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## The effect of conspecific removal on behavioral and physiological responses of dairy cattle

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1 Running Head: SEPARATION EFFECTS IN DAIRY CATTLE

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4 **The effect of conspecific removal on behavioral and physiological responses of dairy**  
5 **cattle**

6

7 **Jessica K Walker\*†, David R Arney‡, Natalie K Waran§, Ian G Handel#, Clive J C**  
8 **Phillips\***

9

10 \*Centre for Animal Welfare and Ethics, School of Veterinary Sciences, University of  
11 Queensland, Australia.

12 †The Animal Welfare and Biodiversity Research Group, Department of Natural Sciences,  
13 Unitec, Private Bag 92025, Auckland 1025, New Zealand.

14 ‡Institute of Veterinary Medicine and Animal Sciences. Estonian University of Life Sciences.  
15 Kreutzwaldi 46, 51006 Tartu, Estonia.

16 §Jeanne Marchig International Centre for Animal Welfare Education. Royal (Dick) School of  
17 Veterinary Studies, the University of Edinburgh, Easter Bush, Roslin, EH25 9RG, UK.

18 #The Royal (Dick) School of Veterinary Studies, The University of Edinburgh, Easter Bush  
19 Campus, Midlothian EH25 9RG, UK.

20

21 Address for correspondence: Dr Jessica K. Walker, Department of Natural Sciences, Unitec,

22 Private Bag 92025, Auckland 1025, New Zealand. E-mail: [jwalker@unitec.ac.nz](mailto:jwalker@unitec.ac.nz)

23 **ABSTRACT**

24 Adverse social and welfare implications of mixing dairy cows or separating calves from their  
25 mothers have been documented previously. Here we investigated the behavioral and  
26 physiological responses of individuals remaining after conspecifics were removed. We  
27 conducted a series of four experiments incorporating a range of types of different dairy cattle  
28 groupings (Experiment 1 [E1], 126 outdoor lactating dairy cows; Experiment 2 [E2], 120  
29 housed lactating dairy cows; Experiment 3 [E3], 18 housed dairy calves, and Experiment 4  
30 [E4], 22 housed dairy bulls) from which a subset of individuals were permanently removed  
31 (E1 n = 7, E2 n = 5, E3 n = 9, E4 n = 18). Associations between individuals were  
32 established using near-neighbor scores (based upon identities and distances between animals  
33 recorded prior to removal) in E1, E2 and E3. Behavioral recordings were taken for 3 to 5 d,  
34 before and after removal on a sample of cattle in all 4 experiments (E1 n = 20, E2 n = 20, E3  
35 n = 9, E4 n = 4). In two experiments with relatively large groups of dairy cows, E1 and E2,  
36 the responses of cows that did and did not associate with the removed cows were compared.  
37 An increase in time that both non-associates and associates spent eating was observed after  
38 conspecific removal in E1. In E2 this increase was restricted to cows that had not associated  
39 with the removed cows. A reduction in ruminating in remaining cattle was observed in E3  
40 and eating in E4. Immunoglobulin A concentrations increased after separation in both E3 and  
41 E4 cattle, but did not differ significantly between associates and non-associates in E2. **Blood**  
42 **and milk cortisol** concentrations were not affected by conspecific removal. These findings  
43 suggest that some animals had affected feeding behavior and IgA concentrations after  
44 removal of conspecifics.

45

46 Key Words: association, dairy cattle, separation, immunoglobulin A, conspecific

47

**INTRODUCTION**

48 Increasingly, animal emotions form the basis of animal welfare definitions (Dawkins, 1990;  
49 Fraser and Duncan, 1998; Mendl and Paul, 2004; Broom, 2010), with public concern for the  
50 welfare of farm animals often arising from the recognition that animals are able to experience  
51 emotions (Špinka, 2012, Boissy and Erhard, 2014). Farm animals are gregarious and their  
52 social environment plays a fundamental role in the individual's welfare status (Keeling and  
53 Gonyou, 2001; Rault, 2012), with many benefits being derived from the presence of a  
54 conspecific (Rault, 2012). Dairy cattle form long lasting social bonds (Reinhardt and  
55 Reinhardt, 1981; Færevik et al., 2006) and show strong affiliation to conspecifics (Holm et  
56 al., 2002). In modern production systems the regrouping of cattle (regrouping is defined here  
57 as a two-step process: 1. Separation from the old group and 2. Introduction to a new group)  
58 occurs frequently in order to create homogenous groups organized by common  
59 characteristics, such as age, milk yield, body condition, reproduction, and health status (Bøe  
60 and Færevik, 2003; Raussi et al., 2005). This regrouping process, in particular step 2, has  
61 been documented to result in social stress evidenced by behavioral changes that include  
62 increased aggression (Raussi et al., 2005), vocalizations (Boissy and Le Neindre, 1997;  
63 Færevik et al., 2006; De Paula Vieira et al., 2010), changes in locomotory behavior  
64 (Hasegawa et al., 1997; von Keyserlingk et al., 2008) and has negative impacts upon  
65 production traits, such as reduced feed intake (von Keyserlingk et al., 2008; De Paula Vieira  
66 et al., 2010; Schirmann et al., 2011; Duve et al., 2012), milk yield (Hasegawa et al., 1997;  
67 von Keyserlingk et al., 2008) and weight gain (De Paula Vieira et al., 2010). These negative  
68 effects have been documented across a range of cattle scenarios including lactating cows,  
69 heifers and bulls (Mench et al., 1990; Hasegawa et al., 1997; Mounier et al., 2006). However,  
70 studies investigating the effect of repeated regrouping show contradictory findings, with  
71 some suggesting cattle do habituate to regrouping over time (Mench et al., 1990) and others

72 providing no evidence of this (Raussi et al., 2005). Conceivably the stability of relationships  
73 between the cattle and the number of animals affected by the regrouping may determine the  
74 ability of cattle to habituate to the practice.

75

76 The majority of studies investigating the regrouping of cattle have focused on the effect on  
77 the individual(s) being regrouped (e.g. Mench et al., 1990; Raussi et al., 2005; Mounier et al.,  
78 2006; von Keyserlingk et al., 2008). In these studies the effects of separation are often hard  
79 to distinguish from the effects of the novel environment (Rault, 2012). Although the impact  
80 of new individuals introduced to a previously established group has been described, the  
81 impact on the individual(s) remaining in the original group has not. In one study by  
82 Schirmann et al. (2011) the difference in response to regrouping between cows that were  
83 moved to a new pen and those that stayed in their home pen was investigated, however due to  
84 the experimental design, the effects of removal of individual cows on those remaining in the  
85 home pen could not be separated from the effects of the newly introduced cows.

