1	Supplementary Information:	
2	Human mate-choice copying is domain-general social learning	
3	Sally E. Street, Thomas J.H. Morgan, Alex Thornton, Gillian R. Brown, Kevin N. Lala	nd,
4	Catharine P. Cross	
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1. Experimental design

1.1 Memorability of initial ratings

Our experimental design involved participants providing initial ratings and viewing social information for a block of images, before re-rating the images ('final ratings') following completion of the block (Methods). Participants completed the experiment in either in 3 blocks of 10 trials (for 4 of 6 experimental groups, N=33 participants) or 6 blocks of 5 trials (for 2 of 6 experimental groups, N=16 participants). Consequently, there was a time delay between providing initial ratings and providing final ratings, of ~10 minutes where longer blocks were used, or ~5 minutes where shorter blocks were used. Shorter blocks were used in addition to longer blocks, in case the greater time delay caused by longer blocks undermined the ability of participants to remember their initial ratings and social information when re-rating images.

In the post experiment questionnaire (SI 1.3), participants were asked to record, using a sliding scale, the extent to which they could remember their initial ratings when providing their final ratings, on a scale from 0 to 100. Participants reported mostly being able to remember their own ratings (mean 72.18, ±20.35). Further, self-reported memorability of initial ratings did not differ between participants who completed the study in 6 blocks of 5 trials (N=16) or 3 blocks of 10 trials (N=33, two sample T-test: T=-0.07, p=0.94). Therefore, we pooled data across the two blocking conditions for all analyses.

1.2 Participant instructions Read aloud to participants: "In this study you will see images of human faces, human hands and works of art. Your task will be to rate these images for attractiveness. You will receive a £5 Amazon voucher for taking part. The study consists of 30 questions arranged into [3 blocks of 10/6 blocks of 5]. Within each block, first you will rate all the images and be told what some of the other participants thought. You will be shown the average rating of some or all of the other participants, but you will not know whose ratings you are seeing. After rating all the images within the block, you will then re-rate all the images, although the order will be different. After you have completed all the blocks the study will end." "During the experiment, please interact only with the white window open on your screen now and please do not talk to other participants until you have been handed a debriefing sheet. You are free to withdraw at any time should you wish. One of us will be sitting in the adjacent room throughout the entire experiment should you have any problems." Ask if they all understand or have questions "Please click the arrow on your screen to continue."

1.3 Post-experiment questionnaire

After completion of the study, participants were asked to complete a short questionnaire.

Participants were reminded that their responses were voluntary, and all questions included a

'prefer not to say' option. Participants had to be aged 18 or over to take part in the study, but we did not request participants' ages in the post-experiment questionnaire. Participants were

asked to report their sexual orientation using a 7-point scale (where 0 indicated exclusively

heterosexual and 6 exclusively homosexual). 42/49 participants reported their sexual

orientation as 0 or 1, while 7/49 reported their sexual orientation as 2-6. Participants were

asked to self-describe their ethnicity using a free response box. Participants were asked

whether they did or did not know any other participants in the group. 15/49 reported that they

did know one or more of the other participants in the group, 32/49 reported that they did not,

and 2/49 reported 'prefer not to say'. Participants were asked to report, using a sliding scale,

to what extent they could remember their initial ratings when providing their final ratings,

where 100=maximum and 0=minimum, reporting a mean and standard deviation of 72.18,

±20.35.

Participants were asked to describe, using a free response box, how they decided to follow or ignore the social information they were shown. Of the 48/49 participants who responded to this question, 23 could be classified as reporting using a mixture of both their own judgement and the social information, while 24 reported using mostly their own judgement, and 1 gave an unclear answer. 15 of these 48 participants felt that they used the social information differently between the image types, and of 10 reporting being influenced most strongly by one image type in particular, 6 reported being most influenced for images of art, 4 for hands, and 2 for faces. Finally, participants were asked to describe, using a free response box, what they thought the intention of the experiment was. Of the 48/49 participants that responded to this question, all but one understood that the point of the experiment was to study social learning, but only 3 showed awareness that the intention was

to compare copying between different types of image. No participant responded in a way suggesting that they misunderstood the experimental task, or did not believe that the social information was genuine. Therefore, we were confident that participants understood the experimental task and treated the social information as genuine.

