

1 Running head: Organization of drinking behavior in birds

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3 **Satiety splits drinking behavior into bouts: the organization of drinking in birds¹**

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5 **J. Rusakovica,^{*2} T. Plötz,[†] V. Kremer,[‡] P. Glover,[‡] I. Kyriazakis^{*}**

6 ^{*}School of Agriculture, Food and Rural Development, Newcastle University, Newcastle

7 upon Tyne, Tyne and Wear, NE1 7RU, UK;

8 [†]Open Lab, School of Computing Science, Newcastle University;

9 [‡]Aviagen Turkeys Ltd, Tattenhall, Cheshire, CH3 9GA, UK;

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18 ² Corresponding author: julija.rusakovica@ncl.ac.uk

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20 **ABSTRACT:** The regulation of the drinking behavior of animals is usually overlooked and
21 the traits associated with it are not well-defined. We used a unique data set to develop ideas
22 about the analysis and regulation of drinking behavior in birds. The data were generated by a
23 custom-made equipment that measures automatically the individual drinking behavior of a
24 large number of turkeys from different genetic lines. We hypothesized that there is a
25 biologically significant unit by which drinking behavior can be expressed and understood. We
26 developed a novel method, based on mixture distribution models, to allow clustering of
27 drinking events and splitting behavior into bouts. Drinking behavior was found to be predicated
28 on the same principles of satiety that underlie feeding behavior. Within bouts, drinking was
29 interrupted by short non-drinking intervals, whereas bouts were separated by long non-drinking
30 intervals, indicative of bird physiological need. Based on this methodology, a number of
31 drinking behavior traits were identified that revealed differences in the organization of drinking
32 behavior between the turkey genetic lines. Similarly, time accumulation patterns of drinking
33 behavior traits within a day differed within and between genetic lines, suggesting that variation
34 in drinking behavior exists and birds use different behavioral strategies to meet their water
35 intake requirements. However, evolution of drinking behavior traits over time was similar
36 between the lines, suggesting conservation of this behavioral organization. As well as
37 providing ideas about the regulation of drinking behavior, the developed behavioral traits may
38 be of practical relevance, as water utilisation, along with feed efficiency, is part of overall
39 biological efficiency. The methodology should be applicable for the definition of drinking traits
40 in other livestock species, and be used for the identification of deviations from ‘normal’
41 drinking behavior.

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43 **Key words:** drinking behavior, modelling behavior, satiety, turkey, water intake.

44 INTRODUCTION

45 Water intake serves several physiological functions, and for some animals drinking may also
46 fulfil behavioral needs. There are also instances where water consumption exceeds
47 physiological needs (McKinley et. al., 2004). Animals of several species do not drink
48 continuously in time, but do so in bouts. Within such bouts actual drinking may be interrupted
49 by non-drinking intervals. The question is whether this behavior of drinking in bouts is
50 underlined by any physiological principles, or occurs randomly in time. There are several
51 advantages in understanding the basis of ‘normal’ drinking behaviors in animals. Feeding
52 behavior, for example, is understood on the basis of the physiological state of satiety.
53 Deviations from the ‘normal’ patterns of feeding behavior may then be indicative of health and
54 welfare problems. However, it has been suggested that the same principle of satiety (substituted
55 by the term of thirst) cannot be applied or understood in animals, presumably because of both
56 the physiological and non-physiological functions of drinking behaviour (Rolls and Rolls,
57 1991). The above arguments have not been helped by the lack of equipment able to record
58 continuously drinking of animals in a social context. As a consequence, the methodology of
59 measuring and analysing drinking behavior in animals has received significantly little attention.
60 In this paper we exploit a novel system to measure individual drinking behavior in birds, kept
61 in commercial groups. Our hypothesis is that their drinking behavior will be predicated on the
62 same principles of satiety that underline feeding behavior. Our expectation is that, within bouts,
63 actual drinking may be interrupted by short non-drinking intervals, while bouts will be
64 separated by long non-drinking intervals indicative of bird physiological need. We
65 subsequently develop traits related to drinking behavior and its regulation, and investigate how
66 these traits may be affected by factors such as bird genotype and age.

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69 MATERIALS AND METHODS

70 *Ethical note*

71 The data used in this study was derived from animals in the primary turkey pedigree breeding
72 programme of Aviagen Turkeys Ltd. Our study was mainly observational and used data
73 routinely recorded on the pedigree farm; birds were individually identified with RFID (Radio-
74 Frequency Identification) tags. Individual identification is the basis of genetic selection on
75 pedigree farms. We have previously shown that these RFID do not cause any adverse effects
76 on behaviour (Howie et. al., 2009a). The water intake recording equipment and the cameras,
77 when used, were installed before the arrival of the turkeys, so no disturbance was caused. A
78 small number of birds were handled to identify individually through video observations. The
79 handling of the birds was done by professional staff to minimise disturbance and subsequent
80 observations suggested that this handling and marking had no effect on their behavior. The
81 latter procedure was approved by the Newcastle University Animal Welfare Ethics Review
82 Board.

83 *Birds, Housing and Water Intake recording equipment*

84 Records of visits to electronic drinkers were obtained for three turkey breeding lines: (1) line
85 A (n = 954 777 events) from 4627 turkeys from 6-9 weeks of age, (2) line B (n = 770 984
86 events) from 2351 turkeys from 10-13 weeks of age and (3) line C (n = 146 170 events) from
87 291 turkeys from 10-13 weeks of age (Table 1).

88 Birds were male turkeys. Birds from line A were from a paternal line, selected with an emphasis
89 on feed efficiency, breast meat yield and growth, whereas lines B and C represented maternal
90 lines, with an emphasis on reproductive performance and feed efficiency. All three lines were
91 selected for leg health and fitness traits. They were part of routine genetic evaluation at Aviagen
92 Turkeys. Testing of lines B and C at the same age allowed us to test whether the drinking
93 behavior of birds differed between genetic lines.

94 Birds from each genetic line were routinely hatched every week. Different hatches were placed,
95 grown and reared in different sheds. Testing for water intake took place in pens equipped with
96 water stations. The test pen for a double station measured 14.8m x 6.1m, which corresponds to
97 a maximum of 52 kg/m² and 2.5 birds/m² at the end of rearing, for the heaviest line. The pen
98 was equipped with conventional group feeders hanging on feeding lines distributed throughout
99 the shed, and 16 electronic drinkers in double stations (8 drinkers in single stations), placed as
100 a line on one side of the pen. Prime quality wood shavings were used as litter. This resulted in
101 \approx 19 birds per drinker with mean drinker occupancy (i.e. percentage of time during which a
102 bird was using a particular drinker) during the experiment of 19.9%, 13.6% and 14.8% for lines
103 A, B and C, respectively. This allowed birds to drink without competition.

104 Birds were placed in the experimental shed one week before the recording started to allow them
105 to adapt to the drinking system. During this period lighting and temperature were maintained
106 in line with commercial husbandry practises, i.e., 14 hours of light at minimum 30 lux and 10
107 hours of dark and 19-23°C. Birds were fed a standard turkey grower diet. An in-house
108 developed electronic drinking system using transponder-based data capture was used to record
109 bird individual drinking behavior. Access to each drinker was regulated by a set of transparent
110 plastic dividers, which were adjusted to bird size as birds grew to ensure that only one bird
111 could use the drinker at a time. Each of the drinkers contained a water bowl connected to
112 weighing scales. Each bird was fitted on their lower leg with a small passive RFID transponder,
113 bearing a unique identification code which was recorded when a bird entered a drinker. The
114 RFID transponders enabled a drinking event (visit) to be ascribed to a specific bird. Extensive
115 bird observations suggested that the leg transponder did not affect bird normal behaviour
116 (Howie et. al., 2010). A visit started when the system detected bird presence in the drinker and
117 finished when the bird left the drinker. Each visit was recorded only when water consumption
118 occurred. The automated system recorded start and end time of each visit, visit duration, water

119 intake per visit and bird identification (ID). Start and stop times were recorded to the nearest
120 second, and water consumed was recorded to the nearest ml. Also recorded for each visit were
121 the date and the identification codes for the hatch, pen and drinker. Intervals between visits to
122 the water station were estimated as intervals between subsequent visits by the same bird.

