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Short communication

## High-performance thin-layer chromatographic determination of 5-methoxypsoralen in serum from patients

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### Abstract

A simple and rapid high-performance thin-layer chromatographic (HPTLC) determination of 5-methoxypsoralen in serum is necessary for the therapeutic survey of patients treated with Puvatherapy (psoralen+UV A). The assay for this biological fluid involves an extraction with heptane–dichloromethane (4:1, v/v). The analytical method is linear from 50 to 250 ng/ml. This assay range is adequate for analysing human serum, as it corresponds to psoralen concentrations measured in serum from patients treated with psoralen and UV A against psoriasis and vitiligo. The limit of detection is 15 ng/ml. The coefficient of variation was less than 7%. © 1997 Elsevier Science B.V.

**Keywords:** 5-Methoxypsoralen

### 1. Introduction

Psoralens are furocoumarins, used for the treatment of dermatological disease [1–4]. Up until now, psoralen in the serum of psoriatic patients has been determined by high-performance liquid chromatography (HPLC) [5], which is the most used technique for psoralen assessment in serum because of its high sensitivity and specificity. Nevertheless, this method requires a long analysis time. This paper describes a high-performance thin-layer chromatography (HPTLC) method that allows the simultaneous analysis of several serum samples.

### 2. Experimental

#### 2.1. Reagents

All chemicals were of the analytical grade. The following substances were used: 5-methoxypsoralen (5-MOP), trimethoxypsoralen (TMP) (Bergaderm, Paris, France), methanol (Carlo Erba, Rueil Malmaison, France), heptane and dichloromethane (Merck, Nogent sur Marne, France).

#### 2.2. Sample preparation

The psoralens were extracted from solvent according to the Stolk method [6]. To 1 ml of serum, 5 ml of heptane–dichloromethane (4:1, v/v) were added. After vortex-mixing for 7 min and centrifugation at

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5500 g for 2 min, 4 ml of the organic layer were transferred to a clean glass tube and evaporated to dryness at 50°C under a stream of nitrogen. The residue was dissolved in 40 µl of methanol under vortex-mixing for 40 s. A 3-µl volume was used for spotting.

### 2.3. Materials

Stationary phase: HPTLC plates RP<sub>18</sub>, (10×10 cm; ref. 13724) from Merck (Darmstadt, Germany) were used [7]. An automatic TLC sampler III (ATS III) Camag (Merck, Strasbourg, France) was used for spotting. HPTLC plate development was performed in a Camag HPTLC vario chamber from a (4:1, v/v) methanol–water solvent [7]. Plates were scanned in fluorescence mode (excitation, 313 nm; emission, 490 nm) with a TLC densitometer scanner II Camag.

### 2.4. Assay validation

No demixing problems were observed and the results were reproducible. The standard curves for serum were obtained by adding 5-MOP to drug-free biological fluids to obtain a concentration of 25–250 ng/ml, and TMP was added as an internal standard [8,9] at 100 ng/ml. No blanks were included in these standard curves. Standard serums were prepared under operating conditions as described above. Accuracy and precision of the 5-MOP assay in the serum were determined by analysing eight replicates for each of six concentrations (25, 50, 100, 150, 200 and 250 ng/ml).

## 3. Results and discussion

The chromatogram shown in Fig. 1 is of the serum from eight patients after 5-MOP and TMP extraction. At the starting spots, there was always residual fluorescence. These “impurities” could arise from serum extraction (Fig. 1).

The calibration curve for 5-MOP was linear over the concentration range of 50 to 250 ng/ml. This range corresponds to psoralen concentrations measured in serum from patients treated with PUVA (psoralen+UVA) against psoriasis and vitiligo. Below 50 ng/ml, the method was non-linear (Table

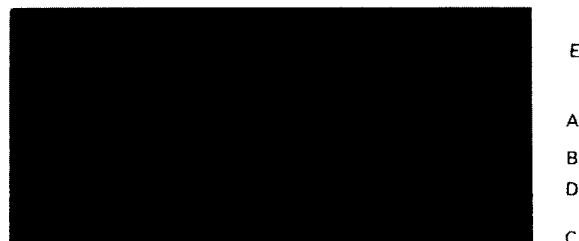


Fig. 1. Chromatogram of eight different serum samples. (A) Migration level of TMP; (B) migration level of 5-MOP; (C) migration level of the solvent (methanol–water); (D) migration level of an impurity of the serum; (E) starting position of the spot.

1). The equation determined from six different concentrations (experiments repeated eight times) was  $y = -53.238 \cdot 10^{-3} + 7.184 \cdot 10^{-3}x$  ( $r^2 = 0.983$ ), where  $y$  is the peak-area ratio of 5-MOP/TMP,  $x$  is the concentration of 5-MOP (in ng/ml) and  $r$  is the correlation coefficient. As usual, the detection limit, 15 ng/ml, was determined with the peak area equal to three times the background and the quantification limit was considered to be 10/3 of the detection limit. We determined only the response,  $y$ , the inter-day precision (expressed as a coefficient of variation, C.V.) and the inter-day accuracy of the 5-MOP assays in serum (Table 1).

The coefficients of variation were less than 7% ( $n=8$ ) (3% in HPLC) [5] over the concentration range 50–250 ng/ml. The accuracy lies between 97.5 and 107.2%. Pharmacokinetic analysis of serum from patients after oral intake of 5-MOP (1.2 mg/kg body weight) was carried on in respect of the calibration curve. The results are similar to those obtained by HPLC [5]. The worst difference between

Table 1  
Inter-day precision and accuracy ( $n=8$ )

Spiked concentration (ng/ml)	Response ( $y$ )	C.V. (%)	Accuracy (%)
25 <sup>a</sup>	0.12	11.28	72.84
50	0.33	2.52	99.78
100	0.67	3.16	100.00
150	0.98	2.23	97.55
200	1.33	6.88	99.40
250	1.80	4.36	107.25

<sup>a</sup>The concentration of 25 ng/ml is not in the linear range, which explains the bad values obtained for C.V. and accuracy.

5-MOP concentrations obtained using these two methods was 13%.

In summary, this newly developed HPTLC method for 5-MOP is highly sensitive and selective and has good reproducibility. It can be used to rapidly measure concentrations of 5-MOP at trace levels in serum for the optimal clinical management of patients. The simple HPTLC method enables, in a limited time, therapeutic drug monitoring to be performed (for the simultaneous analysis of between eight and twelve samples, the total analysis time required was a maximum of 30 min). This is not the case with HPLC, where analysis takes 10 min for a single sample.

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