

# Bayesian reanalysis of Mesoudi et al. (2015)

Alex Mesoudi

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## 1 Introduction

This is a reanalysis of the data presented in Mesoudi, Chang, Murray & Lu (2015), using methods from Richard McElreath’s Statistical Rethinking book and **rethinking** package (McElreath 2016). The reanalysis generates virtually identical results to the original analysis, but is hopefully more useful and transparent to other researchers.

In Mesoudi et al. (2015) we found higher rates of social learning in a computer-based artifact-design task amongst participants from mainland China, compared to participants from the UK, from Hong Kong, and Chinese immigrant students studying in the UK. Participants were presented with 29 opportunities to either copy or not copy others during a season of 30 hunts (participants could not copy on the first hunt). Participants could see others’ success rates, making social learning payoff-biased and therefore highly beneficial in this challenging task. There were 3 seasons, each with 29 opportunities to copy. Seasons 1 and 2 featured different environments but no within-season environmental change. Season 3 featured a different environment *and* within-season environmental change.

## 2 Visualisation of data

The original paper presented bar charts of the proportion of copying per season as in Figure 1 below (Figure 2 in Mesoudi et al. 2015). We can see the higher copying proportion in Chinese Mainland (CM) participants in Seasons 1 and 2, but not in Season 3. In Season 3 the UK, HK and CI participants increased their copying frequency almost up to CM levels.

However, bar charts can obscure variation in data. Ideally visualisations would include representations of the actual data, and counts rather than proportions. Figures 2-4 show some alternatives. These show that there is a broad spread of copying frequencies within each cultural group, but that in Seasons 1 and 2 there are more high-copying CM participants compared to the other three groups.

## 3 Bayesian re-analysis

The original paper used negative binomial regression to compare copying proportions across cultural groups. These regressions are shown in Table 1 of Mesoudi et al. (2015). Here I use the **map** function from the **rethinking** package to compare cultural groups in an aggregated binomial model. The dependent measure is the number of hunts (out of 29) on which a participant copied another participant. This is performed separately for season 1, 2 and 3, each of which featured 29 opportunities to copy<sup>1</sup>. For each season three

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<sup>1</sup>I do not run a single regression with season as a within-participant factor because, as noted above, Season 3 introduces within-season environmental change and so is not comparable to the others. Seasons 1 and 2 could be combined, but the gain in having one fewer model does not seem to me to outweigh the cost of having coefficients that are harder to interpret.

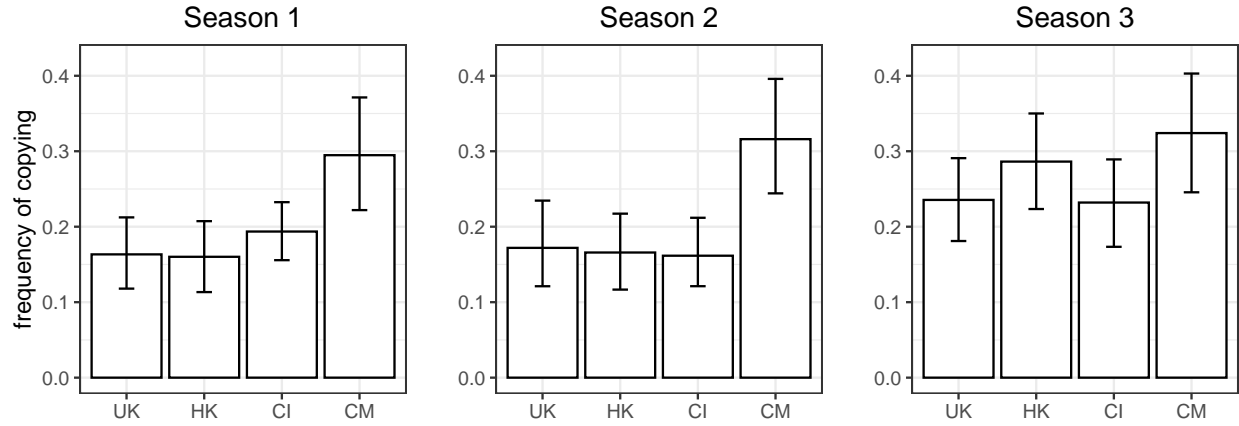


Figure 1: Bar charts as in Mesoudi et al. (2015). UK=British, HK=Hong Kong, CI=Chinese Immigrants, CM=Chinese Mainland.

