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Analysis of Corticosteroids by HPLC-UV and HPLC-MS by Particle Beam Interface

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The Antidoping Lab of Rome is studying a method for the detection and the identification of corticosteroids in human urine.

Since the analysis in GC-MS, very often, require previous derivatization of the samples, with side effects, we prefer to use the HPLC for these kinds of compounds.

For the screening analysis we use the HPLC-UV with diode array detector, on the other side for the confirmations the HPLC-MS by particle beam interface (PB).

The system used by us is shown in figure 1. The switch valve in position 2-1 helps to avoid that a part of the more polar compounds, eluted very quickly, goes into the analytical column. That is very important because it makes the life of the column longer.

For our compounds the RP deactivated column gives better separations compared to the CN and C18 columns. The pre-column (4.6 cm x 2 cm) has the same packing as the analytical column. The detector can be a UV diode array or a mass spectrometer.

To test the sensitivity of the diode array detector we used a mixture of two natural corticosteroids, cortisone and hydrocortisone, by injecting 50 ng of each (figure 2).

In the same analytical conditions we also studied the behaviour of other natural and synthetic corticosteroids. In figure 3 a separation of eight of them is shown.

We also made excretion studies of some drugs containing corticosteroids administering them to non athlete volunteers.

25 mg of cortisone, 25 mg of prednisone and 25 mg of deflazacort were given in a single dose.

Blank urine were collected before the administration.

After administration of prednisone and deflazacort the urine were pooled for 12 hours.

For the cortisone, instead, scheduled sampling were made.

The urine samples were prepared according to the scheme shown in figure 4.

The hydrolysis of the sample did not give other information, compared to the free fraction, and at the same time the background is more disturbed. In figure 5 the free fraction of a male blank urine is shown. Two peaks with the same retention time of cortisone and hydrocortisone are evident at the same instrument sensitivity.

In the same figure the chromatogram of the free fraction of female blank urine is also reported.

In figure 6 the excretion study of cortisone is shown. After three hours it is possible to notice that both the peak of cortisone and hydrocortisone are increased compared to the blank urine while after six hours differences are not seen. The excretion study of prednisone showed that the main metabolite is prednisolone, $R_t=19.2$ min. The unchanged prednisone is evident at $R_t= 14.4$ min. There are also three other peaks, probable metabolites, at $R_t=16.6, 27.2, 32.1$ min respectively of the chromatogram of figure 7.

In figure 8 the conjugated fraction is also reported.

In the case of the administration of deflazacort we can notice, in figure 9, that most of it is excreted unchanged with other two probable metabolites.

The use of HPLC-MS by PB permitted us to identify some of the peaks of the previous chromatograms.

One of the disadvantages of PB is the sensitivity, lower than the GC-MS, but at the same time it is a very simple system and gives the possibility to work in EI and in CI.

The temperature and the helium pressure of the solvation chamber were set at 50°C and 40-45 psi respectively, the source at 200°C .

To test the efficiency of the system we used a standard mixture (1 μg of each). The TIC in full scan of it is reported in figure 10.

To test the sensitivity of the system we injected 50 ng of cortisone in SIM mode (figure 11).

The mass spectra of the investigated synthetic and natural standards are in figure 12.

The mass spectra in full scan of the peak at $R_t= 14.4$ and 19.2 of figure 7 and of the peak at $R_t = 17.6$ of figure 9 are shown in figure 13. They confirmed the results, reported before, in HPLC-UV.

Figure 14 shows the acquisition in SIM mode for the same sample of figure 7.

The sensitivity of the system is increased at least by ten times working in Chemical Ionization.

Good results at the moment have been obtained for cortisone and hydrocortisone using ammonia or methane as reactant gas, working at a source temperature of 130° C (figure 15).

