1	Heritability and maternal effects on social attention during an
2	attention bias task in a non-human primate. Macaca mulatta
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45 Abstract

Social attention is fundamental to a wide range of behaviours in non-human primates. 46 47 However, we know very little about the heritability of social attention in non-human primates, and the heritability of attention to social threat has not been assessed. Here, we 48 49 provide data to begin to fill this gap in knowledge. We tested 67 female rhesus macaques, 50 Macaca mulatta, on an attention bias preferential looking task in which they viewed threatneutral face pairs. We recorded a number of looking time measures of social attention to 51 52 conspecific faces, and attention to conspecific threat faces specifically. In addition, we recorded levels of vigilant scanning in the social group. We quantified heritability and 53 54 maternal effects using pedigree information. Repeatabilities for social attention ranged 55 from 11% - 25%. Repeatability for attention to threat faces was 16%, with zero repeatability 56 for attention bias, calculated by subtracting duration of looking towards the neutral face from duration of looking towards the threat face (a common practice in the literature). 57 58 Heritabilities for social attention were 8% - 14%, with maternal effects 6% - 11%. Heritability for attention to threat was 10%, with maternal effect 4%. This is the first study that we are 59 aware of to test the heritability of attention to threat in a non-human primate. We discuss 60 these findings in light of understanding mechanisms underlying social behaviour in primates, 61 62 evolutionary pathways of social attention in humans, epidemiology of mental health issues such as anxiety, and potential for improving markers of animal emotion and wellbeing in 63 64 captivity.

65

66 Keywords

67 Social attention, attention bias, repeatability, heritability, maternal effects, primate

69 Introduction

70 Social attention underpins a wide range of behaviour in human and non-human primates, 71 allowing individuals to gain rich information about conspecifics – e.g. in relation to their 72 identity, status, behaviour, emotional state and intentions – which then guides decisions about subsequent social interactions (Klein et al., 2009). The importance of social attention 73 in primates' lives is reflected in the expansion of the visual cortex in this lineage (Barton, 74 1998) and by the existence of evolved brain mechanisms for processing social information 75 76 (Chang et al., 2013), including areas specialised for processing faces (Adolphs et al., 1996, 77 Chang et al., 2013) and for rapid assessment of threat (LeDoux, 1996). Studies with primates have demonstrated that patterns of social attention vary markedly between individuals 78 (Bethell et al., 2012, Deaner et al., 2005, Watson et al., 2015), but the causes of such 79 80 variation are not fully understood. Differences between individuals ('personality': Carter et 81 al., 2013, Dall et al., 2012, Dall et al., 2004, Gosling & John, 1999) may be driven by genetic 82 or environmental factors (Dingemanse et al., 2010, Nicolaus et al., 2012) and there is 83 evidence that some of this variation is heritable (Dochtermann et al. 2015; Drent et al. 2003, Winney et al. 2018). 84

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Work on the heritability of social attention in non-human primates has mostly explored this
phenomenon indirectly, for example through gene association studies (Coyne et al., 2015). A
small but growing number of non-human primate studies have more directly quantified
heritability of social attention by including measures of relatedness between individuals
(e.g. Blomquist & Brent, 2014). However, most have not accounted for other potentially

confounding causes of variation such as maternal effects (Cheverud & Wolf, 2009, 91 92 Maestripieri, 2009), and common environmental effects (Blomquist & Brent, 2014, Johnson et al., 2015,). Maternal effects are here defined as influence of the mother on the 93 phenotype of the offspring - prenatal and/or postnatal - aside from those explained by 94 95 inheritance of genetic material, for example through nutritional, behavioural and epigenetic means (Schroeder et al., 2012, Schroeder et al., 2015). In conjunction with the impact of 96 97 sharing a common environment during the dependent phase, offspring behaviour might 98 appear to result from heritability, but could also be attributed to maternal behaviour 99 (Ainsworth & Bell, 1970, Maestripieri, 2009, Mandalaywala et al., 2014) or can magnify the 100 apparent magnitude of the heritability that is present.

