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1	Evolution of steroid concentrations in saliva from immature to pubertal gilts
2	for the identification of biomarkers of gilts receptivity to boar effect.
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25	
26	Abstract

1

Estrus synchronization is necessary for management of gilt reproduction in pig farms. It is usually 27 achieved by using synthetic progestagens, but there is increasing demand for non-hormonal 28 alternative tools with the prospect of sustainability of livestock production. Moreover, in organic 29 30 farms, synthetic hormones are not allowed. Before reaching puberty, gilts exhibit a "waiting period" during which external stimulations, such as boar exposure, could trigger and synchronize the first 31 ovulation. However, practical non-invasive tools for detection of the "waiting period" in pig farms 32 33 are lacking. During this period, estrone levels in urine are high, but urine sampling is difficult in 34 group-housed females. Our objective was to identify among steroids potential biomarkers of this "waiting period" through saliva monitoring from immature to pubertal gilts using gas 35 36 chromatography coupled to tandem mass spectrometry.

Starting between 144 to 147 days of age, six Large White gilts were submitted to ultrasound 37 puberty diagnosis 3 times a week until first ovulation. Urine and saliva samples were collected to 38 analyze weekly estrone and steroidome respectively, until puberty. Urinary estrone concentration 39 significantly increased 2 weeks before first ovulation occurring between 182 and 192 days of age. 40 The period with increasing estrone levels was considered as the "waiting period". Steroidome 41 analysis allowed identifying and quantifying 28 steroids in 500 µl of gilts saliva. Significant 42 decrease of dehydroepiandrosterone and significant increase of 5\alpha-dihydroprogesterone and 17\beta-43 estradiol were detected 2 weeks before puberty, suggesting that these steroids could be potential 44 biomarkers of the "waiting period". 45

46 These results show that painless sampling of saliva could be a non-invasive welfare-friendly tool for 47 the identification of the physiological hormonal status of the gilts and possibly the optimal time for 48 application of the boar effect, a solution to synchronize puberty without exogenous hormones.

49

50 Keywords: steroidome, saliva, puberty, porcine, male effect, GC-MS/MS

51

52 Introduction

2

In pig farms, batch management of sows has developed extensively due to advantages in labor 53 planning, possibilities of piglet adoption when litter size is excessive, all-in all-out practices and 54 health management. The importance of the control of estrous cycle has increased in recent years 55 56 because batch management requires groups of service-ready females for artificial insemination at planned times. Use of synthetic progestagens alone or in combination with gonadotropins has been 57 commonly used to synchronize estrous cycle in gilts in swine farms. However, the wide use of 58 59 synthetic progestagens is a source of steroids releasing into the environment (Liu et al., 2012a; Liu 60 et al., 2012b; Liu et al., 2015). This results in contamination of environment by steroids and their metabolites from animal manure and urine. Thus, our aim is to develop an alternative strategy for 61 62 hormonal pharmacological tools for estrous synchronization in gilts.

Regular boar contact, called the boar effect, is able to stimulate and synchronize puberty attainment 63 in gilts, although its success has been shown to be variable (Hughes et al., 1990). Part of this 64 variability can be linked to differences in sexual maturity, with an effect of age and growth rate of 65 the gilts, breed, season, and physiological stage of the gilts at onset of boar exposure (Hughes et al., 66 1990; Paterson et al., 1991; Kemp et al., 2005; van Wettere et al., 2006; Amaral Filha et al., 2009; 67 Magnabosco et al., 2014). Before puberty, gilts exhibit a "waiting period", defined by a relatively 68 high ovarian follicles development but relatively low gonadotrophin secretion (Camous et al., 1985; 69 Prunier et al., 1993). During this "waiting period", external stimulations, such as boar exposure, 70 could induce and synchronize first ovulation (Hughes et al., 1990). Targeting more effectively the 71 period when gilts are able to respond to boar exposure would enhance the efficacy of the boar 72 73 effect. Thus, the identification of biomarkers of this receptivity period would allow the detection of optimal time for application of the boar effect in pre-pubertal gilts. 74

Steroid hormones are major actors of sexual maturation in mammals such as humans (Mauras et al., 1996; Mauras, 2001; Castellano and Tena-Sempere, 2016), ruminants (Kinder et al., 1995) and pig (Prunier et al., 1993; Kanematsu et al., 2006). Therefore, they could be relevant biomarkers of the receptivity period. An increase of urinary estrone levels has been observed during the "waiting

period" (Camous et al., 1985) and high urinary estrone concentrations could be biomarkers of the 79 receptivity period. However, urine sampling is difficult in group-housed females. On the contrary, 80 saliva can be sampled easily without pain or stress. Several reports have shown that steroid 81 82 hormones could be detected in saliva in human (Manh et al., 2013; Kido et al., 2014; Majewska et al., 2014). In large non-human mammals, salivary concentrations of steroid hormones have never 83 been analyzed, except for cortisol in bovine (Moya et al., 2014; Schwinn et al., 2018), equine 84 85 (Diego et al., 2016; Schwinn et al., 2018), porcine (Devillers et al., 2004; Jama et al., 2016; Casal et al., 2017) and donkey (Rota et al., 2018) and testosterone in porcine (Escribano et al., 2015). 86

The combination of gas chromatography (GC) or liquid chromatography with mass spectrometry 87 88 (MS) or tandem mass spectrometry (MS/MS) has provided reference methods for steroids analysis both in clinical studies and in fundamental research. Thanks to their analytical specificities, these 89 technologies permit to reach extreme sensitivity down to the attomoles and have the advantage to 90 accurately measure several steroids in a biological tissue or fluid of small size, and to establish 91 extended steroid profiles. In particular, GC-MS/MS allows measuring a very large number of 92 93 structurally similar steroids such as stereoisomers and enantiomers (Schumacher et al., 2015; Lamy et al., 2016; Zhu et al., 2017). We developed an analytical protocol, based on GC-MS/MS analysis 94 coupled to a multi-step sample workup for screening and quantification of steroid hormones and 95 96 their precursors and metabolites in small volumes of biological samples (Liere et al., 2004; Liere et al., 2009). 97

98 Thus, the aim of the present study was to identify potential salivary biomarkers of the receptivity 99 period to boar effect through screening and quantification of steroids and their 100 precursors/metabolites in saliva from immature to pubertal gilts using the GC-MS/MS analytical 101 technology.

102

103 Materials and methods

104 *Ethics approval statement*

All procedures on animals were performed in accordance with the "guiding principles for the care
and use of animals in research facilities" from the French Ministry of Agriculture and Research.
The protocol was approved by the ethics committee "Comité d'Ethique en Expérimentation
Animale Val de Loire n° 19".

109 Animals, housing and sample collection

This experiment was conducted from November to December in our experimental farm (PAO, INRA, Nouzilly, France). Nine Large-White 144 to 147-day-old gilts were penned in a group on a concrete floor without straw under natural daylight. They had free access to water and were fed with concentrate once a day (2.4 kg/animal/day; porc Elevage INRA, Sanders Ouest, 35370 Etrelles, France; 16 % crude protein, 2 % crude fat content, 7.1 % cellulose, 6.9 % minerals, 0.62 % phosphore, 1.15 % calcium, 0.25 % sodium, 0.8 % lysine, 0.26 % methionine). No boar contact and no modification of the housing or feeding were performed during the experiment.

Starting at 144-147 days of age, trans-abdominal ultrasonography was carried out three times a week to assess the physiological reproductive status, immature versus pubertal, until puberty detection based on uterus and ovaries development, as previously described (Goudet et al., 2019). Females were considered pubertal when the ultrasonographic images were characteristic of cyclic females with well-defined sections of the uterus and ovulations were detected. Females were considered immature when the uterine images were dark and homogeneous.

