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Inferring binary black holes stellar progenitors with gravitational wave sources

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ABSTRACT

With its last observing run, the LIGO, Virgo, and KAGRA collaboration has detected almost one hundred gravitational waves from compact binary coalescences. A common approach to studying the population properties of the observed binaries is to use phenomenological models to describe the spin, mass, and redshift distributions. More recently, with the aim of providing a clearer link to astrophysical processes forming the observed compact binaries coalescences, several authors have proposed to employ synthetic catalogues for population studies. In this paper, we review how to employ and interpret synthetic binary catalogues for gravitational-wave progenitors studies. We describe how to build multichannel merger rates and describe their associated probabilities focusing on stellar progenitor properties. We introduce a method to quantify the match between the phenomenological reconstruction of merger rates with synthetic catalogues. We detail the implementation of synthetic catalogues for multichannel hierarchical Bayesian inference, highlighting computational aspects and issues related to hyper-prior choice. We find that when inferring stellar progenitors' properties from gravitational-wave observations, the relative efficiency in compact objects production should be taken into account. Finally, by simulating binary black hole detections with LIGO and Virgo sensitivity expected for the O4 observing run, we present two case studies related to the inference of the common envelope efficiency and progenitor metallicity of the binary black holes. We finally discuss how progenitors' properties can be linked to binary black hole properties.

Key words: gravitation – gravitational waves.

1 INTRODUCTION

Since their first detection in 2015 (Abbott et al. 2016), gravitational waves (GWs) have opened a new channel to study our Universe. Besides representing another confirmation of Einstein's General Relativity, GWs also provide us with a new tool for studying stellar evolution, cosmology, and the origin of compact objects. In just 6 yr from their first direct detection, and during just three observing runs, there has been meteoric progress in GW astrophysics. In 2017, the first Binary Neutron Star (BNS) detection with electromagnetic counterpart allowed us to measure the Hubble constant H_0 (Abbott et al. 2017a,b), constrain the speed of gravity, confidently link kilonovae and short γ -ray bursts and observe the formation of heavy elements via r-process (Abbott et al. 2017c). The LIGO and Virgo interferometers observed GW190521, a Binary Black Hole (BBH) merger (Abbott et al. 2020a, c) with masses falling in the Pair Instability Supernova (PISN) gap. Another interesting example is GW190814 (Abbott et al. 2020b), a compact binary merger that includes a BH of $\sim 20 M_\odot$ and a secondary object falling in the expected mass gap between neutron stars and the black holes.

Interesting scientific results have also been achieved by studying the population of Compact Binaries Coalescences (CBCs). Using the GW events from the last Gravitational-Wave Transient catalogues (GWTC) (Abbott et al. 2021d, b), the LIGO/Virgo/KAGRA collab-

oration (LVK) has been able to show that there is a smooth transition between neutron stars and black holes masses, that the preliminary BBH merger rate evolves in redshift and that the BBH mass spectrum presents several features (Fishbach & Holz 2017; Abbott et al. 2021f, c; Tiwari & Fairhurst 2021). The LVK has been able to constrain H_0 using BBHs provided with galaxy catalogues (Abbott et al. 2021e) and astrophysical source mass distributions (Abbott et al. 2021a). All of this has been achieved with a catalogue of 90 GW candidates.

As the number of GW detections rapidly increases, population studies with GW sources are becoming a suitable tool to study the astrophysical formation channels of compact objects. Studying the population of CBCs practically consists in reconstructing the astrophysical merger rate from the observed merger rate (Mandel, Farr & Gair 2019; Vitale et al. 2022) or vice-versa. The astrophysical merger rate is linked to astrophysical processes driving the production of the CBC population. For instance, for BBHs, the presence of a PISN process (Farmer et al. 2019; van Son et al. 2020) prevents the formation of black holes (BHs) in the range $50 M_\odot - 120 M_\odot$. See Mapelli (2021) for an extensive review of the different formation channels for compact binaries. Population studies are also important to understand the nature of any particular 'exceptional' event. In Mandel (2010), Galadage, Talbot & Thrane (2020), Fishbach, Farr & Holz (2020), Moore & Gerosa (2021) the authors present a methodology to recompute the estimation of GW parameters in light of population analyses, while works such as Fishbach & Holz (2020), Farah et al. (2022) try to reconcile GW190814 and GW190521 with the observed population of BBHs.

Currently, two methodologies are employed to reconstruct the astrophysical merger rates. The first one, which was also adopted

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by the LVK (Abbott et al. 2020c, 2021c) reconstructs merger rates in masses, spins, and redshift using inferential statistics and flexible phenomenological models. This approach is widely used in current literature (Fishbach & Holz 2017; Vitale et al. 2019; Callister et al. 2021; Farah et al. 2022) and reconstructs the binary merger rate based on astrophysical assumptions for the phenomenological models. For instance, the BBHs merger rate as a function of redshift is usually approximated at low redshift with $(1+z)^\gamma$, in analogy to the evolution of the star formation rate at low redshift. On one hand, this approach has the advantage of being flexible enough to fit an unknown population. On the other hand, the disadvantage of this approach is that it is not directly connected to the astrophysical processes producing BHs from their progenitors.

In order to provide a more direct astrophysical interpretation of the observed population, a parallel methodology has been employed (e.g. Stevenson, Berry & Mandel 2017; Zevin et al. 2017; Wysocki et al. 2018; Bouffanais et al. 2019; Bouffanais et al. 2021; Delfavero et al. 2021; Mapelli et al. 2022; Ng et al. 2021; Zevin et al. 2021; Delfavero et al., in preparation). This methodology consists in directly reconstructing the merger rate from astrophysical synthesis simulations of binary mergers. The central paradigm of this type of approach is to construct multichannel distributions, where the overall population is the sum of all the astrophysical channels simulated. As an example, one can simulate BBHs formed in isolated stellar binaries and in globular clusters and then define an overall population from them. This type of approach has the advantage of being directly connected to the astrophysical processes forming the binary mergers but has the disadvantage of being less flexible in fitting the observed population.

In this methodological paper, we focus on several aspects related to the interpretation and exploitation of synthetic populations of binaries for studying progenitors of GW sources. The paper is organized as follows. In Section 2 we provide an easy statistical method to quantify the match between phenomenological reconstructed merger rates and binary mergers catalogues. In Section 3 we introduce key concepts for reconstructing and interpreting progenitors of mergers with multichannel analysis. In Section 4 we review and discuss critical issues of using several synthetic catalogues, that either change the astrophysical prescriptions or initial conditions, to fit observed GW events. We refer to this type of analysis as ‘multichannel reconstruction’. Differently from previous literature, we will focus on the reconstruction of stellar progenitors properties from GWs observations, showing how the relative efficiency in producing compact objects can be taken into account. In Section 5, using synthetic BBHs populations, we present two case studies in which the methodologies discussed could be employed: the estimation of the common envelope efficiency and the estimation of the progenitor’s metallicity. We also show how stellar progenitors’ multichannel inference can be related to the multichannel inference of BBHs population present in literature. Finally, in Section 6 we provide our final remarks.

All results presented in this paper are generated with GWPARENTS,¹ a code for the multichannel inference released with this work.

2 MATCHING SYNTHETIC MERGER RATES WITH PHENOMENOLOGICAL RECONSTRUCTIONS

We first discuss in this section a quick method to quantify the agreement between synthetic binary catalogues and binary merger rates reconstructed with phenomenological models. In practice,

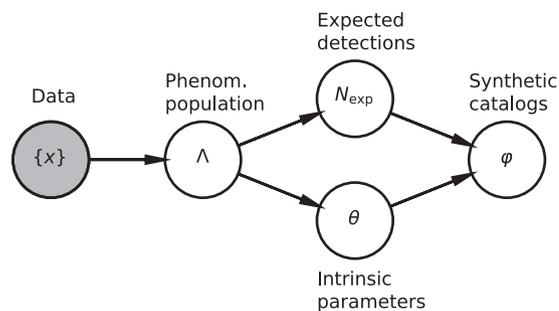


Figure 1. Bayesian graph for comparing synthetic binaries with phenomenological reconstructions of populations from real data. Each node represent a random variable, the shaded node indicates observed data. Each connection between a node and its parents indicates a conditional probability of that node given its parents.

this is the case in which we want to compare a synthetic binary catalogue with a previous analysis reconstructed merger rate using phenomenological models (see Fishbach & Holz 2017; Vitale et al. 2019; Wysocki, Lange & O’Shaughnessy 2019; Abbott et al. 2020c, 2021c; Callister et al. 2021; Farah et al. 2022, as an example) from real GW events. One qualitative avenue that was followed to perform this comparison is to ‘check by eye’ the overlap of the merger rates in terms of masses and redshift of the phenomenological reconstructed rates and the synthetic catalogues.

On one hand, this method offers a quick tool to evaluate the suitability of synthetic binaries from the phenomenological reconstruction. On the other hand, this method does not offer any statistical (or quality factor) indicator and it is hard to visualize in the case that the binary parameters are more than two.

In this section, we introduce for the first time, a more quantitative method to assign a ‘match’ value to each synthetic binary model given the phenomenological reconstruction of astrophysical rates. Let us assume that we have detected $\{x\}$ GW events from which a previous analysis estimated a posterior $p(\Lambda|\{x\})$ on some population-level parameters Λ that describe the phenomenological rate. For instance, a population-level parameter could be the maximum mass of the BBHs mass spectrum or parameters related to the BBHs merger rate as a function of redshift. The population-level parameters, and the phenomenological models, can be used to construct a population distribution $p_{\text{pop}}(\theta|\Lambda)$, where θ represents GW source parameters such as the two masses, and a number of expected detections N_{exp} . In order to assess the suitability of a synthetic population φ_j , we should compare the expected number of detections predicted by φ_j with the one predicted from the phenomenological model. From this comparison, we would like to assign a probability to each model φ_j to fit the observed data, namely $p(\varphi_j|\{x\})$.

