

C8953

NMR strukturní analýza seminář

TOCSY, ROESY, Introduction to heteronuclear correlations

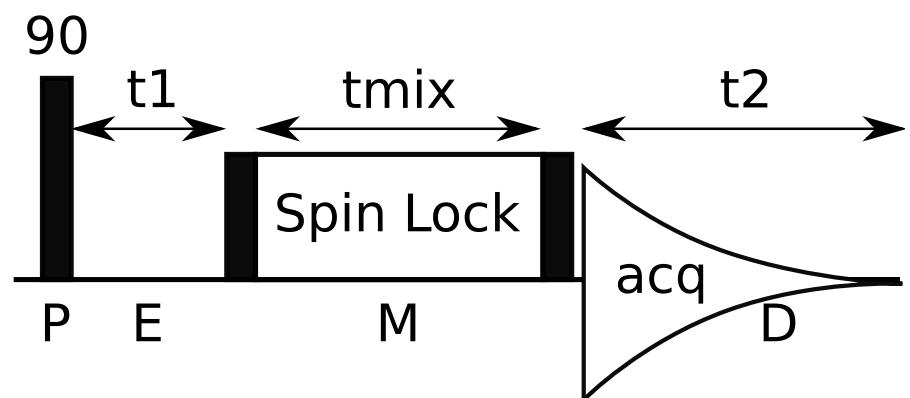
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TOCSY & ROESY

spin lock - isotropic mixing

- ▶ series of 180° pulses
- ▶ various pulse schemes (MLEV-17, DIPSI,...)
- ▶ precession around $B_{Spinlock} < B_0 \rightarrow$ "locking" spins in transversal plane (xy)
- ▶ TOCSY: lower power, offset in the center of spectrum
- ▶ ROESY: higher power, offset on edge
- ▶ crosstalk (ROE in TOCSY, J in ROESY)

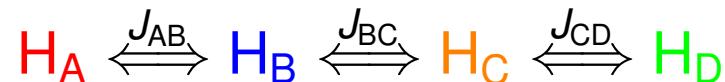


TOCSY (TOtal Correlation SpectroscopY)

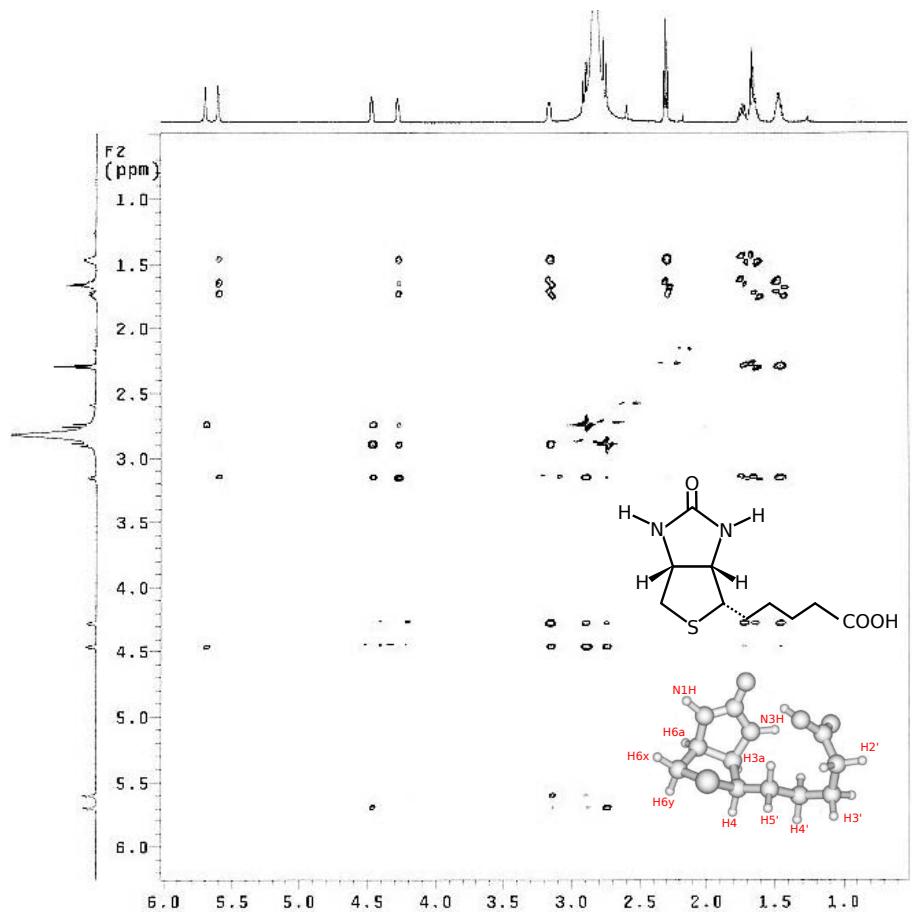
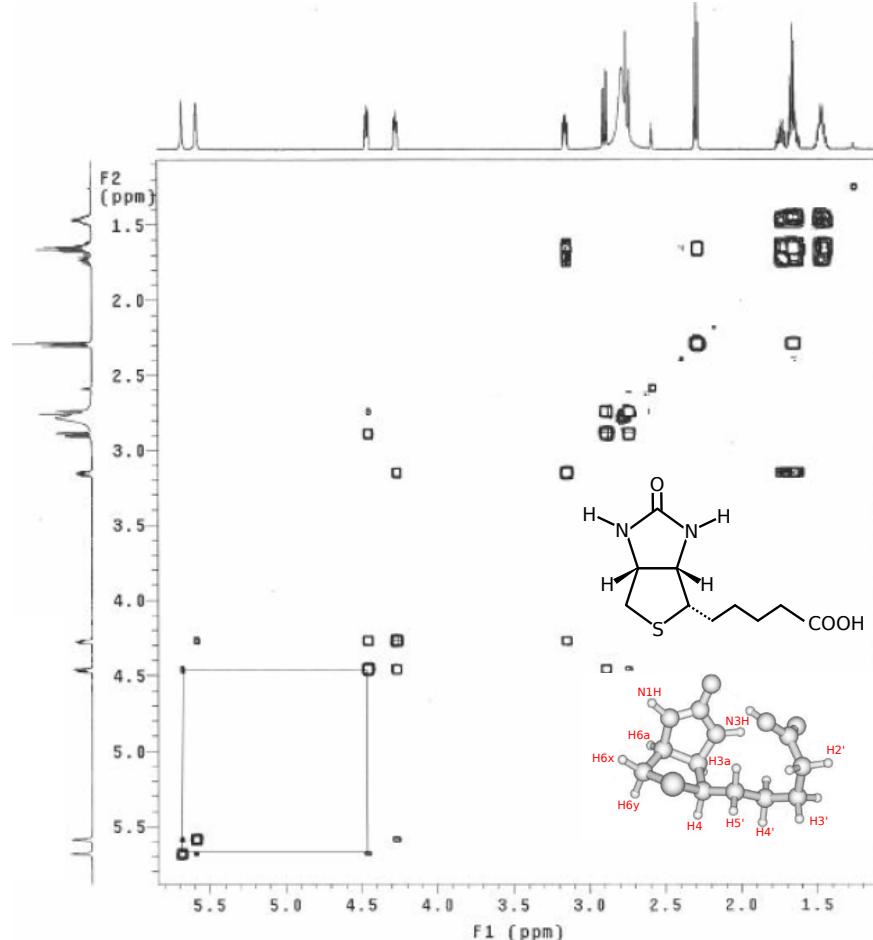
HOHAHA (HOmonuclear HArtmann-HAhn)

correlation based on J , like COSY

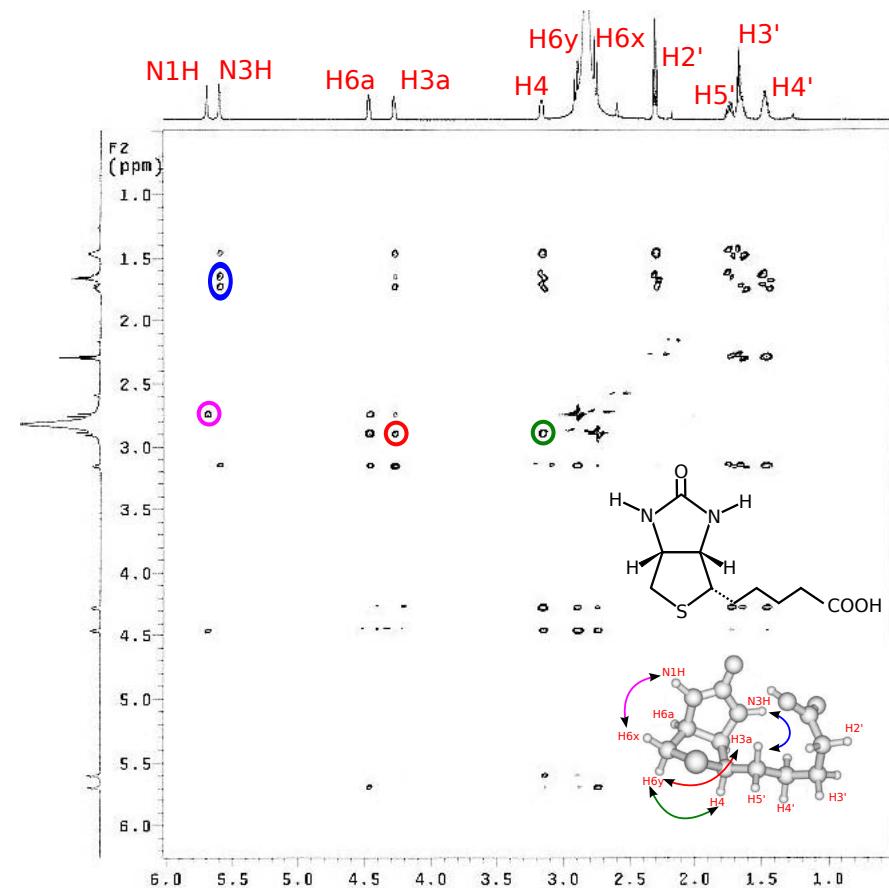
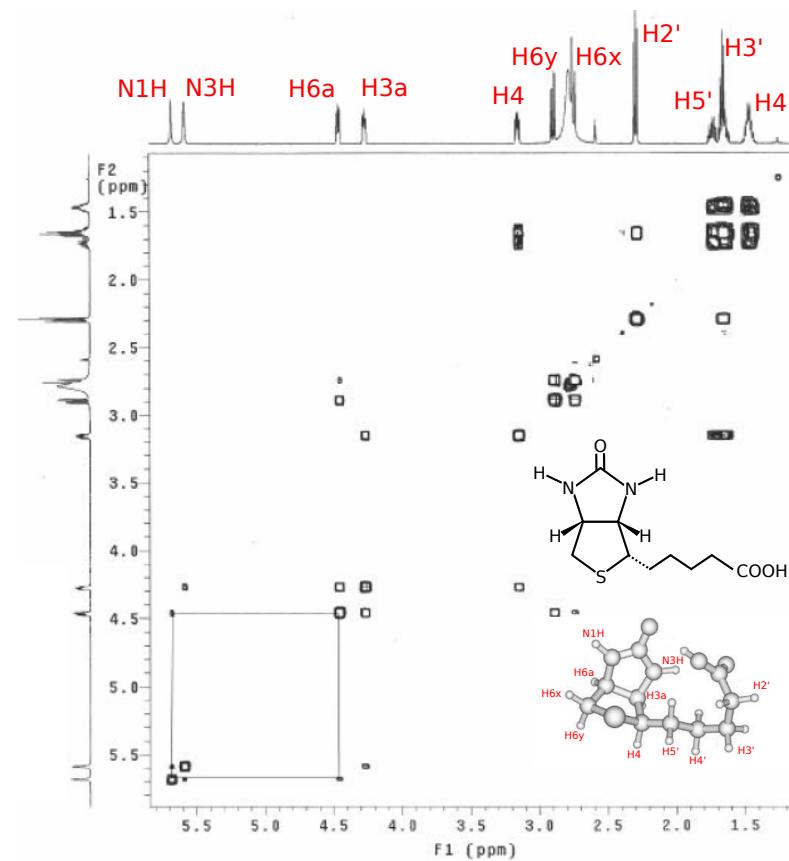
- ▶ correlate mutually *all* protons within a spin system
- ▶ $\tau_{\text{mix}} \approx 20 - 120 \text{ ms}$
- ▶ intensity depends on τ_{mix} and J value



