Crowdsourcing and phylogenetic modelling reveal parrot tool use is not rare

Amalia P. M. Bastos $^{\dagger,1,2}$ , Scott Claessens $^{*\dagger,2}$ , Ximena J. Nelson $^3$ , David Welch $^4$ , Quentin D. Atkinson $^{\dagger,2}$ , & Alex H. Taylor $^{\dagger,2,3,5,6}$ 

- Department of Psychological & Brain Sciences, Johns Hopkins University, Baltimore, MD, United States
  School of Psychology, University of Auckland, Auckland, New Zealand
  - $^3$  School of Biological Sciences, University of Canterbury, Christchurch, New Zealand
    - $^{\rm 4}$  School of Computer Science, University of Auckland, Auckland, New Zealand
      - $^{5}$ ICREA, Pg. Lluís Companys 23, Barcelona, Spain

† indicates shared first authorship, ‡ indicates shared senior authorship

\* Correspondence concerning this article should be addressed to Scott Claessens, Level 2, Building 302, 23 Symonds Street, Auckland, New Zealand. E-mail: scott.claessens@gmail.com

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 $<sup>^{\</sup>rm 6}$ Institute of Neuroscience, Universitat Autònoma de Barcelona, Barcelona, Spain

### Abstract

Studying the prevalence of putatively rare behaviours, such as tool use, is challenging because absence of evidence can arise either from a species' inability to produce the behaviour or from insufficient research effort. Here, we tackle this challenge by combining crowdsourcing and phylogenetic modelling to approximate actual rates of a rarely observed behaviour based on limited data, targeting tool use in parrots. Crowdsourcing on a social media platform revealed novel instances of tool use in 17 parrot species, more than doubling the confirmed number of tool-using parrot species from 11 (3%) to 28 (7%). Phylogenetic modelling ranked additional species that are most likely to be unobserved tool users, suggesting that between 11% and 17% of extant parrot species may be tool users. These discoveries have implications for inferences about the evolutionary drivers and origins of tool use in parrots, revealing associations with relative brain size and feeding generalism and indicating several genera where tool use was likely an ancestral trait. Overall, our findings challenge the assumption that current sampling effort captures the full distribution of putatively rare animal behaviours. Combining our sampling and analysis methods offers a fruitful approach for investigating the distribution, drivers, and origins of other rare behaviours.

Keywords: tool use; technical intelligence; parrots; crowdsourcing; phylogenetic modelling

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Our understanding of the evolution of animal behaviour is built on the assumption
that we have access to sufficient data<sup>1-3</sup>. However, this is not always the case. Data on
behaviours that are rare, fleeting, or otherwise difficult to observe are likely to be patchy
and incomplete<sup>4,5</sup>. Among species for which such behaviours have not been observed, it can
be difficult to differentiate between cases in which the species is truly incapable of
producing the behaviour and cases in which the species is capable of producing the
behaviour but the behaviour has not yet been observed. Such a distinction can be critical
for drawing conclusions about the rarity and evolution of the behaviour in question.

Comparative work on the evolution of tool use is a paradigmatic example of this 9 issue. The initial discoveries of tool use in chimpanzees<sup>6</sup>, birds<sup>7</sup>, dolphins<sup>8</sup>, and octopuses<sup>9</sup> 10 occurred decades after significant advances on other more easily measurable aspects of 11 their biology. Since these initial discoveries, scholars have used the distribution of reported 12 tool use across species to make various claims about the evolutionary drivers of tool use 13 behaviours. For example, based on the observation that bird species with reported tool use 14 tended to have larger brains, researchers have identified higher relative brain size as a likely 15 precondition for tool using capabilities 10-12. These researchers argue that larger brains are 16 better able to integrate visual and somatosensory information when innovating novel 17 behaviours, such as tool use, in changing environments<sup>13,14</sup>. Similarly, researchers have used 18 existing reports of tool use in birds to debate the roles of generalist versus specialist feeding 19 strategies in driving the evolution of tool use, with some arguing that feeding generalists require technical innovations to expand their dietary niche<sup>13,15,16</sup> and others arguing that feeding specialists require technical innovations for extractive foraging of specific foods<sup>17,18</sup>. 22

However, before we can make claims of this kind, we need to know whether current research effort in the literature is sufficient for robust conclusions to be drawn about the evolution of tool use. In fact, evidence suggests that research effort is often systematically

biased towards particular taxonomic groups, parts of the world that are easy to access, and species with life history traits that make them easier to study, such as larger distribution 27 ranges and population sizes<sup>19</sup>. This is a crucial limitation because insufficient observation 28 may lead researchers to miss true instances of tool behaviours and thus draw premature 29 conclusions about the evolutionary drivers and origins of tool use. While researchers have attempted to deal with this problem by controlling for the number of scientific papers 31 published on different species, previous work has not yet attempted to quantify and 32 explicitly model the relationship between actual tool-using behaviour and what is reported in the scientific literature. If more tool-using species exist than previously thought, this could have important implications for theories of the evolutionary drivers and origins of 35 tool use and, more generally, for our understanding of how rare this behaviour actually is.

One potentially powerful method for quantifying actual rates of rare animal
behaviours is crowdsourcing<sup>20</sup>. In a crowdsourcing study, researchers collect reports from
the general public and/or collate and analyse videos posted on social media platforms.

This citizen science approach has been widely used in ecology to monitor the distributional
patterns of species<sup>21</sup>, but has also recently been used to uncover a variety of rare animal
behaviours, including interspecies play in dogs<sup>20</sup>, novel problem-solving behaviours in
horses<sup>22</sup>, and socially-learned foraging innovations in cockatoos<sup>23,24</sup>. By casting the net
wider than the scientific literature, the crowdsourcing method can provide an indication of
the tool-using species that the literature might be missing.

Even after using this crowdsourcing approach, some tool-users could *still* remain unobserved. An additional approach to identify these unobserved species is to specify a causal model of the process that generates the observed data. We propose one such causal model in Figure 1. In this model, we assume that the presence of tool use in the scientific literature (or in crowdsourced reports) is caused by both unobserved tool use capabilities and the number of published studies (or the number of crowdsourced reports) for any given species. Tool users are more likely to be observed if they are well studied, but understudied

tool users may go undetected. Furthermore, based on existing theories of the evolution of tool use<sup>10–18</sup>, we propose that the unobserved presence or absence of tool use is additionally caused by relative brain size, feeding strategy, and shared phylogenetic ancestry. Expressing this causal model as a statistical model can suggest further species which are likely to be unobserved tool-users and, simultaneously, test existing theories of the evolutionary drivers of tool use without incorrectly assuming that absence of evidence is evidence of absence.

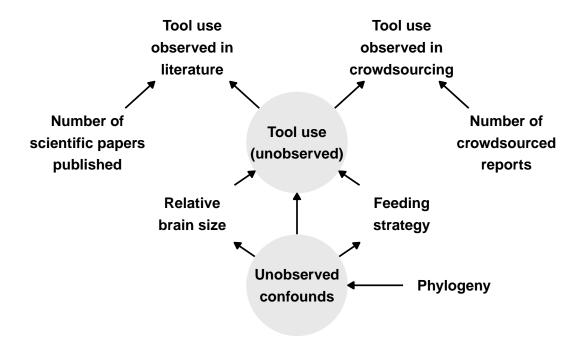


Figure 1. Causal model of observed tool use. Directed acyclic graph of the causal relationships between observed tool use and other variables. Available scientific data on tool use is caused both by unobserved tool use presence and scientific research effort (i.e., number of publications). Available crowdsourced data on tool use is caused both by unobserved tool use presence and crowdsourcing effort (i.e., number of crowdsourced reports). According to theory, unobserved tool use presence should be caused by relative brain size (encephalisation quotient) and feeding strategy (generalist vs. specialist). These variables all share unobserved confounds generated by shared phylogenetic history. Grey circles indicate unobserved variables.

Here, we apply these crowdsourcing and phylogenetic modelling approaches to tool
use in the parrot order. We focus on tool use in parrots for a number of reasons. First, the
scientific literature suggests that only a small proportion of extant parrot species (11 out of
398<sup>25</sup>; 3%) use tools<sup>10,26–38</sup> (note that we do not include in this count species that have

been previously shown to use touchscreen devices as these behaviours were explicitly
trained<sup>39</sup>). Parrot tool use thus provides an ideal test case for examining how robust
sampling is in the scientific literature. Second, parrots are highly popular as pets. Over
70% of all extant parrot species are bred in the aviculture industry and kept as pets
worldwide<sup>40–46</sup>, enabling us to leverage the power of crowdsourcing on a social media
platform to search for evidence of tool use<sup>20</sup>. Third, detailed data on relative brain
sizes<sup>47–52</sup>, feeding strategies<sup>53</sup>, and shared ancestry<sup>54</sup> exist for parrots, allowing us to fit the
statistical model implied by Figure 1 to the entire parrot order.