117 2. Supplementary results 118 119 2.1 Supplementary analysis including non-heterosexual participants 120 121 In our main analysis, we include only those participants (N=42) self-identifying as exclusively 122 or near-exclusively heterosexual (0 or 1 on a 7-point scale where 0 indicates exclusively 123 heterosexual, and 6 exclusively homosexual). When running our analysis on all (N=49) 124 participants, we find highly similar results. 125 126 Model performance 127 128 As in our main analysis, there was a correlation of 0.92 between predicted and observed 129 final ratings (pseudo-R² 0.84, N=1470), confirming that the model was appropriate for the 130 data. 131 132 Chain performance 133 134 Similarly to our main analysis, chain convergence was confirmed by large effective sample 135 sizes (range 3472 to 16538), and Gelman-Rubin statistics (all point estimates = 1, all upper 136 C.l.s = 1.01) across all estimated parameters. 137 138 Effect of condition on social influence 139 140 Similarly to our main analysis, when all participants were included, social information 141 affected participants' final ratings of images of faces (social influence median estimate = 142 0.14, [95% CI: 0.06, 0.22]), hands (social influence = 0.12, [0.04, 0.19]) and abstract art 143 (social influence = 0.15, [0.08, 0.22]). Again, medians and 95% CI for contrasts in social 144 influence between conditions suggested that differences were very close to zero (faces -145 hands = 0.03 [-0.02, 0.07], faces – art = -0.01 [-0.05, 0.04], art – hands = 0.03, [-0.02, 0.08]).

Individual participant effects As in our main analysis, the median variance of the random participant effect was 0.06 [0.04, 0.09], suggesting that relatively little of the variance in social influence was explained by consistent differences in social influence between participants. 153

172 2.2 Supplementary analysis allowing participant effects to differ between conditions 173 174 As in our main analysis, here we include only those participants (N=42) self-identifying as 175 exclusively or near-exclusively heterosexual. 176 177 Model performance 178 179 Very similarly to our main analysis, there was a correlation of 0.92 between predicted and 180 observed final ratings (pseudo-R² 0.85, N=1260), confirming that the model was appropriate 181 for the data. 182 183 Chain performance 184 185 The model included three parallel chains, each of 50,000 iterations thinned by 10 to reduce 186 autocorrelation. At completion, the effective sample size for the different variables ranged 187 from 4,813 to 16,068. Chain convergence was checked using the Gelman-Rubin 188 convergence diagnostic (all point estimates = 1, all upper C.I. = 1, except in one case where 189 the upper C.I. was 1.01). 190 191 Effect of condition on social influence 192 193 Very similarly to our main analysis, when allowing participant effects to vary with condition, 194 social information affected participants' final ratings of images of faces (social influence 195 median estimate = 0.13, [95% CI: 0.04, 0.22]), hands (social influence = 0.15, [0.05, 0.25]) 196 and abstract art (social influence = 0.13, [0.04, 0.22]). Again, 95% CI for contrasts in social 197 influence between conditions suggested that differences were close to, if not precisely, 0 198 (faces - hands = -0.02 [-0.15, 0.12], faces - art = < 0.01 [-0.13, 0.13], art - hands = -0.02, [-0.13, 0.13], art - hands = -0.02, [-0.15, 0.12], faces - art = < 0.01 [-0.13, 0.13], art - hands = -0.02, [-0.15, 0.12], faces - art = < 0.01 [-0.13, 0.13], art - hands = -0.02, [-0.15, 0.12], faces - art = < 0.01 [-0.13, 0.13], art - hands = -0.02, [-0.15, 0.12], faces - art = < 0.01 [-0.13, 0.13], art - hands = -0.02, [-0.15, 0.12], faces - art = < 0.01 [-0.13, 0.13], art - hands = -0.02, [-0.15, 0.15], art - hands = -0.02, [-0.15, 0.15], art - hands = -0.02, [-0.15, 0.15], art - hands = -0.02, [-0.1199 0.15, 0.11]).

Effect of condition on participant effects We find that the variance between participants in social influence is very similar for images of artwork (0.08 [0.05, 0.12]), faces (0.08 [0.05, 0.12]) and hands (0.08, [0.05, 0.14]). All contrasts between conditions in the between-participant variance were very close to zero (faces - hands = -0.01 [-0.06, 0.05], faces - art = >-0.01 [-0.05, 0.05], art - hands = -0.01 [-0.05, 0.05]0.07, 0.05]).

