

S1. Methods and results of maximum parsimony (MP) and maximum likelihood (ML) analyses

S1.1. Methods

We used TNT [1] for maximum parsimony (MP) analyses. Two independent MP analyses were made:

1) the first inference was performed using a heuristic search with 1,000 replications (saving at most 100 trees in each replication) of random (stepwise) addition of taxa followed by Tree Bisection Reconnection (TBR) branch swapping. Two additional runs using trees on memory were performed to refine searches. To evaluate branch support we made 10,000 replicates of standard bootstrap.

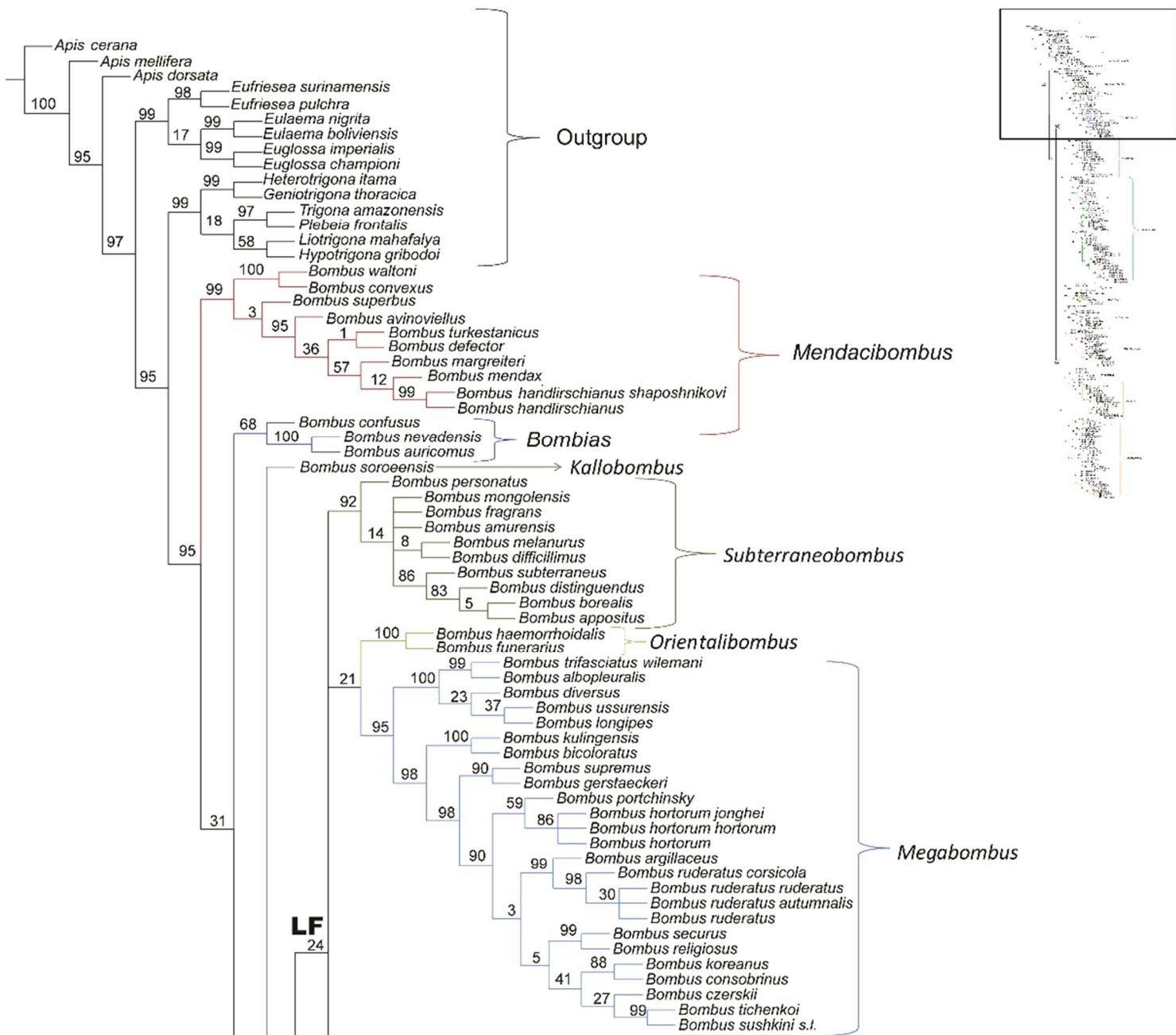
2) the second inference was performed using new technology search (NTS) with sectorial search, ratchet, tree-drifting and tree fusing options and 10 random addition sequences (saving at most 100 trees in each replication). Trees resulting from such analyses were submitted to an additional round of TBR. To evaluate branch support in the resulting trees, we made 10,000 replicates of standard bootstrap.

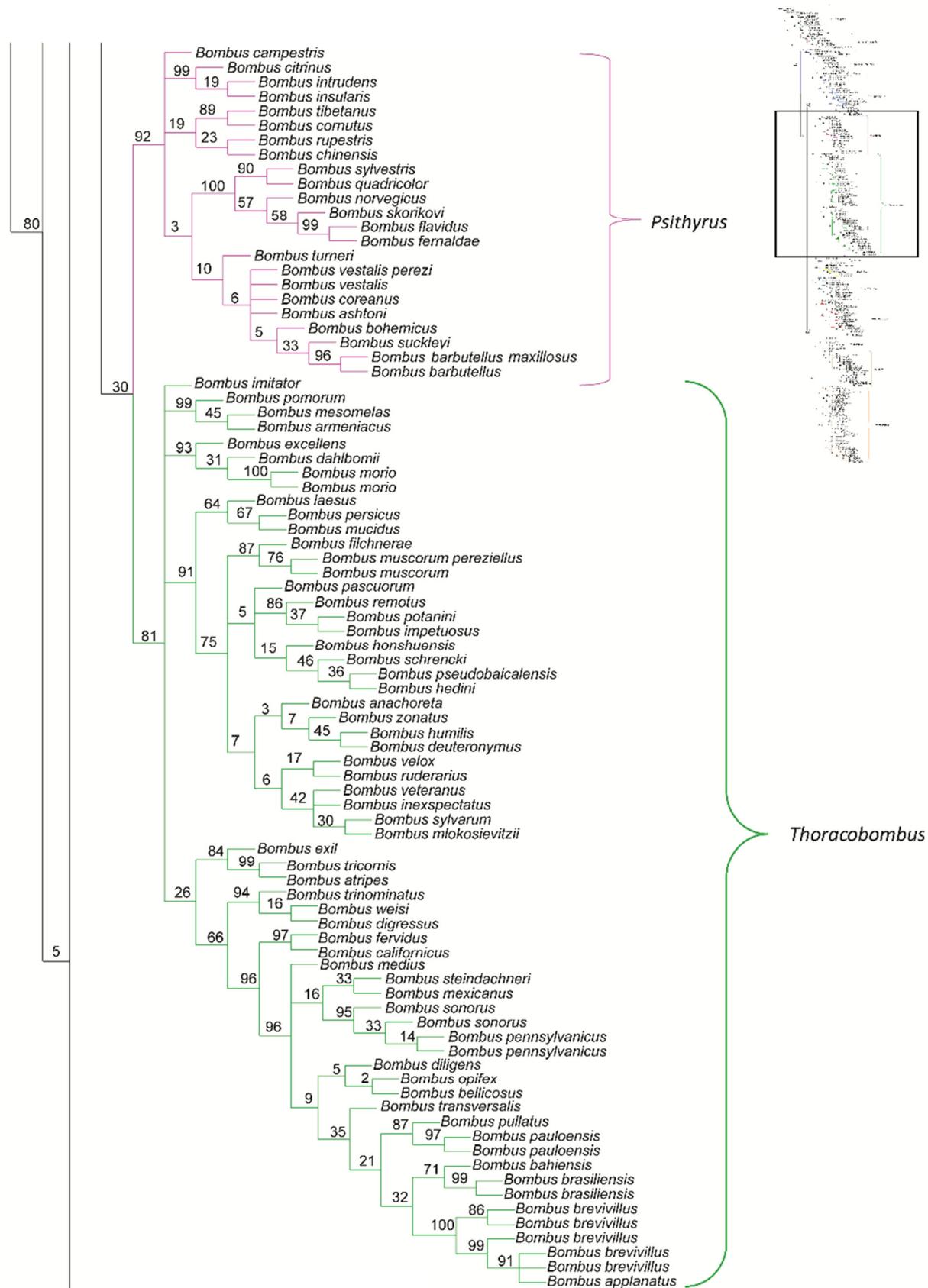
A maximum likelihood analysis (ML) was performed in RAxML version 8 [2]. The model GTR+G was used for all partitions obtained with PartitionFinder v.2.1.1 [3] and 1,000 bootstrap replicates search were used to calculate branch support (ML optimizations were performed every five bootstrap replicates and were followed by a final search for the best-scoring ML tree once bootstrap search was over). We also used the program IQ-TREE [4] for an alternative ML analysis, which differs from RAxML by using flexible models' definitions and a fast bootstrap approximation with algorithm UFBoot2 [5].