We first present the results from our crowdsourcing survey, in which we collated videos of tool use in parrots from an online video platform. This survey reveals a number of previously unidentified tool-using parrot species, which we map onto the phylogeny of the parrot order. We then describe our statistical model in more detail, and use it to (i) rank further parrot species that are likely unobserved tool users and (ii) re-examine key hypotheses regarding the evolutionary drivers and origins of tool use in parrots.

77 Results

### Crowdsourcing video survey

We surveyed the social media platform YouTube for video evidence of tool use in parrots (see Methods for detailed search criteria). In our search, we defined "true" tool use behaviour as the manipulation of an unattached object to achieve a goal<sup>55</sup>, while "borderline" tool use involved the use of an object that was still attached to a substrate<sup>56</sup>.

In total, we found 116 videos of 104 individuals from 25 parrot species performing either true tool use (100 videos of 89 individuals from 22 species) or borderline tool use (16 videos of 16 individuals from 7 species). All videos featured pet parrots in captive settings.

In 68 of these videos, owners did not appear to interfere with the subjects' actions. In 43

videos, there was potential interference, either from the owners being in close physical

contact with the bird (e.g., bird perching on hand), talking to the bird, or handing it the tool (which occurred in only two videos). We could not establish the degree of interference in the remaining 5 videos, as sound had been removed or was substituted by music. None of the videos featured owners directly rewarding tool use behaviours with food. All borderline tool use cases were excluded from further analyses.

Of the 22 parrot species performing true tool use, 13 were represented in our video survey by two or more individuals over multiple independent observations. True tool use always involved the subject using an object for self-scratching (95 videos involved scratching the head and/or neck). The most common tool (53 videos) was a moulted feather.

Human-made objects (e.g., pens, spoons, pieces of wood, cardboard) were also common.

According to YouTube video descriptions and owner comments, 45 of the individuals 98 performing true tool use were males and 18 were females. No sex information was provided 99 for the remaining 26 individuals. As owners provided no information on whether sex had 100 been established through genetic testing, and sexual dimorphism in parrots is rare<sup>57,58</sup>, we 101 could not typically ascertain if descriptions were accurate. It is unclear if the 102 disproportionately large number of males in the sample is a consequence of owners more 103 readily assuming their parrots are male when they have not been genetically tested, owners 104 being more likely to own or film male parrots, or male parrots exhibiting more true tool use 105 behaviours than female parrots. 106

Figure 2 maps the findings from the video survey onto a maximum clade credibility
phylogeny for the parrot order, plotted alongside species previously identified in the
scientific literature. Before the video survey, 11 parrot species (3%) had been identified as
tool users in the scientific literature. Across our video survey, we observed true tool use in
22 species, 5 of which overlapped with the scientific literature and 17 of which were novel
species. All of the species identified in the video survey were cockatoos (Cacatuidae), Old
World parrots (Psittacinae), or neotropical parrots (Arinae). The most common species in

our survey, accounting for 41 videos from 37 individuals, was the green-cheeked conure

(Pyrrhura molinae). In accordance with the scientific literature, the video survey found no

evidence of tool use in any species of Psittaculidae, despite this family containing some of

the most commonly kept pet species, including lovebirds, lorikeets, and Asian parakeet

species. Combining both the video survey and the scientific literature, we can thus identify

28 tool-using parrot species overall (7%), compared to the 11 previously reported.

The identification of new tool-using species in our video survey increases the extent 120 to which phylogeny can explain the distribution of tool use in the parrot order. We 121 estimated phylogenetic signal (Pagel's  $\lambda$ ) of tool use using both the pre-video-survey and 122 post-video-survey data. Pagel's  $\lambda$  varies between 0 and 1, where 0 implies that the distribution of a trait across species is unexplained by phylogenetic relatedness and 1 124 implies that the distribution of a trait across species is fully explained by phylogeny. Using 125 the evidence of tool use from the scientific literature alone (pre-video-survey data; 11 126 tool-using species), we estimated an average posterior Pagel's  $\lambda$  of 0.60 (95% credible 127 interval [0.00 0.90]; total n = 174 species). This estimate was moderate-to-strong, but 128 highly uncertain. In comparison, combining the evidence from both the literature and the 120 video survey (post-video-survey data; 28 tool-using species) resulted in a stronger and more 130 certain estimate of phylogenetic signal. With these data, we estimated Pagel's  $\lambda = 0.65$ 131  $(95\% \text{ CI } [0.50 \ 0.77]; \text{ total } n = 174 \text{ species})$ . Thus, the results of our video survey increase 132 the extent to which the distribution of tool use across parrot species can be explained by 133 shared phylogenetic ancestry. This suggests that we can potentially use phylogenetic 134 information, along with other variables, to identify further tool-using parrot species that 135 may remain undetected.

## Phylogenetic survival cure model

In addition to the 28 tool-using species identified in the literature and our video survey, we fitted a Bayesian phylogenetic survival cure model to rank further species that

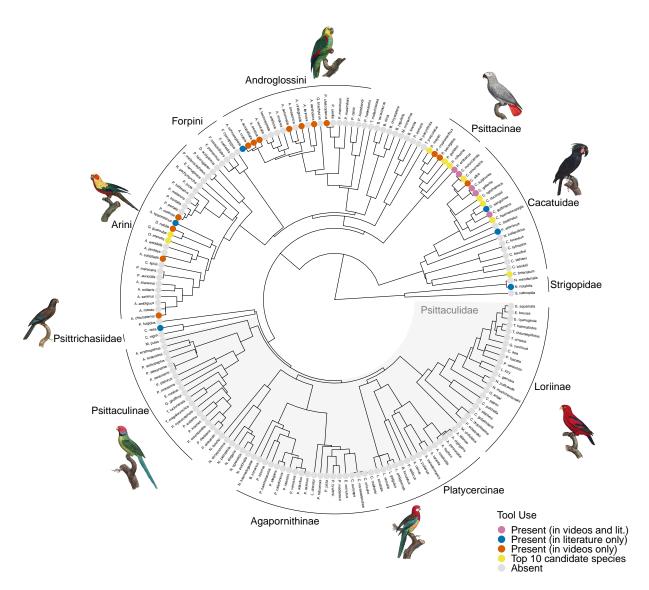


Figure 2. Results of crowdsourcing video survey and phylogenetic survival cure modelling mapped onto a maximum clade credibility phylogeny of the parrot order. Purple tips indicate species observed in the scientific literature and in the video survey. Blue tips indicate species observed only in the scientific literature. Orange tips indicate species observed only in the video survey (note that three species observed only in the video survey are not present in the phylogeny due to a lack of genomic data:  $Psittacara\ erythrogenys,\ Psittacus\ timneh$ , and  $Aratinga\ nenday$ ). Yellow tips indicate the top ten most likely tool-using species from our phylogenetic survival cure model which were not observed in the scientific literature or the video survey. All other tips are labelled as absent (grey). Total n=174 species.

are likely to be undetected tool-users (i.e., tool-using species with no tool use reported in the literature or in crowdsourced videos).

Survival cure models are often used in the medical sciences to analyse the time to
some event, such as cancer relapse, with censored patient data<sup>59</sup>. The data are
right-censored because some patients have relapsed when they are measured, and some
have not yet relapsed when they are measured (i.e., they are censored). The cure aspect of
these models comes from an additional assumption that a certain proportion of the
population is "cured" and can never relapse, no matter how long we measure them for.

Our tool use problem has the same features. We are modelling a time-to-event;
specifically, the amount of "time" (i.e., observation opportunities measured as the number
of published papers or crowdsourced videos) until tool use is identified. This is
right-censored data, because many species will not have had tool use identified when we
measure them. Moreover, we can assume that a certain proportion of the population is
"cured" – that is, they are not tool users, and so we will never identify tool use for them,
no matter how long we measure them for.

In our model, we infer the tool-using status of each species by allowing each species to have their own probability of being "cured" (i.e., not a tool user). Following our causal model (Figure 1), we predict these probabilities based on feeding strategy, encephalisation quotient, and phylogenetic history (see Methods for full model). The model additionally takes research effort into account by allowing that, among species for which tool use is unobserved, all else being equal those with fewer published papers and fewer video search hits have a higher probability of being undetected tool users (Supplementary Figure 1).

We found that this phylogenetic survival cure model was able to adequately
distinguish between species with and without evidence for tool use, with an
area-under-the-curve classification statistic of 0.95 (Supplementary Figure 2). To further
estimate the accuracy of the model's predictions, we also used a leave-one-species-out

approach with known tool users. For each of the 25 tool-using species that were 166 represented on the phylogeny and for which we had brain size and genomic data (we lacked 167 data for three tool-using species), we fitted the model to a modified dataset which set tool 168 use to be absent for the target species in both the scientific literature and the video survey. 169 Across 25 cross-validation models, 18 models (72%) continued to predict the target species 170 as having a median posterior probability of tool use that was within the range of all other 171 tool users. This classification rate was greater than the baseline classification rate of 26%172 for species without evidence of tool use in the full model (38 of 149 species without 173 evidence of tool use had a median posterior probability of tool use that was within the 174 range of the tool-using species). Together, the area-under-the-curve statistic and the 175 leave-one-species-out approach suggest that the model is able to adequately classify known 176 tool users, with some error.

