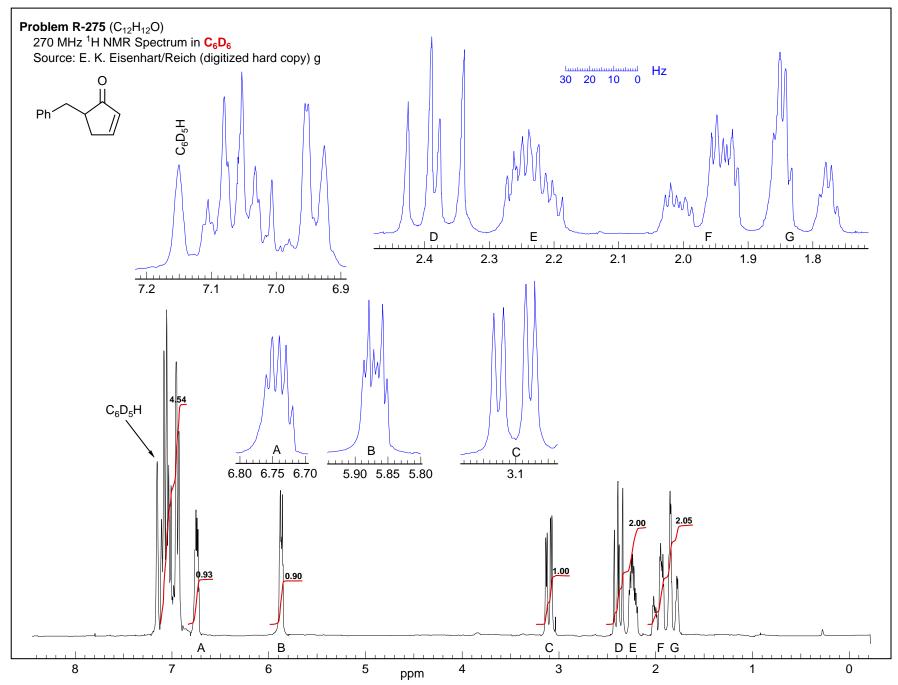
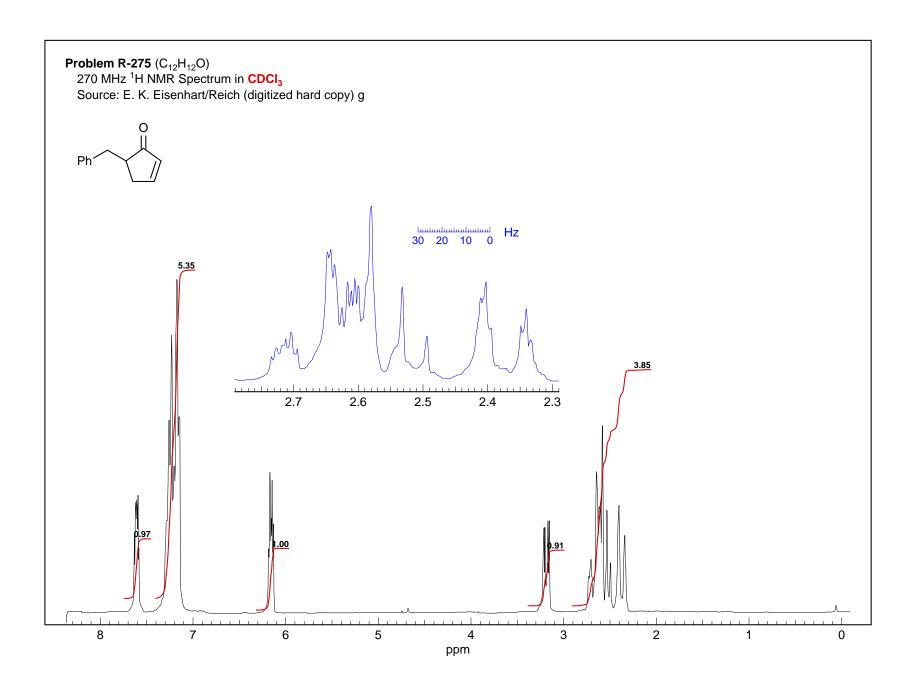
**Problem R-275**: Determine the structure from the <sup>1</sup>H NMR spectrum. Suggest an average conformation for the molecule from the coupling constants and chemical shifts.





