

# Evaluating suitability of saliva to measure steroid concentrations in grey seal pups

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# 1 **Evaluating suitability of saliva to measure steroid concentrations in grey seal pups**

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## 9 **Summary**

10 Measurement of steroids in wild pinnipeds can facilitate assessment of breeding, nutritional and stress status, and  
11 is useful in understanding behavioral responses. Even in young animals, sex steroids may be important in behavioral  
12 interactions and immune modulation. Use of saliva can avoid the large fluctuations seen in some steroids in plasma,  
13 and can negate the need for venipuncture, making it a potentially useful matrix in the wildlife. However, its utility  
14 in estimating steroid levels in wild young pinnipeds has not been evaluated. Here, we investigated the suitability  
15 of saliva for steroid hormone analysis in wild grey seal pups during their suckling and post-weaning fast periods.  
16 We collected saliva (n = 38) and plasma (n = 71) samples during the breeding season on the Isle of May, Scotland,  
17 2012. We investigated success of sample collection, ease of preparation, accuracy and precision of analysis, and,  
18 where possible, comparability of measurements (n = 27) from saliva and plasma. Plasma sampling was rapid,  
19 whereas sampling saliva took up to five times longer. Analytical performance criteria (parallelism, accuracy, and  
20 precision (intra and inter assay co-efficient of variation (% CV)) of commercial ELISA kits to measure estradiol,  
21 testosterone and cortisol in both matrices were assessed. Estradiol and cortisol assays performed well and can be  
22 used in plasma and saliva. However, we could not confidently validate testosterone for either matrix. Saliva  
23 estradiol correlated with levels in plasma. Saliva sample preparation was faster and simpler than plasma preparation  
24 because it did not require extraction. However, given the additional time taken to obtain saliva in the wild, the  
25 possibility of blood contamination from oral damage and the lower success rate in obtaining sufficient sample for  
26 analysis, we recommend that this matrix only be used as an alternative to plasma sampling measurement in  
27 pinnipeds when animals are anaesthetized, tolerate mouth swabbing, or have been trained to accept saliva sampling  
28 in captivity.

29 **Key words:** steroids, testosterone, estradiol, cortisol, grey seal, saliva, ELISA, juveniles, validation

## 30 **1. Introduction**

31 Steroids play a major part in successful reproduction and appropriate stress responses in mammals (Ruiz-Cortes,  
32 2012). Gonadal steroids are also important for neural development, immune system function, energy balance and  
33 behavior in mammals (see Klein (2000) for review). Glucocorticoids indicate nutritional and stress status (Bryan  
34 et al., 2015), and facilitate understanding of behavioral responses (French et al., 2013), immune function and  
35 regulation of energy balance in young animals (Bennett et al., 2013), as well as in adults (Lidgard et al., 2008).  
36 Sex steroid levels are lower in pre-pubertal individuals than adults, but are detectable and are important mediators  
37 of behavior and energy partitioning (Bell, 2018). Measurement of circulating sex steroids in young animals may  
38 thus be important in understanding behavioral and physiological differences within or between sexes (Bell, 2018).  
39 Sample collection methods and hormone analysis procedures should be refined to reduce animal handling time and  
40 stress. Blood samples are often used as the matrix of choice for analysis of steroid samples in pinnipeds (Bennett  
41 et al., 2013; Gardiner and Hall, 1997; Lidgard et al., 2008). However, cortisol levels change rapidly in response to  
42 a range of stressors. Lipemia, and hemolysis can interfere with colorimetric enzyme linked immunosorbent assays  
43 (ELISA) (Holder et al., 2010). Proteins can also negatively affect target availability (Holder et al., 2010).  
44 Differences in adrenal cortex activity means neonates can also produce additional steroid metabolites that cross-  
45 react with some immunoassay antibodies (Makin et al., 2010). Some alternatives to plasma sampling (e.g. urine  
46 (Constable et al., 2006), faeces (Petrauskas and Atkinson, 2006), blubber (Kershaw and Hall, 2016) and hair (Keogh  
47 et al., 2020)) for steroid analysis have been tested in captive and wild pinnipeds, but require substantial processing  
48 time before analysis.

49 Saliva sampling can avoid the need for venipuncture, is a simpler and more stable matrix that can be kept at ambient  
50 temperatures for many hours (Wood, 2009), steroid measurements appear unaffected by salivary flow rate (Vining  
51 et al., 1983) and may show less rapid and lower amplitude changes than plasma levels (Vincent and Michell, 1992),  
52 making them suitable to obtain baseline levels. Salivary samples from captive pinnipeds have been used to measure  
53 testosterone (Theodorou and Atkinson, 1998), estrone sulfate and progesterone (Pietraszek and Atkinson, 1994).  
54 However, wild animals may have oral injuries or infection, or may struggle during collection, and blood-  
55 contaminated saliva leads to elevated steroid values (Granger et al., 2004). Thus, it is important to evaluate the  
56 feasibility, quality and effectiveness of saliva sample collection against the more usual blood sampling from wild  
57 pinnipeds. Moreover, it is important to assess whether alternative approaches to obtaining material for hormone  
58 assays can offer better welfare and logistical advantages while retaining sampling success rate.