The statistical model to compute $p(\varphi_j|\{x\})$ is depicted in Fig. 1. The graph provides a quick tool for evaluating

$$\begin{aligned}
 p(\varphi_j|\{x\}) &= \\
 & \int p(\Lambda|\{x\}) p_{\text{pop}}(\theta|\Lambda) p(N_{\text{exp}}|\Lambda) p(\varphi_j|\theta, N_{\text{exp}}) d\Lambda dN_{\text{exp}} d\theta \\
 &= \int p(\Lambda|\{x\}) p_{\text{pop}}(\theta|\Lambda) p(\varphi_j|\theta, N_{\text{exp}}(\Lambda)) d\Lambda d\theta. \quad (1)
 \end{aligned}$$

In the above equation, we have performed the integral on N_{exp} by using the relation $p(N_{\text{exp}}|\Lambda) = \delta(N_{\text{exp}}(\Lambda) - N_{\text{exp}})$, that is basically representing the fact that for each phenomenological model we can predict an expected number of GW detections. In equation (1), $p(\Lambda|\{x\})$ is the posterior distribution on the phenomenological

¹<https://github.com/simone-mastrogiovanni/gwparents>

population-level parameters inferred from data and $p_{\text{pop}}(\theta|\Lambda)$ the binary parameters distributions that can be reconstructed from them. The term $p(\varphi_j|\theta, N_{\text{exp}}(\Lambda))$ is a probability representing our degree of belief on the astrophysical model φ_j , given a set of binary parameters θ and expected detections N_{exp} . This term can be rewritten using the Bayes Theorem as

$$p(\varphi_j|\theta, N_{\text{exp}}(\Lambda)) = \frac{p(N_{\text{exp}}(\Lambda)|\varphi_j)p_{\text{pop}}(\theta|\varphi_j)p(\varphi_j)}{\sum_k p(N_{\text{exp}}(\Lambda)|\varphi_k)p_{\text{pop}}(\theta|\varphi_k)p(\varphi_k)}, \quad (2)$$

where $p(\varphi_j)$ is a prior belief for the j th formation channel, $p_{\text{pop}}(\theta|\varphi_j)$ is the population prior defined in equation (5) and $p(N_{\text{exp}}(\Lambda)|\varphi_j)$ matching the number of expected detections from the phenomenological model with the number of expected detections from the astrophysical model. When calculating equation (2), one should include also the ‘complementary’ channel $\bar{\varphi}$ that covers the parameter space θ not covered by any of the other channels, i.e. $p(\bar{\varphi}|\theta, N_{\text{exp}}(\Lambda)) = 1 - \sum_j p(\varphi_j|\theta, N_{\text{exp}}(\Lambda))$. Note that equation (2) reduces to the ratio of the population priors in the limit that all the models predict the same number of expected detections. Note also that in this analysis we do not need to include selection biases as they have already been deconvolved by the analysis that fit the phenomenological model. In other words, we are comparing astrophysical rates and not observed rates.

Equation (1) can be computed using the following procedure: if we are provided with a set of N_Λ posterior samples for the population phenomenological parameters Λ_i , for each Λ_i one can compute the expected number of events $N_{\text{exp}}(\Lambda_i)$, then draw N_θ binaries from the population distribution $p_{\text{pop}}(\theta|\Lambda_i)$ and evaluate the integral in equation (1) as

$$p(\varphi_j|\{x\}) = \frac{1}{N_\Lambda N_\theta} \sum_i^{N_\Lambda} \sum_k^{N_\theta} p(\varphi_j|\theta_k, N_{\text{exp}}(\Lambda_i)). \quad (3)$$

Let us give an example. We simulate two populations of BBHs that we refer to ‘isolated’ (φ_{iso}) and ‘globular clusters’ (φ_{gcl}) in analogy with the current BBHs formation channels reviewed in Mapelli (2021). The φ_{iso} population produces a total of 10^5 BBHs with primary mass m_1 distributed according to a truncated power law $p(m_1) \propto m_1^{-2}$ between $5 M_\odot$ and $50 M_\odot$, while m_2 is distributed between $5 M_\odot$ and m_1 with a power law $p(m_2|m_1) \propto m_2$. The φ_{gcl} produces a total of 5×10^3 BBHs with primary mass uniform in $20 M_\odot$ and $90 M_\odot$ and secondary mass uniform in $5 M_\odot$ and m_1 . The overall population of BBHs is defined as the sum of the two channels, i.e. $\varphi_{\text{tot}} = \varphi_{\text{iso}} + \varphi_{\text{gcl}}$. We also assume that a previous analysis using BBHs from the φ_{tot} population has been able to fit the mass spectrum with a broken power and obtained a 10 per cent error on the mass spectrum parameters and overall merger rate. The three populations φ_{iso} , φ_{gcl} , φ_{tot} , and the phenomenological reconstruction of φ_{tot} are represented in Fig. 2. The figure shows how φ_{tot} overlaps with the phenomenological reconstruction. While φ_{iso} , φ_{gcl} fit only the total population in the low and high mass regions with an overlap between $20 M_\odot$ and $50 M_\odot$. We now want to assess the three models φ_{iso} , φ_{gcl} , φ_{tot} with the reconstructed population and find which one is preferred.

The first ingredient that we need, is the evaluation of equation (2) as a function of the BBH masses. In Fig. 3 we show $p(\varphi_j|m_1, m_2)$ computed for all the formation channels. The figure shows the interpretation of $p(\varphi_j|m_1, m_2)$: when we have mass values in the range $m_{1,2} < 20 M_\odot$, the most probable formation channel is φ_{iso} , while when we are looking at binaries with $m_{1,2} > 50$ the most probable formation channel is φ_{gcl} . It is also interesting to note that the complementary formation channel is 100 per cent probable where none of the models considered produces masses, e.g. for the

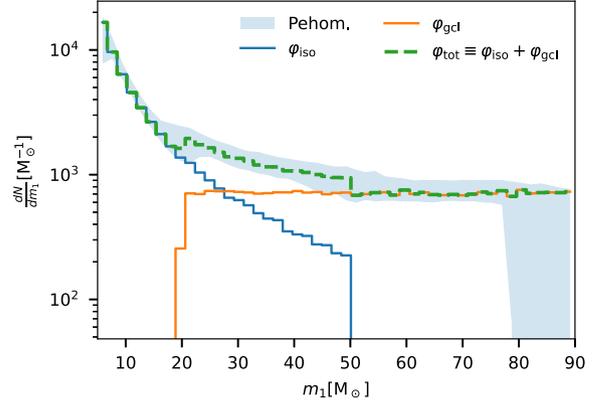


Figure 2. Primary mass distribution for the three synthetic catalogues φ_{iso} , φ_{gcl} , φ_{tot} , and the reconstructed distribution with phenomenological model. The y-axis represent the number of binaries produced per mass bin. The figure shows qualitatively how much the synthetic merger rates overlap with the phenomenological reconstruction of the merger rate.

region $m_2 > m_1$ which is excluded by our simulation. With an evaluation of $p(\varphi_j|m_1, m_2)$, we can now calculate equation (3) by using samples from the phenomenological reconstructed rate. For didactic purposes, let us consider two cases. In the first, we will assume that each formation channel predicts the same amount of BBHs; the preference is solely given by comparing the different mass distributions. In the second, we will include information on how many BBHs each formation channel predicts.

In the first case we obtain $p(\varphi_{\text{iso}}|\{x\}) = 34$ per cent, $p(\varphi_{\text{gcl}}|\{x\}) = 25$ per cent, $p(\varphi_{\text{tot}}|\{x\}) = 34$ per cent, $p(\bar{\varphi}|\{x\}) = 7$ per cent. These probabilities can be used to evaluate how much the population probabilities $p(\theta|\varphi_j)$ overlap with the population probability of the phenomenological rates $p(\theta|\Lambda)$. For instance, φ_{tot} fits 1.36 times better the distribution of masses with respect to φ_{gcl} . If we now include the fact that each formation channel predicts a different amount of BBHs produced, we obtain $p(\varphi_{\text{iso}}|\{x\}) = 0.5$ per cent, $p(\varphi_{\text{gcl}}|\{x\}) = 0.5$ per cent, $p(\varphi_{\text{tot}}|\{x\}) = 91$ per cent, $p(\bar{\varphi}|\{x\}) = 8$ per cent. The clear preference for the φ_{tot} channel is now given by the fact that the number of BBHs produced by φ_{iso} and φ_{gcl} alone is not enough to match the total number of BBHs reconstructed by the phenomenological model.

So far, we have discussed a quantitative method to compare synthetic binary catalogues with phenomenological merger rate reconstructions. This method evaluates the overlap of each model by considering it independent from the others. In the next sections, we will focus on analyses that aim at reconstructing the binary merger rates as a combination of the progenitors’ simulation at our disposal.

3 BUILDING MULTICHANNEL MERGER RATES FROM THE BLACK HOLES PROGENITORS

In this section, we follow a top-to-bottom derivation to show how it is possible to build binary merger rates from synthetic binary catalogues. In the rest of this work, we will focus mostly on BBHs.