86

87 Measurement of stress traditionally involves behavioral observation and physiological  
88 evaluation of, for example, HPA activation (e.g. cortisol) or immunological response (e.g.  
89 immunoglobulin A). Immunoglobulin A (IgA) represents a main element of the humoral  
90 immune response, which provides protection against pathogens at mucosal surfaces (Snoeck  
91 et al., 2006). In its secretory form (S-IgA) it serves to prevent infective agents such as  
92 bacteria and viruses from breaching the mucosal barrier, whilst within serum it functions as  
93 an inflammatory antibody acting on immune effector cells (Snoeck et al., 2006). Relatively  
94 little information is available on the relationship **between IgA and stress responses in farm**  
95 **animals**, with the exception that in pigs S-IgA reportedly increases as a result of chronic  
96 stress caused by social isolation during the first 12 days and declines thereafter (Royo et al.,

97 2005). A similar response has been observed in dogs in the first six days following separation  
98 from a conspecific (Walker et al., 2014) and as a result of stress experienced upon entry into  
99 a kennel environment (Skandakumar et al., 1995). In response to acute stress, S-IgA levels in  
100 rats and dogs have been documented to decrease (Guhad and Hau, 1996; Kikkawa et al.,  
101 2003), and in humans a large body of evidence concludes that negative emotional valence,  
102 resulting from short-term stress, results in decreased S-IgA (reviewed by Segerstrom and  
103 Miller, 2004). Although the influence of emotional states on IgA secretion in cattle has not  
104 been examined, bovine IgA has been quantified in milk (Newby and Bourne, 1977;  
105 Honkanen-Buzalski and Sandholm, 1981), serum, lacteal, saliva, nasal and vaginal secretions  
106 (Duncan et al., 1972). Research has demonstrated that IgA in bovine milk is predominately  
107 serum derived (Newby and Bourne, 1977), suggesting that milk could act as an appropriate,  
108 non-invasive, accessible alternative to serum in the measurement of short and long term  
109 stress. Likewise, cortisol concentrations in milk from cows in established lactation have been  
110 demonstrated to directly relate to cortisol concentrations in blood (Shutt and Fell, 1985),  
111 suggesting that milk is a suitable substitute for serum when measuring cortisol concentrations  
112 in dairy cattle.

113

114 The objective of this study was to investigate the effect of step one of regrouping: **The effect**  
115 **that the removal of individuals from the group** has on remaining group members, utilizing  
116 behavior observations and two physiological measures; cortisol and immunoglobulin A  
117 (IgA).

118

119

## MATERIALS AND METHODS

120 These experiments were approved by the University of Queensland Animal Ethic Committee,  
121 approval numbers CAWE139/10 and CAWE068/11.

122

123 ***Experiment 1.***

124 ***Animals.*** In Experiment 1 (**E1**), Observations were made of a herd of 126 lactating  
125 Holstein-Friesian and mixed breed dairy cows at the University of Queensland (Gatton,  
126 Queensland, Australia). The study was carried out during mid-winter (mean temperature =  
127  $16^{\circ}\text{C} \pm 4.4^{\circ}\text{C}$ ) when the herd was maintained in a 1.93 ha outdoor feedlot area (Figure 1),  
128 with a stocking density of 65.3 cow per ha. Of the total 126 cows, 55% (69/126) were  
129 Holstein-Friesian; 27% (34/126) Holstein-Friesian crossbreed; and one (0.8%) each of Jersey;  
130 Brown Swiss; Brown Swiss cross Jersey; Ayrshire cross and the remaining 15% (19/126)  
131 were of unknown crossbreed.

132

133 The group structure was dynamic with cows temporarily removed from the herd as a result of  
134 cessation of lactation, illness or estrus cycle, as well as for use during agriculture and  
135 veterinary teaching demonstrations and practicals. The cows were milked twice daily in a  
136 herringbone parlor between 0600 to 0800 h and 1500 to 1800 h. Feed was delivered twice  
137 daily at 0800 and 1300 h to a covered feeding trough in a paddock. The cows were  
138 maintained on a TMR consisting of 13% soybean meal, 38% grain mix, 26% barley silage,  
139 14% soybean silage, 6% lucerne silage and 3% mineral mix on a DM basis. The feed bunk  
140 was 60 m long with enough room for all cows to feed comfortably at the same time. Water  
141 was available *ad libitum*.

142

143 ***Near Neighbour Observations.*** A subset of 7 individuals were selected for removal  
144 and subsequent culling due to age-related reductions in milk production. Nearest neighbor  
145 identities and distances were recorded prior to this removal to establish the strength of  
146 association between these removed individuals and others in the group. The distance from



147 each of the 7 individual cows signaled for removal, to up to 5 nearest neighbors was  
148 estimated visually using cow length (mid-point between the shoulder and tail of the cow  
149 shoulder to rump (Gibbons et al., 2010) as a guide. Near-neighbor observations were carried  
150 out on 34 to 39 (SD = 1.7) occasions, for each of the 7 focal animals, over 9 d in three  
151 differing locations: the paddock, order of entry into the milking parlor and during feeding.  
152 Recordings in the paddock were conducted between 0900 to 1300 h (with at least 1 h between  
153 recording sessions) on 21 to 24 (SD = 1.1) occasions/cow over 9 d. Order of entry into the  
154 milking parlor recordings were carried out on 3 to 5 (SD = 0.9) separate occasions per cow at  
155 the p.m. milking, with the cow on either side of each focal cow recorded. Cows at the end of  
156 a milking row had only one recorded neighbor. Feeding recordings followed the methodology  
157 described by Cooper et al. (2008), with cows each observed on 8 to 10 (SD = 0.8) occasions  
158 across 9 d between 1300 to 1500 h, following the delivery of the afternoon feed ration. The  
159 two nearest-neighbors on either side of each focal cow were recorded and considered feeding  
160 partners, providing they were within a distance of 1 cow length. If a cow was at the end of the  
161 feeding line she was considered to only have one feeding partner. Observations were carried  
162 out in these three locations as these were the only locations that the cows had access at set  
163 times across a 24 h day.

164

165 ***Selection of Experimental Subjects.*** To allocate subjects to an associate or a non-  
166 associate group we used the near-neighbor recordings to identify the 10 individuals that  
167 displayed the greatest (associate), and the 10 individuals that displayed the weakest (non-  
168 associate), association to the 7 individuals identified for removal. Probability theorem was  
169 used to establish that these interactions did not occur by chance (see statistical analysis  
170 section). The number ( $n = 10$ ) of individuals allocated to both the associate and non-associate  
171 groups was determined from the natural social structure of cattle which is reported to be a

172 mean group size of 10-11 individuals (Bouissou et al., 2001). The 20 selected cows had mean  
173 age of  $6.1 \pm 1.8$  (SD) yr and weight of  $577 \pm 144.5$  (SD) kg. The mean milk yield per  
174 experimental cow was  $27.1 \pm 7.9$  (SD) L per day during the course of the experiment.  
175 Stocking density after removal was 60 cows per ha.

176

177 **Behavioral Observations.** An ethogram (Table 1) was developed based on previous  
178 studies (Krohn, 1994; Fröberg and Lidfors, 2009; Fogsgaard et al., 2012). Behavioral  
179 observations were carried out by three trained observers, two of whom were recording at any  
180 one time. Observations were made continuously using focal animal sampling for each of the  
181 20 cows, at 1h intervals, for a 10 min per cow duration. Cows were recorded in the same  
182 order each time. Observations were carried out continuously for 8 d between morning and  
183 afternoon milking from 0800 to 1500 h (70 min per day/per cow), 4 d being before removal  
184 of designated cows (d -5, -3, -2, -1) and the remaining 4 d after removal (d +1, +2, +3, +4).  
185 Removal took place at 1900 h, therefore the first observations after separation began after  
186 13h post separation.

