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On following behaviour as a mechanism for collective movement

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6 Abstract

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During collective movement, animals display a wide variety of mechanisms 7 to maintain cohesion. In some species, indviduals rely mainly on following 8 their direct predecessor, thereby forming spectacular processions of individ-9 uals in single file. Despite being the simplest case of following behaviour, it 10 is largely absent from the theoretical literature on collective migrations. The 11 objective of this study is to quantify the efficiency of following the predeces-12 sor, in terms of ensuring cohesion. The situation we consider is a sequence 13 of individuals facing a bifurcation. The choice between left and right is influ-14 enced by the choice of the predecessor. First, we model this situation with 15 a two-state Markov chain with a symmetric transition matrix. Cohesion is 16 quantified as the expected number of individuals on either side, and the ex-17 pected number of consecutive followers. Although cohesion increases with 18 the probability of following, it remains surprisingly low unless the probabil-19 ity is almost equal to one. Furthermore, cohesion decreases with group size 20 regardless of the probability of following. Then, we generalise our model to 21 situations in which individuals have a preference for one of the two choices 22 (asymmetric transition matrix). For some parameter sets, the tendency to 23

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follow each other leads a large fraction of the individuals to the non pre-24 ferred side. Moreover, this fraction increases with the total population size. 25 Finally, we include the possibility to follow N individuals. This provides the 26 link between our model and other collective migration models. If enough in-27 dividuals are perceived, the results shift from symmetrical (low cohesion) to 28 asymmetrical (high cohesion) distribution of the individuals. All in all, our 29 results suggest that following the direct predecessor must be complemented 30 with other cohesive behaviours (involving the perception of more individuals 31 or a navigation system) to guarantee its efficiency. We discuss our findings 32 in the context of the different following behaviours covered in the literature. 33 *Keywords:* 34

³⁵ migration, collective decision making, group cohesion, aggregation, Markov
 ³⁶ chains

37 1. Introduction

Social decision making is common among a wide range of species. In those species, when an individual has to make a choice between a set of options, its decision is influenced by the decision of conspecifics. When imitation¹ in the broad sense is involved, it can lead to complex group behaviour (Deneubourg and Goss, 1989). This often results in some level of cohesion and/or synchronisation (Sumpter, 2006). The range of perception of each

¹By imitation we mean that an individual's probability of choosing one option is increased if other individuals already chose that option. Depending on the specific context it can be referred to as allelomimesis, social facilitation, public information... but here we use it in a very broad sense.

individual determines in the number of individuals by whom it will be influenced. This factor can therefore be of key importance in the resulting group
cohesion.

Many types of decisions can be socially influenced. An example is tem-47 poral decisions (when to do something). For instance, pedestrians are influ-48 enced by other pedestrians as they decide when to cross the road (Faria et al., 49 2010). Sometimes, the immitation of others in temporal decisions can lead 50 to synchronisation. There are countless examples ranging from flashing in 51 fireflies (Buck, 1988, Ramırez Ávila et al., 2003) to synchronised alternation 52 between feeding and vigilance bouts in sheep (Michelena et al., 2006, Gau-53 trais et al., 2007) or to synchronized clapping in human audiences (Neda et 54 al., 2000a, 2000b). Social influence has also been widely studied in the con-55 text of spatial decisions, such as where to forage, where to settle or where to 56 breed (Danchin et al., 2004). Finally, there are directional decisions (which 57 direction to take). Sometimes, directional decisions and temporal decisions 58 are not dissociated. Sheep for example, tend to move when an other sheep 50 moves, in the same direction as the initiator (Pillot et al., 2010). In this 60 paper, however, we focus only on directional decisions, with no particular 61 regard for the context of the displacement (where are they going to or why). 62 More precisely, our interest lies in the mechanisms involved in the decision 63 at the individual level, and their consequences at the collective level. 64

This ambition to link individual behaviour with collective movement patterns lies at the heart of a number of mathematical modelling studies. The behaviour at stake is, on the whole, following the other individuals. Understandably, the different biological realities require different types of models

(for a review see Giardina, 2008). In flocks of birds or schools of fish, each
individual repeatedly updates its position according to the position of others
(Couzin and Krause, 2003). The majority of models of schooling or flocking have been individual based, self-propelled particle (SPP) models (Okubo
1986, Viscek et al., 1995, Czirok et al., 1997, Couzin et al. 2002, Grégoire
and Chaté, 2004).