59 Grey seals (*Halichoerus grypus*) are large marine top predators, whose population status is indicative of the health  
60 status of the wider marine environment (Bäcklin et al., 2011; Jüssi et al., 2008), and, although globally rare, are  
61 regionally important in the North East Atlantic and Baltic Seas (SCOS, 2020). Although cortisol has been measured  
62 in post weaned grey seal pups (Bennett et al., 2013, 2012), there have been no studies on androgens and estrogens  
63 in either dependent or weaned pups of this species, and information is sparse of other young pinnipeds (Ortiz et al.,  
64 2003).

65 In this study, we assessed the suitability of saliva for analysis of three steroid hormones – estradiol (E), testosterone  
66 (T) and cortisol (C) – in wild grey seal male and female pups during their early development. We examined a)  
67 success of sample collection including approximate volume obtained, b) ease of sample preparation and analysis  
68 in the laboratory, c) performance of commercially available ELISA kits for salivary measurements and d) where  
69 available, comparability of hormone levels measured in contemporaneous saliva and plasma samples taken from  
70 pups at different nutritional stages in their first six weeks of life.

## 71 **2. Materials and methods**

### 72 **2.1 Study location and animals**

73 Samples were collected from grey seal pups as part of a long term study of grey seal reproduction during the  
74 breeding season (October-December) of 2012 on the Isle of May, Scotland (56°11'N, 02°33'W) on up to four  
75 occasions (nutritional stages – S1 – early suckling, S2 – late suckling, W1 – early weaning, W2 – late weaning),  
76 when animals were 4 to 38 days old (see Supplementary Table 1 and Supplementary Fig. 1). Samples were taken  
77 at approximately 10 day intervals (mean  $\pm$  SD = 9.86  $\pm$  1.54 days) and no significant difference was found in the  
78 length of time between sample occasions (Kruskal Wallis test (KW),  $p > 0.05$ ).

79 All sample collection was performed by personal license holders/ designated competent personnel under UK Home  
80 Office license PPL 60/4009. This work received ethical approval from the University of St Andrews Animal  
81 Welfare and Ethics Committee (AWEC) and was performed in compliance with Animal (Scientific Procedures)  
82 Act (ASPA) 1986 and the EU directive on the protection of animals used for scientific purposes (2010/63/EU).

### 83 **2.2 Sample collection**

84 *Blood sample collection.* Samples (n = 71) were collected during daylight between 10:50 to 17:15 GMT (no  
85 significant difference was found between time of the day during different nutritional stages KW with Bonferoni  
86 correction,  $p > 0,05$ ). Mothers were anesthetized using intramuscular injection from a blow dart (Zoletil<sub>100</sub>, Virbac,  
87 France) and capture took place once the anaesthesia had taken effect approximately 10 min. later. Mothers and

88 pups were then captured simultaneously. Pups were manually restrained, the blood sampling area cleaned and then  
89 anesthetized before they were weighed and sampled. Pups received approximately 0.1 mg/kg, i.v. of anesthetic  
90 (Zoletil<sub>100</sub>, Virbac, France) on all occasions (n = 65), a low risk dose for animals of this age class and effective  
91 for the sampling regime, to obtain blubber samples (Bennett et al., 2015). Four animals were not anesthetized  
92 because only blood samples were collected. Blood was collected from the extradural vein into a sterile 10 ml  
93 potassium ethylenediaminetetraacetic acid (EDTA) Vacutainer (Becton Dickinson, Oxon, UK).

94 *Saliva sample collection.* Saliva sampling was attempted during every blood sampling (n = 65) when animals were  
95 anaesthetized. Saliva samples were taken with inert polymer, cylindrical swabs (8 mm x 125 mm) designed for  
96 saliva sample collection in humans (Salimetrics, UK). Anesthetized pups were not aware or responsive, which  
97 allowed sample collection without additional restraint. The saliva swab was placed in the cheek where it would  
98 pose no threat to the anaesthetized animal and removed prior to returning the pup to its mother or when rejected  
99 by the animal on recovery from anesthesia. Before removal, if possible, the swab was used to collect saliva from  
100 the corners of the mouth. No samples were collected when signs of blood or abrasions were visible inside the mouth  
101 or around the gums. Any other particles of grass or dirt, if present, were not rinsed away to avoid diluting the saliva  
102 sample. The sample was discarded if any traces of blood on the polymer swab were observed during collection.  
103 The swab was then transferred to a 2 mm sterile syringe, which was used to squeeze the saliva out of the swab into  
104 a 2 ml collection tube (Salicap, IBL International, Germany). Saliva volume recovered was estimated to the nearest  
105 0.1 ml using gradations on the collection tubes. Blood and saliva samples were stored at ambient temperatures after  
106 collection for an average of 2 hours 9 min. (from 25 min. to 8 hours 49 min.) before they were processed.