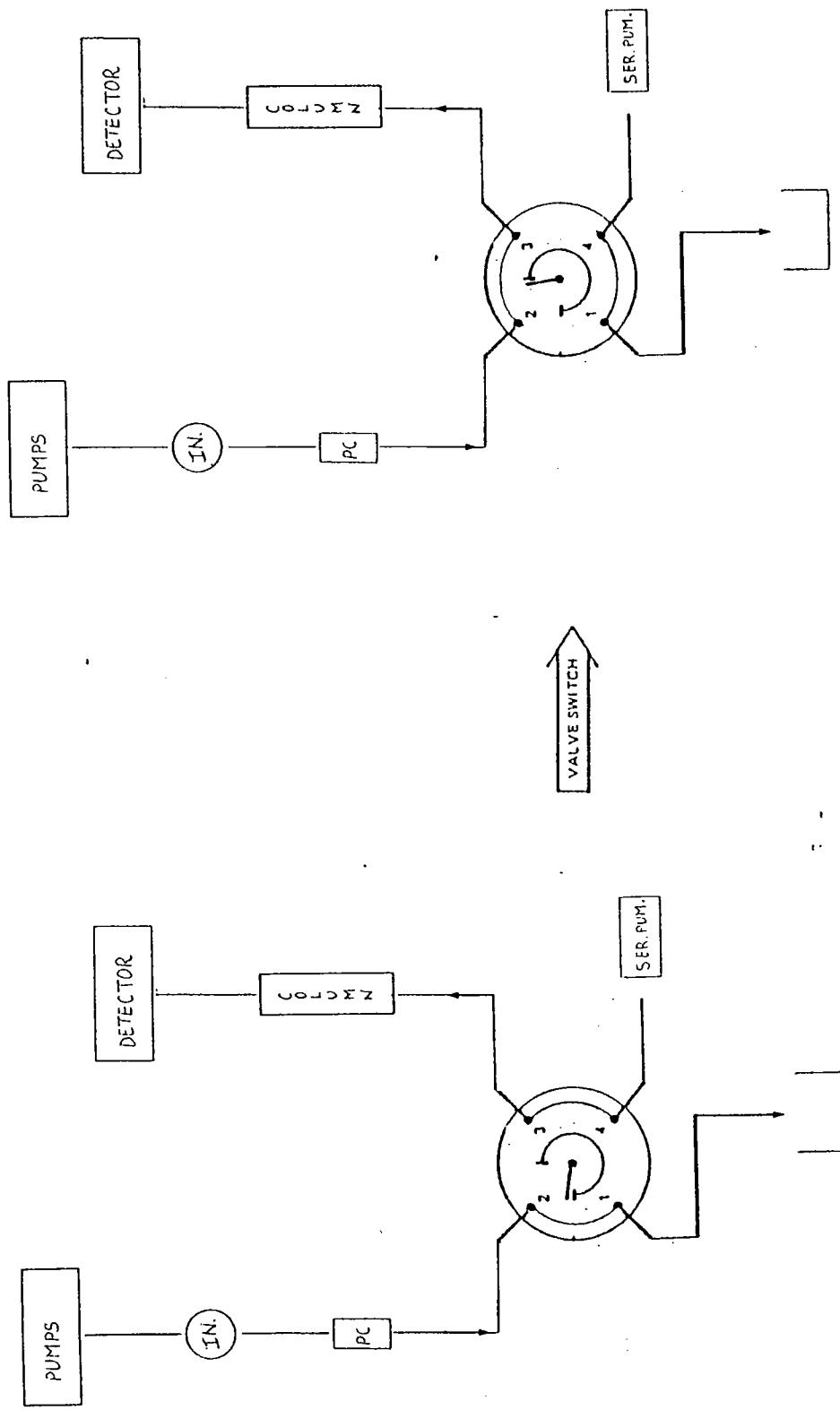


FIG. 1

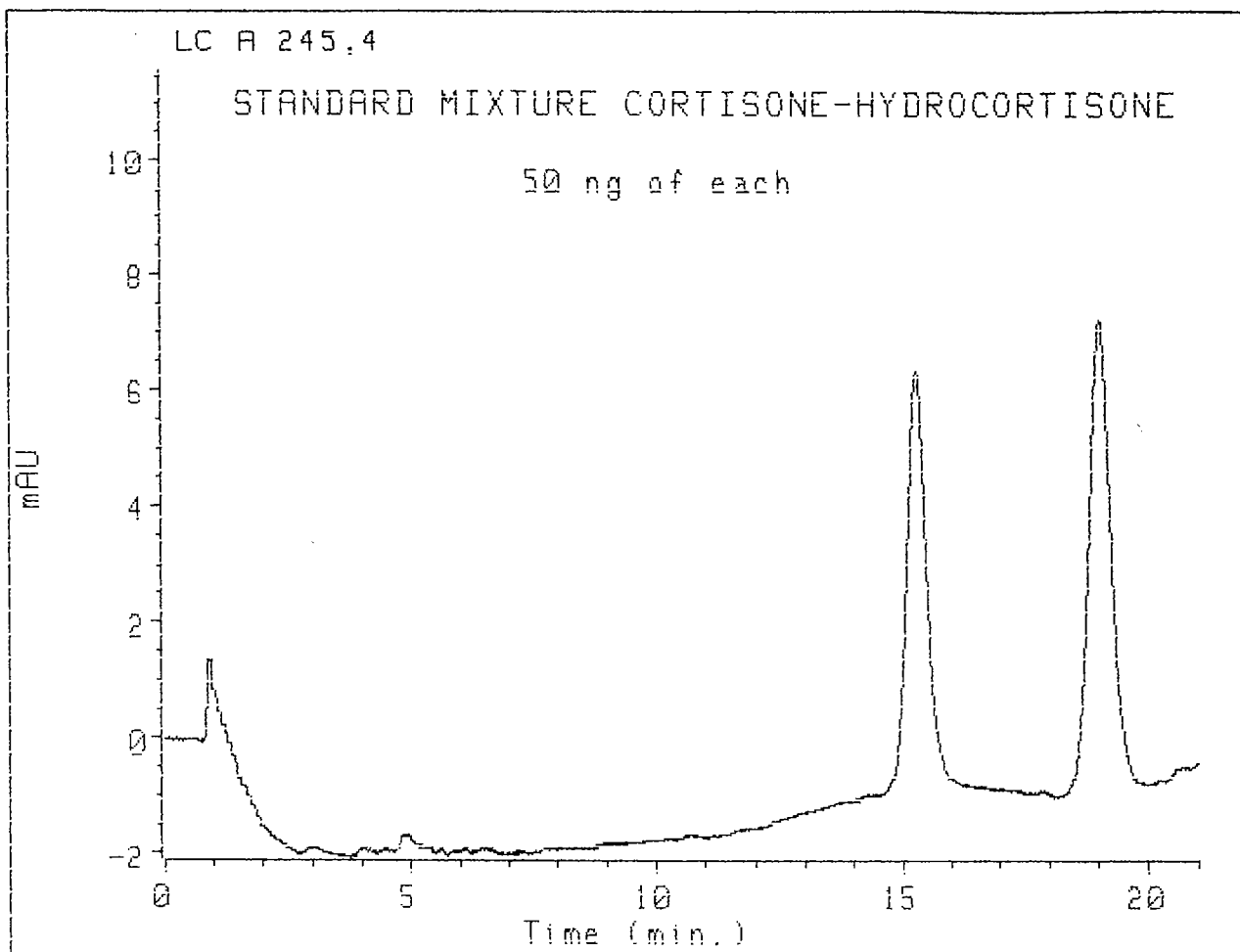


FIG. 2

Column: Supelcosil ABZ 150 mm x 4.6 mm 5 um spherical Flow 0.4 ml/min. UV= 245 nm

Time (min)				
0.01	Solvent	A :	50.0 %	B : 50.0 %
0.01	Contact	4 :	on	
0.01	Column	:	0	
0.80	Column	:	1	
13.00	Solvent	A :	50.0 %	B : 50.0 %
25.00	Solvent	A :	10.0 %	B : 90.0 %
35.00	Solvent	A :	10.0 %	B : 90.0 %
40.00	Solvent	A :	50.0 %	B : 50.0 %
45.00	Solvent	A :	50.0 %	B : 50.0 %

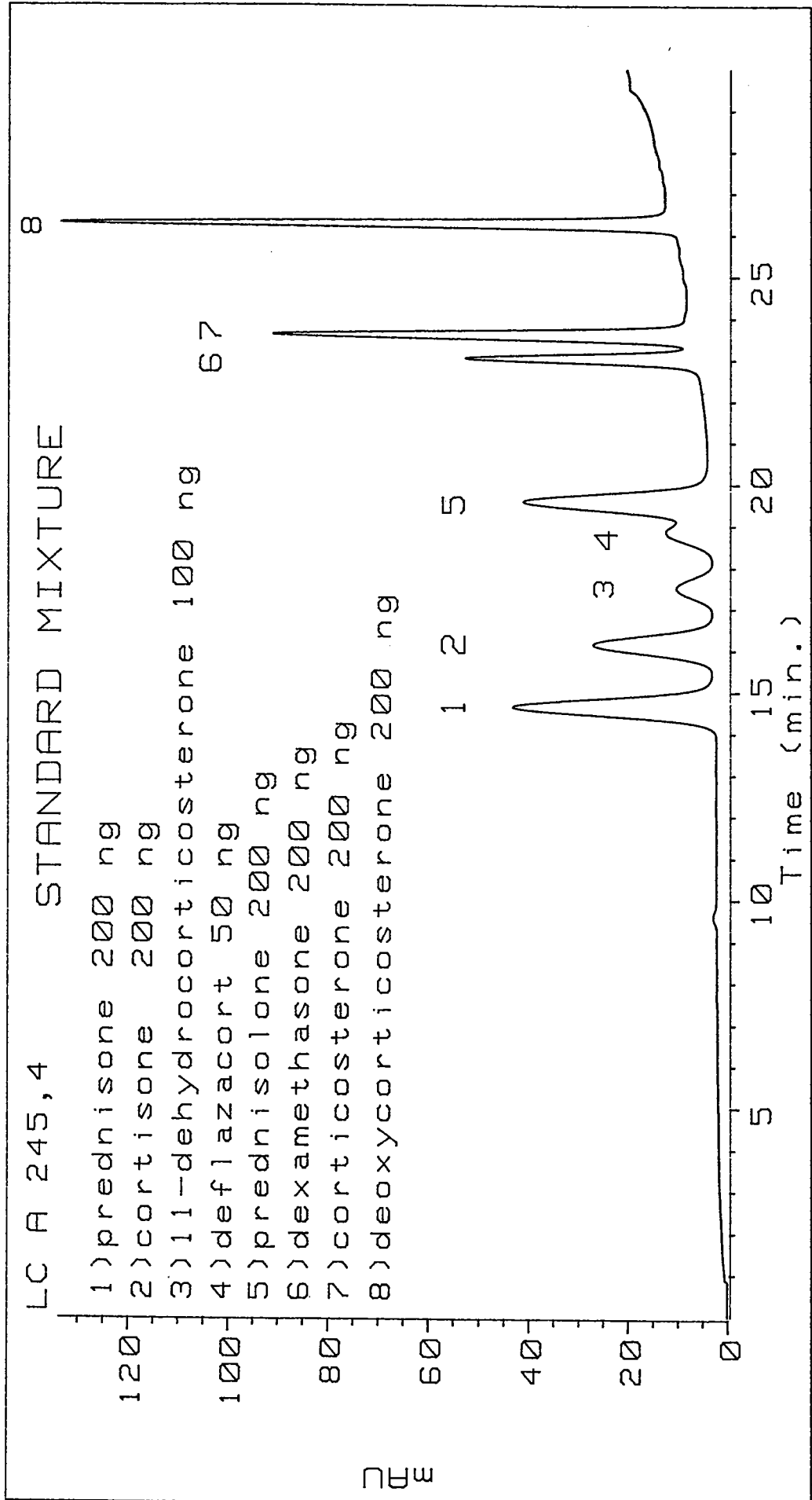


FIG. 3

SAMPLE PREPARATION

FREE FRACTION

5 ml urine



solid carbonate buffer (pH 9.2)



5 ml diethylether (peroxide free)



shaking 15 min, centrifuging



to dryness under N₂ below 40 C



dissolution in 100 ul H₂O:CH₃OH 1:1



25 ul injected

CONJUGATED FRACTION

5 ml urine

β -glucuronidase
hydrolysis



solid carbonate buffer (pH 9.2)