Another important section of the literature on collective movement con-75 cerns situations in which the directional decisions are taken sequentially by 76 each individual, rather than simultaneously. This is typically the case of 77 species that tend to move in single file such as social caterpillars, ants or army 78 worms. Some fish have also been observed moving heads-to-tails (Gudger, 79 1944). Finally, single files occur in any species constrained to move in a se-80 quence due to the geometry of the environment, such as sheep in mountain 81 paths or cows in cattle-handling facilities (Grandin, 1980). The question of 82 social influence on directional choices is typically studied with binary choice 83 set-ups in which a bifurcation presents two identical options. The outcome 84 of such experiments is either a binomial bell-shaped distribution (no social 85 influence) or a U-shaped distribution, with most experiments resulting in the 86 collective selection of one of the two options (social influence). This simple 87 experiment has put forward collective movement in species as diverse as ants 88 (Deneubourg et al., 1990), fish (Ward et al., 2008), dust mites (Mailleux 89 et al., 2011), sheep (Michelena et al., 2010), caterpillars (Dussutour et al., 90 2008), spiders (Jeanson et al., 2004, Saffre et al., 1999) and earth worms 91 (Zirbes et al., 2010). 92

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The mathematical models used to describe such collective choices are

often individual-based simulations coding for the particular mechanism in-94 volved, for example dragline attachment in the case of spiderlings (Jeanson 95 et al., 2004). General models covering a wide range of situations have seldom 96 been formulated. One of these general models, hereafter referred to as the 97 trail model (Deneubourg et al., 1990) expresses the probability of choosing 98 one option as an increasing (potentially non-linear) function of the number 99 of individuals who previously chose that option (see section 4 for more details 100 on this probability function). The major prediction of this model is that in-101 dividuals will distribute themselves asymmetrically between the two options, 102 if the response function is non-linear. Another, perhaps more abstract model 103 is Polya's urn model, which makes similar predictions to the trail model (for 104 given parameter sets). One of the assumptions of these models is that indi-105 viduals are able to perceive the choice of every preceding individual. This 106 assumption is justified in the case of many species, which use modification 107 of the environment (e.g., laying a physical or chemical trail) as means of 108 communication. As long as the modification is permanent at the time-scale 100 considered (no or little evaporation of the trail), then the assumption holds. 110 However, with means of communication such as sight or hearing (e.g. in 111 birds or mammals), the assumption ceases to be valid as soon as the total 112 number of preceding individuals exceeds the range of perception. 113

The extreme case is when perception is limited to one individual, for example when animals in a sequence perceive others through tactile cues (contact). This is the case of processionary caterpillars (Fitzgerald, 2003), earthworms (Zirbes et al., 2010), spiny lobsters (Herrnkind, 1969, Berill 1975), or some self-assembling robots (Levi and Kernbach, 2010). In these species

each animal follows its direct predecessor. Despite being the simplest case
possible of following behaviour, it has never been studied from a theoretical
point of view. Some car traffic models, called "Follow the leader" (Gazis et
al., 1960), consider the influence of the direct predecessor in decisions about
speed and acceleration, but not about direction.

The question we address here, is therefore how does the number of in-124 dividuals perceived affect the cohesion of the moving group? We begin by 125 developing a model for the simplest case: following a single preceding indi-126 vidual. We then extend the model to cover situations in which individuals 127 follow an environmental cue as well as the preceding individual. Finally, the 128 link between our model and the trail model is provided by a general model 129 allowing the perception of N preceding individuals. Our results are discussed 130 in terms of the efficiency of the different following behaviours covered in the 131 literature. 132

133 2. Following the previous individual

The theoretical set-up is a path leading to a bifurcation. n individuals have to take the path and choose between left and right, one after the other. Each individual's information is limited to its direct predecessor's choice. It has a probability p of following its predecessor, and a probability 1 - p of taking a different path from its predecessor (p > 0.5 by definition).

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The process is memoryless, in the sense that every choice depends exclusively on the choice of the predecessor. The choices of everyone else that passed before that previous individual do not have any influence on the focal

individual's choice. Therefore, this system can be assimilated to a two-state
Markov chain, where the two states are the possible choices of the individuals
(left or right). Since the probability of following the predecessor whether he
took left or right is the same, the transition matrix is symmetric:

$$\mathbf{T} = \begin{pmatrix} p & 1-p \\ 1-p & p \end{pmatrix} \tag{1}$$

The entry T(i, j) in the transition matrix represents the probability of transition from state *i* to state *j*. In other words, it is the probability of choosing side *j*, given that the previous individual took side *i*.

