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1 **Quantifying intraherd cattle movement metrics: implications for disease**
2 **transmission risk**

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5

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23 Abstract

24 There is a paucity of data quantifying on-farm management practices such as the frequency of
25 intraherd cattle movements, use of consolidated or spatially fragmented grazing pastures, and
26 duration of time cattle spend at grass with respect to biosecurity and disease transmission. Such
27 movement dynamics are important when attempting to understand the maintenance of chronic
28 infectious disease, such as bovine tuberculosis (bTB). We captured empirical data on daily
29 cattle movements for a sample of eighteen farms throughout one complete grazing season
30 ($n=18,988$ grazing days) and assessed these attributes in relation to herd bTB risk.

31 Dairy herds were stocked at significantly higher densities compared to beef production systems
32 (6.6 animals/ha, 95% confidence intervals (CI) 6.5 – 6.7 and 4.1 animals/ha, 95%CI 4.1 – 4.1
33 respectively, $p<0.001$). Most notably milking cows, were grazed at higher densities than other
34 life stages (e.g. calves, heifers and bullocks) ($p<0.001$) and experienced four times the number
35 of movements between pastures. Beef cattle were more likely to be grazed across multiple
36 (rather than single) fields ($p<0.001$), with greater time spent on fragmented land away from the
37 main/home farm ($p<0.001$). None of the farm or herd attributes analysed (e.g. stocking density,
38 frequency of movement, movement distances or land fragmentation) were associated with herd
39 bovine tuberculosis (bTB) breakdowns during this study. However, there was a weak positive
40 association between bTB breakdowns during the 3 years prior to the study and cattle movement
41 distances ($p=0.05$) and time spent on fragmented land ($p=0.08$). After a bTB breakdown occurs,
42 restrictions on animals moving out of these herds are implemented to control disease spread,
43 yet we argue that more attention is needed on the role of intraherd grazing patterns in modelling
44 disease transmission risk between herds.

45 **Keywords:** Cattle movements; Bovine tuberculosis; Disease Control; Mycobacterium bovis;
46 Biosecurity.

47 **Introduction**

48

49 Bovine tuberculosis (bTB) is a disease of cattle, primarily caused by infection with
50 *Mycobacterium bovis*. The disease mainly affects the respiratory system and transmission
51 between cattle is thought to occur predominately through direct aerosol contact, but indirect
52 contacts may play a role (Griffin et al., 1993; Phillips et al., 2003). Clinical disease is rarely
53 encountered in countries where national disease eradication programmes identify infected
54 animals through ante-mortem testing and removal early on in the course of infection (Collins,
55 2006).

56 Interherd contact of cattle between farms is a known risk factor in the spread of bTB and other
57 infectious diseases (Broughan et al., 2016; Gates et al., 2014), and can facilitate the
58 maintenance of disease despite control (Allen et al., 2018). Previous research of cattle
59 movement networks suggests high connectedness between herds through animal movements
60 and trade can occur (Brown et al., 2019). Direct contact between cattle from different herds
61 can occur at agricultural shows, markets, veterinary practices, shared transport, shared housing
62 and contact ‘across-the-hedge’ (so called ‘nose-to-nose’ contact) between neighbouring herds
63 (Dommergues et al., 2012; Robinson and Christley, 2007). Indeed, at an animal-level
64 movement can be associated with metrics of infection (Byrne et al., 2017; Ramírez-Villaescusa
65 et al., 2009).

66 There is an increased risk of a bTB herd breakdown if local herd prevalence is high (Doyle et
67 al., 2014; White et al., 2013). Current knowledge of intraherd movements (between cattle
68 within the same herd) during the grazing season is scarce, and the implications for the
69 transmission of *M. bovis* is poorly understood (Allen et al., 2018; Brennan et al., 2008; Griffin
70 et al., 1993). Cattle are commonly moved during the grazing season between fields to ensure
71 there is enough forage to meet nutritional requirements. A higher number of intraherd

72 movements between pastures during the grazing season could not only increase intraherd
73 contact rates between different batches of cattle but also interherd contacts rates as cattle may
74 be more frequently adjacent to neighbouring herds. Therefore, grazing movements may play a
75 role in local networks of *M. bovis* transmission. In addition, wildlife can play a part in bTB
76 breakdown risk (notably the European badger, in Britain and Ireland; Allen et al., 2018), and
77 wildlife exposure risk varies spatially depending habitats and resources available across land
78 parcels (Byrne et al., 2015; Campbell et al., 2019). Intraherd movement of animals may
79 modulate this exposure risk, and may also impact on epidemiological studies investigating
80 spatial associations between wild and domesticated hosts (Byrne et al., 2015; Vial et al., 2011).