Starting at 144-147 days of age, urinary and salivary samples were collected in the morning before feeding (7 o'clock), three times a week, on the day of ultrasonography, until puberty detection. Urine samples were collected using a container placed under the urine stream, when possible. Urine samples were obtained one of the two collection attempts. Saliva samples were collected using a cotton swab for saliva collection (Sarstedt Salivette® ref. 51.1534, Sarstedt, Nümbrecht, Germany) held gently with forceps in the gilt mouth that was chewing until it was soaked. Saliva samples were obtained at each collection attempt, 550 µl to 2.25 ml were collected. Urinary and saliva samples

- 130 were brought to the lab on ice, centrifuged at 3000 g for 5 min at 4°C, within 30 minutes after
- 131 collection. The recovered saliva and urine were aliquoted and stored at -80° C until analysis.
- 132 Gilts were slaughtered 7 days after ultrasonographic puberty detection for puberty confirmation and
- 133 genital tract was carefully examined as previously described (Goudet et al., 2019).

134 Urine samples were used for urinary oestrone assay, as mentioned in a previous experiment (Goudet

- et al., 2019). Briefly, urinary oestrone concentrations were measured using the DetectX-Estrone-3-
- 136 sulfate enzyme immunoassay kit (Arbor Assays) and urinary creatinine concentrations were
- 137 measured using the Creatinine Assay kit (R&D Systems) in order to calculate the oestrone-to-
- 138 creatinine ratio and take into account urine dilution.
- 139 Part of the saliva samples was used for metabolome analysis using ¹H-nuclear magnetic resonance
- spectroscopy in a previous experiment (Goudet et al., 2019). The other part was used in the present
- 141 study for steroidome analysis using GC-MS/MS technology.
- 142 Steroids measurement by GC-MS/MS

Steroids were extracted from the saliva (400-550 µl) of gilts with methanol. The saliva was spiked 143 with all the following internal standards for steroid quantification: 2 ng of ${}^{2}H_{6}$ -5 α -DHP (CDN 144 145 Isotopes, Sainte Foy la Grande, France) for $5\alpha/\beta$ -dihydroprogesterone ($5\alpha/\beta$ -DHP) and $5\alpha/\beta$ dihydroandrostanedione ($5\alpha/\beta$ -ADIONE), 2 ng of ²H₄-pregnenolone sulfate (PREGS) for PREGS, 146 5 ng of ²H₄-dehydroepiandrosterone sulfate (DHEAS) for DHEAS, 2 ng of ¹³C₃-testosterone 147 (Isoscience, PA, USA) for testosterone, 2 ng of ${}^{13}C_3$ -androstenedione (ADIONE) for ADIONE, 2 148 ng of ${}^{13}C_3$ -progesterone (P) for P, 2 ng of ${}^{13}C_3$ -17 β -estradiol for 17 β -estradiol, 2 ng of ${}^{13}C_3$ -estrone 149 for estrone, 2 n of ${}^{2}C_{3}$ -deoxycorticosterone (DOC) for DOC, 2 ng of 19nor-PROG for 20 α -DHP 150 and 5\alpha-DHDOC (dihydrodeoxycorticosterone), 2 ng of epietiocholanolone (Steraloids, Newport, 151 152 Rhode Island) for androstenediol (ADIOL), etiocholanolone, epiandrosterone, 5αdihydrotestosterone (5 α -DHT), 3 α/β 5 α -tetrahydrotestosterone (THT), pregnenolone (PREG), 20 α -153 dihydropregnenolone (20 α -DHPREG) $3\alpha/\beta5\alpha/\beta$ -tetrahydroprogesterone (THP), $5\alpha20\alpha$ -THP, 154 $3\alpha/\beta 5\alpha/\beta$ -hexahydroprogesterone (HHP) and $3\alpha 5\alpha$ -THDOC, 2 ng of ${}^{13}C_3$ -17 α -OH PROG for 17 α -155

OH PROG, 17 α -OH PREG and 16 α -OH PROG, and 1 ng of ²H₅-2-methoxyestradiol (2-ME2) for 156 2-ME2. Samples were purified and fractionated by solid-phase extraction with the recycling 157 procedure (Liere et al., 2004; Liere et al., 2009). Briefly, the extracts were dissolved in 1 ml MeOH 158 and applied to the C18 cartridge (500 mg, 6 ml, International Sorbent Technology, IST), followed 159 by 5 ml of MeOH/H₂O (85/15). The flow-through, containing the free and sulfated steroids, was 160 collected and dried. After a previous re-conditioning of the same cartridge with 5 ml H₂O, the dried 161 samples were dissolved in MeOH/H₂O (2/8) and re-applied. The cartridge was then washed with 5 162 ml H₂O and sulfated and unconjugated steroids were eluted with 5 ml MeOH/H₂O (1/1) and 5 ml 163 MeOH/H₂O (9/1), respectively. 164

The unconjugated steroids-containing fraction was then filtered and further purified and 165 166 fractionated by HPLC. The HPLC system was composed of a WPS-3000SL analytical autosampler and a LPG-3400SD quaternary pump gradient coupled with a SR-3000 fraction collector 167 (Thermoscientific, USA). The HPLC separation was achieved with a Lichrosorb Diol column (25 168 169 cm, 4.6 mm, 5 µm) in a thermostated block at 30°C. The column was equilibrated in a solvent system of 90 % hexane and 10 % of a mixture composed of hexane/isopropanol (85/15). Elution 170 was performed at a flow-rate of 1 ml/min, first with 90 % hexane and 10 % of hexane/isopropanol 171 (85/15) for 8 min, then with a linear gradient to 100 % of hexane/isopropanol (85/15) in 2 min. This 172 mobile phase was kept constant for 10 min and a linear gradient to 100 % MeOH was applied. The 173 174 column was washed with MeOH for 15 min.

Two fractions were collected from the HPLC system: $5\alpha/\beta$ -DHPROG and $5\alpha/\beta$ -ADIONE were eluted in the first HPLC fraction (3-13 min) and were next silvlated with 50 µl of a mixture Nmethyl-N-trimethylsilyltrifluoroacetamide/ammonium iodide/dithioerythritol (1000:2:5 vol/wt/wt) for 15 min at 70°C. The second fraction (13-25 min) containing PREG, 20 α -DHPREG, P, DOC, testosterone and their precursors and reduced metabolites, 17 β -estradiol and estrone, was derivatized with 25 µl heptafluorobutyric anhydride (HFBA) and 25 µl anhydrous acetone for 1h at 20°C. Both fractions were dried under a stream of N₂ and resuspended in hexane.

