

A Preferential Attachment Model for the Initial Mass Function

Jessi Cisewski-Kehe *

Department of Statistics and Data Science, Yale University
and

Grant Weller[†]

Savvysherpa, Minneapolis, MN 55430

and

Chad Schafer

Department of Statistics and Data Science, Carnegie Mellon University
and

David W. Hogg

Center for Cosmology and Particle Physics, New York University

January 31, 2018

Abstract

Accurate specification of a likelihood function is becoming increasingly difficult in many inference problems in astronomy. As the sample sizes resulting from astronomical surveys continue to grow, deficiencies in the likelihood function lead to larger biases in key parameter estimates. These deficiencies result from the oversimplification of the physical processes that generated the data, and from the failure to account for observational limitations. Unfortunately, realistic models often do not yield an analytical form for the likelihood. The estimation of a stellar initial mass function (IMF) is an important example. The stellar IMF is the mass distribution of stars initially formed in a given cluster of stars, a population which is not directly observable due to stellar evolution and other disruptions of the cluster. There are several difficulties with specifying a likelihood in this setting, since the physical processes and observational challenges result in measurable masses that cannot legitimately be considered independent draws from an IMF. This work improves inference of the

*The authors gratefully acknowledge *please remember to list all relevant funding sources in the unblinded version*

[†]Corresponding author

IMF by using an approximate Bayesian computation approach that both accounts for observational and astrophysical effects and incorporates a physically-motivated model for star cluster formation. The methodology is illustrated via a simulation study, demonstrating that the proposed approach can recover the true posterior in realistic situations, and applied to observations from the Orion Nebula Cluster.

Keywords: Approximate Bayesian computation, astrostatistics, computational statistics, dependent data

1 Introduction

The Milky Way is home to over 100 billion stars (ADD REF??), many of which are members of *stellar clusters* - gravitationally bound collections of stars. Stellar clusters are formed from low temperature and high density clouds of gas and dust called *molecular clouds*, though there is uncertainty as to how the stars in a cluster form (Beccari et al. 2017). Each theory of star formation yields a different prediction for the distribution of the masses of stars that initially formed in a cluster. Hence, it is of fundamental interest to estimate this distribution, referred to as the *stellar initial mass function (IMF)*, and assess the validity of these competing theories. In fact, research advances in many areas of stellar, galactic, and extragalactic astronomy are at least somewhat reliant upon accurate understanding of the IMF (Bastian et al. 2010). For example, the IMF is a key component of galaxy and stellar evolution and planet formation (Bally & Reipurth 2005, Bastian et al. 2010, Shetty & Cappellari 2014), along with chemical enrichment and abundance of core-collapse supernovae (Weisz et al. 2013).

There is also ongoing discussion surrounding the *universality* of the IMF, i.e., if a single IMF describes the generative distribution of stellar masses for all star clusters (Bastian et al. 2010). Even though the consensus of the astronomical community is that there is a universal IMF, most of the observations had been consistent with universality. With further research and growing sample sizes, however, there is increased theoretical (Dib et al. 2010, Bonnell et al. 2006) and observational (Treu et al. 2010, Spiniello et al. 2014, Dib et al. 2017, Geha et al. 2013) support for an IMF that varies from cluster to cluster.

Salpeter (1955) studied the evolutionary properties of certain populations of stars, and in the process defined the first IMF (which he called the “original mass function”). This work put forth the now-classic model for the IMF, a power law with a finite upper bound equal to the physical maximum mass of a star that could form in a cluster (Salpeter 1955). Recent studies support the validity of this power law form for the IMF for stars of mass greater than half that of our sun (Massey 2003, Bastian et al. 2010, Lim et al. 2013). **JESSI: CHECK PREVIOUS CLAIM.** Similar models have been proposed and used in the astronomical literature for inference of the stellar initial mass function; these will be discussed in the next section. The estimation of the parameters of these proposed models typically relies on the unphysical assumption that the observed stars in a stellar cluster form independently. This work loosens this restriction in order to explore one of several possible physical formation mechanisms of cluster formation (Li et al. 2017).

Despite this seemingly simple form of the power law model, the statistical challenges of estimating the IMF using *observed* stars from a cluster are significant. Many of the limitations are related to observational issues and to the adequate modelling of the evolution of a star cluster after the initial formation. For example, since stars of greater mass die more rapidly, the upper tail of the IMF is not observed in a cluster of sufficient age. Also, the death of massive stars can trigger additional star formation, contaminating the lower end of the IMF with new stars (Woosley & Heger 2015).