### 107 **2.3 Duration and success of sample collection**

108 Plasma was collected within 2 minutes as in Bennett et al. (2012). Saliva sampling took approximately 10 min.,  
109 since this is the typical duration of anesthesia (Moss, pers. obs.). Sufficient plasma was obtained for all assays.

110 Success of saliva sample collection was assessed by comparing the volume collected with the minimum volume  
111 necessary for analysis of three steroids (150 µl for plasma and 250 µl for saliva). ‘Fully collected’ saliva samples  
112 contained the minimum volume of 250 µl (defined as possible to use for analysis of all three steroids), ‘partially  
113 collected’ samples had a volume around 100-250 µl (defined as possible to use for analysis of one or two steroids),  
114 and samples were considered ‘not collected’ when the saliva volume was too low for steroid analysis or sample  
115 was contaminated with blood.

### 116 **2.4 Sample preparation for analysis**

117 Blood samples were centrifuged in a swing-out bench top centrifuge at 2000 g for 15 min. The plasma was then  
118 drawn off using glass Pasteur pipettes, divided in 1.5 ml aliquots, immediately frozen and stored at -20° C until  
119 extraction and analysis. Plasma samples were extracted according to the Steroid Liquid Sample Extraction Protocol  
120 provided by Arbor Assays® (Michigan, USA), which is recommended by ELISA kit various manufacturers  
121 (Kumar, 2020) and validated for prepubertal human serum using EIA and RIA (Ankarberg-Lindgren et al., 2001;  
122 Norjavaara et al., 1996; Raivio et al., 2001). The collected material was evaporated in room temperature using a  
123 vacuum Eppendorf concentrator 5301 (Hamburg, Germany) and kept at -20°C prior to analysis. On the day of the  
124 ELISA the sample was thawed, reconstituted in 300 µl of steroid free serum (Ibl-International, Hamburg, Germany)  
125 and vortexed for 30 min. at room temperature, centrifuged for 10 min. at 3000 g and the supernatant used for  
126 ELISA. Saliva samples were heated at 60°C for 1 h then left to cool to room temperature and centrifuged at 3000 g  
127 for 10 min. prior to ELISA.

## 128 **2.5 Kit validation and sample analysis**

129 Commercially available ELISA kits (IBL-International, Hamburg, Germany, codes for products: TB - RE52151,  
130 TS - RE52631, EB - RE52041, ES - RE52601, CB - RE52061, CS - RE52611) were used according to the  
131 manufacturer's protocol. We attempted to validate each kit for use with saliva and plasma from grey seal pups (see  
132 Table, Mean ± SD) as follows: pooled plasma and pooled saliva samples were each diluted with the zero standard  
133 provided by the manufacturer to 0.75, 0.5 and 0.25 of neat concentration to ensure the measurements remained  
134 within the limits of the standard curve. The concentrations in the dilutions were then compared with expected  
135 concentrations from the neat pooled samples. The pooled extracted and reconstituted saliva and plasma samples  
136 were spiked 1:1 with low and high standard provided by the manufacturer to obtain recovery values. All standards  
137 and samples were analyzed in duplicate within any given plate.

138 Assay validation and sample measurements for plasma samples were performed across two plates, which were  
139 analyzed in two consecutive years, 2014 (Plate 1) and 2015 (Plate 2). Inter-assay CV of plasma was estimated using  
140 two samples, each measured in duplicate on each of the two plates. Mean and standard deviation concentrations  
141 were calculated for each sample in each plates and the standard deviation expressed as a percentage of the mean to  
142 give inter assay CV. The average inter-assay CV of the two samples is reported here. All saliva samples were  
143 measured within one plate; thus only intra-assay CV is reported. Intra-assay CV was calculated from the mean and  
144 standard deviation of the duplicates of each sample in the same plate and the mean intra-assay CV calculated.

## 145 **2.6 Prioritization of hormone assays for analysis**

146 A total number of plasma (for different plates) and saliva samples is provided in Table 1 (and in Supplementary  
147 Table 2 for more detail). When a partial saliva volume was collected and was below 100  $\mu$ l, the cortisol assay was  
148 prioritized since it required less sample ( $2 \times 25 \mu$ l) than the estradiol kit ( $2 \times 50 \mu$ l). When partial saliva volume  
149 collected allowed analysis of two steroids, testosterone was chosen as the second option, because, like cortisol, it  
150 required a lower volume of saliva ( $2 \times 25 \mu$ l) than the estradiol kit ( $2 \times 50 \mu$ l). As a result, the number of samples  
151 analyzed differs between steroids and is lowest for estradiol.