2.3 Supplementary analysis with flatter prior distributions

The careful choice of priors is an essential part of a Bayesian analysis. In the main paper, we present the results of an analysis that used "weakly regularizing priors" as suggested by an anonymous reviewer. These are priors that minimally constrain the output of the analysis by encouraging the model to focus on biologically plausible values. For instance, the social influence parameters for each condition were given, as a prior, a normal distribution with a mean of 0 and a variance of 1. This encourages the model to favour estimates of these parameters with a magnitude close to 0 and gives a very high chance that the magnitude is less than 3. Given that a value of 0 is no social information use, and a value of 1 is total conformity, this seems reasonable. Nonetheless it is important to check that the results are not unduly influenced by the choice of priors, so here we present the results of another analysis in which the priors were extremely flat. In this case the priors for the condition effects are normal distributions with a mean of 0 and a variance of 100. The prior for the variation between participants, which was an exponential distribution with a parameter value of 1 in the main paper, is instead gamma distributed with a shape and rate of 0.001.

Weakly regularizing priors are typically considered a better option than these extremely flat priors, as our anonymous reviewer pointed out. For instance, the extremely broad priors for the condition effects imply that the conditions are likely to be extremely different from each other and characterized by extreme values of social influence. While biologically plausible values will mainly fall between 0 and 1, values such as -200 are treated as perfectly plausible by the model with flat priors. Nonetheless, with enough data the priors can be overwhelmed and results should not be unduly affected. We therefore present these additional results to verify the robustness of our findings.

250	
251	Model performance
252	
253	As with our main analysis, we found a correlation of 0.92 between predicted and observed
254	final ratings (pseudo-R ² 0.84, N=1260), confirming that the model was appropriate for the
255	data.
256	
257	Chain performance
258	
259	Chain convergence was confirmed by large effective sample sizes (range 5334 to 7068) and
260	Gelman-Rubin statistics (all point estimates = 1, all upper C.I.s = 1.01).
261	
262	Effect of condition on social influence
263	
264	Replicating the results of our main analysis, social influence was highly similar for images of
265	faces (median estimate = 0.13, 95% CI: [0.08, 0.17]), hands (0.13, [0.08, 0.18]) and abstract
266	artwork (0.14, [0.10, 0.19]). As before, contrasts in social influence were effectively zero
267	(faces - hands = <-0.01 [-0.05, 0.05], faces - art = -0.02 [-0.06, 0.03], art - hands = 0.01, [-0.05, 0.05])
268	0.04, 0.06])
269	
270	Individual participant effects
271	
272	Similarly to our main analysis, we found a low random participant effect (median estimate =
273	0.01 [<0.01, 0.02]), suggesting little evidence of consistent individual differences in social
274	influence.
275	

276	3. Dataset details and analysis code		
277			
278	3.1 Details of dataset		
279	'Street_et_al_image_preference_data_2017.csv' contains the following columns:		
280			
281	playerID	numerical player identifier (1-49)	
282			
283	trialID	numerical trial identifier (1-1470)	
284			
285	groupID	numerical group identifier (1-6)	
286			
287	nplayers	number of players in group (5-10)	
288			
289	condition	type of image viewed and rated in the trial (art, faces or hands)	
290			
291	questions.per.block	number of questions in block (5 or 10)	
292			
293	initial.rating	initial attractiveness rating (minimum 0, maximum 100)	
294			
295	initial.decision.time	time taken to provide initial attractiveness rating (milliseconds)	
296			
297	social.rating	attractiveness rating of some or all other players (minimum 0,	
298		maximum 100)	
299			
300	social.decision.time	time taken to view social information (milliseconds)	
301			
302	final.rating	final attractiveness rating (minimum 0, maximum 100)	
303			

304	final.decision.time	time taken to provide final attractiveness rating (milliseconds)
305		
306	orientation	participant sexual orientation (minimum 0=exclusively heterosexual,
307		maximum 6=exclusively homosexual)
308		
309	know.anyone	whether participant knew any others in the group (yes, no,
310		prefer_not_to_answer)
311		
312	remember.initial.ratings	to what extent participant reported being able to remember initial
313		ratings when providing final ratings (minimum 0, maximum 100)
314		
315	use.social.info	coded from free responses to question of how participant chose to use
316		or ignore social information (mostly_individual = participant
317		reported using only or primarily individual preferences,
318		both_social_and_individual = participant reported using both social
319		information and individual preferences, not_clear = participant did
320		not provide a clear answer, no_answer = participant provided no
321		answer).
322		
323	experiment.intent	coded from free responses to question of what participant thought
324		was the intention of the experiment (social_influence = participant
325		perceived the intention of the study to be related to social influence,
326		non_social_influence = participant perceived the intention of the
327		study to be unrelated to social influence,
328		social_influence_image_types = participant perceived the intention
329		of the experiment as comparing social influence between image
330		types, no_answer = participant provided no answer).