The observational limitations and the challenge of modelling cluster evolution make *approximate Bayesian computation (ABC)* appealing for estimation of the IMF, as ABC allows for relatively easy incorporation of such effects. In particular, a primary appeal of ABC for this application is the ability to incorporate more complex models for cluster formation. Standard IMF models do not specify the process by which a large mass of gas (the molecular cloud) transforms into a gravitationally bound collection of stars. ABC is based on a simple rejection-sampling approach, in which draws of model parameters from a prior distribution are fed through a simulation model to generate a sample of data. If the generated sample is “close” (based on an appropriately chosen metric) to the observed data, the prior draw that produced that generated sample is retained. The collection of

accepted parameter values comprise draws from an approximation to the posterior. The simulations (the *forward model*) can include any of the complex processes that make it challenging to derive a likelihood function for the observable data.

Hence, with the goal of improving inference on the IMF by incorporating effects that are often otherwise ignored (Kalari et al. 2018, Ashworth et al. 2017, Jose et al. 2017), **THIS IS A VERY IMPORTANT POINT. THESE REFERENCES SHOULD BE CHECKED, AND MAYBE MORE ADDED.** along with connecting a possible formation mechanism with the IMF, an ABC approach is described in this work. This situation is typical of inference challenges that arise in astronomy. See Schafer & Freeman (2012), Akeret et al. (2015), Ishida et al. (2015) for reviews. Recent years have seen a rapid increase in the use of ABC methods for estimation in this field, including specific application to Milky way properties (Robin et al. 2014), strong lensing of galaxies (Killedar et al. 2018, Birrer et al. 2017), large scale structure of the Universe (Hahn, Vakili, Walsh, Hearin, Hogg & Campbell 2017), estimating the redshift distribution (Herbel et al. 2017), galaxy evolution (Hahn, Tinker & Wetzel 2017), weak lensing (Peel et al. 2017, Lin & Kilbinger 2015), exoplanets (Parker 2015), galaxy morphology (Cameron & Pettitt 2012), and supernovae (Weyant et al. 2013).

This paper is organized as follows. In Section 2 we will present background on the IMF along with inference challenges, and introduce ABC. We propose a new stochastic model for stellar formation in Section 3 and discuss our ABC procedure. Section 5 applies the proposed methodology to the estimation of the IMF of the Orion Nebular Cluster, using a recent sample of observations from this scientifically-important Milky Way cluster (Hillenbrand & Hartmann 1998, Da Rio et al. 2012). Finally, Section 6 provides a discussion.

2 Background

2.1 Stellar Initial Mass Function

As noted above, Salpeter (1955) introduced the power law model for the shape of the IMF for masses larger than $0.5M_{\odot}$, where M_{\odot} is the mass of the Sun. Kroupa (2001) extended the range of the IMF by proposing a three-part broken power law model over the range

$0.01M_{\odot} < m < M_{max}$, where M_{max} is the mass of the largest star that could form with nonzero probability. This model postulates different forms for the IMF for stars of masses $0.01M_{\odot} < m < 0.08M_{\odot}$, $0.08M_{\odot} < m < 0.5$ and $m > 0.5M_{\odot}$. Focusing on the upper part, and defining $\theta = (\alpha, M_{max})$, the probability density function for mass x in the upper tail of the stellar IMF is assumed to be given by

$$f_M(m | \theta) = cm^{-\alpha}, \quad m \in [M_{min}, M_{max}], \quad (1)$$

where the constant c is chosen such that f_M is a valid probability density. Alternative models have been proposed that include log-normal distributions, joint power law and log-normal parts, and truncated exponential distributions (Chabrier 2005, Offner et al. 2014). The Kroupa (2001) and Chabrier (2005) models are displayed in Figure 1 along with observational challenges discussed §2.1.1. Power law distributions and log-normal distributions are closely related and may be the result of subtle differences in the underlying formation mechanism (Mitzenmacher 2004). The IMF model we propose will exploit one possible underlying formation mechanism that we show results in power law tails.

2.1.1 Observational Challenges

Observing all stars comprising an IMF is not feasible, as the most massive stars ($m > 10M_{\odot}$) have lifetimes of only a few million years. The lifetime of a star (the time it takes for the star to burn through its hydrogen) depends strongly on its mass: the most massive stars have shorter lives due to the hotter temperatures they must maintain to avoid collapse from the strong gravitational forces. In particular, stellar life is approximately proportional to $m^{-\rho}$ where $\rho \approx 2.5$ (Hansen et al. 2004, p. 28). Chad: I was looking in the book and it seems that $\rho = 2.9$ (which it seems fine to use 2.5 since it is easier to work with later) and it appears on page 30 of the textbook. Can you confirm this? Hence, the mass of the largest star observed in a given cluster is depends on the cluster age.