## 152 **2.7 Statistics**

153 Data analysis was performed using SPSS (IBM SPSS Statistics for Windows, Version 22.0. Armonk, NY: IBM  
154 Corp.) and statistical differences considered to be significant when  $p < 0.05$ . Parallelism was determined by visually  
155 investigating the graphs in which optical density of standard curve and pooled saliva and plasma samples were  
156 plotted against dilution (Supplementary Fig. 3-5). Accuracy of plasma kits was determined by performing linear  
157 regression of diluted pooled samples ( $n = 6$ ) of two plates comparing observed versus expected measurements,  
158 which should show a positive linear relationship with a slope of approximately 1.0 (a slope of 0.8-1.2 and  $r^2 > 0.95$   
159 was considered acceptable) (Hunt et al., 2014; Kershaw and Hall, 2016). Due to the low number of diluted samples  
160 we were only able to inspect recovery rates for saliva. Expected and observed concentrations in high and low spiked  
161 saliva and plasma samples were compared using a paired t-test. Dilution values were considered acceptable if they  
162 were within 30 % of the expected concentrations for low concentrations, and recoveries of spiked samples if they  
163 were within 20 % of expected values in both saliva and plasma (Yan, 2018). Precision of intra- and inter-assay  
164 CV was considered acceptable when  $< 20 \%$ .

165 Spearman rank correlation was used to evaluate whether the saliva sample volume obtained was related to steroid  
166 (ES, TS, CS) concentration. Testosterone was excluded from subsequent comparability analysis due to extremely  
167 high interassay CV. Even though the interassay CV of CB was within an acceptable range, it was removed from  
168 comparability analysis because we were unable to determine how much disturbance animals perceived prior to  
169 sampling, preventing us from reliably comparing CS and CB levels as a result of the potential stress response.

170 Linear regression analysis was used to examine the relationship between estradiol concentrations in saliva and  
171 plasma. Positively skewed ES values were log transformed using natural logarithm prior to regression analysis.

172 A generalized linear mixed model (GLMM) was used to evaluate the possible effect of nutritional stage, pup sex,  
173 body mass, age, and time of the day on EB and ES comparable samples ( $n = 27$ ). A normal distribution model and  
174 a linear link function was used for EB and a gamma distribution model with a log link function was used for ES.

175 Scaled identity covariance structure for nutritional stage as repeated measures for every pup (ID as a Subjects) and  
176 Satterthwaite approximation for estimating degrees of freedom was used (Zuur et al., 2009). Corrected Akaike  
177 information criteria (AICc) was used to select the best model. Residuals plots were inspected to check model fit.

### 178 **3. Results**

#### 179 **3.1 Success of sample collection**

180 Based on previous equivalent sampling (Bennett et al., 2012) the duration of blood collection was up to 5 times  
181 shorter (< 2 min.) than that for taking saliva (< 10 min.). We collected 38 useable saliva samples, which represents  
182 ~54 % of all 71 sampling occasions (Table 1). Unsuccessful saliva sampling events (N = 33) were due to limited  
183 saliva production, which did not provide the minimum amount of saliva during anesthesia, or because samples  
184 were contaminated with blood, and thus had to be discarded. The manufacturer states that a minimum of 500 µl of  
185 saliva is recommended for each ELISA analysis (requiring in this case 1,5 ml for all three steroids), but we never  
186 obtained 1.5 ml from any animal (Supplementary Fig. 2). However, the saliva analysis of all three steroids requires  
187 a minimum of 250-300 µl (TS and CS – 50 µl and ES 25 µl for each well) and we obtained this in 31 (82 %) cases.  
188 From samples that were collected, we had 7 (18 %) partial samples (five samples when only two steroids and three  
189 samples when only one steroid was analyzed). Therefore, we obtained enough saliva to analyze all three steroids  
190 in 44 % of all the animal sampling events. There were also three samples (two from nutritional stage S1 and one  
191 from S2) of TS that were above the limit of detection, and we did not have sufficient sample or reagent to perform  
192 a subsequent dilution. These high samples were not included in further statistical analysis.