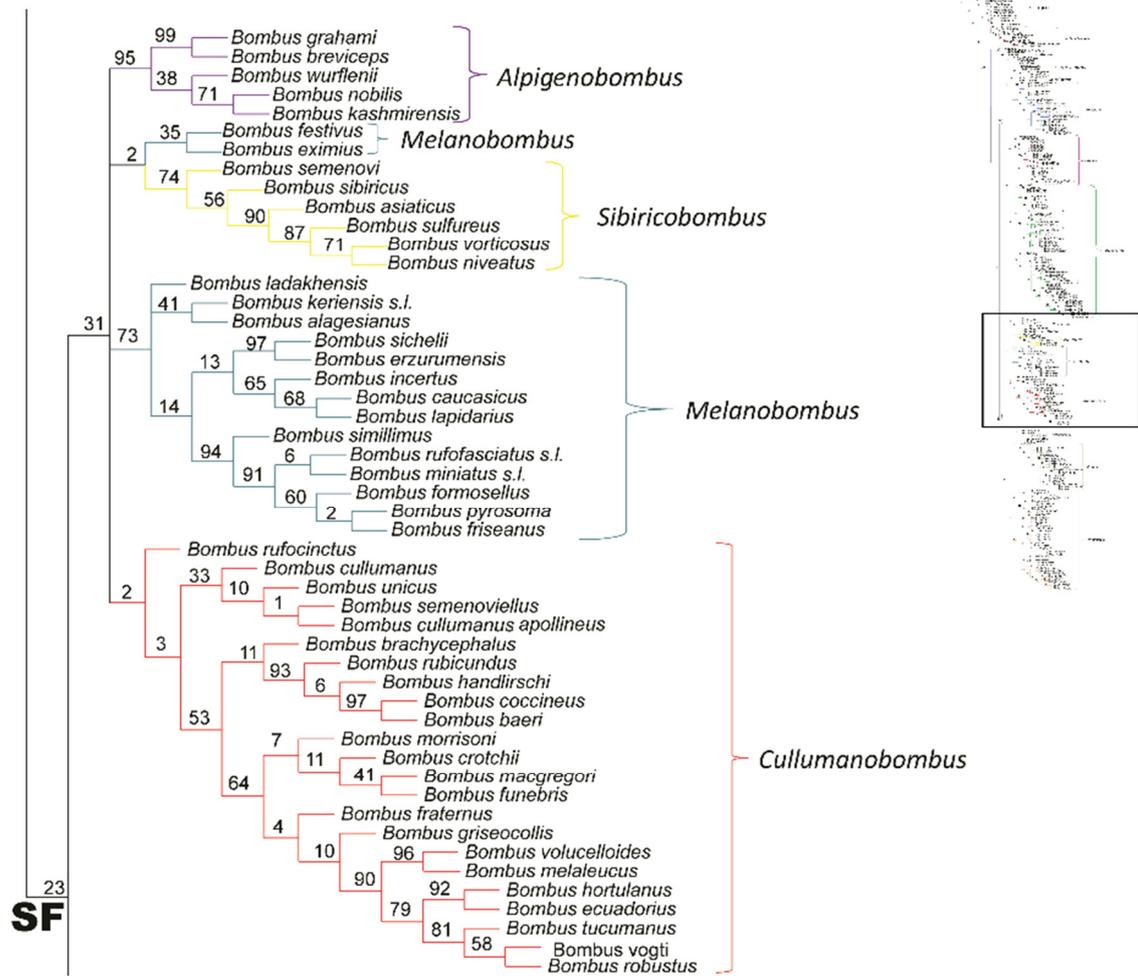
S1.2. Results

The best scores of MP were 13,176 and 14,493 for the analyses without NTS and with NTS, respectively (total rearrangements examined 21,224,652,872 and 21,145,970,952; Figures S7 and S8). The subgenera *Alpigenobombus*, *Melanobombus*, *Sibiricobombus* and *Cullumanobombus* arised in a polytomy, where *B. (Melanobombus) festivus* and *B. (Melanobombus) eximius* grouped with *Sibiricobombus*, making *Melanobombus* paraphyletic. In the MP analyses *Thoracobombus* is paraphyletic, *B. laesus* grouped with *Psithirus* (Figure S7). The ML tree obtained with RaxML recovered *B. rufocinctus* as part of a polytomy, leaving the subgenera *Cullumanobombus* paraphyletic. The polytomy was composed of *Cullumanobombus* + *Sibiricobombus* + *B. (Cullumanobombus) rufocinctus* (Figure S9).

All subgenera were monophyletic in ML analyses obtained with IQ-TREE, although some of them were supported by low bootstrap values. The relationships among species within some subgenera were not well resolved or were weakly supported (Figure S10).