123 *Video Observations*

124 Video observations were set up to (1) validate data recorded by the electronic drinkers, and (2)
125 record feeding events associated with drinking episodes to determine any associations between
126 feed and water intake. Four colour CCTV cameras (Hikvision DS-2CD2132-I) were positioned
127 approximately three meters above the pen area, capturing both electronic drinkers and feeders
128 described above. The clocks on the cameras and the computer collecting data from the
129 electronic drinkers were synchronized at the start of the experiment. Continuous video
130 recording was conducted between 06:00 - 20:00 when lights were on, during which times the
131 majority of all visits occurred (91%, 89% and 98% for lines A, B and C, respectively). Ten
132 birds were randomly selected and colour-marked with black spray in different shapes in order
133 to make them individually identifiable on the video. The other birds in the pen were sprayed
134 with food colouring using non-specific patterns, to avoid too much attention from the rest of
135 the group on the birds of interest.

136 The video analysis was conducted on five days of continuous video recordings. Both feeding
137 and drinking events were measured by frequency and duration. For each visit to the drinker or
138 feeder, the observer recorded bird ID and start and stop times of each visit. The drinking visit
139 was defined as "bird standing on the RFID tag reader (antenna pad) and ingesting water". The
140 start of a visit was when the bird stood with both feet on the tag reader or ingested water. The
141 end of the visit was when the bird left the drinker. Subsequently, visits recorded by the
142 automated system were compared with measures from the video analysis and recorded feeding
143 events were associated with the corresponding interval length between visits to the water

144 station. These data were used to estimate reliability of the electronic water station system by
145 calculating predictability (likelihood that a bird detected by the electronic water station is
146 detected as present at the drinker by video observations) and sensitivity (likelihood that a bird
147 present at the drinker is detected present by the electronic water system).

148 ***Electronic Data Screening***

149 Data screening involved several steps, including elimination of system errors, outlier detection
150 and data flooring. Any visits that were not correctly recorded by the system were removed from
151 the analysis. Visits were classified as outliers based on the water usage per visit to visit duration
152 ratio; any visits with a ratio above or below two standard deviations from the mean were not
153 included in the analysis. This allowed identifying long visits with low water usage and
154 similarly, short visits with abnormally high water records. As there were occasions when birds
155 remained in the drinker after drinking activity took place, such as resting or sleeping, data
156 flooring was performed on the remaining data to include only the visits during which birds
157 were drinking water. This involved limiting the maximum length of a visit to the longest visit
158 length observed during the video analysis. In total, this resulted in elimination of < 1% of visits
159 for lines A and C, and < 3% of visits for line B. The processed data set contained: line A – 948
160 045 visits, line B – 767 950 visits, line C – 144 109 visits.

161 ***Bout Analysis***

162 Because of the clear diurnal rhythm that birds showed, it was decided to use data recorded
163 during the hours when lights were on. We wanted to identify drinking bout criteria consistent
164 with our hypothesis for satiety underlying drinking behavior. A drinking bout criterion was
165 defined as the shortest interval between visits to the drinker that was considered to be part of a
166 bout and was estimated by fitting a mixture model (MM) to the natural log-transformed
167 intervals between visits (Celeux, 2007). According to the estimated criterion, intervals between
168 drinking events could be assigned to either within bout intervals or between bout intervals.

169 Bouts were characterised by duration and frequency, and were defined as time intervals spent
170 in drinking activity. In MMs data records are modelled as separate subpopulations with the
171 overall population being a mixture of the latter, resulting in a model with a finite number of
172 subpopulations. Hence, the first step was to find the number of subpopulations in the data.
173 During video observation it was identified that bird visits to drinkers with short interval length
174 between visits were the result of system oversensitivity to bird movements inside the drinker,
175 which resulted in some visits being fragmented. To reduce this error, an appropriate merging
176 criterion was necessary. This was selected using a Receiver Operating Characteristic (ROC)
177 curve (Hanley and McNeil, 1982), which was used to estimate and visualise true positive and
178 false positive rates for different merging criteria using data from the electronic system and
179 video observations. In total, 60 merging criteria were tested, ranging from 1 to 60 seconds. For
180 each merging criterion, a number of visually observed intervals between drinking episodes was
181 compared with RFID records. If an interval was detected by both methods, it was considered
182 as true positive, while intervals detected only by the RFID system were considered as false
183 positives. Plotted merged visit data showed a clear bimodal distribution, suggesting an initial
184 number of groups. Based on this finding, various combinations of two and three process
185 models, including Gamma, Exponential, Weibull and Normal, were investigated to determine
186 which of them can give both biologically and statistically good description of the data. It was
187 found that a model comprising a truncated log-normal for within bout intervals and a log-
188 normal distribution for between bout intervals gave consistent results for all genetic lines
189 (Equation 1).

$$190 \text{ pdf} = p \left(\frac{1}{\phi \left(\frac{1.39 - \mu_1}{\sigma_1} \right)} \right) \left(\frac{1}{\sigma_1 \sqrt{2\pi}} \right) \exp \left(\frac{-[x - \mu_1]^2}{2\sigma_1^2} \right) + (1 - p) \left(\frac{1}{\sigma_2 \sqrt{2\pi}} \right) \exp \left(\frac{-[x - \mu_2]^2}{2\sigma_2^2} \right) \quad (1)$$

191 where pdf is a probability density function for a Normal mixture model, μ_1, μ_2 and σ_1, σ_2 are
192 means and standard deviation of the truncated log-normal and log-normal distributions, p is the

193 proportion of intervals in the first distribution, ϕ is a correction factor for a truncated
194 distribution, and x is natural log-transformed interval length between visits.

195 The model was fitted for each genetic line and a bout criterion was estimated at the intersection
196 point between the two distributions. Based on the estimated bout criterion, drinking behavior
197 traits were estimated for each line. All estimated drinking behavior traits were tested for
198 statistical significance in statistical software R (R Dev. Core Team, 2014) using a non-
199 parametric Kruskal-Wallis test for the three lines of birds with post-hoc analysis using the
200 package `pgirmess` as proposed by Siegel & Castellan (1988).

201 To investigate whether drinking visits were distributed randomly across time or were guided
202 by physiological principles, such as thirst and satiety, the starting probability (P_{start}) of a bird
203 starting to drink within 30 minutes versus the time since the last visit (t) was calculated for
204 each line. This criterion was chosen to reduce variation in estimated probability, which
205 happened at smaller values. The probabilities were calculated from the data in the following
206 way: number of intervals $> t$ and $\leq t+30$ minutes divided by the number of intervals $\geq t$ minutes.
207 In addition, an empirical cumulative probability distribution function (ecdf) was computed as
208 another method that can be used to compare probabilities of starting the next visit to a drinker
209 within n seconds or less from the last visit. Opposite to the previous approach, this method
210 calculates cumulative probability and can be used to indicate which intervals between drinking
211 episodes are most likely to occur.