GC-MS/MS analysis of the extracts was performed using an AI 1310 autosampler, a Trace 1310 gas 182 chromatograph (GC), and a TSQ 8000 tandem mass spectrometer (MS/MS) (Thermo Fisher 183 Scientific San Jose, CA) using Argon as collision gas. Injection was performed in the splitless mode 184 185 at 250°C (1 min of splitless time). The temperature of the gas chromatograph oven was initially maintained at 50°C for 1 min and ramped between 50 to 200°C at 20°C/min, then ramped to 300°C 186 at 5°C/min and finally ramped to 350°C at 30°C/min. The helium carrier gas flow was maintained 187 constant at 1 ml/min during the analysis. The transfer line and ionization chamber temperatures 188 were 330°C and 180°C, respectively. Electron impact ionization was used for mass spectrometry 189 with ionization energy of 70 eV. GC-MS/MS signals were evaluated using a computer workstation 190 by means of the software Excalibur[®], release 3.0 (Thermoscientific, USA). Identification of steroids 191 was supported by their retention time and according two or three transitions. The precise retention 192 time and the MS/MS parameters of the 35 targeted steroids in gilt saliva are reported in Table 1. 193 Quantification was performed according to the transition giving the more abundant product ion with 194 a previously established calibration curve. The calibration curves were prepared by using 100 µl of 195 196 gilt saliva by adding increasing concentrations of pure targeted steroids (5 calibration points with steroids concentrations ranging from 0.100 to 5 ng/ml) with fixed amounts of internal standards, as 197 described in the analytical work up. Then, steroids were extracted, purified, fractionated and 198 derivatized according to the procedure described in the paper. For each calibration curve, a 199 regression line was calculated using least squares methodology. The calibration curve is considered 200 as satisfactory when the concentrations of the standards were within 15 % of the nominal 201 concentration. 202

The analytical protocol has been validated for all the targeted steroids by using 1 ml from a pool of male rat plasma. The parameters of validation included the limit of detection, linearity, accuracy, intra- and inter-assay precisions and were all described by Lamy et al (Lamy et al., 2016). The limit of detection was determined as the lowest amount of compounds that can be measured by GC-MS/MS with a signal-to-noise ratio greater than 3. It ranged from 5 pg/ml to 0.05 pg/ml. The

linearity was assessed by analysing increasing amounts of rat plasma extracts (20, 50, 100 and 200 208 µl) in triplicate. The linearity was satisfactory for all the steroids with a coefficient of correlation 209 ranging from 0.99 to 0.995. Five aliquots of 1 ml of rat plasma were used at two different times to 210 211 determine the inter-assay coefficient of variation that was estimated at roughly 5-10 % for the targeted steroids. The accuracy of the assay was evaluated by determining the analytical recovery, 212 which was defined as $C/(C_0+S) \times 100(\%)$. C is the concentration of the steroid in the spiked plasma 213 extract (100 μ l), C₀ is the concentration of a steroid in the unspiked plasma extract (100 μ l) and S is 214 215 the spiked concentration. The accuracy was in the range of 94-106 %.

216 Stable isotope dilution analysis combined with extensive sample purification/fractionation and

217 derivatization, sample dilution and the highly specific GC/MS/MS analysis (high resolution gas

218 chromatography coupled to multiple reaction monitoring) are satisfactory to avoid any potential

219 matrix effects with gilt saliva for steroid analysis (Anastassiades et al., 2003).

220 Statistical analysis

The comparison of concentrations of steroids determined by GC-MS/MS between weeks was performed using repeated measures one-way ANOVA, followed by a post-hoc Tukey's multiple comparisons test, using GraphPad Prism version 6 (GraphPad Software, La Jolla California, USA).

224

225 **Results**

226 Uterine tract development and urinary estrone levels

As mentioned in our previous experiment (Goudet et al., 2019), six gilts presented first pubertal ultrasonic images of the uterus and ovary at 182, 189, 190, 190, 191 and 192 days of age (day of puberty detection) and puberty was ascertained at slaughter. Two other gilts were still pre-pubertal at slaughter, and first ovulation was missed for one gilt. Only samples from the six gilts whose onset of puberty was detected at 182 to 192 days were analyzed.

The period from 144-147 days of age to the day of puberty detection (Day 0) was divided into 5 weeks named week-5 to week-1 relatively to the day of puberty detection. For each week, one of the three collected samples was chosen for analyses, so that both urinary and saliva were available on the same day. Then, for each gilt, six salivary samples were analyzed: one from each week, from week-5 to week-1, and one from the day of puberty detection. The mean age of the gilts (\pm sem) on the day of saliva sampling was 154 days (\pm 0.9) for week-5, 163 days (\pm 0.5) for week-4, 170 days (\pm 0.5) for week-3, 178 days (\pm 0.5) for week-2, 186 days (\pm 1.4) for week-1, and 189 days (\pm 1.5) for the day of puberty detection.

As mentioned in our previous experiment (Goudet et al., 2019), mean urinary estrone concentrations \pm sem were 12.9 \pm 2.5 ng/ml (week-5), 10.4 \pm 2.7 ng/ml (week-4), 13.4 \pm 4.1 ng/ml (week-3), 12.7 \pm 2.3 ng/ml (week-2), 62.5 \pm 31.1 ng/ml (week-1) and 121.3 \pm 63.1 ng/ml on the day of puberty detection (Day 0). The estrone-to-creatinine ratio was low from week-5 to week-2 and increased significantly (P < 0.05) from week-2 to the day of puberty detection. The period with increasing estrone levels was considered as the "waiting period".

246 Salivary steroidome analysis

Steroids concentrations were analyzed in six salivary samples from the six gilts, from week-5 to the day of puberty detection. Among the 35 targeted steroids, GC-MS/MS analysis allowed the detection of 28 steroids in 500 μ l of saliva (Table 2). Seven targeted steroids were not detected in saliva : 17 α -OH PREG, 16 α -OH PROG, 17 α -OH PROG, 5 α -DHDOC, 3 α 5 α -THDOC, 3 β 5 α -THT, 2-ME2 (Table 2).

The salivary concentration of pregnenolone, the precursor of all steroid hormones in mammals, tended to slightly increase from 0.9 ± 0.16 ng/ml (week-5 and week-4) to 1.47 ± 0.15 ng/ml (week-1) but without reaching statistical significance (Figure 1). Both direct metabolites of pregnenolone, pregnenolone sulfate (PREGS) and 20 α -dihydropregnenolone (20 α -DHPREG), were also detected in gilts saliva. PREGS concentrations showed important but non-significant variations ranging from 0.67 ± 0.19 to 2.08 ± 0.53 ng/ml, whereas 20 α -DHPREG concentrations were lower and showed little variations around 0.1 ng/ml (Figure 1). Dehydroepiandrosterone (DHEA) is also a metabolite of pregnenolone. Its concentrations showed a significant decrease from week-3 ($0.26 \pm 0.02 \text{ ng/ml}$) to week-2 ($0.11 \pm 0.01 \text{ ng/ml}$, p < 0.01), to week-1 ($0.12 \pm 0.01 \text{ ng/ml}$, p < 0.01) and to the day of puberty detection ($0.09 \pm 0.01 \text{ ng/ml}$, p < 0.01) (Figure 1). DHEA sulfate concentrations tended to decrease from week-5 ($0.58 \pm 0.15 \text{ ng/ml}$) to week-4 ($0.22 \pm 0.03 \text{ ng/ml}$) and then tended to increase progressively to the day of puberty detection ($0.70 \pm 0.22 \text{ ng/ml}$) (Figure 1).

