



Pulsar Timing Array Constraints on the Merger Timescale of Subparsec Supermassive Black Hole Binary Candidates

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Abstract

We estimate the merger timescale of spectroscopically selected, subparsec supermassive black hole binary (SMBHB) candidates by comparing their expected contribution to the gravitational-wave background (GWB) with the sensitivity of current pulsar timing array (PTA) experiments and in particular, with the latest upper limit placed by the North American Nanohertz Observatory for Gravitational Waves. We find that the average timescale to coalescence of such SMBHBs is $\langle t_{\text{evol}} \rangle > 6 \times 10^4$ yr, assuming that their orbital evolution in the PTA frequency band is driven by emission of gravitational waves. If some fraction of SMBHBs do not reside in spectroscopically detected active galaxies, and their incidence in active and inactive galaxies is similar, then the merger timescale could be ~ 10 times longer, $\langle t_{\text{evol}} \rangle > 6 \times 10^5$ yr. These limits are consistent with the range of timescales predicted by theoretical models and imply that all the SMBHB candidates in our spectroscopic sample could be binaries without violating the observational constraints on the GWB. This result illustrates the power of the multimessenger approach, facilitated by the PTAs, in providing an independent statistical test of the nature of SMBHB candidates discovered in electromagnetic searches.

Unified Astronomy Thesaurus concepts: Active galactic nuclei (16); Galaxy mergers (608); Gravitational waves (678); Supermassive black holes (1663)

1. Introduction

Over the past decade spectroscopic searches have identified about 100 supermassive black hole binary (SMBHB) candidates at subparsec orbital separations (Bon et al. 2012, 2016; Eracleous et al. 2012; Decarli et al. 2013; Ju et al. 2013; Shen et al. 2013; Liu et al. 2014; Runnoe et al. 2015, 2017; Li et al. 2016; Wang et al. 2017; Guo et al. 2019). These searches rely on detection and long-term monitoring of the Doppler shift in the optical emission-line spectrum of active galactic nuclei (AGNs), which arise as a consequence of SMBHB orbital motion, under the assumption that at least one of its constituent supermassive black holes (SMBHs) can shine as an AGN (Begelman et al. 1980; Gaskell 1983, 1996).

With a cadence of observations anywhere from days to years, spectroscopic searches are in principle sensitive to binaries with orbital periods in the range ~ 10 – 100 s yr and separations of at most $\text{few} \times 10^4 r_g$ ($r_g = GM/c^2$ and M is the binary mass; Pflueger et al. 2018). For each observed SMBHB with mass $10^8 M_\odot$, a comparable mass ratio, and orbital separation of about $10^4 r_g$, the projection factors (i.e., orientation of the binary orbit relative to the observer’s line of sight) imply a few undetected binaries, and possibly more if some fraction of SMBHBs do not exhibit AGN signatures. Furthermore, for every SMBHB in the “detectable” range, there should be over 200 more gravitationally bound systems with similar properties but at larger separations, where they cannot be detected by optical spectroscopic searches (Pflueger et al. 2018). Thus, any SMBHB detected using this technique would represent the tip of the iceberg of binaries that escape detection because they are either (a) underluminous, (b) have unfavorable orientation, (c) have orbital velocities that are too low, or (d)

reside in a portion of the sky not covered by the search (see Kelley 2020 for a systematic study of these effects).

The main complication of spectroscopic searches is the fact that the velocity shift and modulation of emission lines around their rest-frame wavelength is not unique to SMBHBs (e.g., Eracleous et al. 2012; Popović 2012; Barth et al. 2015; Guo et al. 2019), making it difficult to uniquely identify binaries. This is of importance because if any of detected SMBHB candidates are real binaries, they are direct progenitors of systems that coalesce due to the emission of gravitational waves (GWs). More specifically, they imply some number of SMBHBs inspiraling toward coalescence, whose GW signal is reaching Earth at this very moment. If there were many of them, the stochastic superposition of their GWs would have already been detected by the pulsar timing arrays (PTAs).

PTAs seek to detect GWs by searching for correlations in the timing observations of a network of millisecond pulsars. Currently, there are three such experiments in operation: the North American Observatory for Gravitational Waves (NANOGrav; McLaughlin 2013), the European PTA (EPTA; Desvignes et al. 2016), and the Parkes PTA (PPTA; Hobbs 2013). Together they form the International PTA (IPTA; Verbiest et al. 2016). At this time, PTA searches for an isotropic stochastic GW background (GWB) are starting to reach sensitivities necessary to probe backgrounds of astrophysical origin (Lentati et al. 2015; Shannon et al. 2015; Arzoumanian et al. 2018).