Biotin - COSY vs. ROESY (G. T. Crisp and Yu-Lin Jiang, 2001)



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HMQC (Heteronuclear Multiple Quantum Correlation)

HSQC (Heteronuclear Single Quantum Correlation)

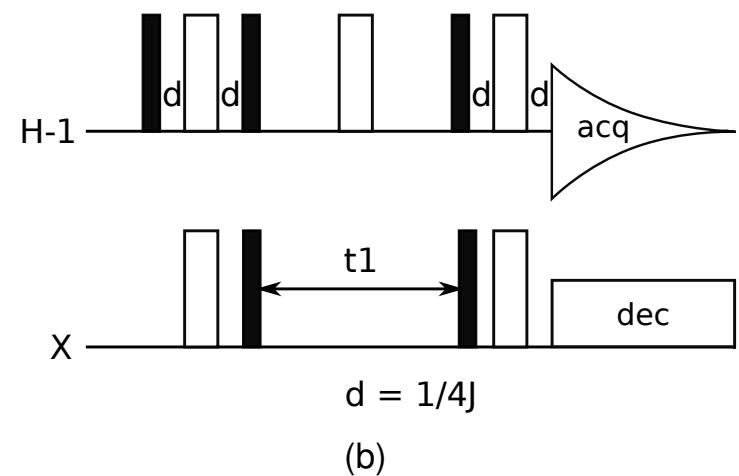
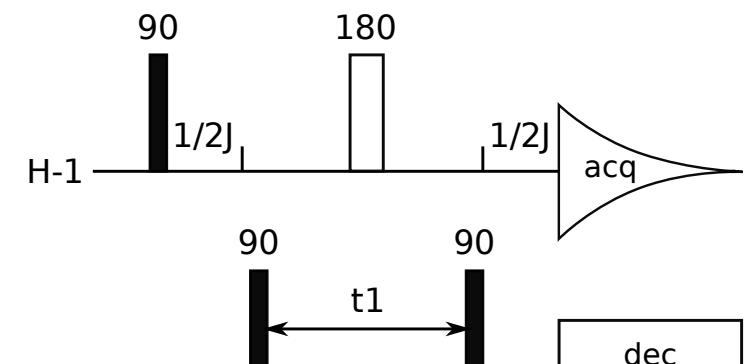
correlate ^1H -X ($\text{X} = ^{13}\text{C}, ^{15}\text{N}, \dots$) based on
 $^1J_{\text{HX}}$

HMQC (a)

- + more robust experiment
- + change of parameters - HMBC
- lower sensitivity and worse resolution

HSQC (b)

- + better resolution, sensitivity
- + part of more complex multidimensional experiments
- less robust



Practical notes ^1H -X HSQC

- ▶ resolution of overlaps
- ▶ routine experiments to control biomolecular sample
- ▶ easy identification of geminal protons
- ▶ indirect determination of protons bonded to NMR inactive heteroatom
- ▶ heteronuclear correlation \Rightarrow no diagonal crosspeak, no symmetry
- ▶ X decoupled during acquisition \Rightarrow singlet crosspeak

HMBC(Heteronuclear Multiple-Bond Correlation)

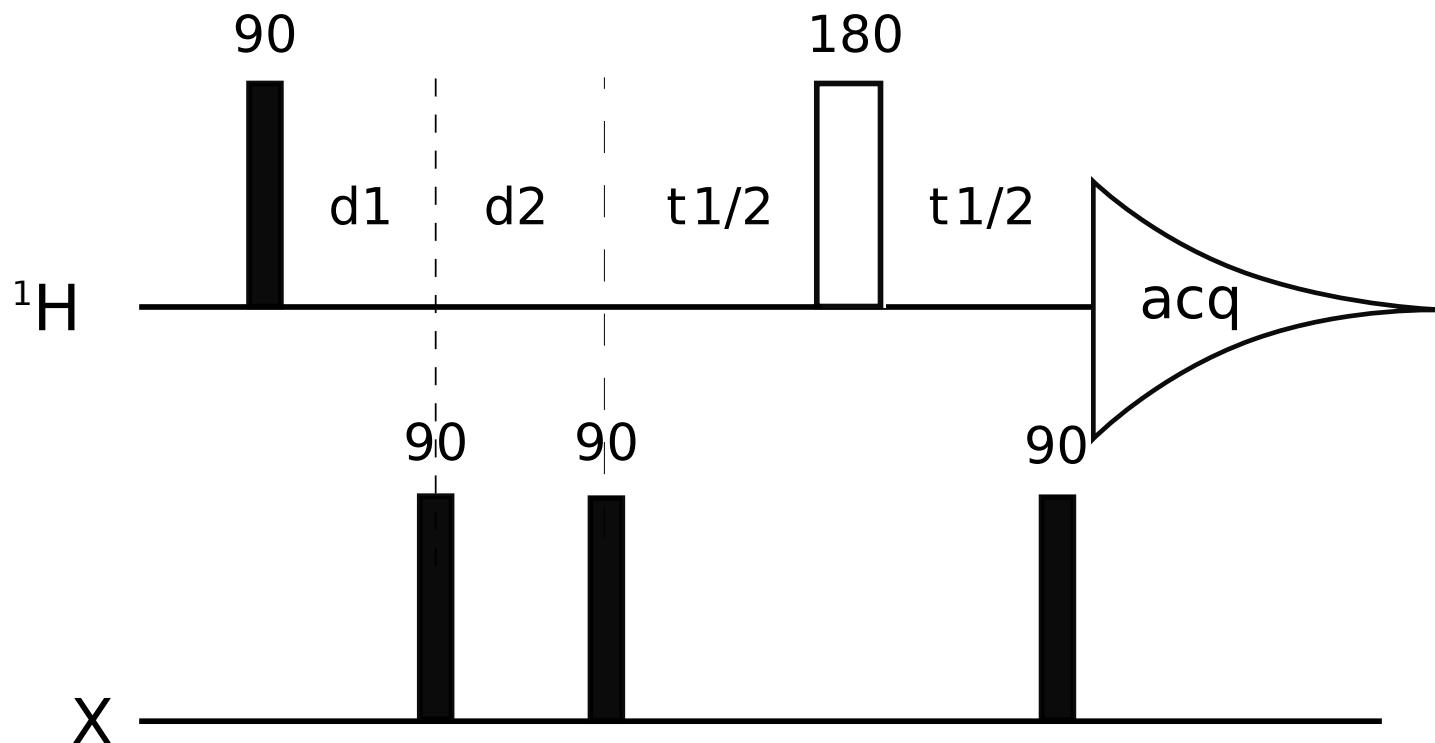
heteronuclear correlation based on long-range H-X spin-spin interaction($^nJ_{HX}$, $n>1$)

- ▶ utilizes polarization transfer from H through 2-5 bonds on heteroatom (^{13}C , ^{15}N)
- ▶ allows to detect quaternary heteroatoms (Cq) or connect signals among isolated spin systems