81 The issue of intraherd movements is likely to vary spatially, explained in large part by local
82 and regional land delineation practices, types of field boundary, patterns of land inheritance,
83 ownership or lease, and frequency of cattle movements (depending on field size etc.). For
84 example, patterns of local land inheritance in Ireland mean many farms have been divided
85 between children, and therefore become fragmented i.e. fields within any one farm may not be
86 spatially contiguous but there may be multiple scattered single fields or contiguous patches
87 (Aalen, 1963). Moreover, some herd owners with small areas of land may purchase or rent
88 additional land (called ‘conacre’) to make farms economically viable (Allen et al., 2018). Such
89 land fragmentation may lead to longer distance and more frequent movements of cattle between
90 consecutively grazed land parcels again potentially increasing intra – and inter- herd contacts,
91 creating opportunities for bTB transmission.

92 High stocking densities may increase disease transmission (Neill et al., 1989). The highest risk
93 of bTB spread within a herd is thought to occur when the animals are housed indoors (O’Reilly
94 and Daborn, 1995). There are recommendations for stocking densities for housed cattle
95 (Herzog et al., 2018) and whilst there are agri-environment scheme recommendations for
96 stocking densities when at grass (e.g. habitat management and energy requirements), there are

97 few recommendations for stocking densities of cattle in the wider countryside in relation to
98 disease spread.

99 Brennan *et al.* (2008) described the difficulties in examining intra- and inter- herd contact rates
100 as there is huge spatial and temporal variability, illustrating a high degree of structural
101 complexity and heterogeneity. Due to the complexity of farming practices and the difficulty in
102 getting on-farm information, there are currently no data available quantifying variation in
103 stocking densities, duration of days spent on contiguous versus fragmented land and intraherd
104 movement rates for a sample of working farms. Furthermore, there is a dearth of basic data in
105 this space, to create rulesets to make better use of large-scale land parcel information systems,
106 where they are available (for example, in Ireland and the UK; (Durr and Froggatt, 2002)).

107 This study aimed to capture empirical data on cattle movements for a sample of farms daily
108 throughout one complete grazing season. The objectives were to quantify: 1) stocking densities
109 of cattle at grass, 2) the amount of time spent on fragmented land and 3) intraherd movement
110 parameters including number of moves, distance moved, and the total extent of grassland used
111 for grazing. Furthermore, comparisons were made between these metrics and the major
112 enterprise types (dairy versus non-dairy herds), and the disease (bTB) histories within these
113 herds. These data will help describe real-world grazing practices and help parameterise future
114 models of disease transmission risk with respect to intraherd movement.

115

116 **Material and Methods**

117

118 *Study site*

119 A total of 25 cattle farms (44% dairy and 56% beef) were surveyed in County Down, Northern
120 Ireland, UK. The area had a high cattle density and was within a so-called ‘bTB hotspot’ with
121 records of herd outbreak for many years (DAERA, 2018; Milne *et al.*, 2019a; Wright *et al.*,

122 2015). The landscape was predominately improved grassland (pasture) grazed by cattle or used
123 for silage production with some sheep grazing and a small proportion of interspersed arable
124 fields.

125

126 *Quantifying time spent at grass*

127 Farmers were given an individual record book with maps of their own farms with each field
128 given a Unique ID. Farmers recorded for every batch of cattle put out to graze the Unique ID
129 of the field (i.e. its spatial location), the date during which cattle were turned out and
130 subsequently moved on, the number of cattle and the batch life stage i.e. calves, heifers,
131 bullocks or cows. Data were recorded daily from May to November 2016. Farmers were
132 (re)engaged weekly by telephone to ensure recording was continuous and consistent. Monthly
133 farm visits (by E.C.) were made for quality assurance to check farmer-recorded data and to
134 ensure there were no problems with data recording. Any persistent issues with data collection
135 or inconsistency in cattle locations that could not be retrospectively validated or corrected,
136 resulted in data from seven farms being excluded from analysis.

137 Throughout the study it was found that some cattle were grazed in multiple fields at a time
138 (gates between adjacent fields being left open). To allow for this in the analysis such fields
139 were combined and treated as one grazing unit with the total summed area used to calculate
140 stocking densities.

141

142 *Bovine tuberculosis*

143 The Department of Agriculture, Environment and Rural Affairs (DAERA), Northern Ireland
144 provided the frequency of bovine tuberculosis herd breakdowns on each study farm for the one,
145 three and five year periods prior to this study. A data sharing agreement was signed between
146 parties with landowner data anonymised such that shared data complied with the Data
147 Protection Act (1998) and General Data Protection Regulation (GDPR) 2016/679. The recent

148 historical status of bTB rather than the future status of the farm was analysed due to the badger
149 Test Vaccinate or Remove study being performed in the area by DAERA which may have
150 influenced future herd bTB tests. Doyle et al. (2016) found farms with historic bTB
151 breakdowns were statistically more likely to have a future breakdown, therefore we argue that
152 using historic bTB status is a reasonable proxy for future risk. It is reasonable to assume that
153 herd management at grass stayed relatively consistent year on year, as none of the study farms
154 advised of new grazing practices.