Furthermore, the IMF will be estimated using a noisy, incomplete view of that cluster. Whether or not a star is observed is dependent on several factors including its mass, its location in the cluster, and its neighbors. Some of these factors are described by a data set's *completeness function*, which quantifies a given star's probability of being observed.

This depends on its luminosity (i.e. intrinsic brightness) since it needs to be sufficiently bright to be observable; in particular, completeness depends on stellar flux in comparison with the flux limits of the observations. There are also issues with *mass segregation*: stars with lower mass tend to be on the edge of the cluster, while the most massive stars are often found in the center (Weisz et al. 2013). Due to *stellar crowding* in the center, stars in this region can be more difficult to observe. Additionally, binary stars (star systems consisting of a pair of stars) are difficult to distinguish from a single star, creating the potential for overstating the mass of an object and understating the number of stars in the cluster.

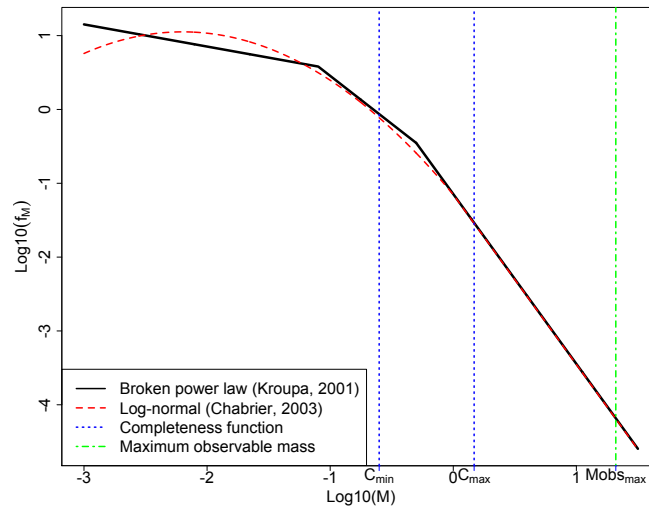


Figure 1: The broken power law model of Kroupa (2001) (black, line) and the lognormal model of Chabrier (2005) (red, dashed) are displayed along with vertical lines representing several observational challenges. The blue vertical dotted lines indicate the range of values (C_{\min}, C_{\max}) on which the completeness function may be defined, and the vertical green dotted and dashed line indicates the maximum observable mass ($M_{\text{obs}_{\max}}$) due to the aging cutoff. The observational challenges are discussed further in §3.2.

ADD ISSUES WITH GETTING MASSES There are additional uncertainties involved in translating the actual observables (e.g. flux) into a mass measurement; that is, observed mass values for stars which are observed are only estimates. **getting mass estimates - using HR diagram, etc. and decisions that are made.**

2.2 Approximate Bayesian Computation

Standard approaches to Bayesian inference, either analytical or built on MCMC, require the specification of a likelihood function, $f(m \mid \boldsymbol{\theta})$, with data $m_{\text{obs}} \in \mathcal{D}$, and parameter(s) $\boldsymbol{\theta} \in \Theta$. In many modern scientific inference problems, such as for some emerging models for the stellar IMF, the likelihood is too complicated to be derived or otherwise specified. As noted previously, ABC provides an approximation to the posterior without specifying a likelihood function, and instead relies on forward simulation of the data generating process.

The basic algorithm for sampling from the ABC posterior is attributed to Tavaré et al. (1997) and Pritchard et al. (1999), used for applications to population genetics. The algorithm has three main steps which are repeated until a sufficiently large sample is generated: *Step 1*, Sample $\boldsymbol{\theta}^*$ from the prior; *Step 2*, Generate m_{sim} from forward process assuming truth $\boldsymbol{\theta}^*$; *Step 3*, Accept $\boldsymbol{\theta}^*$ if some distance function $\rho(m_{\text{obs}}, m_{\text{sim}}) \leq \epsilon$, where ϵ is a tuning parameter that should be close to 0. This last step typically consists of comparing low-dimensional summary statistics generated for the observed and simulated datasets. Adequate statistical and computational performance of ABC algorithms depends greatly on the selection of such summary statistics (Joyce & Marjoram 2008, Blum & François 2010, Blum 2010, Fearnhead & Prangle 2012, Blum et al. 2013).

