Determination of the Origin of Urinary Noranadrosterone-Traces by Gas Chromatography Combustion Isotope Ratio Mass Spectrometry

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Introduction

The steroid 19-norandrosterone (19NA) is an ongoing problem in doping control: On the one hand, it is, besides 19-noretiocholanolone (19NE), the most abundant metabolite of the potent synthetic anabolic steroid 19-nortestosterone (19NT)^[1]. On the other hand, small amounts are biosynthesized by pregnant women and further evidence exists for physiological origin of this compound. Therefore, the World Anti Doping Agency (WADA) introduced a threshold concentration of 2 ng 19NA per mil urine to discriminate 19NT abuse from biosynthetic origin^[2]

Recent findings show however, that formation of 19NA resulting in concentrations above the threshold level might be due to demethylation of androsterone (A) in urine. This steroid is excreted by healthy persons in milligram amounts per day[3]

To avoid false positive results, the origin of unirary 19NA in doping control samples could be elucidated by ¹³C/¹²C-analysis. Synthetic 19NT is derived from C₃-plants by partial synthesis and shows $\delta^{13}C_{VPDB}$ -values of around -28‰. Endogenous steroids are less depleted in ¹³C due to a dietary mixture of C₃- and C₄-plants. Comparing $\delta^{13}C_{VPDB}$ -values of A as endogenous reference compound (ERC) with19NA should endogenet unambiguity to the string of units of the string of the string. MA

should reveal unambiguously the origin of urinary NA.

Samples

11 urine samples containing 19NA from pregnant women and from excretaion studies with different prohormones were analyzed as control samples as well as blank urines spiked with 19NA-glucuronide in different concentrations. 25 19NA-containing samples from routine analysis were then analyzed.

Measurement

¹³C/¹²C-analysis

- Injection Splitless mode: manually up to 3 µl out of at least 5 µl methanol HP 5890 II
- Optima 83 (Macherey/Nagel); 20 m x 0.25 mm ID; film thickness 0.25 µm
- GC Combustion Interface II (Finnigan MAT) oxidation reactor (CuO) at С 940°C

IRMS Delta C (Finnigan MAT)

GC/MS

A similarly equipped GC/MS was used to check each sample for coelutions





 $\delta^{13}C_{VPDB}$ -values of 19NA vs. A of doping control samples Fig. 5



IDG

metabolism of selected steroids and illustration of expected $\delta^{13}C_{VPDB}$ -values Fig. 1: Sample Preparation

After solid phase extraction of 10 ml urine, steroid glucuronides were hydrolyzed Arter solid phase extraction of no milline, steroid glucotindes were hydrolyzed enzymatically and separated at alkaline pH with tert-butyl methyl ether. A semipreparative normal phase HPLC on a $(CH_3)_2N$ -propyl column was applied for purification to separate 19NA and A from E. 19NA and A were further purified and separated by HPLC on C_{18} -material. GC/C/IRMS-measurements were performed without derivatization







 $\delta^{13}C_{\text{VPDB}}\text{-values of 19NA vs.}$ A of control samples from pregnant women Fig. (endogenous) and from excretion studies (exogenous)



Results

Fig. 2 shows GC/C-chromatograms of a blank urine (top) and a 19NA-spiked urine (bottom). Especially the ratio 45/44 indicates, that no confounding coelution is present. All samples were Fig. 4 shows 6¹³C/_{VPDB}-values of 19NA vs. A of control samples. Endogenous values derive from urine of pregnant women whereas exogenous values are from urine of exoretion studies, which is acceptable for concentrations down to 2 ng 19NA vs. A of control samples. Endogenous values derive from urine of pregnant women whereas exogenous values are from urine of exoretion studies, with prohormones. The data obtained allow to discriminate steroid abuse from biosynthetic origin. Fig. 5 is similar to fig. 4, but 19NA-containing doping control samples are presented. Normally, A was the ERC; in three cases, 110H-A was used as reference compound because the δ-value of A was influenced by the abuse of other hormones. Obtained data reveals steroid abuse unambiguously. Fig. 6 shows the Δδ-values of 19NA and ERC of the same samples vs. the concentration of NA. Demethylating activity is marked, if present as well as a heuristic threshold value.

Conclusions

Data of urine samples are presented with a 19NA concentration of more than 5 ng/ml which don't show (any more) demethylating activity but endogenous $\delta^{13}C_{VPDB}$ values. To avoid false positive results in doping control, IRMS is obligate to distinguish between doping or not for 19NA-traces

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