187

## 188 **Experiment 2**

189 **Animals.** Experiment 2 (**E2**) was carried out at the Estonian University of Life  
190 Sciences Maarja Farm, Tartu, Estonia. Observations were made of 2 herds each containing  
191 60 lactating Holstein-Friesians, each with a stocking density of 0.12 cows per m<sup>2</sup>. Cows in  
192 both herds were free-stall housed within the same building and fed cut grass in their housing  
193 (Figure 1). The total housing area for both herds was 1032 m<sup>2</sup>. The group structure was  
194 dynamic with cows temporarily removed for cessation of lactation, illness, or estrus. Milking  
195 occurred twice daily in either a traditional parlor (n = 8) at 0530 and 1500 h or twice daily in  
196 a robotic milking system (n = 12). The two differing milking systems were engaged for

197 teaching demonstrations and research purposes. Cows had *ad libitum* access to water and  
198 were fed a TMR consisting of grass/clover silage 50.8% DM, barley 24.6% DM, wheat 7.8%  
199 DM, rapeseed cake 15% DM and mineral feed 1.8% DM. They were fed twice daily at 1000  
200 and 1700 h. The parlor-milked cows had individual access to bins in their housing (Figure 1)  
201 with varying amounts of added concentrate in TMR. Robot-milked cows received an  
202 additional 0.88-2.64 kg DM/d of concentrate supplement (Baltic Argo AS, Tallinn, Estonia)  
203 from an automated feeder and 2.64-5.28 kg DM/d in the robot, according to yield.

204

205 ***Near Neighbor Observations.*** A subset of 5 individuals (parlor milked herd n = 2;  
206 robot milked herd n = 3), with a mean age of  $3.8 \pm 0.84$  (SD) yr, were identified for removal  
207 due to imminent dry-off and forthcoming parturition. Stocking density after removal in E2  
208 was 0.11 cow per m<sup>2</sup> for both the parlor milked and robot milked herds. An additional subset  
209 of 5 individuals (parlor milked herd n = 2; robot milked herd n = 3), with a mean age of  $4.6 \pm$   
210  $1.1$  (SD) yr, were randomly selected to aid establishment of a control group. Nearest neighbor  
211 distances were determined for all 10 cows prior to removal of the subset of 5 individuals  
212 mentioned above. The near neighbor observations were carried out in loose housing area  
213 which included the feed bunk (Figure 1) utilizing the methodology described in E1, with the  
214 exception that to be considered a near-neighbor the cow had to be within 1 cow length of the  
215 focal animal (rather than three) due to the more intensive housing conditions. Observations  
216 were not carried out separately during milking or at the feed bunk, as was done in E1, due to  
217 the use of the robot milking system and the inclusion of the feed bunk in the loose housing  
218 area. Near-neighbor observations were carried out on 12 occasions at 1300; 1900; 2400 and  
219 0500 h across three consecutive days. These occasions represented even distribution across  
220 each 24 h period whilst also incorporating a 2 h period after milking (parlor side) and feeding  
221 before the start of an observation session.

222

223 ***Selection of Experimental Subjects.*** A subgroup of 20 individuals (parlor n = 8; robot  
224 milkers n = 12) were selected as outlined below and divided into two groups of 10. Selection  
225 utilized near-neighbor recordings to first identify the 10 individuals that displayed the  
226 strongest association to the 5 individuals signaled for subsequent removal. These 10  
227 individuals formed the associate group (mean age  $3.3 \pm 1.4$  (SD) yr). To create a non-  
228 associate group, 5 individuals from the herd (other than those signaled for removal) were  
229 selected at random and the 10 individuals with the strongest near neighbor association to  
230 these formed the non-associate group (mean age  $4.0 \pm 1.10$  (SD) yr). This method of  
231 selection differed from E1 to control for the possibility that our methodology in E1 may have  
232 inadvertently selected cows that were less sociable to all cows, not just the removed cows.  
233 Hence in E2 we chose cows that were equally sociable, but to different cows, for our non-  
234 associate group.

235

236 ***Behavioral Observations.*** The same ethogram detailed in E1 was used in E2.  
237 Behavioral observations of the selected subgroup of 20 individuals were carried out by the  
238 lead researcher from a viewing platform above the housing area for 3 d (d -6, -4, -2) before  
239 removal of pre-selected individuals and 3 d (d+1, +3, +6) after removal. Observations were  
240 conducted using instantaneous scan sampling (Martin and Bateson, 1993) at 10 min intervals  
241 for a 2 h duration at 0200, 1000 and 1800 h, totaling 36 scans per day.

242

243 ***Physiological Sample Collection.*** Milk samples were collected from each of the 20  
244 cows, using polypropylene centrifuge tubes (BRAND GMBH + CO KG, Wertheim,  
245 Germany), during p.m. milking between 1500 to 1600 h on d +1, +3, +6. Samples were

246 centrifuged immediately for 10 min at 3500 rpm, and skim milk was extracted and frozen at -  
247 25°C for subsequent determination of IgA and cortisol concentrations.

248

### 249 ***Experiment 3***

250 ***Animals.*** Experiment 3 (**E3**) was carried out at the same farm as E2 with observations of 9  
251 Holstein-Friesian dairy calves, maintained on the farm within a group of 18 calves (mean age  
252 65 d  $\pm$  16.8 d [SD]). The group of 18 calves were housed together continuously in a straw  
253 and peat-bedded pen (4.5m x 3.5m; stocking density = 1.1 calf per m<sup>2</sup>) (Figure 1) from  
254 approximately 2 wk after birth and maintained on milk replacer (Denka Milk, Voorthuizen,  
255 Netherlands) fed at levels up to 8 L/d at 8 wk of age, obtained from an automatic milk feeder.  
256 Levels of milk intake were not recorded. Hay was provided ad libitum and pellets (Saldus  
257 Labiba, Saldus, Lativa), were provided at up to 1.72 kg DM/d, according to live weight,  
258 accessed from an automatic feeder. All individuals had *ad libitum* access to water. A subset  
259 of 9, 8 wk old, calves were identified for removal into older age groupings. The remaining 9  
260 calves were a mean age of 36 d  $\pm$  9.8 d (SD). The stocking density after removal was 0.6  
261 calves per m<sup>2</sup>. Near neighbor observations were not possible due to the small numbers of  
262 calves in this study.

263

264 ***Behavioral Observations.*** The same ethogram detailed in E1 was used in E3 with the  
265 addition of 'play' behavior. Behavioral observations of the 9 calves were carried out by a  
266 single observer from a viewing area fronting the housing area for 3 d (d -5, -3, -1) before and  
267 3 d (d+1, +3, +6) after removal of pre-selected individuals. Observations were conducted  
268 using instantaneous scan sampling (Martin and Bateson, 1993) at 10 min intervals for 2 h at  
269 1100 and 1700 h, totaling 24 scans per day

270

271 **Physiological Sample Collection.** As milk samples were unable to be collected from  
272 this group, blood samples were collected by a veterinarian from the coccygeal vein on each of  
273 the 9 calves, using heparinized tubes (Venoject<sup>®</sup>, Terumo Corporation, Belgium), between  
274 1000-1100 h on d -1, +1, +3, +6. Samples were centrifuged immediately for 20 min at 3500  
275 rpm, and serum was extracted and frozen at -25°C for subsequent determination of IgA and  
276 cortisol concentrations.

277

#### 278 **Experiment 4**

279 **Animals.** Experiment 4 (**E4**) was conducted on a commercial dairy farm in Rahinge,  
280 Estonia. Subjects comprised 4 16-month old, Holstein-Friesian bulls, maintained in a group  
281 of 22 (stocking density = 0.3 bulls per m<sup>2</sup>). They had been housed together continuously  
282 since around 8 wk of age and were loose housed in a deep straw pen (70 m<sup>2</sup>) within a larger  
283 barn (Figure 1). They had ad libitum access to water and were maintained on a TMR  
284 consisting of grass/clover silage, hay (83% DM) and commercial pellets (300 g/d, 86% DM),  
285 fed at 0900 and 1700 h. A subset of 18 of the bulls were selected for removal based on  
286 qualification for live export. Stocking density after removal was 0.06 bulls per m<sup>2</sup>.

