Isolation, Purification and Structural Identification of Degradation Products from Ketoconazole in a Liquid Formulation: An HPLC and HPLC-MS/MS Study

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Abstract
Ketoconazole is the active ingredient in an antifungal liquid formulation for topical application. A binary gradient HPLC analysis indicated that ketoconazole in the formulation produced four degradation products at minor levels after storage at room temperature for about 3 years. These degradation products were isolated and purified using semi-preparative HPLC procedures followed by analysis using off-line HPLC-ESI-MS and HPLC-ESI-MS/MS. Structures of the degradation products were proposed based on isotope distribution profiles and fragmentation patterns of their molecular ions. Oxidation of ketoconazole appeared to be essential to generate these degradation products. Three of the four degradation products have not been reported in literature.

Keywords
Ketoconazole, Antifungal, Oxidation, Degradation, HPLC-MS, HPLC-MS/MS

Introduction
Ketoconazole (cis-1-acetyl-4-[2-(2,4-dichlorophenyl)-2-[(4H-imidazole-1-ylmethyl)-1,3-dioxolan-4-yl]-methoxy]phenyl)piperazine) (I) (Scheme I) is an imidazole antifungal reagent. The activity spectrum of ketoconazole covers a wide variety of yeasts, dermatophytes, and other fungi [1]. Ketoconazole has potential applications in therapy of dermatomycoses, candidiasis of mouth and vagina, systemic candidiasis, chronic mucocutaneous candidiasis and candiduria, as well as deep and subcutaneous mycoses by dimorphic and other fungi [2]. In contrast to other antymycotic drugs of imidazole type, ketoconazole shows good systemic absorption in animals and humans following oral administration of relatively low doses. Ketoconazole can also be administered topically or by intravenous infusion. In addition, there are some reports on efficacy of topical or oral ketoconazole in dermatological disorders not exclusively related to fungi. Ketoconazole has been shown to be highly effective in reducing erythema and hyperkeratosis [3], and even inhibiting lipopolysaccharide induced activation of nitric oxide synthetase gene in murine macrophage cell line J774 [4]. Interestingly, despite extensive applications and studies of ketoconazole, there is quite limited information published regarding degradation chemistry of ketoconazole although there are some scattered reports regarding its stability [5-7]. As for the biotransformation, Whitehouse LW and co-workers [8] investigated metabolites of ketoconazole from liver extract of male Swiss Webster mice; while Miller JD and co-workers [9] studied metabolites of ketoconazole incubated with cryo-preserved human he-}

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Scheme I: Imidazole antifungal reagent.
Experimental

Material and reagents

Ketoconazole used was a USP reference standard. 2-(2,4-dichlorophenyl)-2-(1H-imidazol-1-ylmethyl)-1,3-dioxalan-4-methanol (II) was obtained from Janssen Biotech through Research Diagnostics, Inc., Pleasant Hill Rd., Fairfield, NJ 07836. HPLC grade methanol, acetonitrile and chloroform were obtained from Fisher Scientific. Ammonium formate, methylene chloride, phosphoric acid and ammonium hydroxide were obtained from J.T. Baker. HPLC grade water was prepared using a Modular Polisher I HPLC Laboratory Reagent Grade Water System (Continental Water System Corp., Bensalem, PA).

Sample extraction

14 g of 1% ketoconazole liquid formulation stored for about three years at room temperature was suspended in 3 mL of methanol and 20 mL of methylene chloride. After stirring at room temperature for 0.5 hours, the mixture was kept at room temperature overnight to produce distinguishable organic and aqueous phases. The organic layer was isolated and evaporated by purging compressed nitrogen (10-20 psi, ~2 hours). The residue was subsequently suspended in 5 mL of methanol and kept at 4 °C for more than 1 hour. The undissolved solid was removed by filtration (Autovial Syringeless Filter, PTFE, 0.45 µm, Whatman) and the filtrate was used for HPLC isolation of the degradation products.