The basic ABC algorithm can be inefficient in cases where the parameter space is of moderate or high dimension. Hence, important adaptations of the basic ABC algorithm incorporate ideas of sequential Monte Carlo (SMC) in order to improve the sampling efficiency (Marjoram et al. 2003, Sisson et al. 2007, Beaumont et al. 2009, Del Moral et al. 2011). A nice overview of ABC can be found in Marin et al. (2012). Here, we use a sequence of decreasing tolerances $\epsilon_{1:T} = (\epsilon_1, \dots, \epsilon_T)$ with the tolerance ϵ_t shrinking until further reductions do not significantly affect the resulting ABC posterior. The improvement in efficiency is due to the modification that happens after the first time step: instead of sampling from the prior distribution, the proposed $\boldsymbol{\theta}$ are drawn from the previous time step's ABC posterior. Using this adaptive proposal distribution can help to improve the sampling efficiency. The resulting draws, however, are not targeting the correct posterior, and so importance weights, W_t , are used to correct this discrepancy.

3 Forward Model for the IMF

Due to their simple interpretations, mathematical ease, and demonstrated consistency with observations, power law IMFs (or similar variants) have been widely adopted in the astronomy literature (Kroupa et al. 2012); however, open questions remain about stellar formation processes. The proposed forward model is a way to link a possible stellar formation process with the realized mass function (MF). One known underlying mechanism for producing data with power-law tails is based on *preferential attachment* (PA). The earliest PA model, the Yule-Simon process, was popularized by Simon (1955), and was originally used to model biological genera and word frequencies. **ADD OTHER USES.** In the stellar formation context, linear PA models generate power law IMFs; sublinear variants produce slight deviations from power laws, and exponential cutoffs can also be induced.

add discussion about PA model, and nonlinear attachment. Proof of power law tails. Use in networks. We need to generalize.

3.1 Preferential attachment for the IMF

describe formation theory we are going to try to capture. Rather than assuming that stellar masses in a cluster arise independently of each other, our PA model proposes a resource-limited mass accretion process between stellar cores whose ability to accumulate additional mass is a function of their existing masses. This dependence feature is particularly important for statistical inference, as models that assume independent observations of stellar masses are vulnerable to incorrect and misleading inference. Additionally, the mass of the largest star to form in a cluster is limited by the total cluster mass.

Our proposed stochastic model for stellar formation is as follows: we first fix a total available cluster mass M_{tot} . This quantity can be physically interpreted as the total mass available for stellar formation in a molecular cloud. At each time step $t = 1, 2, \dots$, a random quantity of mass $m_t \sim \text{Exponential}(\lambda)$ enters the collection of stars; $M_{1,1} = m_1$ becomes the mass of the first star. Subsequent masses entering the system form a new star with probability π_t or join existing star $k = 1, \dots, n_t$ with probability π_{kt} . These probabilities

are specified as

$$\pi_t = \min(1, \alpha) \quad \text{and} \quad \pi_{kt} \propto M_{k,t}^\gamma. \quad (2)$$

The generating process is complete when the total mass of formed stars reaches M_{tot} . The possible ranges of the five parameters are $\lambda > 0$, $\alpha \in [0, 1)$, and $\gamma > 0$.

Proof of power law tails??? Put in appendix.

The parameter α defines the probability that entering mass forms a new star in a cluster. For the growth component, the model allows for linear ($\gamma = 1$), sublinear ($\gamma < 1$), and superlinear ($\gamma > 1$) behavior; the limiting case of $\gamma \rightarrow 0$ gives a uniform attachment model. Finally, the parameter λ acts as a scaling factor which controls the average ‘coarseness’ of masses joining the forming stellar cores.

The proposed PA mass generation model offers considerable flexibility to approximate existing IMF models in the literature. In Figure 2, IMF realizations were drawn assuming the Kroupa (2001) broken power-law model as the truth model and the Chabrier (2005) lognormal model as the true model. Our proposed PA ABC procedure was then used for inference and captures the general shape of the true model.

3.2 Initial Mass Function to the Observed Mass Function

The PA model describes the formation of a star cluster at initial formation. However, we are not generally able to observe the star cluster after initial formation due to observational uncertainties, measurement uncertainties, and aging and dynamical evolution of the cluster. A cluster’s present-day observed mass function (MF) is the observed distribution of the stellar masses of a particular cluster.