155

156 *Statistical analyses*

157 Descriptive statistics (medians, 95% confidence intervals (CI), ranges, Mann-Whitney,
158 Kruskal-Wallis and Fishers Exact tests) were used to summarise patterns in stocking densities,
159 percentage of days spent on fragmented land (land not contiguous to the main land parcel),
160 number of moves, and the distances between consecutive pastures between cattle batch life
161 stages and production systems. The distance of all farm movements was collected for each
162 individual move (field to field, field to housing and vice versa) and then totalled for each farm
163 to give the total distance of farm movements. The centroid of each field was used to calculate
164 the distance of movements. Variation in bovine tuberculosis outbreak history per herd (0/1)
165 was analysed using Generalised Linear Models (GLM) assuming a binomial error structure and
166 a logit link function. Bovine tuberculosis history (one, three and five year) was modelled as the
167 outcome variable and stocking density, area grazed, total number of cattle movements, distance
168 of movements, days spent on fragmented land and length of grazing season modelled separately
169 as explanatory variables. Analyses were conducted using R v3.4.2 (R Core Team, 2018) with
170 the packages *rcompanion* (Mangiafico, 2019) and *ggplot2* (Wickham, 2016). The package
171 *adehabitatHR* (Calenge, 2006) was also used to calculate the extent (hectare) of the 95% and
172 100% Minimum Convex Polygons (MCP) of fields grazed. Farm movement data was

173 calculated at the farm-level only to account for cattle batches maturing and entering the next
174 stage in production during the grazing season.

175 **Results**

176 A total of 18/25 (72%) of farms collected data that met our quality assurance validation and
177 were retained for analysis. Usable data were collected from the 2nd May to the 30th November
178 2016 (213 consecutive grazing days) at 7 dairy farms (39%) and 11 beef farms (61%). A
179 summary of the dairy and beef farm attributes is shown in Table 1.

180 Median stocking density was 4.7 animals/ha (95%CI 4.7 – 4.8, range 0.5 - 143.8). Median
181 stocking densities in dairy production (6.6 animals/ha, 95%CI 6.5 – 6.7, range 0.9 - 143.8)
182 were significantly higher than beef production (4.1 animals/ha, 95%CI 4.1 – 4.1, range 0.5 -
183 45.3; Mann-Whitney $W=33221000$, $p<0.001$) and varied significantly between batch life
184 history stages (Kruskal-Wallis $\chi^2_{df=3}=1990.1$, $p<0.001$) being highest for dairy cows and lowest
185 for beef bullocks (Fig. 1a). Bovine tuberculosis history during one, three and five years prior
186 to the study was unrelated to stocking density (GLM $\chi^2_{df=1}=0.11$ $p=0.76$; $\chi^2_{df=1}=0.18$, $p=0.67$;
187 and $\chi^2_{df=1}=1.25$, $p=0.30$ respectively).

188 Median number of land parcels per farm was 3.5 (95%CI 2 - 4, range 1 - 8) with dairy herds
189 utilising on average 4 land parcels (95%CI 2 – 4, range 2 - 6) and beef herds 3 land parcels
190 (95%CI 2 – 5, range 1 - 8). Cattle movements occurred within an area covering 54km²
191 (100%MCP of all fields) with the majority within 28km² (95%MCP). The extent of the area
192 covered by grazing on each farm did not differ between production systems (100% MCP Mann-
193 Whitney $W=28$, $p=0.38$, 95% MCP Mann-Whitney $W=23$, $p=0.18$, Fig. 2). Bovine
194 tuberculosis history during one, three and five years prior to the study was unrelated to the
195 extent of the area grazed regardless if it was the 100%MCP (GLM $\chi^2_{df=1}=0.96$ $p=0.33$;
196 $\chi^2_{df=1}=0.48$, $p=0.49$; and $\chi^2_{df=1}=0.15$, $p=0.70$ respectively) or 95%MCP that was examined
197 (GLM $\chi^2_{df=1}=2.31$, $p=0.15$; $\chi^2_{df=1}=1.96$, $p=0.18$; and $\chi^2_{df=1}=1.86$, $p=0.20$ respectively).

198 All farms (100%) moved batches of cattle between fields throughout the grazing season with a
199 total of 991 separate movement events across the 18 farms (median = 44 cattle movements per
200 farm per season, range 10 - 149 movements per farm). Number of moves per farm was
201 significantly higher for dairy than beef production systems (Mann-Whitney $W = 2$, $p=0.001$;
202 Table 1). Bovine tuberculosis history during one, three and five years prior to the study was
203 unrelated to number of cattle moves per farm (GLM $\chi^2_{df=1}=3.22$ $p=0.11$; $\chi^2_{df=1}=2.83$, $p=0.12$;
204 and $\chi^2_{df=1}=4.58$, $p=0.07$ respectively).

205 The median total distance of farm movements between consecutively occupied fields was
206 27.5km (range 2.4 - 107.6km) with a total distance of 592.3km across all 18 study farms
207 throughout the grazing season. There was no difference in the total distance of farm moves
208 between production systems (Mann-Whitney $W = 20$, $p=0.10$). The median distance of
209 individual moves for dairy, beef and all farms were 242m, 244m and 242m respectively. There
210 was no difference in the individual length of moves between beef and dairy herds (Mann-
211 Whitney $W = 110030$, $p\text{-value} = 0.38$). Bovine tuberculosis history during one, three and five
212 years prior to the study was unrelated to the distances cattle were moved on each farm (GLM
213 $\chi^2_{df=1}=3.62$ $p=0.09$; $\chi^2_{df=1}=6.52$, $p=0.05$; and $\chi^2_{df=1}=3.76$, $p=0.11$ respectively).