#### 193 **3.2 Validation of ELISA kits for different matrices**

194 *Plasma (EDTA) validation.* We successfully validated plates for estradiol and cortisol, as they showed intra-assay  
195 CV < 10 % and spike recovery rates within 20% of expected (Table 1). Expected vs. observed spiked concentrations  
196 were not different for both steroids (CB:  $t = 0.24$ ,  $df = 3$ ,  $p = 0.83$ , EB:  $t = 0.7$ ,  $df = 3$ ,  $p = 0.54$ ). Dilution curves  
197 were visually parallel to standard curves for CB and EB (Supplementary Fig. 3 and Fig. 5). Inter-assay CV was 8  
198 % for EB and 25 % for CB. Expected vs. observed concentrations of diluted samples showed a linear relationship  
199 with a slope close to 1 for both cortisol ( $r^2 = 0.95$ ,  $df = 5$ ,  $p < 0.01$ , slope = 0.95) and estradiol ( $r^2 = 0.97$ ,  $df = 5$ ,  $p$   
200  $< 0.001$ , slope = 0.92). TB intra-assay CV was < 11 % and average spike recovery was 106 % (Table 1). Expected  
201 vs. observed spiked TB concentrations were not significantly different ( $t = -0.642$ ,  $df = 3$ ,  $p = 0.57$ ). Dilution curves  
202 were visually parallel to the standard curve (Supplementary Fig. 4) and dilution performance was good with a slope



203 close to 1 ( $r^2 = 0.99$ ,  $df = 5$ ,  $p < 0.001$ , slope = 0.99). However, inter-assay CV was unacceptably high (86 %), thus  
204 we did not successfully validate this ELISA for plasma testosterone.

205 *Saliva validation.* Dilution curves were visually parallel to standard curves for all steroids in saliva (Supplementary  
206 Fig. 3-5). TS ELISA overestimated the lowest diluted values up to 18 % and the CS kit by up to 15 %. The ES kit  
207 provided the poorest dilution performance with the lowest diluted values being underestimated by around 40 %,  
208 however such low values were not observed in the samples analyzed and recovery rate was above 80 % for ES  
209 values typically measured. Intra-assay CV was  $< 10\%$  and average recovery of spiked samples ranged from 90 %  
210 for ES to 95 % for CS and 111 % for TS (Table 1). Expected vs. observed spiked concentrations were not different  
211 for all steroids (ES:  $t = -0.89$ ,  $df = 1$ ,  $p = 0.54$ ; TS:  $t = -0.99$ ,  $df = 1$ ,  $p = 0.5$ ; CS:  $t = 1.4$ ,  $df = 1$ ,  $p = 0.4$ ). Saliva  
212 sample volume showed a significant moderate negative correlation with TS (Spearman  $r = -0.35$ ,  $p = 0.047$ ), while  
213 there were no significant correlations between saliva volume and ES (Spearman  $r = -0.32$ ,  $p = 0.08$ ) or CS  
214 (Spearman  $r = -0.03$ ,  $p = 0.86$ ).

215 *Table 1 should be inserted here*

### 216 **3.3 Correlations between steroid concentrations in saliva and plasma**

217 It was possible to compare the relationship of estradiol concentrations between 27 saliva and plasma samples. There  
218 was a significant positive relationship between log normalized ES and EB concentrations ( $r = 0.53$ ,  $r_{adj}^2 = 0.28$ ,  
219  $F_{1,25} = 9.64$ ,  $p < 0,01$ ) (Fig. 1) showing broad agreement between the two matrices.

220 *Fig. 1 should be placed here.*

### 221 **3.4 Sex and nutritional stage effects in different matrices**

222 The best model to describe variation in EB included two significant variables ( $AICc = 260.61$  vs.  $AICc_0 = 314.83$ ).  
223 EB was significantly different between nutritional stages ( $F_{3,22} = 7.18$ ,  $p < 0.01$ ) and higher in females ( $F_{1,22} = 4,42$ ,  
224  $p = 0,047$ ). EB was highest during the early suckling period, remained lower and stable during late suckling – early  
225 post-weaning fast periods and dropped again during the late post-weaning fast (Fig. 2A). No significant interactions  
226 were observed. Individual ID as a random intercept factor was removed, since it was not significant in any of the  
227 models (more details in Supplementary Table 3). Nutritional stage was the only significant effect in the model  
228 explaining variation in ES ( $AICc = 30.64$  vs.  $AICc_0 = 56.23$ ;  $F_{3,23} = 14.44$ ,  $p < 0.001$ ): higher concentrations  
229 occurred during early and late suckling compared with the postweaning fast (Fig. 2B).