287

288 **Behavioral Observations.** Observations were carried out from a viewing platform  
289 above the group utilizing the same ethogram as in E1. The 4 bulls were observed using focal  
290 animal sampling (Martin and Bateson, 1993) for 15 min durations, at 1 h intervals, between  
291 1000 and 1800 h (totaling 2 h of per bull/ per day) on 4 d (d -7, -4, -2, -1) before and after (d  
292 +1, +3, +5, +7) removal. Near neighbor observations were not possible due to the small  
293 numbers of bulls in this study.

294

295 ***Physiological Sample Collection.*** As milk samples were unable to be collected from  
296 this group, blood samples were collected by a veterinarian from the coccygeal vein, using  
297 heparinized tubes (Venoject®, Terumo Corporation, Belgium) on each of the 4 bulls animals  
298 between 1000 and 1100 h on d -1, +1, +3, +5. The bulls were restrained in a holding corner of  
299 their home pen during sample collection. Samples were centrifuged immediately for 20 min  
300 at 3500 rpm, and serum was extracted and frozen at -25°C for subsequent determination of  
301 IgA and cortisol concentrations.

### 302 ***Identification***

303 Individuals in all experiments were numerically identified (on both the head and rump) using  
304 tail paint (FIL Tell Tail, Farmers Industries Limited, New Zealand [E1] or Porcimark  
305 Maerkespray, Kruuse [E2, E3 and E4]).

306

### 307 ***Physiological Sample Analysis***

308 To calculate IgA concentration, the optical density of samples was compared to the optical  
309 density of a standard with a known concentration of IgA, using the Bovine IgA ELISA  
310 Quantitation Kit (E103, Bethyl Laboratories, Montgomery, Texas, USA). ELISA plates were  
311 coated with 100 µl/well of diluted anti-bovine IgA antibody (E10-121, Bethyl Laboratories,  
312 Montgomery, Texas, USA) and incubated at room temperature for 60 min diluted to 1µg/mL  
313 in carbonate-bicarbonate buffer at pH 9.5. Plates were washed five times with wash solution  
314 (50mM/L Tris, 0.14M/L NaCl, 0.5ml/L Tween20, dH2O). Plates were then blocked for 30  
315 min at room temperature with 200 µl/well of blocking solution (50mM/L Tris, 0.14M/L  
316 NaCl, dH2O, 1% BSA) added to each well and incubated at room temperature for 30 min.  
317 Plates were washed five times and 100µl/well of diluted standards or samples were added.  
318 Then 1.5µl of sample was diluted in 1.5mL of diluent, based on the expected concentration

319 (sample dilute: 50mM/L Tris, 0.14M/L NaCl, 0.5ml/L Tween20). Samples were diluted  
320 starting at 1:1000 and extending to 1:156,000. Plates were incubated at room temperature for  
321 a further 60 min and then washed five times, followed by the addition of 100  $\mu$ l/well of  
322 diluted anti-bovine IgA horseradish peroxidase antibody E10-121, (Bethyl Laboratories,  
323 Montgomery, Texas, USA) and incubation at room temperature for a further 60 min. Plates  
324 were washed a further five times and 100  $\mu$ l/well of tetramethylbenzidine substrate solution  
325 was added, with a stop solution (0.18M H<sub>2</sub>SO<sub>4</sub> at 100 $\mu$ l/well) added after 5 min. Optical  
326 density was read at 450 nm with a microplate reader (IL650, Instrumentation Laboratory,  
327 Cheshire, UK). The concentration of IgA in each sample was calculated using linear  
328 regression from a standard curve generated from the standards using IL650 software. IgA  
329 results are reported in mg/dL of serum/milk.

330

331 Cortisol concentrations were quantified with a solid phase competitive chemiluminescent  
332 enzyme immunoassay (Immulite 1000, Siemens Medical Solutions Diagnostics, Camberley,  
333 UK) using a commercial kit (Immulite Cortisol Kit, Siemens Medical Solutions Diagnostics,  
334 Camberley, UK). Results for the cortisol were generated from the standard curves produced  
335 within the Immulite 1000. The cortisol detection limit was 27.6 mM/L.

336

337 All samples were analyzed by CTDS Veterinary Diagnostic Laboratory, Leeds, UK. The  
338 within assay coefficients of variability for serum and skim milk were 6.1 and 1.4% (cortisol)  
339 and 10.1 and 3.9% (IgA), respectively. The between assay coefficients of variability for  
340 serum and skim milk were 8.15 and 4.4% (cortisol) and 16.6 and 5.2% (IgA), respectively



341

342 *Statistical Analysis*

343 All values are reported as means  $\pm$  standard deviations. To establish the nearest neighbors in  
344 E1, a clustering of the observations using a dendrogram was performed. From this we  
345 identified the 10 individual cows (associate group) that had the closest associations with the 7  
346 focal cows signaled for removal. For the non-associate group the 10 cows with the weakest  
347 associations with the focal cows were selected from the cluster analysis. To confirm that  
348 these interactions did not simply occur by chance, the probability of an individual member of  
349 the herd being recorded with one of the 7 focal cows on either 2 or 3 occasions out of the  
350 total number of observations was calculated using probability theorem (Cooper et al., 2008).  
351 To summarize, the chance of each focal cow interacting with any other cow on any  
352 observation was 125/125 (125 cows were used as this was the number of cows any focal cow  
353 could interact with). The chance of the focal cow interacting on the subsequent observation  
354 with the same cow was 1/125, and with a different cow in the herd was 124/125. On the  
355 second observation, the chance of the focal cow interacting with a different cow, except the  
356 first or second, was 123/125 etc. This process continued until the remaining number of terms  
357 was equal to the total number of observations that took place (e.g. 36 for focal cow 1). This is  
358 numerically expressed as:

359

360  $125/125 \times 1/125 \times 124/125 \times 123/125 \times 122/125 \dots 91/125 = 125!/90!$  (! = factorial)

361

362 The number of places cow 1 could appear was  $36!/2!/34!$

363 The probability of 2 cows occurring together on 2 occasions, out of a possible 36 is:

364  $125!/90! \times 1/125^{36} \times 36!/2!/34! = 0.026$

365

366 The probability of 2 cows being observed together on 3 occasions was similarly calculated as  
367  $125!/91! \times 1/125^{36} \times 36!/3!/33! = 0.003$

368 In E2 and E3, Near Neighbor Associations were calculated using the association equation of  
369 Martin and Bateson (1993):

370

371 Near Neighbor Score =

$$\frac{\text{(number of times cow has been near-neighbor of focal cow)}}{\text{(number of focal cow observations)}}$$

372  
373  
374

375 Data was analyzed using Minitab (version 16). For all experiments, descriptive analysis of  
376 the recorded behavioral subcategories was carried out and any behavior occurring  
377 infrequently ( $n < 3$ ) was removed. All behavior subcategories retained for analysis within  
378 each experiment (Table 1) were converted to a proportion per day due to variations in the  
379 total number of observations. The data mostly followed a normal pattern of distribution  
380 (Anderson Darling  $P > 0.05$ ) and where deviations from normality occurred the data was  
381 transformed using Square Root or  $\text{Log}_{10}$  as appropriate. The different measures of behavior  
382 across the experiments precluded the data for being pooled and analyzed together. In E2 and  
383 E3 stand and walk behavior was combined. A General Linear Model (GLM) was used to  
384 investigate significant changes in behavior 'before' versus 'after' removal within each  
385 experiment. In E1 and E2 the model included the variables before or after removal (**B/A**), day  
386 nested within B/A and group (associate or non-associate) nested within B/A. In E3 and E4 the  
387 model included B/A, each individual calf or bull and day nested within B/A. Post-hoc Tukey  
388 was engaged to investigate where significance lay across days in all models. A change in the  
389 concentrations of IgA and cortisol across days was investigated using linear regression.