The massive ($M > 10^8 M_\odot$) and nearby ($z \approx 1$ – 2) SMBHBs are the major contributors to low-frequency GWs sought by PTAs (Sesana et al. 2008). Although current limits are still insufficient to place stringent constraints on the cosmic

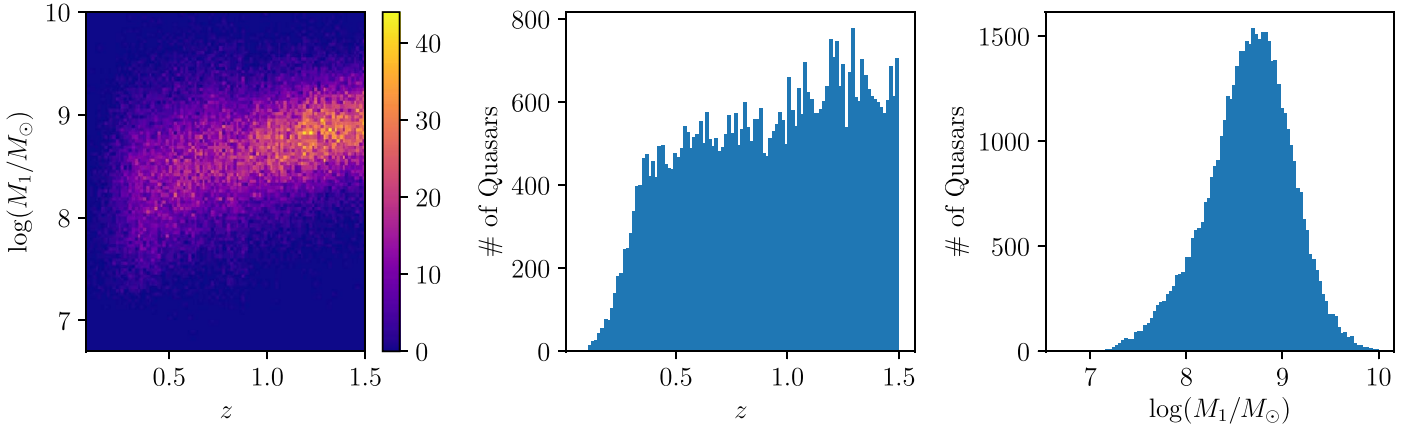


Figure 1. Distribution of SMBHBs in mass and redshift inferred from the SDSS DR7 quasar catalog (left panel) and histograms of the distribution in redshift (middle) and mass (right). The one-dimensional distributions are the projections of the two-dimensional distribution on the mass and redshift axes. We assume that the mass distribution of primary SMBHBs in hypothesized binaries has the same shape as for the SMBHBs in SDSS quasars. The color bar marks the number of quasars.

population of SMBHBs (Middleton et al. 2018), they can be used to test candidates assembled from electromagnetic observations. For example, Sesana et al. (2018) found that the GWB implied by a sample of ~ 150 photometrically selected SMBHB candidates (based on potential periodicity in their light curves) is in tension with the current most stringent PTA upper limits, implying that at least some fraction are false positives. A similar technique was used to place limits on the presence of SMBHBs in periodic blazars (Holgado et al. 2018) and in ultraluminous infrared galaxies (Inayoshi et al. 2018).

In this work, we use a spectroscopic sample of SMBHB candidates from Eracleous et al. (2012, hereafter E12), who searched for $z < 0.7$ Sloan Digital Sky Survey quasars (DR7; Schneider et al. 2010), with broad $H\beta$ lines offset from the rest frame of the host galaxy by $\gtrsim \text{few} \times 100 \text{ km s}^{-1}$. Based on this criterion, E12 selected 88 SMBHB candidates for observational follow-up from an initial group of about 15,900 objects. From this sample of candidates we infer the underlying population of binaries that are inspiraling due to the emission of GWs. Instead of taking a forward-modeling approach to this problem common in the literature, in which we would adopt a particular model for orbital evolution from the subparsec scales to the GW band, we ask: if the GWB signal of the sample of hypothetical SMBHBs that we consider is to be smaller than the sensitivity limit of the PTAs, what is the lower limit on their evolution timescale?

2. Methods

2.1. Merger Rate of SMBHBs

In order to determine the GWB contributed by a population of SMBHBs, we calculate their differential merger rate,

$$\frac{d^5 N}{dM_1 d\tilde{a} dq dz dt_r} = \frac{\nu(M_1; z)}{t_{\text{evol}}(M_1, \tilde{a}, q)} \frac{\rho(\tilde{a}, q, z)}{P_{\text{bias}}}, \quad (1)$$

where $M = M_1 + M_2$ is the binary mass, $q = M_2/M_1 < 1$ is the mass ratio with M_1 (M_2) being the mass of the primary (secondary) SMBH, $\tilde{a} \equiv a/r_g$ is the dimensionless semimajor axis, z is redshift, and t_r is time measured in the rest frame of the SMBHB.

The quantities on the right-hand side of Equation (1) represent the distribution of SMBHB properties inferred from the E12 sample of candidates by correcting for selection effects. Here, $\nu(M_1; z)$ is the mass distribution as a function of z of spectroscopically detectable SMBHBs (see Section 2.2). The

parameter $t_{\text{evol}}(M_1, \tilde{a}, q)$ is the timescale for evolution of a SMBHB from a separation at which it was detected ($\sim 10^4 r_g$ for spectroscopically targeted binaries) to coalescence. It is usually estimated from the merger rate as $dN/dt_r \approx N/t_{\text{evol}}$ and it depends on the SMBHB parameters, as well as the physical mechanisms that drive binary to coalescence (gas, stellar torques, and GW emission; Sesana 2013). The function $\rho(\tilde{a}, q, z)$ is the probability distribution of SMBHB candidates given \tilde{a} , q , and z , introduced in Section 2.2. P_{bias} is a probability that an SMBHB is detected by the E12 spectroscopic search given the selection effects inherent to this technique (see Section 2.3).