## **Problem R-275** (C<sub>12</sub>H<sub>12</sub>O)

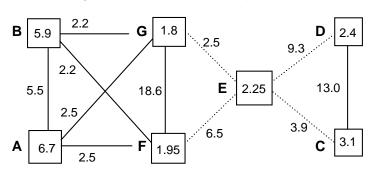
Because of second order effects resulting from several close-coupled protons, the spectrum in CDCl<sub>3</sub> cannot be analyzed. In  $C_6D_6$  the spectrum is mostly first order, and allows complete analysis (see next page for ASIS analysis).

Here are the coupling constants obtained by analysi of the C<sub>6</sub>D<sub>6</sub> spectrum

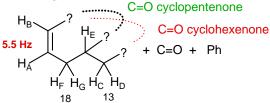
E is hard to analyze a priori, but all coupling to E can be identified from the "left-over" J values in the other multiplets

	Α	В	С	D	Ε	F	G
δ	6.7	5.9	3.1	2.4	2.25	1.95	1.8
	dt	dt	dd	dd	(dddd)	ddt	dq
J	5.5	5.5	13.0	13.0	(9.3)	18.6	18.6
	2.5	2.2	3.9	9.3	(6.5)	6.5	2.5
	2.5	2.2			(3.9)	2.4	2.5
					(2.5)	2.4	2.5

The coupling constants can be matched up as shown in the scheme below



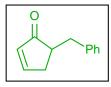
The chemical shifts and couplings lead to the following part structures:



This defines all protons and carbons in the molecule. The three "?" position ae filled by C=O and Ph.

Two structures fit the data:

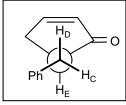
The cyclohexenone would have a ca 10 Hz HC=CH coupling. Since it is only 5.5 Hz, must be the cyclopentenone





$$H_{A}$$
 $H_{B}$ 
 $H_{B$ 

## Conformation of benzyl:



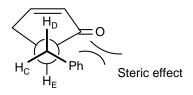
J<sub>DE</sub> large

 $H_{E}$ 

 $J_{DE}\approx J_{CE}$ 

J<sub>CE</sub> small, H<sub>C</sub> downfield (C=O anisotropy)

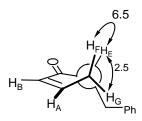
This fits best

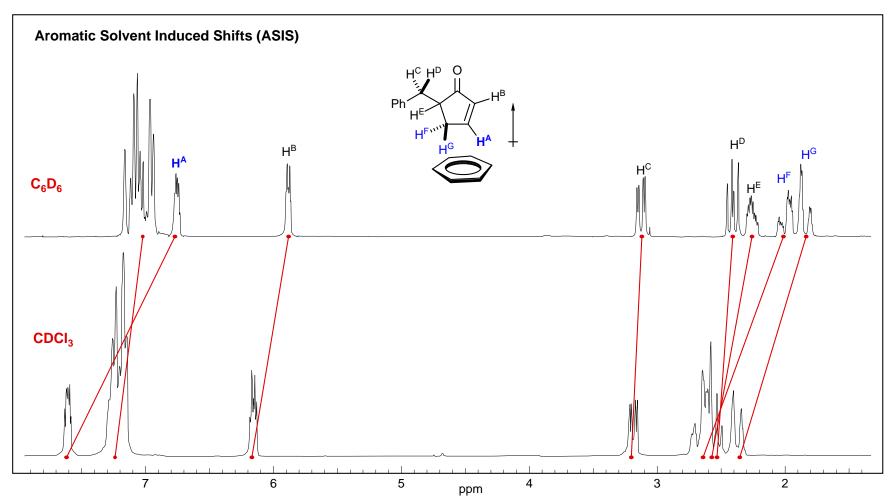


J<sub>DE</sub> large

J<sub>CE</sub> small

Chemical shift of H<sub>C</sub>?





The observed shifts fit well with the "transient pi complex" model for ASIS. H<sup>A</sup> moves the most, H<sup>F</sup> and H<sup>G</sup> also move a lot; H<sup>C</sup> and H<sup>D</sup> move the least - they are furthest from the positive end of the enone dipole.

**Problem R-275**: Determine the structure from the <sup>1</sup>H NMR spectrum. Suggest an average conformation for the molecule from the coupling constants and chemical shifts.