HPLC conditions

Two semi-preparative reversed-phase HPLC methods (HPLC I and HPLC II) and one analytical reversed-phase HPLC method (HPLC III) were developed. HPLC I was used for preliminary isolation of the degradation products. HPLC II was used for subsequent purification of the isolated degradation products and the liquid formulation as well as placebo. The liquid formulation and placebo were diluted into methanol and filtered prior to injection into HPLC system for analysis. HPLC I and HPLC II used a Ranin Rabbit HPLC system equipped with two HPX pumps, a Rhodyne injector, a Knauer variable wavelength monitor at 230 nm, a Waters GCM column oven at 40 °C, and a Develosil UG-3 ODS column (3 µm, 10 × 0.6 cm) with a Develosil UG-5 (1 × 0.4 cm) guard cartridge. For HPLC III, a HP1090L HPLC with detection at 230 nm with Agilent ChemStation for instrument control and diode array detection. Four HPLC mobile phases were used. Mobile phase A was prepared by adding 18.4 g of phosphoric acid (85%) into 4 L of deionized water and the pH was then adjusted to 9.0 by adding ammonium hydroxide (29.6%). Mobile phase B was prepared by mixing 3600 mL of methanol with 400 mL of mobile phase A. Mobile phase C was prepared by adding 3 mL of formic acid into 4 L of deionized water and then pH was adjusted to 4.0 by adding ammonium hydroxide. Mobile phase D was prepared by mixing 900 mL of methanol with 100 mL of mobile phase C. For HPLC I, the following binary gradient was used with mobile phases A and B: 0-1 min., 50% mobile phase A, 1-15 min., linear gradient to 61.6% mobile phase B, 15-45 min., linear gradient to 66.7% mobile phase B, 45-50 min., linear gradient to 100% mobile phase B. Flow rate was set as 1.7 mL/min. For HPLC II, the following binary gradient was used with mobile phases C and D: 0-1 min., 39% mobile phase D, 1-35 min., linear gradient to 60.6% mobile phase D, 35-40 min., linear gradient to 100% mobile phase D. Flow rate was 1.7 mL/min. For HPLC III, the following binary gradient was used with mobile phase A and methanol: 0-1 min., 45% methanol, 1-15 min., linear gradient to 55% methanol, 15-45 min., linear gradient to 60% methanol, 45-50 min., linear gradient to 90% methanol. Flow rate was 1 ml/min.

Off-line HPLC-MS

Mass spectrometric analysis was performed using a Finnigan MAT TSQ 700 triple quadrupole mass spectrometer with an ESI source (San Jose, CA, USA). The spray voltage was set to 4.5 kV and the heated capillary temperature was maintained at 200 °C. The instrument was optimized in the positive ion mode. For tandem spectrometric measurement, argon was used as the collision gas at a pressure of 2.5 mTorr, and the collision energy was maintained at ~40 eV.

The HPLC procedure for introducing samples into mass spectrometric detector used a narrow bore column (YMC BASIC, 250 × 2.0 mm, 5 µm particle size) and Waters Model 600E pump operated at 0.25 mL/min using mobile phases E and F. Mobile phase E was a solution of 5% acetonitrile, 1% methanol with 5 mM ammonium formate adjusted to pH 6.2. Mobile phase F was a solution of 45% acetonitrile, 1% methanol with 5 mM ammonium formate adjusted to pH 6.2. The elution used a binary gradient with 80% of mobile phase E for 3 min initially and then linearly changed to 100% mobile phase F at 25 min.

Results

Chromatographic analysis of the liquid formulation and isolation of the degradation products

HPLC chromatograms of the liquid formulation after stored for about 3 years at room temperature along with corresponding placebo are presented in Figure 1 using method HPLC III. The placebo has the same ingredients as the formulation but without ketoconazole. The liquid formulation and placebo were diluted using methanol and filtered prior to injection into HPLC for analysis. Peak I is due to ketoconazole and at least four suspected degradation products were observed, which were labeled as II, III, IV and V, respectively. Peak II was confirmed to be 2-(2,4-dichlorophenyl)-2-(1H-imidazol-1-ylmethyl)-1,3-dioxalan-4-methanol by comparing its identical HPLC retention time and online UV spectrum with that of an authentic compound under identical chromatographic conditions. The levels of II, III, IV and V were determined to be 0.08%, 0.18%, 0.45% and 0.29%, respectively, normalized with ketoconazole HPLC peak area%. Figure 1 also indicated that there is a peak at about 17 minute which is larger in the aged formulation than in the placebo; this peak co-eluted with the ingredients of placebo and remains to be studied. In this report, only degradation products II, III, V and...
0.2-0.5 mL aliquots of the formulation extract, prepared according to the procedure described in the experimental section of this report, were first chromatographed using HPLC I. The eluents at 12.4, 28.4 and 32.9 min. were collected individually and labeled as III', IV' and V', respectively. Figure 2 presents the analytical chromatograms of III', IV', and V' using HPLC III. A large amount of unwanted material had been removed and the number of chromatographic peaks decreased from more than 16 to less than four for each of the degradation product fractions. For further purification, 0.5-2 mL of III', or IV', or V' solutions were injected and eluted using method HPLC II. The eluents at 27.4, 29.7, and 33 min were individually collected and labeled as III, IV and V. Figure 3 shows the analytical chromatograms of III, IV and V to exhibit single peaks indicating pure degradation products obtained. Peak purity measurements using HP ChemStation indicated that peaks III,
IV and V were homogenous. The above isolation and purification sequence was repeated until adequate amounts of pure degradation products were obtained for following HPLC-MS studies.