214 A total of 16/18 farms (89%) had fragmented land disjunct from the main/home farm and just
215 2/18 (11%) had all their fields within the main/home farm (both beef production systems). A
216 total of 14/18 farms (78%) grazed cattle on fragmented land with 4/18 herds (22%) grazing at
217 the main/home farm only (3 beef and 1 dairy production systems). Two beef herds (11%) did
218 not graze cattle at the main/home farm during the entire grazing season. 6 of the 7 dairy farms
219 (86%) grazed cattle on fragmented land though dairy cows (lactating and dry) spent all their
220 time on the main/home farm (0 days on fragmented land).

221 Median percentage of time herds spent on fragmented land was 38% of days (95% CIs 15.6-
222 55.6%, range = 0-100%). The total time spent on fragmented land differed significantly
223 between production systems ($\chi^2_{df=1}=72.913, p<0.001$, Table 2b) and between batch life history
224 stages ($\chi^2_{df=3}=4098.3, p<0.001$) with dairy cows and heifers spending the greatest time on the
225 main/home farm and beef bullocks most on fragmented land parcels (Table 2b). The median
226 percentage of days batches of cattle on farms spent on fragmented land is shown in Fig. 1b.
227 Bovine tuberculosis history one, three and five years prior to the study was unrelated to time
228 spent on fragmented land (GLM $\chi^2_{df=1}=0.71, p=0.43$; $\chi^2_{df=1}=4.24, p=0.08$; and $\chi^2_{df=1}=3.09,$
229 $p=0.12$ respectively).

230 There was a total of 18,988 grazing days with cattle spending 11,627 days in single field
231 compartments and 7,361 days in combined/multiple field compartments. Time spent in
232 combined/multiple field compartments varied significantly between dairy and beef herds ($\chi^2_{df=1}$
233 $=39.865, p<0.001$) and batch life stages ($\chi^2_{df=3}=1398.2, p<0.001$) with calves, cows and heifers
234 (i.e. dairy animals) spending most time in single field compartments (Table 2a).

235 There was no difference in the grazing season duration between dairy and beef production
236 systems (Mann-Whitney $W=32, p=0.58$; Table 1). Bovine tuberculosis history during one,
237 three and five years prior to the study was unrelated to grazing season duration (GLM
238 $\chi^2_{df=1}=1.14, p=0.34$; $\chi^2_{df=1}=1.32, p=0.28$; and $\chi^2_{df=1}=0.51, p=0.48$ respectively).

239

240 **Discussion**

241

242 There is a paucity of basic data quantifying simple farm management practices such as the
243 frequency of intraherd cattle movements, use of consolidated or spatially fragmented pastures
244 and duration of time at grass. Here, we provide a case study example of cattle movement
245 metrics for a sample of eighteen farms in Northern Ireland highlighting the high frequency of

246 cattle movements and use of fragmented land which hitherto has not been accounted for in
247 analyses of bovine tuberculosis epidemiology. The results from this study highlight the
248 variability in stocking densities and movement metrics even within a small study population.
249 These results would likely vary across landscapes and farming cultures and may not be directly
250 comparable to other geographical locations but should highlight the need to understand local
251 grazing management when trying to model bTB dynamics across time and space. If tailored
252 biosecurity advice is developed on a per herd basis, an understanding of the intra-inter- herd
253 exposure/risk would be beneficial.

254 Dairy production systems had the largest herd size and highest stocking density whilst the
255 higher metabolic demands of milk production compared to other life history stages (MSD Ltd,
256 2017) necessitated twice the number of moves than beef production systems. In dairy systems,
257 available grass was likely to be exhausted sooner than for lower density, less energetically
258 demanding life stages. Dairy cows were stocked at highest densities and are known to be at
259 greater risk of bTB outbreaks (Doyle et al., 2014; Milne et al., 2019b) which may be density-
260 dependent. Dairy farms keep cattle close to the home farm for access to the milking parlour
261 with dairy cows spending zero days on fragmented land, potentially reducing their exposure to
262 neighbouring herds. Whilst this may be expected for lactating cows, even dry cows were
263 retained on home fields on the seven dairy farms likely due to dry cows being near their calving
264 date and farmers keeping them home for close monitoring for potential ill health or calving
265 complications. Since older cattle and dairy herds are risk factors for bTB, grazing dairy cows
266 on non-fragmented land only, may potentially reduce the number of neighbouring herds
267 exposed.

268 All life stages of beef cattle are typically grazed at lower stocking densities than dairy cattle.
269 Some groups of cattle were given access to multiple fields at the same time, this occurred more
270 frequently with batches of beef cattle than with dairy cows, which were grazed in single fields

271 more commonly. Allowing beef cattle access to multiple fields simultaneously may explain
272 their lower stocking densities. By maintaining lower density, more spatially dispersed stocking;
273 may potentially reduce intraherd contact rates (re: disease transmission) and density-dependent
274 physiological stresses that may make animals more vulnerable to infection.