101

Heritability of social attention in non-human primates has been little studied. Johnson et al 102 103 (2015) recorded the time olive and yellow baboons, Papio anubis and P. cynocephalus, spent 104 watching a human observer during a novel object test. Heritability of this behaviour was in 105 the range 18%-38%, with significant effects of sex and age (females tended to be more watchful of the observer while age effects are not fully reported). Watson et al (2015) 106 107 measured levels of vigilant scanning by rhesus macaques while drinking from a waterhole. 108 Heritability for scanning was 12%, with significant effects of age and sex (lower vigilance in 109 females and older monkeys). However, neither of these studies reported the contribution of 110 environmental or (non-genetic) maternal effects. To understand more fully the mechanisms 111 underpinning social attention in primates, it is important to explore the consistency 112 ('repeatability'), narrow-sense heritability (degree of behavioural expression that can be

accounted for by an individual's genes) and influence of non-genetic (including maternal and
other socio-environmental) effects on this behaviour.

115

116 Here we addressed these goals, using data on general attention to social information and attention specifically to social threat, collected from group-housed female rhesus macaques 117 118 Macaca mulatta for whom pedigree information was available. We recorded levels of 119 vigilant scanning while the monkeys were freely interacting with conspecifics, and also measured patterns of attention in a preferential looking task, during which animals were 120 shown pairs of conspecific faces (one with a threatening expression, the other a neutral 121 122 expression: after Bethell et al., 2012). We tested the heritability of attention to social information by measuring the extent to which relatedness explains (a) variation between 123 124 individuals in vigilant scanning rates, and (b) the total time spent looking towards pictures of 125 conspecific faces during the preferential looking task. We tested the heritability of attention 126 to social threat specifically, by measuring the extent to which relatedness explains, during 127 the preferential looking task (a) the time spent looking towards the threat face only, and (b) the bias in attention bias towards to the threat face relative to the neutral face. Finally, we 128 examined the extent to which any heritability may be explained by genetic and non-genetic 129 effects by accounting for variance explained by mother identity, separately. Because the 130 131 rhesus macaque is a female philopatric species with strong mother-daughter bonds, and 132 due to the controlled environmental conditions of captive housing, we treat maternal and 133 permanent environmental effects as a single 'non-genetic' factor labelled here as 'maternalenvironmental' effects. 134

135

- 137 Materials and Methods
- 138 Animals and housing
- 139 Data were collected from 67 adult female rhesus macaques (Macaca mulatta) housed at the
- 140 Centre for Macaques, MRC Harwell Institute, UK (mean age on first day of testing: 8.8 years,
- range 2.5 18.3 years; Table S1). Monkeys were UK-bred from founders of Indian origin.
- 142 Information about and a video of the facility can be seen at:
- 143 <u>https://www.mrc.ac.uk/research/facilities-and-resources-for-researchers/mrc-centre-for-</u>
- 144 <u>macaques/</u> and <u>www.nc3rs.org.uk/macaques</u>. Monkeys were housed in social breeding
- 145 groups comprising 1 adult male and between 3–11 related females, plus infants and
- 146 juveniles, following best practice guidelines (NC3Rs, 2006). Breeding groups had access to
- 147 two home areas: a main home room (dimensions 8.04m long x 3.35m wide x 2.8m height)
- 148 with an adjoining cage room (dimensions 6.12m long x 1.5m wide x 2.8m height); overall
- floor area 35.19m², and total volume 98.54m³. The main room had a large shelf c. 2m above
- the ground and shelves at multiple heights. The floor was covered in deep bedding material,
- and hanging enrichment included ladders, hoses, tunnels, boxes, swings and visual barriers.
- 152 The main room was also fitted with adjustable mirrors which the monkeys could control
- using a handle inside the main room which allowed them to look along the corridor outside
- their room. The cage room typically contained no enrichment devices and was designed so
- that staff could feed directly through the bars, and dividers could be used to separate off an
- 156 individual for veterinary inspection if needed. The main room and cage room were
- 157 connected by four hatches (one high, two medium and one low) which were kept open at all
- times during this study, and were only closed at other times during specific husbandry and