**Figure 4** shows comparison of the analytical chromatograms of aged formulation, placebo, III, IV and V.

**HPLC-MS studies of III, IV and V**

The isolated degradants III, IV and V were analyzed offline using HPLC-ESI-MS in full scan mode and HPLC-ESI-MS/MS on identified molecular ions. The full scan mass spectra of III, IV and V are presented in **Figure 5a**, **Figure 6a** and **Figure 7a**.

Their mass spectra displayed characteristic isotope clusters due to the chlorine atoms and their molecular weights were determined as 602, 556 and 502 for III, IV and V with molecular compositions of C_{26}H_{20}N_{4}O_{9}Cl_{2}, C_{26}H_{22}N_{4}O_{6}Cl_{2} and C_{24}H_{24}N_{4}O_{4}Cl_{2}, respectively. The isotope distribution profiles of the protonated molecular ions are consistent with the theoretical isotope distribution profiles. In order to obtain additional structural information, the fragmentation patterns of the molecular ions of III, IV and V were investigated using HPLC-MS/MS. **Figure 5b**, **Figure 6b** and **Figure 7b** present the observed diagnostic fragments from the corresponding molecular ions along with the proposed structures of III, IV and V. The proposed structures are tentative and were based only on HPLC-MS and HPLC-ESI-MS/MS studies. Due to the complexity of the formulation matrix and the minor levels of the degradants formed plus limited amount of the liquid formulation sample, it was not possible to obtain adequate amounts.
Figure 5a: Full Scan Mass Spectrum of III.

Figure 5b: MS/MS Diagnostic Fragments and Proposed Structure of III.
Figure 6a: Full Scan Mass Spectrum of IV.

Figure 6b: MS/MS Diagnostic Fragments and Proposed Structure of IV.
**Figure 7a:** Full Scan Mass Spectrum of V.

**Figure 7b:** MS/MS Diagnostic Fragments and Proposed Structure of V.
of the pure degradation products for additional structural investigation, such as NMR studies.

**Discussion**

In this report, four degradation products at minor levels from ketoconazole (I) in a liquid formulation sample after storage at room temperature for about 3 years were isolated using semipreparative HPLC and tentatively identified using HPLC-ESI-MS and HPLC-ESI-MS/MS analyses. Three of the four degradation products have not been reported in literature. Based on the comparison between the structures of ketoconazole and that of degradation products II, III, IV, and V, it is concluded that II might be formed from hydrolysis of ketoconazole. However, to generate III, IV and V, likely a series of oxidation, hydroxylation and O-dealkylation need to occur. Oxidation of ketoconazole must be essential. The substituted aromatic amine nucleus of ketoconazole has a pair of free electrons to make it susceptible to oxidation. As a matter of fact, oxidation, cleavage of the imidazole and piperazine rings, O-dealkylation, and hydroxylation were reactions observed in the biotransformation of ketoconazole to yield various metabolites [8,9] although structures of the reported metabolites are different from II, III, IV and V. Besides directly hydrolysis of ketoconazole degradation product II might also be generated from hydrolysis of oxidized ketoconazole since its formation was accelerated after long time storage. A proposed potential mechanism for the formation of the degradation products are presented in Scheme I. No safety evaluation was performed on the degradation products since within 18 month shelf life the degradation product levels were less than 0.1% and the liquid formulation is for topical rinse off application with low systemic absorption.

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