Let us assume that we have generated a population of BBHs progenitors $N_*^{\varphi_j}$ that we evolve through an astrophysical channel φ_j to obtain a certain number of BBH mergers $N_{\text{BBH}}^{\varphi_j}$. The merger rate of BBHs for each astrophysical channel can be written as

$$\frac{dN_{\text{BBH}}^{\varphi_j}}{d\theta dz dt} = \mathcal{T}^{\varphi_j}(\theta, \theta_*, z, z_*, t, t_*) \frac{dN_*^{\varphi_j}}{d\theta_* dz_* dt_*}, \quad (4)$$

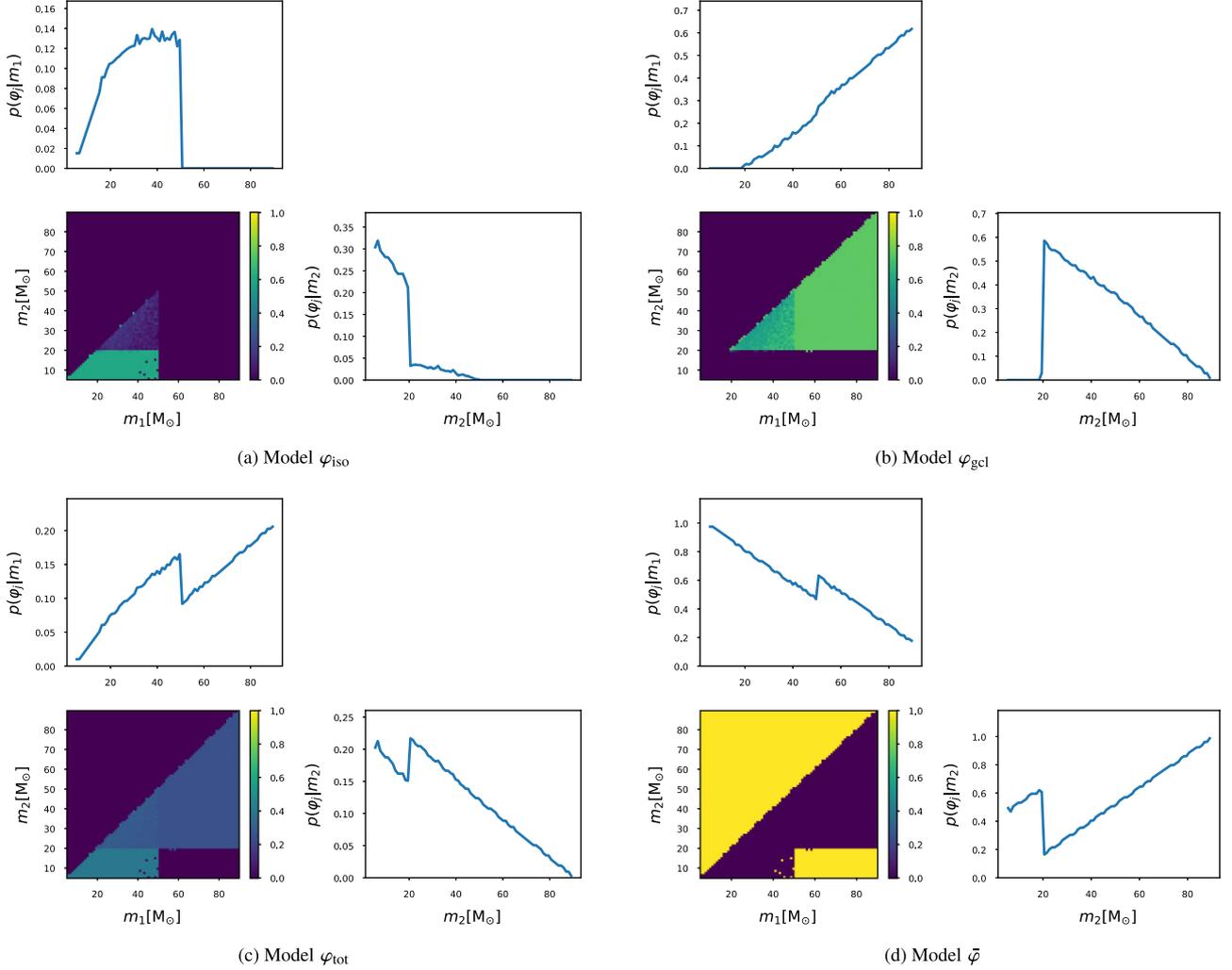


Figure 3. Plots of $p(\varphi_j|\theta, N_{\text{exp}}(\Lambda))$ given in equation (2) for the three synthetic catalogues as a function of the masses. The fourth model is the complementary model and represent the complementary model in the masses space to all the other models. The figures have been generated assuming that all the models predicts the same number of BBHs. This choice was made as in this limit equation (2) reduces to the ratio of the population probabilities of each model. The figures have been generated by dividing the mass space in 6400 bins equally sized. The colourbars indicate the values of models probability given a mass value.

where z_* is the redshift at which the BBH progenitor is formed, θ_* a set of the progenitor parameters such as metallicity and dt_* indicates the time interval at the progenitor redshift. The function \mathcal{T} can be understood as an operator that tells us if a progenitor with parameters θ_* at redshift z_* would produce a BBH with parameters θ at redshift z . For instance, this quantity could be integral over all the various astrophysical properties of the progenitor. A central quantity for many population analyses is the population probability that is built from the binary merger rate as

$$p_{\text{pop}}(\theta, z, t|\varphi_j) = \frac{1}{N_{\text{BBH}}^{\varphi_j}} \frac{dN_{\text{BBH}}^{\varphi_j}}{d\theta dz dt}, \quad (5)$$

where the term $N_{\text{BBH}}^{\varphi_j}$ is the total number of BBHs predicted by the formation channel φ_j .

The idea behind multichannel analysis (Stevenson et al. 2017; Zevin et al. 2017; Wysocki et al. 2018) is to construct (and compare with observed events) an overall BBH merger rate, built as a linear

combination of various formation channels, namely

$$\frac{dN_{\text{BBH}}}{d\theta dz dt} = \sum_j^{N_{\text{syn}}} \lambda_j \frac{dN_{\text{BBH}}^{\varphi_j}}{d\theta dz dt}. \quad (6)$$

The λ_j coefficients are a set of mixture coefficients, that are usually fit in the analysis. The rationale behind this idea is that one single formation channel could not be sufficient to describe the population of observed BBHs (e.g. in the case that BBHs are formed from isolated binary evolution or in globular clusters).

Let us now comment on the physical interpretation for the $\{\lambda\}$ coefficients and their relation to the construction of synthetic binary catalogues. These terms can be understood in terms of progenitors' population. Using equations (4)–(6), the overall BBHs merger rate can be written as

$$\frac{dN_{\text{BBH}}}{d\theta dz dt_s} = \sum_j \mathcal{T}^{\varphi_j}(\theta, \theta_*, z, z_*, t, t_*) \lambda_j \frac{dN_{\text{BBH}}^{\varphi_j}}{d\theta_* dz_* dt_*}. \quad (7)$$

If we assume that the BBHs progenitors distribution is common across all the formation channels considered, then the set of $\{\lambda\}$ should respect the condition $\sum_j \lambda_j = 1$. Namely, the $\{\lambda\}$ represents

the fraction of progenitors that produce BBHs through the formation channels $\{\varphi\}$. In this case we refer to the $\{\lambda\}$ as *fractional mixture coefficients*.

A completely different case can be found when we want to fit the observed BBHs using synthetic catalogues generated from *independent* populations of progenitors. Therefore, the overall population of BBHs will be given by the sum of all the independent sub-populations. In this case, the set $\{\lambda\}$ represents the abundance of each sub-population of BBHs in the observed data. If the observed data are correctly described by the modelled BBHs sub-populations, we would expect each $\lambda_j = 1$. Values of $\lambda_j > 1$ will either indicate that the sub-populations are more numerous or that the transfer function is twice more effective in producing BBHs from the progenitors. The opposite is true for values of $\lambda_j < 1$.

From equation (6) it is possible to define a population probability given as

$$p_{\text{pop}}(\theta, z, t|\{\lambda\varphi\}) = \frac{1}{N_{\text{BBH}}} \frac{dN_{\text{BBH}}}{d\theta dz dt}, \quad (8)$$

where with $\{\lambda\varphi\}$ we indicate a collection of formation channels multiplied by their mixture coefficients. By using equation (6) and the fact that $N_{\text{BBH}} = \sum_j \lambda_j N_{\text{BBH}}^{\varphi_j}$, one can show that the overall population probability is

$$p_{\text{pop}}(\theta, z, t|\{\lambda\varphi\}) = \sum_j \frac{\lambda_j N_{\text{BBH}}^{\varphi_j}}{\sum_k \lambda_k N_{\text{BBH}}^{\varphi_k}} p_{\text{pop}}(\theta, z, t|\varphi_j). \quad (9)$$

The equation above has a direct astrophysical interpretation: if we are provided with a formation channel φ_j that predicts significantly more BBHs than the others, then the overall population probability must be dominated by this channel. The term

$$p(\varphi_j|\{\lambda\varphi\}) = \frac{\lambda_j N_{\text{BBH}}^{\varphi_j}}{\sum_k \lambda_k N_{\text{BBH}}^{\varphi_k}} \quad (10)$$

can be also understood as a probability of the model φ_j given the scalar coefficients $\{\lambda\}$ and the other models $\{\varphi\}$. With this definition, equation (9) can be written as

$$p_{\text{pop}}(\theta, z, t|\{\lambda\varphi\}) = \sum_j p(\varphi_j|\{\lambda\varphi\}) p_{\text{pop}}(\theta, z, t|\varphi_j). \quad (11)$$

Note that there is a fundamental difference between the construction of the above population probability and the one used in several recent works such as Stevenson et al. (2017), Zevin et al. (2017), Bouffanais et al. (2019), Mapelli et al. (2022), Bouffanais et al. (2021), Wong et al. (2021). In these works, the multichannel population probability is built as

$$p_{\text{pop}}(\theta, z, t|\{\Lambda\}) = \sum_j \Lambda_j p_{\text{pop}}(\theta, z, t|\varphi_j), \quad (12)$$

where $\sum \Lambda_j = 1$. The parameters Λ_j effectively represent the fraction of the BBH distribution given by a particular formation channel. Instead, the λ_j defined in this paper represent the fraction of progenitors producing BBHs in a given formation channel. In order to define a progenitor-induced BBH population probability, it is important to take into account the term $p(\varphi_j|\{\lambda\varphi\})$. In Section 5.3.1 we provide an example to discuss how these two quantities are related and can be converted to each other. For now let us give a simple example illustrated in Fig. 4 to better understand the meaning of the λ_j and Λ_j coefficients.