212 *Patterns of Drinking Accumulation Time*

213 We tested whether birds accumulate their drinking activity throughout the day by having time-
214 related traits with same length intervals or a combination of different length intervals. We
215 would expect birds to have variation in the interval length, however the extent of this variation
216 is uncertain both within and between the lines. To test for this expectation we used Lorenz
217 curves (Gastwirth, 1972), which were obtained by plotting the cumulative contribution of the

218 different interval lengths to the total time for four time-related drinking behavior traits: 1)
219 distribution of intervals between visits, 2) bout duration, 3) drinking time within a bout and 4)
220 non-drinking time within a bout. The approach was used to compare patterns of time
221 accumulation between the three genetic lines, which was done by comparing time intervals
222 from the shortest to the longest for each trait expressed in percentages from the overall time
223 length. The advantage of the method is that Lorenz curves can be expressed as a single
224 parameter called the GINI (G) index, which is a standard statistic for comparing patterns of
225 accumulation using this approach. A G index ranges from 0 to 1, where $G = 1$ would indicate
226 that distribution of time intervals is highly unequal in length, with large differences between
227 shortest and longest intervals for a given trait and relatively high proportion of short intervals
228 contributing to the total time. $G = 0$ would indicate that all intervals have similar length and
229 contribute equally to the total time. The higher the G index, the larger the inequality in the
230 distribution of interval length would be. G indices were estimated for each bird within each
231 line for the three genetic lines and tested using Kruskal-Wallis test to estimate variation in the
232 accumulation patterns between the lines with post-hoc analysis using the package `pgirmess` as
233 proposed by Siegel & Castellan (1988).

234 *Evolution of Drinking Behavior*

235 Estimated drinking behavior traits were used to investigate the evolution of drinking behavior
236 during the experiment and the effect of genetic line on bird drinking behavior. One of the
237 objectives was to determine whether birds from different lines have similar underlying
238 structural changes to drinking behavior over time. Principal Component Analysis (PCA)
239 (Jolliffe, 2002) was used to examine inter-relationships of drinking traits within and between
240 genetic lines using R software (R Dev. Core Team, 2014). Separate PCA was performed on
241 overall means of the experiment for each line using the correlation matrix of sample data of the
242 seven drinking behavior traits defined above: five daily traits: 1) bout frequency, 2) bout

243 duration, 3) drinking time, 4) non-drinking time, 5) water usage, and two traits estimated per
244 bout: 6) drinking time per bout, 7) water usage per bout. Principal components that accounted
245 for at least 95% of variability were kept in the analysis. To facilitate comparison of changes in
246 drinking behavior between days of the experiment, estimated principal component loadings
247 from the overall means were multiplied by standardised data computed for each day of the
248 experiment in order to convert daily scores to the same scale. As the length of the experiment
249 differed for some hatches within genetic lines, the number of days for which the analysis was
250 performed was limited to the shortest duration for a hatch within each line (Table 1). First and
251 last days of the experiment were excluded from the analysis, as they did not contain data for
252 the full days.

253 To determine changes in behavior over time, Pearson correlation coefficients were estimated
254 between daily component scores for each component for each line relative to the start of the
255 experiment. To visualise data, correlations computed between daily component scores were
256 added together for each day and compared between days for each line using the mixed-effect
257 model (McCulloch and Neuhaus, 2001), with observation day as a fixed effect and animal as a
258 random effect. Additionally, drinking behavior traits were estimated for each day of the
259 experiment for each line. All comparisons between the genetic lines were made using the linear
260 mixed-effect model, with observation day and genetic line as fixed effect, and animal as a
261 random effect. Subsequently, Tukey post hoc test was used for pairwise comparison of genetic
262 lines.

263 **RESULTS**

264 *Video Observations*

265 During the five days of video observations the electronic system recorded 50% more visits to
266 the water station compared to the manual observations. However, 47.4% of the visits recorded
267 by the automated system occurred within the visit time detected by direct video observations,

268 implying that either the video observations did not distinguish visits with small interval length
269 between visits or more likely the electronic system was giving false breaks within visits. Closer
270 inspection of these intervals revealed that they occurred when a bird was inside the drinker and
271 were the result of bird movement, which included leg, head and body movements while being
272 present inside the drinker. Because of this, birds showed non-uniformity concerning the
273 definition of a drinking visit, as for some birds the system detected each water sip as a separate
274 visit, resulting in a visit being fragmented into many short visits with small between visit
275 intervals, while other birds were more consistent in their drinking behavior. This resulted in
276 different distributions of visit durations and interval length between visits for the electronic and
277 observed data (Figure 1 a, b).

278 Based on data from the video validation, it was decided to combine such intervals together to
279 reduce the error in the analysis. Figure 2 shows the ROC curve with true positive and false
280 positive rates for each tested merging criterion. The merging criterion was chosen at three
281 seconds as the false positive rate at this criterion was zero. When this criterion was applied to
282 the video data set, it resulted in a 58.4% merging rate and a significant reduction of non-
283 uniformity in the data set.

284 The overall reliability of the electronic system was estimated from video observations and
285 resulted in a predictability of 98.8% and sensitivity of 98.6%. In general, the system did not
286 correctly record visits due to bird-drinker interactions: this included situations when the ID tag
287 was not placed correctly on the tag reader, while the bird being inside the drinker, or when two
288 birds were in the drinker at the same time, resulting in two IDs being ascribed to a single visit.

289 ***Bout Analysis***

290 Table 2 shows daily recorded measures made by the electronic drinkers and the estimated water
291 usage rate. All estimates were statistically different between the lines ($p < 0.01$), apart from the
292 daily water usage between lines A and B and mean visit duration between lines B and C ($p >$

293 0.05). Figure 3 shows histograms of the interval length between visits on a logarithmic scale
294 for hours of the day when light was present with the fitted MM for each genetic line. A two-
295 process model gave statistically consistent estimates between the lines. All three genetic lines
296 demonstrated two distinct modes in data distribution with different proportions of interval
297 length between the modes. The fitted probability MM contained one truncated log-normal
298 distribution at four seconds for within bout intervals, as shorter intervals were considered to be
299 the result of bird movement inside the electronic drinker. The second distribution was log-
300 normal for between bout intervals.

301 Drinking behavior traits were estimated using a bout criterion (Table 3). The bout criterion was
302 estimated at the intersection point between the two distributions and resulted in 665 s for line
303 A, 672 s for line B and 602 s for line C. Number of visits per bout varied between 1.11 and
304 1.70, for lines A and C respectively. Water usage per bout, drinking time per bout and number
305 of visits per bout were significantly different between the three lines ($p < 0.01$), while bout
306 duration was not significantly different between lines A and C, and non-drinking time per bout
307 and bout frequency were not significant between lines B and C ($p > 0.05$). According to the
308 video observations, 99% of feeding events occurred between rather than within drinking bouts.
309 The calculated P_{start} of a next visit to the drinking station within 30 minutes after the last visit
310 shows that birds from line A initiated their next visit after a shorter period of time, compared
311 to the two other lines, as their probability curve grew faster than for two other lines (Figure 4
312 a). All three lines demonstrated an initial rapid decrease in P_{start} up to approximately 10
313 minutes, as these intervals were associated with drinking occurring within a bout. Afterwards,
314 P_{start} started to increase over time reaching a marked peak after 1 - 2 hours since the last visit
315 for birds from line A. For lines B and C the peak was not clearly defined, suggesting that birds
316 from these two lines showed more variation in drinking behavior. Cumulative distribution of
317 the interval length between visits (ecdf) showed that birds from line C had a much higher

318 probability of starting the next visit shortly after the previous visit (Figure 4 b). However the
319 probability curves for all three lines converged at around 3 hours, when the majority of visits
320 occurred, constituting 95.7%, 93.5% and 96.8% of the total number of visits for lines A, B and
321 C, respectively. This indicates both that birds usually do not spend more than 3 hours between
322 drinking events and that variation in drinking behavior between the lines is accounted for by
323 short between drinking intervals.