Concentrations of deoxycorticosterone, progesterone and some of its reduced metabolites $(3\alpha 5\alpha$ -265 THPROG, 3β5α-THPROG, $5\alpha 20\alpha$ -THPROG, $3\alpha 5\alpha 20\alpha$ -HHPROG, $3\alpha 5\beta 20\alpha$ -HHPROG, 266 267 $3\beta 5\alpha 20\alpha$ -HHPROG) were lower than 0.2 ng/ml in saliva, whereas concentrations of the other progesterone metabolites (5α-DHPROG, 5β-DHPROG, 20α-DHPROG, 3α5β-THPROG) ranged 268 between 0.1 and 3.9 ng/ml (Figures 2 and 3). The concentrations of progesterone and all of its 269 reduced metabolites (dihydro-, tetrahydro- and hexahydroprogesterone) tended to decrease between 270 week-5 and week-4. Then progesterone levels tended to increase between week-3 and the day of 271 puberty detection and most of its metabolites showed similar pattern (5 α -DHPROG, 5 β -DHPROG, 272 3a5β-THPROG, 3β5a-THPROG, 5a20a-THPROG, 3a5a20a-HHPROG, 3a5β20a-HHPROG, 273 3β5α20α-HHPROG). Concentrations of 5α-DHPROG were significantly different between week-3 274 275 $(0.32 \pm 0.05 \text{ ng/ml})$ and week-2 $(0.66 \pm 0.11 \text{ ng/ml}, \text{p} < 0.05)$ as well as between week-4 (0.32 ± 0.05) 0.06 ng/ml) and week-1 (1.06 \pm 0.16 ng/ml) (Figure 2). Concentrations of 3 β 5 α -THPROG were 276 significantly different between week-1 vs week-4 (Figure 3, p < 0.05). Concentrations of $5\alpha 20\alpha$ -277 THPROG were significantly different between week-2 vs week-4 (Figure 3, p < 0.05). 278

The concentrations of androgens androstenediol, androstenedione, 5α -ADIONE, 5β -ADIONE, epiandrosterone and etiocholanolone ranged from 0.01 to 0.6 ng/ml. Except for androstenediol, all of these androgens showed a similar pattern. They tended to decrease from week-5 to week-4 and increase gradually to the day of puberty detection (Figure 4). Androstenediol tended to increase from week-5 to week-1 and decrease on the day of puberty detection (Figure 4).

11

Testosterone levels were low and did not change significantly with time as for its reduced metabolites 5α -DHT and $3\alpha5\alpha$ -THT (Figure 5).

Estrone salivary levels significantly decreased from week-5 ($0.079 \pm 0.012 \text{ ng/ml}$) to week-4 ($0.015 \pm 0.007 \text{ ng/ml}$, p < 0.01) and increased progressively to the day of puberty detection (0.063 ± 0.021 ng/ml) (Figure 5). The concentrations of the most potent estrogen 17β -estradiol were low from week-5 ($0.035 \pm 0.006 \text{ ng/ml}$) to week-3 ($0.011 \pm 0.001 \text{ ng/ml}$), significantly higher on week-2 ($0.047 \pm 0.005 \text{ ng/ml}$, p < 0.01) and week-1 ($0.055 \pm 0.007 \text{ ng/ml}$, p < 0.01) and decreased on the day of puberty detection ($0.026 \pm 0.006 \text{ ng/ml}$) (Figure 5).

292 When combining progesterone and its $5\alpha/\beta$ -reduced metabolites (5α -DHPROG + 5β -DHPROG + 20α -DHPROG + 3α , 5α -THPROG + 3α , 5β -THPROG + 3β , 5α -THPROG + 5α , 20α -THPROG + 293 $3\alpha, 5\alpha, 20\alpha$ -HHPROG + $3\alpha, 5\beta, 20\alpha$ -HHPROG + $3\beta, 5\alpha, 20\alpha$ -HHPROG), concentrations tended to 294 decrease between week-5 (3.48 \pm 0.92 ng/ml) and week-4 (0.89 \pm 0.17 ng/ml), remained low on 295 week-3 (1.58 \pm 0.33 ng/ml), and increased progressively from week-2 (3.47 \pm 0.98 ng/ml) to the 296 day of puberty detection (8.44 \pm 2.9 ng/ml) (Figure 6). A similar pattern was observed when the 5 α -297 reduced metabolites of progesterone were pooled (5 α -DHPROG + 3 α ,5 α -THPROG + 3 β ,5 α -298 THPROG + 5α , 20α -THPROG + 3α , 5α , 20α -HHPROG + 3β , 5α , 20α -HHPROG) with a significant 299 increase from week-3 ($0.42 \pm 0.07 \text{ ng/ml}$) to week-2 ($0.81 \pm 0.12 \text{ ng/ml}$, p < 0.05), to week-1 (1.26) 300 \pm 0.19 ng/ml) and to the day of puberty detection (1.95 \pm 0.67 ng/ml). The salivary concentrations 301 of 5 β -metabolites of progesterone (5 β -DHPROG + 3 α ,5 β -THPROG + 3 α ,5 β ,20 α -HHPROG) also 302 showed the same tendency with increasing levels from week-3 ($0.45 \pm 0.14 \text{ ng/ml}$) to the day of 303 304 puberty detection $(5.22 \pm 1.8 \text{ ng/ml})$ (Figure 6).

305

306 Discussion

307 Studies about endocrine events during sexual maturation often include analyses of changes in the 308 concentrations of steroid hormones in plasma (Prunier et al., 1993; Kanematsu et al., 2006; Noguchi 309 et al., 2010). However, blood sampling is invasive, stressful and difficult in the pig. On the

contrary, saliva sampling is an interesting alternative since it is easy to collect, without pain or 310 stress. Steroid hormones enter saliva by different mechanisms depending on whether they are 311 charged or neutral, free or bound to proteins. Furthermore, the presence of some steroidogenic 312 313 enzymes in saliva may complicate the relationship between the salivary and plasma steroids. Thus, salivary steroid levels do not always reflect plasma levels (Lewis, 2006). However, some reports 314 have shown that the determination of progesterone, 17α -hydroxyprogesterone and 17β -estradiol in 315 saliva may be useful for assessing the ovarian function and in particular, salivary samples have been 316 shown to allow the differentiation between the follicular and luteal phase in women (Groschl et al., 317 2001; Bao et al., 2003). Saliva sampling might have some disadvantages such as lower hormones 318 concentrations compared to blood or urine, but this inconvenience can be solved by highly sensitive 319 320 detection techniques. Gas chromatography coupled to tandem mass spectrometry (GC-MS/MS) associated with a well-validated extraction, fractionation and purification procedure is a sensitive, 321 precise and selective technique suitable for quantifying numerous steroid hormones and their 322 323 precursors and metabolites in small volumes of biological fluids with high sensitivity and accuracy (Liere et al., 2004; Liere et al., 2009). The analytical protocol used in this study was validated by 324 using mice brain extracts which represent the more challenging complex matrix in an analytical 325 point of view (Liere and Schumacher, 2015; Schumacher et al., 2015). This protocol included the 326 addition of stable isotope internal standards in the extraction step in the beginning of the analytical 327 procedure combined with subsequent multiple purification and fractionation steps such as Solid 328 Phase Extraction and High Performance Liquid Chromatography in order to have a very reliable 329 protocol in terms of steroids identification and quantification whatever the studied biological 330 matrix: brain, plasma (Zhu et al., 2017), oviductal fluid (Lamy et al., 2016), saliva. Many groups 331 having a great expertise in the analytical field have reported that the matrix effect could be bypassed 332 by using highly specific GC-MS/MS stable isotope dilution analysis combined with upstream 333 extensive sample purification/fractionation, derivatization and sample dilution (Schenck and 334 Lehotay, 2000; Anastassiades et al., 2003; Mastovska et al., 2005; Chamkasem and Harmon, 2016) 335

indicating that our GC-MS/MS protocol was suitable and satisfactory to avoid any potential matrix
effects with gilt saliva for steroid analysis (Anastassiades et al., 2003).

Thus, in the present study, we were able to characterize the steroid profile in gilt saliva from immature to pubertal stage by using GC-MS/MS analysis. This technique allowed us the detection of 28 steroids in 500 µl of saliva.