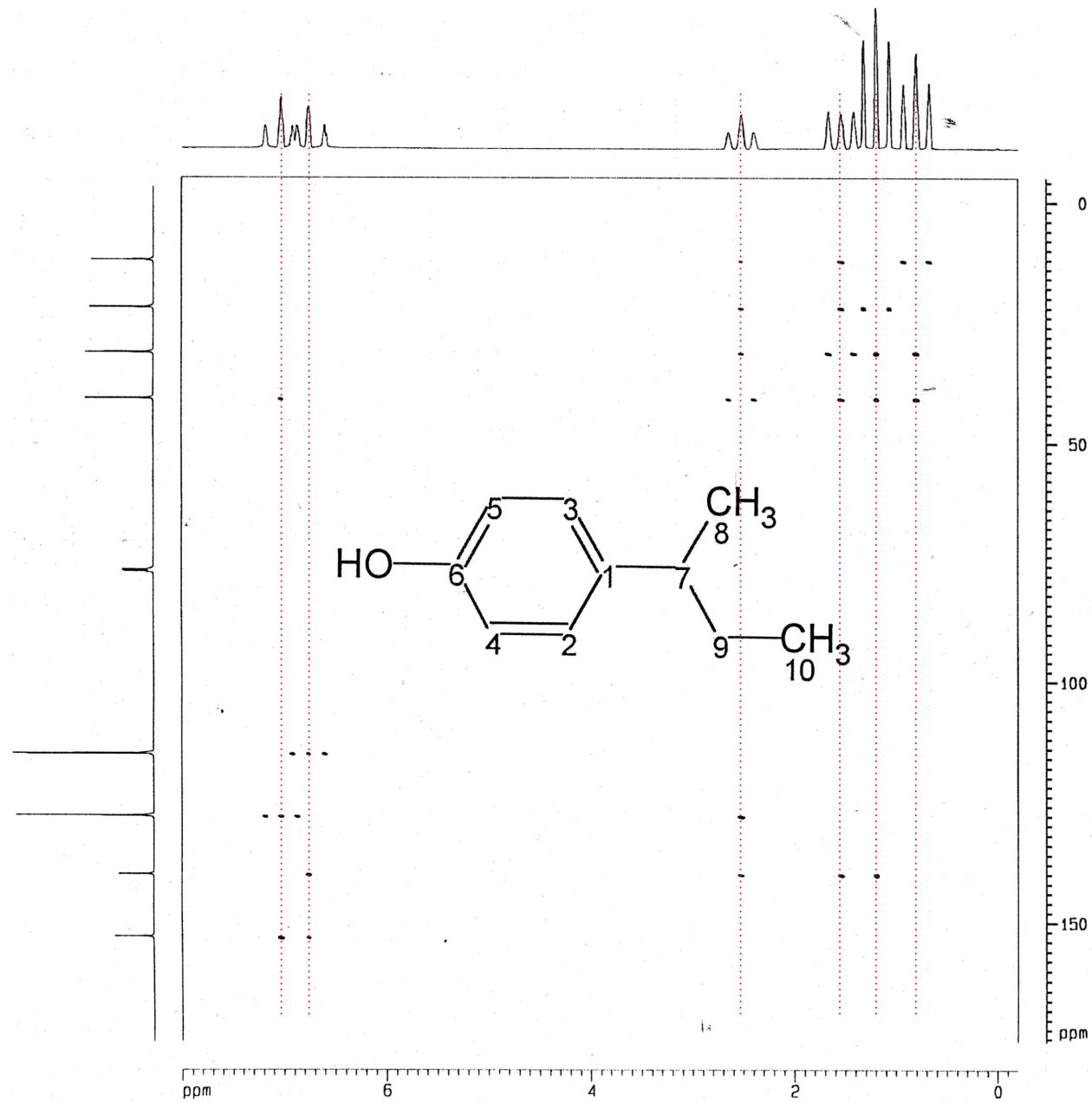
HMBC

correct settings of d1, d2 fo evolution of J -coupling necessary

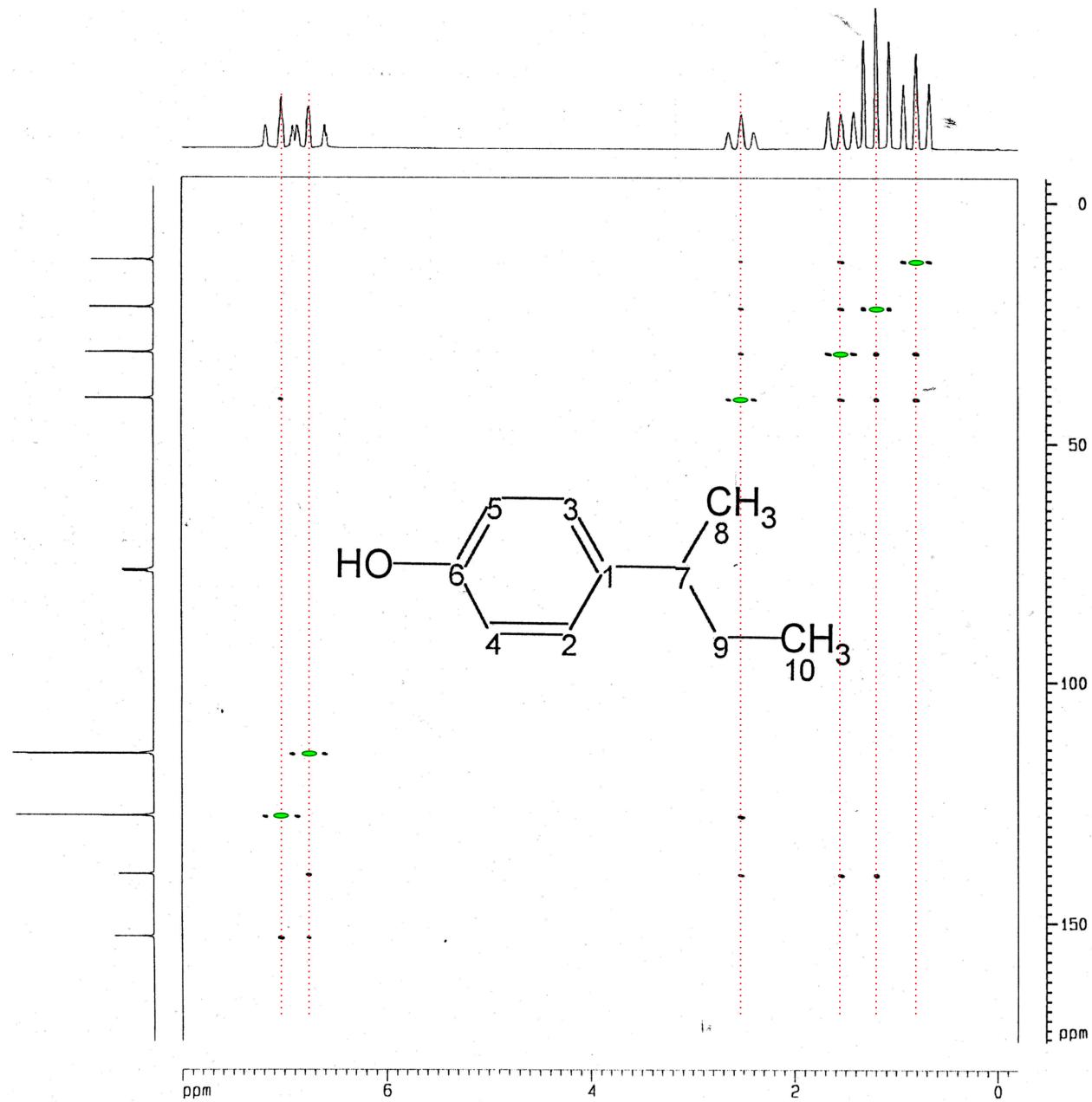
- ▶ $d1=1/2*^1J_{C-H}$ - (120-180 Hz)
- ▶ $d2=1/2*^{2-5}J_{C-H}$ - (3-12 Hz)



