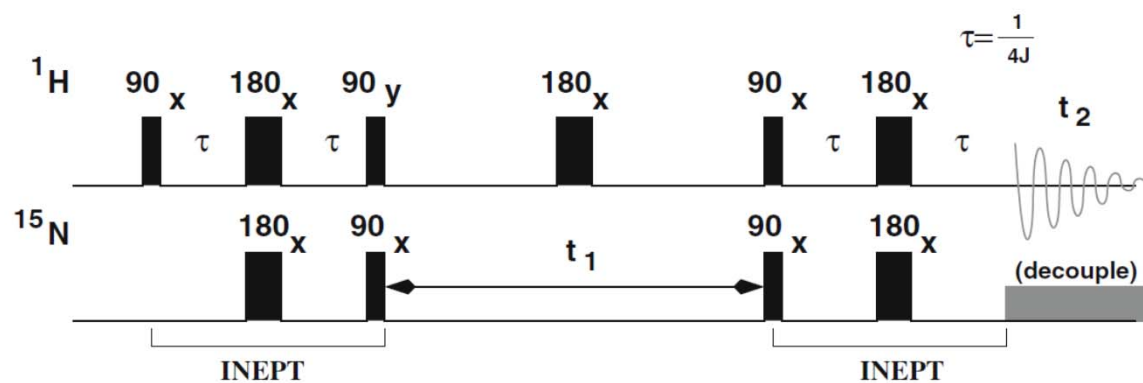
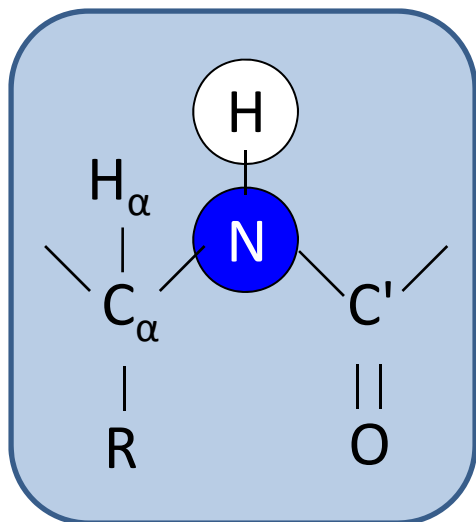
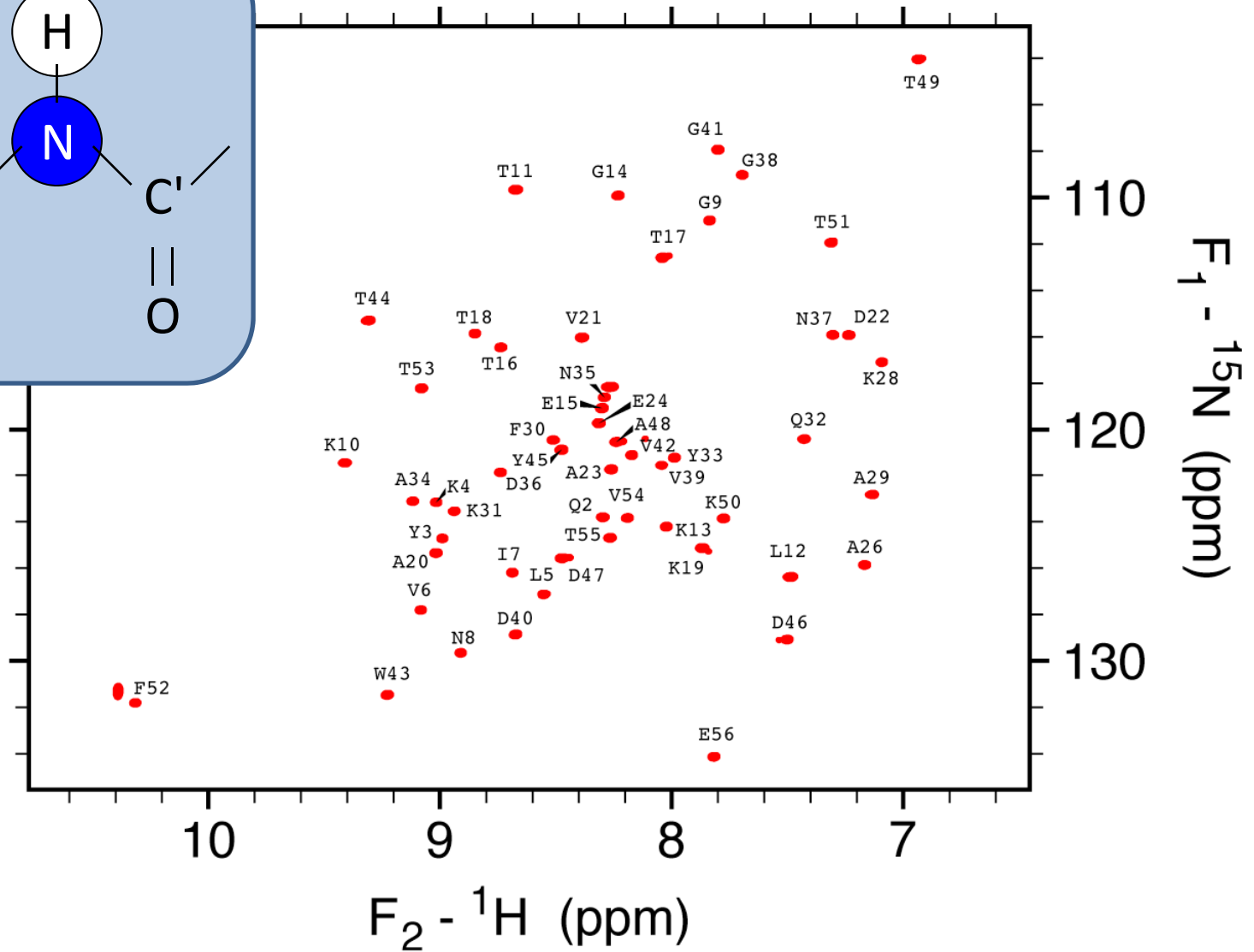
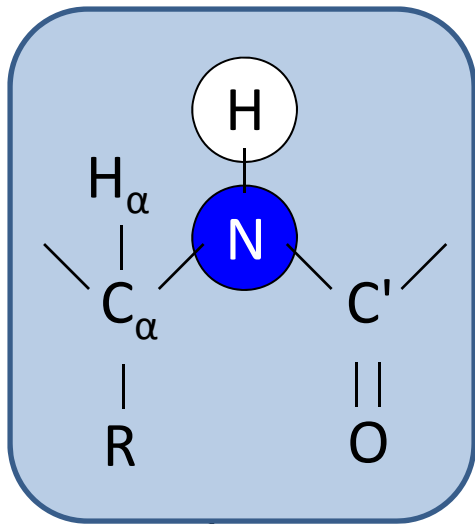


