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Cluster analysis reveals seasonal variation of sperm subpopulations in extended boar semen

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1 **Title:**

2 **Cluster analysis reveals seasonal variation of sperm subpopulations in extended boar**
3 **semen**

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5

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12

13 **Running head:** Sperm subpopulations in boar semen

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17

18 **Abstract**

19 This study aimed to identify motile sperm subpopulations in extended boar semen and to observe
20 the presumptive seasonal variation in their distribution. Data from 4837 boar ejaculates collected over
21 a two-year period were analyzed in terms of kinematic parameters by Computer Assisted Sperm
22 Analysis (CASA). Individual sperm data were used to determine subgroups of motile sperm within
23 the ejaculates using cluster analysis. Four motile sperm subpopulations (SP) were identified, with
24 distinct movement patterns: SP1 sperm with high velocity and high linearity; SP2 sperm with high
25 velocity but low linearity; SP3 sperm with low velocity but high linearity; and SP4 sperm with low
26 velocity and low linearity. SP1 constituted the least overall proportion within the ejaculates ($p < 0.05$).
27 Season of semen collection significantly influenced the different proportions of sperm subpopulations.
28 Spring was characterized by similar proportions of SP1 and SP4 (NS) and higher proportions of SP3.
29 Summer brought a decrease in both subgroups containing fast sperm (SP1 and SP2) ($p < 0.05$). During
30 autumn, increases in SP2 and SP4 were recorded. Winter substantially affected the proportions of all
31 sperm subpopulations ($p < 0.05$) and SP2 became the most represented subgroup, while SP1 (fast and
32 linear) reached its highest proportion compared to other seasons. In conclusion, extended boar semen
33 is structured in distinct motile sperm subpopulations whose proportions vary according to the season
34 of collection. Summer and autumn seem to have a negative impact on the fast and linear
35 subpopulation. Cluster analysis can be useful in revealing differences in semen quality that are not
36 normally detected by classical evaluation based on mean values.

37 **Keywords:** boar, CASA, cluster analysis, season, sperm subpopulations

38

39 **1. Introduction**

40 Computer Assisted Sperm Analysis (CASA) systems are able to record individual values for a
41 variable number of sperm analyzed during semen examination. Using a multi-step statistical analysis,
42 the individual values obtained allow a further, more detailed evaluation of the semen sample, with
43 classification of the sperm within subpopulations based on their kinematic parameters [1]. This offers
44 the possibility of a detailed profile for each ejaculate.

45 The starting point for detecting sperm subpopulations originated from the observation made by
46 W.V. Holt that boar ejaculates contain specific kinematic, relatively homogenous subgroups, which
47 can be deduced from the results offered by CASA profiling [2]. Numerous studies on these so-called
48 motile subpopulations have followed in the boar [3-5], as well as in bulls [6, 7], stallions [8], donkeys
49 [9], goat bucks [10], deer [11], dogs [12], and even fish [13]. Some authors currently believe that the
50 interpretation of CASA results based only on mean values represents an incomplete approach [14-16].
51 As such, stratifying an ejaculate into sperm subpopulations opens a new perspective on sperm
52 evaluation, which may be helpful in improving seminal dose calculations in assisted reproductive
53 programs [17] and to obtain improved information on ejaculates [18].

54 Although a consensus on the physiological role of sperm subpopulations has not yet been
55 achieved, the utility of this procedure has been demonstrated by studies that have revealed interesting
56 correlations, such as the link between the distribution of subpopulations and fertility [19], activation
57 of motility in presence of bicarbonate [4], *in vitro* capacitation and the acrosome reaction [3], or
58 cryoresistance [20]. These findings could be useful in selecting ejaculates with higher cryoresistance
59 and fertility.

60 Seasonal variation of some seminal parameters in farm animals has previously been demonstrated
61 [21-23] and it is thus reasonable to question whether seasonal dynamics may also affect the
62 distribution of motile sperm in specific subpopulations. Although sperm subpopulations analysis
63 based on CASA output has been a well-debated subject during recent years, to date there is no
64 information in the literature regarding seasonal variations in boar. Therefore, the aim of this study was

65 to identify motile sperm subpopulations in extended boar semen using CASA profiling and to
66 describe their seasonal variation.