324 *Patterns of Drinking Accumulation Time*

325 Figure 5 presents the Lorenz curves plotted for each genetic line estimated for four drinking
326 behavior traits during the hours of the day when light was present. The percentage of total time
327 spent in drinking visits shows that the distribution of interval length between drinking visits
328 differs between the genetic lines (Figure 5 a). Fifty percent of the shortest intervals between
329 visits contributed 70% to the total between interval time for birds of line A, whereas for line B
330 and line C they contributed 80% and 90%, respectively. Similarly, distribution of non-drinking
331 time is highly unequal in its accumulation pattern, with a G index close to one, implying high
332 prevalence of short non-drinking intervals within a bout (Figure 5 d). In contrast, accumulation
333 patterns of drinking time and bout duration show greater equality in the contribution of interval
334 length to the accumulation pattern (Figure 5 b, c).

335 Calculated G indices per individual bird for the drinking behavior characteristics are
336 significantly different between the lines ($p < 0.01$), suggesting that there is variation in the
337 distribution of birds with different time accumulation preferences (Table 4). While birds from
338 lines B and C are more similar in the accumulation time over the four characteristics, birds
339 from line A have a more equal distribution across four characteristics, meaning less variation
340 in drinking behavior for this line.

341 *Evolution of Drinking Behavior*

342 The analysis was carried using four principal components, as they accounted for 95%
343 variability in the data. Bird drinking behavior evolved during the experiment; however this
344 change was similar between the lines (Figure 6), as all three lines showed a downward shift in
345 the estimated correlations between the daily scores of the four principle components identified
346 by the PCA. Performing statistical tests on added daily correlation coefficients resulted in
347 significant differences in bird behavior relative to the start of the experiment (line A: $t = -4.242$,
348 $p < 0.01$, line B: $t = -5.357$, $p < 0.001$, line C: $t = -2.592$, $p < 0.05$). Similarly, there was a
349 difference in the absolute correlations between lines A and C ($z = 10.48$, $p < 0.01$), and between
350 B and C ($z = 12.51$, $p < 0.01$) throughout the experiment. However, no significant difference
351 was found between lines A and B ($z = -2.03$, $p > 0.1$).

352 Distribution of drinking behavior characteristics over the experiment for the three lines showed
353 that bird daily bout frequency, bout duration and drinking time per bout decreased over the
354 experiment, while daily water usage and water usage per bout increased (Figure 7).
355 Comparisons of drinking behavior traits using the mixed-effect model showed that all traits
356 were significantly different between the lines ($p < 0.001$), except for mean bout frequency
357 between lines A and B ($z = -1.784$, $p > 0.1$), mean daily non-drinking time per bout between
358 lines B and C ($z = -1.098$, $p > 0.1$) and mean water usage per bout between lines A and B ($z =$
359 -0.818 , $p > 0.1$), A and C ($z = 0.275$, $p > 0.1$), B and C ($z = 1.093$, $p > 0.1$). This suggests that
360 differences in drinking behavior between the lines exist, but these differences are not
361 consistent.

362 Analysis of inter-relationships between the drinking behavior traits showed that daily bout
363 duration had similar loadings across principal component 1 for each line (Figure 8). Other
364 relationships between drinking behavior traits differed between the lines: lines A and B had a
365 more similar relationship between the traits, compared to line C, with similar loadings for daily
366 bout frequency, daily nondrinking time within a bout, daily drinking time and mean water

367 usage per bout across principal components. All three lines showed opposite loadings for bout
368 frequency and water intake per bout over PC1, implying a negative correlation between these
369 two traits, while bout frequency and daily water intake showed no significant correlation,
370 indicating that birds use different strategies to attain the same amount of water.

371 **DISCUSSION**

372 We utilised a novel system that enables the recording of the drinking behavior of turkeys kept
373 in large groups, to develop a modelling methodology for the analysis and interpretation of their
374 drinking behavior. The system allows us to understand the basis of drinking behavior and its
375 regulation in birds selected for different productive traits. Contrary to the measurement and
376 analysis of feeding behaviour (Kyriazakis and Tolcamp, 2011), the analysis of drinking
377 behavior has received significantly less attention, due in part to past limitations of measuring
378 drinking behavior in a social context. The methodology for the analysis and interpretation of
379 drinking behavior was developed on turkeys, but the ideas advanced should have implications
380 for the drinking behavior in other animal species.

381 The study was conducted in a commercial setting, which allowed us to have access to large
382 bird numbers from different genetic lines. The employed recording system has advantages over
383 previous approaches used to record water intake (Maselyne et. al., 2015a), as it allowed to
384 detect drinking events of individual birds continuously and on a large-scale in a group-based
385 environment. This allowed us to extend the analysis beyond simple estimates of water intake,
386 and additionally focus on traits closely related to drinking activity. The system measured total
387 water removed from the drinker which would include both water consumed as well as water
388 spilt (Manning et. al., 2007b). As it was not possible to discriminate between the two, irregular
389 water spillage was monitored by farm staff.

390 Other potential limitations of the system are the actual drinker set up and its position in relation
391 to the feeders. As the water station was designed to record individual drinking behavior and

392 avoid cross-readings, once a bird was inside a drinker, it was separated by side plates from the
393 rest of the flock. In addition, the feeders were located at some distance from the drinkers and
394 this arrangement most likely had an effect on the natural sequence of drinking and feeding
395 events. For example, (Bley and Bessei, 2008) found in a study on individual feeding behavior
396 of group-housed pekin ducks that electronic systems influence the distribution of feeding
397 events in time, as once a bird was inside an electronic feeder/drinker, it was less likely that it
398 would alternate between feeders and drinkers.

399 A first step of the analysis involved the validation of the system used for recording drinking
400 behavior. This revealed system oversensitivity when registering visits due to bird movement,
401 thus fragmenting some of the visits. We introduced a criterion, based on statistical
402 methodology, according to which visits to drinkers could be classified into fragmented visits.
403 In general, data correction from automated recording systems is relatively common, but the
404 methodology used for this purpose varies between studies due to different settings of the
405 recording systems and is frequently based on arbitrary criteria (Brown-Brandl and Eigenberg,
406 2011; Casey et. al., 2005; Maselyne et. al., 2015b; Mendes et. al., 2001). Electronic system
407 reliability showed high agreement with data obtained from the visual observations, with high
408 scores for predictability (98.8%) and sensitivity (98.6%). We conclude that the system used in
409 our study is suitable for recording animal drinking behavior. As in previous studies, most of
410 the actual errors in our study occurred due to bird-drinker interactions, such as multiple birds
411 being in one drinker at the same time. It has been shown that bird density influences system
412 sensitivity and accuracy measures. While the number of such incidences in our study was small,
413 a study on drinking behavior of group-housed pigs (Andersen et. al., 2014) showed that
414 increased competition results in higher number of interrupted visits. This implies that in large
415 scale studies conducted in commercial settings animal density should be taken into account to

416 ensure both high performance of the recording equipment and sufficient access to the system
417 for the animal.

418 Following system validation, four methods were applied to characterise bird drinking behavior.
419 Firstly, a mixture model was used to classify interval length between drinking events into
420 within and between bout intervals. Secondly, conditional and cumulative probability functions
421 were calculated to identify if drinking behavior is random in time or has a physiological basis.
422 Unlike feeding behavior, which has a physiological basis (Howie et. al., 2009b; Tolkamp et.
423 al., 1998), water use may arise from other sources (Howard, 1975; Manning et. al., 2007a). The
424 modelling method allowed identification of novel traits associated with drinking behavior,
425 which could not be calculated without an appropriate bout criterion, and to compare these traits
426 between the different genetic lines. Thirdly, time accumulation patterns were used to
427 characterise bird preferences towards allocating their time to a particular drinking activity.
428 Lastly, evolution of drinking behavior between the genetic lines was examined.