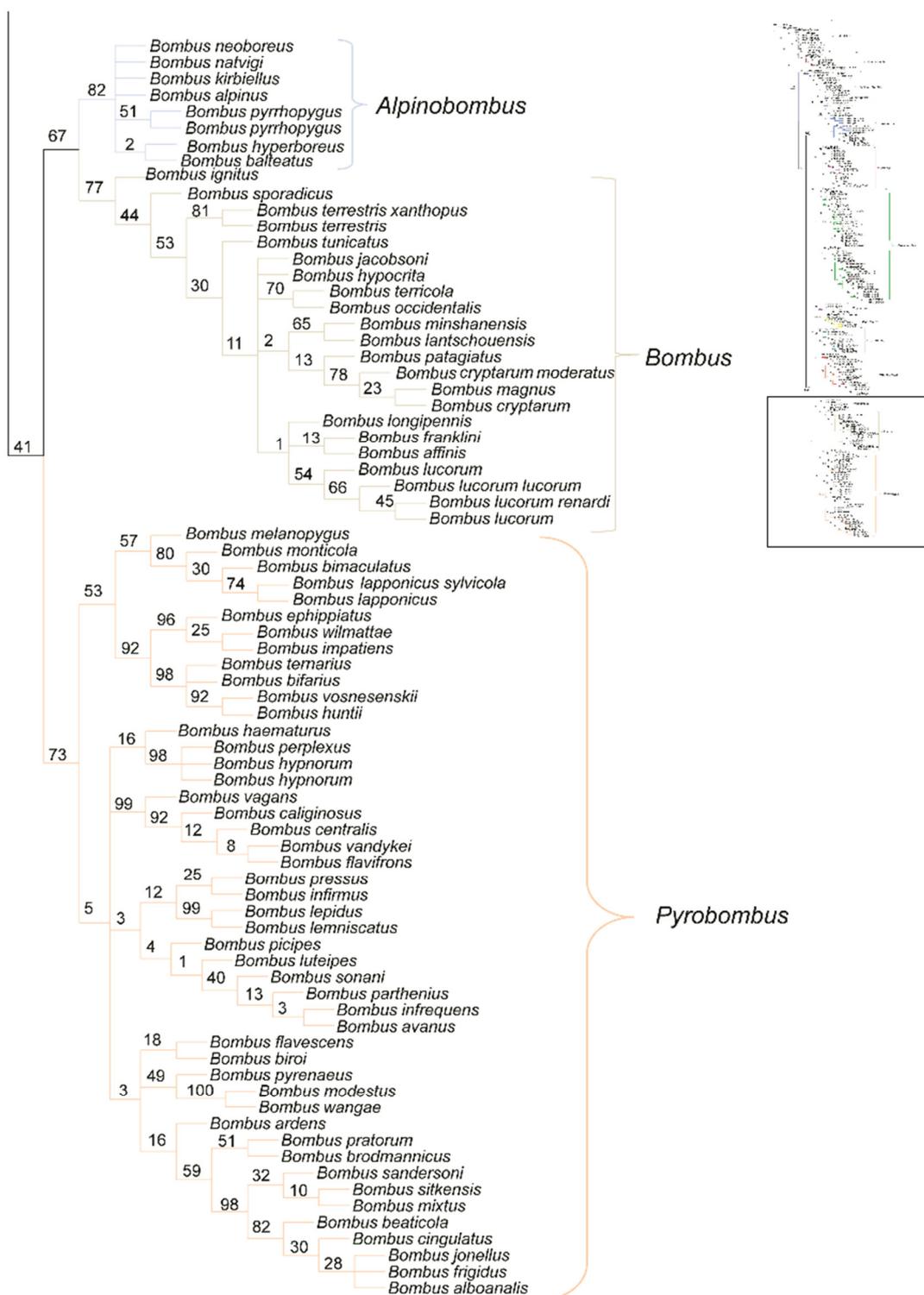
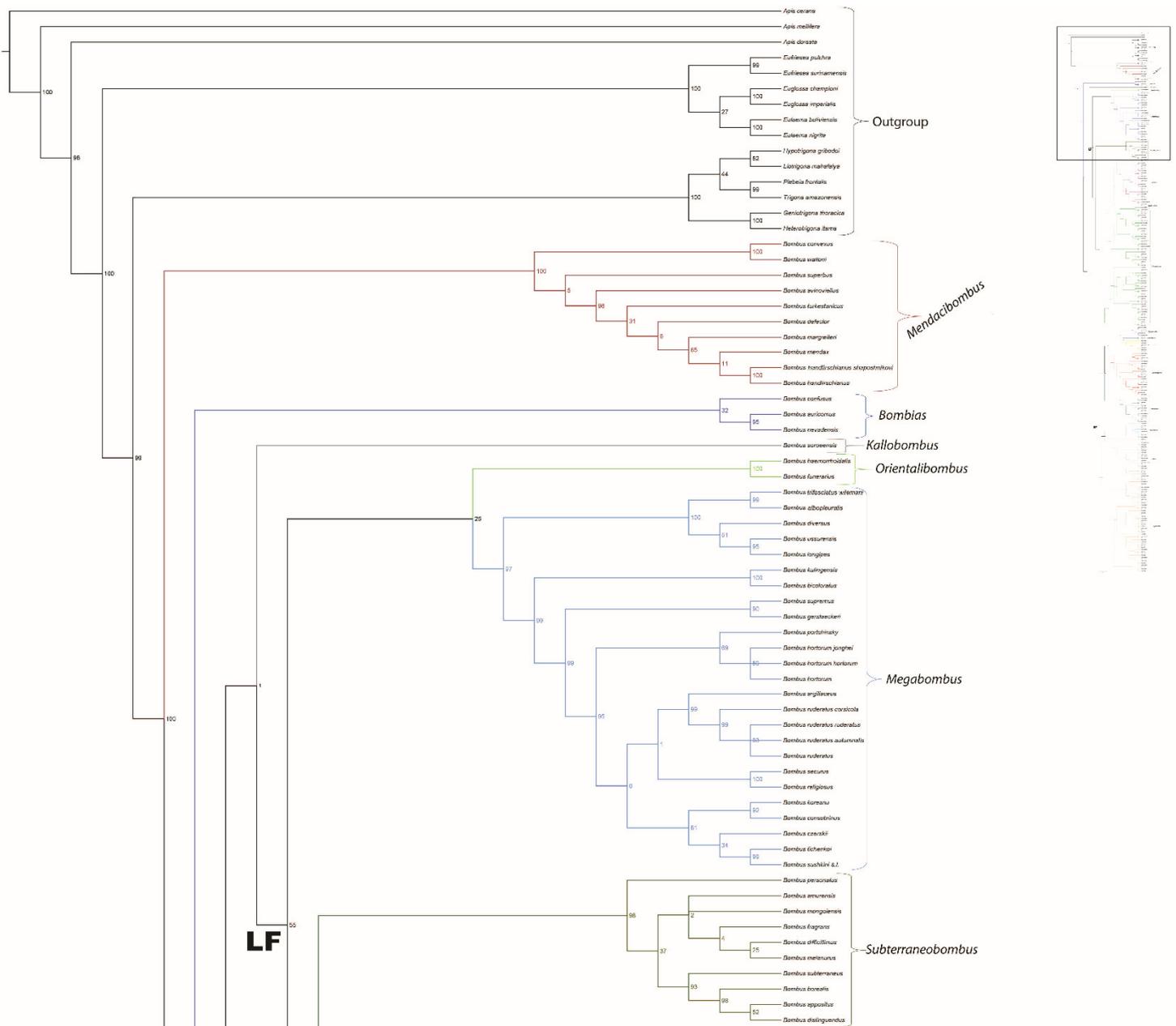
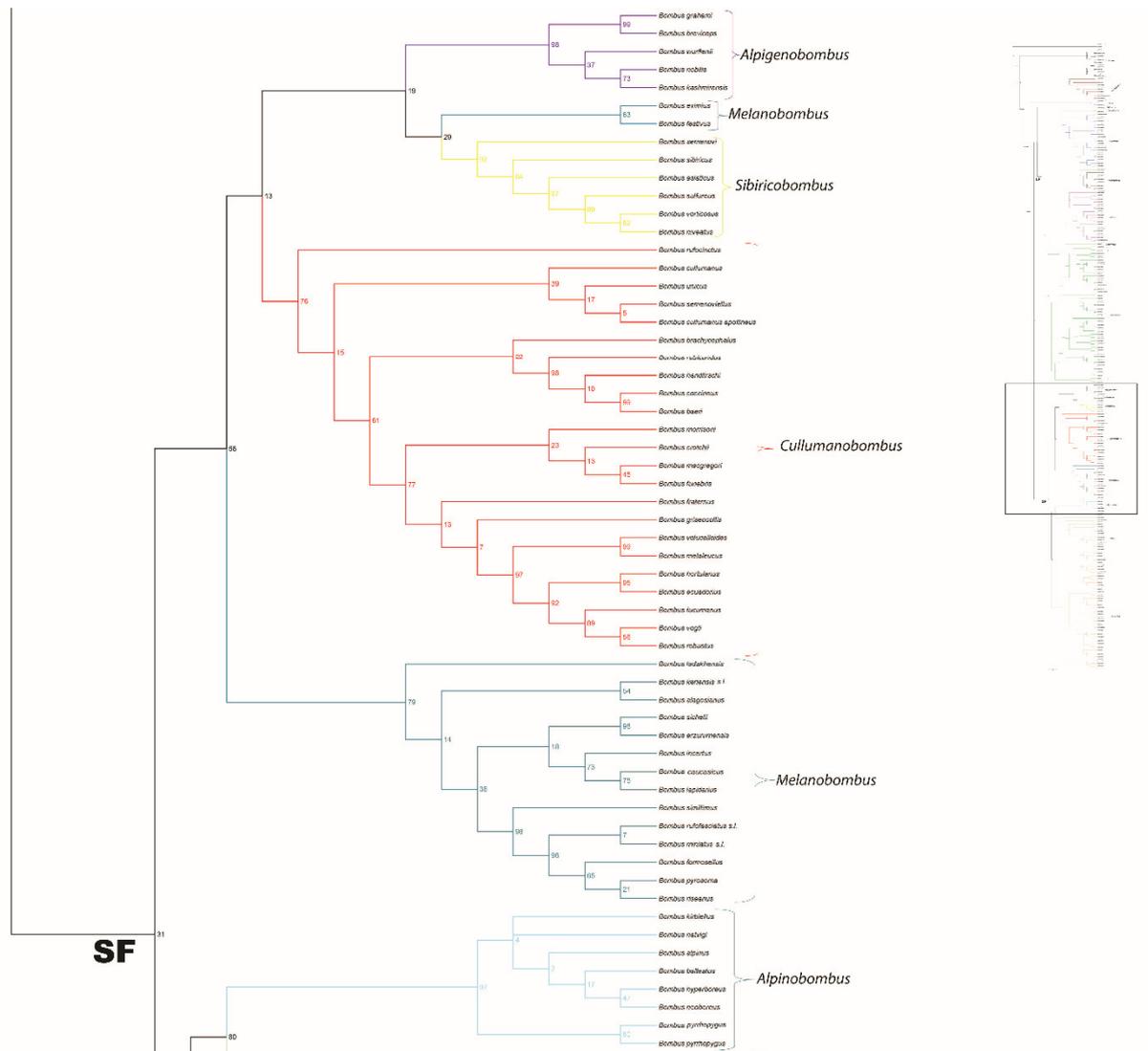


Figure S7. Strict consensus tree of Maximum Parsimony – MP analysis of bumblebees' species. Topologies recovered with the data of matrix presented in Table S5. The values shown above each branch are bootstrap values. Tree performed using a heuristic search with 1000 replications (saving at most 100 trees in each replication) of random (stepwise) addition of taxa followed by Tree Bisection-Reconnection (TBR) branch swapping. Two additional runs using trees on memory were performed to refine searches. To evaluate branch supports we made 10,000 replicates of standard bootstrap. Sister groups of the long-faced (LF) and short-faced (SF) bumblebees were shown.