67

68 **2. Materials and methods**

69 *2.1. Boars and semen*

70 The study evaluated data from 4837 ejaculates obtained from 702 healthy boars, aged between 9
71 months and 7 years, belonging to the Pietrain (90%), Landrace (5%), and Duroc (5%) breeds and
72 housed in an artificial insemination (AI) station in southern Germany. All the boars were commonly
73 used for semen collection for commercial purpose and were held in individual pens within the same
74 farm. Boars were fed a standard diet during the year, received water *ad libitum*, and experienced
75 natural photoperiods through windows facing outdoors. The stables were not furnished with
76 microclimate control systems, so boars were exposed to the natural variations of their environment
77 across seasons. The main climatic data recorded in the region are presented in Table 1.

78

Table 1

79 Semen was collected over a 24-month period, between March 2013 and February 2015 by manual
80 methods and using an artificial vagina. Ejaculates from the same boar were routinely collected weekly
81 within the AI station. After passing a general exam for quality (65% total sperm motility and $\geq 15 \times 10^9$
82 total sperm count), the ejaculates were diluted using Beltsville Thawing Solution (BTS, Minitube,
83 Tiefenbach, Germany) and submitted for liquid-state preservation at 17°C in doses of 100 mL (90 mL
84 extended semen + 10 mL air).

85 *2.2. Analysis of semen*

86 Semen analysis was performed after three days of storage, by means of the CASA system,
87 software SpermVision 3.7 (Minitube of America - MOFA[®], Verona, WI, USA), connected to a Zeiss
88 Axio Scope A1 microscope (Carl Zeiss MicroImaging GmbH, Göttingen, Germany) equipped with a
89 heated stage at 38°C. For the examination, semen samples of 3 μ L were placed in four-chamber slides

90 (Leja, Nieuw Vennepe, The Netherlands) with a chamber depth of 20 μm . The sperm kinematic
91 parameters were recorded from 7 successive fields per sample. For each examination, the field with
92 sperm motility closest to that calculated as mean value per sample was saved. The following
93 kinematic parameters were calculated: total sperm motility (TMot), progressive sperm motility
94 (PMot), average path velocity (VAP), curvilinear velocity (VCL), straight line velocity (VSL),
95 straightness (STR), linearity (LIN), wobble (WOB), amplitude of lateral head displacement (ALH)
96 and beat cross frequency (BCF).

97 Individual data of each motile spermatozoon within the field saved by computer were accessed.
98 Thus, calculated values were obtained for a total of 312,444 sperm analyzed over the two-year period.
99 All data were then incorporated in a single dataset.

100 2.3. Cluster analysis

101 In order to identify the subpopulations, the values were subjected to a series of statistical analyses
102 using IBM SPSS[®] Statistics version 21 (IBM[®] Corporation, Chicago, IL, USA). The first step was a
103 Principal Component Analysis (PCA) of the eight above-mentioned kinematic parameters. The aim of
104 the PCA is to reduce the number of variables in a database, in order to make data easier to visualize
105 and work with. The analysis is based on identifying correlations among the variables, variation
106 patterns, and subsequent extraction of the elements greatly influencing the variation of the values. We
107 used the Kaiser criterion, retaining only the components with eigenvalue greater than 1. After
108 applying PCA, we identified only two components with eigenvalue greater than 1. We then observed
109 two patterns of variation in the matrix of components; the first characterized VCL, VAP, VSL, BCF,
110 and ALH, while the second was specific to STR, LIN, and WOB. Accordingly, we selected the two
111 variables with the highest values in the matrix of components for further analysis, namely VAP
112 (0.973) and LIN (0.967). The next step was the standardization of values for VAP and LIN such that
113 the mean values would be 0 and the standard deviations 1. The purpose of value standardization is to
114 avoid erroneous calculation of the subgroup centers caused by different value scales of the two
115 parameters. Further, the assignment of sperm into subgroups was performed using the k-means cluster
116 analysis based on Euclidean distances, as described in previous studies [6, 7, 12]. Each spermatozoon

117 was assigned to a cluster (subgroup, subpopulation) such that its movement pattern was similar or
118 close to the other sperm belonging to the same cluster, but significantly different to the movement
119 pattern of the sperm belonging to other clusters. After applying k-means cluster analysis, the cluster
120 membership of each spermatozoon was saved and used to subsequently calculate the proportion of
121 each identified subpopulation. To ascertain if the percentages of motile sperm subpopulations were
122 dependent on a specific season from a statistical standpoint, the χ^2 test was applied. Also, to analyze
123 the variance of the share of a certain subpopulation among seasons, the ANOVA test was used. The
124 differences in frequencies between two groups (for example fast sperm vs. slow sperm) were
125 evaluated through the Paired Sample *t*-test.