275 Beef farms had the longest distances moved per farm, whilst beef bullocks spent the most time
276 on fragmented land (though median individual distances moved were similar between beef and
277 dairy systems). These animals may therefore be at greatest risk of disease transmission but are
278 ultimately removed from the herd after being taken for slaughter.

279 We found substantial variation in the extent of grazed land (MCPs of grazed fields). Some
280 farms in the current study had up to 8 spatially disjunct land parcels resulting in greater numbers
281 of boundaries with adjacent farms (likely increasing the time cattle spend adjacent to other
282 herds increasing across-the-hedge (nose-to-nose) contact). Thus, a bTB free herd in greater
283 contact with adjacent herds may be at a higher infection risk than those herds that spend their
284 time on consolidated single land parcels and a bTB infected herd occupying or being moved
285 between numerous disjunct land parcels may come into contact with adjacent bTB free herds
286 exposing them to potential infection. Herds that cover larger areas may come into contact with
287 an increased number of badger social groups, due to the differences in bTB prevalence within
288 different badger social groups (Delahay et al., 2000), the risk of *M. bovis* transmission may be
289 increased.

290 None of the farm or herd attributes in terms of stocking densities, frequency of movements,
291 distance of movements and fragmentation of grazing land influenced a herd's bTB outbreak
292 status, regardless of the window examined (1, 3 or 5 years prior to study). There was a weak
293 positive trend for bovine tuberculosis outbreaks during the 3 years prior to study with distances
294 cattle were moved ($p=0.05$) and time spent on fragmented land ($p=0.08$). This could be due to

295 random chance (at $\alpha=0.95$ about 1/20 tests are likely to be significant by random chance) but
296 it may also be that the small sample size (18 farms) provided low statistical power to quantify
297 relationships; thus greater numbers of observations may resolve these relationships.
298 Nevertheless, Brown *et al.* (2019) suggested that herd connectivity may play an important role
299 in the bTB maintenance in Northern Ireland, mirroring findings from elsewhere (Gilbert et al.,
300 2005; Palisson et al., 2016). The Department of Agriculture, Environment and Rural Affairs
301 (DAERA) restrict herd movements after bTB breakdown to prevent exposing other herds, for
302 example, at cattle sales, markets and shows but there are currently no restrictions on intraherd
303 movement (despite considerable frequency of movement including to fragmented land for
304 some herds). Such restrictions would be logistically hard to enforce but herd owners with a
305 bTB breakdown should be recommended to evaluate and try to minimise intraherd movements
306 to reduce transmission risk to a new herd. Where practical, these approaches may allow a
307 reduction in the spatial exposure to neighbours, and indirectly to wildlife hosts (Campbell et
308 al., 2019).

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Attribute	Beef (<i>n</i> =11)	Dairy (<i>n</i> =7)	Both (<i>n</i> =18)	P-value Dairy Vs. Beef (Test)
Farm size (hectares)	39 (95%CI 23-55, range 11-112)	64 (95%CI 60-119, range 20-140)	50 (95%CI 33-70, range 11-140)	0.06 (Mann-Whitney)
Fields/farm (number)	31 (95%CI 17-38, range 8-84)	40 (95%CI 29-63, range 25-101)	34 (95%CI 25-43, range 8-101)	0.14 (Mann-Whitney)
Herd size (cattle)	43 cattle (95%CI 30-80, range 11-114)	168 cattle (95%CI 123-240, range 65-266)	74 cattle (95%CI 36-151, range 11-266)	<0.001 (Mann-Whitney)
Grazing season duration (days)	204 days (95%CI 183->213, range 180->213)	209 days (95%CI 186->213, range 180->213)	205 days (95%CI 190->213, range 180->213)	0.58 (Mann-Whitney)
Median Stocking Density	4.1 animals/ha (95%CI 4.1-4.1, range 0.5-45.3)	6.6 animals/ha (95%CI 6.5-6.7, range 0.9-143.8)	4.7 animals/ha (95%CI 4.7-4.8, range 0.5-143.8)	<0.001 (Mann-Whitney)
Total number of fields	360	337	697	
bTB outbreak in previous year (% of herds)	9 (1/11)	29 (2/7)	17 (3/18)	0.53 (Fishers Exact Test)
bTB outbreak in previous 3 years (% of herds)	36 (4-/1)	57 (4/7)	44 (8/18)	0.63 (Fishers Exact Test)
bTB outbreak in previous 5 years (% of herds)	46 (5/11)	86 (6/7)	61 (11/18)	0.15 (Fishers Exact Test)
Total number of movements	318	673	991	
Number of moves/farm	22 (95%CI 13-42, range 10-76)	89 (95%CI 70-118, range 68-149)	44 (95%CI 21-83, range 10-149)	<0.001 (Mann-Whitney)
Total distance of all moves (km)	277	315	592	
Distance of moves/farm (km)	12 (95%CI 6-46, range 2-78)	38 (95%CI 36-41, range 17-108)	28 (95%CI 11-41, range 2-108)	0.10 (Mann-Whitney)
Distance of individual moves (m)	244 (95%CI 226-268, range 70-6,339)	242 (95%CI 225-262, range 4-3-,579)	242 (95%CI 229-262, range 43-6,339)	0.38 (Mann-Whitney)
100% MCP for movement extent (ha)	67.3 (95%CI 16-314, range 5-542)	151 (95%CI 41-306, range 40-364)	99.4 (95%CI 34-228, range 5-542)	0.38 (Mann-Whitney)
95% MCP for movement extent (ha)	24 (95%CI 12-138, range 2-381)	123 (95%CI 43-305, range 35-332)	95.4 (95%CI 24-156, range 2-381)	0.18 (Mann-Whitney)

