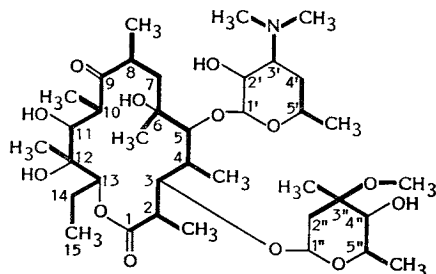


PARTICULAR UTILITY OF THE  
HMBC TECHNIQUE TO  
POLYPROPIONATE DERIVED  
METABOLITES AS EXEMPLIFIED  
BY ERYTHROMYCIN A

Sir:

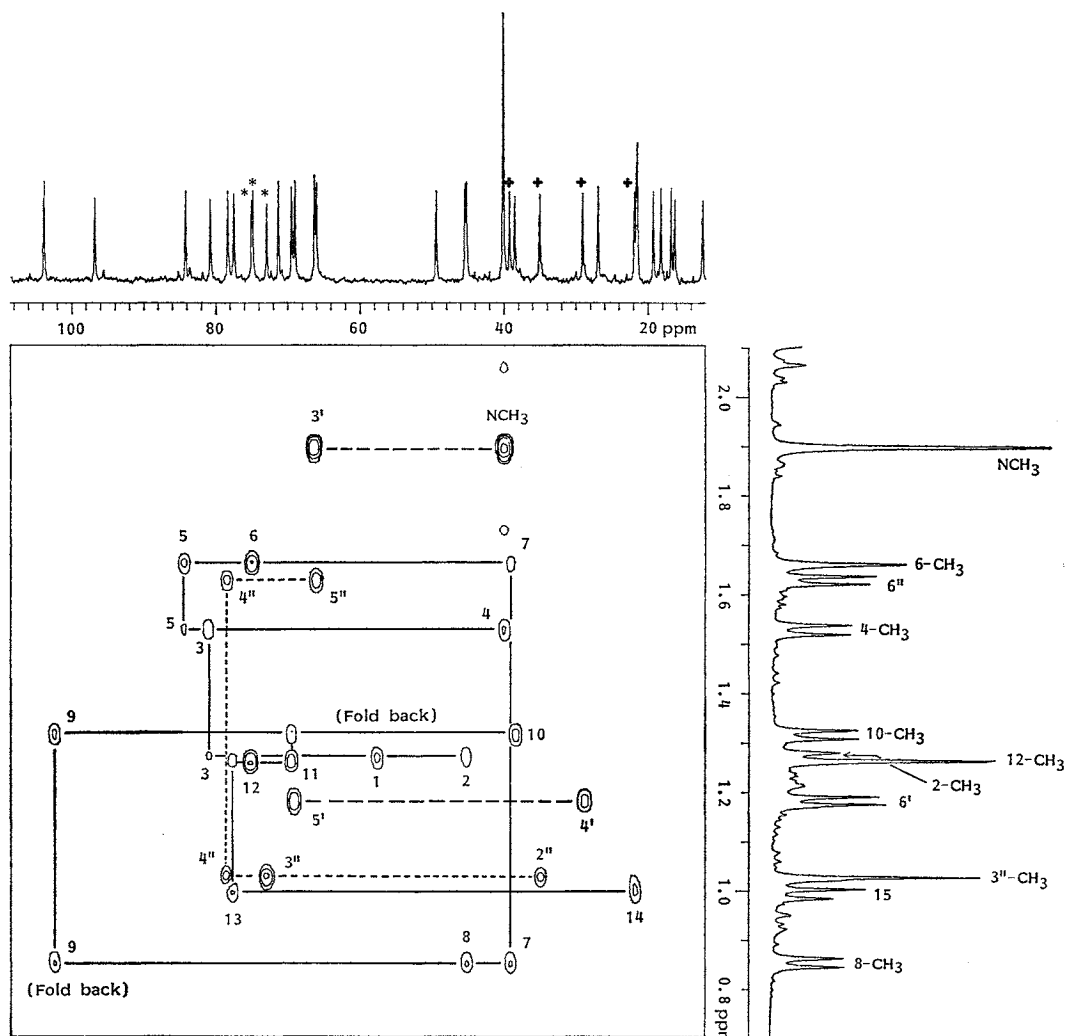
Recently a new 2D NMR technique named Heteronuclear Multiple Bond Connectivity (HMBC) was reported by BAX *et al.*<sup>1-3)</sup> This method reveals <sup>13</sup>C and <sup>1</sup>H connectivities separated by two or three bonds through detection of cross peaks observed with <sup>1</sup>H nucleus. Since

Fig. 1. The structure of erythromycin A.