Let us assume to be provided with a set of progenitors producing BBHs via two formation channels φ_1 and φ_2 , with the first formation channel predicting two times more BBHs than the other. Let us

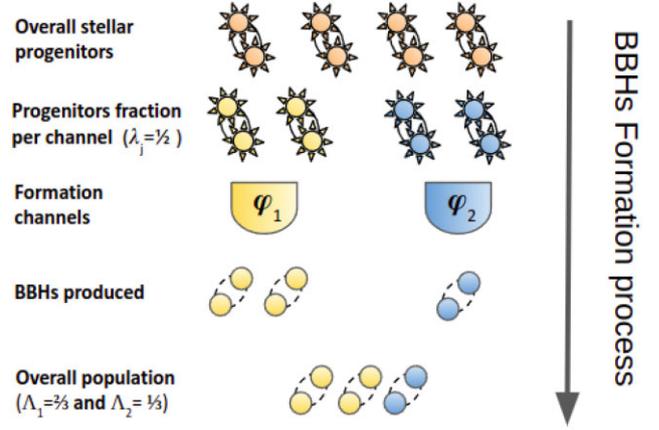


Figure 4. The illustration shows the relation between the coefficients λ_j and Λ_j by following the BBHs from the stellar progenitors. The figure starts from a common population of stellar progenitors, half of which enter a formation channel φ_1 (yellow) and the other half φ_2 (blue). The first formation channel is two times more efficient than the second in producing BBHs. At the end 2/3 of the observable population of BBHs has been produced in φ_1 and 1/3 in φ_2 .

also assume that 1/2 of progenitors enter the first formation channel and 1/2 of the second. In other words $\lambda_1 = \lambda_2 = 1/2$. When we look at the population distribution of BBHs, we would find that 2/3 of the BBHs are produced in the formation channel φ_1 and 1/3 by φ_2 . In other words $\Lambda_1 = 2/3$, $\Lambda_2 = 1/3$. Therefore, if we perform our inference using equation (12), we cannot directly use the Λ_j to draw conclusions about the BBHs progenitors. We can only draw conclusions about the fraction of BBHs produced in a given formation mechanism.

On the one hand, one of the advantages of this method is that the mixture coefficients Λ_j/λ_j are directly linked to the population of BHs and their progenitors. On the other hand, this type of methodology is less flexible than phenomenological models and can only fit population features (e.g. PISN) that are already present in the synthetic catalogues considered. In principle, one could also consider adding a phenomenological model to this approach by defining an additional mixture coefficient $\Lambda_{\text{phenom}}/\lambda_{\text{phenom}}$ associated with the phenomenological model. In this scenario, we could fit a set of mixture coefficients associated with the synthetic catalogues and one mixture coefficient associated with the phenomenological model. However, this approach could have two drawbacks. First, the phenomenological model needs to be fitted with a set of continuous variables thus increasing the dimensionality and computational cost of the analysis. Secondly, phenomenological models are very flexible and thus we expect the inference on the phenomenological models to be less informative. Therefore, we suggest performing two separate analyses. One uses synthetic catalogues and mixing coefficients, and the other uses phenomenological models and then compares Bayes factors.

4 PROGENITORS MULTICHANNEL BAYESIAN ANALYSES

We now discuss the case in which we would like to reconstruct the merger rate for multiple formation channels starting from the observed BBHs. Differently from what we discussed in the previous section, in this case, we will not use phenomenological models and we will rely solely on synthetic binary catalogues. We will

use the mixture model approach presented in Section 3 and write the overall BBHs merger rate as in equation (6). We will discuss in Section 4.1 the statistical background for multichannel analyses based on synthetic catalogues, in Section 4.2 how priors on the mixture coefficients can be chosen and in Section 4.3 computational difficulties related to this kind of analysis.

4.1 Statistical method

The hierarchical likelihood of having N_{obs} GW events from data $\{x\}$ conditioned on the set of models φ_j and the mixture coefficients λ_j is (see Vitale et al. 2022 for a bottom-to-top derivation)

$$p(\{x\}|\{\lambda\varphi\}) \propto e^{-N_{\text{exp}}} \prod_i^{N_{\text{obs}}} T_{\text{obs}} \int \frac{p(x_i|\theta) dN_{\text{BBH}}}{1+z} d\theta dz dt, \quad (13)$$

where $p(x_i|\theta)$ is the GW likelihood and N_{exp} is the number of expected events observable in a given observing time T_{obs} . The GW likelihood quantifies the uncertainties with which the source astrophysical parameters, such as luminosity distance and detector-frame masses, are determined. Equation (13) can be rewritten in the alternative form (Vitale et al. 2022)

$$p(\{x\}|\{\lambda\varphi\}) \propto e^{-N_{\text{exp}}} N_{\text{exp}}^{N_{\text{obs}}} \prod_i^{N_{\text{obs}}} \frac{\int p(x_i|\theta) p_{\text{pop}}(\theta|\{\lambda\varphi\}) d\theta}{\beta(\{\lambda\varphi\})}, \quad (14)$$

where $p_{\text{pop}}(\theta|\{\lambda\varphi\})$ is the population probability defined as in equation (11), and $\beta(\{\lambda\varphi\})$ is the selection effect (see later).

Our aim is to quickly evaluate equation (14) as a function of the mixture coefficients λ_j . We will factorize equation (14) in several terms that can be computed once for each formation channel φ_j and rescaled with λ_j to quickly evaluate the hierarchical likelihood. The numerator factor in the product of equation (14) can be rewritten as,

$$\int p(x_i|\theta) p_{\text{pop}}(\theta|\{\lambda\varphi\}) d\theta = \sum_j p(\varphi_j|\{\lambda\varphi\}) \mathcal{L}_{i,j}, \quad (15)$$

where we have expanded $p_{\text{pop}}(\theta|\{\lambda\varphi\})$ using equation (11) and we have defined

$$\mathcal{L}_{i,j} = \int p(x_i|\theta) p_{\text{pop}}(\theta|\varphi_j) d\theta. \quad (16)$$

The $\mathcal{L}_{i,j}$ can be evaluated numerically once for each i th GW event and j th formation channel. We also recall that $p(\varphi_j|\{\lambda\varphi\})$ can be constructed using equation (10) and using only the number of BBHs predicted by each model and the mixture coefficients λ_j . The selection effect $\beta(\{\lambda\varphi\})$ can be quickly computed by knowing the total number of BBHs predicted by each formation channel and the fraction of BBHs detectable by each channel $\beta(\varphi_j)$, namely

$$\beta(\{\lambda\varphi\}) = \frac{\sum_j \lambda_j N_{\text{BBH}}^{\varphi_j} \beta(\varphi_j)}{\sum_j \lambda_j N_{\text{BBH}}^{\varphi_j}}. \quad (17)$$

Finally, the Poissonian term

$$e^{-N_{\text{exp}}} N_{\text{exp}}^{N_{\text{obs}}}$$

in equation (14) can be easily computed by recognizing that $N_{\text{exp}} = \sum_j \lambda_j N_{\text{BBH}}^{\varphi_j} \beta(\varphi_j)$.

This term is usually marginalized out in multichannel analyses focusing on BBHs population as performed in Zevin et al. (2017), Stevenson et al. (2017), Mapelli et al. (2022), Bouffanais et al. (2019, 2021), Zevin et al. (2021). To do so, we need to introduce a ‘nuisance scaling parameter’ A in common to all the population models such

that $N_{\text{BBH}} = A \sum_j \lambda_j N_{\text{BBH}}^{\varphi_j}$. If we take a prior on A uniformly distributed in logarithmic space, it is possible to marginalize out the Poissonian term and equation (14) reduces to

$$p(\{x\}|\{\lambda\varphi\}) \propto \prod_i^{N_{\text{obs}}} \frac{\int p(x_i|\theta) p_{\text{pop}}(\theta|\{\lambda\varphi\}) d\theta}{\beta(\{\lambda\varphi\})}. \quad (18)$$

Note that, in comparison to population analyses based on phenomenological models (Fishbach & Holz 2017; Vitale et al. 2019; Abbott et al. 2020c, 2021c; Callister et al. 2021; Farah et al. 2022), the parameter A effectively act as a common rescaling for the BBH merger rate density $R_0(\varphi_j)$ identified by each model. In other words, equation (18) reconstructs the BBHs astrophysical distributions in terms of masses and redshift without accounting for the absolute merger rate. While this choice is mathematically correct and reconstructs the correct distribution in masses and redshift of BBHs, one should be careful about the astrophysical interpretation. For instance, the synthetic simulations might predict many more events than the observed ones, while still being able to fit the redshift and mass distribution. This choice is usually done when the rates of the different formation channels are highly uncertain and not correlated with the astrophysical processes that characterize the mass and redshift distributions of BBHs.

To summarize, in order to quickly perform a multichannel analysis using several formation channels φ_j , we need to: (i) Estimate the total number of BBHs produced by each formation channel $N_{\text{BBH}}^{\varphi_j}$ and their detectable fraction $\beta(\varphi_j)$, (ii) for each formation channel and GW event estimate the term $\mathcal{L}_{i,j}$ in equation (16) and (iii) for some values of the set $\{\lambda\}$ use equations (15) and (17) to effectively build the hierarchical likelihood.

4.2 Priors on the mixture coefficients

We now discuss how priors on the mixture coefficients $\{\lambda\}$ can be chosen according to the astrophysical case considered. In Section 3 we considered two cases: the case in which each formation channel has an independent sub-population of progenitors and the case for which the population of BBHs progenitors is in common to each formation channel. In the former, the λ_j are independent of each other and a value of $\lambda_j = 1$ indicates that the BBH formation channel is observed in data as the model predicts. In this case, each prior on λ_j can be chosen independently and from an astrophysical point of view, this case corresponds to changing the initial astrophysical conditions (e.g star formation rate) of the simulation. In the latter, the set of $\{\lambda\}$ must satisfy the constraint $\sum_j \lambda_j = 1$, and these parameters effectively represent the relative fraction of BBHs observed produced by each channel. In terms of astrophysics, this represents the case for which the initial conditions of the simulation are set but the astrophysical evolution prescriptions are changed.