5 ml diethylether (peroxide free)



shaking 15 min, centrifuging



to dryness under N₂ below 40 C



dissolution in 100 ul H₂O:CH₃OH 1:1



25 ul injected

FIG. 4

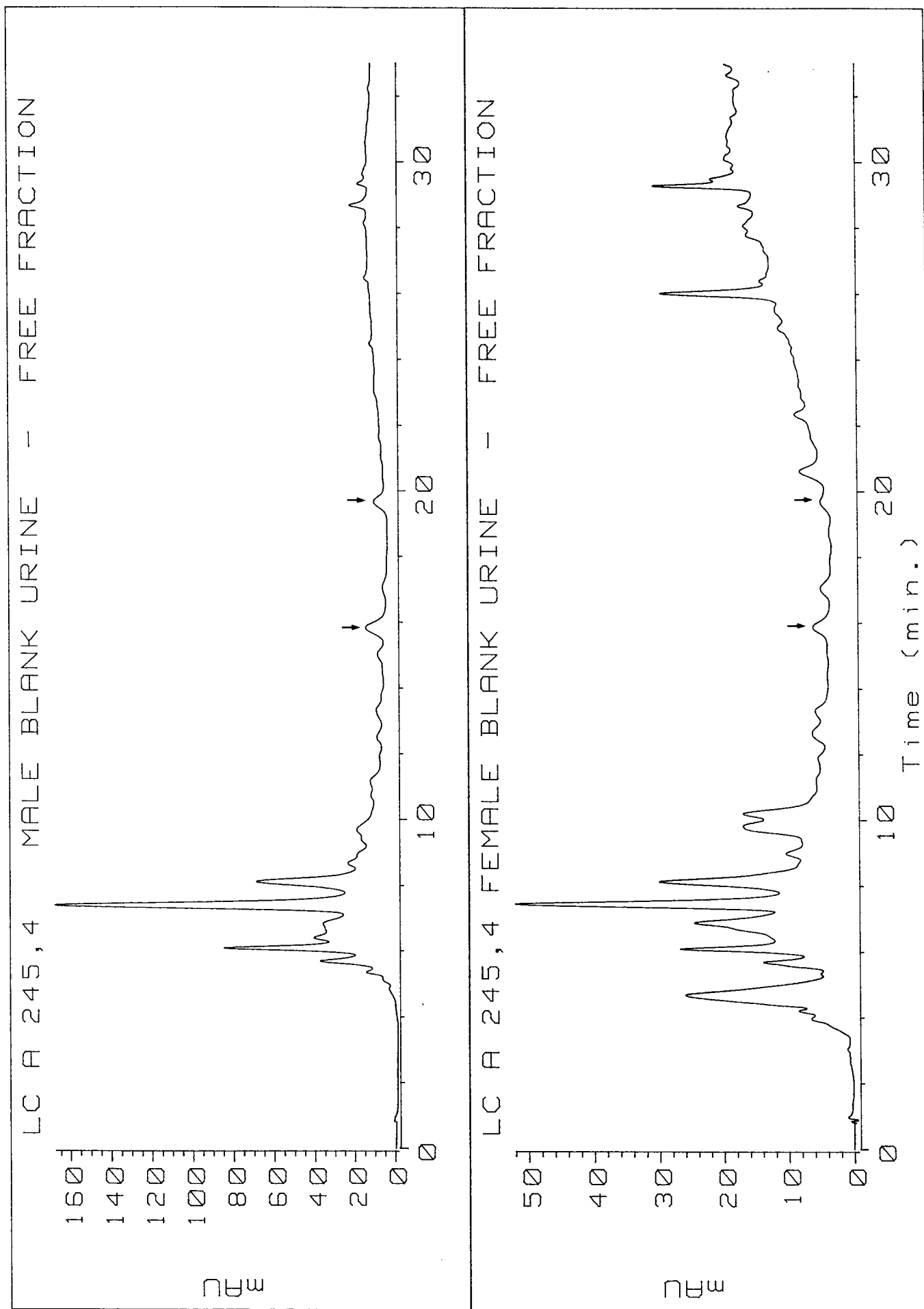


FIG.5

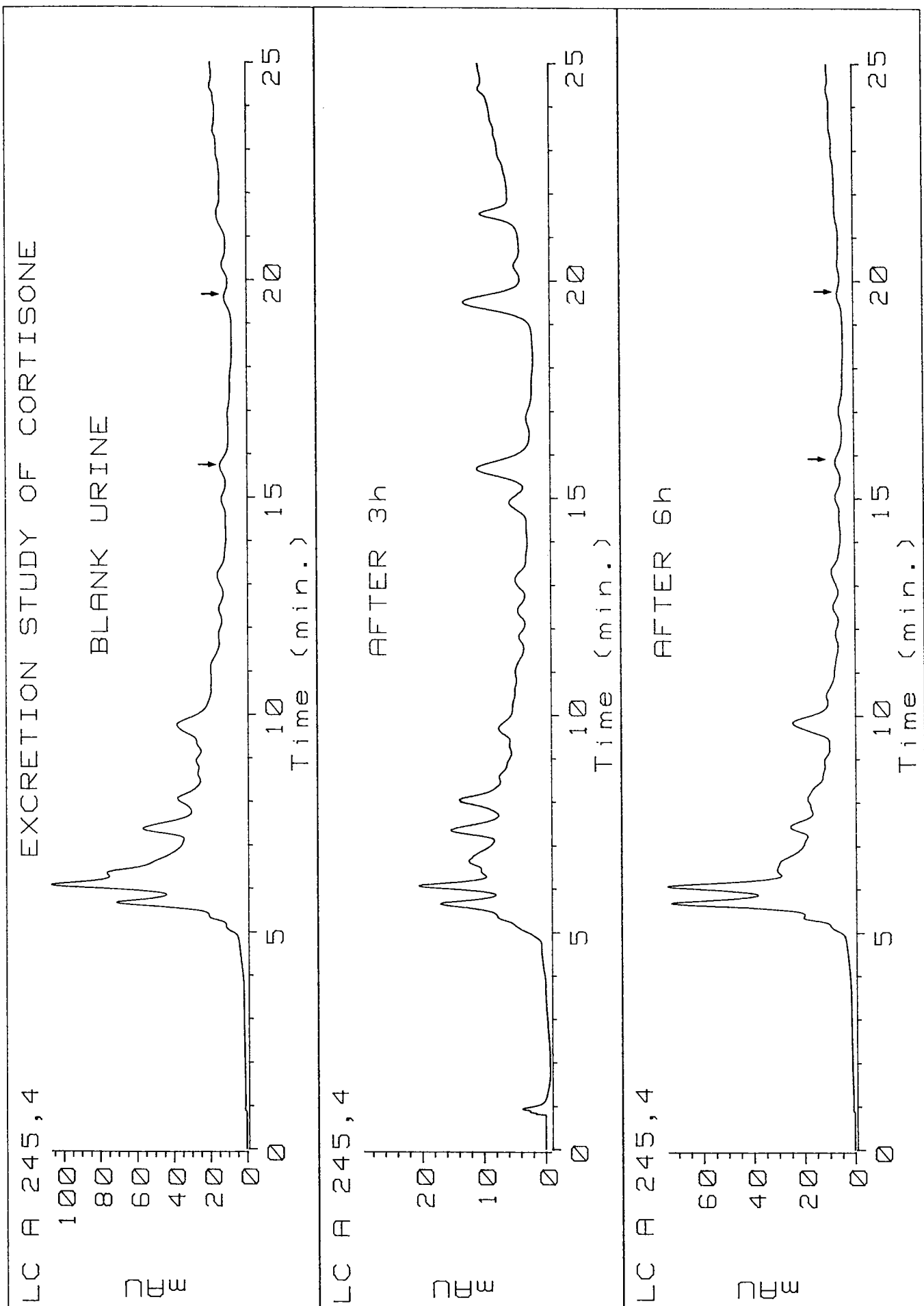
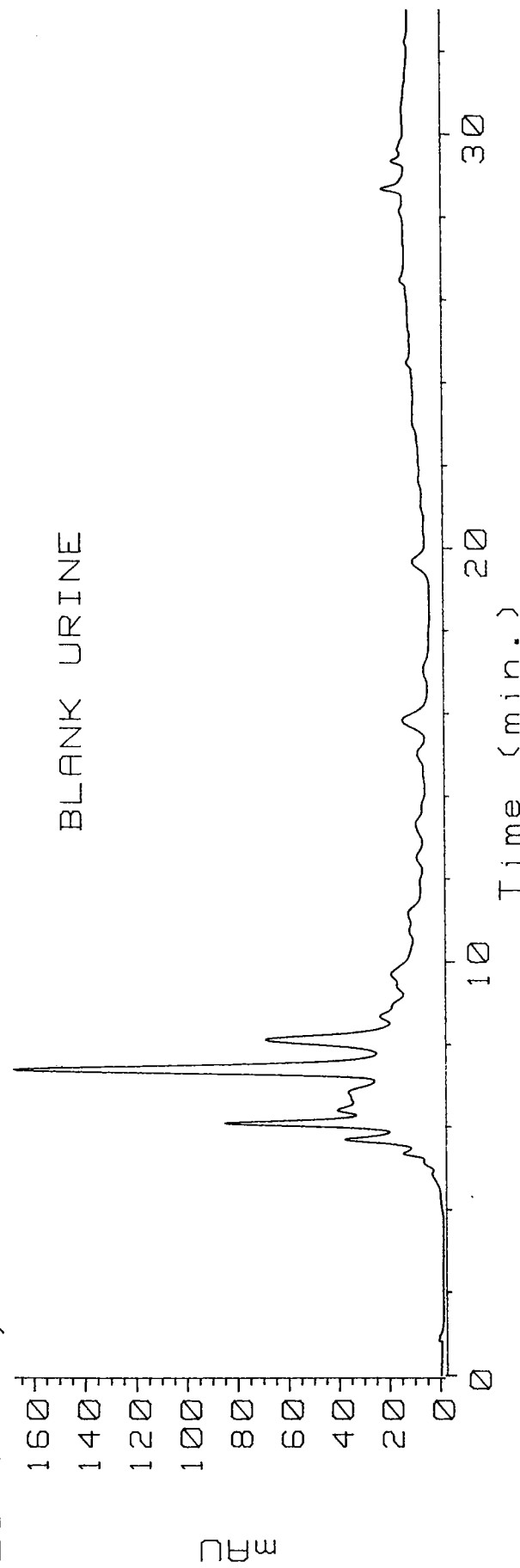