The question we address here is whether this mechanism is efficient in 150 terms of ensuring cohesion. We suggest two ways of answering this question. 151 Firstly, we calculate the proportion of individuals expected to end up on the 152 side with the most individuals (hereafter referred to as the "winning side"). 153 A very cohesive species would have an expected fraction of individuals on 154 the winning side that is relatively high, whereas a poorly cohesive species 155 would have an expected fraction approaching 0.5. Secondly, we calculate the 156 number of individuals that are expected to take the same side successively, 157 thereby forming a chain of followers. The expected length of those chains 158 represents an accurate estimate of cohesion because all of those successive 159 followers will end up in the same place. 160

¹⁶¹ Expected fraction of individuals on the winning side

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Let n be the total number of individuals that face the binary choice. First, we need to calculate the expected absolute number of individuals on

the winning side. This can be done by using the definition of the expected value in probability theory $E(X) = \sum_{k=n/2}^{n} k \cdot P(X = k)$, where k can be any possible number of individuals on the winning side and ranges from n/2to n. P(X = k) is the probability of having k individuals on the winning side.

This probability is difficult to calculate, because all of the possible se-170 quences leading to k individuals on a given side do not necessarily have the 171 same probability (see an example in Fig. 1). Therefore, we calculate this 172 probability P(X = k) by grouping all of those sequences in groups with the 173 same number of switches (an individual taking the opposite side to his prede-174 cessor). The reason for grouping the sequences in such a way is simple: the 175 probability of any given sequence depends only on the number of switches a176 and on the total number of individuals n. Therefore all of the sequences in 177 each group have the same probability, that is $P(X = k) = 0.5 \cdot p^{a-1-s}(1-p)^a$. 178 The sum of the probabilities of all those possible sequences gives us the prob-179 ability of having k individuals on the winning side: 180

$$P(X=k) = \sum_{a=0}^{2(n-k)} 0.5 \cdot p^{n-1-a} (1-p)^a f(n,k,a)$$
(2)

The function f(n, k, a) gives the number of sequences leading to k out of nindividuals on the winning side in which a switches took place. We propose a way to calculate this function f(n, k, a). The sequences can have two possible configurations depending on the side chosen by the first individual (Fig. 2). In the first configuration (a), the number of individuals who chose the left are divided into three subgroups, and in configuration (b) into two subgroups. For the individuals who took the right, it is the opposite.

Fig. 3 shows the exhaustive enumeration of all the sequences in a concrete example involving 4 switches. In that example, the number of sequences of type (a) is simply the number of different ways there are to make 3 subgroups with 3 black elements multiplied by the number of ways there are to make 2 subgroups out of 4 white elements (same reasoning for type (b)).

If we define G(x, y) as the number of ways to make x subgroups out of y elements, then the total number of sequences f(n, k, a) can be calculated with the expression in Eq. 3. We give some details about this G(x, y) function in appendix A. This expression was verified with simulations (appendix C).

$$f(n,k,a) = 2 G\left(\left\lceil \frac{a+1}{2} \rceil, n-k \right) \cdot G\left(\left\lfloor \frac{a+1}{2} \rceil, k \right) + 2 G\left(\left\lfloor \frac{a+1}{2} \rceil, n-k \right) \cdot G\left(\left\lceil \frac{a+1}{2} \rceil, k \right) \right) \right)$$
(3)

With this formula, we calculated the expected fraction of individuals on 197 the winning side, and how it varies with the total number of individuals pass-198 ing through the binary choice set-up. This was done for different values of 199 the probability of following. As is shown in Fig. 4, the system tends towards 200 a symmetric distribution of the individuals between the two sides, for prob-201 abilities of following that are different from 1. In other words, even though 202 individuals follow one another, the fraction of individuals on either side tends 203 towards 0.5. In Markovian terms, this is easily verified by calculating the sta-204 tionary distribution of the system, that is, when the probabilities of finding 205 the system in any given state do not change over time. At the stationary 206 distribution the probability of finding the system in state i is called $\pi(i)$ and 207

is given by the formula $\pi(i) = T(j,i)/(T(j,i) + T(i,j))$, where T(i,j) is the entry of the transition matrix (Eq. 1) containing the probability of taking side j, given that the previous individual took side i, $(i \neq j)$. In our case, $\pi(\text{left}) = \pi(\text{right}) = 0.5$.

