907444908043795.

# Amino Acid Summary

- There are **20 amino acids used in proteins.**
- **Shared features:**
  - All are L-isomers (except glycine)
  - Amphoteric (can act as acid or base)
  - Each has an **amino, carboxylic acid,** and **side chain group** (except proline)
  - Each amino acid can be represented multiple ways, but the **space its electrons take up is always the same.**
- **Unique features:**
  - Cysteine can form **disulfide** bonds
  - Tyrosine and Tryptophan **absorb light** at 280 nm well
  - Serine, Threonine, Tyrosine have hydroxyl groups which **can be modified** to activate/inactivate protein function
  - Glycine and proline each have **unique flexibility or lack of it.**
- Amino acids can be titrated, and their charge state is important.
- **Amino acid shapes constrain bond angles.**



# Next Lecture:

- See how amino acids joined in peptides form repeated elements in protein structure.
- Look at determinants of a protein's final structure.