^{15}N HSQC

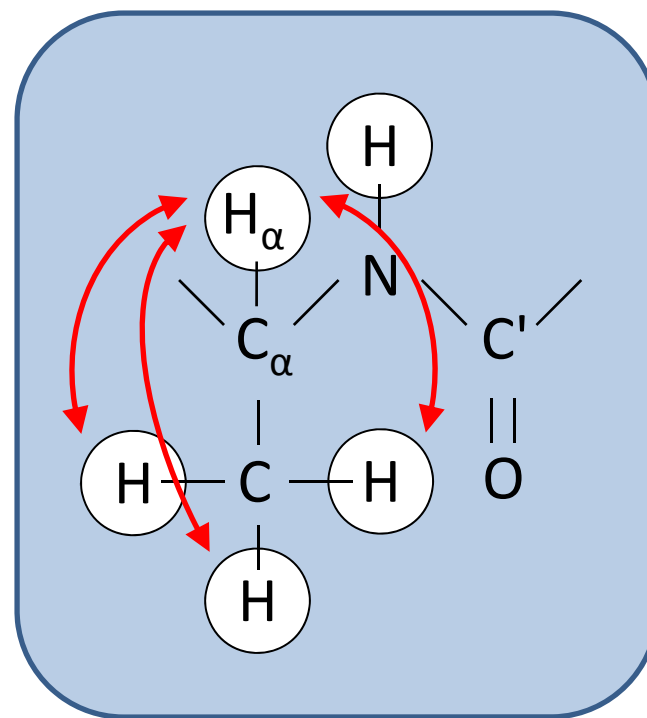
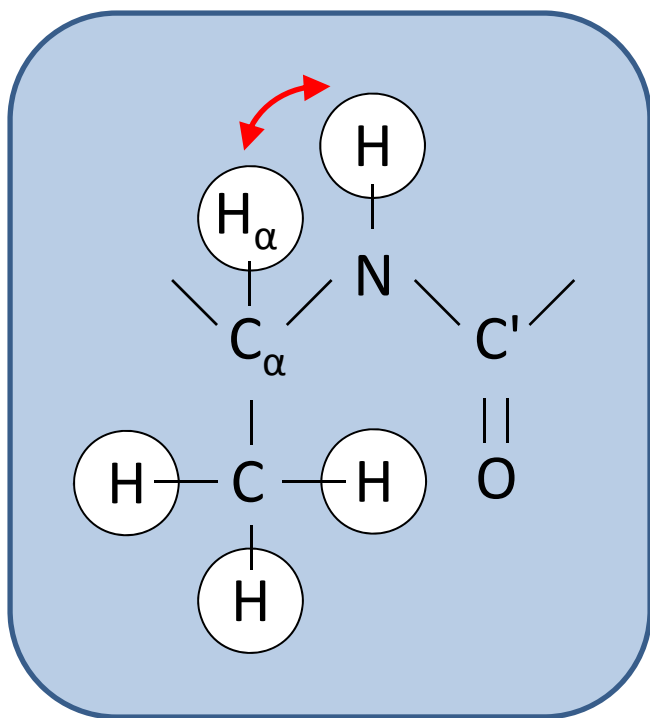