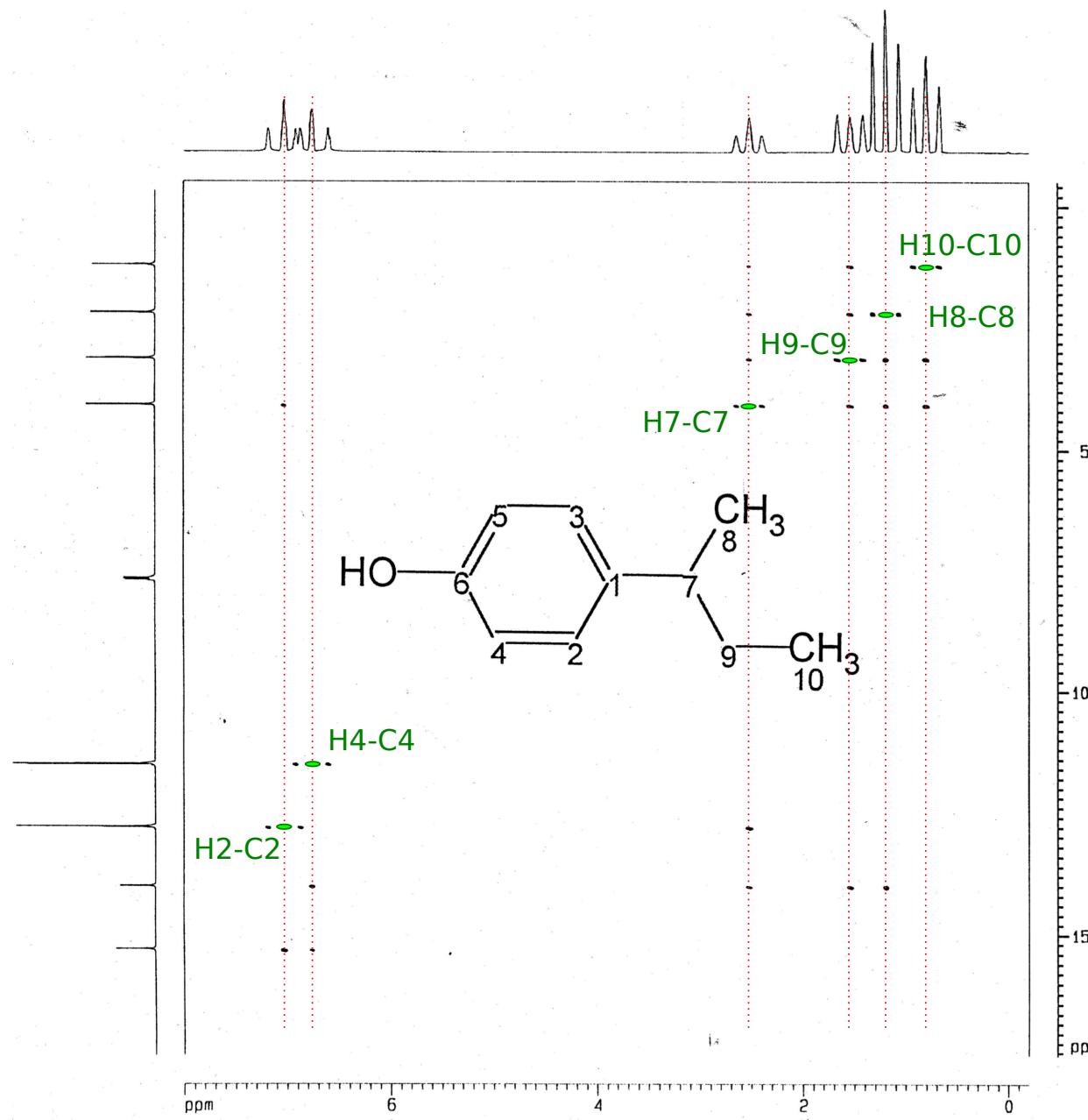
¹H-¹³C HMBC



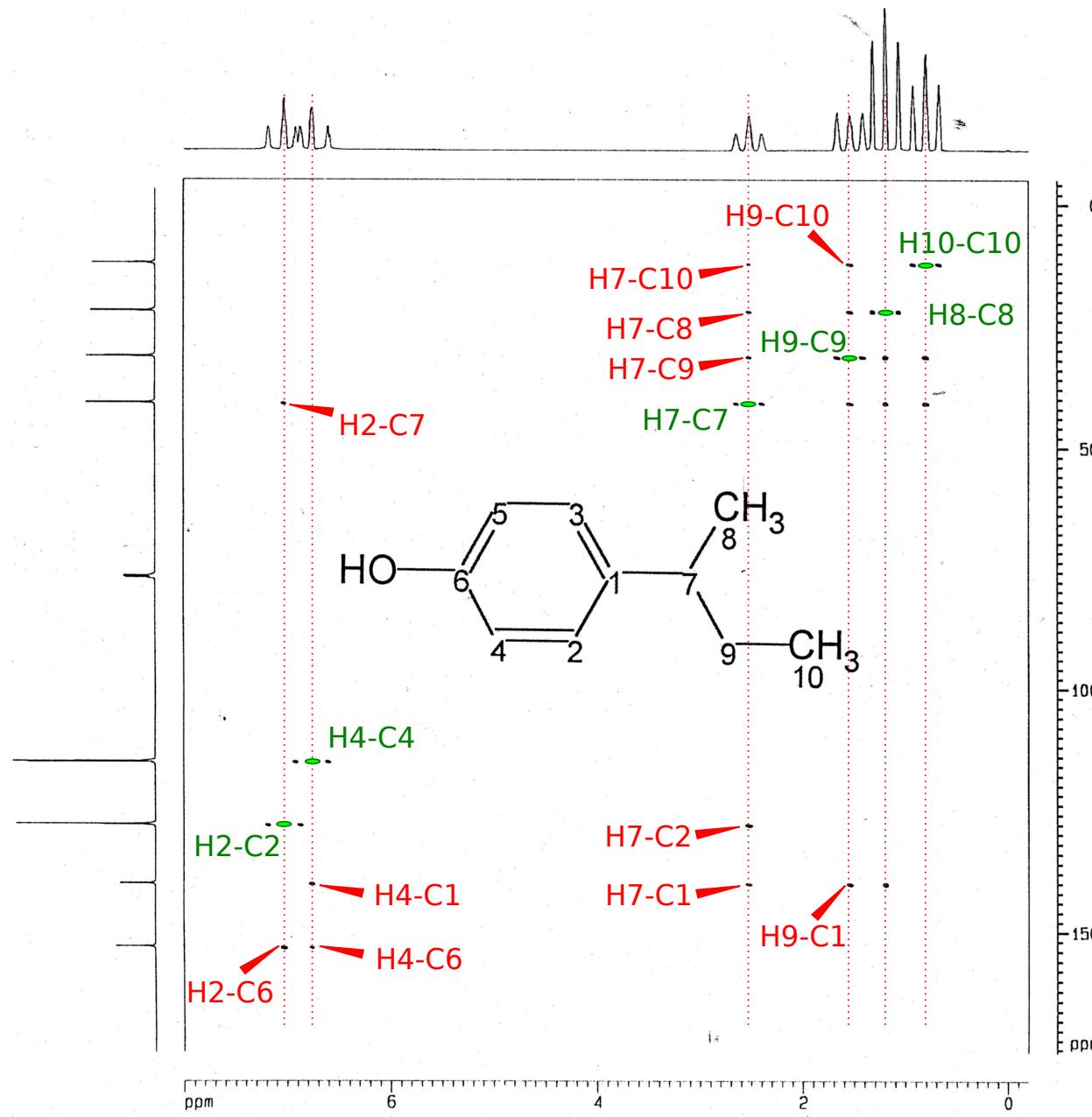
^1H - ^{13}C HMBC



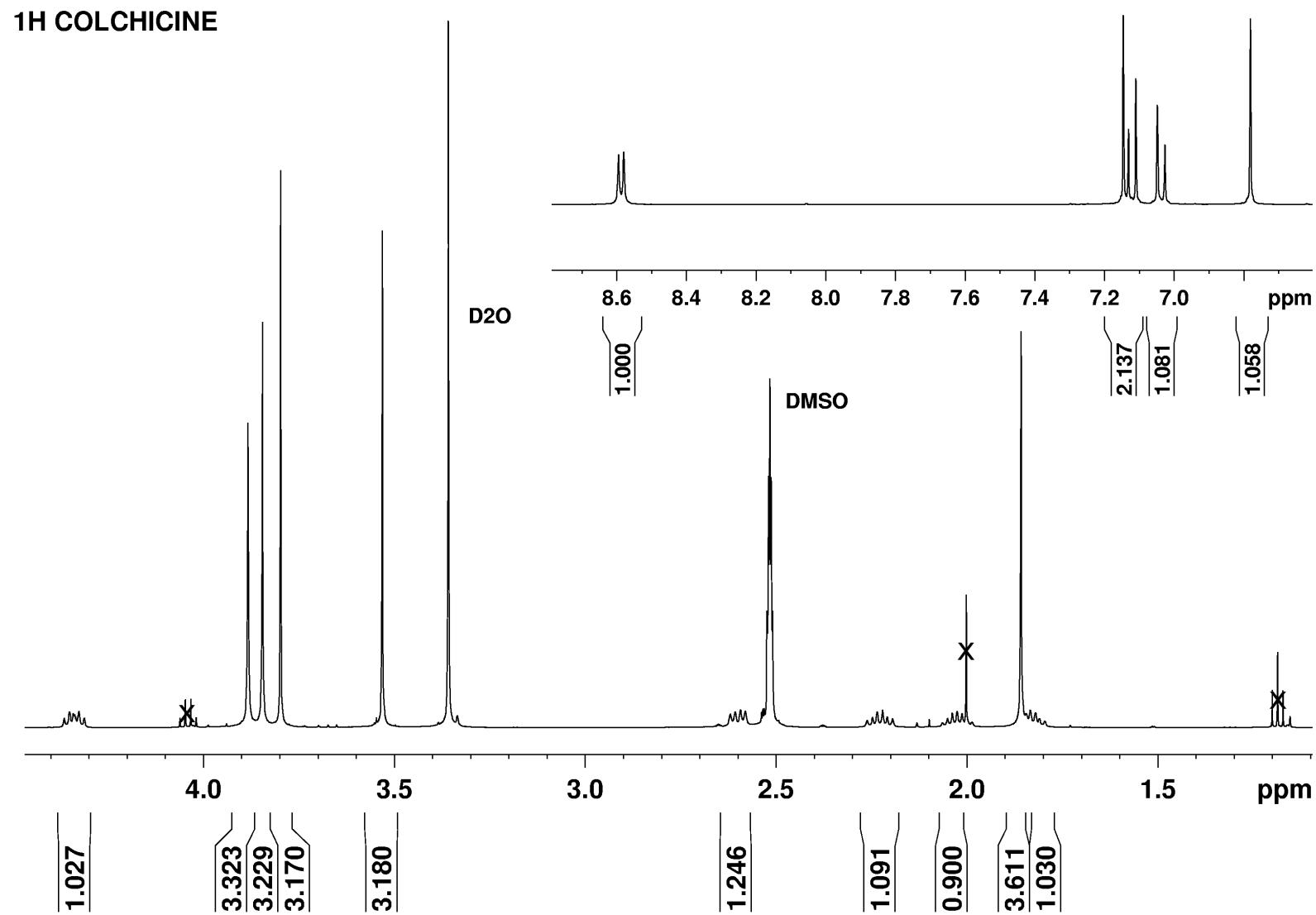
^1H - ^{13}C HMBC



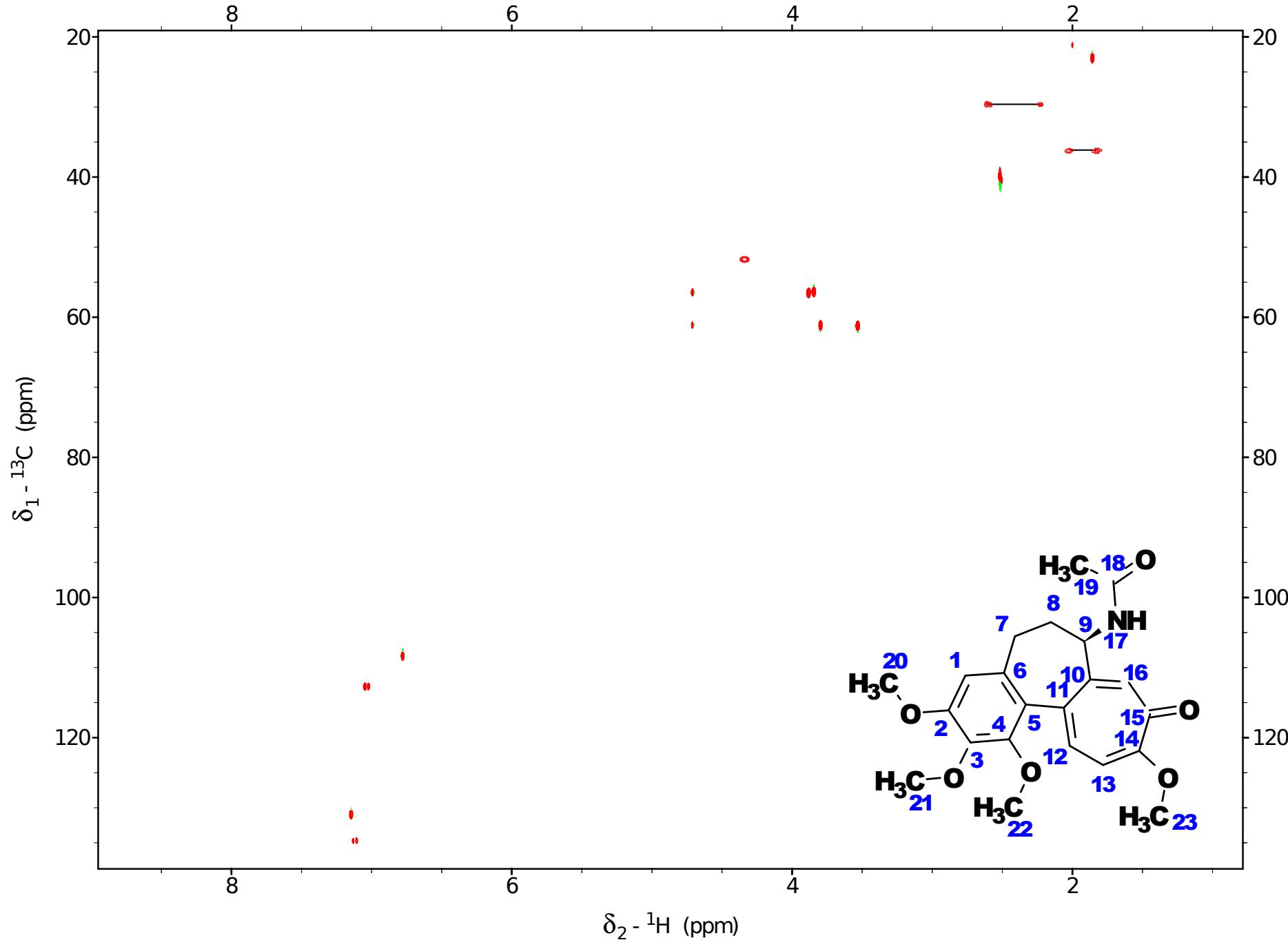
^1H - ^{13}C HMBC