²¹² Expected number of consecutive followers

The individuals that follow each other uninterruptedly form a "chain" of individuals that all end up on the same side. The mean length of those chains gives us an estimate of the cohesiveness of the species: the larger the mean chain length, the greater the cohesion. Note that this is true in general, even for choices that are not binary, which are more representative of the actual nature of most species' environment. The expected size E(C) (or mean size) of these chains of followers is:

$$E(C) = \sum_{c=1}^{n-1} cp^{c-1}(1-p) + np^{n-1}$$
(4)

As is shown in Fig. 5A, this expected chain length increases with the 220 total number of individuals up to a certain point. From there onwards it is 221 independent of the total number of individuals. For an infinite number of 222 individuals, it depends exclusively on the probability of following, according 223 to the function E(C) = 1/(1-p) (Fig. 5B). It is noteworthy that the ex-224 pected chain length is hardly affected by following the predecessor, except for 225 probabilities of following superior to 0.85. For example, in a group of indi-226 viduals that follow each other with a probability of 0.85, the expected number 227 of followers would be of 6.7, whereas in a group moving independently the 228 expected number would be 2. Chains of more than 20 individuals are only 229

²³⁰ obtained with probabilities of following superior to 0.95.

231

²³² 3. Following the previous individual and an environmental cue

So far, we have considered the probability of following to be indepen-233 dent of the direction taken by the preceding individual. However, in a more 234 general case, we can include a bias or an environmental cue that causes the 235 individuals to prefer one side over the other. The preferred side will hereafter 236 be referred to as the biased side. For the first individual, let the probabil-237 ity of taking the biased side be b. From then onwards, we can consider a 238 two-state Markov process in which the Markovian transition matrix becomes 239 asymmetric: 240

$$\mathbf{T} = \begin{pmatrix} p_B & 1 - p_B \\ 1 - p_{NB} & p_{NB} \end{pmatrix}$$
(5)

In this transition matrix, p_B and p_{NB} are the probabilities of following the 241 preceding individual towards the biased and the non-biased side respectively. 242 For the system to be consistent with the scenario we propose, the parame-243 ters require some constraints: (1) the probability of following an individual 244 towards the biased side p_B is the highest of all probabilities (including b); 245 (2) the probability of taking the non-biased side, even though the previous 246 individual took the biased side p_{NB} is the lowest of all probabilities (3) if the 247 probability of following towards the non-biased side is higher than 0.5, the 248 attraction to others is higher than the attraction to the environmental cue 249 causing the bias (and vice-versa). 250

In this situation, the expected number of individuals on the biased side is:

$$E(X) = \sum_{k=0}^{n} k \cdot P(K=k) \tag{6}$$

The probability of having k individuals on the biased side P(K = k) is explained in appendix B.

We calculated the expected proportion of individuals on the biased side for different values of the three parameters p_B , p_{NB} and b. Unsurprisingly, the majority of individuals always ends up on the biased side. The expected proportion tends towards a steady value as the total number of individuals increases (Fig. 6). This steady value depends on p_B , p_{NB} (Fig. 7) and equals the probability of finding the system in state B, once the stationary distribution is reached, that is $\pi(B)$:

$$\pi(B) = (1 - p_{NB})/(2 - p_B - p_{NB})$$
(7)

This value can be greater than the bias (amplification of the bias) or lower 262 than the bias (alleviation of the bias), depending on the values of p_B and p_{NB} 263 (Fig.6). This means that for certain parameter sets (satisfying $\pi(B) < b$), 264 the tendency to follow each other can lead a large fraction of the popula-265 tion to the non-preferred side. At least, this fraction is larger than it would 266 have been if the individuals were not influenced by one another. In these 267 situations, the fraction on the non-preferred side increases with the total 268 population. Another interesting feature is that for certain values of p_B and 269 p_{NB} , satisfying $p_{NB} = (2b - bp_B - 1)/(b - 1)$, the expected fraction of in-270 dividuals on the biased side is b. This result is identical to what it would 271

be, had they chosen independently from one another in a biased set-up. In other words, they can be very strongly influenced by each other, and to the observer who looks only at the end result, rather than at the dynamics, it may seem as if they had just followed a bias.