2.2. Distribution of SMBHBs— $\nu(M_1; z)$ and $\rho(\tilde{a}, q, z)$

We derive the mass distribution of primary SMBHBs in all hypothesized binaries within the redshift range $0 < z < 1.5^5$ by assuming that it has the same shape as the mass distribution of SMBHBs powering SDSS quasars but a different normalization, since only a small fraction of quasars may host binaries. This is reasonable since we expect that the primary SMBHBs in the E12 sample would have formed in the same way as the rest of the SDSS quasars powered by isolated SMBHBs: through prior mergers and accretion. Thus, we describe the primary SMBHBs using the mass distribution of the quasars from the SDSS DR7 catalog. These masses are obtained using the virial SMBH mass estimators, based on the continuum luminosities and the $H\beta$ or Mg II lines (Shen et al. 2011). We use measurements for which the observed line profiles are fit with reduced chi-squared between about 0.8 and 1.5, ensuring a reliable fit, and eliminate quasars with broad absorption lines, which may have inaccurate mass estimates.

The left panel of Figure 1 shows the resulting SMBH mass distribution for the SDSS quasars. This is a distribution whose normalization evolves with redshift (middle panel), with a majority of SMBH masses in the range $10^7\text{--}10^8 M_\odot$ and a median of $\sim 5 \times 10^8 M_\odot$ (right panel). It is worth noting that because SDSS is a flux-limited survey, at every redshift there are active galaxies that are below its detection threshold. This is reflected in a dearth of SMBHBs with masses $\lesssim 10^8 M_\odot$ beyond $z \approx 0.5$ in the left panel of Figure 1. This is of interest because if some

⁵ This expression implies an upper limit in redshift that encloses most of the GWB from SMBHBs detected by PTAs. We justify this assumption in Section 3.

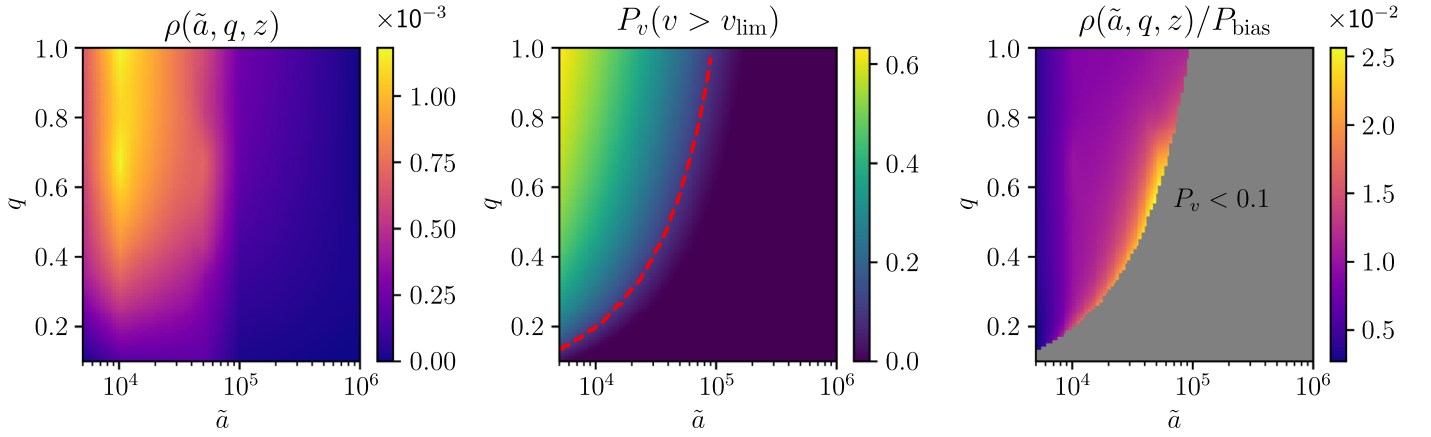


Figure 2. Left panel: probability density distribution of the SMBHB candidates from the E12 sample, $\rho(\tilde{a}, q, z)$, integrated over redshift. Middle panel: probability of detecting a SMBHB with the radial component of orbital velocity greater than $v_{\text{lim}} = 350 \text{ km s}^{-1}$. The red dashed line marks the $P_v = 0.1$ contour. Right panel: the probability density distribution of the inferred SMBHBs population after accounting for the selection effects, $\rho(\tilde{a}, q, z)/P_{\text{bias}}$. The probability density in the grayed-out region is set to zero (see Section 2.3).

fraction of these objects are tracers of inspiraling SMBHBs, they represent a contribution to the GWB that is unaccounted for. We examine the impact of this selection effect on the resulting GWB in Section 3.