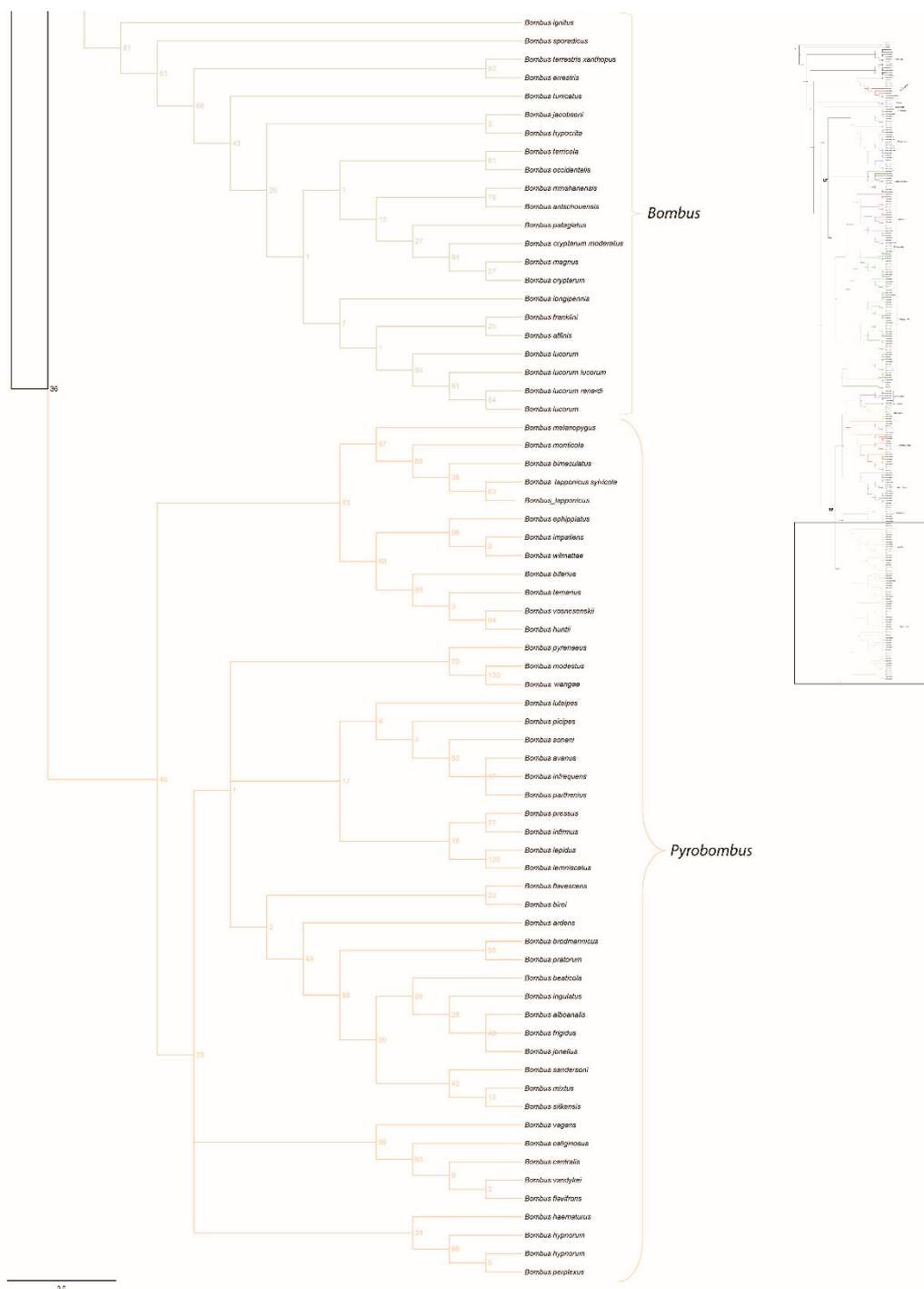
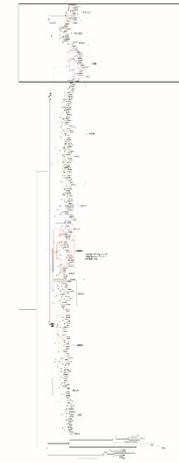
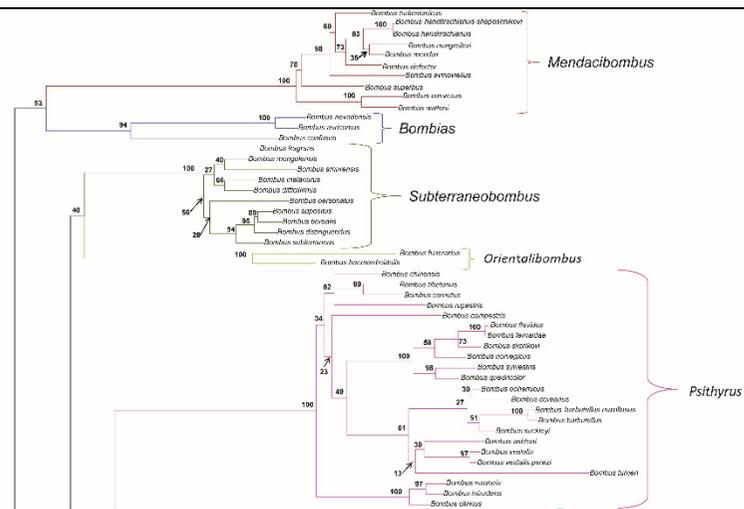


Figure S8. Strict consensus tree of Maximum Parsimony – MP analysis of bumblebees’ species. Topologies recovered with the data of matrix presented in Table S5. The values shown around each branch are bootstrap values. Tree performed using new technology search (NTS) with sectorial search, ratchet, drift and tree fusing options and 10 random addition sequences (saving at most 100 trees in each replication). After analysis with the new technology search were used a round of Tree Bisection Reconnection (TBR). To evaluate branch supports we made 10,000 replicates of standard bootstrap. Sister groups of the long-faced (LF) and short-faced (SF) bumblebees were shown.



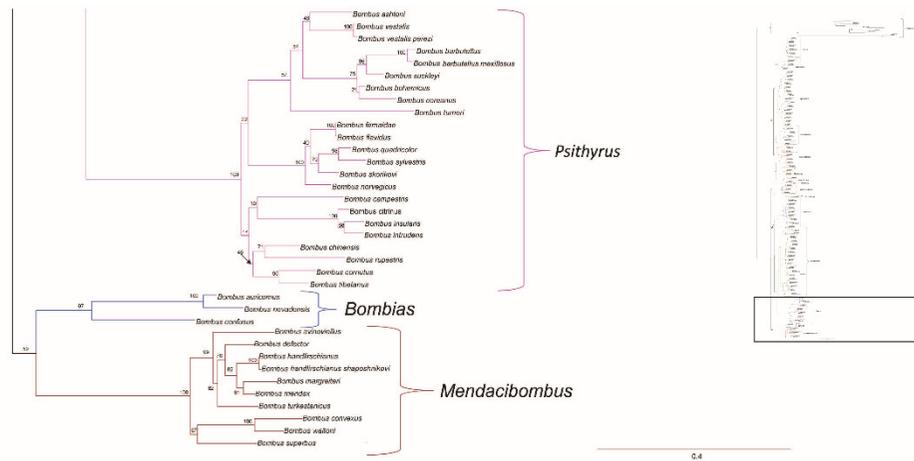


Figure S10. Maximum Likelihood tree – ML search obtained with IQ-TREE. Topologies recovered with the data of matrix presented in Supplementary Table S5. Bootstrap support values are indicated above each branch. Sister groups of the long-faced (LF) and short-faced (SF) bumblebees were shown.

S2. Priors for divergence time estimation

Divergence time estimates. We describe here the paleontological evidence upon which our nine calibration points are based.

Calibration 1 – The fossil bee *Euglossa moronei* Engel, 1999, which is from Miocene Dominican Republic amber [6]. Based on biostratigraphic and paleogeographic data, Dominican amber has been dated from of late Early Miocene through early Middle Miocene age (20.43–13.65 Ma; [7]). For this calibration point, we applied a lognormal distribution with mean of 1.0, Stdev of 0.8 and offset of 15 Ma (95% HPD: 15.6–28 Ma) to the node uniting *Euglossa* subgenus.

Calibration 2 – *Apis lithohermaea* Engel, 2006 is the oldest fossil record for crown group *Apis* (15.97–13.65 Ma). The fossil is from the Chôjabaru Formation of Iki Island, Japan which has been estimated to be from the middle Miocene. Based on some key morphological characters, it was assigned to the *dorsata* species group and was described as being quite similar to modern *Apis dorsata* [8]. This fossil therefore provides a minimum age for the diversification of *Apis dorsata* from *Apis cerana*, a member of the *mellifera* species group. A lognormal distribution with mean of 1.0, Stdev of 0.8 and offset of 10 Ma (95% HPD: 10.6–23 Ma) was applied as a prior for most recent common ancestor between *A. dorsata* and *A. cerana*.

Calibration 3 – The fossil bee *Eulaema (Apeulaema) zigrasi* Engel, 2014 (23.03–15.97 Ma), which is from Early Miocene Mexican amber [9]. For this calibration point, we applied a lognormal distribution with mean of 0.8 and Stdev of 0.9 and offset of 16 Ma (95% HPD: 16.4–29 Ma) to the node uniting *Apeulaema* species.

Calibration 4 – The fossil bee *Paleoeglossa melissiflora* Poinar, 1998 (20.43–13.65 Ma), which is from Dominican Republic [10]. Belongs to *Eufriesea* according to Engel *et al.* [11]. For this calibration point, we applied a lognormal distribution with mean of 0.8 and Stdev of 0.9 and offset of 14 Ma (95% HPD: 14.4–27 Ma) to the node uniting *Eufriesea* species.

Calibration 5 – The fossil bee *Bombus (Bombus) randeckensis* Wappler & Engel, 2012 (18.97–13.65 Ma), which is from the Miocene Randeck Maar of southwestern Germany [12]. For this calibration point, we applied lognormal distribution with a mean of 0.8 and Stdev of 1 and offset of 14 Ma (95% HPD: 14.3–29.8 Ma) to the node uniting *Bombus* s.str.

Calibration 6 – *Oligobombus cuspidatus* Antropov, 2014 (37.2–33.9 Ma), which if from the Late Eocene Bembridge Marls (A'Court Smith Collection) [13]. It likely belongs to stem-group Bombini as well *Calyptapis florissantensis* Cockerell, 1906 [14]. For this calibration point, we applied a lognormal distribution with mean of 1.2 and Stdev of 0.7 and offset of 33 Ma (95% HPD: 33.8–46.1 Ma) to the node uniting *Bombus* species.

Calibration 7 – *Cretotrigona prisca* (Michener & Grimaldi, 1988) (70.6–66.043 Ma) is the oldest fossil bee and the oldest record of eusocial behavior among the Apoidea [15]. This fossil is from Late Cretaceous New Jersey amber, Delaware River, Kinkora [16]. For this calibration point, we applied a lognormal distribution with mean of 1 and Stdev of 0.9 and offset of 66 Ma (95% HPD: 66.5–81.9 Ma) to the node uniting Meliponina.

Calibration 8. The fossil bees *Liotrigonopsis rozeni* Engel, 2001 and *Kelneriapis eocenica* (Kelner-Pillault, 1969) these fossils are from eocenic Baltic amber (AMNH collection) [17]. For this calibration point, we applied a lognormal distribution with mean of 1 and Stdev of 0.8 and offset of 40 Ma (95% HPD: 40.6–53 Ma) to the node uniting *Liotrigona* and *Hypotrigona*.

Calibration 9 – The fossil bee *Bombus (Cullumanobombus) trophonius* Prokop, Dehon, Michez, Engel, 2017 (20.0–16.9 Ma), which is from the Early Miocene deposits of deposits of the Most Basin at the Bílina Mine, Czech Republic [18]. For this calibration point, we applied lognormal distribution with a mean of 1 and Stdev of 0.3 and offset of 15.9 Ma (95% HPD: 17.6–20.4 Ma) to the node uniting *Cullumanobombus*.