159	veterinary protocols. For this study, monkeys were tested in the cage room and all
160	individuals were free to come and go at all times (no dividers were used nor was any
161	monkey separated for testing). For the largest group (n=21 adults, juveniles and infants) the
162	two rooms provided c. 9m ³ volume per adult which is above the minimum of 3.5m ³ for
163	breeding animals under the UK Home Office Code of Practice (HomeOffice, 2014). All other
164	groups were smaller than this and so had more space per individual. Monkeys were fed a
165	regular diet of primate pellets, and a forage mix and various fruit and vegetables which were
166	scattered in the deep bedding to encourage natural foraging behaviour. Food and water
167	were available ad libitum in the main room at all times.
168	
169	
170	Pedigree
171	Pedigree information was available from colony records managed by DF and CW. The full
172	pedigree contained 597 individuals with a maximum depth of 7 generations. Pedigree
173	information was processed in R version 3.4.3 (RCoreTeam, 2018) using the package
174	MasterBayes (Hadfield et al., 2006). Summary statistics were calculated using the R package
175	pedantics (Morrissey, 2014). The full pedigree and informative pedigree are presented in
176	Supplementary materials (Figure S1). Mother identities were assigned from the pedigree.
177	
178	Life history measures
178 179	Life history measures For each female, life history information for three factors was collected through direct

testing (mean=9.7 years, range 2.5 – 18.1), social rank (high n=27, middle n=27, low n=13)

182	and reproductive status (cycling, pregnant, nursing). We assigned social rank in discussion
183	with facility staff based on our observations of which animals were most likely to attack or
184	displace others from preferred locations or food sources. High rank was assigned to
185	monkeys which displaced most others, low rank was assigned to monkeys who were most
186	likely to be displaced, attacked or to avoid others, and those who were displaced by high
187	ranked individuals but displaced low ranked individuals were assigned mid rank.
188	Reproductive status was in some cases confirmed retrospectively from timings of births,
189	assuming a gestation length of 167 days (Silk et al., 1993).

191 Cognitive measures

Attention to social information in general, and social threat specifically, was assessed using 192 193 an attention bias preferential looking task (Bethell et al., 2012). Monkeys had previously been target-trained, using positive reinforcement, to sit next to individual 'targets' in the 194 caged area adjacent to the main enclosure, as detailed in (Kemp et al., 2017). Whilst seated 195 196 by their unique target, each monkey was familiarised with a freestanding apparatus consisting of two picture holders with occluders on a height-adjustable tripod, and with a 197 digital video camera (Panasonic HCV520) positioned centrally between them. During the 198 199 initial familiarisation monkeys were encouraged to look towards the apparatus by 200 presenting food rewards centrally in front of the camera. Once oriented centrally, the occluders were manually opened via a sliding mechanism at the back of the apparatus to 201 202 reveal two pictures which had been pre-loaded, one into each picture holder. Monkeys 203 freely viewed the pictures for 3 seconds until the occluders were closed. During

familiarisation, pictures were of food items and conspecific infants, which were considered
to be of interest to adult female macagues.

206

207 During testing, monkeys underwent the same procedure except that pictures shown were of male conspecific faces (Figure 1) and the monkeys' gaze towards the two images was filmed 208 209 for each 3 second trial. Studies with humans most commonly use angry, disgust and pain 210 facial expressions as social threat cues (Heathcote et al., 2015, Hommer et al., 2014, Pérez-Edgar, 2010, Perez-Edgar et al., 2011, Schofield et al., 2013,). Here we used images of seven 211 212 unknown male macaques taken during aggressive encounters and during resting states. 213 Each picture pair contained one frontal view of the unfamiliar male macaque face with direct gaze and mouth open baring teeth in a tense, aggressive expression (threat face) and 214 215 one frontal view of the same male with the eyes and mouth closed and face relaxed in a 216 presumably neutral expression (neutral face). Pictures were cropped around the face and 217 superimposed on a rectangular grey background. Within each picture pair faces were 218 equated for luminance and contrast energy (full details given in Bethell et al., 2012). Location of the threat face on the left or right of the picture pair was counterbalanced 219 220 across trials for each monkey. This counterbalancing allowed us to control for the enhanced processing of emotional information in the right hemisphere of the primate brain (i.e. for 221 222 information presented to the left visual field: Adolphs et al. 1996) in our analyses.