449 **Table 1** Descriptive statistics (totals, medians, ranges and confidence intervals (CI)) summarising farm, cattle
450 herd and movement attributes of beef and dairy production systems for *n*=18 in County Down, Northern Ireland
451 during the grazing season (2nd May – 30th November 2016). Units for each metrics are stated in brackets.

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Life stage / Production	a) Field arrangement			Test of association		
	Single	Multiple	Total	χ^2	df	p
i) Batch						
Bullocks	3,725 (48%)	4,098 (52%)	7,823 (41%)	1398.2	3	<0.001
Calves	1,656 (73%)	605 (27%)	2,261 (12%)			
Cows	3,378 (80%)	824 (20%)	4,202 (22%)			
Heifers	2,868 (61%)	1,834 (39%)	4,702 (25%)			
<i>Sub-total</i>	<i>11,627 (100%)</i>	<i>7,361 (100%)</i>	<i>18,988 (100%)</i>			
ii) Production						
Beef	5,677 (59%)	3,941 (41%)	9,618 (51%)	39.9	1	<0.001
Dairy	5,950 (63%)	3,420 (37%)	9,370 (49%)			
<i>Sub-total</i>	<i>11,627 (100%)</i>	<i>7,361 (100%)</i>	<i>18,988 (100%)</i>			
b) Field contiguity						
i) Batch						
	Main/home	Fragmented	Total	4098.3	3	<0.001
Bullocks	2,570 (33%)	5,253 (67%)	7,823 (41%)			
Calves	1,353 (60%)	908 (40%)	2,261 (12%)			
Cows	3,922 (93%)	280 (7%)	4,202 (22%)			
Heifers	2,746 (58%)	1,956 (42%)	4,702 (25%)			
<i>Sub-total</i>	<i>10,591 (100%)</i>	<i>8,397 (100%)</i>	<i>18,988 (100%)</i>			
ii) Production						
Beef	5,072 (53%)	4546 (47%)	9618 (51%)	72.9	1	<0.001
Dairy	5,519 (59%)	3851 (41%)	9370 (49%)			
<i>Sub-total</i>	<i>10,591 (100%)</i>	<i>8,397 (100%)</i>	<i>18,988 (100%)</i>			

463 **Table 2** Frequency (total number of days) of cattle use of **a)** field arrangements (single verses multiple fields per
464 patch) and **b)** field contiguity (home verses fragmented) for $n=18$ farms.