Figure 3 visualises the ranked posterior probabilities of tool use from the phylogenetic 178 survival cure model for all parrot species. As expected, the known tool users are ranked 179 towards the top of this list. However, several "tool use absent" species also rank highly on 180 the list, despite not being identified as tool users in the scientific literature or in our video 181 survey. In fact, according to the model, the most likely tool user is a species for which tool 182 use is unobserved in our data: the blue-eyed cockatoo (Cacatua opthalmica). This species 183 is endemic to Papua New Guinea and is relatively understudied, with only 6 published 184 papers and 596 video search hits, which is fewer than the model expects are necessary to 185 discover tool use when it is present (Figure 4). This species is also found in the Cacatua 186 genus, a clade containing several known tool users. This prediction makes sense given the 187 high phylogenetic signal for tool use reported by the model (Supplementary Figures 3 and 188 4). Beyond the blue-eyed cockatoo, other highly ranked species without observed evidence 189 of tool use are the Meyer's parrot (*Poicephalus meyeri*), the golden parakeet (*Guaruba* 190 guarouba), the long-billed corella (Cacatua tenuirostris), the Solomons cockatoo (Cacatua 191 ducorpsii), the red-fronted parrot (Poicephalus qulielmi), the Cape parrot (Poicephalus 192

robustus), the yellow-eared parrot (Ognorhynchus icterotis), the red-vented cockatoo 193 (Cacatua haematuropyqia), and the gang-gang cockatoo (Callocephalon fimbriatum). Figure 194 2 plots these species on the parrot phylogeny, using the top ten highest ranked species 195 without observed evidence of tool use as an abitrary cutoff for visualisation purposes. 196 The posterior probabilities shown in Figure 3 are estimated with uncertainty, so it is 197 difficult to "identify" any particular species as an undetected tool user. Nevertheless, 198 taking the sum of all the posterior probabilities for the 149 species without observed evidence of tool use, we can estimate that around 26 of those species are likely to be undetected tool users (median sum of probabilities = 25.68, 95\% CI [15.15 41.33]). When 201 combined with the species known to use tools, this implies that between 11% and 17% of

# 204 Implications for the evolutionary drivers and origins of tool use

extant parrot species may be tool users.

203

The predicted probabilities from our phylogenetic survival cure model have 205 implications for inferences about the evolutionary drivers and origins of tool use in the 206 parrot order. Regarding the drivers of tool use hypothesised in Figure 1, the phylogenetic 207 survival cure model revealed that encephalisation quotient strongly positively predicted the 208 probability of tool use (median posterior log odds slope = 1.12, 95\% CI [0.39 2.00]; total n 209 = 174 species; Figure 5). This helps explain the ranking in Figure 3: the blue-eyed 210 cockatoo has the largest relative brain size in the dataset. We also found that feeding 211 generalist species were slightly more likely to be tool users, though the posterior difference 212 between generalists and specialists was quite uncertain (median posterior log odds difference = 0.33, 95% CI [-1.13 1.76]; total n = 174 species). These results from the 214 survival cure model differed from the results of models fitted to pre-video-survey and post-video-survey data without the survival cure component, which found no effect of 216 relative brain size and no difference between feeding strategies, respectively 217 (Supplementary Figure 5). 218

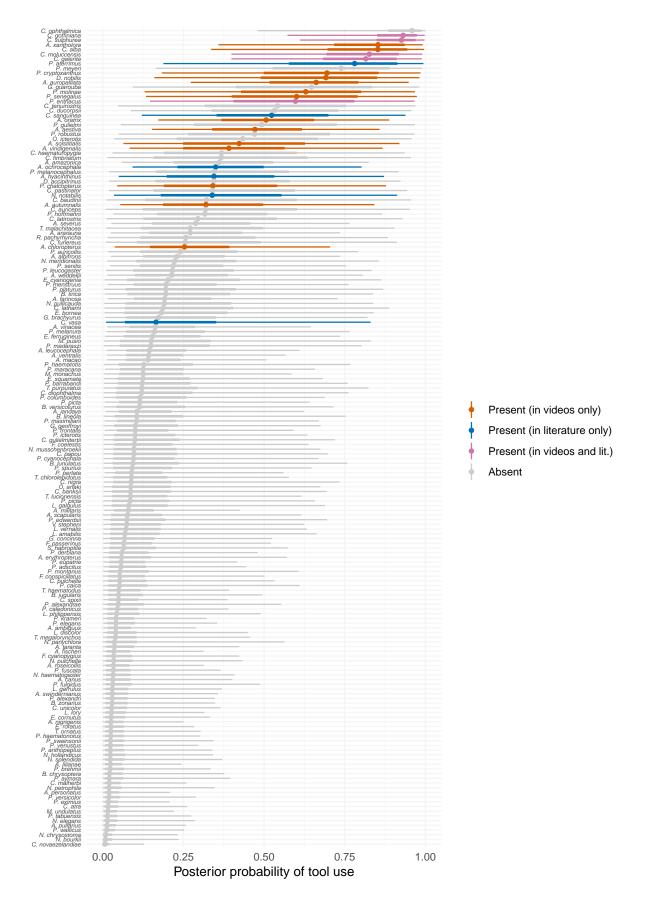


Figure 3. Posterior predicted probabilities of tool use for each species from our phylogenetic survival cure model. Points are posterior medians and lines are 50% and 95% credible intervals. Total n=174 species.

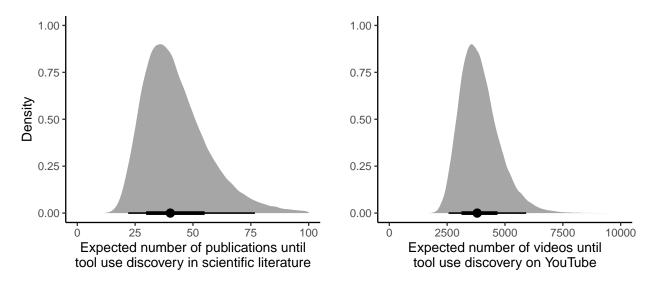


Figure 4. Expected number of published papers and videos until tool use discovery, according to the survival component of the phylogenetic survival cure model. Densities are full posterior distributions, points are posterior medians, and lines are 50% and 95% credible intervals.

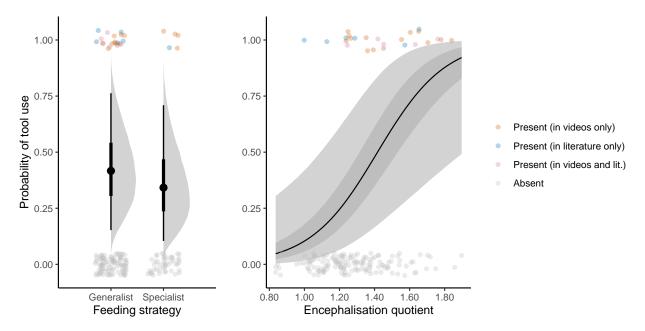


Figure 5. Posterior predictions for the effects of feeding strategy and encephalisation quotient on the probability of tool use from the phylogenetic survival cure model. In the left plot, points and lines represent posterior medians and 50% and 95% credible intervals, with densities representing full posterior distributions. In the right plot, the line and shaded area represents the posterior median regression line with 95% credible intervals. In both plots, individual species are coloured according to the presence / absence of tool use in the video survey and the scientific literature. Total n = 174 species, generalist n = 121 species, specialist n = 53 species.

Table 1 Estimated probabilities of tool use for most recent common ancestors of several parrot genera. Probabilities estimated using exploratory ancestral state reconstruction models fitted to the pre-video-survey data, post-video-survey data, and predicted probabilities from the phylogenetic survival cure model. Total n = 174 species.