341 Conversion of cholesterol to pregnenolone is the first step in the synthesis of all steroid hormones in gonads and adrenal cortex (Payne and Hales, 2004; Robic et al., 2014). Pregnenolone can then be 342 converted to 17α -OHPREG and 20α -DHPREG or sulfated to pregnenolone sulfate (Payne and 343 Hales, 2004; Schuler et al., 2014). In our study, pregnenolone and pregnenolone sulfate levels 344 ranged between 0.7 and 2 ng/ml, 20\alpha-DHPREG levels were lower than 0.11 ng/ml and 17\alpha-345 OHPREG was undetectable in gilts saliva. In human, similar levels were observed in pre-pubertal 346 girls (Majewska et al., 2014). To our knowledge, the salivary concentrations of these steroid 347 348 hormones had never been analyzed in large domestic mammals. Plasma concentrations of pregnenolone, pregnenolone sulfate, 20α -DHPREG and 17α -OHPREG were analyzed in cyclic 349 gilts (Stone and Seamark, 1985), cyclic ewes (McKay et al., 1987) and women (Deng et al., 2017). 350 Pregnenolone levels in gilts saliva were similar to plasma levels in cyclic gilts or ewes, but 351 pregnenolone sulfate, 20α -DHPREG and 17α -OHPREG levels in gilts saliva were lower than 352 plasma levels in gilts, ewes and women. In our study, pregnenolone, pregnenolone sulfate and 20α -353 DHPREG concentrations did not significantly vary between weeks. They were not influenced by 354 the entrance into the "waiting period", evidenced by increasing urinary estrone levels. Thus, these 355 precursors of steroid hormones may not be relevant biomarkers of the period of receptivity to the 356 boar effect. 357

DHEA is synthesized from 17α-OHPREG in gonads and adrenal cortex, and sulfated to DHEAS in
adrenals (Payne and Hales, 2004; Robic et al., 2014). In gilts saliva, DHEA levels were around 0.3
ng/ml from week-5 to week-3 and significantly dropped to 0.1 ng/ml from week-2 to the day of
puberty detection. DHEAS levels did not significantly vary between weeks. Salivary concentrations

of DHEA and DHEAS had never been analyzed in large domestic mammals. In human saliva from 362 pre-pubertal girls, DHEA levels were similar to levels from young gilts, but DHEAS levels were 363 higher (Majewska et al., 2014). Plasma from cyclic gilts contains 0.03 ng/ml of DHEA, suggesting 364 365 that DHEA levels continue to decrease after puberty (Stone and Seamark, 1985). In our study, DHEA levels in gilts saliva significantly dropped between week-3 and week-2, around 175 days 366 old. The drop of DHEA levels is concomitant with the increase of urinary oestrone concentration 367 368 and could correspond to the beginning of the "waiting period" during which gilts are receptive to the boar effect. Thus, DHEA may be a potential biomarker of the period of receptivity to the boar 369 effect. Further experiments comparing gilts that respond or not to the boar effect are in progress to 370 371 confirm that DHEA is a relevant biomarker.

Progesterone is synthesized from pregnenolone in gonads and adrenal cortex and is ubiquitously 372 converted to its reduced metabolites (dihydroprogesterone, tetrahydroprogesterone, 373 hexahydroprogesterone) (Payne and Hales, 2004; Meffre et al., 2007; Zhu et al., 2017). The use of 374 GC-MS/MS as a tool to analyze multiple steroids with very closed chemical structures with high 375 specificity and sensitivity provides opportunities to explore for the first time the presence and 376 concentration of progesterone metabolites in saliva of large domestic animals. In our study, 377 progesterone levels from gilts saliva ranged between 0.02 ng/ml and 0.2 ng/ml. Saliva progesterone 378 379 was analyzed in some domestic and wild animals with concentrations ranging from 0.05 to 0.1 ng/ml in cyclic cows (Gao et al., 1988), from 2 to 8.5 ng/ml in cyclic buffaloes (Lasheen et al., 380 2018), and 2.4 ng/ml in adult female alpacas (Volkery et al., 2012). No data are available in saliva 381 from non-pubertal females. Blood progesterone concentrations in pre-pubertal gilts were similar to 382 saliva levels (Kolesarova et al., 2010; Zhuo et al., 2014). Thus, salivary progesterone levels could 383 384 reflect the circulating level, and salivary progesterone could be a relevant biomarker for assessing ovarian function, as shown in human (Lewis, 2006). In our study, concentrations of the 385 progesterone metabolites ranged between 0.001 and 4 ng/ml. In human saliva of pre-pubertal girls, 386 $3\alpha 5\alpha$ -THPROG levels were similar to levels in gilts saliva (Majewska et al., 2014). Recent studies 387

have analyzed the concentration of progesterone metabolites in oviductal fluid of cows (Lamy et al., 388 2016), in plasma of mare (Legacki et al., 2016), in brain and plasma rats and mice (Meffre et al., 389 2007; Zhu et al., 2017). However, to our knowledge, no data are available on the saliva of domestic 390 391 mammals. In our study, progesterone levels from gilts saliva tended to increase between week-3 and week-2, close to the beginning of the "waiting period", suggesting a relationship between salivary 392 393 progesterone levels and ovarian activity. However, this increase was not significant. Interestingly, most of the progesterone metabolites showed a similar increase from week-3 to the day of puberty 394 detection (5 α -DHPROG, 5 β -DHPROG, 3 α 5 β -THPROG, 3 β 5 α -THPROG, 5 α 20 α -THPROG, 395 3\alpha5\alpha20\alpha-HHPROG, 3\alpha5\beta20\alpha-HHPROG, 3\beta5\alpha20\alpha-HHPROG) meaning that they are reliable 396 markers of the ovarian progesterone synthesis. Moreover, 5α -DHPROG concentrations 397 significantly increased from week-3 to week-2, at the beginning of the "waiting period", and 398 remained high until the day of puberty detection. When adding together the concentrations of all 399 5α -metabolites of progesterone and all 5 β -metabolites of progesterone, we showed an increase of 400 the concentrations between week-3 and week-2, and this increase was significant for the 5α -401 metabolites of progesterone. These metabolites analyzed all together represent an interesting target 402 for investigating the ovarian function. Thus, the 5α -metabolites of progesterone and 5α -DHPROG 403 could be potential biomarkers of the "waiting period" during which gilts are receptive to the boar 404 effect. Further experiments are in progress to confirm that this biomarker could be relevant. 405

Androstenediol is synthesized from DHEA while androstenedione is synthesized from both 17α-406 OHPROG and DHEA in gonads and adrenal cortex (Payne and Hales, 2004; Robic et al., 2014; 407 Kovac et al., 2016). Androstenedione is then reduced to 5α -ADIONE which is converted to 408 epiandrosterone, and 5β-ADIONE which is converted to etiocholanolone (Anizan et al., 2011; Luu-409 The, 2013; Robic et al., 2014). Their concentrations in gilts saliva ranged from 0.01 to 0.63 ng/ml. 410 To our knowledge, the salivary concentration of these steroid hormones had never been analyzed in 411 domestic mammals. In human saliva from pre-pubertal girls, similar levels were observed for 412 413 androstenediol, androstenedione, etiocholanolone and epiandrosterone (Majewska et al., 2014).

Plasma from cyclic gilts contained 0.15 ng/ml androstenedione (Stone and Seamark, 1985), whereas salivary levels on the day of puberty detection reached 0.6 ng/ml, suggesting a decrease after puberty. Finally, our data showed a progressive increase from week-4 to puberty for androstenedione and its metabolites. Androstenedione is a precursor of 17β -estradiol in ovaries and its increase could also reflect the beginning of the ovarian maturation. But the sensitivity as a biomarker appears to be too low so that androstenedione may not be a relevant biomarker of the period of receptivity to the boar effect.