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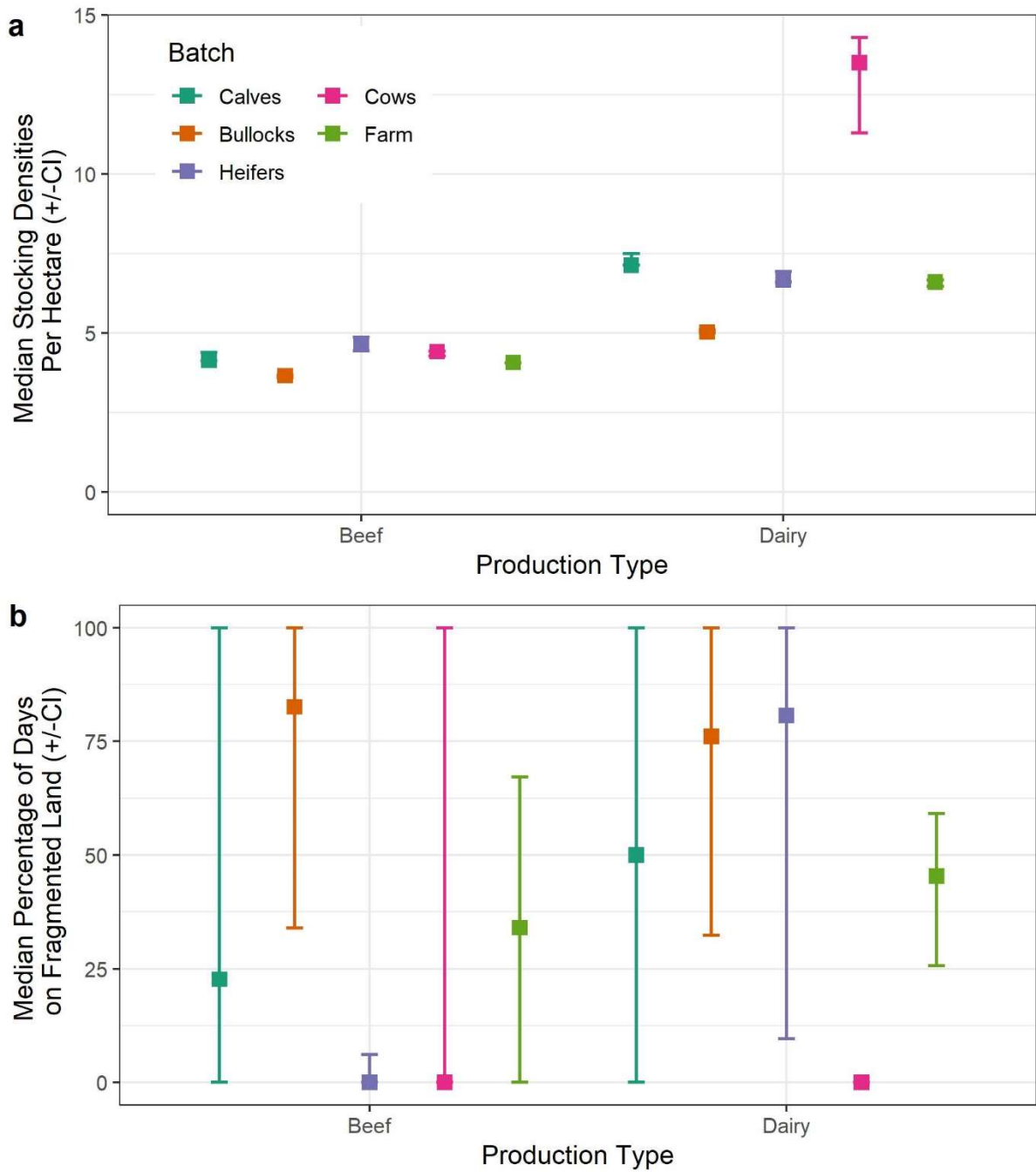
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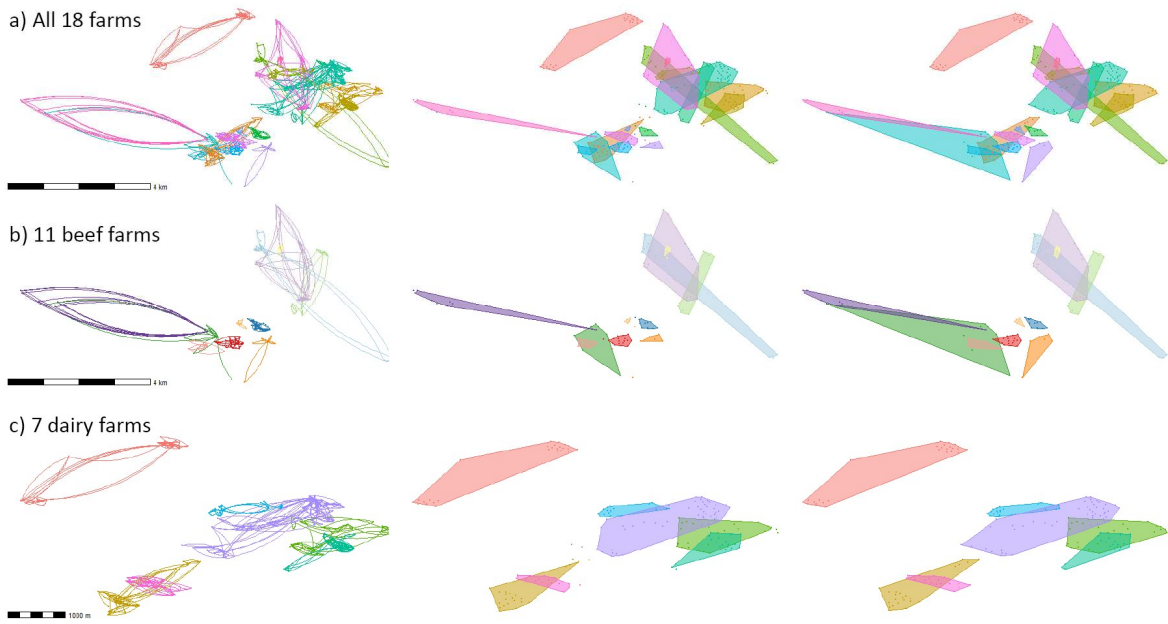
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478 **Fig. 1** Median \pm 95% confidence intervals (CI) for **a)** stocking densities for each cattle life history stage
 479 (number of cattle present per hectare) and **b)** percentage of days spent on fragmented land during the grazing
 480 season

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483 **Fig 2** Networked cattle movements (lines) between centroid of fields (dots) for **a)** $n=18$ all farms, **b)** $n=11$ beef
 484 farms and **c)** $n=7$ dairy farms (left column) and corresponding Minimum Convex Polygons (both 95% and 100%)
 485 for fields within the same categories. Each colour represents an individual herd.