126

127 **3. Results**

128 *3.1. Overall kinematic parameters of semen as determined by CASA profiling*

129 All selected kinematic parameters were affected by the season of semen collection, although
130 similarities between some seasons in terms of certain parameters could be observed (Table 2).

131 TMot registered the highest mean value (\pm SD) during winter ($80.1\% \pm 10.4$) and lowest during
132 summer ($73.4\% \pm 13.3$). The mean value registered in spring ($80.1\% \pm 10.5$) was similar to that
133 obtained in winter (NS), whereas autumn values ($73.7\% \pm 14.4$) were not significantly different from
134 those of the summer (NS). The same was observed for PMot, with higher values during winter and
135 spring ($75.2\% \pm 12.3$ and $74.1\% \pm 13.2$ respectively), and lower values in autumn and summer
136 ($64.6\% \pm 17.8$ and $64.9\% \pm 16.5$, respectively).

137 Significant variability of sperm velocity was observed among seasons. VCL was the parameter
138 most responsive to the annual changes, showing significant oscillations from one season to another (p
139 <0.05). The highest values for sperm velocity, reflected by VAP, VCL, and VSL parameters were
140 observed during winter ($p <0.05$). No differences between summer and autumn were detected with
141 regards to VAP (NS).

142 Although characterized by a lower overall velocity the sperm showed a less corrugated trajectory
143 during summer, as described by the parameters STR, LIN, and WOB. Similar values for STR, LIN,
144 and WOB were recorded during summer and spring (NS), and these values were both lower during
145 the autumn and winter months ($p <0.05$).

146 Table 2

147 *3.2. Sperm subpopulations*

148 The cluster analysis revealed a clear heterogeneity of sperm populations, with the coexistence of
149 four distinct subgroups of motile sperm characterized by specific values of velocity and linearity. The
150 mean values of kinematic parameters for the four subpopulations are presented in Table 3. Briefly,
151 their main characteristics were as follows (Fig. 1):

- 152 - Subpopulation 1 (SP1) represented sperm with a high VAP and a high LIN (Fig. 2A). About 15%
153 of the sperm was assigned to this subgroup;
- 154 - Subpopulation 2 (SP2) included sperm with a high VAP but a low LIN (Fig. 2B). This subgroup
155 contained 28% of the sperm;
- 156 - Subpopulation 3 (SP3) was defined as sperm showing low velocity but high linearity (Fig. 2C).
157 The share of this subgroup was about 35%.
- 158 - Subpopulation 4 (SP4) consisted of sperm with a low VAP and a low LIN (Fig. 2D).
159 Approximately 22% of the sperm were part of this cluster.

160 Figure 1

161 Figure 2

162 Besides the previously mentioned differences in sperm velocity and linearity, there were also
163 differences regarding ALH and BCF (Table 3). The SP2 and SP1 showed higher values for ALH,
164 along with higher values of VAP and BCF. The SP3 contained sperm with the lowest ALH, while the
165 SP4 was not only defined by a low VAP, but also by a low BCF.

166 Table 3

167 The season in which semen was collected influenced the distribution of SPs, as revealed by the χ^2
168 test ($p < 0.05$) (Fig. 3). Spring was characterized by similar proportions (NS) of SP1 (16.9%) and SP4
169 (17.9%) and higher proportions of SP3 (37.1%).