3.2 R code for main analysis

```
332
333
     # load packages
334
     library(rjags)
335
     library(coda)
336
337
     # load data
338
     data<-read.csv("Street et al image preference data 2017.csv",
339
340
     str(data) # data file contains 1470 observations from total 49
341
     players
342
343
     # for the main analysis, include only participants who are
344
     exclusively or near-exclusively heterosexual (0 or 1 on the response
345
     scale)
346
     data2<-subset(data, orientation<2)</pre>
347
     str(data2) # data file contains 1260 observations from total 42
348
     players
349
350
     # re-assign player ID numbers (must be numbered 1:42 for model to
351
     run)
352
     data2$playerIDnew<-as.numeric(as.factor(data2$playerID))</pre>
353
354
     # select the variables for analysis
355
     initial <-data2 $ initial.rating # participants' initial rating
356
     social<-data2$social.rating # social information</pre>
357
     final<-data2$final.rating # participants' final rating</pre>
358
     player<-data2$playerIDnew # player identity</pre>
359
     condition<-as.numeric(data2$condition) # content type (art=1,</pre>
360
     faces=2, hands=3)
361
362
     # extract the sample sizes
363
     N<-length(final) # total number of trials
364
     N players<-length(unique(data2$playerID)) # total number of
365
     participants
366
367
     # transform all ratings to fall between 0 and 1
368
     p.final<-(final/100)*0.999+0.0005
369
     p.initial<-(initial/100)*0.999+0.0005
370
     p.social<-(social/100)*0.999+0.0005
371
372
     # load the model, setup to run for 3 chains, with a burn-in period
373
     of 5000 iterations
374
     model<-jags.model("Street et al 2017 main model JAGS code.bug.txt",</pre>
     data=list('p.initial'=p.initial, 'p.social'=p.social,
375
376
     'p.final'=p.final, 'player'=player, 'N players'=N players,
377
     'condition'=condition, 'N'=N), n.chains=3, n.adapt=5000, quiet=F)
378
379
     # run the model (takes around 25 minutes, runs 50000 iterations,
380
     sampling every 10 generations for each chain)
```

```
381
382
     # RUN ONLY ONE OF THE FOLLOWING TWO OPTIONS:
383
384
     # OPTION 1: this line monitors the predicted final ratings, the
385
     social influence parameter, the effect of image condition, the
386
     random participant effect and the variance of the random participant
387
     effect
388
     results<-coda.samples(model, c('final_pred', 'social_influence',
389
     'condition influence baseline', 'random player influence effect',
     'tau_players_social'), n.iter=50000, thin=10)
390
391
     save(results, file="model samples full.txt")
392
393
     # OPTION 2: this line is the same as the above but does not monitor
394
     predicted final ratings, social influence, or random player effects
395
     (although these are still in the model). It will take up less space
396
     on your hdd.
397
     results <- coda. samples (model, c('condition_influence_baseline',
398
     'tau players social'), n.iter=50000, thin=10)
399
     save(results, file="model samples lite.txt")
400
401
     # check the minimum effective sample sizes for all parameters
402
     sample size <- effectiveSize(results)</pre>
403
     range(sample size)
404
405
     # check the Gelman convergence diagnostic for all parameters
406
     # note this may break if using model samples full, suggest using
407
     model samples lite instead
408
     rhats <- gelman.diag(results)</pre>
409
     rhats
410
411
     # combine all three chains into a single data frame
412
     combined results <- rbind(as.data.frame(results[[1]]),</pre>
413
     as.data.frame(results[[2]]), as.data.frame(results[[3]]))
414
415
     # medians & 95% CI for condition effects on social learning #
416
     quantile (combined results $ 'condition influence baseline[1]',
417
     c(0.025, 0.5, 0.975)) # art
418
     quantile(combined results$'condition influence baseline[2]',
419
     c(0.025, 0.5, 0.975)) # faces
420
     quantile(combined results$'condition influence baseline[3]',
421
     c(0.025, 0.5, 0.975)) # hands
422
423
     # plot showing effect of condition (as used in Figure 2)
424
     par(mfrow=c(3,1))
425
     par(mar=c(7.5,6,2,2))
426
     hist(combined results$'condition influence baseline[1]',
     col=rgb(1,0,0,0.75), xlab="", main="", cex.axis=2.5, xlim=c(-0.05,