When the $\{\lambda\}$ must be normalized (common BBHs progenitors), a non-trivial bound is introduced in the joint prior of the mixture coefficients. One possibility is to build a joint prior that satisfies the normalization constraint by drawing sequentially the values of λ_j from a cascade of conditional probabilities. Namely, we write the joint prior as

$$p(\{\lambda\}) = \prod_j p(\lambda_j|\{\lambda\}_{i<j}), \quad (19)$$

where $\{\lambda\}_{i<j}$ indicates a set of λ_i with index lower than j . By choosing uniform conditional priors, the above equation can be

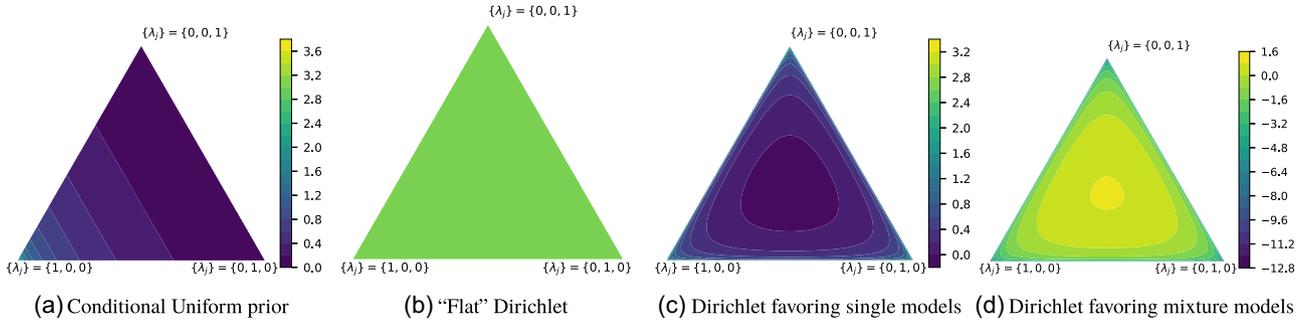


Figure 5. Representation on a 2-simplex of the logarithm of the prior probability distribution on the fractional mixture coefficients $\{\lambda\}$ of three models. From left to right: Conditional uniform prior with ordering preference, ‘Flat’ Dirichlet prior with concentration parameters 1, Dirichlet prior favouring single models with concentration parameters 0.5 and Dirichlet prior favoring mixture models with concentration parameters 3.0.

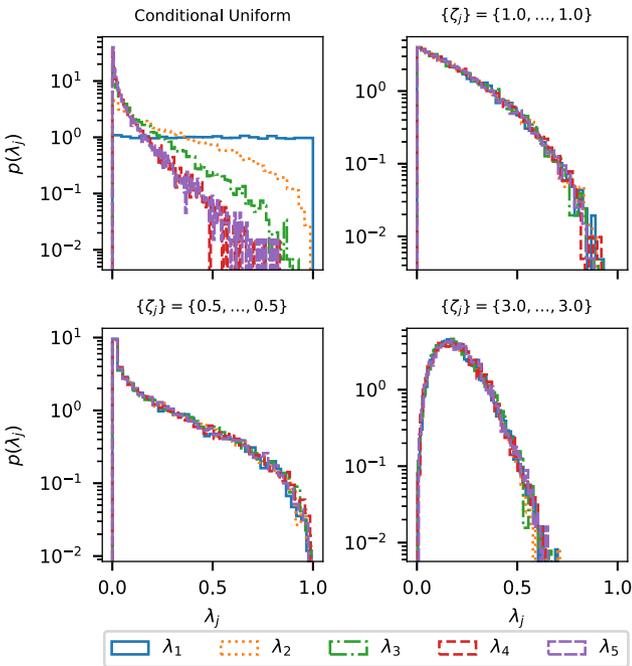


Figure 6. Marginal priors on five mixture models. The different panels correspond to the marginal priors for different choices as described in Fig. 5 and Section 4.1. The conditional uniform prior is given in equation (19).

written as

$$\begin{aligned}
 p(\lambda_i) &= p(\lambda_1) \\
 p(\lambda_j | \{\lambda\}_{i < j}) &= \begin{cases} p(\lambda_j | \{\lambda\}_{i < j}) & \text{if } 0 \leq \lambda_j \leq 1 - \sum_{i < j} \lambda_i \\ 0 & \text{if } \lambda_j > 1 - \sum_{i < j} \lambda_i \end{cases} \\
 p(\lambda_N | \{\lambda\}_{i < N}) &= \delta \left(1 - \sum_{i < N} \lambda_i \right).
 \end{aligned}$$

We note that this prior choice is not optimal for multichannel studies as it introduces an ordering preference. In Fig. 5 (left-hand panel), we show the logarithm of a joint prior to built in this way for the case that we are provided with three astrophysical formation channels. As it can be seen from the figure, this prior naturally prefers the first ordered model. In the case that multiple models are provided, this type of prior will strongly disfavour models that are ordered in the last. We display this effect in Fig. 6 by showing the marginal prior

distributions in the case that we are provided with five astrophysical channels.

A more natural choice that removes the problem of model ordering is to use a Dirichlet distribution on the $\{\lambda\}$ as done in Stevenson et al. (2017), Zevin et al. (2017), Wysocki et al. (2018). The Dirichlet distribution ensures the normalization of the $\{\lambda\}$ and also provides a set of concentration parameters $\{\zeta\}$ governing how the probability is distributed on the plane identified by $\sum \lambda_j = 1$. Fig. 5 shows the logarithm of the Dirichlet prior to different choices of the $\{\zeta\}$ parameters. If $\zeta_j = 1$, the prior probability is uniform across the combination of all the formation channels. If $\zeta_j < 1$, the prior will prefer to build the overall BBH rate using a single formation model. Finally, if $\zeta_j > 1$, the prior will prefer to build the BBH rate as a superposition of all the models. In general, as we will show in Section 5.3, the $\{\zeta\}$ parameters can also be treated as free parameters to infer. The marginal priors on the λ_j in the cases presented for a Dirichlet distribution are shown in Fig. 6. One can observe that the marginal priors are equal for all the models.

We therefore argue that Dirichlet priors should be used when performing this type of analyses.

4.3 Evaluating Monte Carlo integrals

The calculation of the hierarchical likelihood in equation (13) requires the evaluation of several numerical integrals. A first implicit integral is given by the calculation of the fraction of BBHs that we expect to detect. This integral is given by the product of the detection probability as a function of the BBHs parameters with the BBHs population distribution and merger rate. The integral is not evaluated analytically and a common technique to estimate it is by using injection studies. The idea is simply to generate GW injections in noise from the desired BBH population and estimate what is the detectable fraction. Farr (2019) showed that the number of detectable injections should be at least four times higher than the GW events considered in the analysis. Otherwise, the evaluation of the hierarchical likelihood is not numerically stable.

The other term that requires an integral over the BBH population is equation (16). This integral is usually evaluated as a *Monte Carlo integral* by using N_s samples from the posterior of each BBH detected. This approach consists in approximating the integral as

$$\mathcal{L}_{i,j} \approx \frac{1}{N_s} \sum_j^{N_s} \frac{p_{\text{pop}}(\theta | \varphi_j)}{p_0(\theta)}, \quad (20)$$

where $p_0(\theta)$ is a prior applied to calculate the BBH posteriors on the binary parameters and $p_{\text{pop}}(\theta|\varphi_j)$ is the population prior associated to the formation channel. Alternatively, one can decide to perform the Monte Carlo integral by summing over simulated BBHs from the formation channel $p_{\text{pop}}(\theta|\varphi_j)$ and write

$$\mathcal{L}_{i,j} \approx \frac{1}{N_s} \sum_j^{N_s} \frac{p(\theta|x_i)}{p_0(\theta)}. \quad (21)$$

In principle, we would expect equations (20) and (21) to return the same result. Both approaches have in common one necessity, either the BBH population of the formation channel or the GW posterior of observed events should be known as a function of the parameters θ . These analytic functions are not usually known, in fact, we are usually provided with either a list of posterior samples from $p(\theta|x_i)$ or a list of BBHs simulated from $p_{\text{pop}}(\theta|\varphi_j)$. One possibility to compute analytically this probability from a set of samples, is by using kernel density estimates, histograms, or non-parametric fitting such as the ones proposed in Wysocki et al. (2018), Golomb & Talbot (2022), Del Pozzo et al. (2018), Tiwari, Fairhurst & Hannam (2018), Sadiq, Dent & Wysocki (2022), Delfavero et al. (2021), Rinaldi & Del Pozzo (2022), Delfavero et al. (in preparation). If equation (20) is used, the sum is performed over GW posterior samples and the fit is on the BBH population of the formation channel. If equation (21) is used, the sum is over the BBHs predicted by the formation channel and the GW posterior is evaluated by the fit. In both cases, a possible Gaussian kernel fitting should always be validated against the original distribution.

A rule of thumb to decide what is best suited to evaluate the Monte Carlo Integral is the following. Equation (20) can be used if the formation channel has a phenomenological (or semi-analytical) model and no fitting is needed. Equation (20) can also be used when the BBH parameters from GW data are measured with a precision significantly lower than the typical ranges covered by the BBH formation channels. Equation (21) can be used in this case that the range spanned by the BBH formation channels is comparable or significantly lower than the precision with which we are able to measure BBH parameters from data.

Current population studies are based on BBHs, for which we expect formation channels to cover a wider range in masses and redshift to the typical error budgets estimated from GW data. That is why so far Monte Carlo Integrals are mostly evaluated with equation (20).