223

Monkeys took part in one cognitive trial per day for four consecutive days from Tuesday –
 Friday in a given week. As part of a larger study investigating the relationship between
 emotion state and attention to social threat, monkeys were tested during weeks in which

veterinary health checks took place (presumed to be stressful, referred to here as 'stress' 227 228 condition), and weeks during which no presumed stressors occurred (referred to here as 229 'baseline' condition). During baseline weeks, when unexpected disruptions occurred (e.g. a 230 monkey had given birth, or a fight had occurred that morning) testing was delayed until the 231 next available day; this is because we intended to conduct trials with monkeys in a presumed 'non-stressed' state. To control for these potential confounding factors we 232 included condition as a control variable. As reproductive hormone levels may also affect 233 234 patterns of social attention (Lacreuse & Herndon, 2003) we also included reproductive status as a control variable. 235

236





237

238 Figure 1. A threat – neutral face pair

- 240 Video was coded in JWatcher+ Video V1.0 (Blumstein et al., 2000). Two coders (CK and HT)
- 241 blind coded video on a frame-by-frame basis for direction of eye gaze towards the left and

242	right picture locations during each trial. Coders achieved good agreement for coding a
243	subsample of the videos (Cohen's <i>k</i> =0.87). Once coded for direction of gaze on each frame,
244	trials were matched with records for location of the threat face (left/right). Number of trials
245	each monkey completed was recorded, as was time of day at which testing occurred.
246	
247	Behavioural observations
248	In the afternoon following a monkey's cognitive trial, her levels of vigilant scanning in the
249	social group were recorded for 5 minutes using continuous focal animal sampling (Altmann,
250	1974). We recorded duration of time engaged in vigilant scanning behaviour, defined here
251	as 'predominant behaviour is observing surrounding environment, moving the eyes and or
252	head actively scanning surroundings'. We considered vigilant scanning to be a suitable proxy
253	for social vigilance behaviour since social groups were generally highly socially active.
254	
255	Data preparation
256	Five variables functioned as response measures. There were three measures of social
257	attention. Duration of vigilant scanning in the social group (VIG), was recorded in
258	seconds/min; duration of time looking at both face pictures during the preferential looking
259	task (SOC), was recorded in ms per trial; and duration of time spent looking at faces
260	presented in the left location (i.e. to the left visual field, LVF), was recorded in ms per trial.

There were two measures of *attention to threat*. Duration of time looking at the threat face (THREAT), was recorded in ms per trial. Attention bias for threat (BIAS), was calculated as [THREAT – NEUT] where NEUT was the time spent looking towards the neutral face during the trial, providing bias scores on a scale from -3000ms to 3000ms.

266

We visually inspected plots of all response and predictor variables to check for a normal distribution using the R packages 'tidyverse' (Wickham, 2017) and 'car' (Fox & Weisberg, 2011). We transformed variables where this improved the distribution. For the response variables, we square-root transformed THREAT and LVF. For the predictor variables time of day and trial number required log transformation. All covariates were then scaled to a mean of 0 and SD±1 to provide more comparable estimates (Aiken & West, 1991, Schielzeth, 2010).

273

274 Statistical analysis

Statistical analyses were performed in R v. 3.4.3 (RCoreTeam, 2018). We estimated additive 275 276 genetic and maternal-environmental effects using an 'animal model' (Falconer & Mackay, 1995) with a Bayesian approach (Hadfield, 2010). The animal model is a generalised linear 277 278 mixed model (GLMM) which includes the pedigree as a random effect. This allows 279 heritability to be estimated and accounts for its effect on the statistical relationships being 280 tested. Specifically, we first calculated repeatability (with individual monkey as a random 281 effect) to establish how much variation in each attention measure was due to betweenindividual differences. The proportion of variance explained by between-individual 282 differences is the repeatability – typically considered the upper limit for heritability of a trait 283

(Lynch & Walsh, 1998). Therefore we subsequently only assessed heritability and maternalenvironmental effects for measures with a repeatability >0. We built an appropriate model
using the function 'Imer' in the package 'Ime4' (version 1.1-15; (Bates et al., 2015). We
checked all potential predictor variables for autocorrelation. The three life-history variables
(age, rank and reproductive status) showed no evidence of autocorrelation and were
retained as fixed effects (all r<0.04).