429 Distribution of interval length between visits showed a well-defined separation in the interval
430 length between visits to drinkers for the three lines, and appeared to follow a similar pattern to
431 the distribution of visits to a feeder observed in birds (Howie et. al., 2009b; Howie et. al., 2010).
432 There was a population of short intervals which was considered to be intervals within a drinking
433 bout and a population of intervals which was considered to represent intervals between drinking
434 bouts. We applied several statistical distributions to model intervals between drinking visits
435 and we investigated which combination of distributions could provide an appropriate
436 description of data. Since we had left-truncated data, we fitted a truncated log-Normal
437 distribution to describe intervals within a drinking bout. For the second distribution, we
438 attempted to fit distributions such as Normal, Gamma and Weibull that have been previously
439 applied to similar types of data (Lundy et. al., 2012; Yeates et. al., 2001). In particular, we were
440 interested in the fit of the Weibull distribution, as it can apply on skewed data and appears to

441 have some properties consistent with the concept of satiety (Yeates et. al., 2001). However, the
442 application of the alternative functions to Normal distribution did not improve the description
443 of the populations of longer intervals. The above mixture model enabled us to identify a bout
444 criterion, which resulted in similar estimates of a bout criterion between the lines, i.e. a
445 truncated normal and normal distributions. We consider this to be an important outcome of our
446 study. Despite having substantial amounts of “free” time birds spent only small proportion of
447 time in drinking behavior, which is in agreement with the concept of satiety.

448 Feeding was not considered to be within a drinking bout, but according to the analysis, occurred
449 between drinking bouts. This is in contrast with the previous studies on feeding behavior, where
450 drinking was considered to be within a feeding bout (Howie et. al., 2009a; Yeates et. al., 2001),
451 and a bout included both feeding and drinking. In general, this can be explained by the
452 difference in the length between feeding and drinking bout criteria. According to previous
453 studies, feeding bout criteria may be longer than drinking bout criteria (Huzzey et. al., 2005;
454 Tolkamp et. al., 2011), and as feeding in general is more frequent in time, it is more likely that
455 drinking will occur within feeding bouts. However, this could also have been influenced by our
456 system settings, as it took longer for birds to move between drinkers and feeders, thus limiting
457 the probability that drinking would be associated with feeding.

458 We hypothesised that drinking behavior is predicated on the physiological principles of satiety
459 (Fitzsimons, 1998; McKinley and Johnson, 2004). The separation of visits to the drinkers into
460 two populations enabled us to calculate the conditional probability of initiating a visit
461 immediately after the previous visit (P_{start}). P_{start} was initially low, but in general increased
462 over time for all turkey lines with gradual decline for longer intervals. As it has been argued
463 previously by Tolkamp and Kyriazakis (1999), physiological regulation predicts that the
464 probability of an animal initiating a behavior of a visit to a feeder or drinker to be low after the
465 completion of a previous one, but will increase over time. If drinking were not to follow

466 physiological regulations, Pstart would have a uniform flat line shape, whereas the determined
467 Pstart of drinking behavior was almost identical to the one for feeding behaviour (Howie et.
468 al., 2009b; R Dev. Core Team, 2014) suggesting that both feeding and drinking behavior have
469 a similar physiological basis, that of satiety.

470 The above methodology enabled us to identify a number of drinking behavior traits, such as
471 bout frequency, duration of a bout, number of visits per bout, drinking time per bout, non-
472 drinking time per bout and water usage per bout. Importantly, these traits were based on
473 biological principles and are likely be of potential value (Kyriazakis and Tolkamp, 2011), and
474 therefore, of potential interest to turkey breeders. For example, according to the European Food
475 Safety Authority (EFSA), currently there is an increased interest to identify risk factors
476 associated with the drinking behavior traits in poultry that could lead to increased health risks
477 (EFSA, 2010). In particular, traits related to the efficient water use by birds are associated with
478 decreased litter moisture and better gut health (Swalander et. al., 2013), while overall time
479 spent in drinking activity, bout duration and non-drinking time per bout may be indicators of
480 time spent in proximity to drinkers and be associated with the risk of health challenges
481 associated with wet litter, such as food pad dermatitis.

482 Drinking bouts consisted of a small number of visits (maximum of 1.70 visits for line C). Most
483 of the calculated traits were significantly different between the lines. Birds from line A
484 exhibited different behavior compared to the other two lines, which can partly be accounted
485 for by their younger age. However since we have two differences for line A, their age and
486 genetics, we could not make any reliable conclusions when comparing this line to the other
487 two. Differences in the traits, such as water usage per bout and drinking time per bout between
488 lines B and C suggested that there is a true difference in the drinking behavior between these
489 two lines, which were tested at similar ages. Birds from line B tended to have a higher drinking
490 rate in comparison to birds from line C, which also explains their shorter bout duration. In

491 contrast, non-drinking time per bout was shorter for line A, implying that birds from this line
492 at the age considered do not split their water intake into many visits. Another trait that differed
493 between the lines was the number of visits per bout, which was higher for line C compared to
494 lines A and B. The difference in this trait between the lines was mostly due to higher proportion
495 of short intervals between visits to the same drinker with a length of up to 13 seconds. Bird
496 densities in the pens were similar making it unlikely that these intervals resulted from
497 interrupted visits; the differences may be true behavioral differences between the lines.

498 Accumulation patterns of time-related traits were used to obtain information regarding drinking
499 activities throughout the day and investigate whether birds of different ages or selected for
500 different traits have similar organisation of drinking activity. Overall, the G indices were
501 statistically different between lines for all traits considered. Birds from line A had smaller
502 variation in estimated drinking behavior traits, compared to the two other lines, suggesting a
503 more equal, on average, organisation of drinking behavior within this line. Lines B and C which
504 consisted of same age birds were more similar in the distribution of drinking traits during the
505 day, with line C having the highest variation of the estimated traits.

506 Bird drinking behavior had a tendency to change with time for the estimated traits. In general,
507 birds showed a decrease in the number of bouts over time accompanied by an increase in the
508 amount of water used per bout. Similarly drinking time and duration per bout in general
509 decreased over time, the extent of which varied between the lines. This is consistent with the
510 time related trends seen in the feeding behavior of turkeys (Howie et. al., 2010). One interesting
511 finding is that daily bout frequency, daily total water usage and daily total drinking time were
512 uncorrelated or weakly correlated in all three lines. As estimates of daily bout frequency
513 differed between birds in each line, it suggested that birds have different strategies in meeting
514 their water requirement. These results are supported by a similar study on bird feeding
515 behavior, where it was found that birds vary in the distribution of bouts and intervals between

516 them (Howie et. al., 2009b), and as this was heritable it is perhaps useable. However, overall
517 change in drinking behavior was similar between the lines, suggesting that while difference
518 between the lines in the organisation of drinking exists, the more fundamental aspects of
519 behavior, such as regulation of drinking and drinking behavior evolution over time is conserved
520 between the lines. This means that birds have the flexibility to adapt their behavior without
521 compromising essential body functions and regulations.

522 In conclusion, we have developed a methodology that enables the analysis of drinking behavior
523 in birds. The methodology suggests that drinking behavior is underlined by the principles of
524 satiety, and for this reason may have applicability across different animal species. The
525 methodology also enabled us to identify a number of drinking behavior traits that arise from
526 biologically defined criteria of drinking bouts. There seemed to be differences in these traits
527 between bird genotypes selected for different productive traits. Therefore, these drinking
528 behavior traits may be of potential relevance to turkey breeders, as water utilisation, along with
529 feed efficiency, is part of overall biological efficiency. Furthermore, drinking behavior could
530 provide further insights on the link between water usage and health, and the environmental
531 impact of turkey production systems.