Genus	Pre-video-survey	Post-video-survey	Survival cure probabilities
Amazona	0.06, 95% CI [0.01 0.17]	0.21, 95% CI [0.04 0.66]	0.48, 95% CI [0.05 0.84]
Cacatua	0.08, 95% CI [0.03 0.20]	0.17, 95% CI [0.03 0.45]	0.81, 95% CI [0.23 0.99]
Nestor	0.11, 95% CI [0.05 0.29]	0.21, 95% CI [0.12 0.37]	0.53, 95% CI [0.29 0.66]
Poice phalus	0.06, 95% CI [0.02 0.11]	$0.19, 95\% \text{ CI } [0.12 \ 0.41]$	$0.72, 95\% \text{ CI } [0.16 \ 0.90]$

Regarding the origins of tool use, we fitted exploratory ancestral state reconstruction 219 models to the pre-video-survey data, the post-video-survey data, and the predicted 220 probabilities from the phylogenetic survival cure model. The discoveries from our video 221 survey and from our phylogenetic modelling increased the likelihood that tool use was 222 present in the most recent common ancestor for several parrot genera. These include the 223 amazon parrots native to the Americas (Amazona), the true white cockatoos and corellas 224 found in South East Asia and Australasia (Cacatua), the kea and the kākā from New 225 Zealand (Nestor), and the Poicephalus genus native to Africa (Table 1; Supplementary 226 Figures 6 - 8). These findings suggest that species from each of these genera share their 227 tool use capabilities via common descent from their respective common ancestors, rather 228 than via independent evolution within each genus. 229

230 Discussion

Since the earliest anecdotal reports of parrots using tools in the 1970s<sup>60</sup>, only 11
parrot species (3% of all extant parrot species) have been documented as tool users in the
scientific literature. Our study used crowdsourcing to identify 17 additional tool-using
parrot species that are new to science, more than doubling the overall count to 28 species
(7%). These species consisted of cockatoos (Cacatuidae), Old World parrots (Psittacinae),

236 and neotropical parrots (Arinae).

We found strong phylogenetic signal in our extended dataset, demonstrating that the
distribution of tool use across parrot species is affected by shared phylogenetic ancestry.

This allowed us to use phylogenetic information, along with other variables, to infer the
unobserved probabilities of tool use across the parrot order. Our phylogenetic survival cure
model incorporated information on phylogenetic history, research effort, relative brain size,
and feeding specialisation to rank parrot species that were most likely to be undetected
tool users. The sum of probabilities from this model implied that between 15 and 41 of the
species without observed evidence of tool use are likely to be undetected tool users,
suggesting that the true proportion of tool users may be as high as 17%.

These findings have a number of important implications. First, our findings show 246 that current research effort in the scientific literature is insufficient to capture the real 247 world occurrence of parrot tool use. If the scientific literature had sampled the natural 248 world sufficiently, we would expect to see close correspondence between those species 249 reported as tool users in the literature and those species the public have filmed performing 250 tool use. Instead, we discovered a large discrepancy between these two data sources, both 251 in the prevalence of tool use and the species identified. This raises the possibility that 252 insufficient research effort is a general issue across the scientific literature, both for tool use 253 in other groups and for other rare behaviours. 254

Second, in terms of the evolution of tool use in parrots, our study challenges a key
assumption made in the literature to date: that only a minority of parrots are tool
users<sup>30,31,34,37,38</sup>. The paucity of evidence for tool use across parrots in the literature
initially implied that tool use may have evolved independently in different parrot species.
Our discovery of the widespread distribution of tool use across the parrot phylogeny, along
with the strong phylogenetic signal in this expanded dataset, challenges this and suggests
that, at least for some parrot clades, the capacity for tool use might be a homologous trait

that has been evolutionarily conserved. Our exploratory ancestral state reconstruction
analysis provides preliminary support for this hypothesis, revealing an increased
probability of tool use among the most recent common ancestors for the *Amazona*, *Cacatua*, *Nestor*, and *Poicephalus* genera. Even at this preliminary stage, our analysis
therefore raises an alternative hypothesis for the observed tool use in *Cacatua*<sup>30,38</sup> and *Nestor*<sup>26–29,31,34</sup>, namely that tool behaviours have arisen due to the common ancestor
being a tool user, rather than from behavioural innovation within a species.

Third, our results support existing theories of the drivers of tool use. We found that 269 encephalisation was strongly positively related to the probability of tool use in our phylogenetic model, supporting previous theories linking relative brain size to increased tool innovation in  $birds^{10-12}$  and  $primates^{61}$ . Nevertheless, we recognise that cross-species correlations between brain size and behaviour are challenging to interpret causally<sup>3,4</sup>, and 273 thus encourage further work on the specific neural correlates of technical intelligence in 274 parrots (see e.g. 62). In our phylogenetic model, we also found that tool use was somewhat 275 more likely among feeding generalists compared to feeding specialists, although this 276 difference was uncertain. This trend supports previous suggestions that increased cognitive 277 abilities and technical innovation rates are required to expand a generalist species' dietary 278 niche<sup>15,17,63</sup>. However, the trend contradicts theories linking tool use to dietary 279 specialisation, whereby species eating specific foods that require extractive foraging have 280 higher cognitive ability and are especially prone to using tools (reviewed in ref<sup>17</sup>). 281

We believe that the observations of tool use in our video survey reflect intentional
tool use, rather than accidental object manipulations or trained behaviours. In the
crowdsourcing video survey, individual parrots often used tools slowly and repetitively over
long periods of time, even across multiple different videos, suggesting that their behaviour
was not accidental<sup>64</sup>. Moreover, for 60% of the species we report on, we found two or more
videos of repetitive and sustained scratching by different individuals in separate
households. Following the assumption that accidental tool use (e.g., briefly holding an

object and attempting to scratch oneself simultaneously) is rare<sup>65</sup>, the repeated observation 289 of the same tool use behaviours for scratching by multiple animals of the same species 290 across multiple households suggests that these manipulations are intentional and recurring 291 events rather than rare "accidents" and likely do not represent unusual stereotypies of a 292 single individual. We also found little evidence to suggest that the tool use behaviours were 293 trained or unintentionally cued by the birds' owners. Over half of the videos coded 294 contained no evidence of human interference with the parrots' tool use aside from filming 295 the behaviour. Humans only handed parrots their tools in two of the videos, and none of 296 the videos featured owners directly rewarding tool use behaviours with food. Finally, and 297 perhaps most importantly, the high levels of phylogenetic signal in our data provides strong 298 evidence that the observations from our video survey reflect biologically-endowed capacities 299 for tool use rather than accidents or trained behaviours, which would likely appear uniformly across the phylogeny. 301

While missing data imputation is becoming more common in phylogenetic analyses<sup>66</sup>, 302 the important distinction between absence of evidence and evidence of absence has not 303 been given as much attention. Our phylogenetic analysis provides one approach to this 304 problem by distinguishing between true absences of tool use and absences of tool use due 305 to a lack of research effort in the scientific literature or in crowdsourced videos. To achieve 306 this, we explicitly modelled the measurement of the outcome variable along a research 307 effort time series, such that species with lower research effort in the literature or in videos 308 were likely to be censored. In line with our causal model, we also included relative brain 300 size, feeding strategy, and phylogenetic history as predictors of unobserved tool use. We 310 encourage researchers to test this model by directing future study efforts towards the 311 parrot species with the highest probabilities of being undetected tool-users. Future work 312 should also refine the causal model in Figure 1 to provide more certain estimates of tool 313 use probabilities, either by including additional predictor variables or modelling further 314 causes of measurement error in the taxonomic record, such as species abundance and

316 geographic accessibility<sup>19</sup>.

In conclusion, we have shown that the scientific literature has insufficiently captured 317 the full distribution of tool use in the parrot order. Our crowdsourcing survey has more 318 than doubled the number of known tool using parrot species from 11 to 28, and our 319 phylogenetic inference suggested that the true proportion of parrot tool users could be as 320 high as 17%. These discoveries have implications for theories of the evolutionary drivers 321 and origins of tool use in parrots. Beyond parrot tool use, the crowdsourcing and phylogenetic methods used in this study have the potential to be applied to other rarely 323 observed behaviours, including tool use in other taxa<sup>67</sup>, rhythmic entrainment in birds<sup>68–71</sup>, 324 teasing behaviours in primates<sup>72</sup>, and tactical deception across the animal kingdom<sup>73–75</sup>. 325 We hope that these methods will continue to uncover a diverse array of ephemeral 326 behaviours that have as yet gone undetected in the scientific literature. 327

328 Methods

### Video searches and coding

Our video search was conducted on YouTube in July 2020. Search terms included 330 "parrot using tool" and variants (e.g., "macaw using tool", "lorikeet using tool", "parakeet using tool"), "tool use in parrot", "parrot tool use", "parrot scratching itself" (included 332 after we found several videos demonstrating self-care tool use in previous searches) and 333 equivalent terms (e.g., "parrot preening itself", "parrot grooming itself", "parrot 334 scratching"). For all species that did not display results including object manipulation or 335 scratching behaviours, we also searched the species' common name(s) + "tool use", as well 336 as the species' common name(s) + "scratching". We also searched for translations of the 337 terms "parrot tool use" and "parrot scratching" in languages for all countries where bird 338 ownership was reported as  $>5\%^{76}$ , namely, Turkish, Czech, Polish, French, Italian, Dutch, 339 German, Russian, Spanish, Portuguese, and Mandarin. 340

When we found a relevant video, we also searched for similar content uploaded by the
same person/channel. For each YouTube search conducted, we watched all relevant videos
until we reached five consecutive videos that did not feature any parrots. At this point, we
ended that search and initiated the next search. In line with previous recommendations<sup>20</sup>,
we planned to exclude any videos that consisted of four or more shots edited together so as
to ensure the behaviours being observed were not edited or manipulated, but none of the
videos obtained qualified for exclusion.