390

**RESULTS****391 Behavioral Responses**

392 In E1, one cow was removed from analysis due to sickness in the final days of observation.  
393 Based on the number of times the focal (removed cow) was seen with one of the associate  
394 cows, the probability of association for each of the focal (removed) cows was follows: Focal  
395 cow (FC) 1 = 0.03, FC2 = 0.01, FC3 = 0.01, FC4 = 0.04, FC5 = 0.01, FC6 = 0.01, FC 7  
396 =0.0002. None of the non-associate cows were recorded with the focal cows on more than  
397 one occasion. The chance that any 2 cows would be observed together on 2 or 3 occasions  
398 were small, 0.025 and 0.0035, respectively, therefore the nearest neighbor associations were  
399 considered real and not by chance. There was an increase ( $F = 9.47$ ,  $df = 1$ ,  $P = 0.003$ ) in time  
400 spent eating (Table 2) after the separation, with no difference between the response of  
401 associate and non-associate cows ( $P = 0.96$ ), which suggests that the removal of cows  
402 influenced remaining cows' eating behavior independent of measured associations between  
403 individuals (Table 2). There were no differences ( $P \geq 0.24$ ) in time spent standing, walking,  
404 lying, sleeping, ruminating or butting after separation (Table 2). There was no effect of day  
405 ( $P > 0.05$ ) on time spent performing any of the recorded behavior.

406

407 In E2, there was also an overall increase ( $F = 4.37$ ,  $df = 1$ ,  $P = 0.04$ ) in time that remaining  
408 cows spent eating (Table 3) after separation, however, this was only in the non-associate  
409 cows (Tukey  $P = 0.02$ ), not the associate cows (Tukey  $P = 0.99$ ). There was no change ( $P \geq$   
410  $0.34$ ) in time spent in any other behavior after separation (Table 3). There was no effect of  
411 day ( $P > 0.05$ ) on the occurrence of any behavior recorded.

412

413 In E3, a reduction in ruminating behavior was observed after separation ( $F = 7.97$ ,  $df = 1$ ,  $P =$   
414  $0.007$ ) (Table 4), with no change in time spent in any other behavior after separation. There  
415 was no effect of day ( $P > 0.05$ ) on the occurrence of any behavior recorded in E3.

416

417 In E4, a decrease in eating was observed after separation in E4 ( $F = 4.94$ ,  $df = 1$ ,  $P = 0.037$ )  
418 (Table 5). Walking was increased on d+1 following separation, compared to d+3, +5, +7, -7  
419 and -1 ( $F = 7.63$ ,  $df = 6$ ,  $P < 0.0001$ ; all Tukey tests  $P \leq 0.02$ ) (Figure 2). An effect of day  
420 was observed for standing ( $F = 2.69$ ,  $df = 6$ ,  $P = 0.04$ ) and sleeping ( $F = 2.76$ ,  $df = 6$ ,  $P =$   
421  $0.04$ ), however post hoc Tukey tests were all non-significant ( $P \geq 0.10$ )

422

### 423 *Physiological Responses*

424 There was no difference in IgA concentration after separation between the non-associate and  
425 the associate group in E2 (mean non-associate group = 113.8 mg/dL; mean of associate group  
426 = 106.2 mg/dL,  $P = 0.42$ )

427 There was an increase in IgA over the first 6 days after separation in both E3 (mean before  
428 separation = 68.2 mg/dL; mean after separation = 83.7 mg/dL; SED = 0.78 mg/dL;  $F = 93.58$ ,  
429  $df = 1$   $P = 0.01$ ) and E4 (mean before separation = 56.0 mg/dL; mean after separation = 88.3  
430 mg/dL; SED = 3.96 mg/dL;  $F = 27.80$ ,  $df = 1$   $P = 0.03$ ) (Figure 3a and 3b). The regression  
431 equations, with adjusted  $r^2$  and P values for the coefficients for day, were:

432 E3:  $y = 62.0 (\pm 2.02) + 7.13 (\pm 0.74) x$ ,  $r^2$  (adjusted) = 96.9%,  $P = 0.01$

433 E4:  $y = 43.0 (\pm 6.30) + 13.2 (\pm 2.51) x$ ,  $r^2$  (adjusted) = 89.9%,  $P = 0.03$

434 where  $y =$  IgA concentration in serum (mg/dL) and  $x =$  day

435

436 Cortisol concentrations did not differ in E2 between non-associate (mean = 1.8 mM/L) and  
437 associate groups (mean = 1.7 mM/L; SED = 0.25 mM/L;  $P = 0.79$ ), E3 (mean before  
438 separation = 33.9 mM/L; mean after separation = 30.1 mM/L; SED = 1.33 mM/L;  $P = 0.19$ )  
439 or E4 (mean before separation = 27.6 mM/L; mean after separation = 32.3 mM/L; SED = 2.87  
440 mM/L;  $P = 0.30$ ).

441

442

## DISCUSSION

443 Social behavior is a major determinant of farm animal welfare (Keeling and Gonyou, 2001;  
444 Rault, 2012). In cattle, social behavior is characterised by the formation and maintenance of  
445 cohesive social groups (Gibbons et al., 2010). In the present study, across 4 experiments, we  
446 removed both small groups of individuals from larger groups (less than 10% of the total  
447 group) and large groups of individuals (50%-80% of the total group) to study the effect of  
448 removal on remaining group members. Overt behavioral responses were limited and  
449 physiological responses were restricted to increases in IgA concentrations in small groups of  
450 cattle, with no evidence of cortisol responses. This indicates that, unlike moving individual  
451 cattle into a new-group, acute stress was not experienced by the remaining cattle, even when  
452 the majority of animals were removed and supports previous studies suggesting that cattle,  
453 moving to new groups, habituate to repeated regroupings (e.g. Mench et al., 1990)

454

455 The observed increase in time spent eating after removal in our first two experiments (E1 and  
456 E2) may be the result of some dominant cows being removed from the herd, allowing  
457 subordinates more opportunities and time to feed than previously. Heifers, for example, have  
458 been demonstrated to spend more time eating and ingest greater quantities of feed when they  
459 are kept separate from older cows (Bøe and Færevik, 2003). Age is an important determinant

460 in social positioning (Kabuga, 1992) and different types of separation, e.g. dividing a feed  
461 trough with protective barriers, have been used to improve the feeding time of subordinate  
462 cows (Bouissou et al., 2001). In E2 removed cows were on average six months older than the  
463 remaining associate cows. It is also possible that this increase could have resulted from more  
464 space being available at the feed trough or that the cows were more active, resulting in  
465 increased energy expenditure and subsequently intake requirements. However, this was not  
466 supported by increased time spent walking.