170 Figure 3

171 During the summer months the proportion of SP1 (12.2%) and SP2 (21.7%) decreased compared
172 to spring, while SP3 (39.7%) and SP4 (26.4%) showed the opposite changes during this season (p
173 < 0.05). Summer had a negative effect on the velocity of sperm and their linearity, as the proportion of
174 both subgroups of rapid sperm (SP1 + SP2) and the subgroup of linear sperm (SP1 + SP3) decreased
175 ($p < 0.05$). It must be pointed out that the changes in sperm linearity were not detected by the overall
176 analysis of mean values of the LIN parameter (NS) (Table 2). Most sperm during summer were from
177 SP3 and the second most abundant was the SP4 group.

178 The autumn season also brought some changes in the distribution of motile sperm within
179 subpopulations compared to summer ($p < 0.05$). SP3 (33.7%) remained the most numerous, but

180 significant increases in SP2 and SP4 were also recorded ($p < 0.05$). SP1 (sperm with high velocity and
181 high linearity) showed similar values to those of summer (NS). Both in summer and autumn, the SP1
182 group recorded a much lower share when compared to the other three subpopulations (Fig. 3).
183 Although the overall analysis based on mean values detected no changes (NS) in sperm VAP (Table
184 2), significantly more sperm were considered to be fast in autumn compared to summer (SP1 + SP2 =
185 37.6% in autumn versus 33.9% in summer) ($p < 0.05$).

186 The cold season significantly changed the proportions of all sperm subpopulations. The SP1 (fast
187 and linear) group increased substantially (21.1%), reaching the highest proportion among the different
188 seasons ($p < 0.05$). At the same time, the SP4 (slow, non-linear) group decreased considerably (p
189 < 0.05), and was the subgroup with the lowest number of sperm (13.9%). A remarkable rise was also
190 shown by SP2 (35.6%), which became the most frequent subgroup during winter ($p < 0.05$). The
191 increases in overall sperm velocity (Table 2) during winter were reflected by increases of both
192 subgroups with high velocity (SP1 and SP2). In contrast, the increases in overall linearity (Table 2)
193 were associated only with significant increases in the subgroup with high linearity and high velocity
194 (SP1), while the subgroup with high linearity and low velocity (SP3) showed a lower share compared
195 to autumn (Fig. 3).

196

197 **4. Discussion**

198 In our study, we observed the coexistence of four subpopulations of sperm with different
199 movement patterns. The presence of four distinct subpopulations in semen has also reported by other
200 authors, both in boar [3, 24] and in other species [8, 25-27]. The subpopulation containing sperm with
201 higher velocity and higher linearity was less numerous than the other three subpopulations; this is
202 consistent with previous results [3] and suggests a relatively poor representation of this kind of sperm
203 within boar semen. In comparison, ram semen seems to be mostly constituted of rapid, linear sperm
204 [28].

205 Seasonal variations of seminal parameters in boar have been studied for years and it is now widely
206 accepted that summer causes a decrease in semen quality and quantity. Previous studies observed the
207 effects of the hot season on a variety of parameters, such as semen volume [29], total number of
208 sperm [30], sperm motility [31], sperm viability [32], sperm morphology [33], sperm agglutination
209 [34], cryoresistance [35], protein content [36], and acrosin activity [37]. All these fluctuations gave
210 rise to discussions of “seasonal infertility”, described in boars, but also in sows, which is defined
211 mostly as reduced fertility occurring during the summer months and in early autumn [38, 39].

212 Our investigation revealed that the collection season exerts a great influence on the distribution of
213 sperm subpopulations in extended boar semen. The subgroup of fast and linear sperm, which is
214 considered by some authors to be the one with the highest fertilizing potential [12, 28], was best
215 represented during the winter, and was poorly represented during the autumn and summer months. On
216 the other hand, the subgroup of slow and non-linear sperm, which theoretically have lower chances of
217 reaching the oviduct, was more numerous during the autumn and summer and was less numerous
218 during the spring and winter. These data suggest that the previously described decrease in swine
219 fertility during the hot season [39], could also be also related to changes in motile sperm
220 subpopulations.