FIG.6

LC A 245, 4 EXCRETION STUDY OF PREDNISONONE - FREE FRACTION



LC A 245, 4 POOL OF 12h

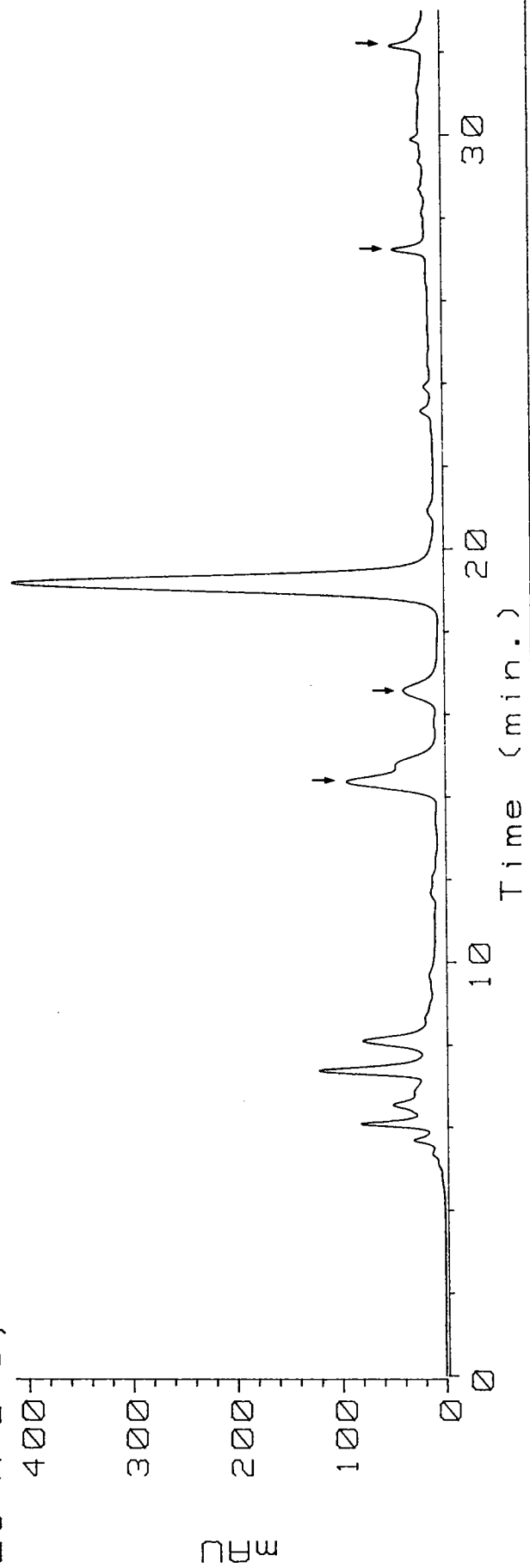


FIG.7

LC A 245,4 EXCRETION STUDY OF PREDNISONONE - CONJUGATED FRACTION

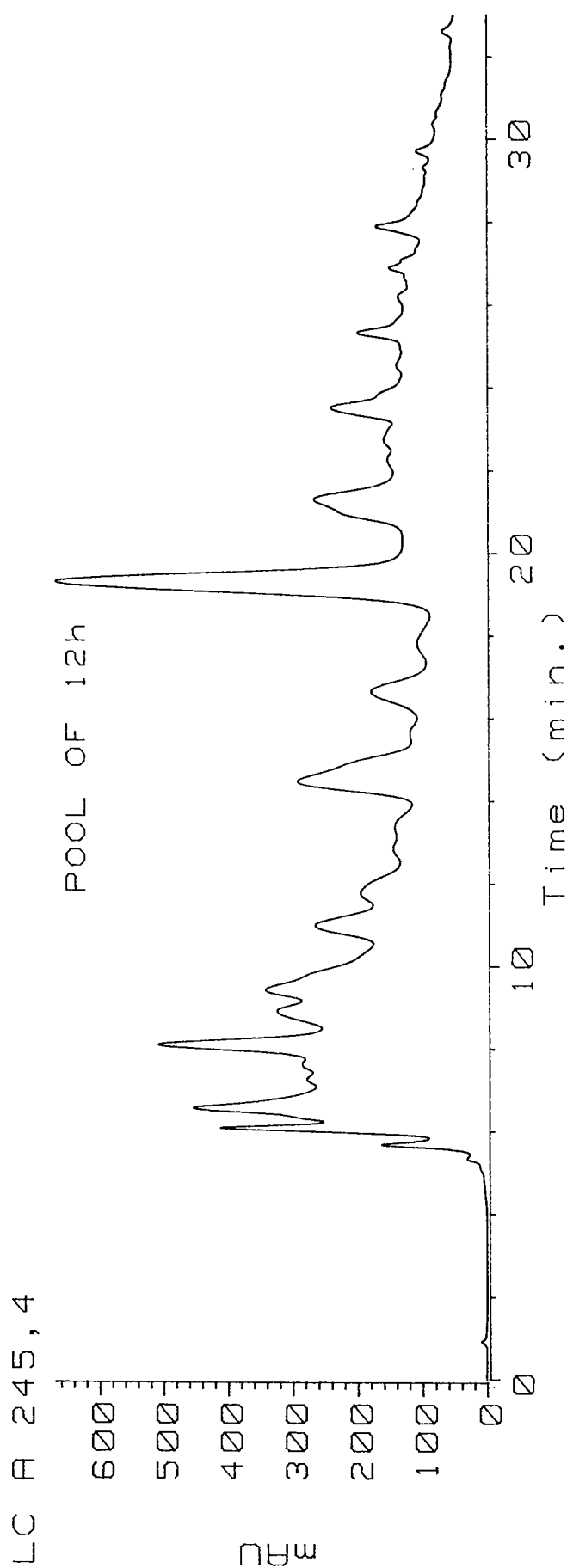
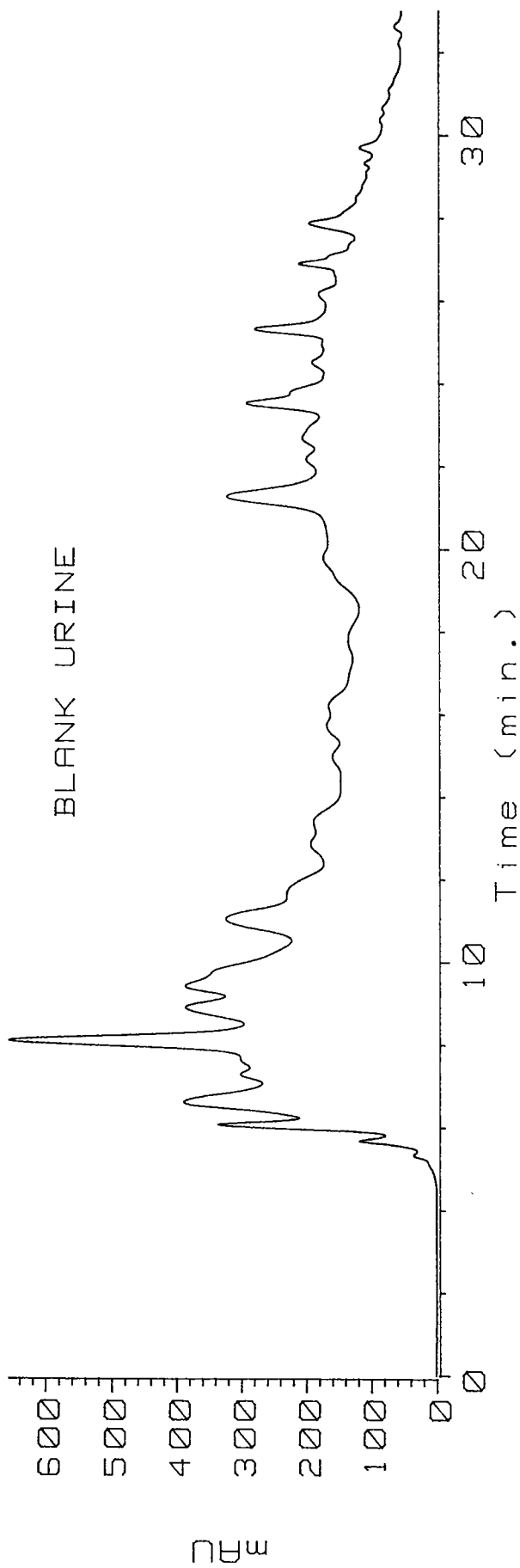