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$_{277}$ 4. Following the N previous individuals

So far, we have assumed that the range of perception of indidviduals was limited to their direct predecessor. In order to allow for larger perception ranges, we generalise our model by defining N, the number of predecessors perceived by the focal individual. The situation we have discussed so far is in fact an extreme case where N = 1, the other extreme being the ability to perceive all of the preceding individuals (N = n), which is the case of the trail model when the trail does not evaporate.

Let us go back to a non-biased environment. Each individual perceives N preceding individuals, and chooses preferentially the side where most of them went. The choice is made according to the choice function of the trail model (Deneubourg et al., 1990):

$$p_L = \frac{(\kappa + N_L)^{\epsilon}}{(\kappa + N_L)^{\epsilon} + (\kappa + N_R)^{\epsilon}} \quad \text{and} \quad p_R = \frac{(\kappa + N_R)^{\epsilon}}{(\kappa + N_L)^{\epsilon} + (\kappa + N_R)^{\epsilon}} \tag{8}$$

In this choice function, p_L and p_R are the probabilities of choosing the left and the right, respectively. N_L and N_R are the number of perceived individuals that went to the left and to the right, respectively. Parameter κ represents the inherent attractiveness of each side. For high values of κ , the

branches are so attractive that the attractiveness added by the social term 293 $(N_L \text{ or } N_R)$ does not greatly influence the probability. Therefore, when κ 294 tends towards infinity, both probabilities tend towards 0.5, and the resulting 295 probability distribution is the binomial distribution. The exponent ϵ gives 296 an estimate of the system's amplification potential. If $\epsilon = 0$, the probability 297 distribution is also binomial. This choice function was previously used to 298 model trail following in ants (Deneubourg et al., 1990) and in dust mites 299 (Mailleux et al. 2011). Actually, in the latter study the function used is a 300 slightly modified version, but it shares the same properties, namely the gener-301 ation of great asymmetries in the distribution of individuals when $\epsilon > 1$. For 302 this trail model, it is possible to calculate the exact probability distribution, 303 that is the probability of all possible outcomes, expressed as the probability 304 p(l,r) of having l individuals to the left and r individuals to the right. The 305 initial probabilities are p(0,1) = p(1,0) = 0.5. The probabilities of having a 306 negative number of individuals on either side (p(x, y) with x < 0 or y < 0) 307 are naturally equal to zero. From there onwards the probability distribution 308 is given by: 309

$$p(l,r) = p(l-1,r)p_L + p(l,r-1) * p_R$$
(9)

For numbers of perceived individuals lower than the total number of individuals, we used Monte Carlo simulations to generate the probability distributions. This was done for various values of the parameters N, κ , and ϵ . Fig. 8 shows a sample of the the parameter sets we used. As expected, the lower the inherent attractiveness of the branches κ , the higher the asymmetry. Also, when $\epsilon = 1$, the average number of individuals on the winning

side is relatively low, even when many preceding individuals are perceived (high values of N). Very low values of κ ($\kappa \leq 1$) might make the expected number on the winning side higher, but this is due to a high probability of every individual taking the same side. In fact, for amplification to appear, the exponent ϵ has to be higher than one. As Fig. 9 shows, the asymmetries increase with the number of perceived individuals (N).

322 5. Discussion

In this paper we study the efficiency (in terms of group cohesion) of fol-323 lowing the preceding individual. This question is largely absent from the 324 theoretical literature on collective movements and socially influenced deci-325 sions. As we show in this paper, the question is far less trivial to answer 326 than to ask. Our main results may seem somewhat counter-intuitive: cohe-327 sive behaviour at the individual level (following the preceeding individual) 328 does not lead to cohesion at the collective level. Eventually, it leads to a 320 homogeneous dispersal of the group. Some other results are more intuitive, 330 such as the positive relationship between cohesiveness and the probability of 331 following, or the cohesion obtained from increasing the number of perceived 332 individuals, but they remain worth quantifying. By providing a quantitative 333 description of the relationship between parameters and variables, our model 334 offers a set of predictions that experimenters can compare their results with. 335 This study concerns binary-choice experiments made on groups, which 336 are extremely popular in ethology and psychology. The set-up of these ex-337 periments (reviewed in Jeanson and Deneubourg, 2009) can either present 338 two identical options (as in Mailleux et al. 2011, or Zirbes et al., 2010) or 339