221 Seasonal analysis of sperm subpopulations could contribute to a better definition of semen quality
222 throughout the year and could thus have a significant economic impact, as more efficient semen
223 evaluation will lead to an improvement of production of AI doses [40]. Furthermore, seasonal analysis

224 may reveal aspects that may be ignored by usual evaluation methods. For example, in our study the
225 overall percentage of motile sperm was similar in winter and spring. However, when we analyzed
226 individual motile sperm subpopulations we observed a clear difference in the structure of ejaculates,
227 with a significantly higher percentage of rapid sperm present during the winter months. Given the fact
228 that the percentage of motile sperm is still the main criterion for assessing semen quality in production
229 centers [41, 42], one may mistakenly think after a quick look, that the ejaculates collected during
230 spring were similar to those collected during winter, when in fact they were not. Moreover, while the
231 overall analysis based on mean values did not detect significant differences in sperm VAP between
232 summer and autumn, cluster analysis revealed that during the autumn months significantly more
233 sperm could be classified as fast. This means that the succession of seasons does indeed have an effect
234 on the velocity of sperm, but probably not for all subpopulations and not to the same extent.

235 The fact that seasons modify the distribution of sperm within subpopulations could influence the
236 perception we have on seasonal variations of kinematic parameters in boar semen. Thus, seasonal
237 dynamics in kinematic parameters might be caused by changes in the proportion of sperm assigned to
238 the different subpopulations and not by an overall increase/decrease in values for all the ejaculated
239 sperm. For example, boar sperm velocity seems to be lower during summer [31]. Our study revealed
240 that the subpopulations with fast sperm are less numerous during summer (Fig. 3), which may suggest
241 that the overall decrease of VAP is not necessarily caused by a decrease in velocity of all sperm, but
242 rather by changes in the distribution of sperm within subpopulations, with some sperm “passing” from
243 fast subpopulations to slow subpopulations. This would mean that not all the sperm analyzed suffered
244 during the hot season, but only a certain percentage of sperm whose decreased velocity led to a
245 decline in overall velocity. Furthermore, this might suggest that boar testicles produce “resistant”
246 sperm, which will retain their characteristics even in less favorable environmental conditions and
247 “sensitive” sperm, which will be easily affected by different factors, such as temperature. This
248 hypothesis is also supported by the fact that in our study, the subpopulation containing sperm with
249 high velocity and high linearity remained stable over the summer and autumn, while all the other
250 subpopulations suffered significant changes (Fig. 3).

251 The main cause for the decrease in the number of fast and linear sperm during the summer months
252 was probably heat stress, as it is well known that the high temperatures specific to the hot season
253 affect spermatogenesis [43-45]. There have been a few theories proposed that have attempted to
254 explain the overall seasonal variation in boar semen. Based on previous studies, the main factors
255 implicated in this variation include temperature [22], photoperiod [30], and humidity [36]; some
256 authors have indicated the existence of an ancestral mechanism inherited from the wild boar [37, 46].
257 A useful step in determining the cause of seasonal variations would be that of finding the basis of the
258 impairment. Sperm gain their mobility during the maturation phase in the epididymis, but it is
259 currently difficult to determine whether the reduction in the number of fast and linear sperm reflects
260 impairment of the epididymal function or that of specific testicular segments involved in
261 spermatogenesis. For example, membrane integrity is strongly correlated with the functional status of
262 sperm mitochondria [47], so disorders in the plasma membrane might result in dysfunctions of the
263 mitochondrial sheath, leading to decreased velocity.

264 The seasonality of reproduction in the swine is once again confirmed by this study. In addition to
265 all the previous studies indicating not only variability in fertility of sows, but also in a large number of
266 seminal parameters in the boar, our study describes the seasonal changes occurring in the distribution
267 of motile sperm in distinct subpopulations based on their movement patterns (Fig. 3). What still
268 remains unclear, however, is the underlying algorithm determining the changes in the proportion of
269 sperm subpopulations. Apparently, some fast and linear sperm pass to another subgroup under
270 stressful conditions. We cannot specify yet in which direction are they migrating. Are they losing
271 velocity by going to the subgroup with slow and linear sperm? Or on the contrary, are they sacrificing
272 the uniformity of their movement to move towards a fast and non-linear subgroup? We cannot
273 exclude that the algorithm of migration differs according to the factors that cause it. For example, in
274 our study the proportion of fast and linear sperm, and also of fast and non-linear sperm, decreased
275 from spring to summer (Fig. 3). At the same time, the proportion of slow and linear, but also of slow
276 and non-linear sperm, increased. This suggests a loss in sperm velocity while maintaining sperm
277 linearity. On the other hand, in the study by Ramió et al. [3], adding progesterone in the *in vitro*
278 capacitating medium caused an increase in the proportion of rapid, but non-linear sperm indicating

279 that some sperm lose their linearity by passing into the non-linear subgroup and thus follow a
280 different migration algorithm than that observed in our study.

