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A NEW SERIES OF NATURAL PROSTAGLANDINS IDENTIFICATION OF 1a,1b-DIHOMO-2,3-DIDEHYDRO-PGE₂ IN RAM SEMINAL VESICLES

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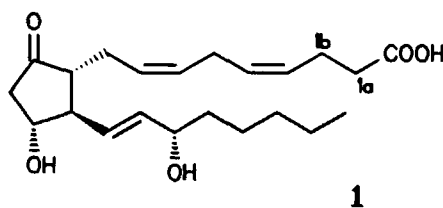
Abstract: A novel natural E-prostaglandin - 1a,1b-dihomo-2,3 didehydro-PGE₂ (**1**) was isolated from ram seminal vesicles and its structure was elucidated by MS and NMR analysis.

Key Words: Ram seminal vesicles, natural E-prostaglandin

Ram seminal vesicles (RSV) contain a large amount of the primary E-prostaglandins, viz. PGE₁, PGE₂ and PGE₃¹, derived from dihomog- γ -linolenic (C_{20:3} ω 6), arachidonic (C_{20:4} ω 6) and eicosapentaenoic (C_{20:5} ω 3) acids, respectively. In addition, a strong selective inhibitor of platelet aggregation, 5,6-dihydro-PGE₃², was recently described in the same tissue³. The present work reports the identification of a new natural prostaglandin, 1a,1b-dihomo-2,3-didehydro-PGE₂ (**1**), in RSV. Its obvious precursor, all cis-4,7,10,13,16-docosapentaenoic acid (C_{22:5} ω 6), can be biosynthesized in mammals by the chain elongation of arachidonic acid and subsequent delta-4 desaturation of adrenic acid (C_{22:4} ω 6). Previously it was found that both the docosaenoic acids mentioned above are present in the phospholipids of seminal vesicles⁴ and, therefore, they can serve as natural precursors of prostaglandins *in vitro*⁵. Furthermore, it was shown that RSV microsomes convert adrenic acid to 1a,1b-dihomo-prostaglandins E₂ and F_{2 α} *in*

*vitro*⁶. Thus, it seemed plausible that C₂₂ω6 prostaglandins occur in RSV in detectable amounts.

The E-type prostaglandins were separated from the lipid fraction⁷ of RSV (5 kg) on silica gel using a MeOH/CHCl₃ gradient. The PGE fraction (1.1 g) eluted with 3.5% MeOH in CHCl₃ was fractionated by repeated normal and reversed phase preparative HPLC⁸ to give, in addition to PGE₁, PGE₂, PGE₃, 1a,1b-dihomo-PGE₂, 5,6-dihydro-PGE₃, 8-iso-PGE₁, 8-iso-PGE₂, 8-iso-PGE₃, 11β-8-iso-PGE₁, 15β-PGE₂ and 15β-PGE₁, also 5.2 mg of pure compound **1** as colorless oil, [α]_D²² -31° (c 0.6, CHCl₃). **1** was converted with 0.4 N KOH/MeOH to the corresponding PGB compound with ε=27,000 at λ_{max}=278.



The molecular formula C₂₂H₃₅O₅ was established by negative ion CI-MS (details of MS study are given below) and NMR study⁹.

To elucidate the structure of **1** mass-spectrometrically, the PFB-(TMS)₃-9-enol derivatives of **1** and authentic 1a,1b-dihomo-PGE₂, PGE₂, PGE₁, PGE₃ and 5,6-dihydro-PGE₃ were prepared. The NICI spectrum of 9-enol-**1**-(TMS)₃-PFB showed a base peak at m/z 593[M-PFB] and two minor peaks 503[M-(PFB+90)] and 413[M-(PFB-2*90)], indicating the presence of one carboxylic and two hydroxyl groups. The EI-MS of the same derivative of **1** showed a molecular ion at m/z-774, as well as ions at m/z 703[M-71]⁺, 684[M-90]⁺, 613[M-(71+90)]⁺, 594[M-2*90]⁺, 523[M-(71+2*90)]⁺, 504[M-3*90]⁺, 483, 441[M-(C1-C7)]⁺, 352, 313, 269, 217 (C9-C11), 181 (PFB), 173 (C15-C20), 73 (a base peak). The fragmentation pattern was similar to that of 1a,1b-dihomo-PGE₂ with the exception of the fact that all high-mass ions above m/z 441 were two mass units lower than the corresponding ones in the spectrum of the latter, indicating the presence of an additional double bond in the molecule of **1**. The presence of ions [M-71]⁺, [M-(71+90)]⁺ and 173 demonstrates that the ω-chain of **1** is identical to that of

1a,1b-dihomo-PGE₂ and PGE₂. The fragment ions at m/z 441 (loss of C1-C7) observed in both the spectra indicate that an additional double bond of compound **1** is located in the α -chain. The fragment [M-333]⁺ (loss of α -chain) was dominating in the spectrum of the TMS-PFB derivative of B type PG formed from **1** in alkali.

The location and configuration of the double bonds and substituents in the molecule of **1** were established by 2D ¹H-¹H and ¹H-¹³C COSY correlations⁹ on a Bruker AMX-500 instrument. Data on PGE₁, PGE₂ and other prostaglandins^{8,10} confirm unambiguously the substitution pattern and relative configurations of substituents on a 5-membered ring, the 15-hydroxyl group and the C-13,14 E double bond (³J_{HH}=15.1 Hz). The positions of the two remaining double bonds were elucidated from 2D correlation diagrams. The Z-configuration of the Δ 2 bond follows from the carbon chemical shifts of model 4Z- and 4E-nonenoic acids (cf. C-1b at 22.7 ppm with 22.6 for Z- and 27.8 for the E-isomer of the corresponding acid)¹¹. The Z-configuration of the Δ 5 bond is confirmed by the carbon chemical shift of C-7 at 25.1 ppm (cf. C-7 of PGE₂ at 25.2 ppm^{8,10}) and by the chemical shift of C-4 at 25.5 ppm (cf. C-10 of 8Z,11Z,14Z-hexadecatrienoic acid at 25.7 ppm¹²).

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