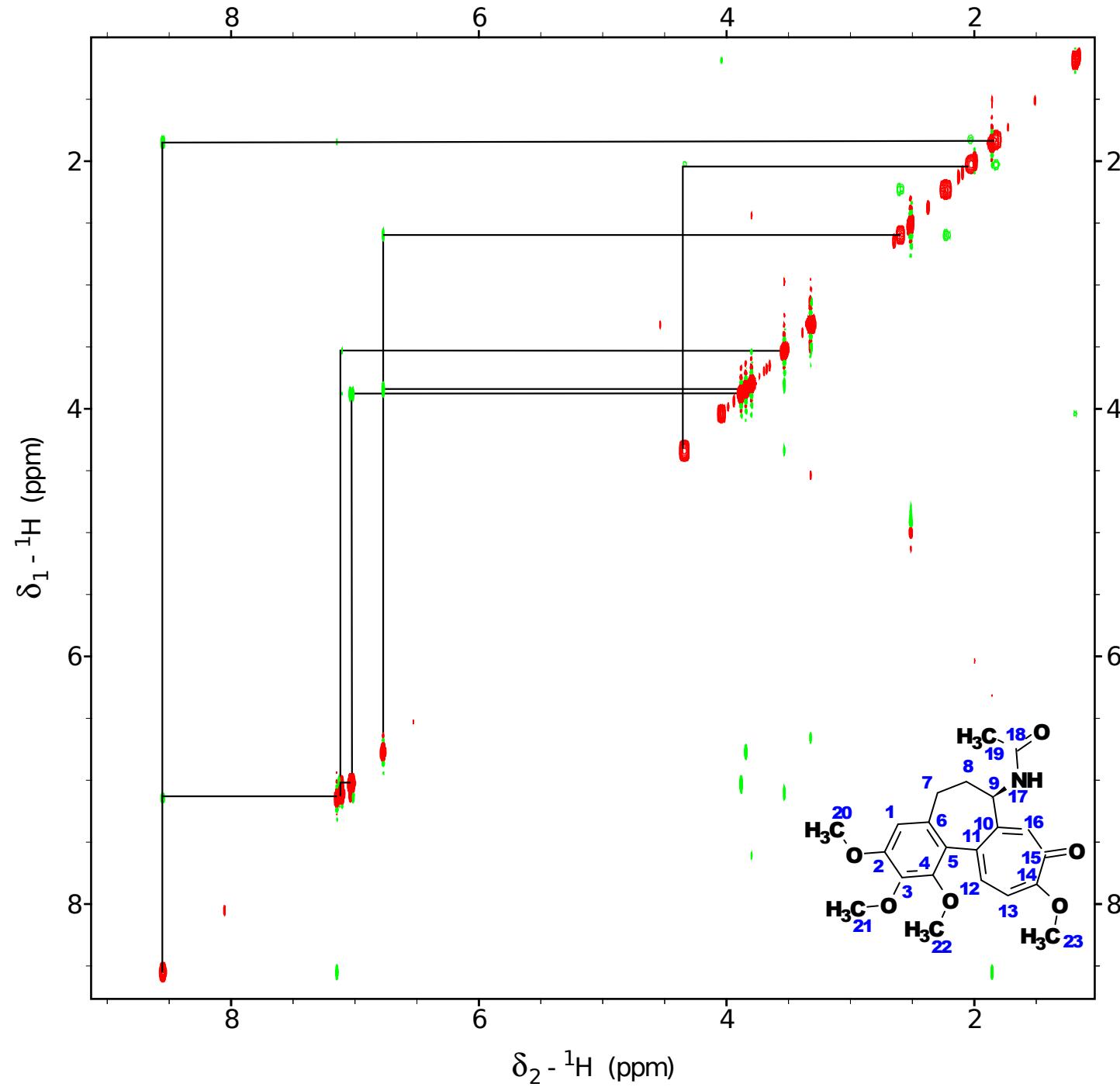
Colchicine 1D-¹H



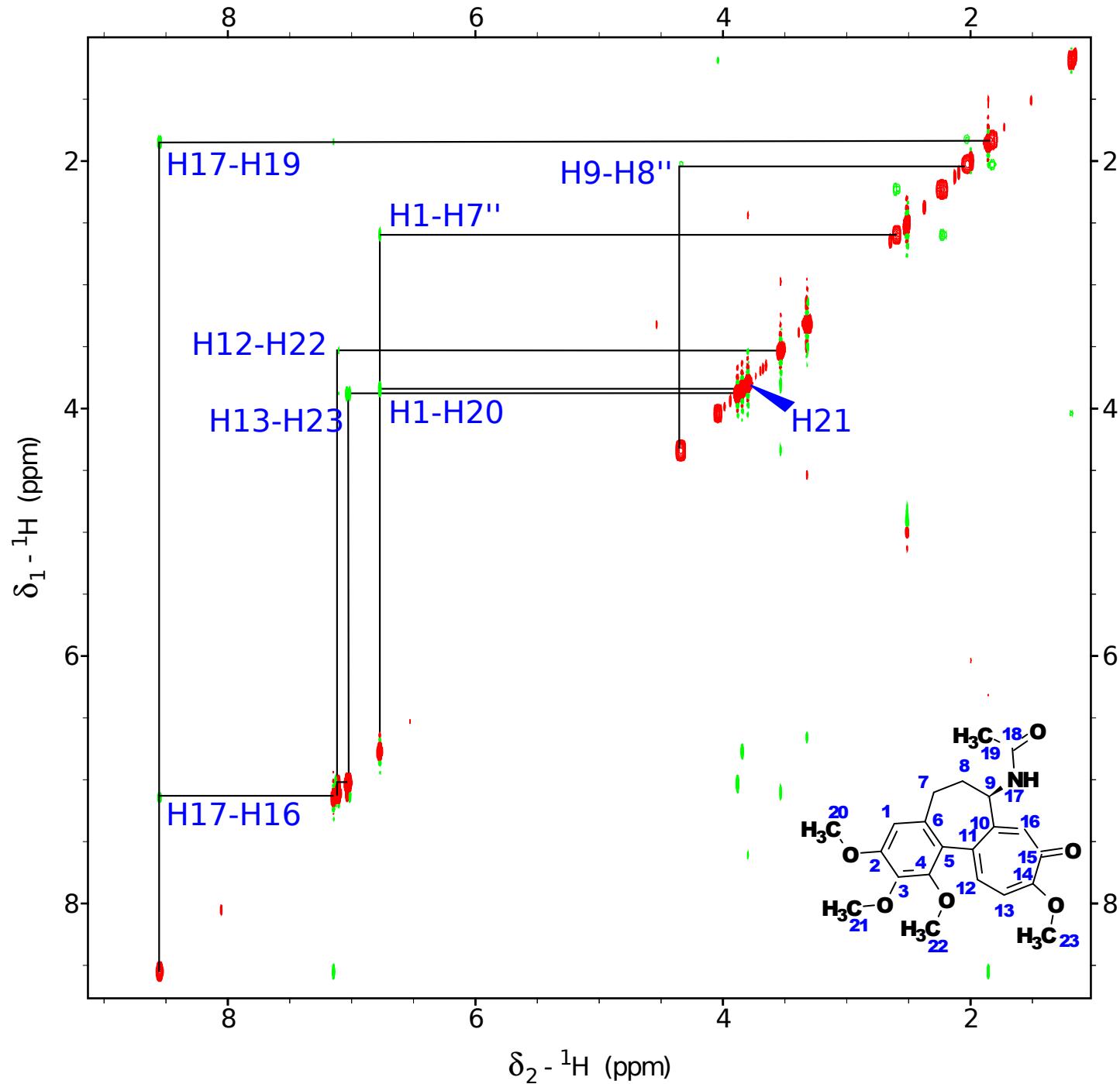
Colchicine - ^1H - ^{13}C HSQC



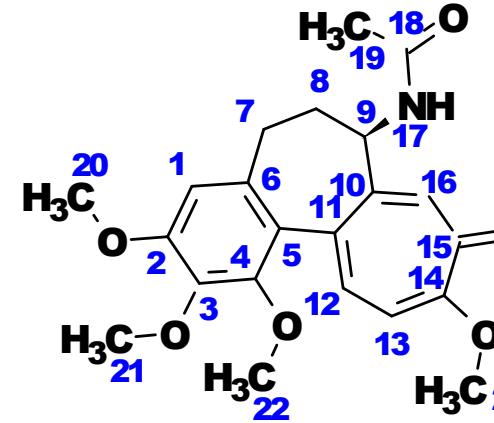
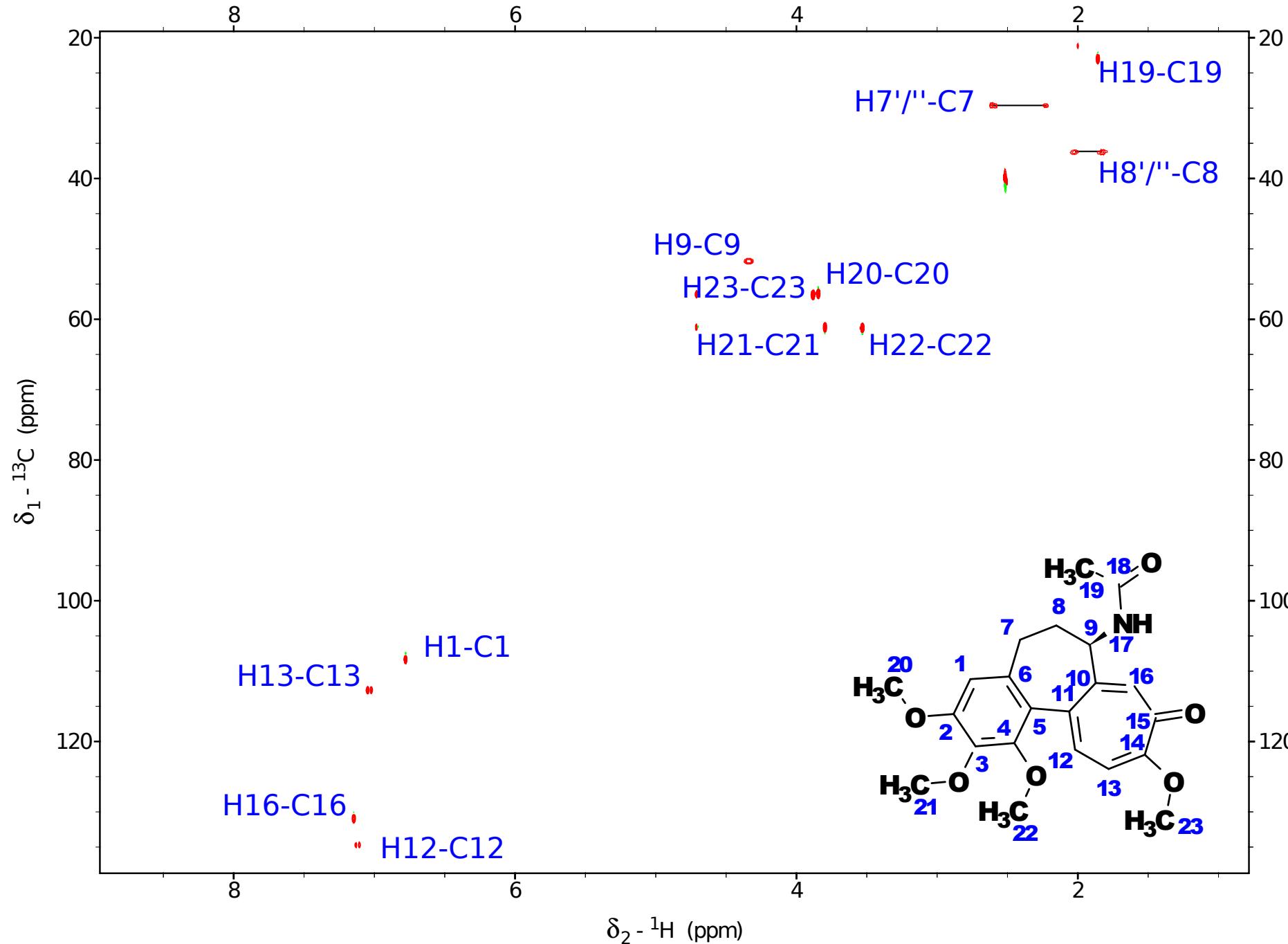
Colchicine - NOESY