290

291 We controlled for experimental factors expected to contribute to within-individual variation 292 in the measures of attention. Condition (baseline or stress) was included to account for the 293 effects of current emotion state on attention to threat faces (Bethell et al., 2012). Time of day (recorded as 1-hour time blocks between 9am and 1pm) was included to control for 294 possible influence of circadian rhythm on alertness (Foster & Kreitzman, 2014); and trial 295 296 number (1-12) was included to control for habituation effects following repeated 297 presentations of stimuli over time (Bethell et al., 2019). For analysis of data from the 298 cognitive trials, side on which the threat face was shown (left or right) was included to control for left visual field priority of processing for emotional information. Stimulus ID 299 300 (seven stimulus pairs were used) was included to account for variation in features of the stimulus monkeys' faces that might influence attention (Waitt et al., 2003). 301

302

303 Repeatability

We ran GLMMs with measurements of the response variable with repeated observations
within individuals, one model for each response variable, using the package MCMCgImm

(Hadfield, 2010). To estimate repeatability (R) for each response variable we modelled an
identifier for each individual animal as a random factor on the intercept. We then calculated
the repeatability as

309 (1) $R = V_{\rm ID}/(V_{\rm ID}+V_{\rm R})$

310

311 where V_{ID} stands for the variance explained by between-individual differences, and V_R for 312 the residual variance. This approach is common in animal personality research (Nakagawa & 313 Schielzeth 2010).

314

315 Heritability and maternal effects

316 For those traits in which we identified non-zero repeatability, we ran animal models to estimate additive genetic variance (V_A), and maternal-environmental (V_M) effects using the 317 318 MCMCglmm package in R (Hadfield et al., 2006, Hadfield, 2010). Models were structured as for the repeatability, but we added additional random effects. Firstly, we included the 319 inverse relatedness matrix calculated from the pedigree as a random effect to estimate 320 321 additive genetic variance (V_A) (Wilson et al. 2010). Secondly, we added maternal identity as 322 a random effect to estimate maternal effects ($V_{\rm M}$). In these models (compared to the 323 repeatability models above) variance component estimated by the individual identity was named VPE, because now this part only estimates the variance explained by permanent 324 environment effects estimated through repeated measures on the individual, excluding any 325 additive genetic effects (Kruuk & Hadfield 2007). Note that VPE also accounts for the 326 327 pseudoreplication introduced by the repeated measures, and as such we always kept this parameter in the model. We calculated the heritability h^2 , the proportion of variance 328

explained by permanent environment *PE*, and the proportion of variance explained by
maternal effects *M* of each behavioural trait as

331 (2) $h^2 = V_A / (V_{PE} + V_A + V_M + V_R)$

332 (3) $PE = V_{PE}/(V_{PE}+V_A+V_M+V_R)$

333 (4)
$$M = V_{\rm M} / (V_{\rm PE} + V_{\rm A} + V_{\rm M} + V_{\rm R})$$

334

(Dochtermann et al., 2015) suggested that, to better understand the heritability of 335 behavioural traits, any within-individual variance is not relevant as it is not informative for 336 between-individual differences, which we are interested in. For transparency and to aid 337 meta-analytic approaches, we present both types of quantitative genetic estimates. Hence, 338 339 following Dochtermann et al. (2015), we also calculated the heritability of our looking 340 measures treated as components of 'personality' differences by assessing the proportion of the variance due to between-individual differences (disregarding V_R), and present the 341 342 proportion of the repeatable variance in contrast to the total variance in equations 2-4 as: (5) $h_P^2 = V_A / (V_{PE} + V_A + V_M)$ 343 (6) $PEp = V_{PE}/(V_{PE}+V_A+V_M)$ 344 (7) $Mp = V_M / (V_{PE} + V_A + V_M)$ 345 346 We fitted Bayesian GLMMs using the R package MCMCglmm (Hadfield 2010). We used 347 348 default priors where possible, and if needed we used proper priors with a low degree of

belief. We used visual inspection of plots of posteriors to ensure that the chain converged,

and assured empirically that autocorrelation at the specified thinning interval was low (i.e.

351 <0.1).

349

353 Ethical Statement

- This work was approved by the CFM Animal Welfare and Ethical Review Body and LJMU 354 355 ethics panel approval #EB/2014-1. All research was carried out in accordance with ethical 356 guidelines for work with non-human primates (NC3Rs, 2006) and reported following ARRIVE guidelines (Kilkenny et al., 2010). The general health of the animals was monitored daily by 357 the care staff via visual checks, and annually with a full veterinary examination. If during the 358 359 course of the project a researcher observed an injury or a concerning pattern of behaviour 360 they reported their concerns to the CFM care staff promptly so that any necessary treatment could take place. 361 362 363 364 Results Data summary 365 Vigilant scanning 366 Data on vigilant scanning in the social group were collected from 67 monkeys on 400 367 occasions over a period of 15 months (mean = 7.01 observations per monkey, range 2-12). 368 Monkeys spent an average of 26.65 seconds/minute engaged in vigilant scanning (range 0 – 369 59.60 seconds). 370 371
- 372 Cognitive trials (SOC, LVF, THREAT and BIAS)