Testosterone is synthesized from androstenedione and androstenediol in testis (Robic et al., 2014; 421 Kovac et al., 2016). It is then reduced to dihydrotestosterone which can be converted to 422 tetrahydrotestosterone (Meffre et al., 2007; Robic et al., 2014; Kovac et al., 2016). In our study, 423 testosterone levels in gilts saliva were low. Testosterone levels were evaluated in the saliva of 424 young gilts (< 150 days old) (Escribano et al., 2014), heifers (10-11 months old) (Geburt et al., 425 2015), and young girls (< 1 year old) (Fang et al., 2017). A correlation was shown between saliva 426 and blood concentration of testosterone in young girls (Fang et al., 2017). However, no data were 427 available on the evolution of testosterone levels during pre-pubertal development of female 428 mammals, whether in saliva or in blood. Since no significant changes of testosterone, 5α -DHT and 429 $3\alpha 5\alpha$ -THT levels were observed around the beginning of the "waiting period", these steroid 430 hormones were not considered to be relevant biomarkers of the period of receptivity to the boar 431 effect. 432

Estrone is synthesized mainly from androstenedione and converted to estradiol in ovaries (Payne and Hales, 2004; Robic et al., 2014). In gilts saliva, estrone and 17β -estradiol levels ranged between 0.01 and 0.08 ng/ml, with a decrease from week-5 to week-4 and a progressive increase to week-1. In human saliva from young girls, estradiol and estrone concentrations reached similar levels (Fang et al., 2017). To our knowledge, the salivary concentration of these steroid hormones has never been analyzed in domestic mammals, except in saliva from buffalo females during estrus and diestrus (Ravinder et al., 2016). Serum 17 β -estradiol concentrations in gilts (0.02 ng/ml before

puberty and 0.03 ng/ml at puberty) (Cosgrove et al., 1993; Li et al., 2016) were close to salivary 440 levels measured in our study. Moreover, a correlation has been shown between saliva and serum 441 concentrations of estradiol and estrone in adult women and young girls (Dielen et al., 2017; Fang et 442 443 al., 2017). Finally, the increase of estradiol levels before puberty was also observed in serum from young girls (Sehested et al., 2000) and plasma from heifers (Melvin et al., 1999). Thus, salivary 444 17β-estradiol might be considered as a surrogate marker for blood estradiol. In our study, 17β-445 estradiol levels in gilts saliva significantly increased between week-3 and week-2, around 175 days 446 old. Since this period was considered as the beginning of the "waiting period", the increase in 447 estradiol levels could be related to the "waiting period" during which gilts are receptive to the boar 448 effect. Thus, 17β-estradiol may be a potential biomarker of the period of receptivity to the boar 449 effect. 450

Salivary biomarkers of the period of gilts receptivity to the boar effect, such as steroids whose 451 concentration significantly vary at the beginning of this period, would be of great help to improve 452 detection of the gilts to stimulate. Overall, our study showed that the beginning of the "waiting 453 period", during which gilts are sensitive to boar exposure, is concomitant with a significant decrease 454 of salivary DHEA levels, a significant increase of 5α -DHPROG concentrations, and a significant 455 increase of 17β -estradiol levels. This increase of estradiol levels could coincide with the appearance 456 457 of antral follicles in the porcine ovary, with increasing estrogen production (Camous et al., 1985). In prepubertal heifers, increasing circulating concentrations of 17^β-estradiol were associated with 458 development of ovarian follicles (Bergfeld et al., 1994). Since DHEA is the precursor of androgens 459 such as androstenedione, which is the main precursor of estrone and estradiol, as mentioned above, 460 the decrease of DHEA levels may be linked to an increase of its consumption for the synthesis of 461 estradiol. The increase of the concentration of progesterone and its metabolites also suggests a 462 relationship with the increase of ovarian activity, since luteinization of follicles and increase of 463 circulating progesterone levels of ovarian origin has been observed during the peri-pubertal period 464 in heifers and ewes (Kinder et al., 1995). Thus, the combined determination of DHEA, 17β -465

466	estradiol, 5α -dihydroprogesterone and/or 5α -reduced metabolites of progesterone in gilts saliva
467	appears to give a precise "picture" of the ovarian function and thus could help to identify the
468	"waiting period" during which gilts are receptive to boar effect. However, their low concentrations
469	in saliva require expensive and sensitive analysis and may limit their use in pig farms. Moreover,
470	whether these potential steroidal biomarkers are relevant to identify the period of gilts receptivity to
471	boar effect has to be checked. For this purpose, experiments comparing gilts that respond or not to
472	boar exposure are in progress. They will help to confirm relevant biomarkers of the period of gilts
473	receptivity to boar effect.
474	
475	Conclusion
476	These results show that painless sampling of saliva could be a non-invasive welfare-friendly tool for
477	the identification of the physiological hormonal status of the gilts. DHEA, 17β -estradiol, 5α -
478	dihydroprogesterone and 5α -reduced metabolites of progesterone could be potential salivary
479	biomarkers of the receptivity period to boar effect. They could allow the identification of the
480	optimal time for application of the boar effect. Further studies are in progress to ascertain their
481	relevance. These studies participate in the development of non-hormonal alternative breeding tools
482	for optimal management of gilt reproduction.
483	
484	List of non-standard abbreviations
485	GC: gas chromatography
486	MS: mass spectrometry
487	MS/MS: tandem mass spectrometry
488	PREG : Pregnenolone
489	PREGS : Pregnenolone sulfate
490	20α-DHPREG : 20α-dihydropregnenolone
491	17α-OH PREG : 17α-hydroxypregnenolone

19

- 492 PROG : Progesterone
- 5α -DHPROG : 5α -dihydroprogesterone
- 5β -DHPROG : 5β -dihydroprogesterone
- 20α -DHPROG : 20α -dihydroprogesterone
- $3\alpha, 5\alpha$ -THPROG : $3\alpha, 5\alpha$ -tetrahydroprogesterone (allopregnanolone)
- $3\alpha,5\beta$ -THPROG : $3\alpha,5\beta$ -tetrahydroprogesterone (pregnanolone)
- 3β , 5α -THPROG : 3β , 5α -tetrahydroprogesterone (iso-allopregnanolone)
- 5α , 20 α -THPROG : 5α , 20 α -tetrahydroprogesterone
- $3\alpha, 5\alpha, 20\alpha$ -HHPROG : 5α -pregnane- $3\alpha, 20\alpha$ -diol ($3\alpha, 5\alpha, 20\alpha$ -hexahydroprogesterone)
- $3\alpha,5\beta,20\alpha$ -HHPROG : 5β -pregnane- $3\alpha,20\alpha$ -diol ($3\alpha,5\beta,20\alpha$ -hexahydroprogesterone)
- 3β , 5α , 20α -HHPROG : 5α -pregnane- 3β , 20α -diol (3β , 5α , 20α -hexahydroprogesterone)
- 16α -OH PROG : 16α -hydroxyprogesterone
- 17α -OH PROG : 17α -hydroxyprogesterone
- 505 DOC : Deoxycorticosterone
- 5α -DHDOC : 5α -dihydrodeoxycorticosterone
- 3α , 5α -THDOC : 3α 5 α -tetrahydrodeoxycorticosterone (allotetrahydrodeoxycorticosterone)
- 508 DHEA : Dehydroepiandrosterone
- 509 DHEAS : Dehydroepiandrosterone sulfate
- 510 ADIOL : Androstenediol (Δ 5-androstene 3 β ,17 β -diol)
- 511 ADIONE : Androstenedione (Δ 4-androstene 3,17-dione)
- 5α -ADIONE : 5α -androstane-3,17-dione
- 5β -ADIONE : 5β -androstane-3,17-dione
- 514 Epiandrosterone : 5α -androstane- 3β -ol-17-one
- 515 Etiocholanolone : 5β -androstane- 3α -ol-17-one
- 516 T : Testosterone