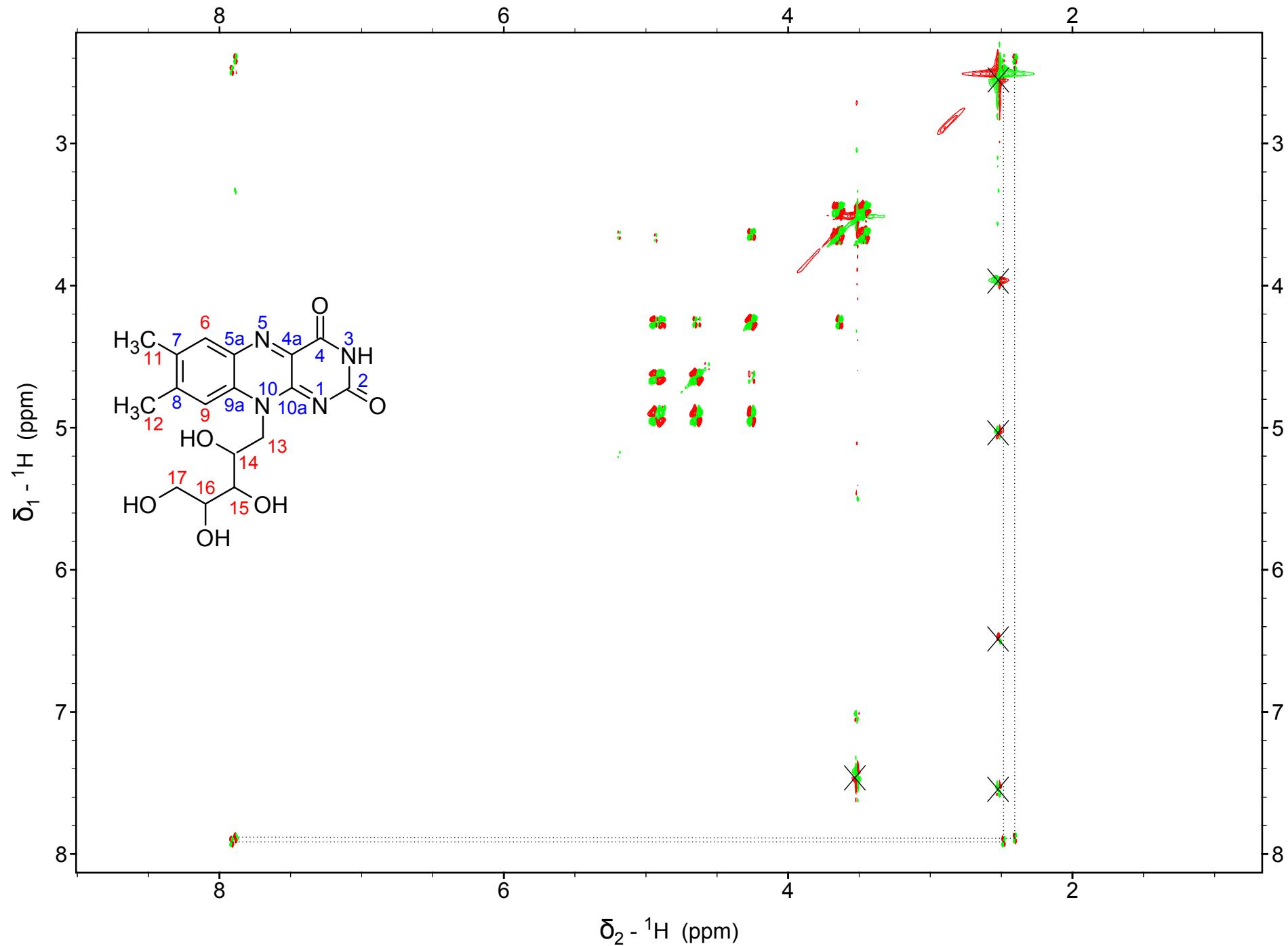
Colchicine - NOESY



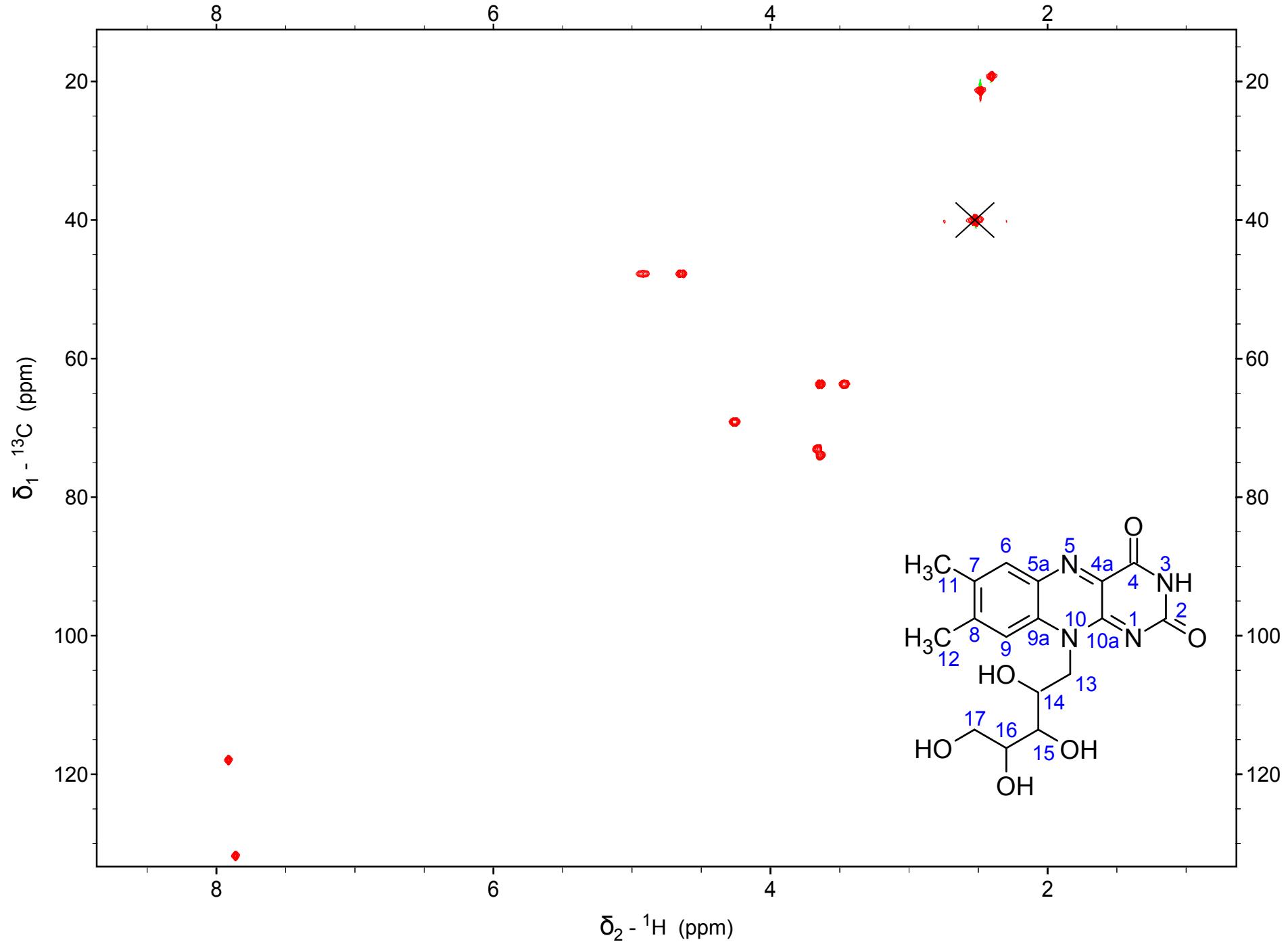
Colchicine - ^1H - ^{13}C HSQC



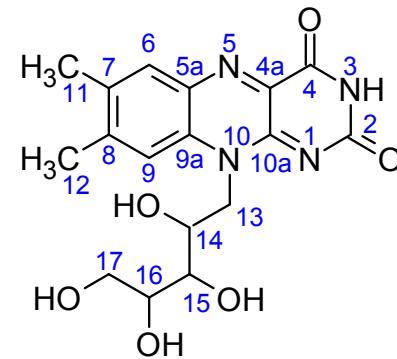
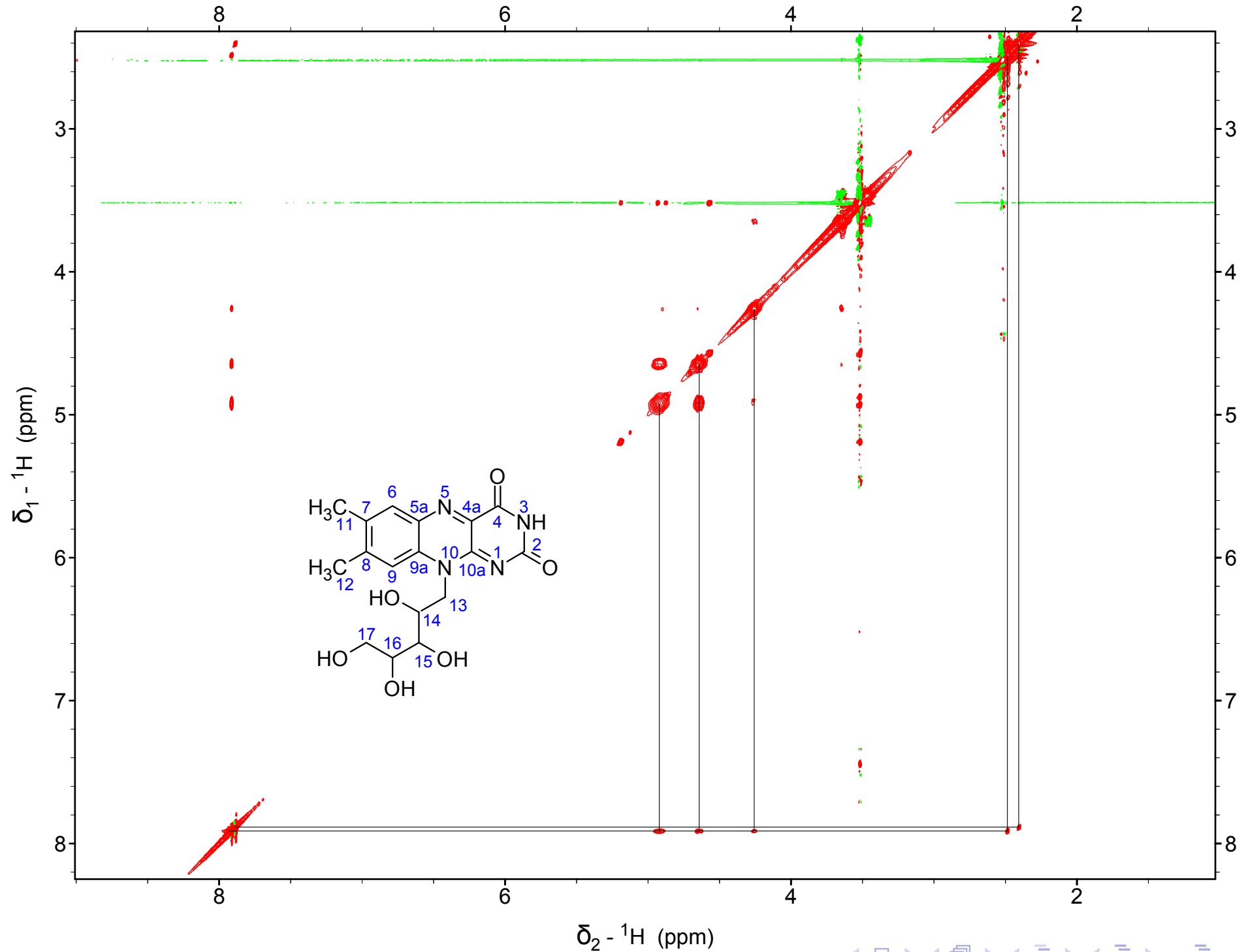
Riboflavin: DQF-COSY