467

468 In E1, the increase was observed in both the non-associate and associate groups and suggests  
469 a herd effect. It was evident that the cows in the non-associate group did not associate with  
470 the removed cows nor did they associate much with all other cows. Hence the decision was  
471 made to change the non-associate group in E2 to be cows that associated with the same  
472 frequency with cows that were not removed. In this instance the increase in feed intake was  
473 confined to these non-associate cows, which may suggest feeding behavior in the associate  
474 cows was suppressed, comparative to the non-associate cows, as a result of the removal of  
475 associated cows. In addition, social buffering and/or emotion contagion could explain in part  
476 the apparent absence of observed behavioral changes. Central to sustaining good welfare for  
477 herd-living animals is the maintenance of synchronicity of behavior (Miller and Wood-Gush,  
478 1991). From an evolutionary perspective, similarity in emotional states achieved via the  
479 sharing of emotions can be seen as advantageous, as it results in efficient coordination of  
480 behavior (Špinko, 2012). Social buffering refers to observed reduced arousal, during stressful  
481 events, as a result of social grouping (Bouissou et al., 2001). Emotion contagion causes  
482 animals to shift their own affective state to that of other animals in a particular state (Špinko,  
483 2012). Although understudied, animals have been evidenced to both emit and detect  
484 emotional signals, and during stressful events the social group can lower the individual's

485 arousal (Bouissou et al., 2001). The emotional sharing and synchronicity of behavior has  
486 been most extensively studied in the social transmission of fear, which can be prevented by  
487 the presence of companions that do not show fear or vice versa (Veissier and le Neindre,  
488 1992; Mounier et al., 2006). For example heifers show less avoidance of unusual noise in the  
489 presence of pen mates (Boissy and Le Neindre, 1990), appear less fearful in novel  
490 environments when social partners are present (Veissier and le Neindre, 1992), display  
491 increased fear in either feeding or explorative situations when exposed to urine from stressed  
492 conspecifics and show a lower tendency to feed in the presence of a stressed partner, than in  
493 the presence of an unstressed one (Boissy et al., 1998). In the present study it is possible that  
494 individuals in the associate groups E1 and E2 did not show overt behavioral signs of social  
495 stress because the majority of the group did not experience social separation or display  
496 behavioral changes.

497

498 The natural social structure of cattle is that of a group with a mean size of 10-11 individuals  
499 (Bouissou et al., 2001). Group living involves the formation of social relationships and  
500 preferential interactions with certain companions (Nicol, 2011), which suggests cattle are  
501 likely to form positive social relationships with more than one other individual. Social  
502 support often provides a single partner to an individual (Færevik et al., 2006; De Paula Vieira  
503 et al., 2010; Duve et al., 2012), however, as cattle naturally maintain larger groups, they  
504 require more peers to benefit from social support (Boissy et al., 1998). It is possible that the  
505 impact of the removal of a group of animals was diminished by the presence of other socially  
506 important individuals that remained in the herd.

507

508 In E3 and E4 we observed behavioral changes that are more commonly associated with social  
509 stress. In E3 and E4 ruminating and eating times were reduced, respectively. Unlike the first

510 two experiments, in E3 and E4 the groups were less fluid and the majority of group members  
511 were removed so that the remaining group members (e.g. E3 n=9, E4 n=4) experienced a  
512 more similar situation to previous isolation studies where one or a small number of  
513 individuals are isolated from a larger group (von Keyserlingk et al., 2008). Consequently,  
514 remaining peers may not have been present in large enough numbers to effectively buffer the  
515 negative effect of separation, and previous experiences of separation may not have occurred  
516 with enough frequency to facilitate habituation. Following regrouping, calves housed in large  
517 groups (group size n=16) have been shown to modify their behavior to indicate improved  
518 welfare, compared to calves housed in small groups (group size n=4) (Færevik et al., 2007).  
519 Furthermore, in E3 feed intake was restricted by automated feeding equipment which may  
520 explain why a reduction in rumination was observed and not feed intake.

521

522 The possibility that the behavioral observations in E3 and E4 were indicative of stress is  
523 supported by the approximately linear increase in IgA observed up to 6 days after separation.  
524 Although more research is required to confirm the comparative IgA concentrations in bovine  
525 milk versus serum, published research in other mammal species including, sows, dogs and  
526 rats, has demonstrated that milk is rich in IgA (e.g. Heddle and Rowley, 1975; McGhee et al.,  
527 1975; Klobasa et al., 1987). One study by Näslund et al., (2000) demonstrated a high  
528 correlation between IgA titres in milk and serum, supporting the possibility of milk acting as  
529 a non-invasive alternative to serum. Additionally, IgA levels in milk appear to be consistent  
530 across lactational stages (Näslund et al., 2000). This suggests that it was the experimental  
531 design (group size and separation), rather than the substrate within which IgA was quantified,  
532 that was responsible for the observed (or lack of) changes in IgA after removal. Conversely,  
533 as we did not include a control in this study we cannot conclusively rule out the possibility  
534 that other variables (e.g. age, health status or environmental changes) may account for the



535 increase in IgA observed. To the authors knowledge all subjects were in healthy condition,  
536 had not recently received vaccination, nor were there extreme weather changes during the  
537 course of the experiment or any other notable stimuli that may have influenced IgA levels.

538

539 We also measured cortisol both between groups and before and after removal and found no  
540 significant changes, which may be because social separation is a chronic stressor. Cortisol  
541 has previously been used as a measure of the physiological impact (through HPA arousal) of  
542 social separation in a range of species and across a range of social relationships (Hennessy,  
543 1997). These have predominately involved brief separations and have not always resulted in  
544 activation of the HPA system (Hennessy, 1997). As serum collection was obtained more than  
545 12 h after removal in the present study, it is possible that HPA arousal had occurred but  
546 ended.

547

548

### CONCLUSIONS

549 Behavioral responses to removal of predetermined individuals were observed in four  
550 experiments to be context specific. In two experiments with relatively large groups of dairy  
551 cows, social stress post-separation appears to have been diminished by social buffering or  
552 previous habituation to separation. In two experiments with smaller groups of cattle, IgA  
553 concentrations and nutritional responses suggest that social separation resulted in stress over  
554 several days. These responses suggest that social separation can be detrimental to the welfare  
555 of those remaining in the group but that the impact can be ameliorated by the presence of  
556 unstressed group members.

557

558

559

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563

564

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**Tables***Table 1: Behavioral categories and subcategories and their definitions*

Behavior	Definition
Locomotive Behavior	
Walk <sup>1,2,3,4</sup>	Forward movement with three legs remaining in contact with the ground.
Stand <sup>1,2,3,4</sup>	All four hooves on ground and legs upright and extended to support body.
Lie <sup>1,2,3,4</sup>	Recumbent, with body on floor.
Sleep <sup>1,2,3,4</sup>	Sternally recumbent with head tucked backwards towards shoulder, resting on body.
Maintenance Behavior	
Eat <sup>1,3,4</sup>	Cow ingests (or pokes with muzzle) food provided at feed bunk or from automatic feeder. Calf ingests milk provided by automated milker.
Drink <sup>4</sup>	Cow imbibes from automated water system.
Defecate <sup>5</sup>	Cow passes a faecal motion in standing or lying position.
Urinate <sup>5</sup>	Cow passes urine in a standing or lying position.
Ruminate <sup>1,2,3,4</sup>	Regurgitation, chewing, and swallowing of food bolus.
Vocal Behavior	
Vocalise <sup>5</sup>	Audible noise is produced by the cow.
Oral Behavior	
Lick <sup>5</sup>	Part of the tongue is protruded and moved over surfaces (e.g. pen bars).
Sniff <sup>5</sup>	Muzzle close to an object (or other individual) and inhaling air.
Social Interaction	
Play <sup>3</sup>	Running, trotting, galloping or jumping, alone or with other calves.
Human Interaction <sup>5</sup>	Physical contact with human.
Butting <sup>1</sup>	Cow uses head to head or head to body contact in an attempt to physically push another cow, or cause a cow to rise from lying position.
Other	
Groom – Auto <sup>2,3,4</sup>	Rubbing parts of the body or head against other body parts or fixtures of the stall (including automated groomer), or licking body parts.
Groom – Allo <sup>2,3,4</sup>	Rubbing parts of the body or head against another individually or licking the other individual.