FIG.8

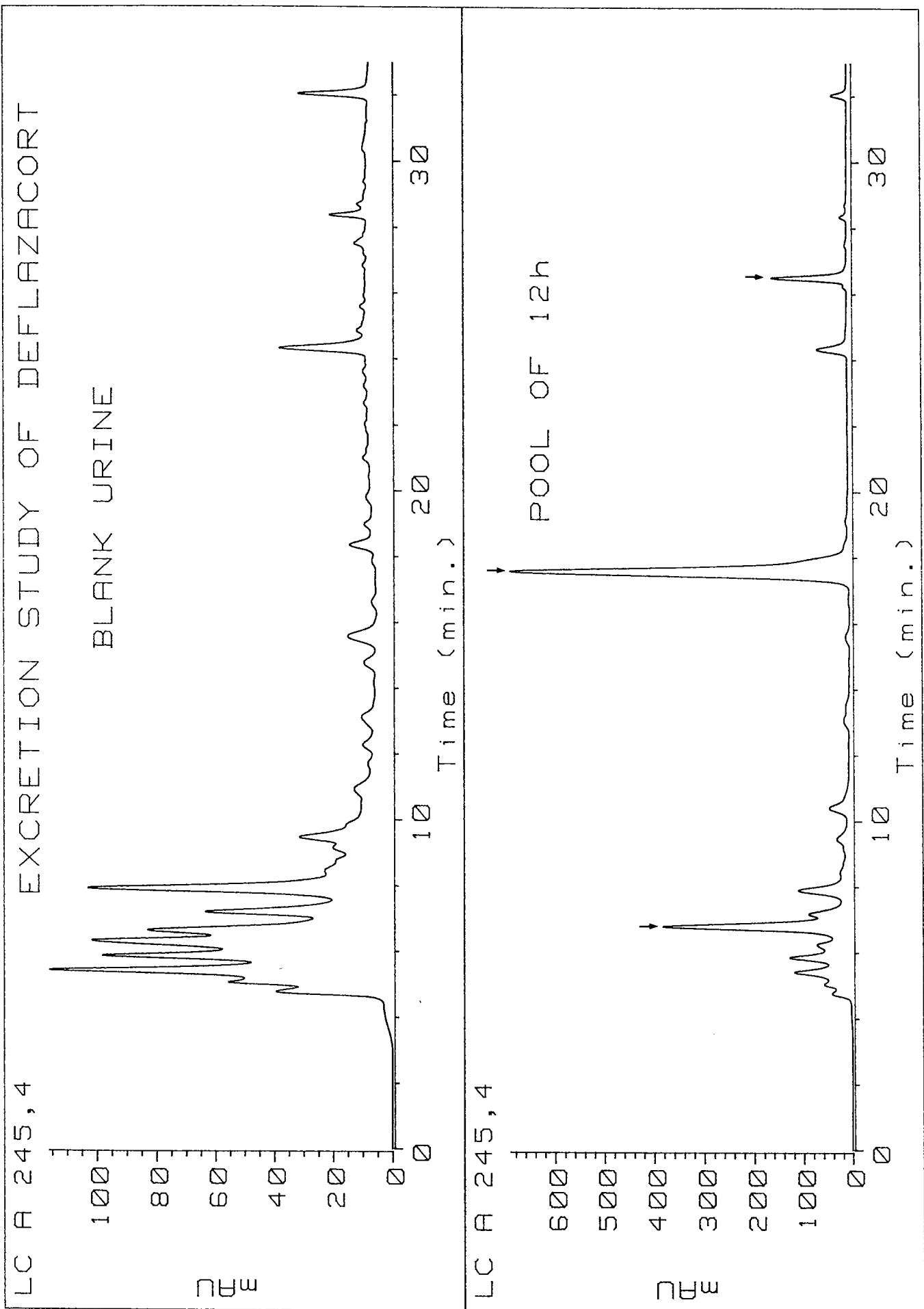


FIG.9

TIC OF STANDARD MIXTURE

1 ug of each

- 1) prednisone
- 2) cortisone
- 3) deflazacort
- 4) prednisolone
- 5) hydrocortisone
- 6) dexamethasone
- 7) corticosterone