5 CASE STUDIES: THE COMMON ENVELOPE EFFICIENCY AND PROGENITORS METALLICITY

In this section, we present case studies to show how synthetic catalogues of BBH mergers can be used with GW population studies. For all the test cases (unless specified), we consider the BBH rates with the hierarchical likelihood in equation (14). We use the model by Srinivasan et al. (in preparation) for our binary population. In this simple model, the binary population is based on the code COSMIC (Breivik et al. 2020) to simulate BBH mergers. The star formation rate is parametrized in terms of star metallicity, galaxy mass, and redshift of formation of the binary following Lamberts et al. (2016). For every value of metallicity, a population of BBH merger progenitors is generated by using COSMIC to evolve zero-age main-sequence stars, selecting those that form BBH mergers. The overall population of BBH mergers is obtained by re-weighting the cosmic BBHs progenitors by the star formation rate.

5.1 COSMIC simulations

COSMIC simulates binary evolution based on prescriptions that model physical processes such as stellar winds, mass transfers between the binary, and supernovae kicks. One of the prescriptions of interest is the unstable mass transfer during the binary evolution that results in a common envelope (CE) phase parametrized by an efficiency α_{CE} . Depending on the value of the CE efficiency, stellar binaries can be more or less efficient in producing BBH mergers (Barrett et al. 2018). We explore the effect of CE efficiencies. Specifically, we consider CE efficiency values of $\alpha_{\text{CE}} = \{0.3, 0.5, 1.0\}$. The other prescriptions of the COSMIC simulations are set to their default values reported on COSMIC 3.4.0.² As this study focuses on statistical inference, we choose not to optimize the model to fit observed distributions and rates. For each simulation, COSMIC provides us with the distribution of time delays between the progenitor formation and the BBH merger. The procedure of building the population depends on the type of multichannel analysis we consider (see later).

5.2 Generation of the GW mock catalogue

To build a mock catalogue of observed GW events, we use an approach similar to Fishbach, Holz & Farr (2018), Farr et al. (2019) to simulate the detection of GW events and the estimation of source masses and redshift for each detected binary. For each binary, we calculate the matched filter SNR ρ as

$$\rho = 8 \left(\frac{\mathcal{M}_c}{26 M_\odot} \right)^{5/6} \left(\frac{d_L}{1500 \text{ Mpc}} \right) w, \quad (22)$$

where \mathcal{M}_c is the binary redshifted chirp mass and d_L is the binary luminosity distance (calculated using a Planck cosmology; Planck Collaboration XIII 2016). The scaling factors for the chirp mass and the luminosity distance are chosen to assume a network composed by LIGO Hanford, Livingston, and Virgo with typical detection ranges for O4 (Abbott et al. 2018). The luminosity distance scaling is calculated with the single-detector reach distances reported in Abbott et al. (2018). The parameter w is a scaling factor that takes into account the fact that not all the detectors in the network are optimally oriented with respect to the source position (Dominik et al. 2015). The cumulative distribution of w for a three-detector network is publicly available.³

Once the optimal SNR is calculated for each binary, we draw a ‘observed’ SNR ρ_{obs} from a non-central χ^2 -square distribution (with non-centrality parameter ρ) and with a 6 degrees of freedom since we have three detectors in the network. Binaries are detected if they exceed an observed SNR of 12. For each detected binary, we then draw an ‘observed’ chirp mass $\mathcal{M}_{c,\text{obs}}$ and symmetric mass ratio η_{obs} using the same likelihoods in Appendix B of Farr et al. (2019). Once we are provided with a set of ‘observed’ chirp masses, symmetric mass ratio, and SNR, we generate mock posterior samples on their original ‘true’ values using the likelihood models from which they were generated. These mock posterior samples are then converted to posterior samples in source frame masses and in redshift using equation (22) (and correcting for the change of variable $\{\rho, \mathcal{M}_c, \eta\} \rightarrow \{d_L, m_1, m_2\}$) that we then use to generate mock posterior samples on redshift and source-frame masses for each detected signal.

We study the reconstruction of the BBH formation channels by extracting 64, 128, 256, 512, 1024, and 2048 binaries from the set

²<https://cosmic-popsynth.github.io/docs/stable/>

³<https://pages.jh.edu/eberti2/research/>

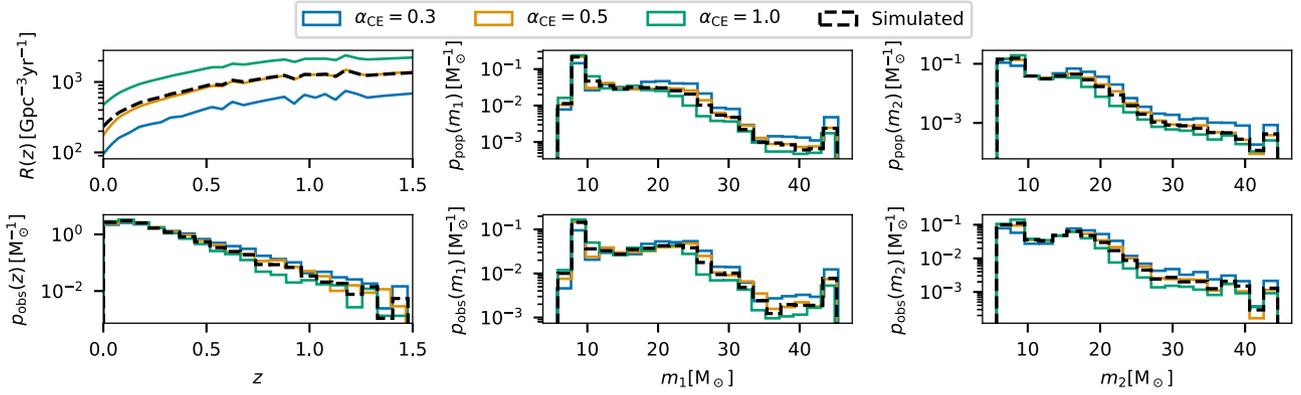


Figure 7. Left-hand panel: BBH merger rate redshift evolution for the three common envelope models we consider. Centre: Distribution of the primary source mass for the models we consider. Right-hand panel: Distribution of the secondary source mass for the models we consider. In all the panels, the simulated mixture population generated with fractions $\lambda = \{0.4, 0.3, 0.3\}$ is indicated with a black dashed line. The bottom panels show the observed distributions of BBHs in redshift and masses once an SNR cut of 12 is applied.

of detected signals. We use the statistical approach described in Section 4 and the likelihood function in equation (13) to find posterior distributions on the mixture coefficients. This study makes use of the BILBY code (Ashton et al. 2019; Romero-Shaw et al. 2020) and its nested sampling DYNESTY implementation (Higson et al. 2019) to sample from the posterior distribution of the mixture coefficients.

5.3 Measuring the progenitors common envelope efficiency from BBHs ($\sum_j \lambda_j = 1$)

In our first case study, we would like to infer CE efficiency from the observed GW events. In this case, the formation channels from which we build our BBH catalogues are the COSMIC simulations with $\alpha_{\text{CE}} = \{0.3, 0.5, 1.0\}$. As described in Section 3, this is the case for which the progenitor population of BBHs is in common between the formation channels. The $\{\lambda\}$ represent the fraction of BBHs produced from progenitors with a given CE. For our case study, we assume that 40 per cent of the BBH population is produced with $\alpha_{\text{CE}} = 0.3$, 30 per cent with $\alpha_{\text{CE}} = 0.5$, and 30 per cent with $\alpha_{\text{CE}} = 1.0$. Note that this is a toy model to show how the method can work, as massive binaries might not display different CE efficiencies (Wong et al. 2022).

By using equation (6) and the chosen mixture coefficients, we build the overall BBHs merger rate. In Fig. 7 we show the BBHs rate evolution in terms of redshift and masses for the three models and for the overall population that we simulate. While the three simulations predict similar mass distributions in shape, they significantly differ in terms of absolute merger rates (Hurley, Tout & Pols 2002; Ricker et al. 2018; Mandel & Broekgaarden 2022). We run the multichannel analysis using two sets of priors for the fractional mixture coefficients and we consider also the BBHs merge rates in the hierarchical likelihood. In the first analysis, we use a Dirichlet prior with concentration parameters fixed to $\{\zeta\} = \{0.5, 0.5, 0.5\}$ favouring single models, while in the second analysis we also allow the concentration parameters to vary in a uniform distribution between $[0.01, 100]$. Fig. 8 shows the reconstructed marginal posteriors of the mixture fractions between the three CE models. From the plot, we can see that when we are provided with few GW events, the constraints on the fractional mixture coefficients are weak. As more and more GW events are detected, the constraints on the mixture coefficients improve. With 2048 events a precision of $\sim 4 - 5$ per cent is reached on the determination of the mixture fractions. In Appendix A we also

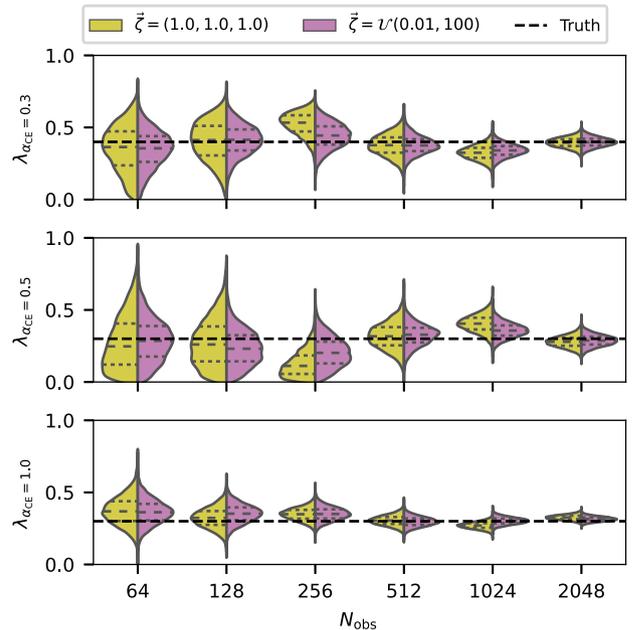


Figure 8. Posterior distributions on the fractional mixture coefficients for the three CE efficiency models as a function of the number of detected events. The horizontal black dashed line indicates the true value used for the simulation (that is always included in the 90 per cent credible intervals). The grey dashed lines in the posterior indicate the posteriors median and symmetric quartiles (50 per cent credible intervals). The yellow posteriors are generated by fixing a Dirichlet prior to distribution. The pink posteriors are generated allowing the Dirichlet parameter to change.

provide a more detailed discussion about the correlations among the various fractional coefficients and the Dirichlet concentration parameters.