Table 1 provides behavioral descriptors utilized for analysis

<sup>1</sup> = Behaviors analyzed in Experiment 1

<sup>2</sup> = Behaviors analyzed in Experiment 2

<sup>3</sup> = Behaviors analyzed in Experiment 3

<sup>4</sup> = Behaviors analyzed in Experiment 4

<sup>5</sup> = Behaviors not included in analysis due to infrequency of occurrence

Table 2: The mean % (and SED) of time for each behavior occurring across all days before and all days after removal for non-associate cows ( $n = 10$ ) and associate cows ( $n = 10$ ) in Experiment 1 (investigation of the effects on remaining cows as a result of removal of a small group of individuals [ $n = 7$ ] from the larger herd [ $n = 126$ ]) and the probabilities of differences before and after removal. There were no significant ( $P \leq 0.05$ ) associate Vs non-associate effects or individual day effects

Behavior	Before Removal		After Removal		SED	Before-After P-Values
	non-associate	associate	non-associate	associate		
Standing	46.0%	55.5%	49.8%	56.3%	2.02	0.40
Walking	2.2%	2.4%	2.4%	2.9%	1.01	0.41
Lying	49.1%	41.1%	44.4%	39.3%	2.01	0.24
Sleeping	2.2%	1.0%	1.0%	1.5%	0.40	0.55
Eating	19.9%	19.8%	26.0%	26.7%	1.5	0.003
Ruminating	30.9%	31.6%	36.9%	30.5%	1.48	0.24
Butting	0.08%	0.10%	0.09%	0.11%	0.01	0.74

Table 3: The mean % (and SED) of recordings for each behavior occurring both before and after cow removal for non-associate cows ( $n = 10$ ) and associate cows ( $n = 10$ ) in Experiment 2 and the probabilities of differences before and after removal. Post hoc (Tukey P-Value) associates Vs non-associate effects are reported. There were no significant ( $P \leq 0.05$ ) individual day effects.

Behavior	Before Removal			After Removal			SED	Before-After P-Values
	non-associate	associate	Tukey P-Value	non-associate	associate	Tukey P-Value		
Standing	17.2%	17.2%	-	16.4%	15.7%	-	4.52	0.48
Lying	50.2%	50.9%	-	46.6%	49.4%	-	1.88	0.34
Sleeping	5.6%	4.4%	-	5.7%	3.5%	-	0.45	0.89
Eating	13.5%	16.2%	0.02	19.5%	16.7%	0.99	1.18	0.04
Ruminating	30.2%	32.4%	-	30.7%	32.1%	-	1.41	0.99
Grooming	1.16%	1.16%	-	1.35%	0.83%	-	0.24	0.82



Table 4: The mean % (and SED) of recordings for each behavior occurring in calves ( $n = 49$ ) both before and after removal of conspecifics ( $n = 18$ ) in Experiment 3. There were no significant ( $P \leq 0.05$ ) individual day effects.

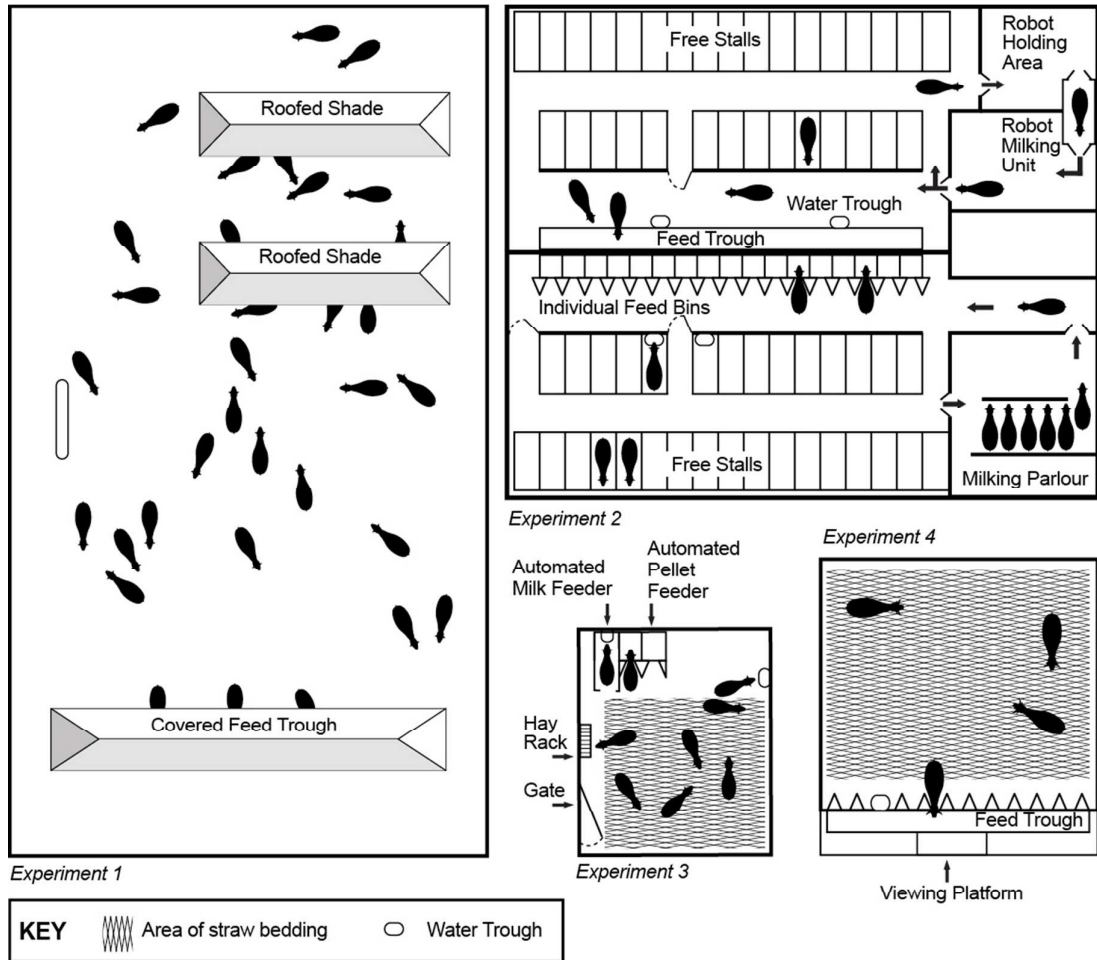
Behavior	Before Removal	After Removal	SED	Before-After P-Values
Standing	20.1%	18.7%	2.37	0.50
Lying	51.7%	56.9%	2.96	0.17
Sleeping	28.0%	24.0%	2.64	0.40
Eating	9.4%	8.4%	3.17	0.99
Ruminating	28.85	20.1%	2.24	0.007
Grooming	2.12%	3.23%	2.45	0.29

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Table 5: The mean % ( $\pm$ SED) of time for each behavior occurring in bulls ( $n = 4$ ) both before and after removal in Experiment 4 and the probabilities of differences before and after removal and day.