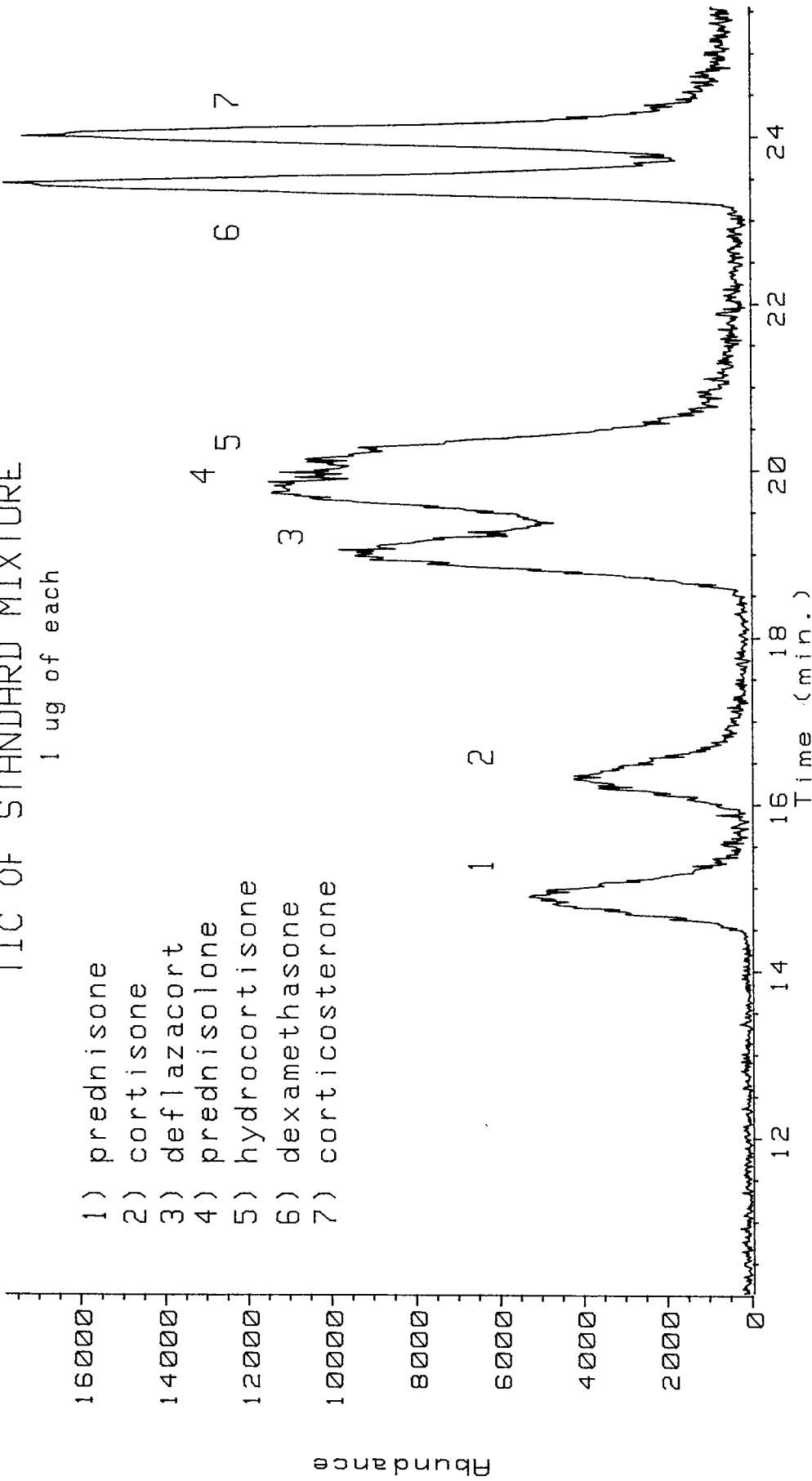


FIG.10

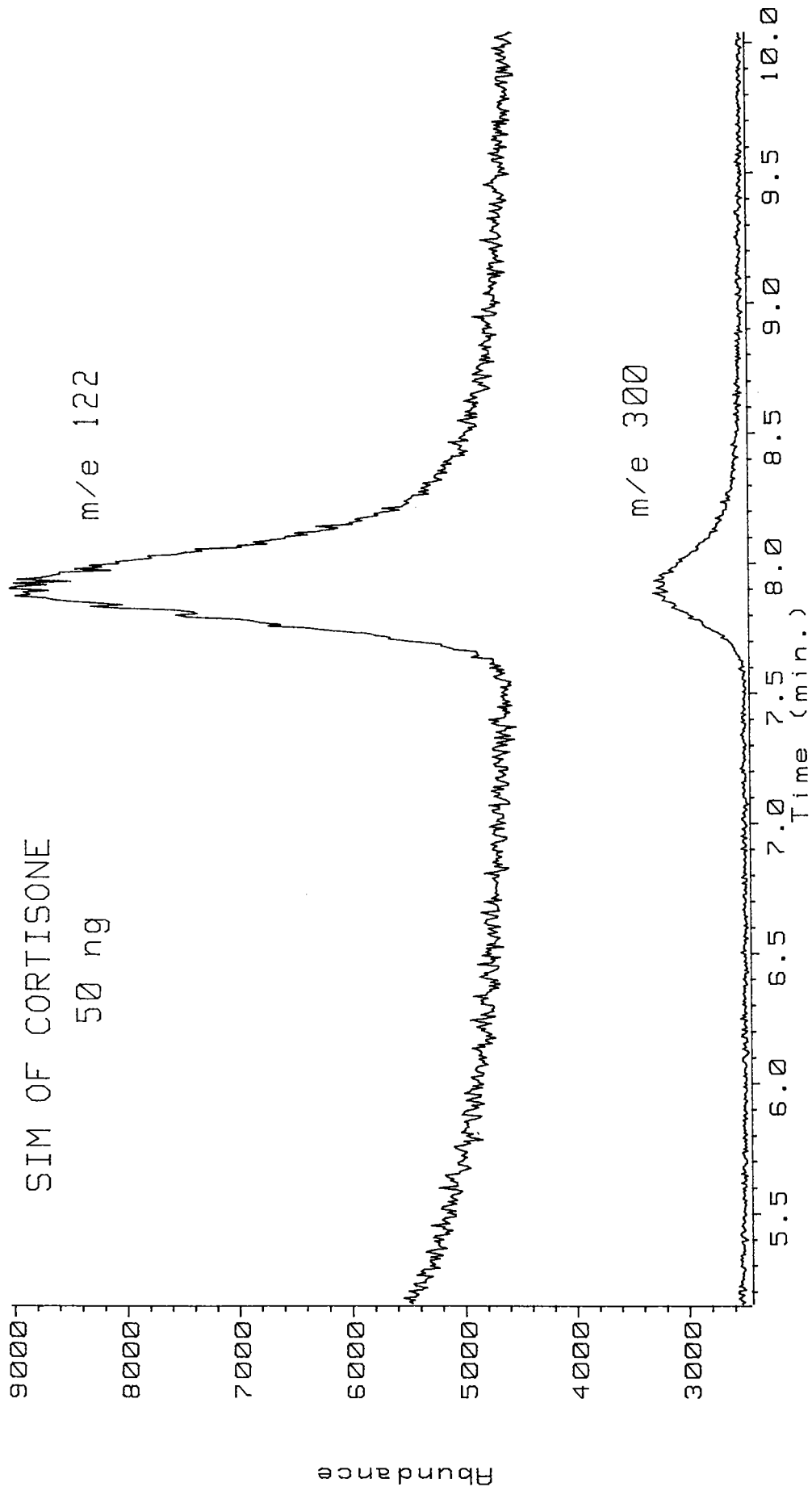


FIG.11

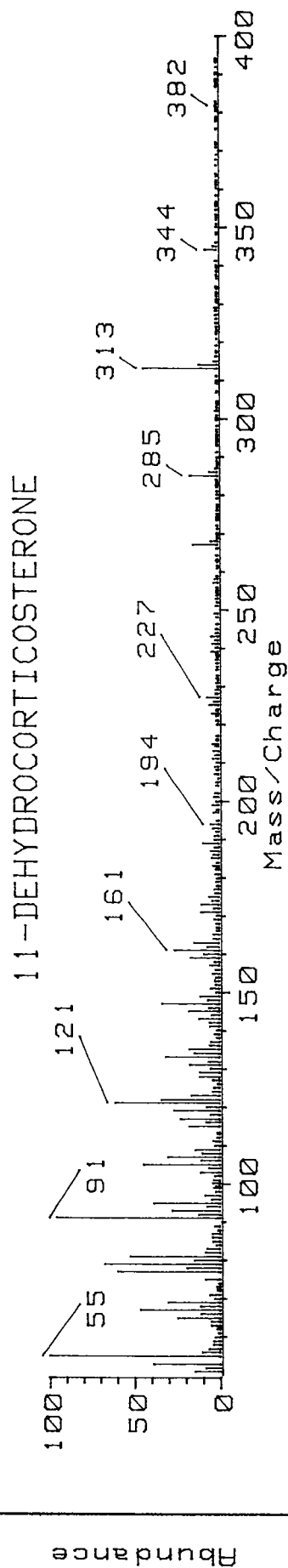
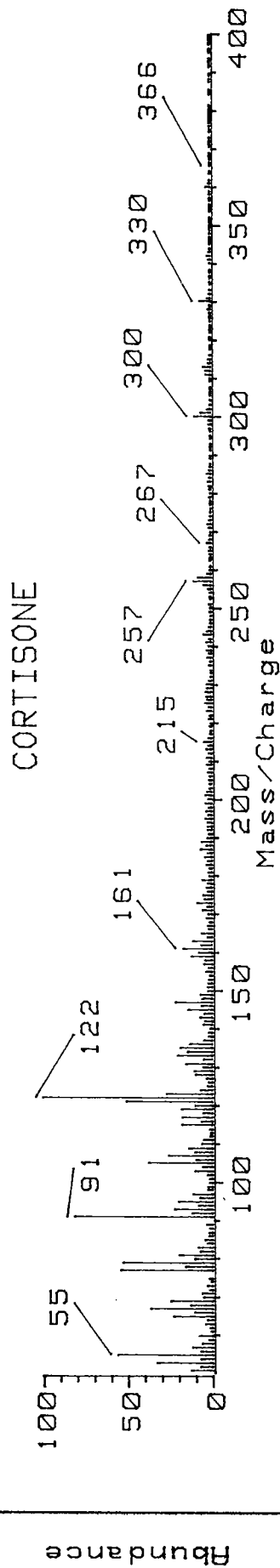
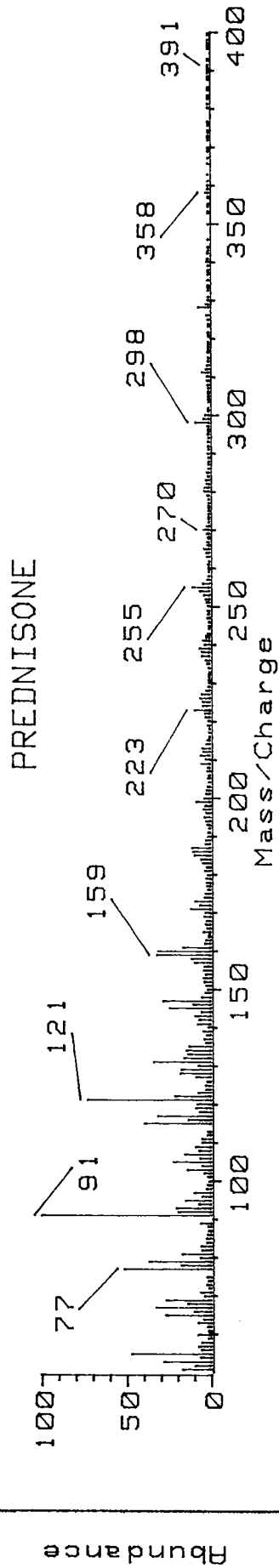