230 *Fig. 2 should be placed here.*

## 231 **4 Discussion**

232 **4.1 Ease and success of saliva sample collection and sample preparation**

233 Saliva sample preparation was faster by up to one day and simpler because it required less equipment, chemicals  
234 and steps of extraction, than preparation of plasma samples. However, while saliva may appear to be an attractive  
235 option for measuring E and C in grey seals, there are a number of caveats. First, saliva samples take substantially  
236 longer (up to five times) to collect than plasma samples in wild grey seal pups, even when, as here, the animals are  
237 anaesthetized. As this type of anesthesia still requires venipuncture, the involvement of a trained professional is  
238 required to administer the anesthetic agent and obtain the sample and there is risk of extended apnea from the  
239 anesthesia. In unanesthetized animals handling time and stress, and the risk of oral damage and bleeding if the  
240 animal struggles can be substantially greater. Thus this method cannot be considered non-invasive for this species.  
241 In addition, it was often difficult to obtain enough sample for all three steroids to be analyzed: we were only able  
242 to obtain sufficient saliva to use in 44 % of sampling events (full samples), compared to 100 % for blood collection.  
243 We only obtained 0.5 ml or more – the volume recommended by the manufacturer for one of the steroid assays -  
244 in 32 % of saliva samples collected. We did not obtain an overall 1.5 ml, which was recommended by the  
245 manufacturer for all three assays to be performed, from any of the animals. Saliva may therefore be more  
246 appropriate for use in pinniped species that will tolerate mouth swabbing, in captive individuals that have been  
247 trained to accept saliva sampling, where anesthesia is deemed appropriate and safe for the species and age group,  
248 and if sufficient sample is guaranteed to be obtained for the analysis. These conditions are less likely to be met in  
249 a wild context. Our study shows that sufficient and uncontaminated saliva samples are unlikely to be collected with  
250 sufficient success to complete an intended analysis in grey seals, even when animals are anesthetized and when  
251 adequate time is given for saliva collection.

252 **4.2 Validity of commercial ELISA kits for steroid analysis from grey seal saliva and plasma**

253 This study demonstrates that estrogen can be measured with acceptable accuracy in both plasma and saliva of grey  
254 seal pups. Commercial ELISA kits may thus be suitable for estimating estradiol in grey seal pup plasma, however  
255 efforts need to be made to minimize between plate variation and ensure inter-plate confounds are minimized,  
256 because small biologically important differences might be too small to detect, especially when intra and inter assay  
257 variability are high. Saliva and plasma estradiol concentrations measured here were comparable with those reported  
258 previously in other marine mammals (Amaral et al., 2015; Theodorou and Atkinson, 1998). However, the range of  
259 steroid concentrations in blood vary more between studies than concentrations reported in saliva. Plasma estradiol  
260 was higher than concentrations reported in harbor seal young and adults by Lydersen & Kovacs (2005), northern

261 fur seals (*Callorhinus ursinus*) (Dierauf and Gulland, 2001) and southern elephant seals (*Mirounga leonina*)  
262 (Ferreira et al., 2005), but was similar to concentrations in harbor seal reported by Reijnders (1990).

263 For CB, lack of specificity due to high cross reactivity with progesterone and its metabolites could be one of the  
264 reasons for high CV. While T assay dilution performance for saliva and blood was from acceptable to excellent,  
265 the repeatability of TB concentrations was very poor. It is hard to identify what could have caused the differences  
266 between plates analyzed at different times. To the best of our knowledge this assay has not been validated for other  
267 wildlife in either saliva or plasma so it is difficult to establish whether the performance issues here were assay, age  
268 class or species specific.

#### 269 **4.3 Steroid concentration differences between sex and nutritional stages**

270 Saliva provided similar detail on estradiol differences between nutritional states compared with plasma. However,  
271 no sex differences were observed in saliva, whereas females consistently had higher estradiol than males in the  
272 matched plasma samples. These early sex differences have not been previously reported. The lack of detectable sex  
273 differences in saliva estradiol may limit its utility for measuring sex steroids in young animals. Higher  
274 concentrations of steroids during suckling than fasting may result from a reduction in metabolic processes after  
275 weaning when animals need to conserve energy (Reilly, 1991), are not metabolized rapidly after birth and still  
276 resemble concentrations transferred from mother *in utero* or are transferred to pups through mother's milk (Pundir  
277 et al., 2020). Steroid levels may track changes seen in the mother, which have high concentrations of estradiol at  
278 birth followed by a drop postpartum and increasing levels during the second half of lactation, possibly due to estrus  
279 (Mellish and Iverson, 2005). Higher concentrations of estradiol during the suckling period are related to immune  
280 system activity and cause sex-associated resistance to helminth parasite infection (Guzmán et al., 2009).  
281 Differences in steroid concentrations between nutritional categories and sex may thus have consequences for pups'  
282 behavior, immunity (Hall et al., 2002), energy balance (Bennett et al., 2013) and sensitivity to endocrine disrupting  
283 chemicals (Troisi et al., 2020) and may contribute to reported sex differences in their survival (Hall et al., 2001).