281 Examiners might consider introducing clustering analysis in their studies on semen. Although the
282 new trend among researchers is to use flow cytometry analysis, CASA systems are still widely used.
283 We believe it would be helpful if the use of CASA for research purposes were extended to the
284 determination of motile sperm subpopulations. This might complicate studies somewhat, but at the
285 same time it would make them more comprehensive. Conceivably the behavior of sperm based on
286 specific parameters should not be generalized to the entire sample, and we should rather study the
287 effects on each subpopulation, identifying which is the most affected. Clustering analysis is no longer
288 a novelty for theriogenologists, and its inclusion as an additional feature of semen examination would
289 only represent a logical step.

290 In conclusion, extended boar semen seems to be structured in distinct motile sperm
291 subpopulations, defined by specific movement patterns. Sperm could be classified as fast and linear,
292 fast and non-linear, slow and linear, and respectively, slow and non-linear. The proportion of each
293 class varies greatly according to the season, and the hot season seems to have a negative impact on the
294 percentage of fast and linear sperm. These findings might change the way we perceive seasonal
295 variations in overall values of the kinematic parameters in boar semen. These variations might be
296 caused by changes in the number assigned to each subpopulation and not by increases or decreases in
297 the values of all ejaculated sperm. We recommend introducing clustering analysis in studies on semen
298 where possible. Sperm subpopulations respond differently to environmental conditions, and the values
299 of kinematic parameters should not be generalized to the entire sample as a whole.

300

301 **Conflict of interest**

302 There are no conflicts of interest associated with this publication.

303

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308

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Table 1

Seasonal mean values for the main climatic factors recorded in the region^a and period of study.

Season ^b	Temperature (°C)	Relative humidity (%)	Duration of daylight (h)	Atmospheric pressure (hPa)
Spring	9.3	73.3	13.7	1013.7
Summer	18.4	70.0	15.4	1016.0
Autumn	10.4	87.8	10.8	1016.5
Winter	2.6	89.1	9.1	1017.5

^a Sources of data: www.timeanddate.com/sun/germany/nuremberg;

<http://umweltdaten.nuernberg.de/wetterdaten/messstation-nuernberg-flugfeld/archiv/>

^b Spring = March–May; Summer = June–August; Autumn = September–November; Winter = December–February

Table 2

Overall mean values of selected kinematic parameters of boar semen recorded during the four seasons. Values are reported as means \pm SD of 4837 ejaculates from 702 boars.

Parameter ^c	Spring ^f	Summer	Autumn	Winter
TMot (%) ^c	80.1 \pm 10.5 ^a	73.4 \pm 13.3 ^b	73.7 \pm 14.4 ^b	80.1 \pm 10.4 ^a
PMot (%)	74.1 \pm 13.2 ^a	64.9 \pm 16.5 ^b	64.6 \pm 17.8 ^b	75.2 \pm 12.3 ^a
VAP (μ m/s)	66.4 \pm 12.2 ^a	59.0 \pm 12.7 ^b	60.1 \pm 14.2 ^b	73.5 \pm 9.8 ^c
VCL (μ m/s)	126.8 \pm 28.7 ^a	112.9 \pm 29.1 ^b	119.6 \pm 32.4 ^c	143.4 \pm 25.6 ^d
VSL (μ m/s)	52.7 \pm 9.4 ^a	47.1 \pm 9.9 ^b	46.5 \pm 10.8 ^b	57.2 \pm 8.53 ^c
ALH (μ m)	2.98 \pm 0.63 ^a	2.86 \pm 0.65 ^b	3.05 \pm 0.72 ^a	3.33 \pm 0.61 ^c
BCF (Hz)	36.6 \pm 3.2 ^a	34.5 \pm 3.5 ^b	33.8 \pm 4.5 ^c	37.3 \pm 2.94 ^d
STR (VSL/VAP)	0.79 \pm 0.08 ^a	0.80 \pm 0.07 ^a	0.77 \pm 0.08 ^b	0.78 \pm 0.08 ^b
WOB (VAP/VCL)	0.53 \pm 0.06 ^a	0.53 \pm 0.06 ^a	0.51 \pm 0.06 ^c	0.52 \pm 0.06 ^d
LIN (VSL/VCL)	0.42 \pm 0.08 ^a	0.43 \pm 0.08 ^a	0.40 \pm 0.08 ^b	0.41 \pm 0.08 ^c