FIG. 12 a

Mass Spectra of corticosteroids . Full scan EI

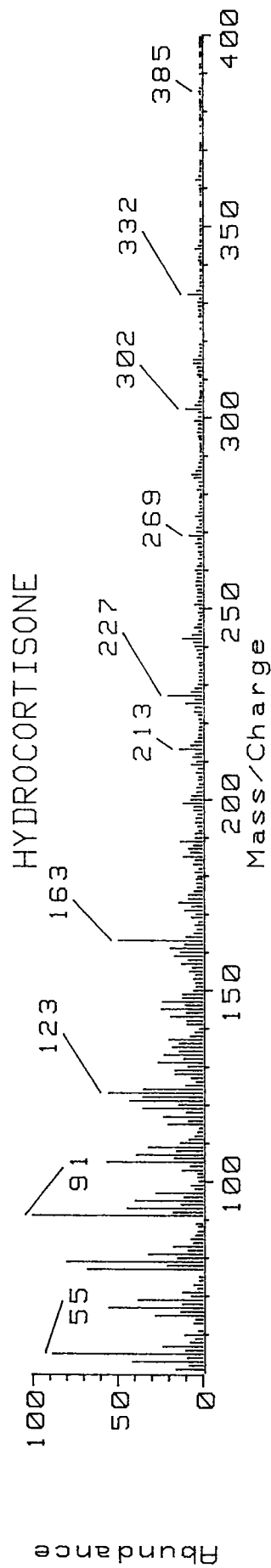
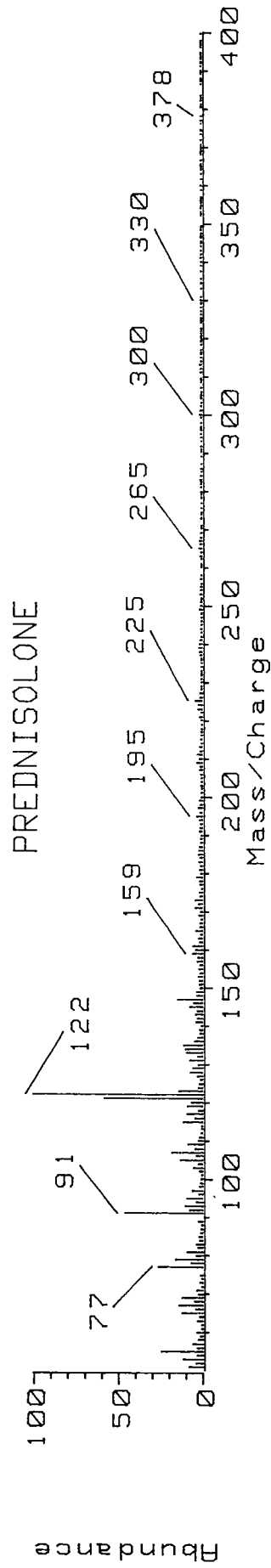
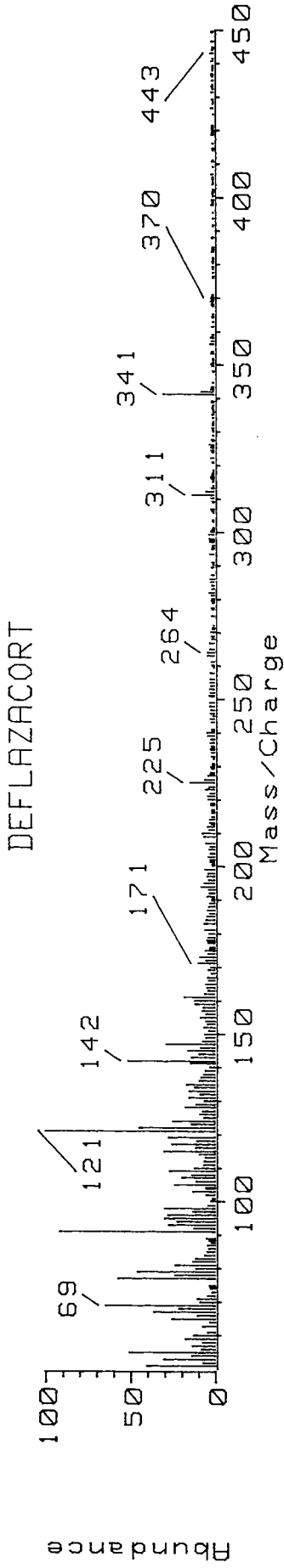