5.3.1 Reconstructing progenitors fractions from BBH fractions

As we argue in Section 3, there is a fundamental difference between constructing the population probability using equation (11) or equation (12). In the former case, we are inferring the *fraction of progenitors entering a formation channel* (λ_j), while in the latter we are inferring the *fraction of BBHs produced in a formation channel*

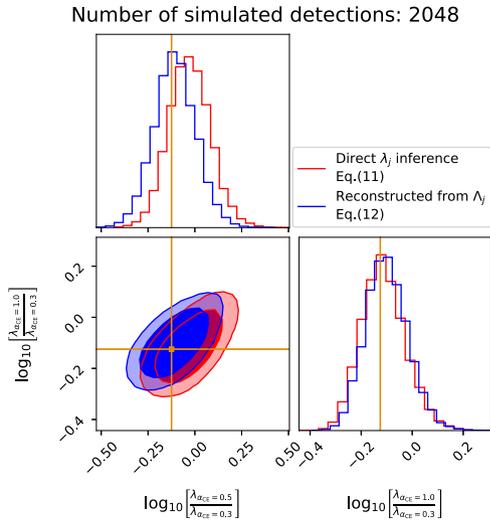


Figure 9. Joint posterior distribution on the progenitors λ_j ratios for the CE simulations using 2048 GW detections. The figure compares the direct inference of the progenitor fractions (red contour) with the indirect reconstruction from BBHs population fractions (blue contour). The solid lines mark the 1σ and 2σ contours. The yellow solid lines mark the simulated population.

(Λ_j). These two quantities can be related a posteriori comparing equations (11) and (12) and noting that

$$\frac{\lambda_j N_{\text{BBH}}^{\varphi_j}}{\sum_k \lambda_k N_{\text{BBH}}^{\varphi_k}} = \Lambda_j. \quad (23)$$

From the above relation, it follows that

$$\frac{\lambda_j}{\lambda_i} = \frac{\Lambda_j}{\Lambda_i} \frac{N_{\text{BBH}}^{\varphi_i}}{N_{\text{BBH}}^{\varphi_j}}, \quad (24)$$

i.e. the ratio of the progenitors fraction entering the formation channel j and i can be calculated by scaling the ratio of BBH fractions produced in the formation channel j and i (and vice versa). Indeed it is interesting to note that the two ratios coincide when the formation channels have the same efficiency in producing BBHs.

For instance, in our previous example, we constructed a BBH population that was composed by 40 per cent, 30 per cent, and 30 per cent of progenitors with CE efficiency of 0.3, 0.5, and 1.0, respectively. We perform again the multichannel analysis but this time using equations (11)–(12) implemented in the hierarchical likelihood in equation (18) (marginalizing out the rates). In Fig. 9, we compare the distribution of the progenitor ratios λ_j/λ_i obtained in Section 5.3 and the ones reconstructed using equation (24) and the BBHs ratios Λ_j/Λ_i . We can see that the progenitors ratios can be effectively reconstructed from the BBHs ratios.

5.4 Measuring the BBHs progenitors metallicity ($\sum \lambda_j \neq 1$)

In the second case study, we use the simulation with CE efficiency 1.0 and we divide the population of BBHs progenitors according to their metallicity, uniformly divided in base 10 logarithm between $Z = 5.0 \times 10^{-3} Z_\odot$ and $Z = 1.6 Z_\odot$. We are therefore in presence of independent sub-populations of BBHs progenitors, as described in Section 3. Each sub-population of progenitors provides us with a sub-population of BBHs. The total BBH merger rate is the sum of the rates of these ten sub-populations. For this case study, we assume that the sub-populations are not present as predicted by the COSMIC simulation. Instead, we assume that each sub-population contributes

to the overall BBHs merger rate with multiplicity coefficients $\lambda = \{0.5, 0.6, 0.7, 0.8, 0.9, 1.0, 1.1, 1.2, 1.3, 1.4\}$ (ordered in terms of increasing metallicity bins). As an example, the BBHs produced from a progenitor with metallicity $Z_\odot = 2Z_\odot$ produce 40 per cent more BBHs with respect to the initial model predictions.

Fig. 10 shows the BBH merger rate and mass distributions for these sub-populations and the true population created from their sum. We observe that in this simple model the progenitor metallicity introduces many different features in the mass spectrum and merger rate. We perform the reconstruction of the mixture coefficients for the sub-populations using priors on λ_j independent from each other and uniform between 0.5 and 1.5. Fig. 11 shows the marginal posterior distribution among all the mixture coefficients. We can see that not all of the fractional parameters can be constrained in the proposed prior range. However, we can observe that in general, models which predict more BBHs are better constrained than models that predict less. This is expected as progenitors entering channels predicting more BBHs are easier to constrain. This case study shows that synthetic binary catalogues can be used with GW events to probe the BHs progenitor metallicity but also the evolution of the star formation rate.

6 CONCLUSIONS

In this paper, we have described in detail how to employ and statistically interpret synthetic compact binaries in light of GW detections. In particular, we have focused on analyses trying to infer and constrain the presence of BBHs progenitors in multiple channels.

In Section 2, we have presented for the first time an efficient method to evaluate the ‘match’ between synthetic catalogues of binaries and phenomenological reconstructed astrophysical rates. Given the phenomenological rate reconstruction, the method is able to assign a probability to each of the synthetic catalogues to be representative of the estimated rate. The probabilities can be used to quickly evaluate how much a model fits the phenomenological rates with respect to the other.

In Section 3, we have formalized how different progenitor populations can be used to build multichannel population models. We have discussed how an overall BBHs merger rate should be built and interpreted in terms of progenitors mixture coefficients $\{\lambda\}$. We have argued that in the case that the progenitor population is common across the different BBHs formation channels, then one can use fractional mixture coefficients to infer the percentage of progenitors undergoing through each formation channel ($\sum_j \lambda_j = 1$). We have also discussed the case for which we are in presence of multiple and independent sub-populations, showing that in this case the $\{\lambda\}$ can be assumed to be independent of each other. From an astrophysical perspective, normalized mixture coefficients can be used when building multichannel progenitors models for which only the stellar evolution is modified. While independent mixture coefficients can be used when stellar evolution models are fixed, but the original progenitor rates (initial conditions) are varied.

In Section 4, we have reviewed the hierarchical statistical method used to employ synthetic populations with observed GW events, describing the method in light of the multichannel analyses presented in Section 3. In Section 4, we have also described technical aspects related to the computational implementation of this methodology and the prior choice that should be made on the mixture coefficients in order to not introduce an ‘ordering preference’.

In Section 5, we have presented two case studies for BBH progenitors’ multichannel analyses. In the first case, we discussed a possible measure of the CE efficiency parameter. Based on an astrophysical model for BBH formation, we show that binary evolution criteria,

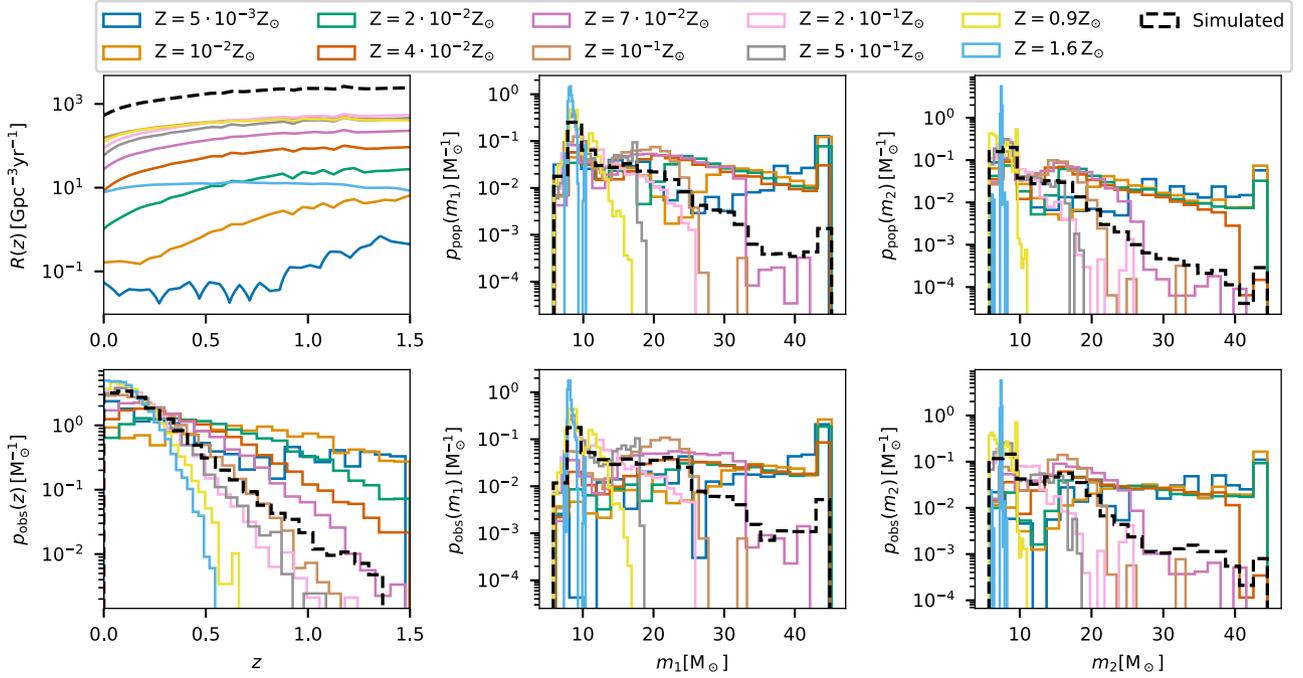


Figure 10. Left-hand panel: BBH merger rate redshift evolution for the metallicity sub-populations models considered. Centre: Distribution of the primary source mass for the models considered. Right-hand panel: Distribution of the secondary source mass for the models considered. The ‘simulated’ curve shows the injected population and is obtained summing all the 10 metallicity bins sub-populations. The bottom panels show the observed distributions of BBHs in redshift and masses once an SNR cut of 12 is applied.