Observation limitations can be easily incorporated into the ABC framework. For simplicity, we adopt a “linear ramp” completeness function describing the probability of ob-

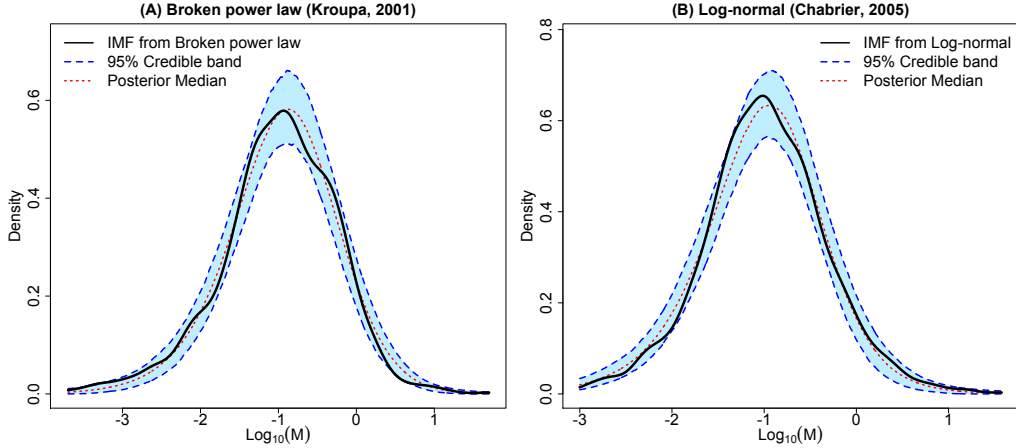


Figure 2: The solid black lines display the IMF with 1,000 stars drawn from (A) the broken power-law model of Kroupa (2001) and (B) the log-normal model of Chabrier (2005). The proposed PA ABC model was used with $N = 1,000$ particles, and 95% point-wise credible bands are displayed (blue, dashed lines) along with the posterior median (red, dotted) for each data set. The preferential attachment model provides flexibility to approximate both existing models.

serving a star of mass m :

$$\Pr(\text{observing a star} \mid m) = \begin{cases} 0, & m < C_{\min} \\ \frac{m - C_{\min}}{C_{\max} - C_{\min}}, & m \in [C_{\min}, C_{\max}] \\ 1, & m > C_{\max}. \end{cases} \quad (3)$$

We assume that the values C_{\min} and C_{\max} are known, though we note that selecting an appropriate completeness function is a difficult process which requires quantification from the observational astronomers for each set of data. Different models for the completeness function could also be considered, including those which allow for spatially-varying observational completeness. A benefit of ABC is the ease at which a new completeness function can be incorporated – it amounts to a simple change in the forward model.

Due to measurement error and practical limitations in translating luminosities into masses, the masses of stars are not perfectly known. This uncertainty can be incorporated in different ways; following Weisz et al. (2013), we assume that the inferred mass of a star

m_i is related to its true mass M_i via

$$\log m_i = \log M_i + \sigma_i \eta_i, \quad (4)$$

where η_i is a standard normal random variable, and σ_i is known measurement error. The model for mass uncertainties in (4) is simple and could be extended to account for other sources of uncertainty (e.g. redshift).

As noted previously, the lifecycle of a star depends on certain characteristics such as mass. In the proposed algorithm, stars generated in a cluster are aged using a simple truncation of the largest masses. That is, the distribution of stellar masses for a star cluster of age τ Myr is given by

$$f_M(m \mid \theta, \tau) = f_M(m \mid \theta) \mathbb{I}\{M \leq \tau^{-2/5} \times 10^{8/5}\}, \quad (5)$$

corresponding to stellar lifetimes of $10^4 M^{-2.5}$ Myr (Hansen et al. 2004). More sophisticated models that account for effects such as binary stars and stellar wind mass loss can be inserted into this framework.

4 Methods

We propose an ABC framework to make inferences on the IMF given a cluster’s present-day observed MF. The proposed ABC algorithm is displayed in Algorithm (1), where N is the desired particle sample size to approximate the posterior distribution, and is motivated by the adaptive ABC algorithm of Beaumont et al. (2009). The forward model, F , in Algorithm (1) is where observational limitations and uncertainties, stellar evolution, and other astrophysical elements can be incorporated as outlined in Section 3.

Algorithm (1) is initialized using the basic ABC rejection algorithm at time step $t = 1$ using a distance function $\rho(m_{\text{sim}}, m_{\text{obs}})$ to measure the distance between the simulated and observed datasets, m_{sim} and m_{obs} , respectively. The first tolerance, ϵ_1 , is adaptively selected by drawing kN particles for some $k > 0$. Then the N particles that have the smallest distance are retained, and ϵ_1 is defined as the largest of those N distances retained. For subsequent time steps ($t > 1$), rather than proposing a draw, θ^* , from the prior, $\pi(\theta)$,

the proposed θ^* is selected from the previous time step's $(t - 1)$ ABC posterior samples. The selected θ^* is then moved according to some kernel, $K(\theta^*, \cdot)$, to ameliorate degeneracy as the sampler evolves. In order to ensure the true posterior (which requires sampling from the prior) is targeted, the retained draws are weighted according to the appropriate importance weights, W_t – this incorporates the proposal distribution's kernel.