427
428
     0.30), ylim=c(0, 1500), breaks=50, ylab="", las=1, xaxt="n") # art
429
     axis(1, at=c(-0.05, 0, 0.05, 0.10, 0.15, 0.20, 0.25, 0.30),
430
     labels=F)
431
     abline(h=0)
```

```
432
     abline(v=median(combined results$'condition influence baseline[1]'),
433
     lty=2, lwd=3)
434
     abline(v=quantile(combined results$'condition influence baseline[1]'
435
      0.025, lty=2)
436
     abline(v=quantile(combined results$'condition influence baseline[1]'
437
     , 0.975), 1ty=2)
438
     hist(combined results$'condition influence baseline[2]',
439
     col=rgb(0,0,1,0.75), breaks=50, main="", cex.axis=2.5, xlim=c(-0.05,
     0.30), ylim=c(0, 1500), ylab="", las=1, xlab="", xaxt="n") # faces
440
441
     axis(1, at=c(-0.05, 0, 0.05, 0.10, 0.15, 0.20, 0.25, 0.30),
442
     labels=F)
443
     abline (h=0)
444
     abline(v=median(combined results$'condition influence baseline[2]'),
445
     lty=2, lwd=3)
446
     abline(v=quantile(combined results$'condition influence baseline[2]'
447
     , 0.025), 1ty=2)
448
     abline(v=quantile(combined results$'condition influence baseline[2]'
449
     , 0.975), 1ty=2)
450
     hist(combined results$'condition influence baseline[3]',
451
     col=rgb(1,1,0,0.75), breaks=50, main="", xlim=c(-0.05, 0.30),
452
     ylim=c(0, 1500), xlab="", ylab="", las=1, xaxt="n", cex.axis=2.5) #
453
     hands
454
     axis(1, at=c(-0.05, 0, 0.05, 0.10, 0.15, 0.20, 0.25, 0.30),
455
     labels=T, cex.axis=2.5, padj=1)
456
     abline(h=0)
457
     abline(v=median(combined results$'condition influence baseline[3]'),
458
     lty=2, lwd=3)
459
     abline(v=quantile(combined results$'condition influence baseline[3]'
460
     0.025, lty=2)
461
     abline(v=quantile(combined results$'condition influence baseline[3]'
462
     , 0.975), 1ty=2)
463
     mtext("Social influence by condition", side=1, line=5, cex=1.75)
464
465
     # contrasts in effects of condition on social learning
466
     Faces v Hands<-(combined results$'condition influence baseline[2]'-
467
     combined results$'condition influence baseline[3]') # faces vs.
468
     hands
469
     quantile(Faces v Hands, c(0.025, 0.5, 0.975))
470
471
     Faces v Art<-(combined results$'condition influence baseline[2]'-
472
     combined results$'condition influence baseline[1]') # faces vs. art
473
     quantile(Faces v Art, c(0.025, 0.5, 0.975))
474
475
     Art v Hands<-(combined results$'condition influence baseline[1]'-
476
     combined results$'condition influence baseline[3]') # art vs. hands
477
     quantile (Art v Hands, c(0.025, 0.5, 0.975))
478
479
     # individual participant effect
480
     par(mfrow=c(1,1))
481
     par(mar=c(5.1, 4.1, 4.1, 2.1))
482
     hist(1/combined results$tau players social)
```

```
483
     1/median(combined results$tau players social)
484
     1/quantile(combined results$tau players social, c(0.025, 0.975))
485
     1/quantile(combined results$tau players social, c(0.025, 0.5,
486
     0.975))
487
488
     # IF OPTION 1 WAS SELECTED, CHECK CORRESPONDENCE OF PREDICTED AND
489
     OBSERVED FINAL RATINGS
490
491
     # extract predicted final ratings from all chains
492
     results means <- col Means (combined results) # take the column means,
493
     to get the mean predicted final rating for each trial across the
494
     posterior distribution
495
     results predicted final <- results means [grep("final pred",
496
     names(results means))] # extract only the columns with predicted
497
     final ratings
498
499
     # Pseudo R^2 - Pearson's correlation of predicted vs. observed final
500
501
     cor.test(results predicted final, p.final, method="pearson") #
502
     Pearson's corr
503
     cor.test(results predicted final, p.final,
504
     method="pearson")$estimate^2 # pseudo R squared
505
506
     # distribution of modelled final ratings compared to observed final
507
     ratings (as used in Figure 3)
508
     par(mfrow=c(2,1))
509
     par(mar=c(7.5, 6, 4, 2))
     hist(p.final, cex.lab=2, col=rgb(0.3,0.6,0.3,0.25), main="",
510
511
     cex.axis=2, ylim=c(0, 60), breaks=50, ylab="", las=1, xlab="Observed
512
     final ratings")
513
     hist(results predicted final, col=rgb(0.3,0.6,0.3,0.75), ylim=c(0,
514
     60), breaks=50, xlab="Predicted final ratings", cex.axis=2,
515
     cex.lab=2, main="", las=1, ylab="")
516
517
518
```