Bold lines show the connectivities revealed by analysis of the HMBC spectrum.

Fig. 2. The pertinent region of the HMBC spectrum of erythromycin A (in C<sub>6</sub>D<sub>6</sub>).



+ and \* represent methylene and quaternary carbons, respectively. The remaining signals below 30 ppm are due to methine carbons.

the sensitivity of this method is dependent on the signal intensities of the proton signals, its application to polyketide antibiotics such as macrolides and polyethers with many methyl groups, which are observed as strong sharp signals in the  $^1\text{H}$  NMR spectra, is highly promising as evidenced by our previous work.<sup>4,5)</sup>

Another advantage of the application of HMBC for structural studies of complicated molecules with many methyl groups is that the cross peaks of the methyl carbons can be very easily analyzed, since the carbon next to a methyl residue (*i.e.*  $^2J_{\text{C-H}}$ ) is easily identified by analysis of  $^1\text{H}$ - $^1\text{H}$  correlation spectroscopy (COSY) and  $^{13}\text{C}$ - $^1\text{H}$  COSY spectra. Thus, the distinction of  $^2J_{\text{C-H}}$  and  $^3J_{\text{C-H}}$ , which is very difficult in most cases, can easily be accomplished.

In this paper, we wish to report the application of HMBC to the assignment of the  $^{13}\text{C}$  NMR spectrum of erythromycin A (Fig. 1).

Fig. 2 shows the HMBC spectrum of erythromycin A dissolved in  $\text{C}_6\text{D}_6$ . In order to improve digital resolution, the relevant region of  $^{13}\text{C}$  NMR spectrum ( $\delta_{\text{C}}$  10~110) was measured resulting in folding over of the ester (C-1, appeared at  $\delta_{\text{C}}$  175.9 in the 1D  $^{13}\text{C}$  NMR spectrum) and ketone (C-9, appeared at  $\delta_{\text{C}}$  220.6 in the  $^{13}\text{C}$  NMR spectrum) resonances, which are observed

at approximately 58 and 102 ppm, respectively.

Analysis of this spectrum could be made very straightforwardly. For example, a methyl proton doublet at  $\delta_{\text{H}}$  1.27 (2- $\text{CH}_3$ ) showed cross peaks with an ester carbonyl (observed as a fold back signal at *ca.*  $\delta_{\text{C}}$  58 (C-1)), a methine at  $\delta_{\text{C}}$  45.4 (C-2) and an oxymethine at  $\delta_{\text{C}}$  80.8 (C-3), which are unambiguously assigned to the partial structure,  $\text{OOC1-C2H}(\text{CH}_3)\text{-C3H}(\text{O})$  by taking account of their  $^{13}\text{C}$  chemical shifts. The direct combination of this methyl group to the non oxygenated methine ( $\delta_{\text{H}}$  3.0,  $\delta_{\text{C}}$  45.4 (C-2)) was based on analysis of  $^1\text{H}$ - $^1\text{H}$  COSY and  $^{13}\text{C}$ - $^1\text{H}$  COSY spectra (data not shown).

As shown in Fig. 2, another cross peak is observed between C-3 and a methyl doublet at  $\delta_{\text{H}}$  1.525 (4- $\text{CH}_3$ ), which shows additional cross peaks with a methine ( $\delta_{\text{C}}$  40.1 (C-4)) and an oxymethine ( $\delta_{\text{C}}$  84.2 (C-5)). These results proved the connectivity of  $\text{C3H}(\text{O})\text{-C4H}(\text{CH}_3)\text{-C5H}(\text{O})$  extending the relationship from C-1 to C-5. The complete connectivity of the carbon skeleton of the aglycone moiety (C-1 to C-15) could easily be established by repeating the same procedure. In addition, the connectivities  $\text{C2''H}_2\text{-C3''}(\text{CH}_3)(\text{OCH}_3)\text{-C4''H}(\text{O})\text{-C5''H}(\text{O})\text{-CH}_3$  (shown by dotted lines) in cladinose, and  $(\text{CH}_3)_2\text{N-C3''H}$  and  $\text{C4''H}_2\text{-}$

Table 1.  $^{13}\text{C}$  Chemical shifts of erythromycin.