A total of 432 cognitive trials were completed by the 67 monkeys (mean = 8.27 trials, range 373 374 4-12). We removed data related to cognitive trials as follows: trials involving monkeys which 375 had given birth in the preceding 24 hours, or which had been injured in the last 48 hours 376 (n=6 trials, 1.39% of the data); trials involving monkeys which had experienced a change in 377 group structure in the preceding 7 days (n=15 trials, 3.47% of the data); trials that occurred 378 within 24hours after room cleaning (n=10 trials, 2.31% of the data); and trials that occurred 379 after 13:00hrs (n=29 trials, 6.71% of the data). We also removed data from 10 females 380 which were moved to a non-breeding group at the start of the study period (n=72 trials, 16.67%). This resulted in 371 cognitive trials from 57 monkeys for inclusion in the analysis 381 382 (mean = 6.50 trials per monkey, range 4-12). Model output for cognitive trials are given in 383 Supplementary Table S2.

384

385 Pedigree

There were 144 individuals in the informative pedigree, with 49 unique mothers for 67 monkeys (Figure S1).

388

389 Repeatability

Repeatabilities are shown in Table 1. Vigilant scanning in the social group was repeatable

391 (VIG, R= 0.11), as was attention to social information during cognitive trials (SOC, R=0.25),

duration of looking at faces presented in the left visual field (LVF, R=0.13), and time spent

looking towards the threat face (THREAT, R=0.16). There was zero repeatability for attention

bias score (BIAS, R=0.00).

Table 1. Repeatability (R), heritability (h²), and maternal-environmental effects (*M*) for behaviour (B) and personality (P) for the four measures 395 of attention for social information and social threat with non-zero repeatability (95% Confidence Intervals shown in brackets). R: repeatability 396 of behaviour within individuals; h²B: heritability of behaviour; MB: maternal effects on behaviour; h²P: heritability of personality; MP: maternal 397 effects on personality. VIG: duration of vigilant scanning in the social group; SOC: duration of time looking at both faces; LVF duration of time 398 399 looking at both faces when presented to the left visual field only; THREAT: duration of looking towards the threat face; BIAS: attention bias score calculated as [duration of looking towards threat face - duration of looking towards the neutral face]. *Confirmatory analyses revealed 400 zero repeatability (R=0) for duration of looking at faces presented to the right visual field. ⁺Confirmatory analysis revealed there was zero 401 repeatability (R=0) for attention towards the NEUT face. 402 403

		R	h^2	M _B	h^2	M _P		
Target of attention	Response	(95CI)	Behaviour	Maternal effects on Behaviour	Personality	Maternal effect on Personalit		
Social information	ion VIG 0.11 (0.09	0.06	0.55	0.32		
		(0.00-0.24)	(0.02–0.18)	(0.02 – 0.16)	(0.19-0.78)	(0.24-0.49)		
	SOC	0.25	0.14	0.11	0.59	0.38		
		(0.13 - 0.38)	(0.04–0.26)	(0.02 - 0.23)	(0.17-0.83)	(0.08 - 0.61)		
	LVF*	0.13	0.08	0.10	0.23	0.38		
		(0.06-0.26)	(0.02-0.18)	(0.02 – 0.20)	(0.08-0.60)	(0.11-0.64)		
Social threat		0.16	0.10	0.04	0.63	0.30		
	THREAT	(0.06–0.30)	(0.04 - 0.23)	(0.02-0.15)	(0.22-0.85)	(0.11 - 0.59)		

406 Heritability and Maternal Effects

- 407 Heritabilities and maternal-environmental effects for all measures with non-zero
- 408 repeatability are shown in Table 1. Heritability for attention to social information (VIG, SOC
- and LVF) and social threat (THREAT) was present but low (VIG, $h^2 = 0.09$; SOC, $h^2 = 0.14$; LVF,
- 410 h^2 =0.08; THREAT, h^2 = 0.10). Maternal-environmental effects were of comparable
- 411 magnitude to heritability for all four measures, with highly overlapping confidence intervals
- 412 (VIG, $M_B = 0.06$; SOC, $M_B = 0.11$; LVF, $M_B = 0.10$; THREAT, $M_B = 0.04$).

