

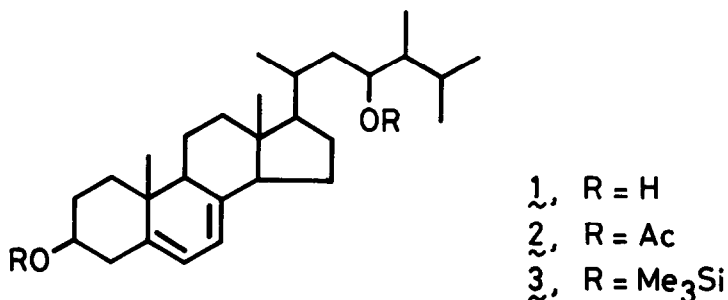
ISOLATION OF ERGOSTA-5,7-DIENE-3 β ,23 ξ -DIOL FROM YEAST

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Abstract: A novel hydroxylated sterol was isolated from a *Saccharomyces cerevisiae* mutant defective in the Δ^{22} -desaturation of sterol side chain, and it was identified as ergosta-5,7-diene-3 β ,23 ξ -diol.

In spite of the development of studies on sterols in yeast, scanty reports have been published on hydroxylated sterol(s) in it. In this communication, we report the isolation of a novel hydroxylated sterol from a *Saccharomyces cerevisiae* mutant N22,¹ which is defective in the Δ^{22} -desaturation of sterol side chain. The compound was identified as ergosta-5,7-diene-3 β ,23 ξ -diol (1).



From N22 cells grown on a synthetic medium without shaking, sterols were extracted and acetylated.^{1b} One of the acetates which were detected in fraction I,^{1b} on AgNO₃-TLC was shown to have a longer retention time than most sterols on GLC. The compound was suggested to be a diacetate ester (2)² of hydroxylated C₂₈-sterol by GC-MS. Compound 2 was further purified by TLC on silica gel plate (solvent, PhH-MeOH = 98 : 2; R_f, 0.51) and was hydrolyzed to obtain 1.³ UV spectrum³ of 1 showed absorption bands specific for $\Delta^{5,7}$ -conjugated diene.⁴ NMR spectrum³ of 1 showed that 1 had 3 β -hydroxyl group. Moreover, the incubation of N22 cells with [methyl-¹⁴C]-L-methionine (1.83 x 10⁷ dpm, 53 Ci/mol) as described in the previous paper^{1b} enabled the incorporation of radioactivity into 1 (2.85 x 10⁴ dpm; specific radioactivity, 1.0 Ci/mol). This indicates that 1 has C-28 carbon⁵ which is known to originate from the methyl group of L-methionine. In addition, an intense peak⁶ of m/z 173 corresponding presumably to fragment

$C_{21}H_{42}OSi$ formed by the cleavage of the bond between C-22 and C-23 was observed in MS of di-trimethylsilyl ether (**3**).⁷ This indicates that **1** has a hydroxyl group at 23 position. Other analytical data were also consistent with the literature values⁷ of synthetic (23RS)-ergosta-5,7-diene-3 β ,23-diol.

The cellular contents of **1** in strain M10, the parent strain of N22, and in a wild-type *S. cerevisiae* ATCC 12341, if present, were less than 2% of that found in N22 (about 70 μ g/g dry cells). This suggests that the accumulation of **1** in N22 cells is due to the genetic characteristics of the mutant. Recently we reported the involvement of a cytochrome P-450 in the Δ^{22} -desaturation.⁸ That paper seems to be the first one reporting the involvement of cytochrome P-450 in desaturation reaction. The Δ^{22} -desaturation is considered to proceed in two stages — hydroxylation of sterol accompanied by dehydration of the product. It seems that **1** accumulates owing to the lack of the enzyme involved in the dehydration.

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References and Notes

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2. Retention time of **2** relative to cholesteryl acetate on 3% SE-30 column was 2.6. m/z 498 (M^+), 438 (100%), 378, 363, 253.
3. Analysis of **1** by GLC on 3% SE-30 column showed that purity of **1** (retention time relative to cholesterol, 2.3) was about 90%. Analytical data of **1**: λ_{max}^{EtOH} 271 (ϵ 13,800), 282 (15,000), 293 (8,700); m/z 414 (M^+), 381 (100%), 211; 1H NMR (400 MHz, $CDCl_3$) δ 3.63 (br m $C_{3\alpha}H$ and $C_{23}H$), 5.39 (m C_7H), 5.57 (m C_6H); ν_{max} (KBr) 3350 (OH), 1120 (C-O), 830 (trisubstituted double bond), 800 (trisubstituted double bond).
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6. Retention time of **3** relative to cholesteryl trimethylsilyl ether on 3% SE-30 column was 2.4. m/z 558 (M^+), 453, 397, 307 (100%), 251, 173 (68%).
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