3.3 JAGS code for main analysis

```
521
522
     ### Model
523
524
     model{
525
526
      for (i in 1:N) {
527
528
          p.final[i] ~ dbeta(a final[i], b final[i]) # final ratings are
529
     modelled as a beta distribution
530
531
          a_final[i] <- final_pred[i] * phi_final # alpha shape parameter</pre>
532
     for the beta distribution
533
534
          b final[i] <- (1 - final pred[i]) * phi final # beta shape</pre>
535
     parameter for the beta distribution
536
537
          logit(final pred[i]) <- final lp[i] # logit link function</pre>
538
539
          final lp[i] <- logit(p.initial[i]) + social influence[i] *</pre>
540
     social deviation[i] # final ratings are predicted by the amount of
541
     social influence, relative to the amount that initial ratings and
542
     social information differ
543
544
          social deviation[i] <- logit(p.social[i]) - logit(p.initial[i])</pre>
545
     + 0.00001 # social deviation is the difference between the social
546
     information and the initial ratings, with a tiny constant added to
547
     avoid zero values
548
549
          social influence[i] <-</pre>
550
     condition influence baseline[condition[i]] +
551
     random player influence effect[player[i]] # social influence can
552
     vary by image condition (faces, hands or art), and by a random
553
     effect of participant identity
554
555
      }
556
557
     ### Priors
558
559
       phi final ~ dexp(1) # exponential prior for the shape parameters
560
     for the beta distribution of modelled final ratings
561
562
       for (i in 1:3) {
563
         condition influence baseline[i] ~ dnorm(0,1) # normal prior for
564
     the effect of image condition
565
       }
566
567
       for (i in 1:N players) {
```

```
random_player_influence_effect[i] ~ dnorm(0, tau_players_social)

for the random participant effect

for the random participant effect

tau_players_social ~ dexp(1) # exponential prior for the variance

for the random participant effect

for the variance

for the random participant effect

for the variance

for the variance

for the variance

for the random participant effect

for the variance

for the
```

```
578
     3.4 R code for supplementary analysis
579
     # load packages
580
     library(rjags)
581
     library(coda)
582
583
     # load data
584
     data<-read.csv("Street et al image preference data 2017.csv",
585
     header=T)
586
     str(data) # data file contains 1470 observations from total 49
587
     players
588
589
     ### supplementary analysis: does participant self-reported ability
590
     to remember initial ratings differ between blocking conditions (3 \times
591
     10 vs. 6 x 5 blocks)?
592
593
     ppt data<-aggregate(data[,c("playerID", "questions.per.block",</pre>
594
     "remember.initial.ratings")], by=list(data$playerID), FUN="mean") #
595
     collapse data to participant level. For convenience we aggregate by
596
     the function 'mean'.
597
598
     mean(ppt data$remember.initial.ratings) # overall mean and SD for
599
     self-reported memorability of initial ratings
600
     sd(ppt data$remember.initial.ratings)
601
602
     t.test(ppt data$remember.initial.ratings~as.factor(ppt data$question
603
     s.per.block), var.equal=T) # t-test assuming equal variances
604
605
     library("car") # check if the assumption of equal variances is
606
     violated using function in the 'car' package
607
     leveneTest(remember.initial.ratings~as.factor(questions.per.block),
608
     data=ppt data) # assumption of equal variances not violated
609
610
     #### supplementary analysis: replicating main analysis without
611
     excluding any participants based on sexual identity
612
613
     # select the variables for analysis
     initial<-data$initial.rating # participants' initial rating</pre>
614
615
     social<-data$social.rating # social information</pre>
616
     final<-data$final.rating # participants' final rating</pre>
617
     player<-data$playerID # player identity</pre>
618
     condition<-as.numeric(data$condition) # content type (art=1,</pre>
619
     faces=2, hands=3)
620
621
     # extract the sample sizes
622
     N<-length(final) # total number of trials
623
     N players<-length(unique(data$playerID)) # total number of
624
     participants
625
626
     # transform all ratings to fall between 0 and 1
627
     p.final<-(final/100)*0.999+0.0005
```

```
628
     p.initial<-(initial/100)*0.999+0.0005
629
     p.social<-(social/100)*0.999+0.0005
630
631
     # load the model, setup to run for 3 chains, with a burn-in period
632
     of 5000 iterations
633
     model<-jags.model("Street et al 2017 main model JAGS code.bug.txt",</pre>
634
     data=list('p.initial'=p.initial, 'p.social'=p.social,
635
     'p.final'=p.final, 'player'=player, 'N players'=N players,
636
     'condition'=condition, 'N'=N), n.chains=3, n.adapt=5000, quiet=F)
637
638
     # run the model (takes around 25 minutes, runs 50000 iterations,
639
     sampling every 10 generations for each chain)
640
641
     # RUN ONLY ONE OF THE FOLLOWING TWO OPTIONS:
642
643
     # OPTION 1: this line monitors the predicted final ratings, the
     social influence parameter, the effect of image condition, the
644
645
     random participant effect and the variance of the random participant
646
     effect
647
     results <- coda. samples (model, c('final pred', 'social influence',
648
     'condition influence baseline', 'random player influence effect',
649
     'tau players social'), n.iter=50000, thin=10)
650
     save(results, file="model samples full all ppts.txt")
651
652
     # OPTION 2: this line is the same as the above but does not monitor
653
     predicted final ratings, social influence, or random player effects
654
     (although these are still in the model). It will take up less space
655
     on your hdd.
656
     results <- coda. samples (model, c ('condition influence baseline',
657
     'tau players social'), n.iter=50000, thin=10)
658
     save(results, file="model samples lite all ppts.txt")
659
660
     # check the minimum effective sample sizes for all parameters
661
     sample size <- effectiveSize(results)</pre>
662
     range(sample size)
663
664
     # check the Gelman convergence diagnostic for all parameters
665
     # note this may break if using model samples full, suggest using
666
     model samples lite instead
667
     rhats <- gelman.diag(results)</pre>
668
     rhats
669
670
     # combine all three chains into a single data frame
671
     combined results <- rbind(as.data.frame(results[[1]]),</pre>
672
     as.data.frame(results[[2]]), as.data.frame(results[[3]]))
673
674
     # medians & 95% CI for condition effects on social learning #
675
     quantile (combined results $ 'condition influence baseline[1]',
676
     c(0.025, 0.5, 0.975)) # art
677
     quantile(combined results$'condition influence baseline[2]',
678
     c(0.025, 0.5, 0.975)) # faces
```

```
679
     quantile(combined results$'condition influence_baseline[3]',
680
     c(0.025, 0.5, 0.975)) # hands
681
682
     # contrasts in effects of condition on social learning
683
     Faces v Hands<-(combined results$'condition influence baseline[2]'-
684
     combined results$'condition influence baseline[3]') # faces vs.
685
686
     quantile(Faces_v_Hands, c(0.025, 0.5, 0.975))
687
688
     Faces v Art<-(combined results$'condition influence baseline[2]'-
689
     combined results$'condition influence baseline[1]') # faces vs. art
690
     quantile(Faces v Art, c(0.025, 0.5, 0.975))
691
692
     Art v Hands<-(combined results$'condition influence baseline[1]'-
693
     combined results$'condition influence baseline[3]') # art vs. hands
694
     quantile (Art v Hands, c(0.025, 0.5, 0.975))
695
696
     # individual participant effect
697
     1/quantile(combined results$tau players social, c(0.025, 0.5,
698
     0.975))
699
700
     # IF OPTION 1 WAS SELECTED, CHECK CORRESPONDENCE OF PREDICTED AND
701
     OBSERVED FINAL RATINGS
702
703
     # extract predicted final ratings from all chains
704
     results means <- colMeans (combined results) # take the column means,
705
     to get the mean predicted final rating for each trial across the
706
     posterior distribution
707
     results predicted final <- results means [grep ("final pred",
708
     names(results means))] # extract only the columns with predicted
709
     final ratings
710
711
     # Pseudo R^2 - Pearson's correlation of predicted vs. observed final
712
713
     cor.test(results predicted final, p.final, method="pearson") #
714
     Pearson's corr
715
     cor.test(results predicted final, p.final,
716
     method="pearson")$estimate^2 # pseudo R squared
717
```