Behavior	Before Removal	After Removal	SED	Before-After P-Values	Day P-Values
Standing	33.1%	32.8%	3.60	0.95	0.04
Walking	1.5%	1.2%	0.22	0.39	<0.0001
Lying	59.0%	55.1%	3.84	0.48	0.29
Sleeping	4.4%	6.6%	1.21	0.21	0.04
Eating	15.7%	8.8%	2.22	0.04	0.15
Drinking	1.3%	1.3%	0.48	0.98	0.46
Ruminating	36.3%	25.5%	4.59	0.11	0.41
Butting	0.9%	2.9%	1.06	0.19	0.30
Grooming	1.7%	3.1%	0.52	0.65	0.07

Walker Figure 1



view

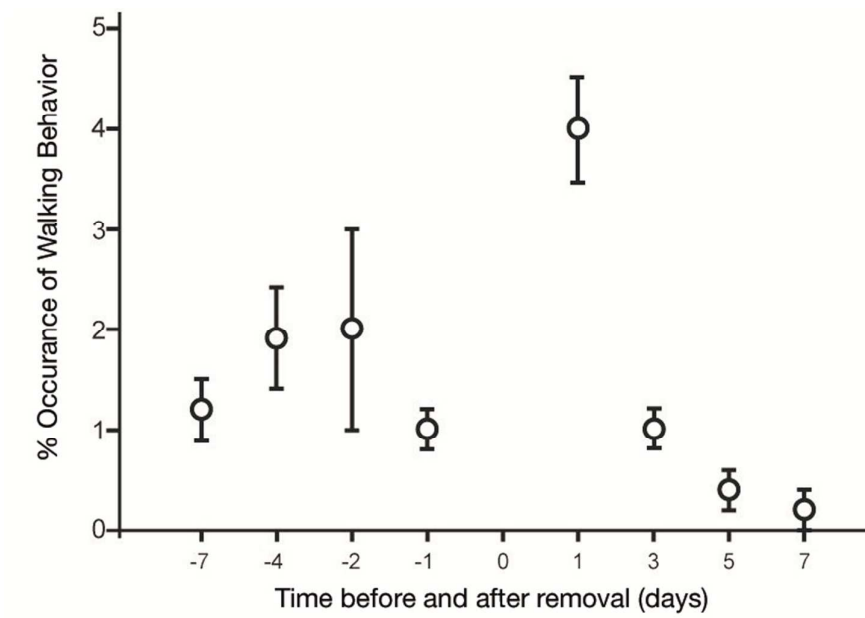
*Walker Figure 2*

Figure 2, Manuscript Number , 34

Walker Figure 3

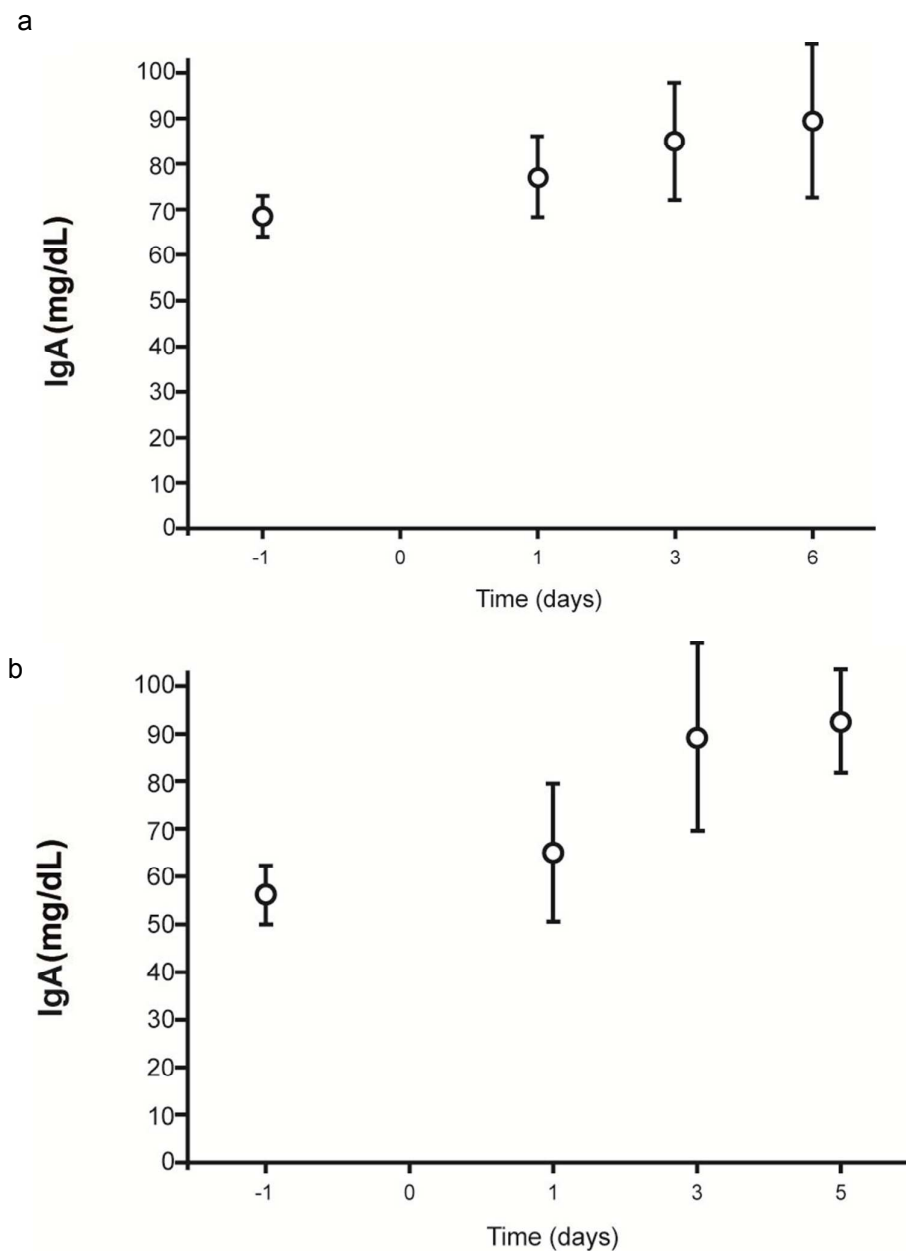


Figure 3, Manuscript Number , 35

***Figure Captions***

Figure 1: Overhead view of experimental facilities. Number of cattle does not represent exact experimental numbers in Experiment 1 and 2.

Figure 2: The % of walking behavior displayed across the 8 days of observations in Experiment 4.

Figure 3: Mean IgA concentrations in Experiment 3 a ( $F = 93.58$ ,  $df = 1$   $P = 0.01$ ) from d -1 (before removal) to d +1, +3, and +6 (after removal) and in Experiment 4 b ( $F = 27.80$ ,  $df = 1$   $P = 0.03$ ) across d -1 (before removal) to d+1, +3, and +5 (after removal).