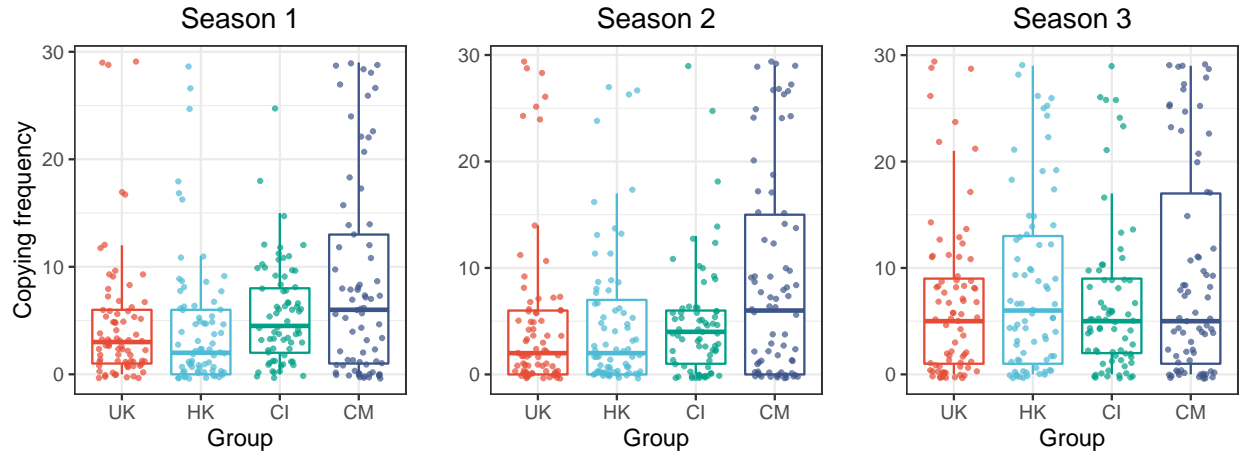


Figure 2: Jitter and box plots

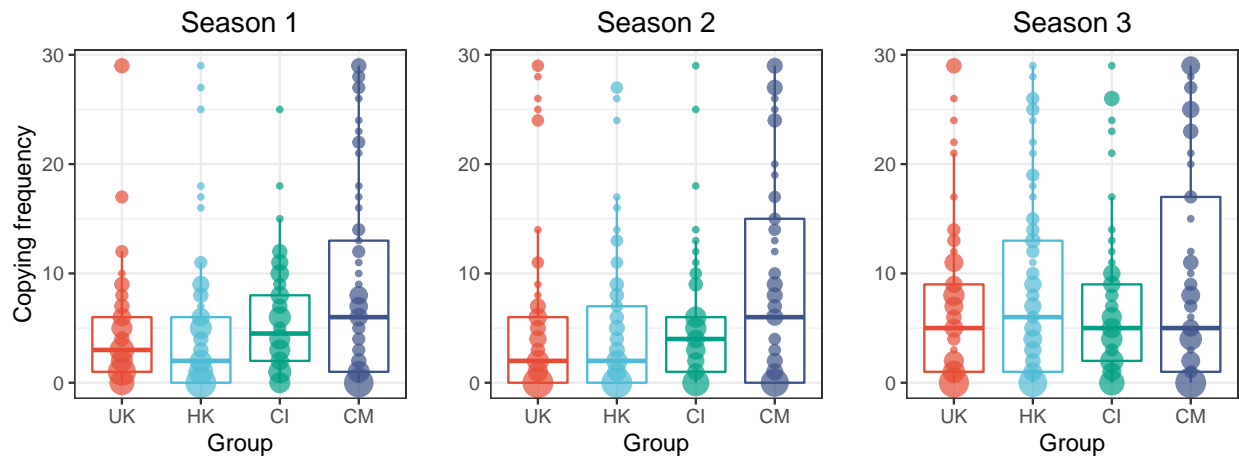


Figure 3: Count and box plots. The size of the circle indicates the number of participants at that value.