The virial SMBH mass measurements, like the ones obtained from the SDSS DR7 catalog, are known to be subject to Malmquist bias (Shen et al. 2008). This effect arises because the underlying SMBH mass distribution in the mass range of interest is bottom heavy (i.e., there are more SMBHBs toward lower masses), and as a result more objects scatter from the low-mass bins to high than the other way around. Thus, the observed virial mass distribution for the SDSS sample is biased high by about 0.55 dex relative to the “true” underlying distribution. We evaluate the impact of this effect by performing calculations of the merger rate with $\nu(M_1; z)$ (a) uncorrected for Malmquist bias, as shown in Figure 1, and (b) corrected for this bias by shifting the distribution to lower masses by 0.55 dex. The median SMBH mass of the corrected distribution is then about $10^8 M_\odot$.

Finally, we obtain the normalization of the SMBHB mass distribution in either scenario by scaling down the SMBH mass distribution function in Figure 1 in such a way, that in the redshift range $0 < z < 0.7$ the number of objects corresponds to 88, the number of SMBHB candidates in the E12 sample. The resulting number of SMBHBs out to $z = 1.5$ inferred in this way is 285 (see, however, the discussion of selection effects in Section 2.3).

In order to aid the interpretation of spectroscopic SMBHB candidates, Nguyen & Bogdanović (2016) and Nguyen et al. (2019) developed a semianalytic model to calculate the broad emission-line profiles emitted from circumbinary accretion flows associated with subparsec SMBHBs. They found that the modeled profiles show distinct statistical properties as a function of the binary semimajor axis and mass ratio and that, as a result, broad emission lines can be used to infer their distribution. A subsequent analysis presented in Nguyen et al. (2020) showed that as a population, the E12 SMBHB candidates favor an average value of the semimajor axis corresponding to $\log \tilde{a} \approx 4.20$ with a standard deviation of 0.42, and comparable mass ratios, $q > 0.5$.

The left panel of Figure 2 shows the resulting probability density distribution for the E12 sample of SMBHB candidates from Nguyen et al. (2020), $\rho(\tilde{a}, q, z)$, integrated over redshift.

The distribution shown in the figure is normalized in such a way that when integrated with respect to \tilde{a} , q , and z returns 88, the total number of the E12 SMBHB candidates. In the absence of other information about the properties of the SMBHB candidates with redshift $z \geq 0.7$, we assume that they are characterized by the same distribution, $\rho(\tilde{a}, q, z)$, as the E12 sample.

One can show that each hypothetical SMBHB, characterized by the distribution of M_1 , \tilde{a} , and q described here, is more likely to have an evolution timescale longer than a Hubble time, if its evolution was driven solely by the emission of GWs. Therefore, these SMBHBs can evolve into the PTA frequency band only if their evolution at larger separations is driven by gas and/or stars.

2.3. Probability of Detection— P_{bias}

If all objects in the E12 sample are true binaries, one would expect an underlying population larger than 88, given the selection effects of the search. We account for two such effects: one is a probability of detection given a partial sky coverage of the SDSS DR7 spectroscopic survey, which corresponds to $P_{\text{sdss}} \approx 1/4$. The other is a probability, P_v , that an SMBHB has the radial component of orbital velocity greater than some threshold value that defines the sensitivity of the search, $v > v_{\text{lim}}$. Note that the latter probability accounts for the fact that some fraction of SMBHBs escape detection either because they have unfavorable orientation or because their orbital velocity is lower than v_{lim} regardless of orientation, as mentioned in Section 1. The total probability of detection is then $P_{\text{bias}} = P_{\text{sdss}} P_v$. Note that so far we do not account for the fact that some unknown fraction of SMBHBs may reside in systems that do not exhibit AGN signatures (see the discussion in Section 4).

Assuming for simplicity SMBHBs are on circular orbits and that the measured radial velocity traces the motion of the primary SMBH, P_v can be expressed analytically as (Pflueger et al. 2018)

$$P_v(v > v_{\text{lim}}) = 1 - \frac{2}{\pi} \left(\arcsin \zeta + \zeta \ln \left[\frac{1 + \cos(\arcsin \zeta)}{\zeta} \right] \right), \quad (2)$$

where $\zeta = \tilde{a}^{1/2}(1/q + 1)(v_{\text{lim}}/c)$ is a dimensionless parameter. Note that the premise that v is associated with the primary SMBH marks a departure from that commonly adopted by spectroscopic searches, which assume that v is associated with the secondary instead. This is supported by modeling that indicates that in most SMBHB configurations the accretion disk around the primary makes the dominant contribution to the $H\beta$ broad emission-line flux (Nguyen et al. 2020). See Section 3 for a description of how our results are affected by this assumption.

The middle panel of Figure 2 shows P_v calculated for $v_{\text{lim}} = 350 \text{ km s}^{-1}$. This value corresponds to the smallest velocity offset measured in the E12 sample, in the first epoch of observations, and is representative of the sensitivity achieved by the search. Figure 2 illustrates that the probability of detection increases with q , as the orbital speed of the primary SMBH becomes more pronounced. Similarly, P_v decreases with a , as the binary orbital velocity decreases with separation.