517	5α -DHT : 5α -dihydrotestosterone
518	$3\alpha, 5\alpha$ -THT : $3\alpha, 5\alpha$ -tetrahydrotestosterone
519	3β , 5α -THT : 3β , 5α -tetrahydrotestosterone
520	E1 : Estrone
521	17β -E2 : 17β -estradiol
522	2-ME2 : 2-methoxyestradiol
523	
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526	
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528	
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530	
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533	
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709

710 **Figure captions**

Figure 1. GC-MS/MS measurements of pregnenolone, pregnenolone sulfate, 20α -

712 dihydropregnenolone, dehydroepiandrosterone and dehydroepiandrosterone sulfate concentrations

- (ng/ml + sem) in gilts saliva from week-5 to the day of puberty detection. For each steroid the y-
- axis was standardized to the maximal concentration, so that the variations are easy-to-read.
- 715 Statistical differences between weeks were analyzed with one-way repeated measures ANOVA
- followed by Tukey post-tests. * significantly different from week-5, # from week-4, \$ from week-3.

717 * # \$: P < 0.05, \$\$ P < 0.01.

- 718
- Figure 2. GC-MS/MS measurements of progesterone, 5α -dihydroprogesterone, 5β -
- dihydroprogesterone, 20α -dihydroprogesterone, $3\alpha5\alpha$ -tetrahydroprogesterone and $3\alpha5\beta$ -

tetrahydroprogesterone concentrations $(ng/ml \pm sem)$ in gilts saliva from week-5 to the day of

puberty detection. For each steroid the y-axis was standardized to the maximal concentration, so

that the variations are easy-to-read. Statistical differences between weeks were analyzed with one-

724 way repeated measures ANOVA followed by Tukey post-tests. # significantly different from week-

- 725 4, \$ from week-3. # \$: P < 0.05.
- 726
- Figure 3. GC-MS/MS measurements of $3\beta 5\alpha$ -tetrahydroprogesterone, $5\alpha 20\alpha$ -
- tetrahydroprogesterone, $3\alpha 5\alpha 20\alpha$ -hexahydroprogesterone, $3\alpha 5\beta 20\alpha$ -hexahydroprogesterone,
- 729 $3\beta 5\alpha 20\alpha$ -hexahydroprogesterone and deoxycorticosterone concentrations (ng/ml + sem) in gilts
- rais saliva from week-5 to the day of puberty detection. For each steroid the y-axis was standardized to
- the maximal concentration, so that the variations are easy-to-read. Statistical differences between

weeks were analyzed with one-way repeated measures ANOVA followed by Tukey post-tests. #

significantly different from week-4, P < 0.05.

734

735	Figure 4. GC-MS/MS measurements of androstenediol, and rost enedione, 5α -and rost ane-3, 17-
736	dione, 5 β -androstane-3,17-dione, 5 α -androstane-3 β -ol-17-one and 5 β -androstane-3 α -ol-17-one
737	concentrations (ng/ml \pm sem) in gilts saliva from week-5 to the day of puberty detection. For each
738	steroid the y-axis was standardized to the maximal concentration, so that the variations are easy-to-
739	read. Statistical differences between weeks were analyzed with one-way repeated measures
740	ANOVA followed by Tukey post-tests.
741	
742	Figure 5. GC-MS/MS measurements of testosterone, 5α -dihydrotestosterone, $3\alpha5\alpha$ -
743	tetrahydrotestosterone, estrone and 17 β -estradiol concentrations (ng/ml ± sem) in gilts saliva from
744	week-5 to the day of puberty detection. For each steroid the y-axis was standardized to the maximal
745	concentration, so that the variations are easy-to-read. Statistical differences between weeks were
746	analyzed with one-way repeated measures ANOVA followed by Tukey post-tests. * significantly
747	different from week-5, # from week-4, \$ from week-3. ** ## $\$ P < 0.01$.
748	
749	Figure 6. GC-MS/MS measurements of progesterone and its main reduced metabolites (5 α -
750	DHPROG + 5 β -DHPROG + 3 α ,5 α -THPROG + 3 α ,5 β -THPROG + 3 β ,5 α -THPROG + 5 α ,20 α -
751	THPROG + 3α , 5α , 20α -HHPROG + 3α , 5β , 20α -HHPROG + 3β , 5α , 20α -HHPROG), 5α -
752	metabolites (5 α -DHPROG + 3 α ,5 α -THPROG + 3 β ,5 α -THPROG + 5 α ,20 α -THPROG +
753	3α , 5α , 20α -HHPROG + 3β , 5α , 20α -HHPROG) and 5β -metabolites (5β -DHPROG + 3α , 5β -
754	THPROG + 3α , 5β , 20α -HHPROG) concentrations (ng/ml \pm sem) in gilts saliva from week-5 to the
755	day of puberty detection. The y-axis was standardized to the maximal concentration, so that the
756	variations are easy-to-read. Statistical differences between weeks were analyzed with one-way

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757	repeated measures ANOVA followed by Tukey post-tests. * significantly different from week-5, #
758	from week-4, \$ from week-3. * # \$: P < 0.05.
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764	Tables
765	