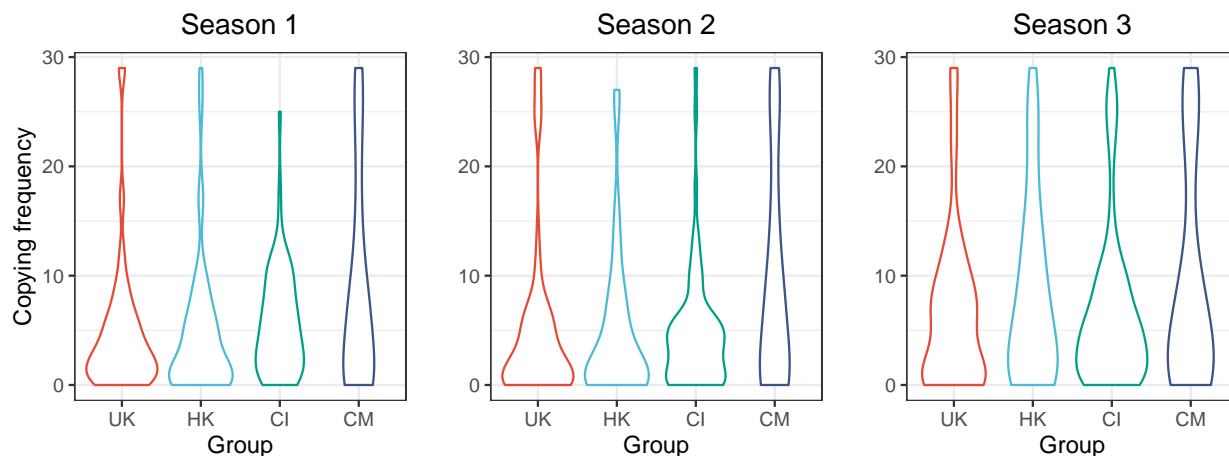


Figure 4: Violin plots, showing narrower bases for CM in Seasons 1 and 2, compared to other groups which are bottom-heavy.

models are compared: an intercept-only model, a model with cultural group as a predictor, and a full model with cultural group, age and sex as predictors. UK is the reference group. The code for the full model of Season 1 is:

```
s1.full.model <- map(
  alist(
    season1_copies ~ dbinom( 29 , p ) ,
    logit(p) <- a + bHK*cultureHK + bCI*cultureCI + bCM*cultureCM + bA*age + bF*female,
    a ~ dnorm(0,1) ,
    bHK ~ dnorm(0,1) ,
    bCI ~ dnorm(0,1) ,
    bCM ~ dnorm(0,1) ,
    bA ~ dnorm(0,0.5) ,
    bF ~ dnorm(0,0.5)
  ) ,
  data=d )
```

with Season 2 and 3 identical except for the dependent measures. I also ran these models with MCMC using `map2stan` but the results were virtually identical, so I do not report this here.

### 3.1 Seasons 1 and 2

For both Seasons 1 and 2 the full model is overwhelmingly best supported:

	WAIC	SE	dWAIC	dSE	pWAIC	weight
s1.full.model	3542.020	289.9277	0.00000	NA	55.28215	9.999997e-01
s1.culture.model	3571.973	292.0297	29.95292	45.67245	39.40754	3.131881e-07
s1.null.model	3688.161	304.2416	146.14044	98.69676	10.22784	1.845042e-32

	WAIC	SE	dWAIC	dSE	pWAIC	weight
s2.full.model	4107.125	320.0513	0.00000	NA	68.39998	9.983338e-01
s2.culture.model	4119.916	318.6873	12.79114	42.62932	47.31743	1.666153e-03
s2.null.model	4285.966	330.6072	178.84066	114.39152	12.25825	1.460560e-39