All videos featuring parrots manipulating objects were investigated for potential tool 348 use or borderline tool use. We defined tool use as the manipulation of an unattached object as an extension of the beak or foot to achieve a goal towards another object, individual, or 350 oneself<sup>55</sup>. Borderline tool use was similarly defined, except that it involved the use of an 351 object that was still attached to a substrate<sup>56</sup>. For example, if individuals used a fallen 352 feather or stick for self-scratching this was defined as tool use, but using one's currently 353 attached tail feathers or cage furnishings for the same purpose was defined as borderline 354 tool use. Self-scratching had to involve slow and repeated movements of touching an object 355 to one's body (or, in the case of borderline tool use, rubbing repetitively against an 356 attached object<sup>64</sup>). 357

All relevant videos were coded for video length, species, tool use presence

(yes/borderline), tool use type (e.g., scratching, feeding), the object being used (e.g.,

feather, stick), tool use target, human interference with the action (talking or handing

object to parrot, holding parrot), and the number of shots within each video. Our complete

dataset also includes the name for each video, link, subject name, sex (as declared by

owner, as most parrot species are not sexually dimorphic), publishing date, and dates

found and coded.

## Data for parrot species

We collected data for 194 parrot species (Supplementary Figures 9 - 11). We 366 gathered feeding strategy data as a dichotomous variable ("generalist" or "specialist") from 367 the EltonTraits ecological database<sup>53</sup>. As per the database, specialists were defined as 368 species whose diet comprised at least 70% of a single food source. To calculate relative 369 brain size, we collated data from the literature for all known body mass (g) and brain mass 370 (g) values across parrots<sup>47–52</sup>. For all species for which we obtained body and brain mass 371 data, we calculated the encephalisation quotient (EQ) using the following formula<sup>77</sup>: 372  $BrainWeight/(0.12*BodyWeight(\frac{2}{3}))$ . We found body mass and brain mass data for a 373 total of 194 parrot species. This included all tool-using species in our video dataset, with the exception of three species: Diopsittaca nobilis, Psittacara erythrogenys, and Coracopsis vasa. For the latter, we used values for the closely related Coracopsis nigra. The other two species were excluded from the final dataset. 377

For modelling purposes, we coded research effort in both the scientific literature and 378 the crowdsourced videos. For the scientific literature, we operationalised research effort as 379 the number of papers published for each species' Latin name up to and including the first 380 paper containing tool use for that species. If no tool use had been identified in the 381 scientific literature for a species, then we coded the total number of papers published to 382 date. We used the scientific database Scopus for coding the number of published papers. 383 For the crowdsourced videos, we coded research effort as the number of search hits for each 384 species on YouTube. If tool use had been identified on YouTube, we estimated the number 385 of search hits when the first video of tool use was published on YouTube, assuming linear growth of search hits since the inception of YouTube. If tool use had not been identified, 387 we used the current number of search hits. 388

For phylogenetic data, we used the phylogenetic tool at www.birdtree.org<sup>54</sup> to compile 1000 posterior draws of phylogenetic trees for 174 of the 194 parrot species for

which both EQ and genomic data exist. A single maximum clade credibility tree was
generated from these posterior draws for visualisation purposes. In our analyses, we
iterated over posterior draws of the phylogeny to account for phylogenetic uncertainty.

## 394 Phylogenetic signal

We used the *fitDiscrete* function in the *ape* R package<sup>78</sup> to calculate phylogenetic signal, for both the pre-survey and post-survey tool use data. We iterated the model over 100 posterior parrot phylogenies to incorporate phylogenetic uncertainty.

# Causal model of tool use

To infer unobserved probabilities of tool use across parrots, we proposed a causal 399 model of observed tool use (Figure 1). We assumed that observed tool use in the scientific 400 literature and in the crowdsourced videos is caused by both the unobserved presence or 401 absence of tool use and research effort, proxied by the number of papers published on a 402 species and the number of videos published on a species. Tool users are more likely to be 403 observed if they are well studied, but understudied tool users may go undetected. In 404 addition, based on theory, we also assumed that unobserved tool use is caused by feeding 405 strategy and relative brain size<sup>10–13,15–17</sup>. Finally, we assumed that shared phylogenetic 406 history causes unobserved confounding and non-independence in unobserved tool use, 407 feeding strategy, and relative brain size across the parrot phylogeny.

# Bayesian phylogenetic survival cure model

Given our proposed causal model, we constructed a statistical model to impute unobserved probabilities of tool use and test existing theories of the evolution of tool use in parrots. To understand the model, suppose that we have the following observed variables for parrot species i. For the scientific literature, we declare  $N_{\text{Lit},i}$  as the number of papers

published before and up to tool use identification for species i (or, if tool use has not been 414 identified, the total number of papers published for species i) and  $\mathcal{T}_{\mathrm{Lit},i}$  as a binary variable 415 stating whether (1) or not (0) tool use has yet been observed in the scientific literature for 416 species i. For the crowdsourced videos, we declare  $N_{Vid,i}$  as the number of videos published 417 before and up to tool use identification for species i (or, if tool use has not been identified, 418 the total number of videos published for species i) and  $T_{Vid,i}$  as a binary variable stating 419 whether (1) or not (0) tool use has yet been observed in the crowdsourced videos for 420 species i. Additionally,  $F_i$  and  $EQ_i$  are feeding strategy and encephalisation quotient values 421 for species i and we have a phylogenetic distance matrix D that describes the patristic 422 distances between all parrot species. 423

We assume that species i is a non-tool-user (i.e. "cured") with some probability  $p_i$ .

We also assume that tool use is identified in the scientific literature and the crowdsourced videos at constant rates  $\lambda_{\text{Lit}}$  and  $\lambda_{\text{Vid}}$  following exponential survival functions. Given these assumptions, we can then describe the different ways in which variables  $N_{\text{Lit}}$  and  $N_{\text{Vid}}$  can be distributed. Focusing on the scientific literature, if tool use has been observed  $(T_{\text{Lit},i}=1)$ , then the likelihood for  $N_{\text{Lit},i}$  is:

$$\Pr(N_{\text{Lit},i}|T_{\text{Lit},i} = 1, p_i, \lambda_{\text{Lit}}) = (1 - p_i) \cdot \text{Exponential}(N_{\text{Lit},i}|\lambda_{\text{Lit}})$$
(1)

On the other hand, if tool use has not yet been observed ( $T_{Lit,i} = 0$ ), there are two ways that the outcome variable could have been realised. First, the species could be a non-tool-user with probability  $p_i$ . Second, the species could be a tool-user with probability ( $1 - p_i$ ) that has been censored and has not had its tool use measured yet. Together, then, the likelihood for  $N_{Lit,i}$  is:

$$\Pr(N_{\text{Lit},i}|T_{\text{Lit},i} = 0, p_i, \lambda_{\text{Lit}}) = p_i + ((1 - p_i) \cdot \text{Exponential-CCDF}(N_{\text{Lit},i}|\lambda_{\text{Lit}}))$$
(2)

The Exponential-CCDF function allows for the censored nature of the data. The same data generating process is assumed to underlie the crowdsourced videos.

We define the mixture likelihood SurvivalCure as the distribution above, with parameters p (the probability of being a "cured" non-tool-user) and  $\lambda$  (the rate of the exponential distribution). We use an Ornstein-Uhlenbeck Gaussian process<sup>79</sup> to model phylogenetic covariance. Below, we specify the full model with priors:

$$N_{\text{Lit},i} \sim \text{SurvivalCure}(\lambda_{\text{Lit},i}, p_i)$$

$$N_{\text{Vid},i} \sim \text{SurvivalCure}(\lambda_{\text{Vid},i}, p_i)$$

$$\lambda_{\text{Lit},i} = 1/\exp(\gamma_{\text{Lit}})$$

$$\lambda_{\text{Vid},i} = 1/\exp(\gamma_{\text{Vid}})$$

$$\log \text{it}(p_i) = \alpha_{\text{FEEDING}[i]} + \beta \text{EQ}_i + k_{\text{SPECIES}[i]}$$

$$\begin{pmatrix} k_1 \\ k_2 \\ \dots \\ k_n \end{pmatrix} \sim \text{MVNormal} \begin{pmatrix} 0 \\ 0 \\ \dots \\ 0 \end{pmatrix}, \mathbf{K}$$

$$\begin{pmatrix} \mathbf{K}_{\text{I}} \\ \mathbf{K}_{\text{I}} \\ \dots \\ \mathbf{K}_{\text{I}} \end{pmatrix} = \eta^2 \exp(-\rho^2 D_{ij})$$

$$\gamma_{\text{Lit}}, \gamma_{\text{Vid}}, \alpha_{1,2}, \beta \sim \text{Normal}(0, 1)$$

$$\eta^2, \rho^2 \sim \text{Exponential}(0.5)$$

The priors for this model produce reasonable prior predictions of the probabilities of