^{a,b,c,d} Within the same row, different superscripts show significant difference at $p < 0.05$;

^c Abbreviations: TMot = total sperm motility; PMot = progressive sperm motility; VCL = curvilinear velocity; VAP = average path velocity; VSL = straight-line velocity; ALH = amplitude of lateral head displacement; BCF = beat cross frequency; STR = straightness of track; WOB = wobble; LIN = linearity of track;

^f Spring = March–May; Summer = June–August; Autumn = September–November; Winter = December–February

Table 3

Kinematic parameters of the four sperm subpopulations (SP1 to SP4). Values are means \pm SD of 4837 ejaculates from 702 boars.

Parameter ^e	SP 1	SP 2	SP 3	SP 4
VAP ($\mu\text{m/s}$)	100.1 \pm 18.2 ^a	81.3 \pm 18.9 ^b	50.5 \pm 16.5 ^c	25.4 \pm 14.5 ^d
VCL ($\mu\text{m/s}$)	167.1 \pm 43.4 ^a	183.6 \pm 51.4 ^b	81.8 \pm 30.7 ^c	68.5 \pm 40.7 ^d
VSL ($\mu\text{m/s}$)	89.0 \pm 17.0 ^a	51.9 \pm 16.3 ^b	45.1 \pm 15.7 ^c	15.9 \pm 10.6 ^d
ALH (μm)	3.57 \pm 1.30 ^a	4.17 \pm 1.48 ^b	2.21 \pm 1.13 ^c	2.38 \pm 1.54 ^d
BCF (Hz)	37.5 \pm 9.1 ^a	37.1 \pm 8.1 ^b	36.5 \pm 12.6 ^c	22.27 \pm 13.3 ^d
STR (VSL/VAP)	0.89 \pm 0.08 ^a	0.66 \pm 0.19 ^b	0.89 \pm 0.07 ^c	0.63 \pm 0.21 ^d
WOB (VAP/VCL)	0.62 \pm 0.10 ^a	0.45 \pm 0.07 ^b	0.64 \pm 0.12 ^c	0.39 \pm 0.13 ^d
LIN (VSL/VCL)	0.55 \pm 0.11 ^a	0.30 \pm 0.10 ^b	0.57 \pm 0.11 ^c	0.24 \pm 0.11 ^d

^{a,b,c,d} Within the same row, different superscripts show significant difference at $p < 0.05$.

^e Abbreviations: VCL = curvilinear velocity; VAP = average path velocity; VSL = straight-line velocity; ALH = amplitude of lateral head displacement; BCF = beat cross frequency; STR = straightness of track; WOB = wobble; LIN = linearity of track