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714

715 **TABLES**

716 **Table 1.** General overview of the data available for drinking behaviour analysis with number
 717 of hatches available for each line. Duration of the drinking recording experiment and number
 718 of birds per hatch are presented using minimum, maximum and median values to show
 719 variation in these characteristics within and between the lines.

Genetic Lines	Number of hatches	Experiment duration (days)			Number of birds per hatch		
		Min	Max	Median	Min	Max	Median
A	29	14	16	15	131	292	146
B	15	23	28	24	135	224	142
C	2	22	24	23	141	144	143

720

721

722 **Table 2.** Medians with interquartile ranges of the number of visits, water usage per visit and
 723 visit duration, and total drinking time and total water usage estimated per day per individual
 724 bird for the three genetic lines of turkeys.

Recorded parameters	Line A	Line B	Line C
Number of visits	11.17 ^{a,b} (10, 13.7)	12.17 ^{a,c} (10.21, 14.42)	15.32 ^{b,c} (12, 20.45)
Water usage per visit (ml)	68.03 ^{a,b} (57.02, 80.39)	64.96 ^{a,c} (54.02, 77.67)	45.90 ^{b,c} (34.24, 56.71)
Mean visit duration (s)	77.95 ^{a,b} (66.37, 91.22)	51.82 ^a (42.5, 61.84)	48.21 ^b (34.47, 61.17)
Daily drinking time (s)	903.91 ^{a,b} (777.87, 1054.99)	623.39 ^{a,c} (542.96, 706.7)	699.68 ^{b,c} (600.86, 821.95)
Daily water usage (ml)	790.29 ^b (701, 890.45)	785.62 ^c (725.61, 847.30)	687.29 ^{b,c} (624.21, 760.81)
Water usage rate (ml/s)	0.88 ^{a,b} (0.75, 1.02)	1.26 ^{a,c} (1.11, 1.45)	0.99 ^{b,c} (0.84, 1.16)

725 ^a P < .01 between the lines A and B.

726 ^b P < .01 between the lines A and C.

727 ^c P < .01 between the lines B and C.

728

729 **Table 3.** Medians with interquartile ranges of drinking behavior traits per day per individual
 730 bird for three turkey genetic lines A, B and C, estimated after grouping visits to the water
 731 station into bouts.

Traits	Line A	Line B	Line C
Bout criterion (s)	665	672	602
N of bouts	10.64 ^{a,b} (9.25, 12.27)	10.22 ^a (8.83, 11.65)	9.64 ^b (8.19, 11.87)
N of visits per bout	1.08 ^{a,b} (1.05, 1.13)	1.17 ^{a,c} (1.11, 1.26)	1.49 ^{b,c} (1.26, 1.95)
Bout duration (s)	93.89 ^a (81.82, 107.84)	80.48 ^{a,c} (70.51, 92.68)	94.7 ^c (82.82, 106.97)
Drinking time per bout (s)	85.36 ^{a,b} (73.11, 99.17)	61.78 ^{a,c} (52.17, 72.32)	73.98 ^{b,c} (61.18, 85.74)
Non-drinking time per bout (s)	7.24 ^{a,b} (3.69, 12.25)	17.10 ^a (10.73, 24.79)	18.58 ^b (11.42, 28.23)
Water usage per bout (ml)	74.50 ^{a,b} (63.52, 86.91)	77.15 ^{a,c} (66.50, 89.86)	68.92 ^{b,c} (57.75, 85.57)

732 ^a P < .01 between the lines A and B.

733 ^b P < .01 between the lines A and C.

734 ^c P < .01 between the lines B and C.

735

736 **Table 4.** GINI indices estimated per line, and medians with interquartile ranges measured per
 737 individual bird within each line (A, B and C) for intervals between visits, bout duration, non-
 738 drinking time and drinking time per bout for individual birds.

Traits	Line A	Line B	Line C
	0.39	0.47	0.63
Intervals between visits	0.36 ^{a,b} (0.33,0.4)	0.44 ^{a,c} (0.4, 0.49)	0.56 ^{b,c} (0.46, 0.67)
	0.33	0.45	0.42
Bout duration	0.29 ^{a,b} (0.25,0.33)	0.41 ^{a,c} (0.36,0.47)	0.37 ^{b,c} (0.33, 0.44)
	0.97	0.94	0.88
Non-drinking time per bout	0.97 ^{a,b} (0.95,0.98)	0.94 ^{a,c} (0.91, 0.96)	0.88 ^{b,c} (0.82,0.93)
	0.28	0.33	0.37
Drinking time per bout	0.24 ^{a,b} (0.22, 0.26)	0.30 ^{a,c} (0.28, 0.33)	0.32 ^{b,c} (0.29, 0.36)

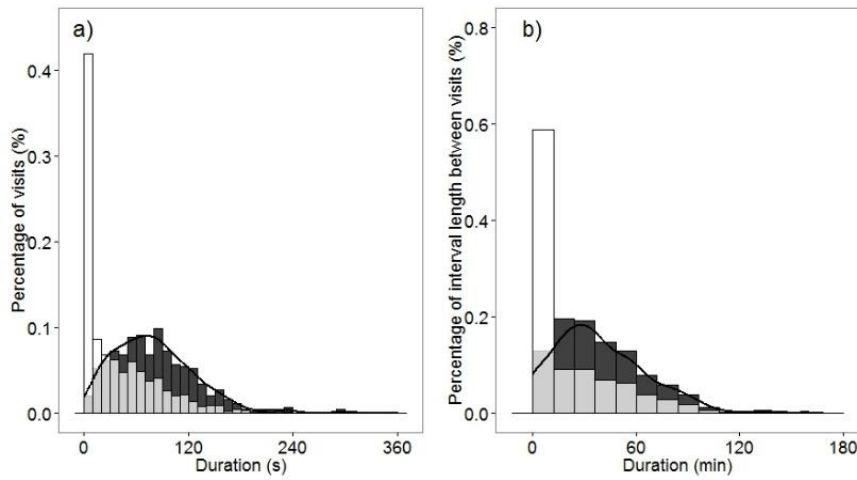
739 ^a P < .01 between the lines A and B.

740 ^b P < .01 between the lines A and C.

741 ^c P < .01 between the lines B and C.

742

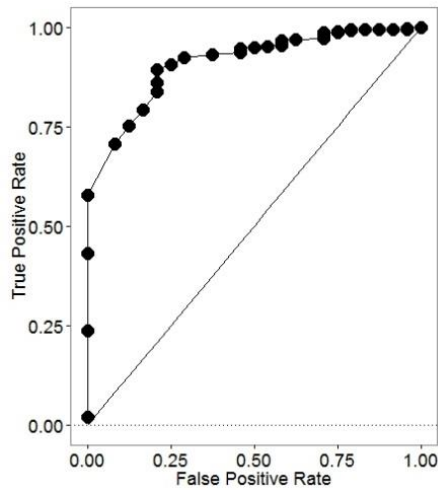
743 **FIGURES**



744

745 **Figure 1.** Data distribution following the validation step, where intervals less than 4 seconds
746 have been excluded. (a) Frequency of visit durations to the water station with a bin size of 10
747 seconds and (b) frequency of interval length between visits to the water station with a bin size
748 of 20 minutes, estimated by the electronic system (white bars) and manual observations (black
749 bars), with the overlaid density line for the manual observations estimated from the
750 corresponding histogram.