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tool use for each parrot species (Supplementary Figure 12), but a sensitivity analysis 442 revealed that the ranking and posterior probabilities reported in the main text were robust 443 to modifying these priors (Supplementary Figure 13). We estimated the posterior 444 distribution of this model using Hamiltonian Monte Carlo as implemented in Stan version 445 2.26.180. We ran the model for 4000 samples, with 2000 warmup samples, and iterated the 446 model over 100 posterior parrot phylogenies to incorporate phylogenetic uncertainty. R-hat 447 values and effective sample sizes suggested that the model converged normally. Trace plots 448 are reported in Supplementary Figure 14. We report equal-tailed credible intervals to describe the posterior distribution of this model in the main text. 450

To validate our method, we fitted the model to 100 simulated datasets with known parameters. The model was able to successfully recover those parameters (Supplementary Figure 15). We also ran a leave-one-species-out exercise to ensure that we could accurately predict known tool users. We repeated this approach for each known tool user by setting observed tool use to zero. Cross-validation results are reported in the main text.

#### 456 Ancestral state reconstruction

To determine whether the identification of novel tool-using species has implications
for our understanding of the evolutionary origins of tool use in parrots, we fitted three
exploratory ancestral state reconstruction models. We used the ancThresh function from
the phytools R package<sup>81</sup>, iterating the function over 100 posterior parrot phylogenies. This
function estimates discrete ancestral states by assuming the evolution of a latent
continuous variable following an Ornstein-Uhlenbeck process. We fitted this model to three
different outcome variables: (i) presence vs. absence of tool use in scientific literature only,
(ii) presence vs. absence of tool use in literature and/or videos, and (iii) the median
predicted probabilities of tool use from the phylogenetic survival cure model.

# Reproducibility

All analyses were conducted in R v4.2.1.<sup>82</sup>. Visualisations were produced using the  $ggplot2^{83}$  and  $cowplot^{84}$  packages. The manuscript was reproducibly generated using the  $targets^{85}$  and  $papaja^{86}$  packages. Code to reproduce all analyses and figures can be found here: https://github.com/ScottClaessens/phyloParrot

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### **Author Contributions**

All authors contributed to the conceptualisation of the paper. A.P.M.B., X.J.N., and A.H.T. developed the video search methodology. S.C., D.W., and Q.A.D. developed the statistical models and analysed the data. All the authors wrote the manuscript and approved the final version for submission.

# **Competing Interests**

The authors declare no competing interests.

### Data Availability

All data used in this study are publicly available on GitHub: https://github.com/ScottClaessens/phyloParrot

# Code Availability

All code to reproduce the analyses in this study are publicly available on GitHub: https://github.com/ScottClaessens/phyloParrot

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## **Supplementary Information**

Crowdsourcing and phylogenetic modelling reveal parrot tool use is not rare Amalia P. M. Bastos<sup>1,2</sup>, Scott Claessens<sup>2</sup>, Ximena J. Nelson<sup>3</sup>, David Welch<sup>4</sup>, Quentin D. Atkinson<sup>2</sup>, Alex H. Taylor<sup>2,3,5,6</sup>

<sup>&</sup>lt;sup>1</sup> Department of Psychological & Brain Sciences, Johns Hopkins University, Baltimore, MD, United States

<sup>&</sup>lt;sup>2</sup> School of Psychology, University of Auckland, Auckland, New Zealand

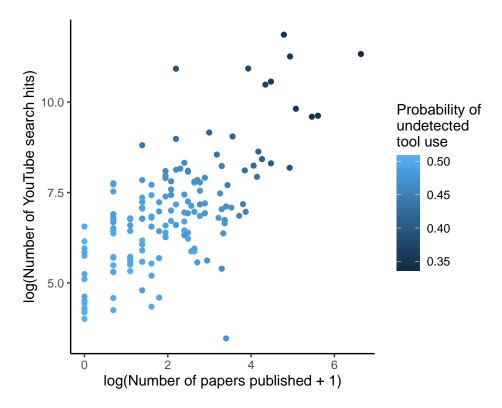
<sup>&</sup>lt;sup>3</sup> School of Biological Sciences, University of Canterbury, Christchurch, New Zealand

<sup>&</sup>lt;sup>4</sup> School of Computer Science, University of Auckland, Auckland, New Zealand

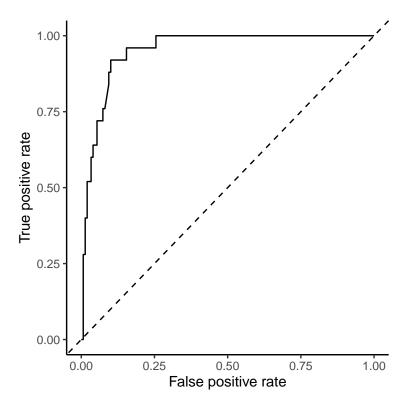
<sup>&</sup>lt;sup>5</sup> ICREA, Pg. Lluís Companys 23, Barcelona, Spain

<sup>&</sup>lt;sup>6</sup> Institute of Neuroscience, Universitat Autònoma de Barcelona, Barcelona, Spain

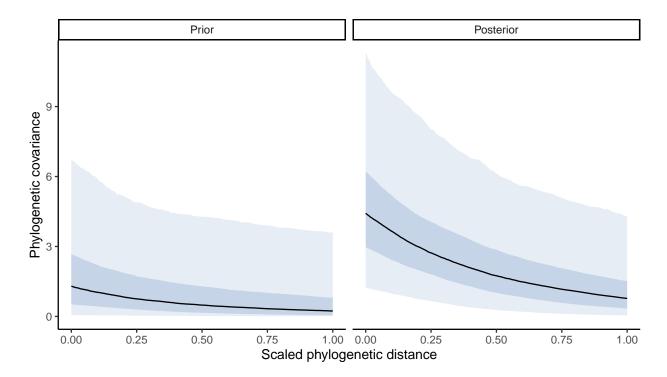
## Supplementary Figures



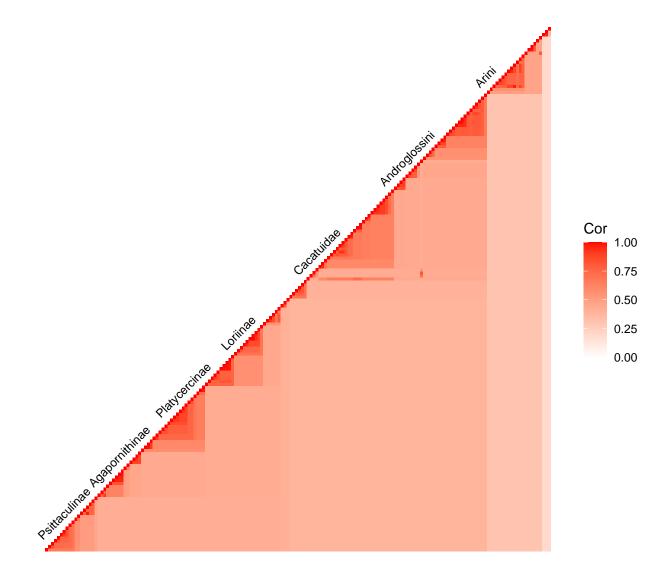
Supplementary Figure 1. Median posterior probabilities of undetected tool use for each parrot species without observed evidence of tool use from reduced model. This reduced version of the phylogenetic survival cure model does not contain relative brain size or feeding strategy as predictors, nor does it contain any phylogenetic covariance. The only information included in the model is the number of papers published and the number of YouTube search hits for each species. Each point is a parrot species without observed evidence of tool use, and the colour of the points scales with the probability of undetected tool use. All else being equal, those species with fewer published papers and fewer YouTube search hits have a higher probability of being undetected tool users.



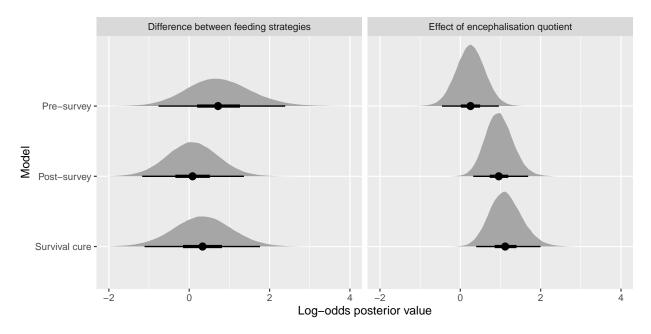
Supplementary Figure 2. Receiver operating characteristic (ROC) curve for the phylogenetic survival cure model. The area-under-the-curve in this plot is 0.95, suggesting that the model is able to adequately classify observed tool users and non-tool users.