From *Fundamentals of Protein NMR Spectroscopy*
 Rule & Hitchens, Chapt. 10, p. 205

2D ^{15}N HSQC: GB3

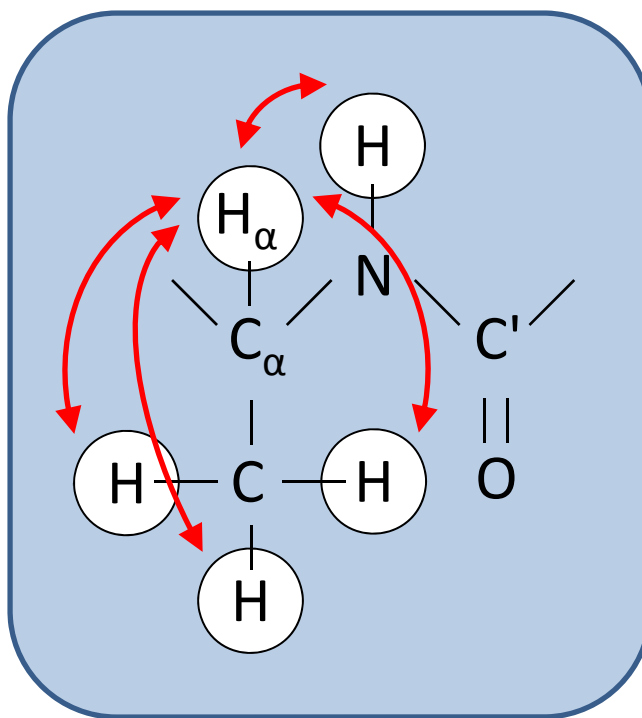


^1H COSY



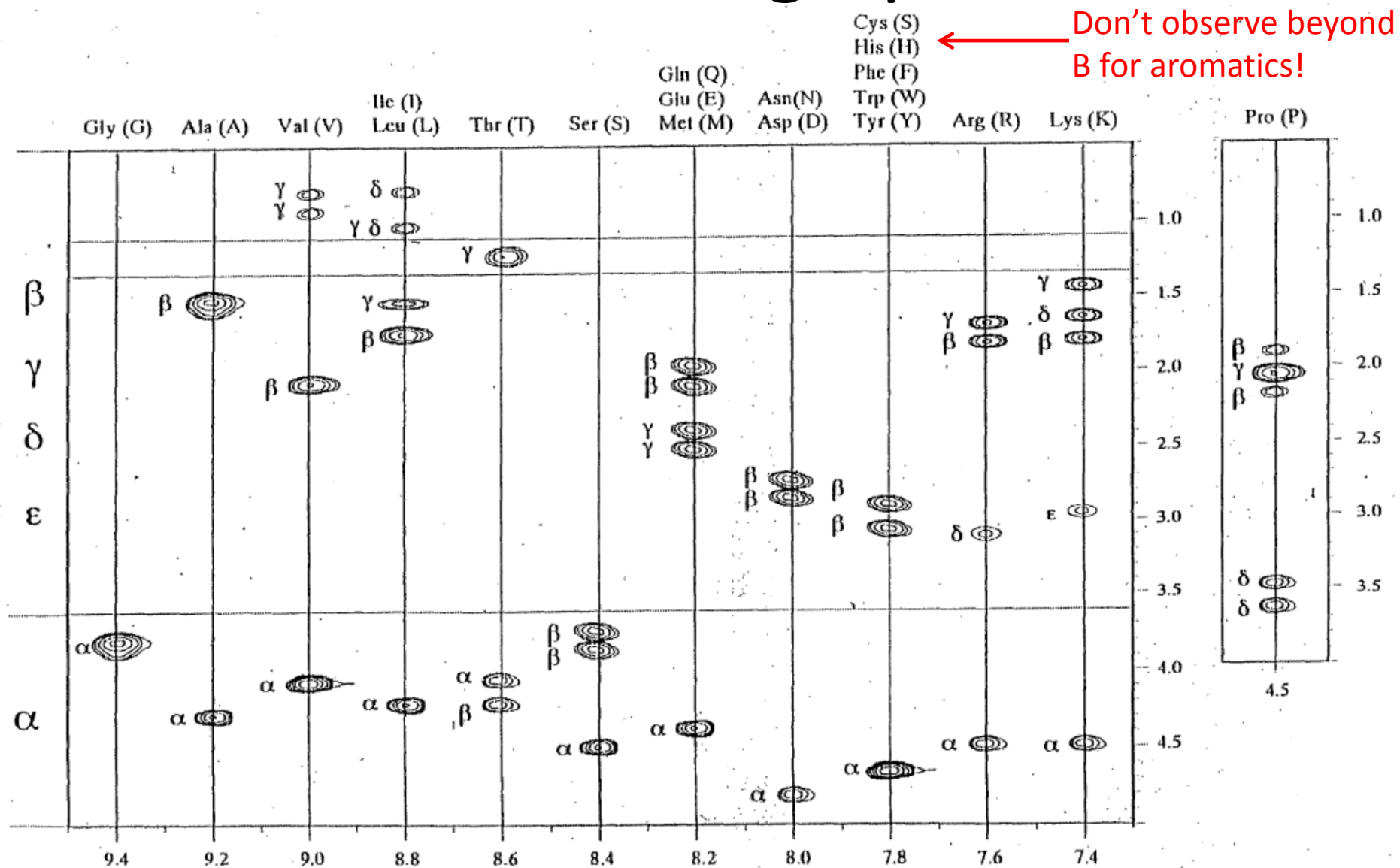
- Only one “hop” at a time between nuclei observed.
- Example: $\text{H}_\text{N} \leftrightarrow \text{H}_\alpha$, $\text{H}_\beta \leftrightarrow \text{H}_\alpha$ (but not both!)

^1H TOCSY

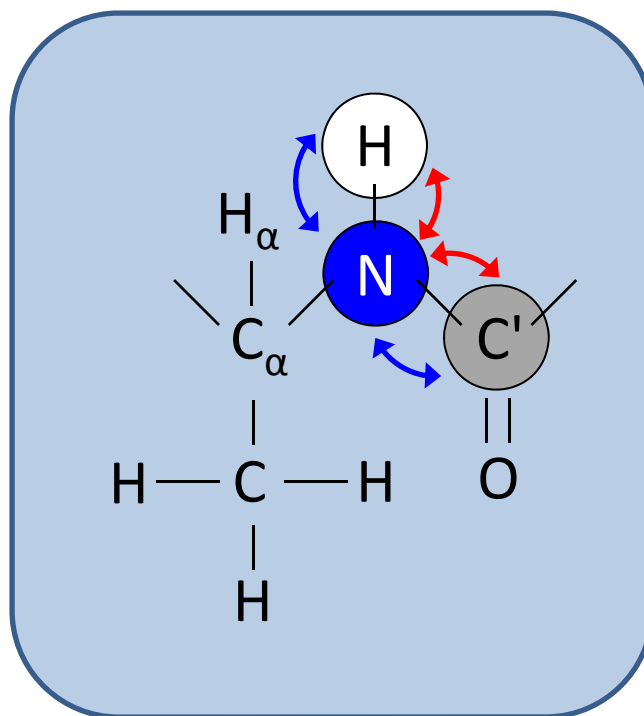


- Magnetization “hops” from nucleus to nucleus
- Example: Crosspeaks observed between H_N and: H_α , H_β , H_γ (gets weaker for more hops)

^1H TOCSY Fingerprints



3D HNCO



Out: Red transfers occur during preparation period

Back: Blue transfers occur between chemical shift labeling steps on the way back

- See peaks at H_N , N, and C'_{i-1} (i.e. the C' of the prior residue)
- Requires ^{13}C and ^{15}N labeled protein

Summary

- NOESY allows transfer of magnetization through space; distance geometry can reconstruct the structure
- COSY and TOCSY correlate homonuclear spins using J-coupling through one hop or multiple hops
- HSQC uses J-coupling to correlate bonded, heteronuclear spins
- HNCO transfers magnetization “out and back” to previous residue’s C’.
- Combining multidimensional experiments allows assignment of entire proteins