#### 284 **5 Conclusions**

285 This study is a first attempt to validate assays to measure sex steroids in grey seal pups in two different matrices.  
286 Estradiol and cortisol assays performed well and can be used in plasma and saliva, even in young grey seal pups.  
287 However, we could not confidently validate testosterone for both matrices. Although saliva is a promising  
288 analytical matrix for some steroids in some wildlife species, saliva collection is time consuming compared to rapid  
289 blood collection and easily affected by blood from gums or tongue in grey seals. The need for longer restraint and

290 anesthesia of animals during sampling compared with blood collection, and the lack of reliability in obtaining  
291 sufficient sample volume means this method is not ‘non-invasive’ and is not justifiable in this context for this study  
292 species. More importantly, saliva does not provide the same resolution of detail provided by plasma estradiol  
293 concentrations, since differences between sexes and the pattern of change between developmental states is not  
294 observed in saliva. We therefore recommend that salivary sampling is not performed in wild grey seals, unless  
295 animals are anesthetized for another reason; saliva can be obtained during that time in a way that minimizes blood  
296 and other contamination; and a large enough volume is obtained for the intended analysis. Saliva samples could be  
297 appropriate for analysis of estrogen and cortisol when collection occurs in captivity in trained animals that will  
298 tolerate mouth swabbing, where the stress effect of disturbance can be quantified and minimized, and where  
299 sufficient sample can be obtained, but may not be a good matrix for testosterone analysis, at least in young  
300 individuals.

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307

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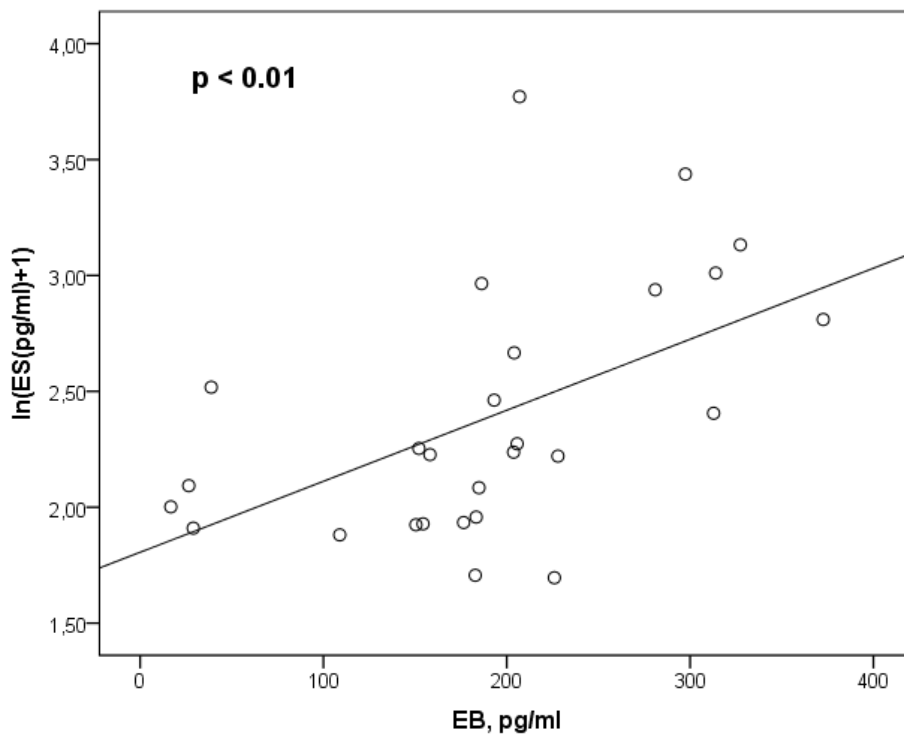
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429

430 **Table 1.** Descriptive statistics of ELISA analysis of three steroid hormones – testosterone, estradiol and cortisol – measured in plasma and saliva from suckling and  
 431 fasting grey seal pups.