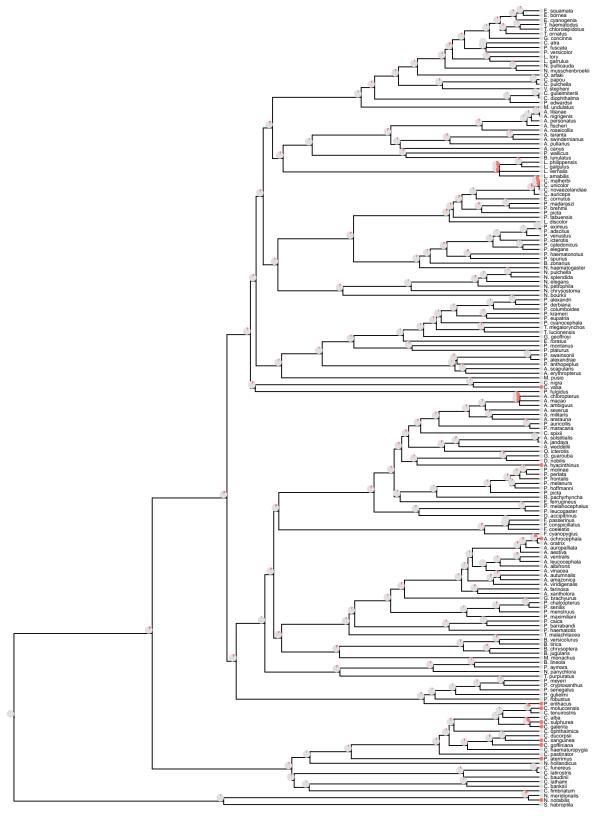
Supplementary Figure 3. Prior and posterior phylogenetic covariance functions from the Bayesian survival cure model fitted to the full dataset. Lines are median posterior functions and shaded areas are 50% and 95% credible intervals.



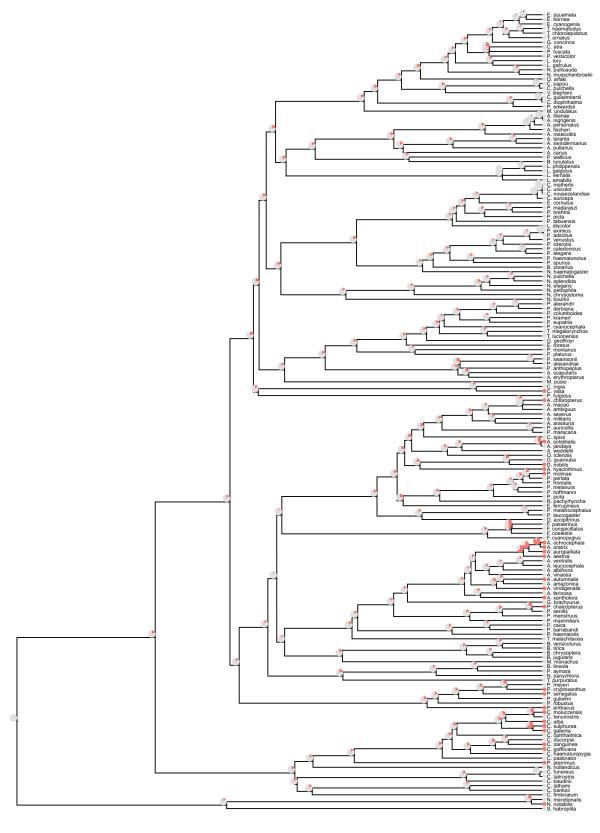
Supplementary Figure 4. Between-species correlation matrix implied by the posterior phylogenetic covariance function from the Bayesian survival cure model. Correlations are median posterior estimates. Individual species names omitted for space reasons.



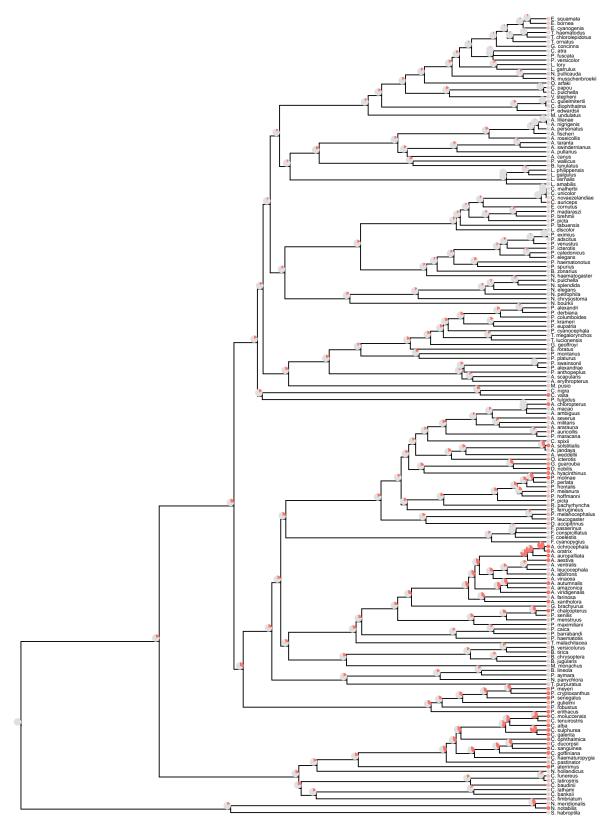
Supplementary Figure 5. Comparing results between the survival cure model and models fitted to the pre-survey and post-survey data without any survival cure component. Densities are full posterior distributions from three separate models iterated over 100 posterior parrot phylogenies. Points represent posterior medians, and lines represent 50% and 95% credible intervals.



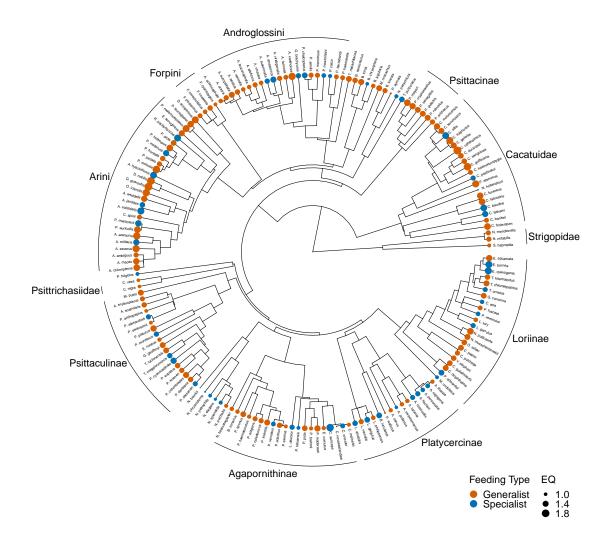
Supplementary Figure 6. Results of exploratory ancestral state reconstruction analysis fitted to pre-video-survey data, represented on a maximum clade credibility tree. Tip nodes represent the presence (red) or absence (grey) of observed tool use in the scientific literature. Pie charts represent the posterior probability of tool use presence at each ancestral node.



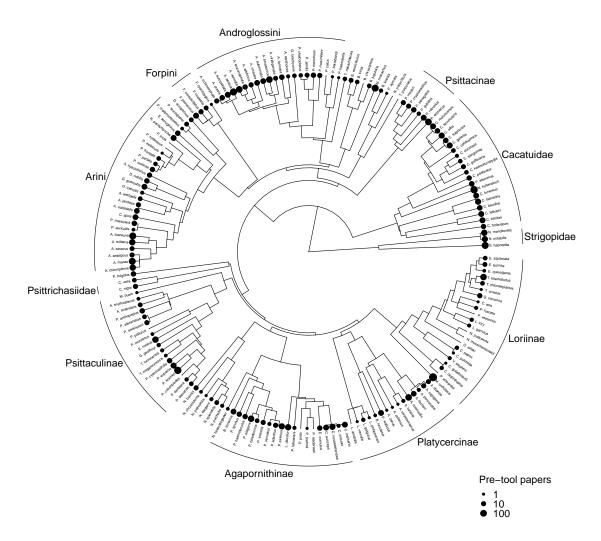
Supplementary Figure 7. Results of exploratory ancestral state reconstruction analysis fitted to post-video-survey data, represented on a maximum clade credibility tree. Tip nodes represent the presence (red) or absence (grey) of observed tool use in the scientific literature and the video survey. Pie charts represent the posterior probability of tool use presence at each ancestral node.