two different options (as in Canonge et al., 2009 or Ward et al. 2008). The 340 former is an elegant way of putting forward social influence on the choice 341 of individuals. The latter can be used to study the interplay between social 342 interaction and individual preference. Our predictions cover both situations, 343 and can provide insight into the interpretation of such experimental results. 344 For example, a homogeneous distribution of individuals between two identi-345 cal options is typically interpreted as an absence of social interaction. Here, 346 we show that a homogeneous distribution can happen, even in the presence 347 of strong social influence, provided that the perception range is limited (by 348 the nature of the communication used, or by the experimental set-up) to 349 a relatively small number of predecessors. Therefore, questions on percep-350 tion mechanisms and population density in the set-up must be taken into 351 account. Furthermore, the models we propose can be used to validate (or 352 invalidate) hypotheses on the mechanisms by which animals influence one 353 another. Take for example the hypothesis that individuals follow each other 354 by contact. It can be tested by calculating the probability of following the 355 direct predecessor, based on the behaviour of the second individual. From 356 that probability and our model it is possible to predict the behaviour of the 357 subsequent individuals (Fig. 4). The comparison between this prediction 358 and the experimental results could confirm (or invalidate) the hypothesis. 359 Another conclusion that can be drawn from our results is that it is useful 360 to calculate the mean length of chains of consecutive followers. Indeed, this 361 variable can be crucial to distinguish between a situation in which there is 362 no social interaction and one in which each individual follows its the direct 363 predecessor. It can also be used to make the difference between following the 364

³⁶⁵ predecessor and following a trail.

The model we study in this paper is general enough to cover a wide range 366 of situations. Animals that use tactile cues to perceive and follow their pre-367 decessor, such as processionary caterpillars (Fitzgerald, 2003), spiny lobsters 368 (Herrnkind, 1969) or earthworms (Zirbes et al., 2010) are the obvious exam-369 ples. But our predictions can extend to self-assembling robots designed to 370 form chains, or even to humans in situations where information is limited to 371 one's direct predecessor, such as some situations in car traffic. In fact, the 372 model can apply to socially influenced decisions that are not directional deci-373 sions. For example, it is well known that conformism can make humans very 374 influenced by other people's decisions (Asch, 1951). Therefore, situations 375 in which humans have to make decisions sequentially and can perceive the 376 decisions of others (typically consumer behaviour) can be studied through 377 the theoretical perspective we propose. 378

More generally, the mathematical quetions raised in this study are rele-379 vant to any situation that can be assimilated to a two-state Markov chain. 380 These situations can be found even outside the context of decisions. For ex-381 ample, a classical model to predict precipitation is a two-state Markov chain 382 (Todorovic et al., 1975). Other studies have tackled some of the mathemat-383 ical questions raised here in completely different contexts. The question of 384 the exact expression for the probability distribution discussed in the first 385 section (Eqs. 2 and 3) is explored from a purely mathematical point of view 386 in the context of a correlated random walk through the integers (Renshaw 387 and Henderson 1981). The more general case of the asymmetric environment 388 was studied from a purely abstract perspective by Gabriel (1959). The exact 380

expressions derived in these two studies are in essence very similar to ours. However, our objective here is to use these mathematical explorations to infer properties to a biological system, that is to a group of animals in motion.

One interesting feature of our results is the fragility of the cohesion of 393 groups of followers. Indeed, we show that for probabilities of following 394 p < 0.85, the expected chains are not drastically longer than they would 395 be if individuals acted independently of one another (6.7 versus 2). Fur-396 thermore, what we define as the probability of following p, contains in fact 397 the probability of perceiving multiplied by the probability of following the 398 preceding individual. In our theoretical set-up, we have implicitly considered 399 the probability of perceiving the preceding individual to be equal to one. In 400 natural situations, where the linear density of individuals can be low, this 401 assumption does not necessarily hold. This potentially decreases the cohe-402 sion of groups. In addition to this, even for high probabilities of perceiving 403 and of following (different from one), large groups will end up homogeneously 404 distributed across the options. Cohesion actually decreases with the size of 405 the group. 406