To derive the probability density of the underlying SMBHB population, and factor out the selection effects described above, we divide $\rho(\tilde{a}, q, z)$ by P_{bias} and show the result in the right panel of Figure 2. This distribution indicates an increasing number of SMBHBs at larger orbital separations, as expected if wider binaries are evolving more slowly. This approach, however, cannot be used to reliably extrapolate the number of SMBHBs in the region where the sensitivity of the search drops significantly. This region is marked by dark blue colors in the left and middle panels of Figure 2 and is outlined by the red dashed line in the middle panel with a $P_v = 0.1$ contour. In order to mitigate the uncertainty caused by small number statistics we set $\rho(\tilde{a}, q, z) = 0$ where $P_v < 0.1$ and make no predictions for the underlying SMBHB population in the grayed-out area in the right panel of Figure 2. We account for the effect of truncation in $\rho(\tilde{a}, q, z)$ by rescaling its normalization to ensure that when integrated in terms of \tilde{a} , q , and z it still returns 88.

The inferred number of SMBHBs with $z < 0.7$ obtained in this way, calculated by integrating the distribution $\rho(\tilde{a}, q, z)/P_{\text{bias}}$ shown in the right panel, is around 1492, indicating that for every SMBHB detected in this parameter space there are about 16 more that escape detection on average, because of selection effects. Extending the same reasoning to SMBHBs with $z < 1.5$ implies about $285 \times 17 = 4845$ binaries in this redshift range.

It is worth mentioning that an additional selection effect introduced by spectroscopic searches is a probability that an SMBHB has a change in radial velocity, measured as an epoch-to-epoch modulation in the velocity offset of the broad emission lines, larger than some threshold value, $\Delta v > \Delta v_{\text{lim}}$ (see Pflueger et al. 2018). We neglect this effect as it was not used to eliminate any SMBHBs in the E12 sample thus far.

2.4. Calculation of the Gravitational-wave Background

We calculate the GWB strain as a function of the observed frequency, $h_c(f)$, following the approach described in Phinney (2001) and Sesana et al. (2004, 2008):

$$h_c^2(f) = \frac{4G}{\pi c^2 f^2} \int dz \frac{n(z)}{1+z} \frac{dE_{\text{GW}}}{d \ln f_r}, \quad (3)$$

where $f_r = f(1+z)$ is the GW frequency in the rest frame of the binary. E_{GW} is the energy emitted in GW, which for a

circular SMBHB can be expressed as

$$\frac{dE_{\text{GW}}}{d \ln f_r} = \frac{\pi^{2/3}}{3G} (G\mathcal{M})^{5/3} f_r^{2/3}, \quad (4)$$

and $\mathcal{M} = M_1 q^{3/5}/(1+q)^{1/5}$ is the chirp mass. In this calculation, Equation (4) represents SMBHBs emitting in the frequency band of NANOGrav that evolve primarily due to the emission of GWs, as opposed to gas and stellar torques. We discuss the implications of this assumption in Section 4. $n(z)$ represents the number of binary mergers per unit comoving volume per unit redshift, $n(z) = d^2 N/dz dV_c$, and thus

$$n(z) = \iiint dM_1 d\tilde{a} dq \frac{d^5 N}{dz dM_1 d\tilde{a} dq dt_r dV_c}, \quad (5)$$

where the relationship between time and comoving volume is given by $dt_r/dV_c = [4\pi c(1+z)d_M^2(z)]^{-1}$. The comoving distance is given by

$$d_M(z) = \frac{c}{H_0} \int_0^z \frac{dz'}{\sqrt{\Omega_M(1+z')^3 + \Omega_\Lambda}}, \quad (6)$$

where we assume a flat universe with $\Omega_M = 0.315$, $\Omega_\Lambda = 0.685$, $\Omega_k = 0$, and $H_0 = 67.4 \text{ km s}^{-1} \text{ Mpc}^{-1}$ (Planck Collaboration et al. 2018). Combining Equations (3)–(6) with Equation (1) we obtain

$$h_c^2(f) = \frac{G^{5/3}}{3\pi^{4/3}c^3} \frac{1}{f^{4/3}} \iiint dM_1 dz d\tilde{a} dq \times \frac{1}{t_{\text{evol}}(M_1, \tilde{a}, q)} \frac{\nu(M_1; z)}{(1+z)^{4/3}} \frac{\rho(\tilde{a}, q, z)}{P_{\text{bias}}} \frac{\mathcal{M}^{5/3}}{d_M^2(z)}, \quad (7)$$

and subsequently,

$$h_c^2(f) = \frac{G^{5/3}}{3\pi^{4/3}c^3} \frac{1}{f^{4/3}} \frac{1}{\langle t_{\text{evol}} \rangle} \iiint dM_1 dz d\tilde{a} dq \times \frac{\nu(M_1; z)}{(1+z)^{4/3}} \frac{\rho(\tilde{a}, q, z)}{P_{\text{bias}}} \frac{\mathcal{M}^{5/3}}{d_M^2(z)}. \quad (8)$$

Equating Equations (7) and (8) yields a definition of $\langle t_{\text{evol}} \rangle$, a characteristic merger timescale for evolution of the ensemble of SMBHBs in the redshift range $0 < z < 1.5$, from the separations at which the E12 candidates are typically detected ($\sim 10^4 r_g$) to coalescence. $\langle t_{\text{evol}} \rangle$ is calculated as an average over the distributions in M_1 , z , \tilde{a} , and q , and weighted by the factors in Equation (3), of which $dE_{\text{GW}}/d \ln f_r$ puts weight on the loudest binaries in a given frequency interval.