Carbon	$\delta_{\text{C}}$	Carbon	$\delta_{\text{C}}$
C-1 (COO)	175.9	C-1' (O-CH-O)	103.8
C-2 (CH)	45.4	C-2' (CH-O)	71.3
C-3 (CH-O)	80.8	C-3' (CH-N)	66.3
C-4 (CH)	40.1	C-4' ( $\text{CH}_2$ )	29.1
C-5 (CH-O)	84.2	C-5' (CH-O)	69.0
C-6 (C-O)	74.9	C-6' ( $\text{CH}_3$ )	21.6
C-7 ( $\text{CH}_2$ )	39.2	N- $\text{CH}_3$	40.1
C-8 (CH)	45.2	C-1'' (O-CH-O)	96.8
C-9 (C=O)	220.6	C-2'' ( $\text{CH}_2$ )	35.0
C-10 (CH)	38.5	C-3'' (C-O)	72.9
C-11 (CH-O)	69.4	C-4'' (CH-O)	78.3
C-12 (C-O)	75.0	C-5'' (CH-O)	66.0
C-13 (CH-O)	77.5	C-6'' ( $\text{CH}_2$ )	19.3
C-14 ( $\text{CH}_2$ )	21.9	3''- $\text{CH}_3$	21.5
C-15 ( $\text{CH}_3$ )	11.1	O- $\text{CH}_3$	49.4
2- $\text{CH}_3$	16.3		
4- $\text{CH}_3$	9.6		
6- $\text{CH}_3$	26.9		
8- $\text{CH}_3$	18.2		
10- $\text{CH}_3$	12.4		
12- $\text{CH}_3$	16.8		

Taken in  $\text{C}_6\text{D}_6$ .

C5'H(O)-CH<sub>3</sub> (shown by broken lines) in desosamine are also revealed as shown in Fig. 2. The connectivities revealed by analysis of the HMBC spectrum are indicated by bold lines in Fig. 1.

In addition to the structural information just explained, the HMBC spectrum revealed the relationships between 1'-H and C-5, 1'-H and C-5', 1''-H and C-3, and 1''-H and C-5'' (data not shown). However, the linkage between C-1 and C-13 through an oxygen could not be proved due probably to the very small long range coupling ( $^3J_{O-H}$ ) between 13-H and C-1. Additional structural information such as C1''-C2'' and C1'-C2'-C3'-C4' could also be obtained by HMBC. These connectivities, however, were more easily obtained by usual <sup>1</sup>H NMR techniques such as COSY.

These results show clearly the particular usefulness of HMBC for structural elucidation of complicated antibiotics with many methyl groups.

In order to get good results by HMBC experiments, it is important to choose a solvent which gives good separation of methyl signals. As far as our experiences are concerned, C<sub>6</sub>D<sub>6</sub> or pyridine-*d*<sub>5</sub> gave better results than CDCl<sub>3</sub> in most cases.

The <sup>13</sup>C chemical shift data thus obtained are summarized in Table 1. In view of the difference of the solvent employed in this experiment, this assignment is in good agreement with those reported previously.<sup>6-9)</sup>

The experimental conditions were as follows; spectral width F<sub>1</sub> (<sup>13</sup>C) 10,000 Hz, F<sub>2</sub> (<sup>1</sup>H) 3,000 Hz, data matrix (F<sub>1</sub> × F<sub>2</sub>) = 128 × 1,024 (F<sub>1</sub> data were zero filled prior to Fourier transformation), 240 scans per t<sub>1</sub>, total measuring time 14 hours. The sample (10 mg) was dissolved in 0.35 ml of C<sub>6</sub>D<sub>6</sub>.

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