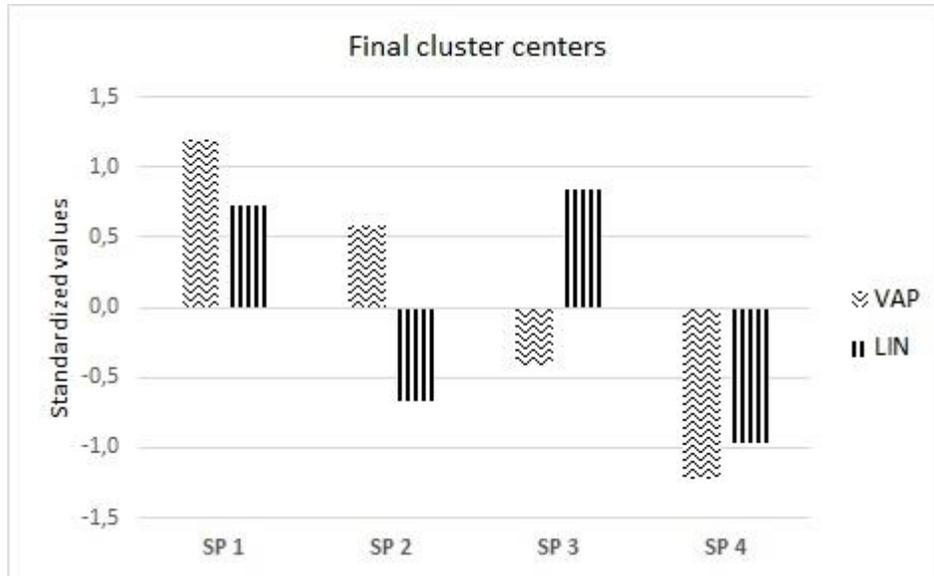


Fig. 1. Characteristics of four motile sperm subpopulations identified in boar semen in terms of average path velocity (VAP) and linearity (LIN). Values are standardized such that the mean values would be 0 and the standard deviations 1.

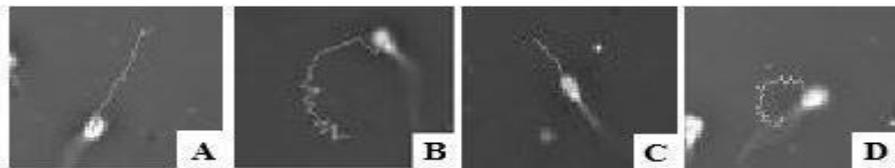


Fig. 2. Characteristic motility tracks of boar sperm of the four motile subpopulations (A: subpopulation1; B: subpopulation 2; C: subpopulation 3; D: subpopulation 4).

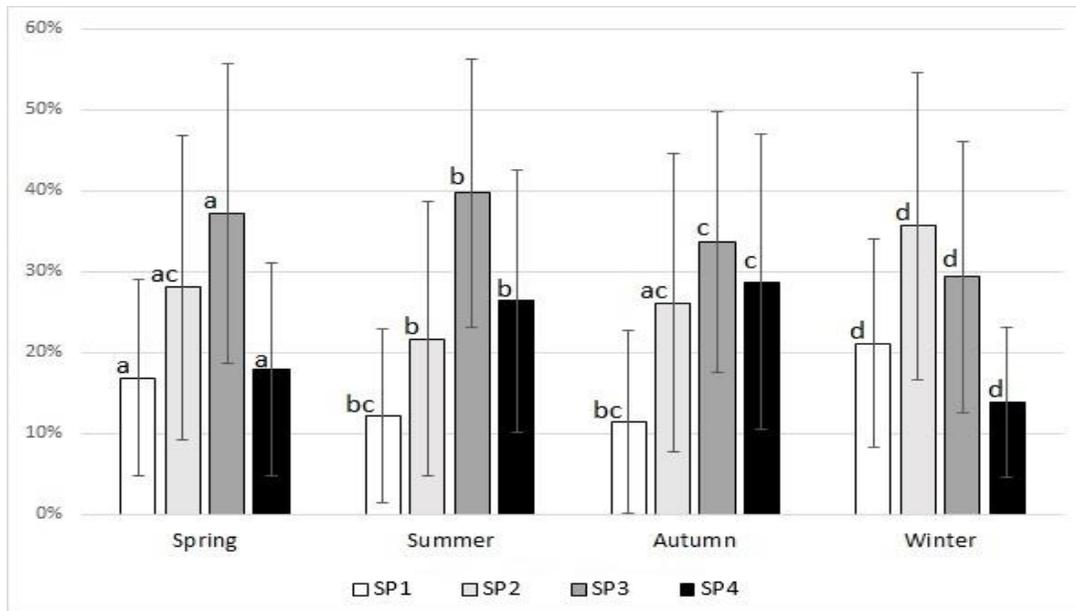


Fig. 3. Distribution of sperm subpopulations in boar semen during the four seasons. Values are means \pm SD of 4837 ejaculates of 702 boars. Different superscripts indicate significant differences among seasons within the same subpopulation.

Spring = March–May; Summer = June–August; Autumn = September–November; Winter = December–February.