Table 1 : GC-MS/MS parameters used for steroid identification and quantification

Steroids (Molecular weight)	Derivatized steroids (molecular weight)	Retention time (min.)	Precursor ions (<i>m/z</i>)	Product ions (m/z)	Collision energy (eV)
Steroids (Molecular weight)Derivatized steroids (molecular weight)Retention time (mi.)Precursor ions (m/z)Product ions (m/z)Collis ener (eX)HPLC fraction # 1 5α -DHPROG (316) 5α -DHPROG-3,20-TMS2 (460)16.3746044510 5β -DHPROG (316) 5β -DHPROG-3,20-TMS2 (460)15.0344535512 5α -ADIONE (288) 5α -ADIONE-3,17-dione (432)18.2343227518 5β -ADIONE (288) 5β -ADIONE-3,17-dione (432)16.854324178PREG (316)PREG-3-HFB (512)23.312982838 20α -DHPREG (318) 20α -DHPREG-3,20-HFB2 (710)19.8549621110 17α -OH PREG (32) 17α -OH PREG-3,20-HFB2 (510)23.4251014710 17α -OH PREG (330) 17α -OH PROG-HFB (526)21.9646536914 16α -OH PROG (330) 16α -OH PROG-3,20-HFB (526)21.4972249310 20α -DHPROG (318) $3\alpha5\alpha$ -THPROG-3,20-HFB (526)21.4972249310 20α -DHPROG (318) $3\alpha5\beta$ -THPROG-3,20-HFB (526)21.4972249310 20α -DHPROG (318) $3\alpha5\beta$ -THPROG-3,40-HFB (514)22.2249648111 $3\alpha5g20\alpha$ -HHPROG (318) $3\alpha5g20\alpha$ -HHPROG-3,40-HFB (514)23.734964678 $3\alpha5g20\alpha$ -HHPROG (318) $3\alpha5g20\alpha$ -HHPROG-3,20-HFB (712)19.4171242910 $365\alpha20\alpha$ -HHPROG (320) $3\alpha5g20\alpha$ -HHPROG-				(0.1)	
5α-DHPROG (316)	5α-DHPROG-3,20-TMS ₂ (460)	16.37	460	445	10
5β-DHPROG (316)	5β-DHPROG-3,20-TMS ₂ (460)	15.03	445	355	12
5α-ADIONE (288)	5α-ADIONE-3,17-dione (432)	18.23	432	275	18
5β-ADIONE (288)	5β-ADIONE-3,17-dione (432)	16.85	432	417	8
	HPLC fraction	# 2			
PREG (316)	PREG-3-HFB (512)	23.31	298	283	8
20α-DHPREG (318)	20α -DHPREG-3,20-HFB ₂ (710)	19.85	496	211	10
17α-OH PREG (332)	17α -OH PREG-HFB (528)	21.88	467	253	8
PROG (314)	PROG-3-HFB (510)	23.42	510	147	10
17α-OH PROG (330)	17α-OH PROG-HFB (526)	21.96	465	369	14
16α-OH PROG (330)	16α-OH PROG-HFB (526)	21.49	722	493	10
20α-DHPROG (316)	20α-DHPROG-3,20-HFB ₂ (708)	20.38	708	693	12
3α5α-THPROG (318)	3α5α-THPROG-3-HFB (514)	22.22	496	481	10
3α5βTHPROG (318)	3α5β-THPROG-3-HFB (514)	22.40	496	481	12
3β5α-THPROG (318)	3β5α-THPROG-3-HFB (514)	23.73	496	467	8
5α20α-THPROG (318)	5α20α-THPROG-3-HFB (514)	25.06	514	231	10
3a5a20a-HHPROG (320)	3α5α20α-HHPROG-3,20-HFB ₂ (712)	19.19	712	429	10
3α5β20α-HHPROG (320)	$3\alpha 5\beta 20\alpha$ -HHPROG-3,20-HFB ₂ (712)	19.41	712	429	10
3β5α20α-HHPROG (320)	3β5α20α-HHPROG-3,20-HFB ₂ (712)	20.66	712	429	10
11-deoxycorticosterone (330)	11-deoxycorticosterone-3,21-HFB ₂ (722)	23.91	722	707	12
5α-DHDOC (332)	5α-DHDOC-HFB (528)	28.34	301	255	6

3α5α-THDOC (334)	3α5α-THDOC-3,21-HFB ₂ (726)	22.74	499	257	6
DHEA (288)	DHEA-3HFB (484)	21.15	270	199	8
ADIOL (290)	ADIOL-HFB2-3,17-HFB2 (682)	17.57	468	453	10
ADIONE (286)	ADIONE-3-HFB (482)	21.20	482	268	8
Epiandrosterone (290)	Epiandrosterone-HFB (486)	21.61	442	213	16
Etiocholanolone (290)	Etiocholanolone-HFB (486)	20.33	442	213	16
Testosterone (288)	Testosterone-3,17-HFB ₂ (680)	17.61	680	665	12
5α-DHTestosterone (290)	5α-DHTestosterone-17-HFB (486)	22.16	414	399	10
3a5a-THT (292)	3α5α-THT-3,17-HFB ₂ (684)	16.42	455	241	8
3β5α-THT (292)	$3\beta 5\alpha$ -THT-3,17-HFB ₂ (684)	17.94	470	241	14
17β-estradiol (272)	17β -estradiol-3,17-HFB ₂ (664)	18.32	664	237	10
Estrone (270)	Estrone-3-HFB (466)	21.79	466	422	10
2-ME2 (302)	2-ME2-3,17-HFB ₂ (694)	21.19	694	267	10
Steroid sulfates fraction					
PREGS, Na (418)	PREG-3-HFB (512)	23.31	298	283	8
DHEAS, Na (390)	DHEA-3HFB (484)	21.15	270	199	8

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Table 2 : targeted steroids analyzed by GC-MS/MS in gilt saliva and their abbreviation. Steroids are

classified according to their chemical structure, i.e. their carbon number. Among the 35 targeted

steroids, 28 steroids were detected and quantified.

Targeted steroids	Abbreviation	Detected
C21-Steroids		
Pregnenolone	PREG	Yes
Pregnenolone sulfate	PREGS	Yes
20α-dihydropregnenolone	20α-DHPREG	Yes
17α -hydroxypregnenolone	17α-OH PREG	No
Progesterone	PROG	Yes
5α-dihydroprogesterone	5α-DHPROG	Yes
5β-dihydroprogesterone	5β-DHPROG	Yes
20α-dihydroprogesterone	20α-DHPROG	Yes
3α , 5α -tetrahydroprogesterone (allopregnanolone)	3α,5α-THPROG	Yes
3α ,5 β -tetrahydroprogesterone (pregnanolone)	3α,5β-THPROG	Yes
3β , 5α -tetrahydroprogesterone (iso-allopregnanolone)	3β,5α-THPROG	Yes
5α,20α-tetrahydroprogesterone	5α,20α-THPROG	Yes
5α -pregnane- 3α ,20 α -diol (3α , 5α ,20 α -hexahydroprogesterone)	3α,5α,20α-HHPROG	Yes
5β-pregnane-3α,20α-diol (3α,5β,20α-hexahydroprogesterone)	3α,5β,20α-HHPROG	Yes
5α -pregnane- 3β ,20 α -diol (3β ,5 α ,20 α -hexahydroprogesterone)	3β,5α,20α-HHPROG	Yes
16α-hydroxyprogesterone	16 <i>α</i> -OH PROG	No
17 <i>α</i> -hydroxyprogesterone	17α-OH PROG	No
Deoxycorticosterone	DOC	Yes
5α-dihydrodeoxycorticosterone	5α-DHDOC	No

$3\alpha 5\alpha$ -tetrahydrodeoxycorticosterone (allotetrahydrodeoxycorticosterone)	3α,5α-ΤΗDOC	No
C19-Steroids		
Dehydroepiandrosterone	DHEA	Yes
Dehydroepiandrosterone sulfate	DHEAS	Yes
Androstenediol (Δ 5-androstene 3 β ,17 β -diol)	ADIOL	Yes
Androstenedione (Δ 4-androstene 3,17-dione)	ADIONE	Yes
5α-androstane-3,17-dione	5α-ADIONE	Yes
5β-androstane-3,17-dione	5β-ADIONE	Yes
5α-androstane-3β-ol-17-one	Epiandrosterone	Yes
5β-androstane-3α-ol-17-one	Etiocholanolone	Yes
Testosterone	Т	Yes
5α-dihydrotestosterone	5α-DHT	Yes
3α,5α-tetrahydrotestosterone	3α,5α-THT	Yes
3β ,5 α -tetrahydrotestosterone	3 <i>β,</i> 5 <i>α</i> -THT	No
C18-Steroids		
Estrone	E1	Yes
17β-estradiol	17β-Ε2	Yes
2-methoxyestradiol	2-ME2	No





PREGS



Week

DHEAS



20α-DHPREG



P=0.0033



0.5

0.0

-5

-4



20α-DHPROG







-3

Week

-2

-1

Puberty





 $3\alpha 5\alpha 20\alpha$ -HHPROG



^{P=0.08} $3\alpha 5\beta 20\alpha$ -HHPROG



3β5α20α-HHPROG



DOC



ADIOL







5α-ADIONE





5β-ADIONE





Etiocholanolone



3α5α-THT

0.015 -

0.010

0.005

0.000

-5

-4

-3

Week

-2

lm/gu



ESTRONE



17β-ESTRADIOL P=0.02



Puberty

-1

Week

TESTOSTERONE

5α-DHT



PROG + 5\alpha/\beta-metabolites





P = 0.06

 5β -metabolites of PROG