The full model for Season 1 is shown below, along with exponentiated coefficients to give relative odds (e.g. CM participants are 2.24 times more likely to copy compared to UK participants). HK participants are virtually identical in copying frequency to UK participants. CI are slightly higher, and CM are much higher. Women copy more than men, and age has little effect.

	mean	sd	5.5%	94.5%
a	-0.89138218	0.29352959	-1.36049916	-0.42226520
bHK	-0.01087069	0.08240649	-0.14257219	0.12083080
bCI	0.15896675	0.08072107	0.02995890	0.28797461
bCM	0.80460477	0.07596445	0.68319892	0.92601063
bA	-0.04516066	0.01402102	-0.06756896	-0.02275236
bF	0.31842881	0.05673902	0.22774889	0.40910872

	a	bHK	bCI	bCM	bA	bF
	0.4100885	0.9891882	1.1722990	2.2358127	0.9558439	1.3749657

The full model for Season 2 is shown below and is almost the same as for Season 1. HK participants are again virtually identical to UK participants, CI slightly lower, and CM much higher. Women again copy more than men, and age again has little effect.

	mean	sd	5.5%	94.5%
a	-1.37305798	0.28874802	-1.8345331	-0.911582863
bHK	-0.03179234	0.08095486	-0.1611738	0.097589167
bCI	-0.12815177	0.08276089	-0.2604197	0.004116118
bCM	0.81721331	0.07416131	0.6986892	0.935737400
bA	-0.01821570	0.01369759	-0.0401071	0.003675703
bF	0.32000461	0.05662660	0.2295044	0.410504852

	a	bHK	bCI	bCM	bA	bF
	0.2533311	0.9687077	0.8797199	2.2641815	0.9819492	1.3771341

### 3.2 Season 3

For Season 3, the full model and culture model both received support but the full model slightly more:

	WAIC	SE	dWAIC	dSE	pWAIC	weight
s3.culture.model	4614.164	284.9116	0.00000	NA	46.07585	9.999918e-01
s3.full.model	4638.532	287.9174	24.36795	15.96332	72.17189	5.111671e-06
s3.null.model	4639.510	289.3924	25.34681	57.09625	13.08106	3.133331e-06

The full model is shown below. CI participants are virtually identical to UK participants, while HK and CM are higher. Both age and sex have very weak effects.

	mean	sd	5.5%	94.5%
a	-0.61815013	0.25551463	-1.02651186	-0.209788409
bHK	0.25896841	0.06928701	0.14823439	0.369702435
bCI	-0.02243465	0.07288596	-0.13892049	0.094051189
bCM	0.46124558	0.06894550	0.35105735	0.571433807
bA	-0.02788373	0.01209865	-0.04721971	-0.008547748
bF	0.02007690	0.05048880	-0.06061396	0.100767760

	a	bHK	bCI	bCM	bA	bF
	0.5389405	1.2955929	0.9778151	1.5860483	0.9725014	1.0202798

## 4 Summary

The coefficients and confidence intervals shown here are very similar to the original regression model results shown in Mesoudi et al. (2015), but hopefully more straightforward and understandable. I will soon incorporate task performance into the above, to complete the reanalysis of the original paper.

The data file HKdata.csv and the RMarkdown file containing code for running these reanalyses and producing this document is available at:

[https://github.com/amesoudi/mesoudi\\_chang\\_murray\\_lu\\_2015](https://github.com/amesoudi/mesoudi_chang_murray_lu_2015)

## 5 References

McElreath, R. (2016). Statistical rethinking: A Bayesian course with examples in r and stan. CRC Press.

Mesoudi, A., Chang, L., Murray, K., & Lu, H. (2015). Higher frequency of social learning in China than in the West shows cultural variation in the dynamics of cultural evolution. *Proceedings of the Royal Society B*, 282, 20142209.