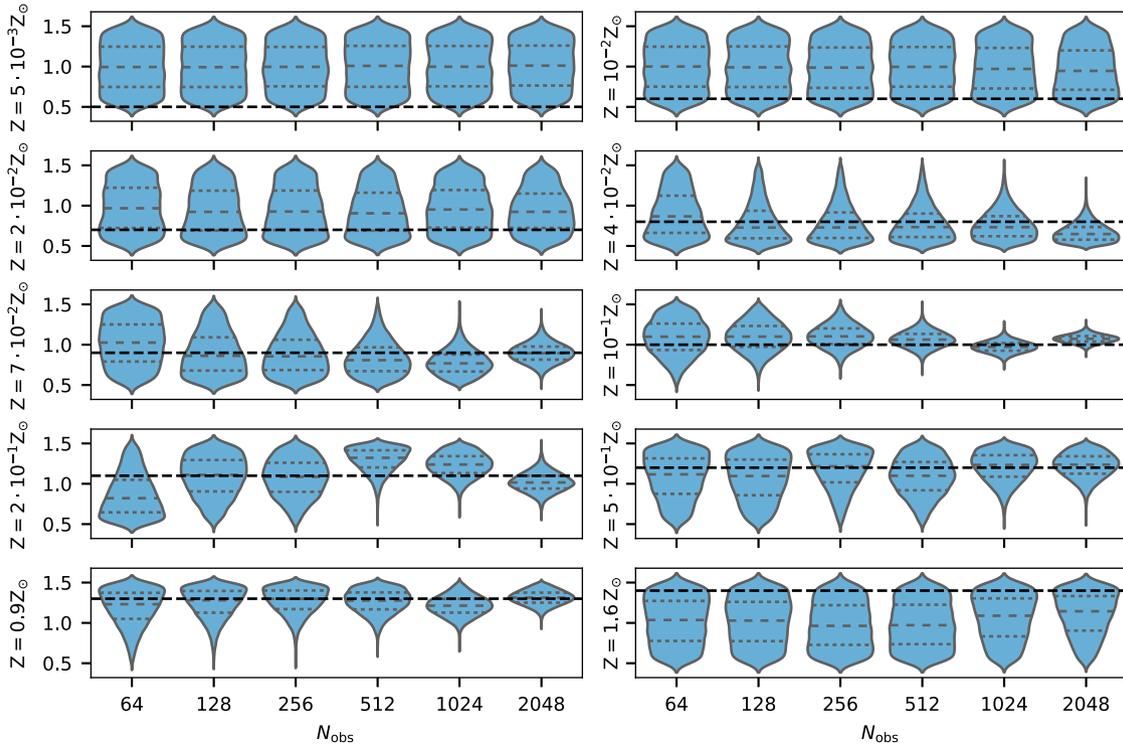


Figure 11. Posteriors of the mixture coefficients associated with the metallicity-dependent sub-populations as a function of the number of detected events. The horizontal black dashed line indicates the true value used for the simulation (that is always included in the 90 per cent CI). The grey dashed lines in the posterior indicate the posteriors’ median and symmetric quartiles (50 per cent CI).

such as the CE efficiency, could be constrained to good precision with a few thousand of detection (or in the coming years). The second case that we discussed, made use of BBH sub-populations divided into progenitor metallicity bins. We have shown that, provided the astrophysical model and star formation rate, some of the BBHs progenitors' metallicity can be constrained with thousands of GW detections.

With the next two observing runs O4 and O5, the LIGO, Virgo, and KAGRA detectors will reveal thousands of BBHs and possibly hundreds of BNSs (Abbott et al. 2018). Using this observed population it will be possible to probe the progenitor properties of the GW sources and unveil the astrophysical processes bringing to compact object formation.

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DATA AVAILABILITY

Results and simulations obtained in this paper are generated using the GWPARENTS python package, which can be download at <https://github.com/simone-mastrogiovanni/gwparents>.

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APPENDIX A: COMMON ENVELOPE EFFICIENCY WITH DIRICHLET CONCENTRATION PARAMETERS

In this appendix, we run the same inference on the progenitor fractions as in Section 5.3 but using also priors on the Dirichlet concentration parameters. It is interesting to see what are the correlations in the determination of the mixture fractions and the concentration parameters α . In Fig. A1, we show their joint posterior distribution for 2048 GW detections. We note that the fractional mixture parameters of the CE efficiency 0.5, 1.0 show a non-negligible anticorrelation. This is due to the fact that these two formation channels predict similar values (and higher with respect to $\alpha_{\text{CE}} = 0.3$) of the BBH merger rate, see Fig. 7. These two models

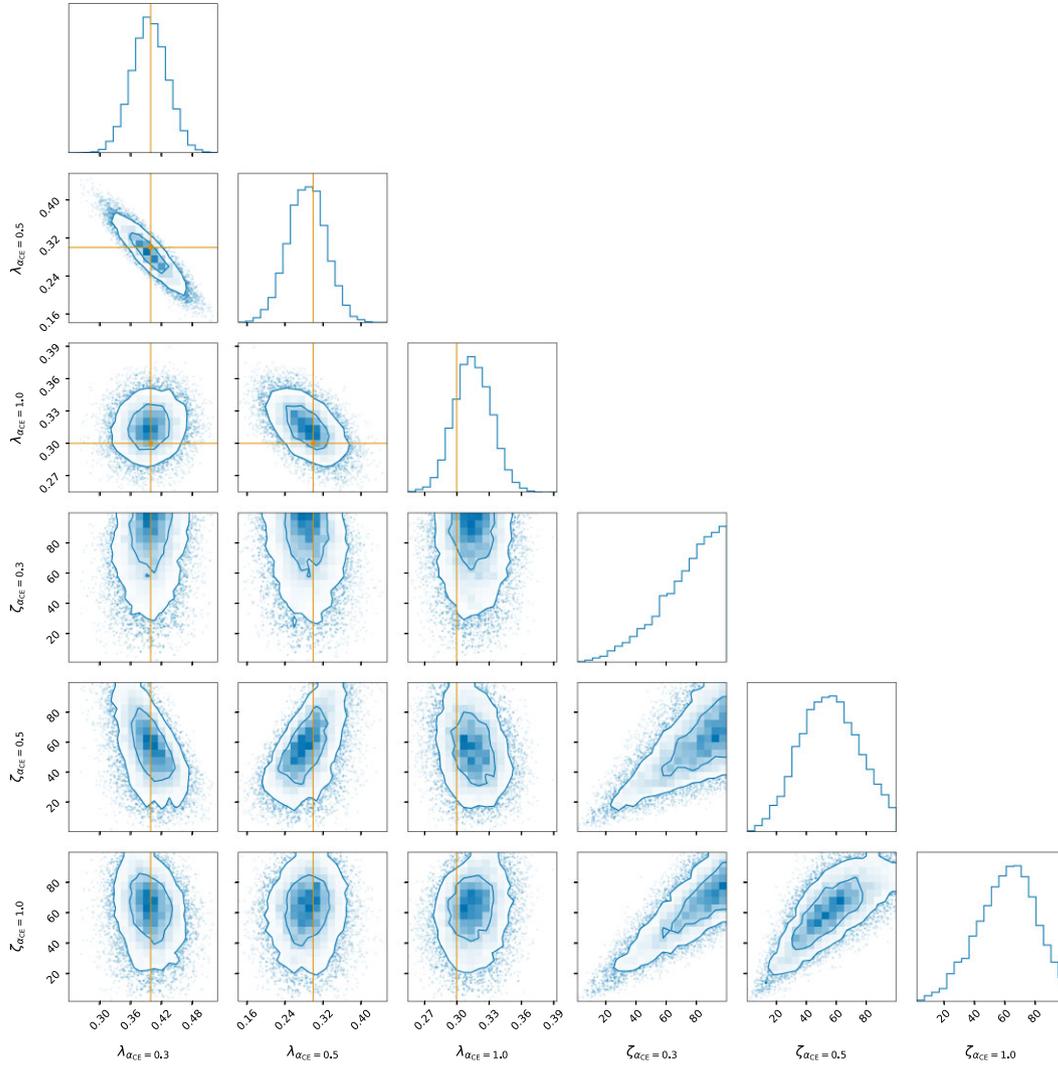


Figure A1. Corner plots of the posterior on the fractional mixture coefficients and concentration parameters of the Dirichlet distribution obtained for the CE efficiency run and 2048 GW detections. The blue solid lines indicate the injected values for the fractional mixture coefficients.

are anticorrelated as they cannot both be present with high fractions, otherwise, they would overestimate the overall merger rate. On the other hand, one can see that the coefficient corresponding to $\alpha_{\text{CE}} = 0.3$ does not show any significant correlation with the others. This is due to the fact that the CE population has a negligible BBH merger rate if compared to the other two. The concentration parameter $\{\zeta\}$ acts as a ‘nuisance’ parameter for determining the prior weights on

the mixture coefficients. It is interesting to note however that all the concentration parameters are correlated. This is due to the fact that, given a Dirichlet distribution on λ_j parameters with concentration parameters ζ_j , the expected values of $E[\lambda_j] = \zeta_j / \sum_k \zeta_k$.

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