751



752

753 **Figure 2.** Receiver Operating Characteristics (ROC) curve showing true and false positive rates

754 for each merging criterion tested (0 to 60 seconds) with markers from left to right, estimated

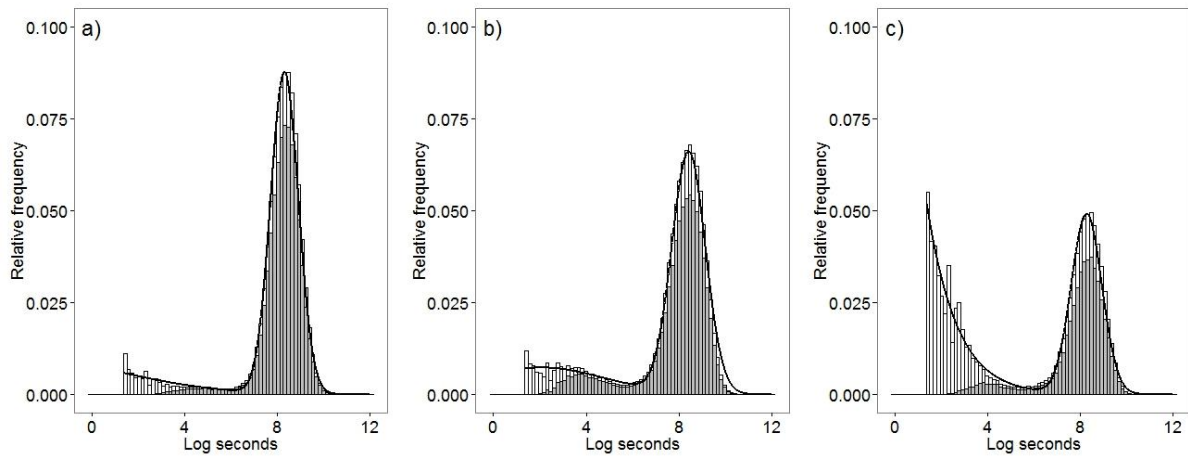
755 from automated (via radio-frequency identification) versus manually recorded visits. The true

756 positive rate measures the proportion of positives (visits recorded by the RFID system) that are

757 confirmed through direct video observations. The false positive shows the proportion of

758 positives detected by the RFID system that are rejected by video observations.

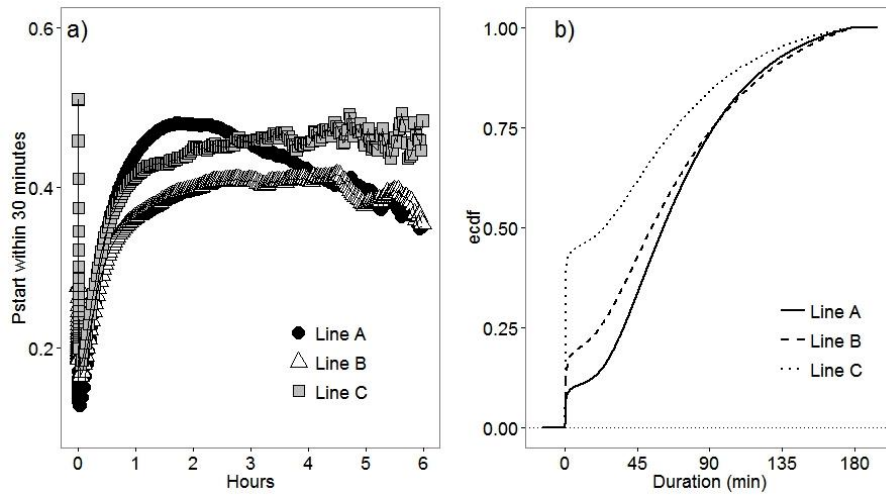
759



760

761 **Figure 3.** Frequency distribution of intervals between visits to the same drinker (white bars)
 762 and different drinkers (grey bars) whilst light was present in the turkey sheds (06:00-20:00).
 763 Intervals are expressed on a natural-log scale (bin size = 0.15 log units). The solid lines are
 764 fitted mixed models to the natural-log transformed interval length between visits; they contain
 765 a truncated log-normal distribution for within bout intervals and a log-normal distribution for
 766 between bout intervals for three turkey genetic lines A, B and C, graphs (a), (b) and (c)
 767 respectively.

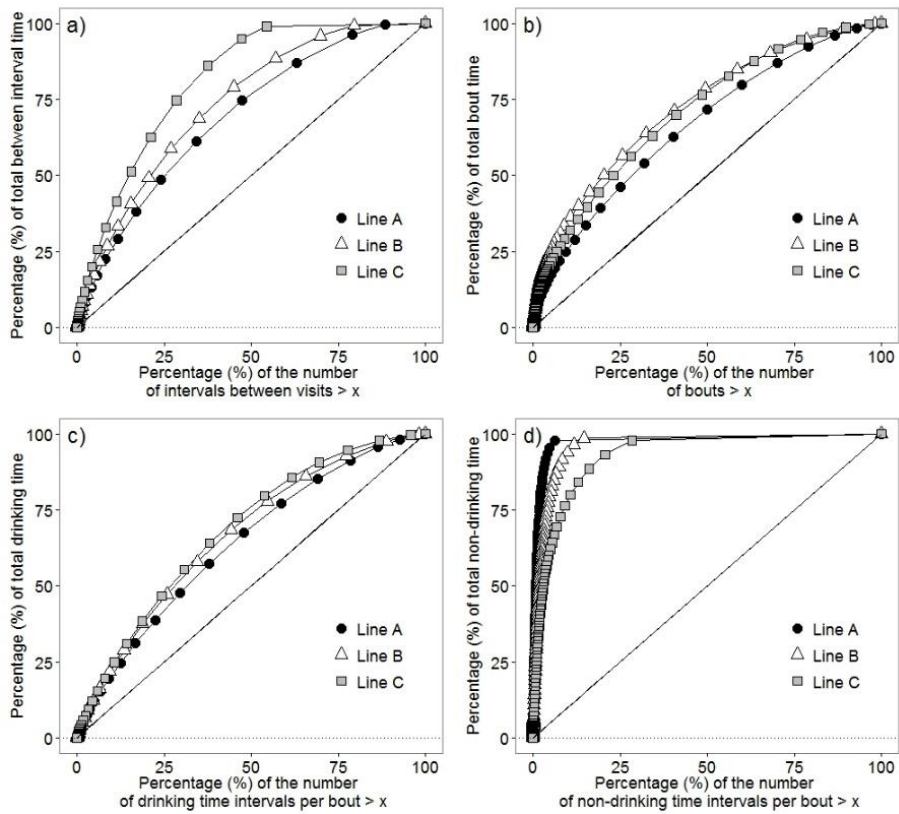
768



769

770 **Figure 4.** Probabilities of a next visit. (a) Probability of a bird starting a next visit to an
 771 electronic drinker (P_{start}) within the next 30 minutes since the last visit and (b) empirical
 772 cumulative probability distribution function (ecdf) of interval length between visits to the
 773 electronic drinker for three genetic lines of turkeys. Graph (b) shows the cumulative probability
 774 of a next visit within n seconds or less for the three genetic lines.