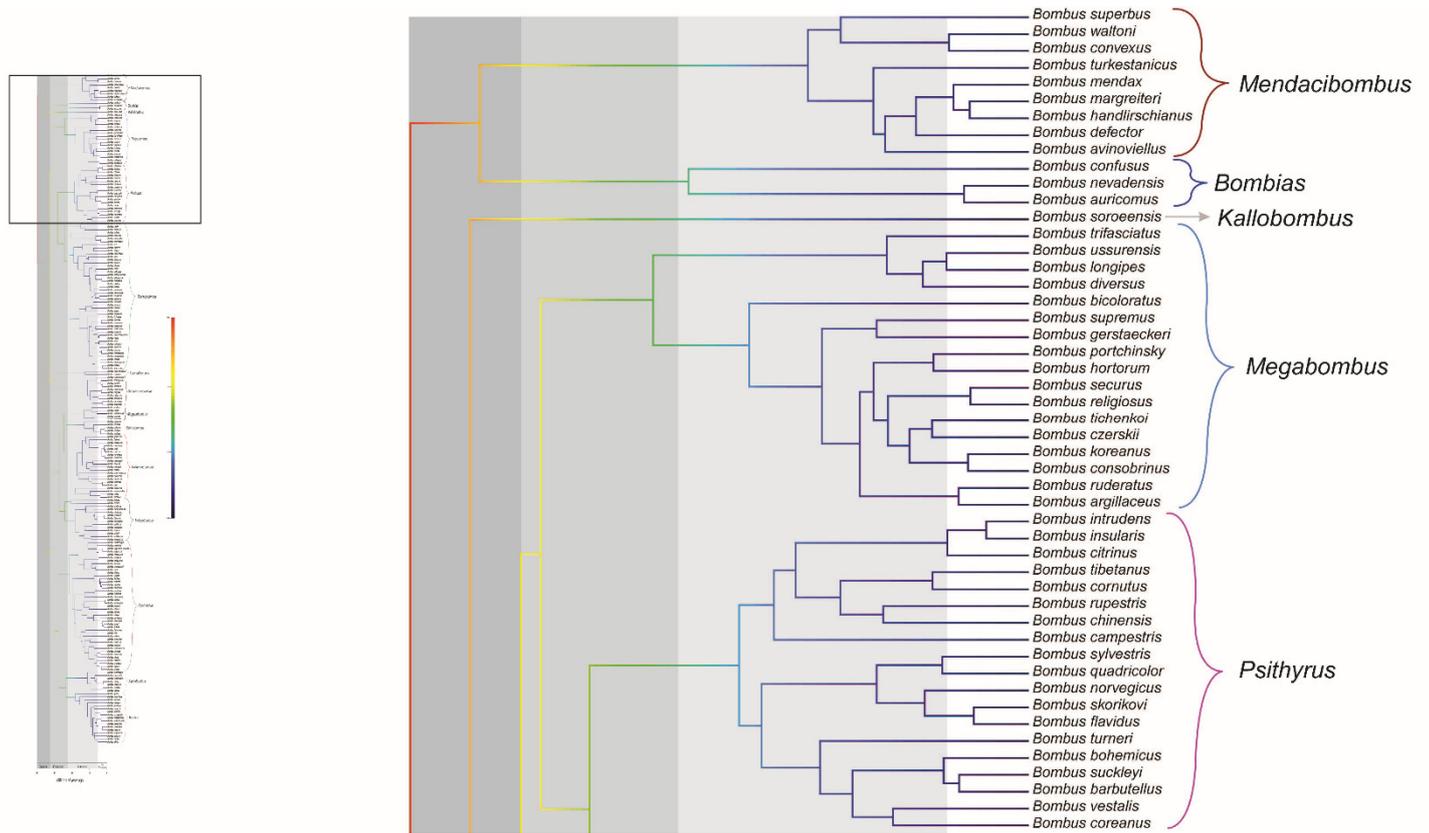
S3. Methods and results of diversification patterns

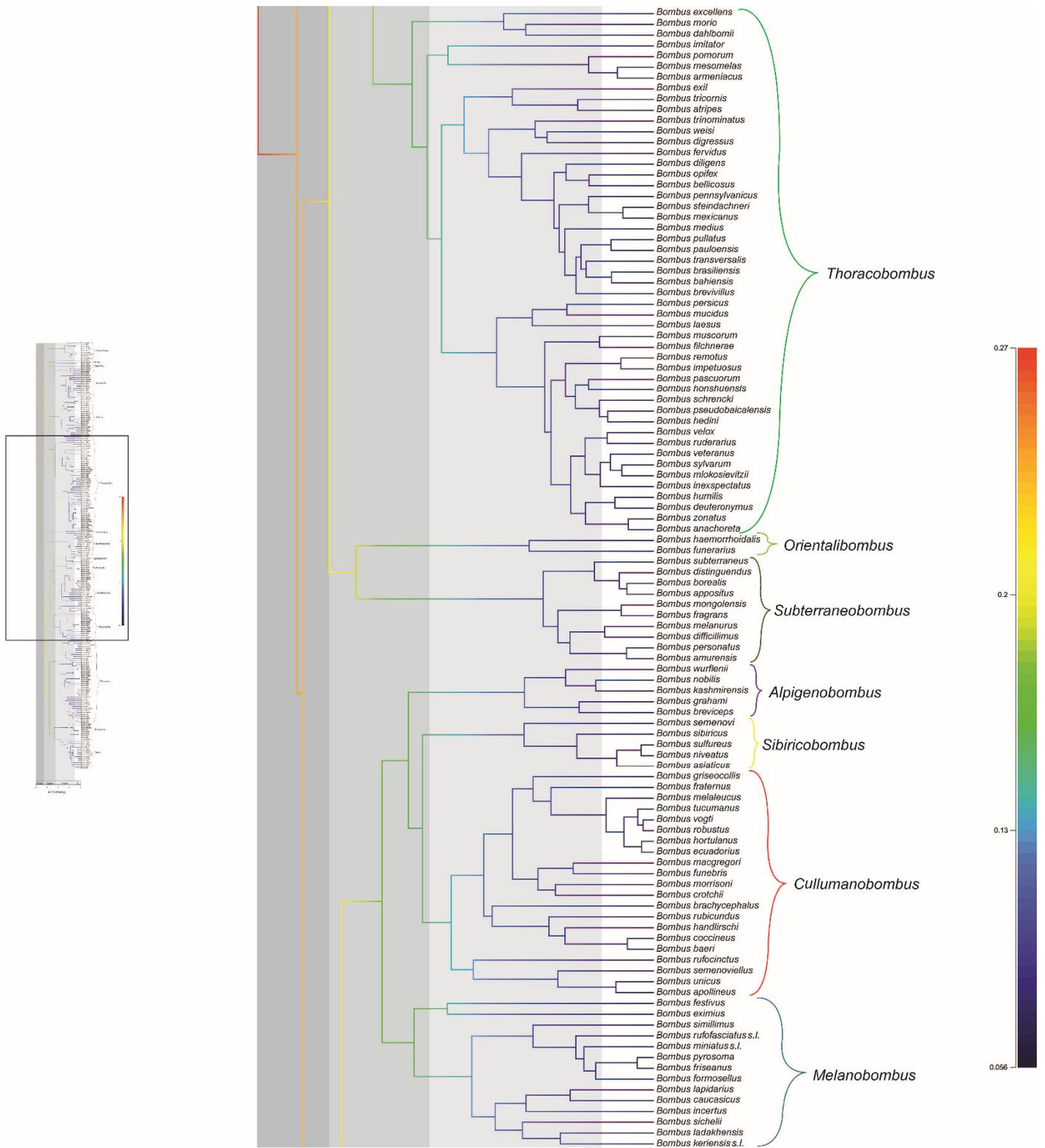
S3.1. Methods

In order to understand the diversification of bumblebee lineages through time and the variation in rates of speciation and extinction, we used Bayesian Analysis of Macroevolutionary Mixtures (BAMM, [19]). This software employs reversible jump MCMC to explore the space of models of lineage diversification, and test hypotheses about the number of divergence events responsible for a given phylogenetic pattern [19]. This is a flexible Bayesian framework for inferring the number and location of shifts in macroevolutionary rates across a dated phylogenetic tree. We used the modified MCC tree as the input phylogeny (see section Biogeographic analyses and Figure S2 for the modified MCC tree and MCC tree, respectively). Rates were inferred across the phylogeny and variation in rates of speciation and extinction of bumblebee species through time were plotted in graphs. The analyses were made with 10 million generations, with a burn-in of 20%. Priors were set by BAMMtools package in R and all output from BAMM were also processed in BAMMtools [1].

S3.2. Results

Sampling models with up to eight different evolutionary regimes were tested, however, only schemes with zero-shift models were consistently selected in the BAMM analysis (genus *Bombus*, PP = 1; Figures S11 and S12). BAMM results showed an overall decreasing pattern (Figures S11 and S12) of marginal Bayesian posterior densities of macroevolutionary rates of diversification along the time and, also, that this variation was similar across the different clades.