At such large initial separations the evolution of SMBHBs headed for coalescence is driven by stellar and gas torques. This allows us to decouple $\langle t_{\text{evol}} \rangle$ from the calculation of the GW signal of such SMBHBs in the NANOGrav band, where we assume that GW emission dominates their evolution (Equation (4)). Hence, in Equation (8) $\langle t_{\text{evol}} \rangle$ appears as a parameter in front of the integral. At $z < 0.7$ the integral turns into a summation over 88 objects with individual redshifts and mass distribution described in Section 2.2. At $z \geq 0.7$ we integrate over the mass and redshift distribution of the SDSS quasars shown in Figure 1, and normalize it relative to the number of SMBHB candidates at $z < 0.7$.

Table 1
GWB Strain at $f = 1 \text{ yr}^{-1}$

$\langle t_{\text{evol}} \rangle / \text{yr}$	h_{c1}	h_{c2}	h_{c3}
10^9	2.80×10^{-17}	3.24×10^{-17}	1.13×10^{-17}
10^8	8.85×10^{-17}	1.02×10^{-16}	3.57×10^{-17}
10^7	2.80×10^{-16}	3.24×10^{-16}	1.13×10^{-16}
10^6	8.85×10^{-16}	1.02×10^{-15}	3.57×10^{-16}
10^5	2.80×10^{-15}	3.24×10^{-15}	1.13×10^{-15}
10^4	8.85×10^{-15}	1.02×10^{-14}	3.57×10^{-15}

Note. $\langle t_{\text{evol}} \rangle$ —merger timescale. h_{c1} , h_{c2} —GWB strain amplitudes for SMBHBs at $z < 0.7$ and $z < 1.5$, respectively, uncorrected for Malmquist bias. h_{c3} —GWB strain amplitude for SMBHBs at $z < 1.5$, corrected for Malmquist bias. See Section 3 for more detail.

3. NANOGrav Constraints on the Merger Timescale

We use Equation (8) to calculate the GWB strain from the population of putative SMBHBs inferred from the E12 sample given a merger timescale, $\langle t_{\text{evol}} \rangle$. Specifically, we calculate the strain at a reference frequency $f = 1 \text{ yr}^{-1}$ and summarize the results in Table 1. The first column of the table shows the value of $\langle t_{\text{evol}} \rangle$ and the second shows the corresponding GWB strain, h_{c1} , calculated for a population of SMBHBs in the redshift range $0 < z < 0.7$, equivalent to that of the E12 sample. In this scenario the mass distribution of SMBHBs was not corrected for Malmquist bias. Note that for all values $h_c^2 \propto 1/\langle t_{\text{evol}} \rangle$, so the table illustrates how different evolution times of SMBHBs affect the resulting amplitude of the GWB strain. Namely, longer $\langle t_{\text{evol}} \rangle$ implies slower inspiral of binaries from subparsec scales to coalescence and, consequently, lower GWB.

The third column of Table 1 shows the strain amplitude, h_{c2} , calculated for a population of SMBHBs in the full redshift range, $0 < z < 1.5$, with masses uncorrected for Malmquist bias. Comparison of models h_{c1} and h_{c2} shows that when the contribution to the GWB from binaries with $z \geq 0.7$ is included, the overall strain amplitude increases by about 16%. Low-redshift SMBHBs therefore dominate the stochastic GWB at $f = 1 \text{ yr}^{-1}$ by a large margin. Hence, even if there is a population of low-luminosity or higher-redshift SMBHBs not captured by the flux-limited SDSS spectroscopic survey, their contribution to the GWB should be small.

The fourth column shows h_{c3} , calculated for a population of SMBHBs with $0 < z < 1.5$ with masses corrected for Malmquist bias. Because the corrected mass distribution is characterized by a lower median value, in this case the overall GWB amplitude decreases by a factor of approximately 3 relative to h_{c2} .

In the next step, we compare the calculated strain amplitudes in Table 1 to the latest constraints provided by the 11 yr NANOGrav data set, which sets a 95% upper limit on the GW strain amplitude of $A_{\text{GWB}} < 1.45 \times 10^{-15}$ for SMBHBs emitting at a frequency of 1 yr^{-1} (Arzoumanian et al. 2018). Although this limit is a factor ~ 1.5 less stringent than that published by Shannon et al. (2015), it includes a self-consistent Bayesian model of the solar system ephemeris, making it more robust.

