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Phytochemistry, Vol. 23, No. 4, pp. 921-923, 1984. Printed in Great Britain.

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24α -METHYL- 5α -CHOLEST-7-EN- 3β -OL FROM SEED OIL OF *HELIANTHUS ANNUUS*

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(Received 6 September 1983)

Key Word Index—Helianthus annuus; Compositae; seeds; 24-alkylsterols; 24α -methyl- 5α -cholest-7-en- 3β -ol.

Abstract—24-Methyl- 5α -cholest-7-en- 3β -ol (24-methyllathosterol) isolated from the seed oil of *Helianthus annuus* was shown to have 24α -configuration by ¹H NMR spectroscopy. The stereochemistry at C-24 of some other 24-alkylsterols isolated from this plant material also was determined.

24-Methylcholest-5-en-3 β -ol (24-methylcholesterol, 6) [1-3] and 24-methyl-trans-22-dehydrocholesterol (9) [4-6] found in higher plants often consist of a mixture of epimers at C-24, i.e. campesterol (24a) and 22-dihydrobrassicasterol (24 β) for 6, and trans-22-dehydrocampesterol (24 α) and brassicasterol (24 β) for 9. 24-Methyl- 5α -cholest-7-en-3 β -ol (24-methyllathosterol, 1), a Δ^7 analog of 6, also occurs in some higher plants including Compositae plants such as Helianthus annuus and Carthamus tinctorius [7]. However, the stereochemistry at C-24 of this sterol (1) from these higher plants remains undetermined to the best of our knowledge. Therefore, we have conducted here the determination of C-24 stereochemistry with the aid of 400 MHz ¹H NMR spectroscopy for sterol 1 and some other 24-alkylsterols isolated from the seed oil of H. annuus and have undertaken thorough analysis of the total sterol fraction of this plant material.

The Δ^5 - ($R_f = 0.33$, fraction A, 291 mg) and Δ^7 - ($R_f = 0.31$, fraction B, 50 mg) sterol fractions, separated by silica gel TLC from the unsaponifiable lipid (1.5 g) of H. annus seed oil (160 g), were acetylated, and the resulting acetate fractions were subjected to silver nitrate-silica gel TLC. The acetylated fraction A gave three further fractions. Fraction A-1 from the least polar band ($R_f = 0.60$) was a mixture of the acetates of 6 and 24-ethylcholesterol (7). Fraction A-2 from the band of $R_f = 0.44$ was 24-

ethyl-trans-22-dehydrocholesteryl (8) acetate. Fraction A-3 from several bands of $R_f = 0.1-0.4$ consisted of the acetates of 24-methyl-trans-22-dehydrocholesterol (9), 24Z-ethylidenecholesterol (isofucosterol, 11) and 24methylene-25-methylcholesterol (12). The acetylated fraction B separated into two fractions on argentation TLC. Fraction B-1 from the least polar band $(R_f = 0.60)$ consisted of the acetates of 1 and 24-ethyllathosterol (2). Fraction B-2 from several bands of $R_f = 0.1-0.4$ was a mixture of the acetates of 24-methylenelathosterol (episterol, 3), 24Z-ethylidenelathosterol (Δ^7 -avenasterol, 4) and 24-ethyl-24(25)-dehydrolathosterol (peposterol, 5). The identifications were performed by GC and GC/MS analyses of the acetate derivatives. Acetate fraction A-2 was then hydrolysed to give free sterol 8 (mp 163-166°). Acetate fractions A-1 and B-1 were also hydrolysed and further fractionated by reverse-phase HPLC. Fraction A-1, on HPLC, yielded 6 ($RR_t = 1.10$, mp 145-148°) and 7 $(RR_t = 1.18, \text{ mp } 136-140^\circ), \text{ and fraction B-1 gave 1 } (RR_t = 1.18, \text{ mp } 136-140^\circ)$ = 1.09, mp 144-146°) and 2 ($RR_t = 1.18$, mp 144-148°). In order to determine the configuration at C-24 of 1 as well as 2 and 6-8, these sterols were subjected to 400 MHz ¹H NMR spectroscopy. Table 1 shows the chemical shifts of the methyl group signals of these sterols accompanied with three authentic sterols, 24β -methyllathosterol, 24α ethyl-trans-22-dehydrocholesterol (stigmasterol) and its 24β -epimer (poriferasterol), for which assignments were 922

Table 1. Methyl group ¹H NMR chemical shifts (400 MHz, CDCl₃)* of some 24-alkylsterols