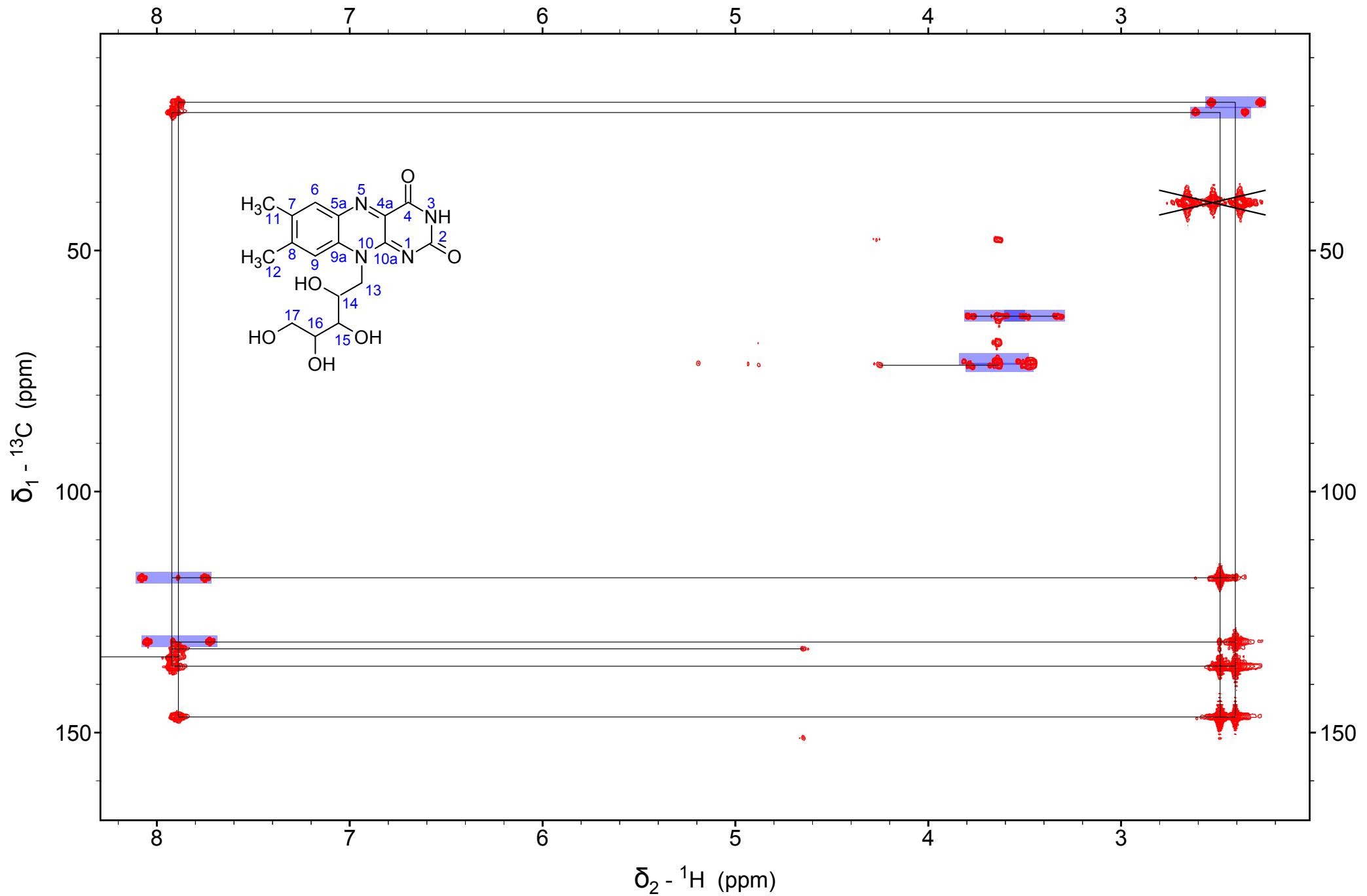
Riboflavin: ^1H - ^{13}C HSQC



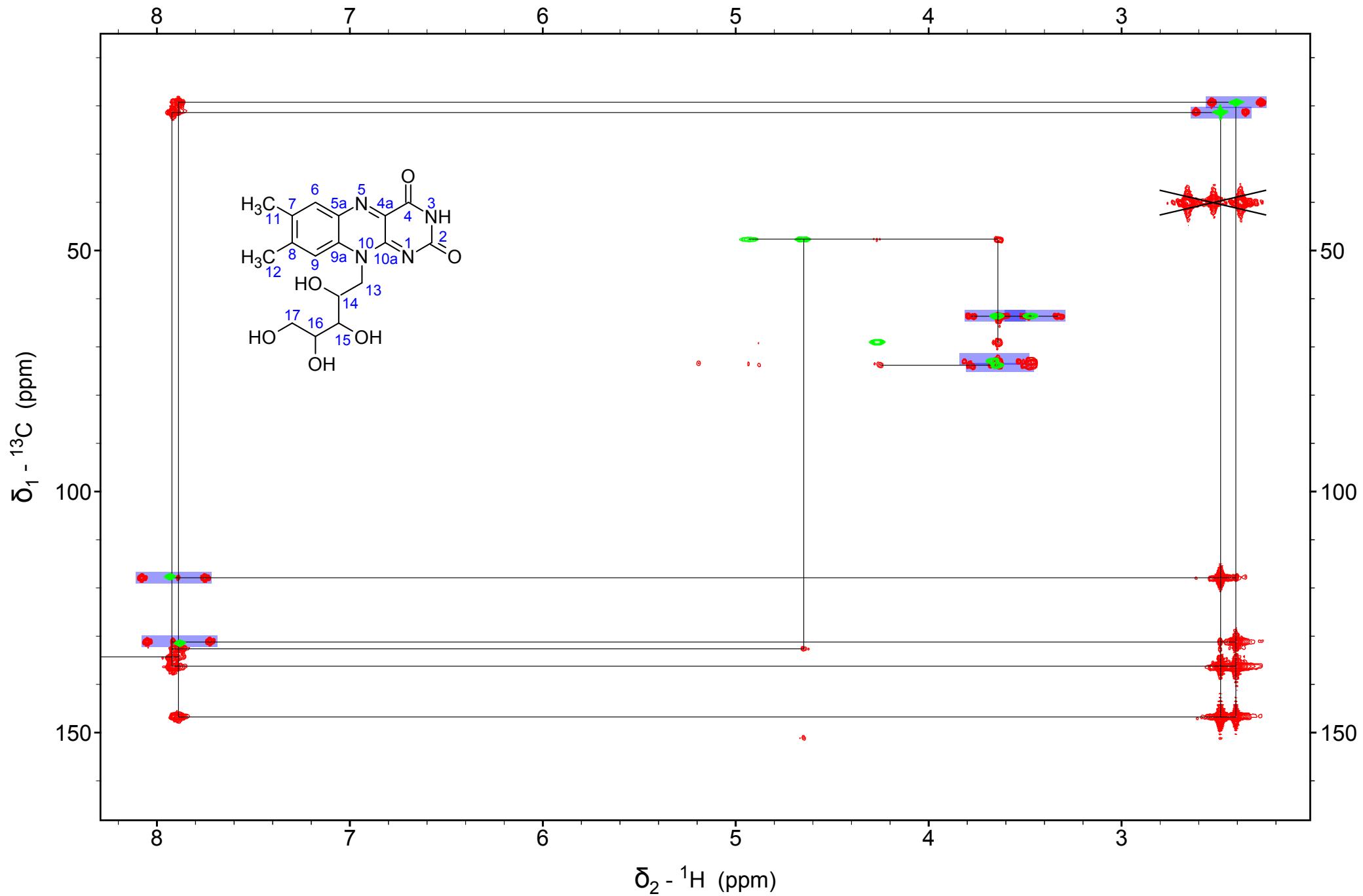
Riboflavin: NOESY



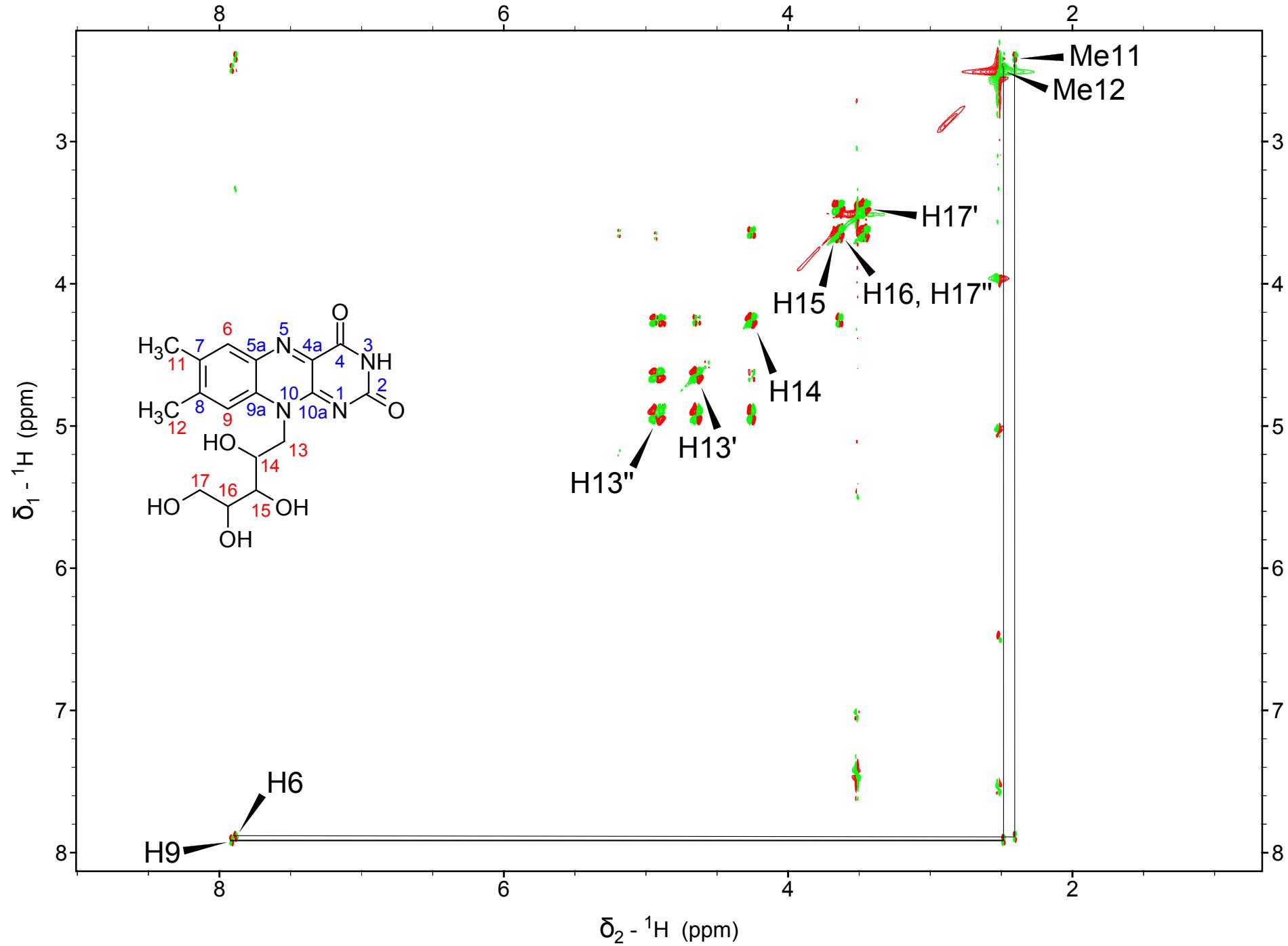
Riboflavin: ^1H - ^{13}C HMBC