3.5 JAGS code for supplementary analysis

```
720
721
     ### Model
722
723
     model{
724
725
       for (i in 1:N) {
726
727
          p.final[i] ~ dbeta(a_final[i], b_final[i]) # final ratings are
728
     modelled as a beta distribution
729
730
          a_final[i] <- final_pred[i] * phi_final # alpha shape parameter</pre>
731
     for the beta distribution
732
733
          b final[i] <- (1 - final pred[i]) * phi final # beta shape</pre>
734
     parameter for the beta distribution
735
736
          logit(final pred[i]) <- final lp[i] # logit link function</pre>
737
738
          final lp[i] <- logit(p.initial[i]) + social influence[i] *</pre>
739
     social deviation[i] # final ratings are predicted by the amount of
740
     social influence, relative to the amount that initial ratings and
741
     social information differ
742
743
          social deviation[i] <- logit(p.social[i]) - logit(p.initial[i])</pre>
744
     + 0.00001 # social deviation is the differences between the social
745
     information and the initial ratings, with a tiny constant added to
746
     avoid zero values
747
748
          social influence[i] <-</pre>
749
     condition influence baseline[condition[i]] +
750
     random player influence effect[condition[i], player[i]] # social
751
     influence can vary by image condition (faces, hands or artwork), and
752
     by a random effect of participant identity. Additionally, the random
753
     participant identity effect can differ by condition.
754
755
      }
756
757
     ### Priors
758
759
       phi final ~ dexp(1) # exponential prior for the shape parameters
760
     for the beta distribution of modelled final ratings
761
762
       for (i in 1:3) {
763
         condition influence baseline[i] ~ dnorm(0,1) # normal prior for
764
     the effect of image condition
765
       }
766
767
       for (i in 1:3) {
```

```
768
      tau_players_social[i] ~ dexp(1) # exponential prior for the
769
    variance of the random participant effect
770
         for (j in 1:N players) {
           random player influence effect[i, j] ~ dnorm(0,
771
772
     tau players social[i]) # normal prior for the random participant
773
     effect dependent on image condition (faces, hands or art).
774
        }
775
      }
776
     }
777
778
779
780
```