413

414

415 Discussion

- 416 We tested for heritability of, and maternal-environmental effects on, attention for social 417 information in general, and social threat specifically, in adult female rhesus macaques. Using data on levels of vigilant scanning while animals were in their social group, and their 418 duration of looking towards threat-neutral stimulus pairs presented during an attention bias 419 420 preferential looking task, we found evidence that both attention for social information in general, and for social threat specifically, are heritable and also shaped by maternal-421 422 environmental effects. These findings provide the first measures of both genetic and nongenetic contributions to social attention in a non-human primate. 423 424 The heritability of vigilant scanning in the social group in this study, at 9%, is slightly lower 425 than, but generally consistent with values reported in the few other published findings on 426
- 427 social attention within social groups: rhesus macaques 12% (Watson et al., 2015); baboons

16-38% (Johnson et al., 2015). In both of these studies maternal and other permanent 428 environmental effects were not directly accounted for in the analysis. Here we found that 429 430 vigilant scanning was also partly explained by maternal-environmental effects, which 431 accounted for 6% of the variability in this behaviour. Both genetic and non-genetic factors 432 therefore influence vigilant scanning in the social group. We also found genetic and nongenetic contributions to social attention in the attention bias preferential looking task -433 434 heritability and maternal-environmental effects being 14% and 11% respectively for 435 attention to social stimuli overall, 8% and 10% for attention to social stimuli presented to the left visual field and 10% and 4% for attention to threat faces. 436

437

Our findings contribute to a growing literature demonstrating heritability of social attention 438 in non-human primates. Hopkins et al. (2014) assessed heritability of joint attention in 439 440 chimpanzees (Pan troglodytes), finding this to be between 22%-25%, and more strongly 441 heritable in males than females. Maternal effects were not tested in that study. Work with 442 rhesus macaques highlights how important maternal effects are and their potential for interaction with genetics: Golub et al. (2012) tested infant male rhesus macaques between 3 443 -14 months for the effect on several measures of social attention of MAOA genotype and 444 iron deficiency in the mother's diet during the third trimester. Offspring with the low-445 446 expressing MAOA genotype whose mothers had the iron deficient diet showed the poorest 447 level of emotion regulation; they were most vigilant towards video of aggressive 448 conspecifics and exhibited the most fearful behaviours during a number of social response tests. Where data are available, the inclusion of mother identity in studies of heritability of 449

450 behavioural traits will help elucidate the relative contribution and interaction between451 genotype and environment.

452

453 Repeatability values were slightly lower than, but broadly consistent with, the distribution of reported values for a wide range of behaviours in the animal behaviour literature (our 454 range: 0.11-0.25; literature mean=0.37 and mode = 0.2, Bell et al., 2009). In their meta-455 456 analysis of animal studies Bell et al (2009) found lower repeatability for behaviours expected to be flexible across contexts, for data collected in the laboratory compared to the field, and 457 for measurements taken further apart in time. They found no relationship between number 458 459 of repetitions of a measure and repeatability, although the majority of studies in the analysis used only a single or a few repeats of a measure. Our more conservative values for 460 repeatability may be partly explained by our previous work showing attention for threat 461 varies across contexts (Bethell et al., 2012), testing of animals in a captive environment, and 462 the relatively large number of repeated trials (up to 11 repeats) spread across a 10 month 463 464 period.

465

472

There was zero repeatability in our study for attention bias for threat (BIAS). Looking time towards the threat face alone may therefore be a more reliable measure of threat bias than calculating difference scores, as is typical in the human literature (Bar-Haim et al., 2007) and seen in earlier attention bias studies with non-human primates (Bethell et al 2012; Mandalaywala et al. 2014). Indeed, a confirmatory check revealed there was zero repeatability for looking time towards the neutral face (NEUT: result not presented here),

indicating that the calculation for BIAS diluted the repeatability for THREAT.