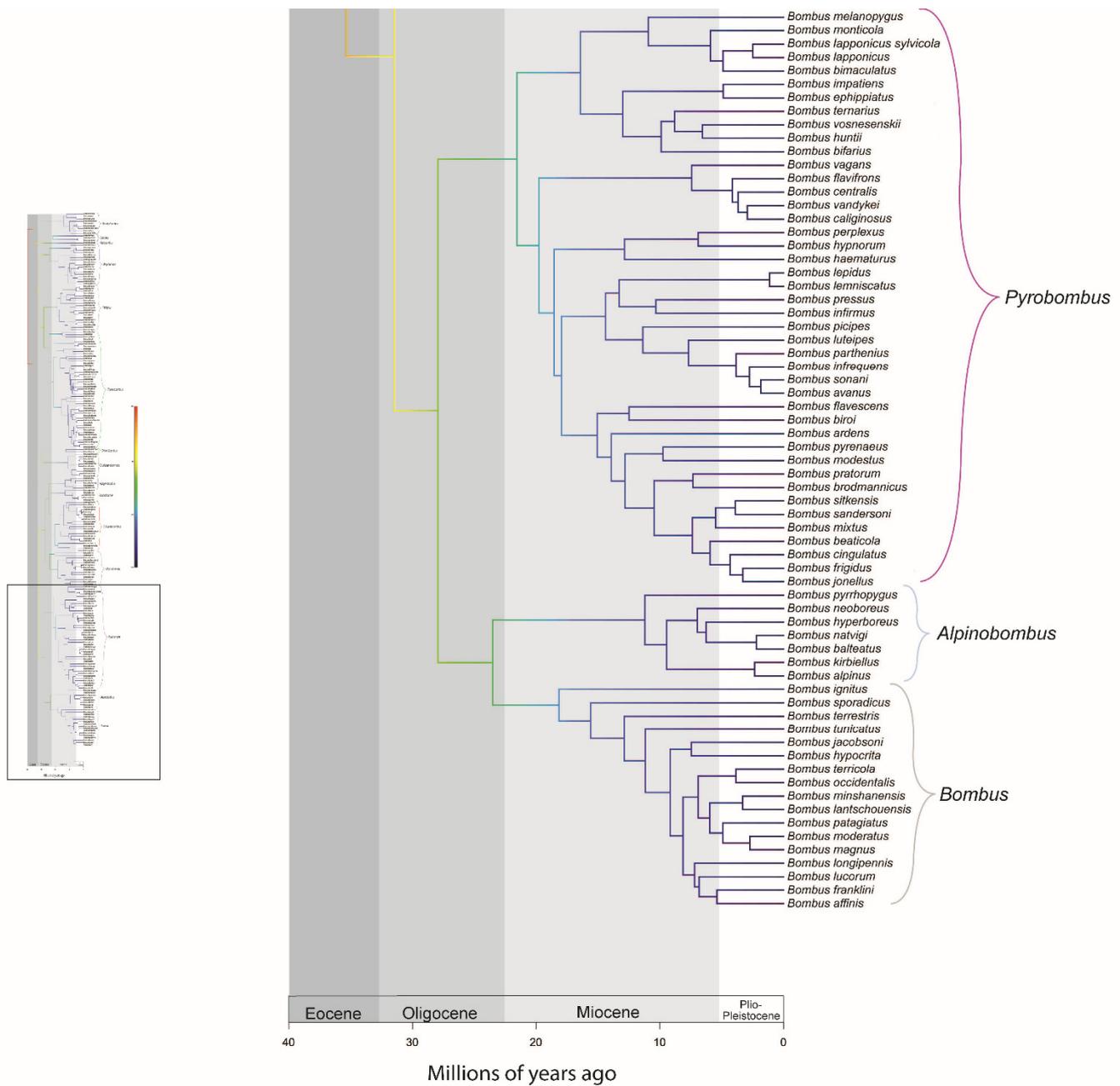


Figure S11. Bumblebee species chronogram with shading of branches reflective of estimated diversification rates (see scale at right) estimated in BAMM. Diversifications are the means of the marginal densities of the rates.

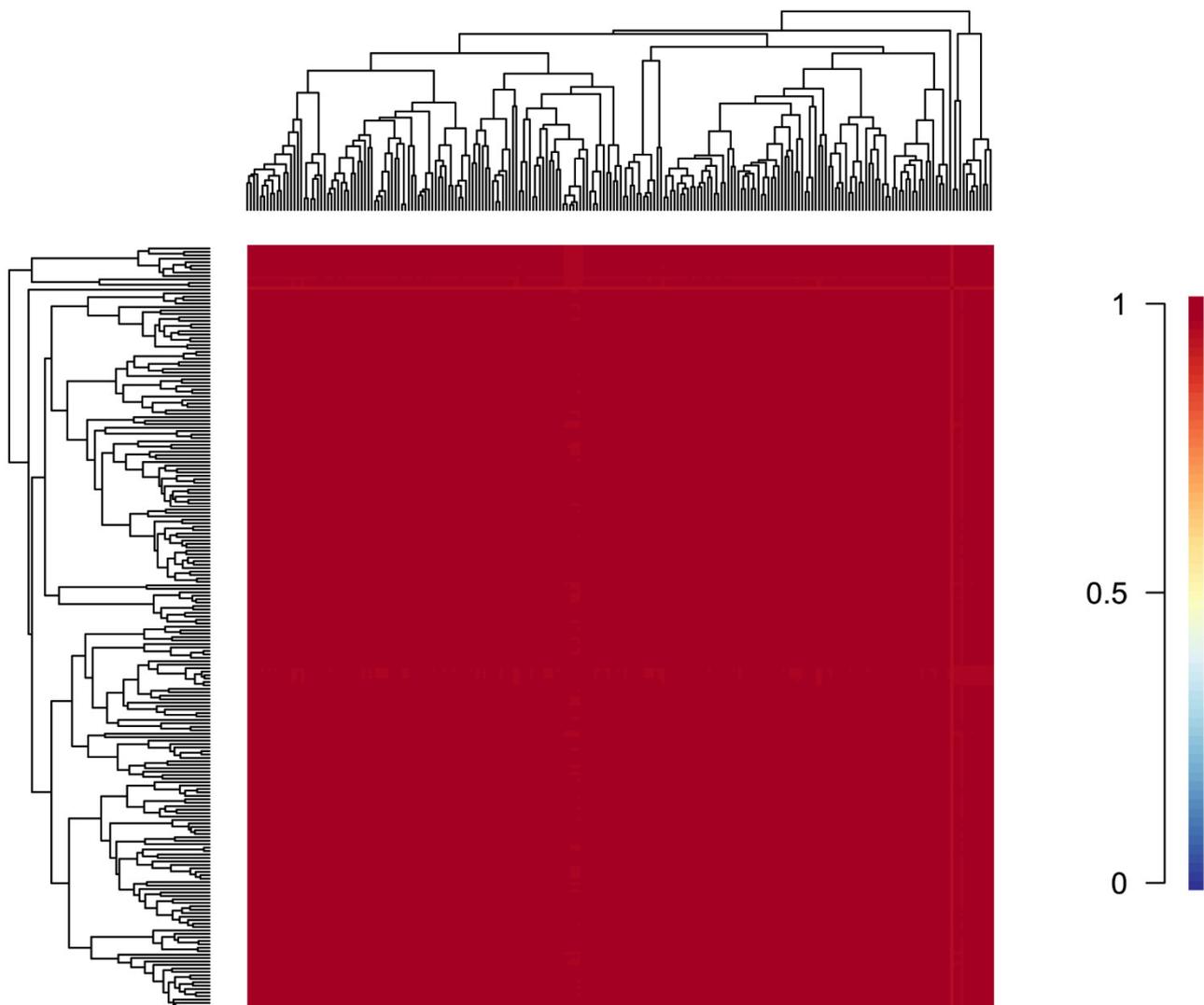


Figure S12. Macroevolutionary cohort matrix for diversification in bumblebees. Pairs of taxa with highly similar rates are depicted in hotter colors (red) and dissimilar rates are shown as colder colors (blue).