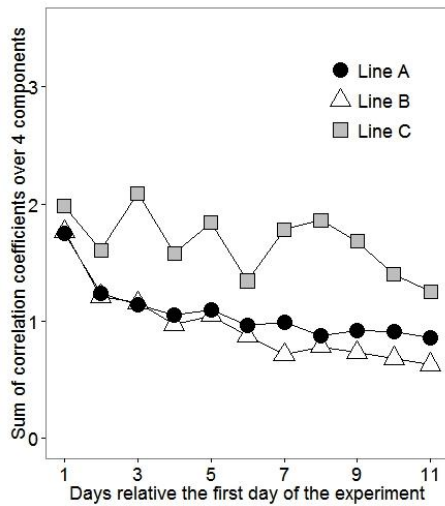
775



776

777 **Figure 5.** Accumulation patterns of four time-related drinking behavior traits for the turkey
 778 genetic lines A, B and C: (a) intervals between visits to the water station, (b) bout duration, (c)
 779 drinking time per bout and (d) non-drinking time per bout. The black line is an equality line
 780 (GINI = 0), where intervals of different length contribute equally to the accumulation pattern.
 781 Y-axis shows the cumulative percentage of total time for a given trait, while x-axis shows the
 782 cumulative percentage of unique interval length, where x = interval length for a given trait
 783 (ordered from the smallest to the largest).

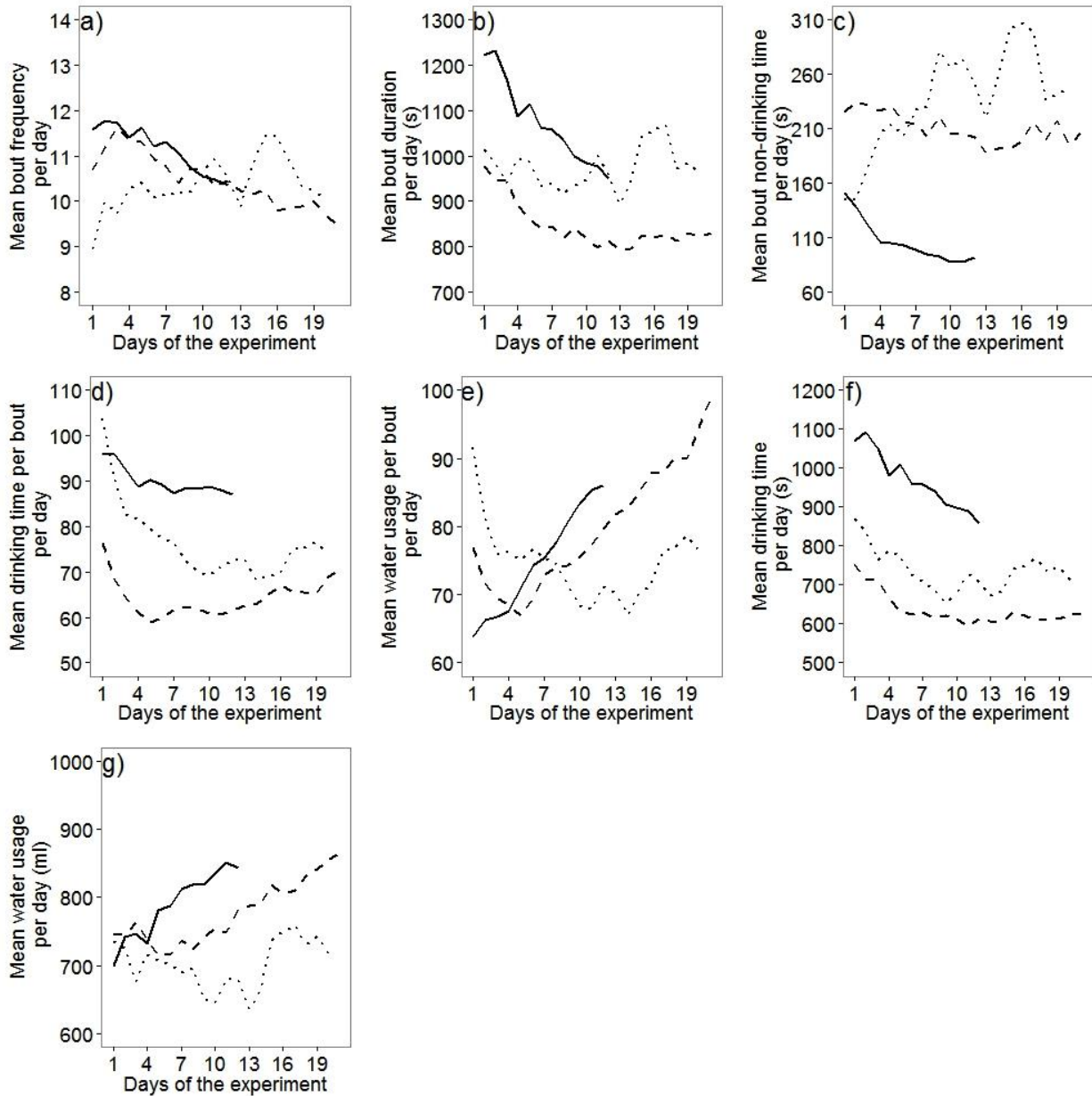
784



785

786 **Figure 6.** Pearson correlation coefficients relative to the first day of the experiment and added
 787 over four principal components for each day from Principal Component Analysis (PCA) that
 788 accounted for 95% of variability in the data, for lines A, B and C. The correlation coefficients
 789 were estimated for each principal component separately and added together for visualisation
 790 purposes.

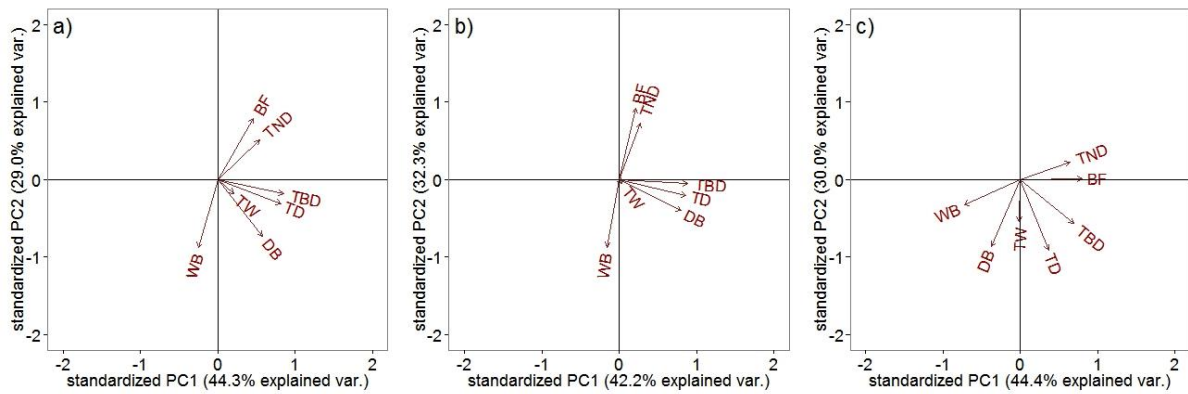
791



792

793 **Figure 7.** Distribution of bout and daily related drinking behavior characteristics during the
 794 experiment calculated per each day of the experiment for line A (solid line), line B (dashed
 795 line) and line C (dotted line): (a) mean bout frequency, (b) bout duration, (c) non-drinking time
 796 within a bout, (d) mean drinking time per bout, (e) mean water usage per bout, (f) drinking
 797 time, (g) water usage. The number of days was taken as a shortest experiment length for a given
 798 genetic line.

799



800

801 **Figure 8.** Principal component analysis showing inter-relationships between 7 drinking
 802 behavior traits over the first two principal components. Traits include: BF – daily bout
 803 frequency, TBD – daily bout duration, TD – daily drinking time, TND – daily nondrinking
 804 time within a bout, TW – daily water usage, DB – mean drinking time per bout, WB – mean
 805 water usage per bout, for genetic lines A (a), B (b) and C (c), respectively. The angle between
 806 the vectors representing traits indicates correlation between traits. For highly correlated traits
 807 vectors point in the same direction (or opposite directions for highly negatively correlated
 808 traits), whereas uncorrelated traits have a right angle between the vectors.