Hormone	Testosterone			Estradiol			Cortisol		
	Plasma (ng/ml)		Saliva (pg/ml)	Plasma (pg/ml)		Saliva (pg/ml)	Plasma (ng/ml)		Saliva (µg/dl)
<b>Matrix</b>	Plasma (ng/ml)		Saliva (pg/ml)	Plasma (pg/ml)		Saliva (pg/ml)	Plasma (ng/ml)		Saliva (µg/dl)
<b>Plate number (year of analysis)</b>	Plate 1 (2014)	Plate 2 (2015)	Plate 1 (2014)	Plate 1 (2014)	Plate 2 (2015)	Plate 1 (2014)	Plate 1 (2014)	Plate 2 (2015)	Plate 1 (2014)
<b>N of samples of grey seal pups analyzed</b>	32	29	32	32	29	32	32	29	37
<b>Mean ± SE</b>	0.7 ± 0.02	0.2 ± 0.04	131.37 ± 14.23	233.45 ± 12.98	98.9 ± 12.12	11.77 ± 1.44	107.62 ± 4.95	49.4 ± 4.13	0.212 ± 0.02
<b>Min – Max</b>	0.51 – 1.12	0.05 – 1.26	31.07 – 313.42	150.43 – 385.9	16.7 – 236.88	4.45 – 42.45	63.97 – 161.25	16.56 – 99.98	0.065 – 0.479
<b>Obtained/expected % of sample dilution:</b>									
1 is the measured concentration of pooled sample	1 = 0.56	1 = 0.18	1 = 111.80	1 = 105.43	1 = 49.43	1 = 11.61	1 = 224.47	1 = 141.49	1 = 0.25
0.75 is 75 % sample mix	95.63 %	105.84 %	100.69 %	95.05 %	82.13 %	83.97 %	97.45 %	102.10 %	92.31 %
0.5 is 50 % sample mix	99.55 %	87.57 %	116.63 %	83.18 %	95.07 %	66.17 %	73.05 %	92.59 %	114.62 %
0.25 is 25 % sample mix	72 %	72.32 %	118.36 %	71.23 %	109.79 %	59.28 %	81.87 %	81.49 %	106.15 %
<b>RR % low spiked</b>	115.58 %	91.30 %	119.74 %	89 %	119.37 %	104.59 %	104.68 %	100.02 %	91.97 %
<b>RR % high spiked</b>	122.56 %	92.61 %	101.57 %	94.14 %	99.26 %	76.13 %	103.8 %	90.69 %	98.21 %
<b>Intra-assay CV (%)</b>	8.54 %	11.04 %	3.89 %	5.15 %	2.84 %	4.45 %	6.35 %	3.93 %	8.35 %
<b>Inter-assay CV (%)</b>	86.24 %		N/A	8.49 %		N/A	25.22 %		N/A

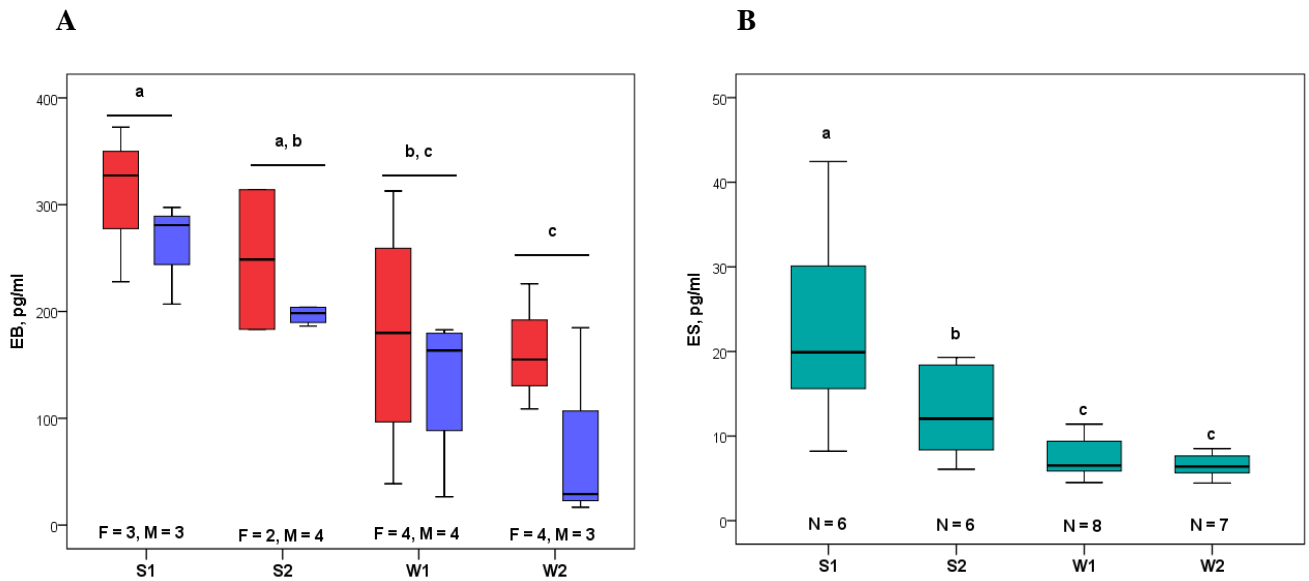
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433

434 **Fig. 1.** A linear relationship between the concentrations of lognormalized values of ES and concentrations of EB.

435 Line indicates linear relationship.



**Fig. 2.** Concentrations of EB (A) and ES (B) from grey seal pups during different nutritional stages (S1, S2, W1, W2). Red boxes denote EB concentrations of female, blue – of male grey sea pups. Letters indicate significant ( $p < 0.05$ ) differences between steroid concentrations during different nutritional stages. Middle line denotes median, outer box – interquartile range; whiskers – 95 % confidence intervals of steroid levels. Sample size (N) for each nutritional stage is given below each bar in the graphs.

436