Sterol	H-18 (s)	H-19 (s)	H-21 $(d, J = 6.5)$	H-26 $(d, J = 7)$	H-27 $(d, J=7)$	H-28 $(d, J=7)$	H-29 $(d, J=7)$
24-Methyllathosterol (1)	0.536	0.800	0.916	0.852	0.804	0.775	
24β-Methyllathosterol†	0.532	0.796	0.924	0.857	0.784	0.778	
24-Ethyllathosterol (2)	0.536	0.796	0.926	0.837	0.814	_	0.846
24-Methylcholesterol (6)	0.680	1.009	0.911‡	0.851	0.803	0.772	_
			0.919‡	0.855	0.783	0.774	
24-Ethylcholesterol (7)	0.680	1.009	0.921	0.834	0.813	_	0.845
24-Ethyl-trans-22-dehydrocholesterol (8)	0.700	1.013	1.022	0.847	0.797		0.806
Stigmasterol (24a)†	0.699	1.012	1.021	0.846	0.797	_	0.805
Poriferasterol (24β)†	0.697	1.011	1.025	0.844	0.791		0.811

^{*}Chemical shifts given in δ values from TMS; coupling constants in Hz.

made by comparison with literature data [8-10]. Stereoisomers at C-24 of 24-methylsterols are reported to show significantly different chemical shifts for the methyl doublets of H-21 and H-27 in the high-resolution ¹H NMR [1,8–10]. The 24-methyllathosterol (1) gave H-21 and $\bar{\text{H}}$ -27 methyl doublets at δ 0.916 and 0.804, respectively, which were consistent with the corresponding signals reported for the 24α-epimer [8]. Taking into consideration that the authentic 24β -epimer had quite different chemical shifts for these doublets (H-21, δ 0.924; H-27, δ 0.784), sterol 1 was reasonably identified as 24 α methyllathosterol. 24-Methylcholesterol (6) was, on the other hand, shown to be a mixture of 24α - (H-21, δ 0.911; H-27, δ 0.803) and 24*B*- (H-21, δ 0.919; H-27, δ 0.783) epimers. Based on the relative heights of the H-21 doublet arising from each of the two diastereoisomers, the relative ratio was estimated as $24\alpha : 24\beta = 7 : 3$. The methyl group proton signals of sterol 8 were consistent with those of the authentic stigmasterol (24a) and differed enough in the chemical shifts of H-27 and H-29 methyl signals to distinguish it from its 24β -epimer (poriferasterol). Hence, sterol 8 was identified as stigmasterol. 24-Ethylsterols 2 and 7 showed almost identical ¹HNMR spectra with those of the authentic 24a-ethyllathosterol (22-dihydrospinasterol) and 24\alpha-ethylcholesterol (sitosterol), respectively, described in our previous article [11]. Furthermore, the ¹H NMR spectra of these H. annuus sterols were considerably different from those of the corresponding 24β -epimers, 24β -ethyllathosterol and 24β ethylcholesterol [11], and hence sterols 2 and 7 were regarded as the 24α-epimers, i.e. 22-dihydrospinasterol and sitosterol, respectively. The quantitative composition of the total sterol fraction of H. annuus seed oil was determined from the GC, argentic TLC and ¹H NMR data as 0.6% (1), 7.3% (2), 0.2% (3), 4.6% (4), 0.1% (5), 6.2% (6, 24α), 2.7% (6, 24β), 60.2%(7), 8.4% (8), 0.1% (9), 1.5% (10), 5.9% (11), 0.5% (12), and 1.7% of several unidentified components.

Thus, 24-methyllathosterol (1) isolated from H. annuus seed oil was identified as the 24α -epimer. 24-Methylcholesterol (6) was, on the other hand, shown to be a mixture of the 24α - and 24β -epimers, consistent with the previous observations on several other higher plants [1,8-10]. Three 24-ethylsterols, 2, 7 and 8, were indicated

to have the 24α -configuration. Sterol 9 remained stereochemically undetermined in this study. 24-Methylene-25-methylcholesterol (12) identified here as a minor sterol was recently isolated for the first time from *Brassica juncea* and its unusual side chain structure has been characterized [12].

EXPERIMENTAL

Mps were uncorr. Silica gel TLC (0.5 mm) was developed ×3 with n-hexane-EtOAc (6:1), and AgNO₃-silica gel (1:4) TLC (0.5 mm) was developed \times 4 with CH₂Cl₂-CCl₄ (1:5). HPLC was carried out on a Partisil 5 ODS-2 column (Whatman, 20 cm × 8 mm i.d.; packed by Erma Optical Works, Tokyo) using a UV detector monitoring at 212 nm (mobile phase, MeOH). RR, on HPLC was expressed relative to cholesterol. GC on OV-17 SCOT glass capillary column was under the conditions already described [13]. MS (70 eV) were taken with a GC/MS (2 % OV-17, 2 m × 3 mm i.d. glass column). ¹H NMR (400 MHz) spectra were determined in CDCl₃ with TMS as internal standard. Other techniques used in this study have been described previously [4]. Degummed H. annuus seed oil was courteously supplied by Dr. A. Nishioka (Ajinomoto Co., Yokohama). Origin and GC data of sterol 12 [12] and the other sterols [13] used here as the reference specimens have been described in our previous articles.