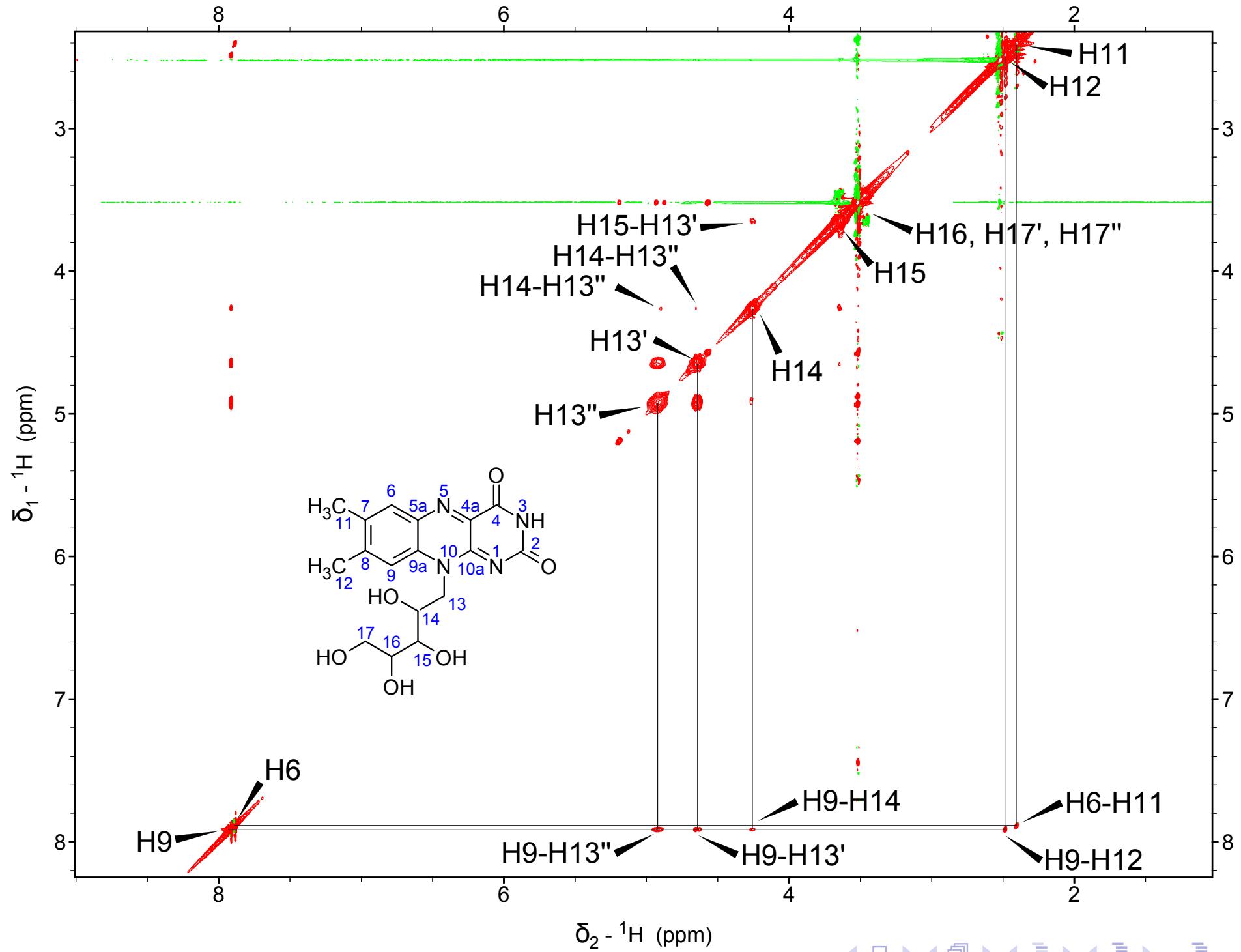
Riboflavin: ^1H - ^{13}C HMBC + HSQC



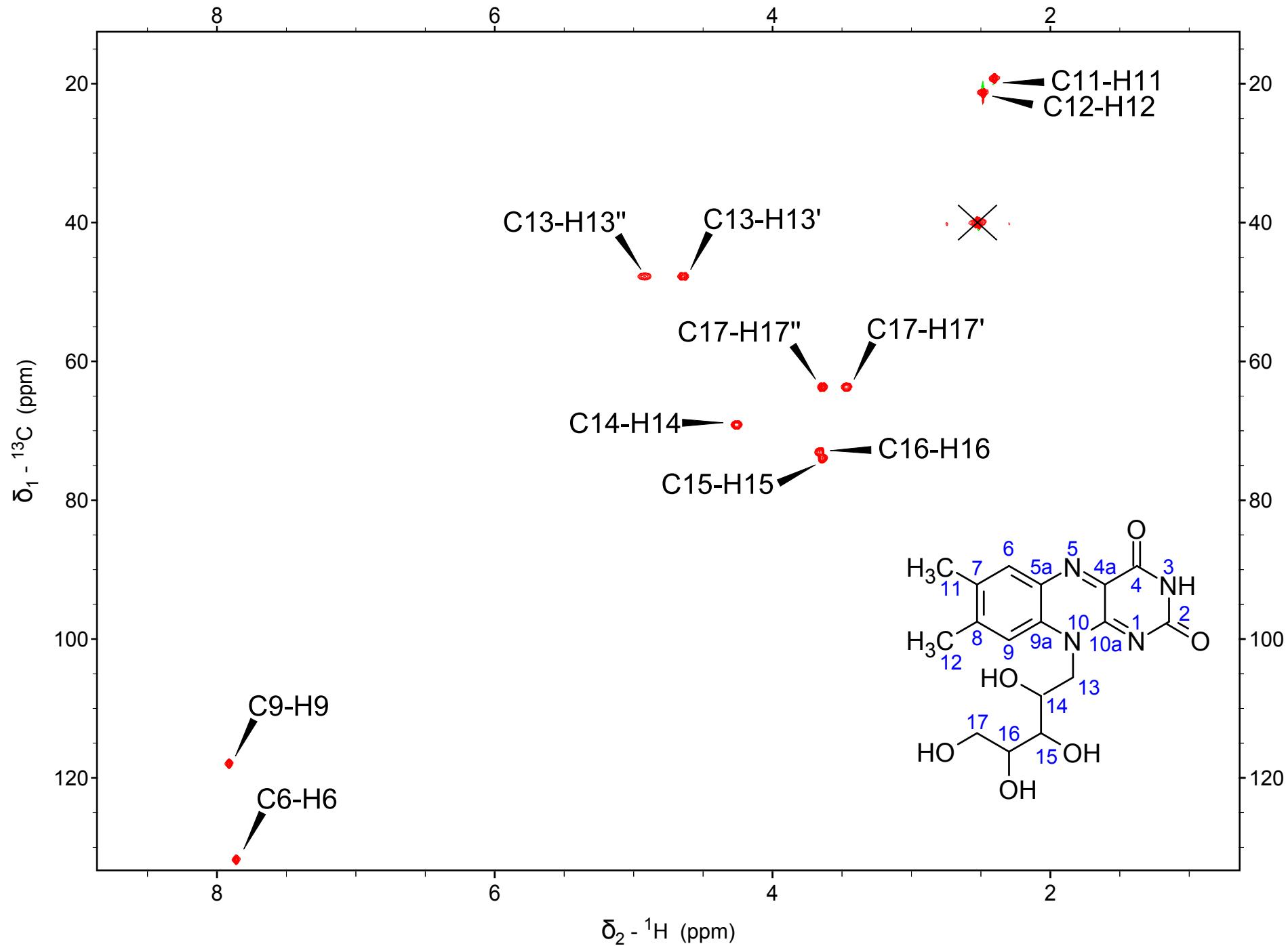
Riboflavin: DQF-COSY



Riboflavin: NOESY



Riboflavin: ^1H - ^{13}C HSQC



Next session:

Midterm test + Complex exercises