Data: Observed stellar masses, (m_{obs})

Result: ABC posterior sample of θ

At iteration $t = 1$:

for $j = 1, \dots, kN$ **do**

 Propose $\theta_t^{(j)}$ by drawing $\theta_t^* \sim \pi(\theta)$

 Generate cluster stellar masses m_{sim} from $F(m \mid \theta_t^*)$

 Calculate distance $\rho_t^{(j)} \leftarrow \rho(m_{\text{sim}}, m_{\text{obs}})$

end

$\theta_t^{(j)} \leftarrow \theta_t^{(l)}$, $l = \text{index of } N \text{ smallest } \rho_t^{(q)}, q = 1, \dots, kN$

$W_t^{(j)} \leftarrow 1/N, j = 1, \dots, N$

At iterations $t = 2, \dots, T$:

for $j = 1, \dots, N$ **do**

while $\rho(m_{\text{sim}}, m_{\text{obs}}) > \epsilon_t$ **do**

 Select $\theta^{(j)}$ by drawing from the $\theta_{t-1}^{(i)}$ with probabilities $W_{t-1}^{(i)}, i = 1, \dots, N$

 Generate $\theta^{*(j)}$ from transition kernel $K(\theta^{(j)}, \cdot)$

 Generate cluster stellar masses m_{sim} from $F(m \mid \theta^{*(j)})$

end

$\theta_t^{(j)} \leftarrow \theta^{*(j)}$

$W_t^{(j)} \leftarrow \frac{\pi(\theta_t^{(j)})}{\sum_{i=1}^N W_{t-1}^{(i)} K(\theta_{t-1}^{(i)}, \theta_t^{(j)})}$

end

$W_t^{(j)} \leftarrow \frac{W_t^{(j)}}{\sum_{l=1}^N W_t^{(l)}}, j = 1, \dots, N$

Algorithm 1: Stellar IMF ABC algorithm with sequential sampling

A key step in the implementation of an ABC algorithm is to quantify the distance

between the simulated and observed stellar masses. We define a bivariate summary statistic and distance function that captures the shape of the present-day MF and the random number of stars observed, displayed in Equations (6) and (7), respectively. For the shape of the present-day MF, we use a kernel density estimate of the \log_{10} masses (due to the heavy-tailed distribution of the initial masses), and an L_2 distance between the simulated and observed \log_{10} mass function estimates. The number of stars observed is the other summary statistic, with the distance being the absolute value of the difference in the ratio of the counts from 1. The bivariate summary statistic is defined as

$$\rho_1(m_{\text{sim}}, m_{\text{obs}}) = \left[\int \left\{ \hat{f}_{\log m_{\text{sim}}}(x) - \hat{f}_{\log m_{\text{obs}}}(x) \right\}^2 dx \right]^{1/2} \quad (6)$$

$$\rho_2(m_{\text{sim}}, m_{\text{obs}}) = \max \{ |1 - n_{\text{sim}}/n_{\text{obs}}|, |1 - n_{\text{obs}}/n_{\text{sim}}| \}, \quad (7)$$

where the \hat{f} are kernel density estimates, and n_{sim} and n_{obs} are the number of stars comprising the observed MF. These summary statistics were selected based on performance of a simulation study using the high-mass section of the broken power-law model because the true posterior is known in this setting. Results and additional discussion of the simulation study can be found in Appendix A.

With the bivariate summary statistic, we use a bivariate tolerance sequence, $(\epsilon_{1t}, \epsilon_{2t})$, for $t = 1, \dots, T$ is such that $\epsilon_{i1} \geq \epsilon_{i2} \geq \dots \geq \epsilon_{iT}$ for $i = 1, 2$. At time step t , the tolerances are determined based on the empirical distribution of the retained distances from time step $t - 1$ (e.g. the 25th percentile). As noted previously, the tolerance sequence is initialized adaptively by selecting kN proposals from the prior distributions, then the N proposals that result in the N smallest distances were selected.¹

In practice, M_{tot} is an unknown quantity of interest. A prior can be assigned to M_{tot} and an additional summary statistic and tolerance sequence can be used. The summary statistics we select in this case is

$$\rho_3(m_{\text{sim}}, m_{\text{obs}}) = \left| \sum_{i=1}^{n_{\text{sim}}} m_{\text{sim}}^i - \sum_{j=1}^{n_{\text{obs}}} m_{\text{obs}}^j \right|, \quad (8)$$

¹The kN sampled distances were scaled, squared, and then added together; the N smallest of these combined distances were retained.

where m_{sim}^i and m_{obs}^j are the masses of the individual simulated and observed stars, respectively.