Not only does cohesion decrease with group size, but in some cases ac-407 curacy also decreases with group size. Indeed we show that in situations in 408 which individuals have a preference for one of the options, a large fraction 409 can end up choosing the non-preferred option, if the probability of following 410 is large enough. More importantly, we show that this fraction increases with 411 group size. This is directly opposed to the outcome of other types of collec-412 tive decision making models, such as the trail model (Deneubourg 1990) or 413 quorum responses (Sumpter and Pratt, 2009). In the latter, cohesiveness and 414

most often accuracy increase with group size. The key difference is that in 415 these models, there are non-linearities that arise from positive feedbacks. In 416 our memoryless situation, it is impossible for amplification to happen, hence 417 the importance of perceiving more than one individual, and of non-linear re-418 sponse functions to maintain cohesion. Indeed, as we show in section 4, stable 419 asymmetries in the distribution happen only when the number of perceived 420 individuals increases and when the exponent of the probability function is 421 higher than one. 422

Nevertheless, the theoretical results we present must be put into perspec-423 tive with the biological reality that they represent. First of all, the cohesive 424 species that we have talked about display behaviours that potentially increase 425 the probability of following, or more precisely the probability of perceiving 426 the predecessor. For example, they can synchronise departures, which in-427 creases the potential linear density. It is true that the larger the group, 428 the lower the cohesion. Nevertheless, the groups in some cases are small 420 enough to ensure sufficient cohesion. The smallest groupe size are groups of 430 two, which is the case of ant recruitment tandems. This type of recruitment 431 requires constant antennal contact between the following ant and the lead-432 ing ant's abdomen (Franks and Richardson, 2006, Möglich et al 1974). It is 433 very efficient, and most of the time both ants end up in the same food source. 434 Caterpillars move in larger groups: Edwards (1910) reports processions of up 435 to 300 individuals. Spiny lobsters also seem to migrate in large groups: every 436 autumn, in the Bahamas, thousands of Spiny lobsters (Herrnkind, 1969) mi-437 grate to the South-West in parallel chains of 3 to 30 individuals (Herrnkind 438 and Cummings, 1964). It is necessary for them to rely on complementary 439

mechanisms to ensure cohesion. caterpillars use silk and chemical cues as a
trail (Fitzgerald, 2003), and spiny lobsters are capable of true navigation in
a magnetic field (Boles and Lohmann, 2003). However it has been shown
in both cases that tactile cues are the most important cues for maintaining
cohesion (Fitzgerald 2003, Herrnkind, 1969).

In this paper we formulate a model that provides answers to questions concerning the efficiency of following behaviour. By studying the simplest case possible of following behaviour, we put forward three important factors in ensuring cohesion: the number of perceived individuals, the potential for amplification, and group size.

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456 Appendix A. The number of combinations resulting in A switches

Although intuitive, the G(x, y) function requires a few definitions in order to be calculated correctly:

- There is one way to make 0 subgroups with 0 elements

- There is no way to make x subgroups with y elements if x > y
- There is no way to make 0 subgroups with y elements if $y \neq 0$

- If x and/or y are negative, G(x, y) = 0

463 - Otherwise,
$$G(x, y) = C_{x-1}^{y-1} = \frac{(y-1)!}{(y-x)!(x-1)!}$$

⁴⁶⁴ Appendix B. Generalisation to any two-state Markov chain

Let $P_{(a)}(K = k)$ be the probability of having k individuals on the biased side given that the first one chose the biased side and $P_{(b)}(K = k)$ the probability, given that the first one chose the non-biased side. The probability P(K = k) of having k individuals on the biased side after n individuals have passed is the sum of those two probabilities.

$$P(K = k) = P_{(a)}(K = k) + P_{(b)}(K = k)$$

$$P_{(a)} = b \sum_{a=0}^{2(n-k)} p_B^{k-\left\lceil \frac{a+1}{2} \right\rceil} p_{NB}^{n-k-\left\lfloor \frac{a+1}{2} \right\rfloor} (1-p_B)^{\left\lceil \frac{a}{2} \right\rceil} (1-p_{NB})^{\left\lfloor \frac{a}{2} \right\rfloor}$$

$$G\left(\left\lceil \frac{a+1}{2} \right\rceil, k\right) \cdot G\left(\left\lfloor \frac{a+1}{2} \right\rfloor, n-k\right)$$

$$P_{(b)} = (1-b) \sum_{a=0}^{2(n-k)} p_B^{k-\left\lfloor \frac{a+1}{2} \right\rfloor} p_{NB}^{n-k-\left\lceil \frac{a+1}{2} \right\rceil} (1-p_B)^{\left\lfloor \frac{a}{2} \right\rfloor} (1-p_{NB})^{\left\lceil \frac{a}{2} \right\rceil}$$

$$G\left(\left\lfloor \frac{a+1}{2} \right\rfloor, k\right) \cdot G\left(\left\lceil \frac{a+1}{2} \right\rceil, n-k\right)$$

⁴⁷⁰ The validity of this expression was verified (appendix C).