Acknowledgements—We thank Dr. Y. Fujimoto, Inst. Phys. Chem. Res. (Saitama), for ¹H NMR spectra. Our thanks are also due to A. Maeda for technical assistance.

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[†] Authentic sterols.

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Phytochemistry, Vol. 23, No. 4, pp. 923-925, 1984. Printed in Great Britain.

0031-9422/84 \$3.00 + 0.00 © 1984 Pergamon Press Ltd.

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SPRUCE DIE-BACK: ISOLATION OF *p*-HYDROXYACETOPHENONE FROM DISEASED SHOOTS OF *PICEA ABIES*

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(Revised received 29 September 1983)

Key Word Index—*Picea abies*; Pinaceae; Norway spruce, spruce die-back; forest decline; shoots; identification; *p*-hydroxyacetophenone.

Abstract—p-Hydroxyacetophenone as a new growth substance exhibiting inhibitory activity was isolated from shoots of *Picea abies* damaged by die-back. Its identity was established by chromatographic and spectroscopic methods.

INTRODUCTION

In continuation of our search for the biologically active compounds present in Norway spruce damaged by spruce die-back, we detected a number of compounds differing both qualitatively and quantitatively from healthy plants [1-3]. The present investigation was directed to the identification of growth inhibitors present in the neutral diethyl ether extract of diseased shoots.

RESULTS AND DISCUSSION

About 2.7 kg of diseased Picea abies shoots (10 year old trees) were harvested, extracted and purified according to the procedure described [1]. The neutral diethyl ether extract was subjected to silica gel partition chromatography $(32 \times 2.4 \text{ cm})$ to give a strong phytotoxic zone (wheat and rice seedling bioassay), which was eluted with chloroform-ethyl acetate, 9:1-3:1. The phytotoxic zone was purified on DEAE-Sephadex A-25 (132 \times 1.7 cm) to produce a strong phytotoxic fraction (eluted with 80% MeOH). TLC of this fraction after double developments (SS 1 and SS 2) yielded a strong biologically active fluorescent quenching band ($R_f = 0.46-0.96$). Re-chromatography of this band after double developments (SS 3 and SS 4) using TLC gave a strong phytotoxic zone (R_f = 0.37-0.73). This phytotoxic zone was further purified by TLC ($R_f = 0.61-0.91$, SS 5). The purified phytotoxic zone was subjected to silica gel partition column and the subsequent bioassay showed a main active zone (eluted with n-hexane-chloroform, 7:3). The main active zone was purified further by silica gel adsorption chromatography using the same gradient as described for partition

chromatography. The adsorption chromatography yielded a crystalline substance, which was sparingly soluble in water and thus, purified again with repeated washings with water. The purified substance was again subjected to TLC (silanized silica gel F_{254} ; SS 6, $R_f = 0.05-0.35$). The HPTLC (silica gel F_{254} : SS 6, $R_f = 0.19$) showed a single strong fluorescent spot, which gave an intense purple coloration after treatment with 5% ethanolic sulphuric acid and subsequent heating at 120° for 30 min. This spot was found to be identical with authentic p-hydroxyacetophenone. UV, IR, mass and 1H NMR spectroscopy further confirmed the identity of the isolated compound.

This is the first identification of p-hydroxyacetophenone as a growth substance from the diseased shoots of Picea abies. In the past p-hydroxyacetophenone has been synthesized and is still used for the synthesis of different compounds of industrial importance. The first isolation of p-hydroxyacetophenone was reported from the ether soluble oil of buds of *Populus trichocarpa* [4], the plant growth regulating activity of which was not reported. At lower concentrations, p-hydroxyacetophenone was found to show weak gibberellin-like activity as observed in the dwarf rice seedling bioassay (Table 1). At higher concentrations, p-hydroxyacetophenone inhibited the growth of rice (roots) and wheat seedlings (> 809 μ g/5 seedlings), and showed needle-browning and death of spruce seedlings (3 weeks old, $> 81 \mu g/plant$; 6 weeks old > 132 μ g/plant, Table 1) and saplings (2 years old, $> 324 \mu g/plant$) when applied to the tip of the apical growing part. Spruce seedlings (3 weeks old) showed dramatically the needle-browning ca 1 hr after application-time. The application of $> 1200 \mu g$ via roots