A simulation study was carried out to check the performance of the proposed model in the case where the solution is known using the summary statistics in Equations (6)-(8).

Number	λ^{-1}	α	γ	M_{tot}
1	0.25	0.3	0.5	1000
2	0.25	0.3	1.0	1000
3	0.25	0.3	1.5	1000
4	0.25	0.7	0.5	1000
5	0.25	0.7	1.0	1000
6	0.25	0.7	1.5	1000

Table 1: Parameters for the simulation study of Section 4.

Figure 3: ADD PLOT OF RESULTING POSTERiors

Figure 4: ADD PLOT OF SAMPLED IMF

4.1 Astrophysical simulation data

Bate (2012)

In Bate - should not have significant super-linear attachment. Check to see if he has examples where there is super-linear attachment.

5 Application

6 Discussion

Accounting for complex dependencies in observations, such as the initial masses of stars forming from a molecular cloud, is a challenging statistical problem. A possible, but unsatisfactory, resolution is simply to proceed as though the dependency is weak enough to assume independence. Instead, we draw from ideas in the preferential attachment literature proposing a new forward model to be used in an ABC algorithm. Though the new generative model was motivated by inference on stellar IMFs, it is generalizable to other applications that operate on a similar principle of dependent data. First we verified that the ABC approach works under the simplifying assumption of independence to ensure the true posterior was recoverable using ABC with the various observational effects. Then we introduced the new generative model, which starts with the total mass and stochastically builds individual stars of particular mass. The new model is general in the sense that it captures the dependencies among the masses of the stars, and also can accommodate standard models proposed in the astronomical literature.

The ONC is an interesting object of study among astronomers. The proposed preferential attachment model fitted via ABC captures the fluctuations in the observed mass function, as well as accounting for varying uncertainty as a function of available data in different regions of the mass range. While the proposed model is able to account for the dependency among the masses (along with the aging, completeness function, and observation uncertainty effects), it would be scientifically and statistically interesting to generalize the generative model to capture the spatial dependency among the observations. Intuitively, such an approach could account for a mechanism that limits the formation of multiple very massive stars relatively near to each other. Understanding the spatial distribution of masses of stars during formation would help advance our understanding stellar formation and evolution. Additionally, we did not attempt to capture the possible disturbances to the observed mass function as stars die, with the exception of deaths of the most massive stars.

Overall, the proposed generative model and ABC methodology provide a useful framework for dealing with complex physical processes that are otherwise difficult to work with in a statistically rigorous fashion. As increasing computational resources allow for greater model complexity in studies of astronomy and other scientific fields, the proposed and other ABC algorithms may open new opportunities for Bayesian inference in challenging problems.

A ABC summary statistic selection

In order to select effective summary statistics for the proposed model, we first employ the ABC methodology in a simplified study that focuses on the posterior of the power law parameter α from Equation (1). We generate a cluster of $n = 10^3$ stars from an IMF with slope $\alpha = 2.35$ (Salpeter 1955), $M_{min} = 2$, and $M_{max} = 60$, a uniform prior distribution for $\alpha \in (0, 6)$, and with M_{max} considered known. This simple model was used in order to check the method against the true posterior of α after the observational and aging effects have been incorporated into the forward model. We use the bivariate summary statistic and distance function of Equation (6). Defining the two-dimensional tolerance sequence as $(\epsilon_{1t}, \epsilon_{2t})$ where the subscript t indicates the algorithm time step, and ϵ_{11}^2 and ϵ_{21} were selected using an adaptive started discussed in Section 4 using an initial number of draws of $10 \times N$ with $N = 10^3$. The algorithm ran for $T = 5$ time steps. At steps $t = 2, \dots, T$, ϵ_{1t} and ϵ_{2t} were set equal to the 25th percentile of the distances retained at the previous step from their corresponding distance functions.

The pseudo-data were aged 30 Myrs with the measurement error of $\sigma = 0.25$ and observation completeness defined by the linear-ramp function in (3). Figure 5 displays the ABC posterior resulting from the ABC algorithm along with the true posterior for α . The