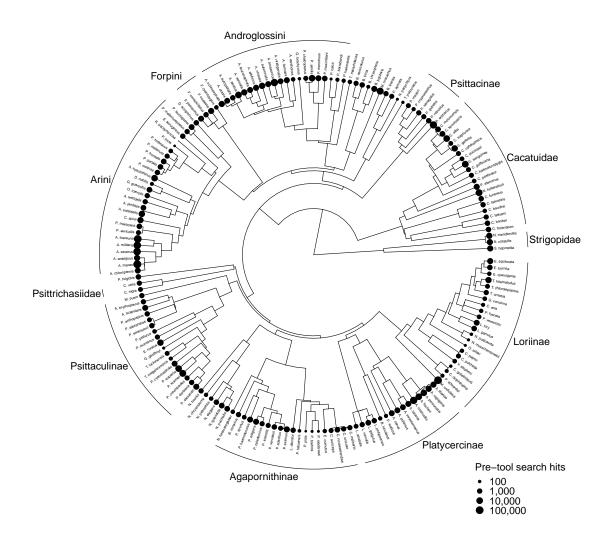
Supplementary Figure 8. Results of exploratory ancestral state reconstruction analysis fitted to predicted probabilities from the phylogenetic survival cure model, represented on a maximum clade credibility tree. Tip nodes represent the median posterior predicted probabilities of tool use from the phylogenetic survival cure model, with more red indicating an increasing probability of tool use presence and more grey indicating a decreasing probability of tool use presence. Pie charts represent the posterior probability of tool use presence at each ancestral node.



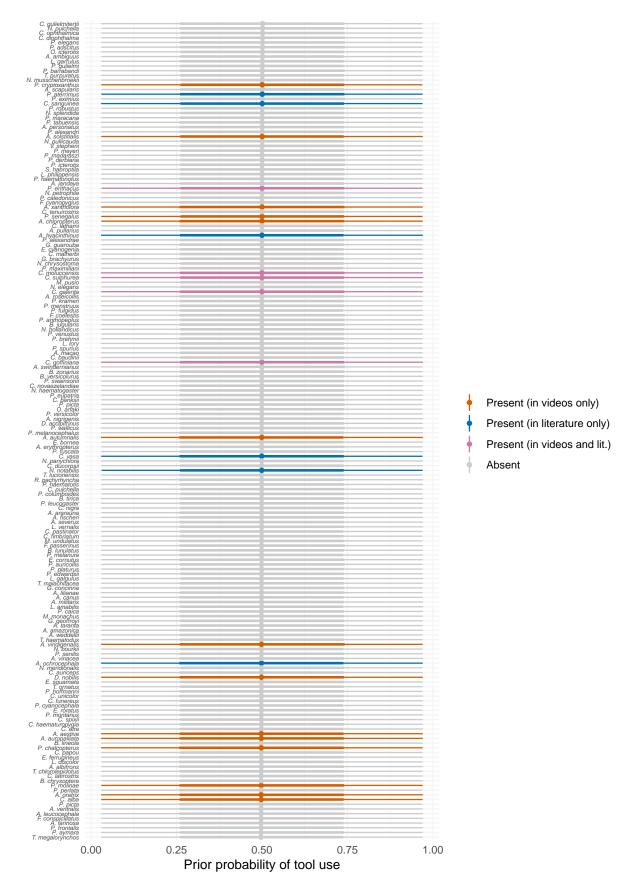
Supplementary Figure 9. Data on encephalisation quotient and feeding strategy for all parrots, presented on a maximum clade credibility tree. Tip points are coloured according to feeding generalism (orange) and specialism (blue), and scaled according to encephalisation quotient (EQ).



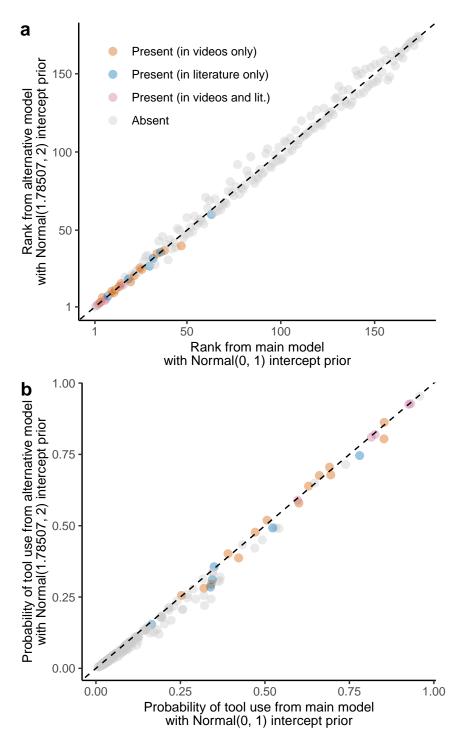
Supplementary Figure 10. Data on number of scientific publications until tool use discovery for all parrots, presented on a maximum clade credibility tree. Tip points are scaled according to the number of published papers up until tool use discovery (or, if tool use has not been observed, the current number of published papers).



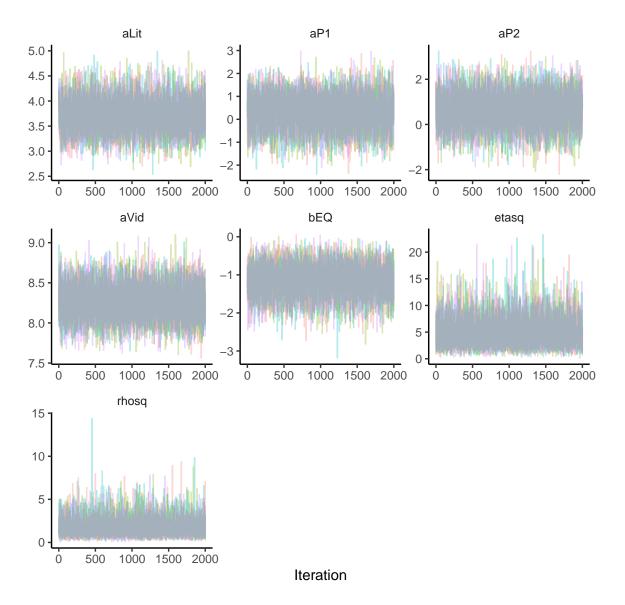
Supplementary Figure 11. Data on number of video search hits until tool use discovery for all parrots, presented on a maximum clade credibility tree. Tip points are scaled according to the estimated number of video search hits up until tool use discovery (or, if tool use has not been observed, the current number of video search hits).



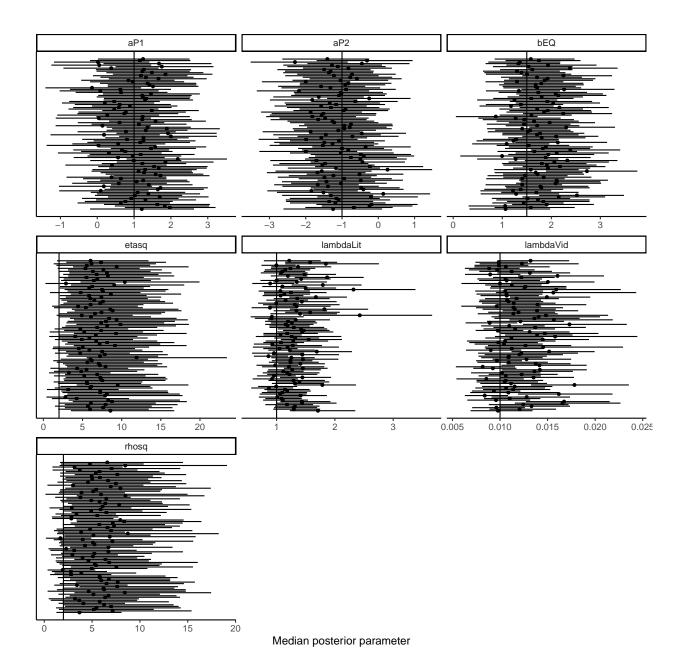
Supplementary Figure 12. Prior predicted probabilities of tool use for each species from our phylogenetic survival cure model. Points are prior medians and lines are 50% and 95% credible intervals.



Supplementary Figure 13. Results of sensitivity analysis. The phylogenetic survival cure model was fitted with either a standard Normal(0, 1) prior on the intercept or an alternative Normal(1.78507, 2) prior on the intercept. This latter prior is wider on the logit scale and roughly converts to a 0.86 prior probability of non-tool-use (or a 0.14 prior probability of tool-use, which is the proportion of tool users in the dataset). The sensitivity analysis showed that changing this intercept prior did not have a marked impact on (a) the posterior rankings of parrot species from 1st to 174th or (b) the median posterior probabilities of tool use for parrot species.



Supplementary Figure 14. Trace plots for the Bayesian phylogenetic survival cure model. Only four chains are shown for ease of presentation.



Supplementary Figure 15. Posterior estimates from Bayesian survival cure models fitted to 100 datasets simulated with known parameters. Each dataset consisted of 100 species. Known parameters are presented as solid vertical lines, whereas points and horizontal lines represent posterior medians and 95% credible intervals. The models were successfully able to recapture the parameters from the simulated datasets.