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STROBE 2007 (v4) checklist of items to be included in reports of observational studies in epidemiology*

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Checklist for cohort, case-control, and cross-sectional studies (combined)

Section/Topic	Item #	Recommendation	Reported on page #
Title and abstract	1	(a) Indicate the study's design with a commonly used term in the title or the abstract	3
		(b) Provide in the abstract an informative and balanced summary of what was done and what was found	3
Introduction			
Background/rationale	2	Explain the scientific background and rationale for the investigation being reported	4-6
Objectives	3	State specific objectives, including any pre-specified hypotheses	6
Methods			
Study design	4	Present key elements of study design early in the paper	6
Setting	5	Describe the setting, locations, and relevant dates, including periods of recruitment, exposure, follow-up, and data collection	6-7
Participants	6	(a) <i>Cohort study</i> —Give the eligibility criteria, and the sources and methods of selection of participants. Describe methods of follow-up <i>Case-control study</i> —Give the eligibility criteria, and the sources and methods of case ascertainment and control selection. Give the rationale for the choice of cases and controls <i>Cross-sectional study</i> —Give the eligibility criteria, and the sources and methods of selection of participants	6-7
		(b) <i>Cohort study</i> —For matched studies, give matching criteria and number of exposed and unexposed <i>Case-control study</i> —For matched studies, give matching criteria and the number of controls per case	
Variables	7	Clearly define all outcomes, exposures, predictors, potential confounders, and effect modifiers. Give diagnostic criteria, if applicable	7-8
Data sources/ measurement	8*	For each variable of interest, give sources of data and details of methods of assessment (measurement). Describe comparability of assessment methods if there is more than one group	6-7
Bias	9	Describe any efforts to address potential sources of bias	7

Study size	10	Explain how the study size was arrived at	6-7
Quantitative variables	11	Explain how quantitative variables were handled in the analyses. If applicable, describe which groupings were chosen and why	8
Statistical methods	12	(a) Describe all statistical methods, including those used to control for confounding	8
		(b) Describe any methods used to examine subgroups and interactions	
		(c) Explain how missing data were addressed	
		(d) <i>Cohort study</i> —If applicable, explain how loss to follow-up was addressed <i>Case-control study</i> —If applicable, explain how matching of cases and controls was addressed <i>Cross-sectional study</i> —If applicable, describe analytical methods taking account of sampling strategy	7
		(e) Describe any sensitivity analyses	
Results			
Participants	13*	(a) Report numbers of individuals at each stage of study—eg numbers potentially eligible, examined for eligibility, confirmed eligible, included in the study, completing follow-up, and analysed	8-10, Table 1
		(b) Give reasons for non-participation at each stage	8
		(c) Consider use of a flow diagram	
Descriptive data	14*	(a) Give characteristics of study participants (eg demographic, clinical, social) and information on exposures and potential confounders	8-10
		(b) Indicate number of participants with missing data for each variable of interest	
		(c) <i>Cohort study</i> —Summarise follow-up time (eg, average and total amount)	8
Outcome data	15*	<i>Cohort study</i> —Report numbers of outcome events or summary measures over time	8-10
		<i>Case-control study</i> —Report numbers in each exposure category, or summary measures of exposure	
		<i>Cross-sectional study</i> —Report numbers of outcome events or summary measures	

Main results	16	(a) Give unadjusted estimates and, if applicable, confounder-adjusted estimates and their precision (eg, 95% confidence interval). Make clear which confounders were adjusted for and why they were included	8-10, Table 1 and 2
		(b) Report category boundaries when continuous variables were categorized	
		(c) If relevant, consider translating estimates of relative risk into absolute risk for a meaningful time period	
Other analyses	17	Report other analyses done—eg analyses of subgroups and interactions, and sensitivity analyses	
Discussion			
Key results	18	Summarise key results with reference to study objectives	8-10
Limitations	19	Discuss limitations of the study, taking into account sources of potential bias or imprecision. Discuss both direction and magnitude of any potential bias	8-10
Interpretation	20	Give a cautious overall interpretation of results considering objectives, limitations, multiplicity of analyses, results from similar studies, and other relevant evidence	12
Generalisability	21	Discuss the generalisability (external validity) of the study results	12-13
Other information			
Funding	22	Give the source of funding and the role of the funders for the present study and, if applicable, for the original study on which the present article is based	13

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491 *Give information separately for cases and controls in case-control studies and, if applicable, for exposed and unexposed groups in cohort and cross-sectional studies.

492 **Note:** An Explanation and Elaboration article discusses each checklist item and gives methodological background and published examples of transparent reporting. The

493 STROBE checklist is best used in conjunction with this article (freely available on the Web sites of PLoS Medicine at <http://www.plosmedicine.org/>, Annals of Internal

494 Medicine at <http://www.annals.org/>, and Epidemiology at <http://www.epidem.com/>). Information on the STROBE Initiative is available at www.strobe-statement.org.

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