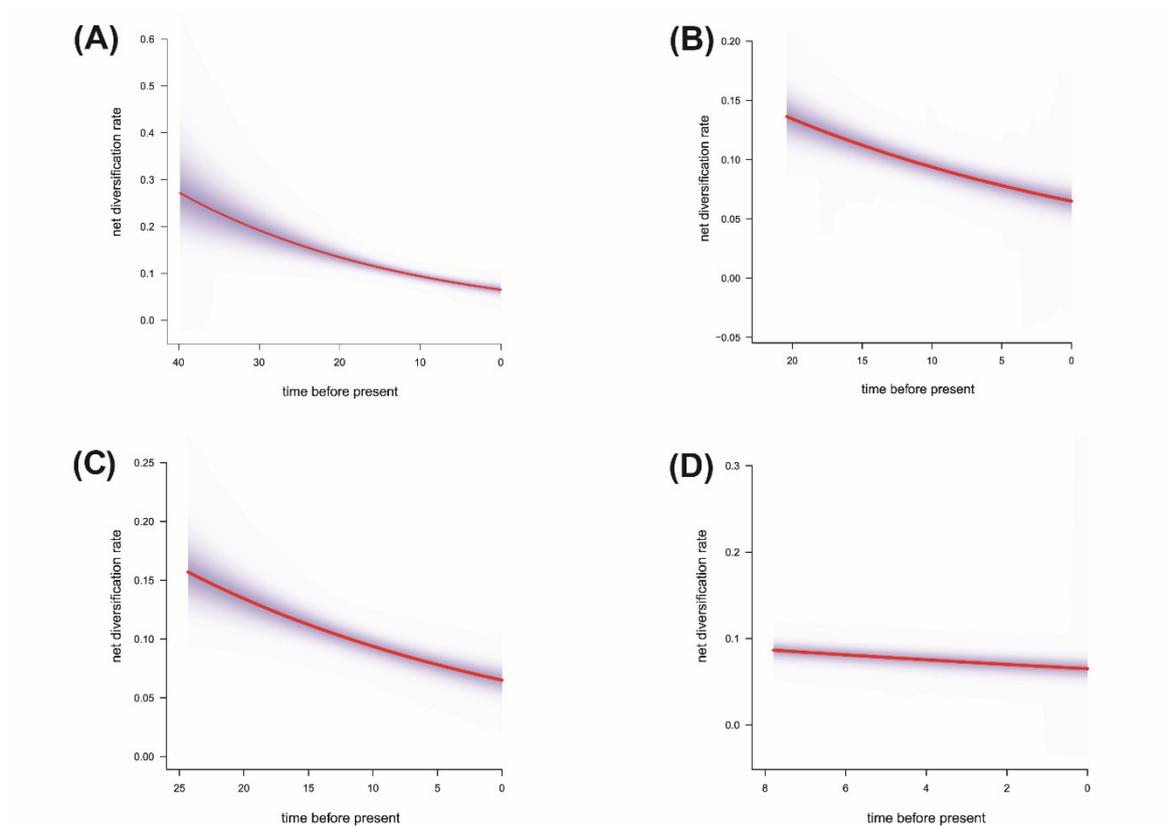


Figure S13. Estimates of speciation rate variability across the bumblebees (red line), with average global climate overlain [20]; purple line. (A)—All species; (B)—Brazilian species; (C)—*Cullumano-*bombus and (D)—*Thoracobombus* subgenus.

S4. Methods and results of missing data analyses

We removed 12S and CytB genes and 31 taxa (which do not share three or more genes with most of the other taxa) to assess the problem missing data in phylogenetic reconstruction and in divergence time estimates, Tables S9 and S10. While the Data set 1 has all characters for all 279 taxa (43.2% of missing data), the Data set 2 has 943,888 characters for 248 taxa (10.2% of missing data), Table S9. We analyzed Data set 2 as done for Data set 1, see sections “Phylogenetic inference” and “Divergence time estimation” in the main text for method descriptions.

The full-length concatenated alignment data employed for phylogenetic inference had a total length of 3,805 bp (see Table S10 for specimens used and amount of missing data). Best partitioning schemes and substitution models are presented in Table S11. The topologies and divergence time estimates found in the analyses of Data set 2 were very similar to those found in the analyses of Data set 1, Figures S14 and S15. Therefore, the differences among our results and the topologies/divergence time estimates found by previous works (*i.e.* Hines [21] and Condamine & Hines [22] results) appears not to be a result of missing data bias in reconstructing trees. Our results strongly support the phylogenetic inferences about topology because informative data were analysed. Furthermore, many studies suggest that missing data are not generally problematic for Bayesian inferences and that the practice of excluding characters simply because they contain missing data may reduce accuracy [23].

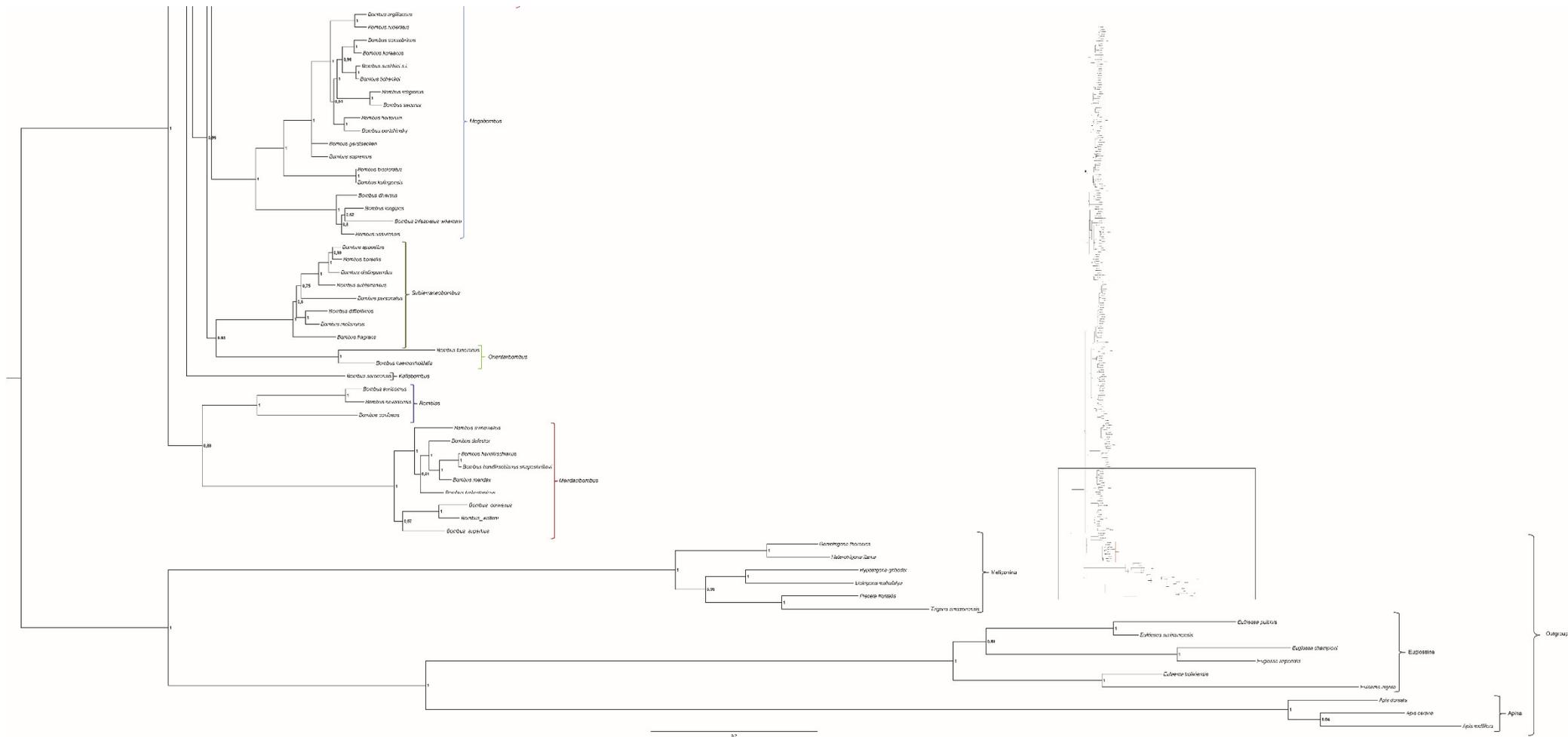
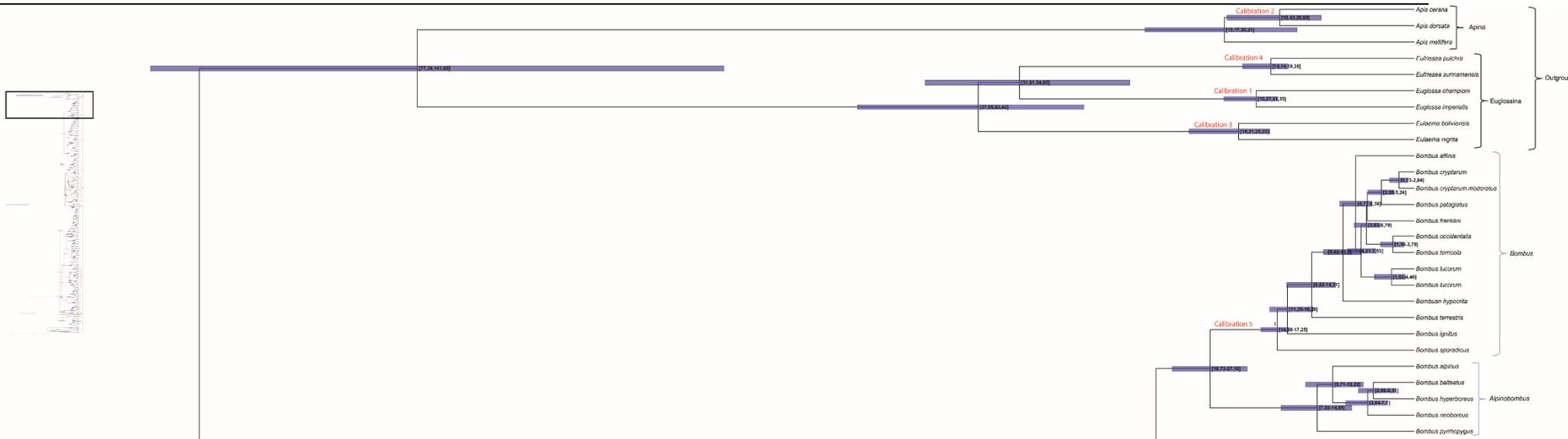
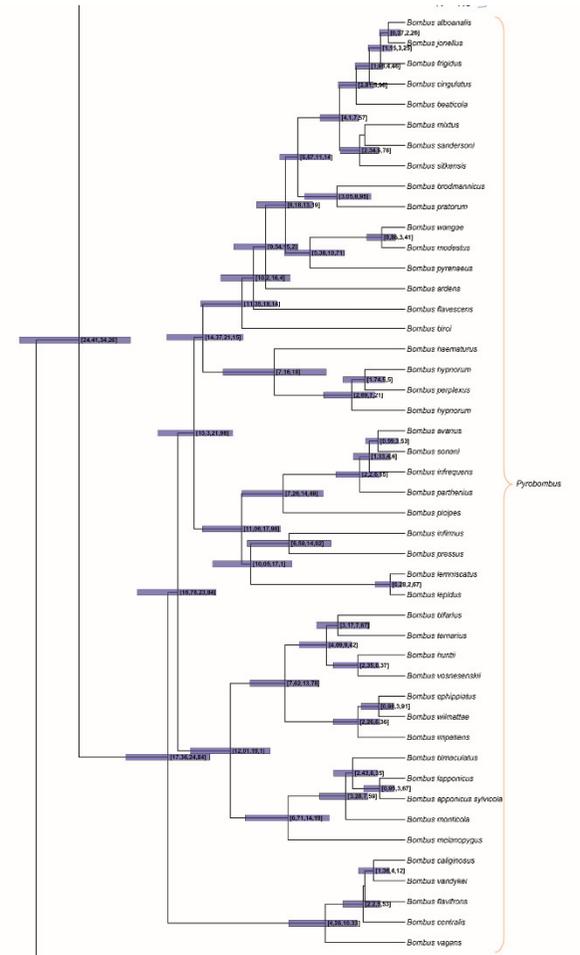
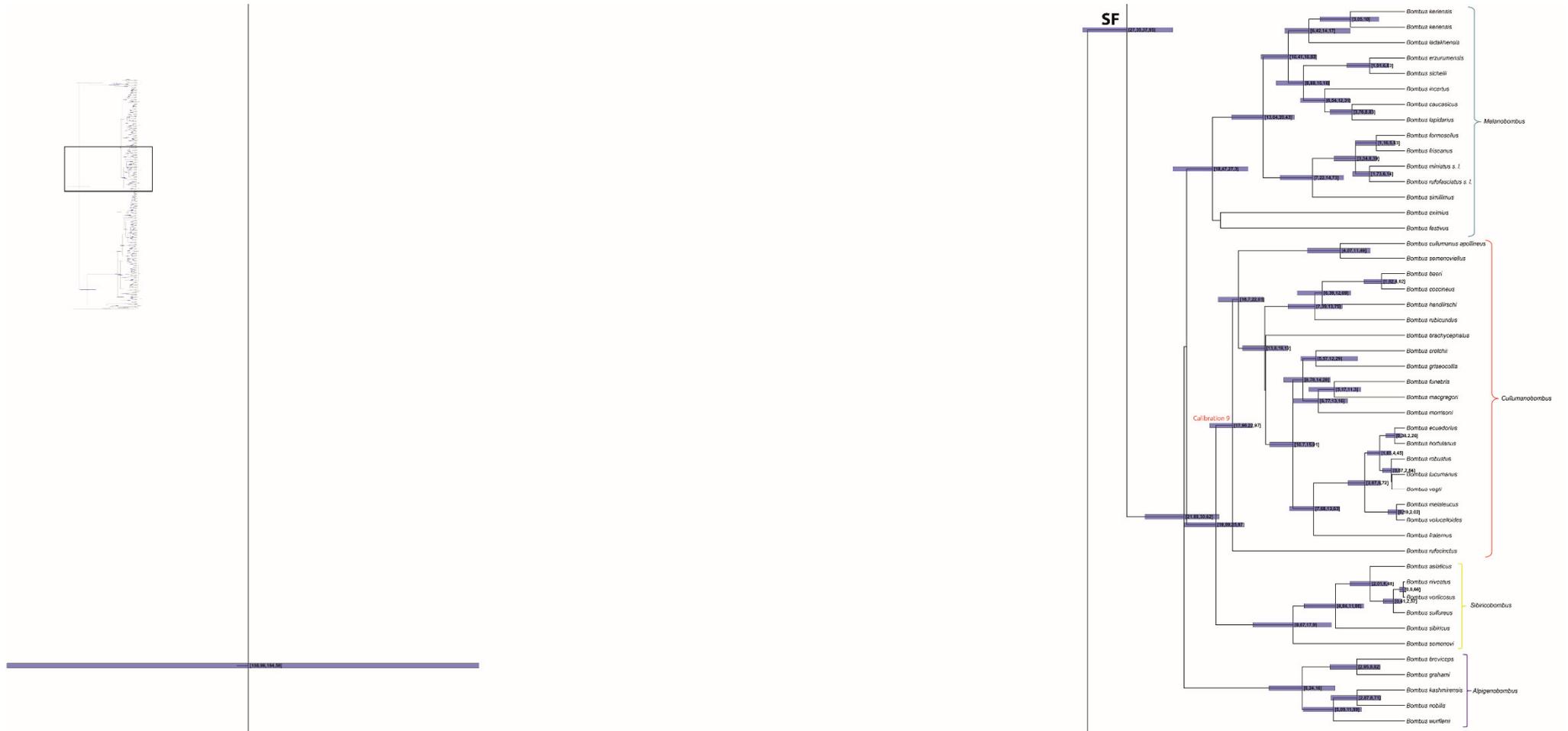