The top left panel of Figure 3 shows the probability density function (PDF), corresponding to the model corrected for Malmquist bias (h_{c3}), which specifies the probability that A_{GWB} falls within a particular range of values. It is commonly

modeled by a Fermi-like function (e.g., Chen et al. 2017),

$$\text{PDF}(h_c) = \frac{C_1}{1 + \exp\left(\frac{h_c - A_{95}}{C_2}\right)}, \quad (9)$$

where $C_1 = 6.90 \times 10^{14}$ and $C_2 = 1.05 \times 10^{-16}$ are constants determined from PDF normalization and a requirement that the 95th percentile value of the strain amplitude is $A_{95} = 1.45 \times 10^{-15}$, respectively. The bottom left panel of Figure 3 shows the resulting cumulative distribution function (CDF), which indicates the probability that A_{GWB} is less than or equal to a given strain amplitude shown on the x -axis. The vertical line marks A_{95} , which corresponds to the sensitivity limit of NANOGrav at $f = 1 \text{ yr}^{-1}$ (Arzoumanian et al. 2018).

The top right panel of Figure 3 shows a PDF for model h_{c3} , of the merger time corresponding to a given value of h_c , such that $\text{PDF}(\langle t_{\text{evol}} \rangle) = \text{PDF}(h_c) |dh_c/d\langle t_{\text{evol}} \rangle|$. The inferred distribution for $\langle t_{\text{evol}} \rangle$ peaks at about 10^5 yr and indicates that there cannot be many subparsec binaries that evolve to merger on timescales $\ll 10^5 \text{ yr}$, since they would produce strain amplitudes $h_c \gg A_{95}$, and would already be detected by NANOGrav. Similarly, low strain amplitudes ($h_c \ll A_{95}$) can be produced by a small number of relatively slowly evolving binaries with $\langle t_{\text{evol}} \rangle > 10^7 \text{ yr}$, illustrated by the extended tail of the distribution.

Similarly to $\text{PDF}(h_c)$, which provides an upper limit on the GWB strain created by inspiraling SMBHBs, $\text{PDF}(\langle t_{\text{evol}} \rangle)$ can be used to infer a lower limit on $\langle t_{\text{evol}} \rangle$ for the same population of binaries. The lower right panel of Figure 3 shows the CDF for $\langle t_{\text{evol}} \rangle$ for model h_{c3} and indicates that 95% of the SMBHBs would have to evolve on timescales $\langle t_{\text{evol}} \rangle > 6 \times 10^4 \text{ yr}$ in order to be consistent with the sensitivity limit of NANOGrav. In comparison, the model where the SMBH mass distribution was not corrected for Malmquist bias (h_{c2}) predicts the peak of the distribution at about $8 \times 10^5 \text{ yr}$ and $\langle t_{\text{evol}} \rangle > 5 \times 10^5 \text{ yr}$ for 95% of the SMBHBs.

4. Discussion and Conclusions

In this Letter we consider the contribution to the strain of a stochastic GWB from an expected population of inspiraling SMBHBs with redshift $z < 1.5$, inferred from a sample of 88 subparsec SMBHB candidates discovered by the E12 spectroscopic search. We find that the average timescale for evolution of such SMBHBs from subparsec separations to coalescence must be $\langle t_{\text{evol}} \rangle > 6 \times 10^4 \text{ yr}$ in order for the amplitude of their GWB to be consistent with the upper limit placed by NANOGrav. This limit is in agreement with a range of timescales ($\sim 10^6$ – 10^9 yr) predicted by theoretical models for SMBHBs of similar properties, that evolve due to interactions with stars and/or gas in their host galaxies, and eventually merge due to the emission of GWs (e.g., Haiman et al. 2009; Lodato et al. 2009; Rafikov 2013). This implies that, based on this test alone and within the uncertainties of theoretical models, all 88 SMBHB candidates from the E12 sample are presently consistent with being true binaries. It is of course plausible that only a fraction (or none) of the E12 candidates are actual SMBHBs—if so, $\langle t_{\text{evol}} \rangle$ would be reduced proportionally. Our results are subject to several assumptions that we discuss below.