Our finding for non-heritability of attention bias for threat (BIAS) is in line with the few 474 475 studies that have explored the heritability of attention bias for threat in humans, typically by 476 comparing responses on cognitive tasks between monozygotic and dizygotic twins (Savage 477 et al., 2017). These studies all found no evidence for heritability of attention bias for threat 478 faces. For example, Rijsdijk et al. (2009) tested 125 pairs of female monozygotic and dizygotic twins on an emotion-face dot-probe study (Mogg & Bradley, 1999), in which 479 480 threat-neutral face pairs were presented for 19ms (i.e. subliminally), and reaction times to 481 subsequently presented probes recorded. Participants showed an overall attention bias for threat (measured as significantly faster responses to probes occurring at the location of 482 previously shown threat versus neutral faces). There was no evidence for heritability of 483 484 attention bias when data were collapsed across both visual fields ($h^2 = 0$). Brown et al. (2013) conducted an emotion-face dot-probe task with 600 eight-year-old mono- and 485 486 dizygotic twins, in which threat-neutral face pairs were shown for 1,000ms. Attentional 487 avoidance of threat was evident in the most anxious children, and this was not influenced by either degree of relatedness between twins or shared environment. The authors concluded 488 489 that attentional processes probably do not mediate the link between genetic risk and the 490 development of anxiety disorders in children, as has been widely proposed in the literature (Macleod et al., 1986, Mogg & Bradley, 1999). Finally, (Elam et al., 2010) conducted a dot-491 492 probe task with 22 twins aged five years in which threat-neutral face pairs were shown for 493 500ms. While monozygotic twins were more similar than were dizygotic twins in their responses to probes, this difference failed to reach significance. Attention bias for threat 494 495 therefore appears to be a highly plastic behaviour in both human and no-human primates. By calculating attention bias scores we lost any of the repeatability seen for looking towards 496

the threat face on its own. It is interesting to consider that calculating difference scores, as
is typically done in human cognitive psychology research, introduces noise that may mask
any signal of heritability or maternal effects.

500

Also in line with the human attention bias literature, we found heritability of attention to 501 social information presented to the left visual field, regardless of emotional content, in the 502 503 range 2-18%. Rijsdijk et al. (2009) reported heritability for attention to social information 504 among female twins tested on a dot probe task using threat-neutral face pairs. Heritability 505 was only found for reaction times to probes presented to the left visual field, regardless of 506 which face type they followed (41% and 49% for threat and neutral faces respectively). This finding was interpreted as evidence for a right hemisphere sensitivity for subliminal 507 biologically relevant cues under substantial genetic control; response times to probes in the 508 509 right visual field were driven by environmental effects only (both $h^2 = 0$). Elam et al. (2010) 510 also found evidence for heritability of a bias for responding to probes following faces in the 511 left visual field (~40%), while again heritability was 0% for probes following faces in the right visual field. 512

513

514 Our results will also be of interest to researchers interested in social attention for threat as 515 an underlying mechanism in human mental health conditions (e.g. Savage et al., 2017) from 516 an evolutionary perspective (Tremblay et al., 2017). There has been a paradigm shift in 517 approaches to mental health in the last decade, moving away from categorising phenotypes 518 under the traditional diagnostic categories presented in the DSM (American Psychiatric 519 Association, 2013), and towards an approach informed by genetics, neurobiology and

experimentally-derived behavioural measures, including attention bias (see Savage et al., 520 521 2017, for a review). As an example, attention bias for positive and negative faces has been 522 proposed to be a heritable mechanism underlying emotional resilience and vulnerability to 523 anxiety in humans, mediated by serotonin transporter (5-HTTLPR) polymorphism (Fox et al., 2009). Attention bias for threat words has been proposed to mediate the link between 524 neurobiology and behavioural pathology in trauma survivors (Pietrzak et al., 2014). The data 525 526 presented here may be of value for interpreting human data in an evolutionary context (e.g. 527 Green & Phillips, 2004).

528

529 Finally, our results will also be of value to animal cognition, behaviour and welfare researchers, trying to unpick the relative contribution of genetic and environmental factors 530 531 influencing susceptibility to stress. Work with mammals (primates: (Bethell et al., 2012), 532 sheep: (Monk et al., 2018)) and birds (Brilot et al., 2009) has shown that attention bias for 533 threat has potential for tracking changes in affective state (reviewed in (Crump et al., 2018). 534 Providing new and adaptable measures of animal emotion are essential for refinement in animal welfare practice (NC3Rs, 2006, Mendl et al., 2010). We hope the findings presented 535 here will help welfare researchers in these fields to better understand the role of heritable 536 and environmental factors in shaping the development and expression of social attention, 537 538 and its relationship to wellbeing.

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