782 4.1 STAN code supplied by an anonymous reviewer 783 The reviewer who suggested we use weakly regularizing priors also kindly provided code to 784 run our analysis in STAN, another piece of Bayesian analysis software. STAN runs the same 785 kinds of analyses as JAGS, but uses different sampling algorithms which can greatly 786 increase the efficiency with which models run. If readers are familiar with both JAGS and 787 STAN and wish to reproduce our analyses they will likely find that STAN is the faster way to 788 do this. 789 790 # Stan model 791 library(rstan) 792 793 stan_model_code <- " 794 data{ 795 int<lower=1> N; 796 int<lower=1> N_condition; 797 int<lower=1> N_player; 798 real final[N]; 799 real initial[N]; 800 real social[N]; 801 int condition[N]; 802 int player[N]; 803 } 804 parameters{ 805 vector[N_condition] b_condition; 806 vector[N_player] b_player; 807 real<lower=0> sigma; 808

781

4. Additional code

real a;

```
809
       real<lower=0> phi;
810
       }
811
       model{
812
       vector[N] b;
813
       vector[N] p;
814
       phi ~ exponential(1);
815
       a \sim normal(0,1);
816
       sigma ~ exponential(1);
817
       b_player ~ normal( 0 , sigma );
818
       b_condition ~ normal(0,1);
819
       for ( i in 1:N ) {
820
       b = a + b_condition[condition] + b_player[player];
821
       p = (1 - b) * logit(initial) + b * logit(social);
822
       p = inv_logit(p);
823
       }
824
       final ~ beta( p*phi, (1-p)*phi);
825
       }
826
       generated quantities{
827
       vector[3] social_influence_condition;
828
       for (i in 1:3) social_influence_condition = a + b_condition;
829
       }
830
831
832
       data_list <- list(
833
       N=N,
834
       N_condition=3,
835
       N_player=N_players,
836
       final=p.final,
```

```
837
       initial=p.initial,
838
       social=p.social,
839
       condition=condition,
840
       player=player
841
       )
842
843
       stan_fit <- stan( model_code=stan_model_code , data=data_list , chains=3 , cores=3 )
844
845
       # diagostics and such
846
       print(stan_fit,probs=c(0.025,0.975))
847
848
       # extract permuted samples
849
       post <- extract(stan_fit)</pre>
```