FIG. 12 b
Mass Spectra of corticosteroids. Full scan EI

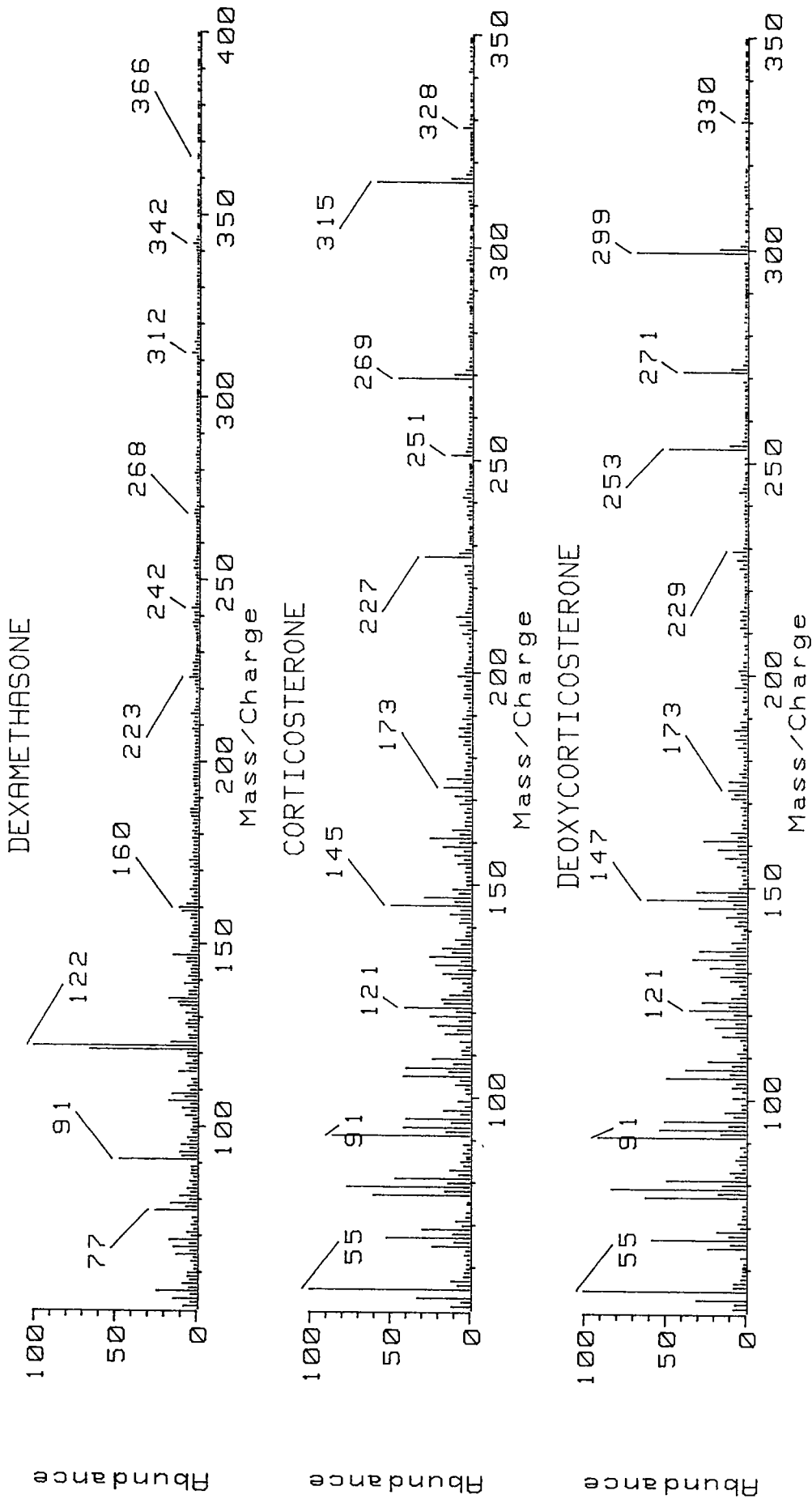


Fig. 12 c
Mass Spectra of corticosteroids. Full scan EI

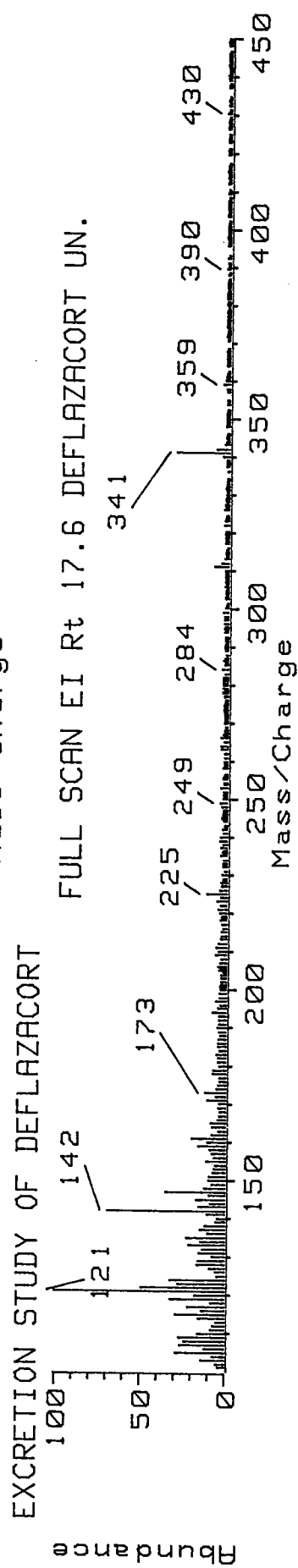
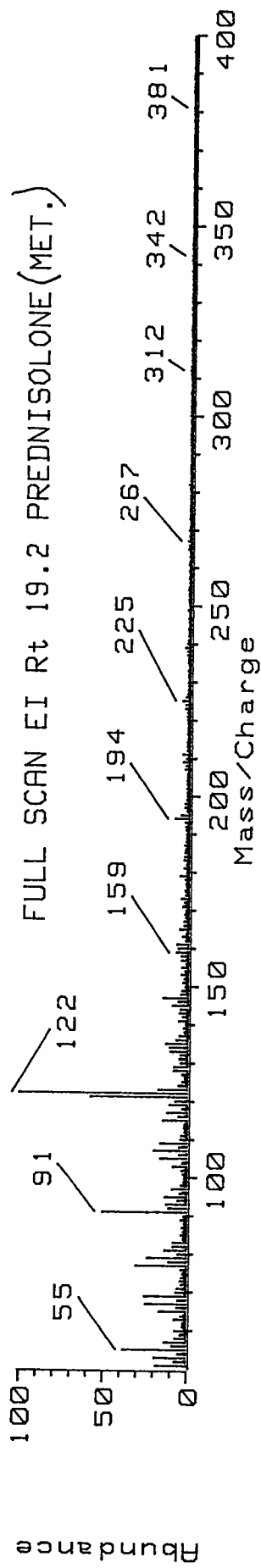


FIG. 13

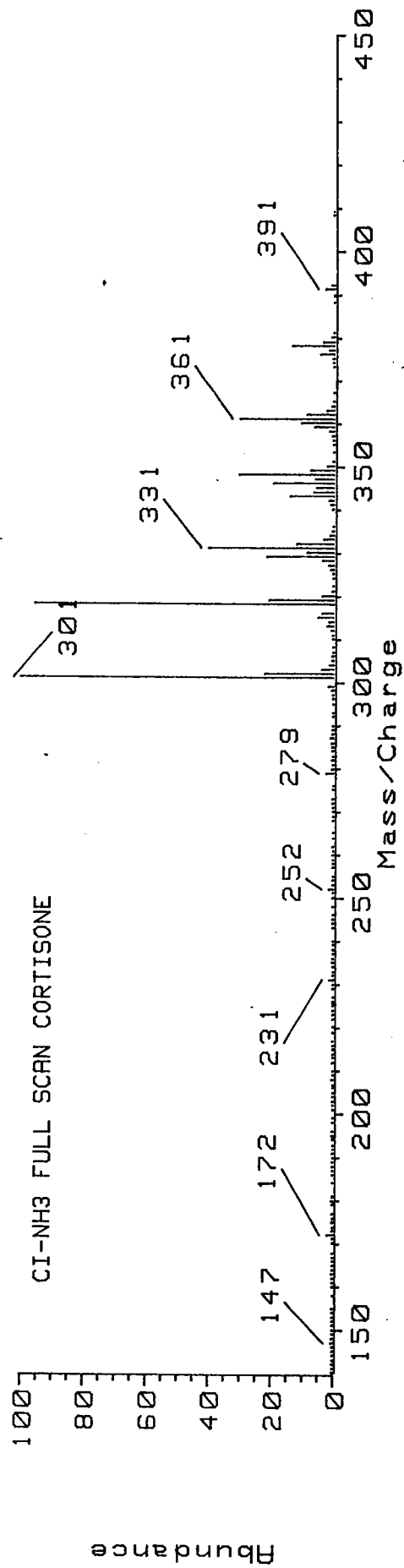
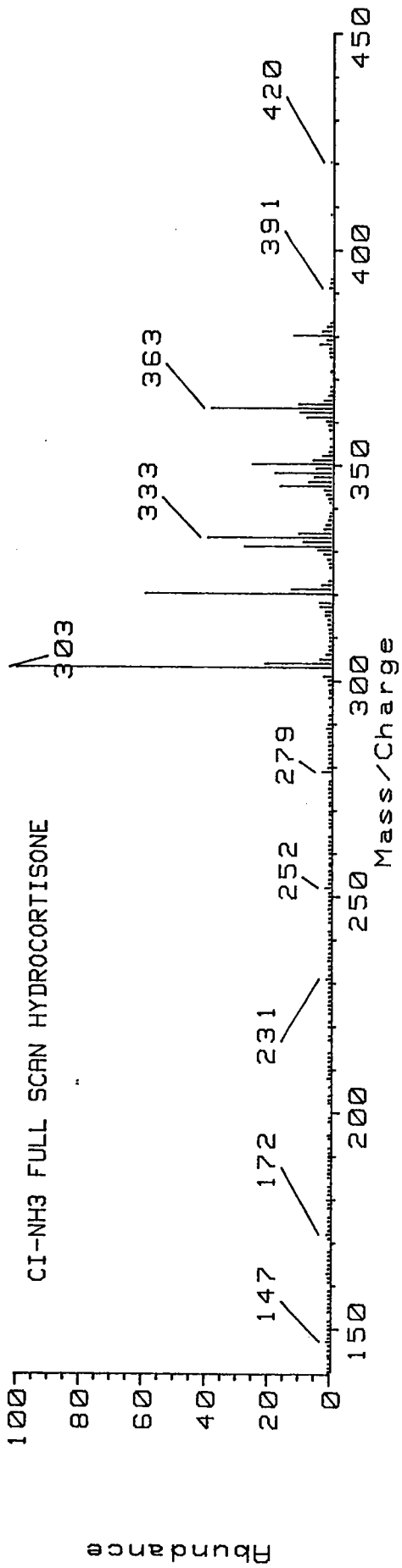


FIG.15