1. In this work we consider a population of hypothetical subparsec SMBHBs that appear as luminous SDSS

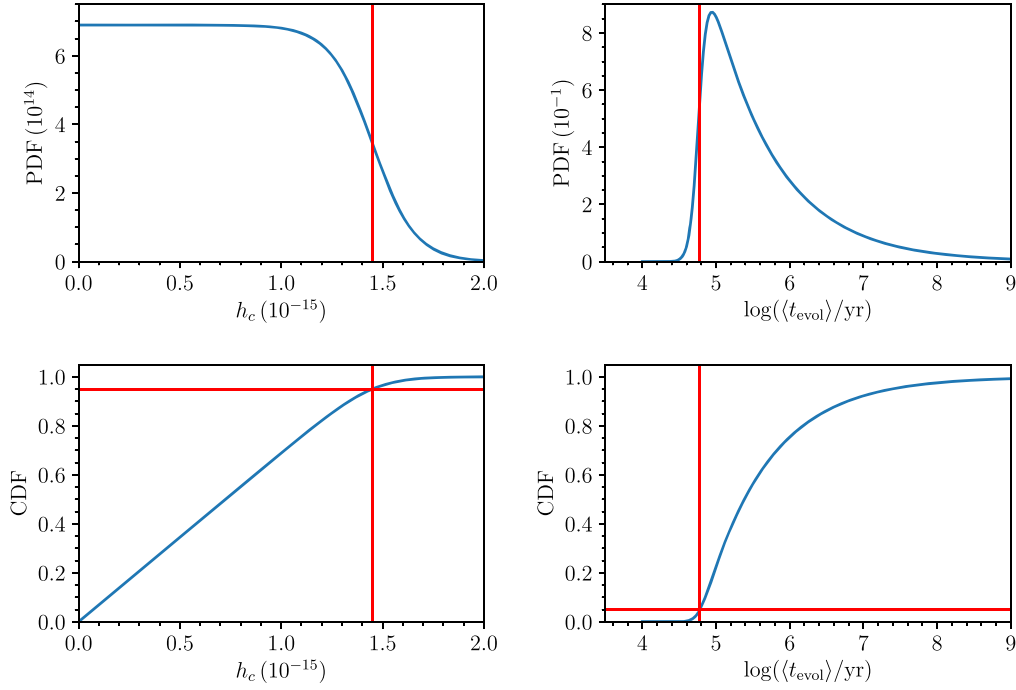


Figure 3. Top row: PDFs for the GW strain amplitude, contributed by a population of inspiraling SMBHBs at a frequency $f = 1 \text{ yr}^{-1}$ (h_c ; left) and the average merger time for the same population ($\langle t_{\text{evol}} \rangle$; right). Both refer to the model h_{c3} , in which SMBHB masses are corrected for Malmquist bias. Bottom row: CDFs corresponding to the PDFs in the top row. Red lines mark the 95th and 5th percentile values of h_c and $\langle t_{\text{evol}} \rangle$, respectively.

quasars but do not account for the presence of SMBHBs in inactive galaxies. If SMBHBs in inactive galaxies are common, they could contribute to the stochastic GWB even if they are not found by the electromagnetic searches. For example, if the frequency of SMBHBs in inactive galaxies is similar to that in AGNs, then the underlying population of binaries could be ~ 10 times larger than the number inferred from the EM searches. If so, this would imply ~ 10 times longer merger timescale, $\langle t_{\text{evol}} \rangle > 6 \times 10^5 \text{ yr}$.

2. We assume that the mass distribution of primary SMBHBs in binaries that contribute to the GWB is the same as that of the SMBHBs that power SDSS quasars. This approach allows us to sidestep complications related to single-epoch virial mass measurements in potential SMBHBs, as those methods may not be applicable to binaries. Even so, the distribution of virial SMBH masses adopted in this work is subject to Malmquist bias, which shifts the distribution of measured masses to higher values by a factor of about 3 relative to the true underlying distribution. We find that if correction for this effect is omitted, the resulting merger timescale is ~ 8 times longer ($\langle t_{\text{evol}} \rangle > 5 \times 10^5 \text{ yr}$) than that for the scenario where this correction is applied. This example illustrates that somewhat different assumptions about the SMBHB mass function can lead to uncertainties of one order of magnitude in the limit on $\langle t_{\text{evol}} \rangle$, and still be consistent with the upper limit on stochastic GWB.
3. Another assumption we adopt is that the radial velocities measured in spectroscopic searches for SMBHBs trace the motion of the primary SMBHBs. If instead the motion traced is that of the secondary, it is more natural to assume that the mass distribution of the secondary (as opposed to the primary) SMBHBs is represented by that of

the SDSS quasars. Because binaries with higher mass ratios are favored, the total mass of such systems would be similar (within a factor of 2) to the case when the primary's motion is traced. This results in $\langle t_{\text{evol}} \rangle$ that is also within a factor of 2 of the value calculated for that scenario. Thus, we do not expect our results to be very sensitive to the assumption that spectroscopic searches trace the motion of the primary SMBHBs.

4. An important assumption of this work is that SMBHBs that contribute to the GWB in the frequency band of NANOGrav inspiral only due to the emission of GWs. For example, this implies that the evolution of $\sim 10^8 M_\odot$ SMBHBs with comparable mass ratios is dominated by GW emission when they reach separations of $\sim \text{few} \times 10^{-3} \text{ pc}$. While this assumption is justified for some binaries, the possibility that the evolution of SMBHBs at these separations is driven by gas or stellar torques cannot be eliminated for all. If so, such SMBHBs would evolve faster through the PTA frequency band, emitting with a lower strain amplitude relative to the scenario in which GW emission dominates. Therefore, the presence of additional physical mechanisms results in a lower, more conservative lower limit $\langle t_{\text{evol}} \rangle$ than that based on the GW emission alone. Along similar lines, gas and stellar torques can in principle excite eccentricity of the SMBHB orbits, in which case our assumption of circular binaries would need to be revised.

In summary, this work illustrates an important place occupied by PTAs and observatories that can provide independent tests of the nature of SMBHBs. While subparsec SMBHBs are still challenging to unambiguously identify, constraints like the one presented here keep narrowing down the range of possibilities for these objects.

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