ABC posterior matches the true posterior, defined as

$$\begin{aligned}
\pi_F(\alpha \mid m_{\text{obs}}, M_{\text{min}}, M_{\text{max}}, n_{\text{obs}}, n_{\text{tot}}, T_{\text{age}}) \propto & \quad (9) \\
& \left\{ \Pr(M > T_{\text{age}}) + \left(\frac{1 - \alpha}{M_{\text{max}}^{1-\alpha} - M_{\text{min}}^{1-\alpha}} \right) \int_2^4 M^{-\alpha} \left(1 - \frac{M-2}{2} \right) dM \right\}^{n_{\text{tot}} - n_{\text{obs}}} \\
& \times \prod_{i=1}^{n_{\text{obs}}} \left\{ \int_2^{T_{\text{age}}} (2\pi\sigma^2)^{-\frac{1}{2}} m_i^{-1} e^{-\frac{1}{2\sigma^2}(\log(m_i) - \log(M))^2} \left(\frac{1 - \alpha}{M_{\text{max}}^{1-\alpha} - M_{\text{min}}^{1-\alpha}} \right) M^{-\alpha} \right. \\
& \left. \times \left(I\{M > 4\} + \left(\frac{M-2}{2} \right) I\{2 \leq M \leq 4\} \right) dM \right\}
\end{aligned}$$

where $T_{\text{age}} = \tau^{-2/5} \times 10^{8/5}$ is the upper-tail mass cutoff due to aging.

Acknowledgements

The authors thank the reviewers and associate editor for helpful comments that lead to significant improvements of this work. The authors also benefited from feedback from Ewan Cameron. Jessi Cisewski and Grant Weller were partially supported by the National Science Foundation under Grant DMS-1043903. Chad Schafer was supported by NSF Grant DMS-1106956. David W. Hogg was partially supported by the NSF (AST-0908357) and the Moore–Sloan Data Science Environment at NYU. Any opinions, findings, and conclusions or recommendations expressed in this material are those of the authors and do not necessarily reflect the views of the National Science Foundation.

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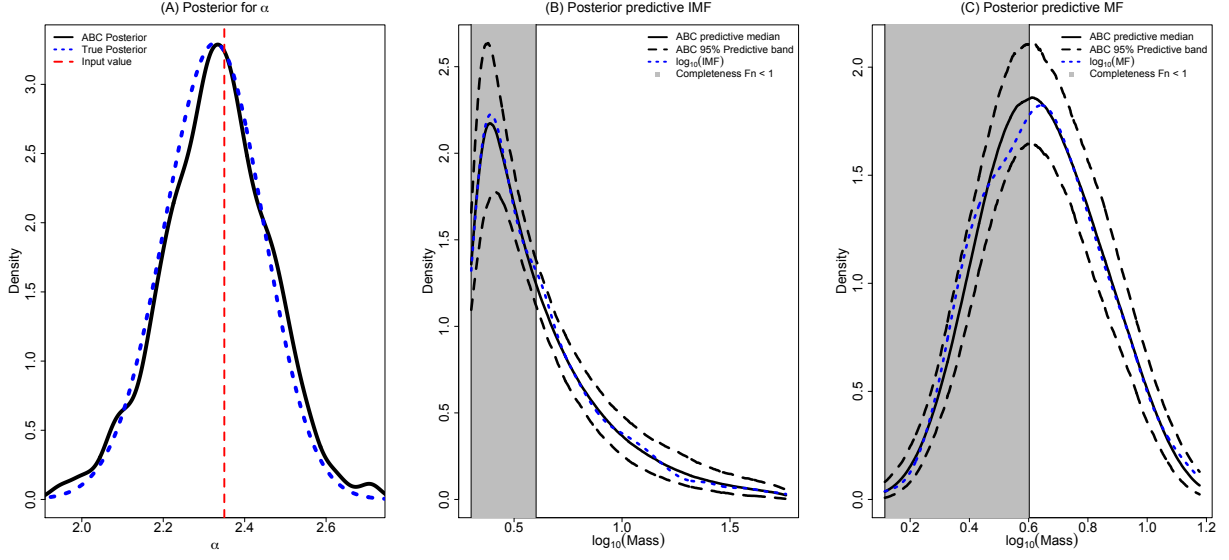


Figure 5: Validation of summary statistics with simple model. (A) The ABC posterior for α (solid black) compared to the true posterior (dotted blue) of Equation (9) using an input value of 2.35 (dashed vertical red). (B) The median of the posterior predictive IMF (solid black) with a corresponding 95% point-wise predictive band (dashed black) compared to the true IMF (blue dotted) which was simulated dataset before aging, completeness, or uncertainty were applied, and the gray shaded region indicates where the completeness function was less than 1. (C) The median of the posterior predictive MF (solid black) with a corresponding 95% point-wise predictive band (dashed black) compared to the observed MF (dotted blue) which was the simulated dataset after aging, completeness, and uncertainty were applied. For the posterior predictive IMF, 1000 draws were made from the ABC posterior of (A) and then 1000 cluster samples were drawn from the power law simulation model. For the posterior predictive MF, the 1000 cluster samples used for (B) were then put through the forward model to apply the aging, completeness, and measurement error effects.

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