⁴⁷¹ Appendix C. Verification of the exact expressions for the probabil⁴⁷² ity distributions

We conducted Monte-Carlo simulations in order to verify that the exact expressions for the probability distributions in sections 1 and 2 are correct. In

each simulation n virtual individuals were sequentially faced with the choice 475 between left and right. The probability of taking left and right depended on 476 the choice of the preceeding individual, and on the environment (homoge-477 neous or heterogeneous) according to the transition matrices (Eqs. 1 and 5). 478 For each individual, a random number was generated. If the random number 479 was lower than the probability of taking left, the individual's choice would be 480 left, otherwise it would be right. Repeating this simulation a million times 481 gave a good approximation of the frequency distribution of each possible out-482 come (analogous to a probability distribution). Fig. 10 shows that the exact 483 expression and the simulations converge to the same distribution. 484

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Figure 1: Sequences of 5 individuals leading to 3 individuals on the right. The sequences of individuals are represented by a sequence of squares, where each square is the choice of one individual: black means it chose the left, white means it chose the right. The different sequences can have different probabilities (P), depending on the order of the squares. More precisely, the probability depends on the number of individuals that took a different path than their predecessor (number of switches a). the function f(n, k, a) gives the number of sequences in each subgroup.

Figure 2: If there are four switches, it means that the first one(s) go to the left, then some go to the right, then some more to the left again ending with some on the right, or the opposite starting with the right

Figure 3: All the sequences satisfying n = 7, k = 4, and a = 4, in which the winning side is the right. It has to be multiplied by two, in order to obtain the total number of sequences f(n, k, a)

Figure 4: Expected fraction of individuals on the winning side according to the total number of individuals. As the total number of individuals increases, the system tends towards symmetry (with the number on the winning side approaching 50%)

Figure 5: Mean length of the chains according to (A.) the total number of individuals and (B.) the probability of following the preceding individual. For every probability of following different from 1, after the total number of choosers exceeds a certain size, the mean length of the chain does no longer depend on the number of choosers. This dependency is strong with probabilities higher than 0.9.

Figure 6: Example of the evolution of the system as individuals pass. For a given value of the bias (b = 0.6) and a given value of the tendency to follow the previous individual towards the biased side ($p_B = 0.7$), the bias can be amplified (if $\pi(B) > b$, so $p_{NB} < 0.55$) or weakened ($p_{NB} > 0.55$)

Figure 7: Proportion of individuals on the biased side for an infinite total number of passing individuals, for various values of p_B and p_{NB} and an environmental bias of b = 0.6.

Figure 8: The expected percentage of individuals on the winning side after 2000 individuals have passed. Different sets of parameters were used in the Monte Carlo simulations. For each parameter set, we ran 2500 simulations and what appears in the figure is the average of the 2500 simulations.

Figure 9: The distribution of the simulation results for a total number of individuals n = 200, and for a fixed attractivity of each branch ($\kappa = 8$). The distributions change as we move from one extreme (N = 1) to the other (N = n = 200). For $\epsilon = 1$, the distribution remains bell shaped, but spreads around the most likely result (half of the individuals on each side). For $\epsilon = 2$, however, the distribution changes from bell-shaped to U-shaped. The most likely results become the asymmetric results (most individuals on one of the two sides).

Figure 10: Probability distributions calculated with exact expressions (solid line) and with simulations (histogram). The parameters taken were n = 40, p = 0.75, b = 0.6, $p_B = 0.8$ and $p_{NB} = 0.7$.





(a)

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(a)



total number of individuals





total number of individuals







number of perceived individuals N



Number of individuals on a given side



Highlights

- The potential for collective movement is evaluated in groups of indivduals that tend to follow their direct predecessor.
- This behaviour is quantified with a two-state Markov chain model.
- The results show that the cohesion resulting from following the direct predecessor is weak.
- A generalized model shows that the perception of more predecessors is needed to ensure cohesion.