Figure S14. Phylogenetic relationships among bumblebees using Bayesian inference – BI and all species available for each subgenus. Topology recovered with the data of matrix presented in Table S10. Values shown under each branch are posterior probabilities. The standard deviation was 0.009617. Sister group of the long-faced (LF) and short-faced (SF) bumblebees were shown.







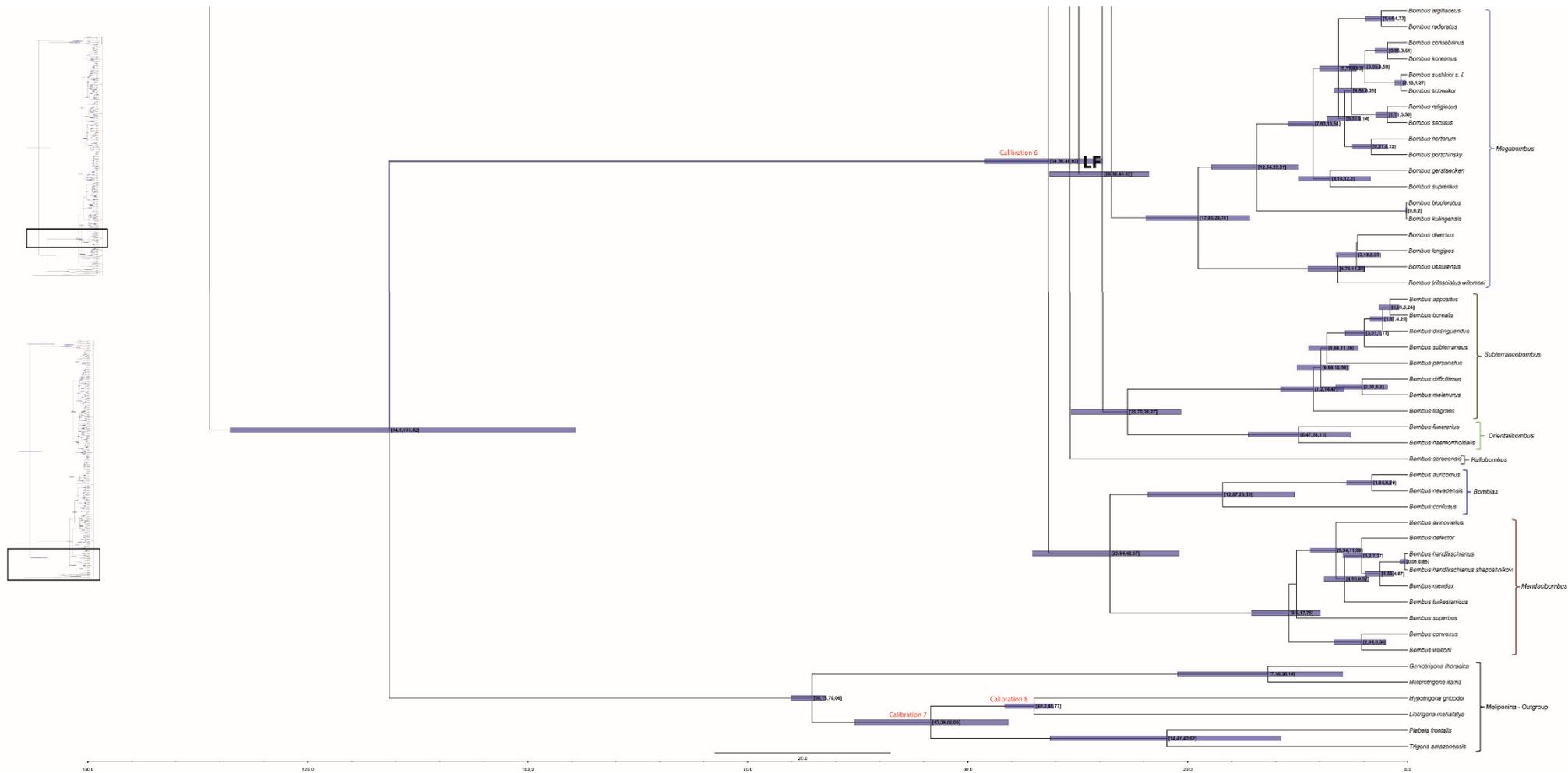


Figure S15. Estimated divergence times using Uncorrelated Lognormal Relaxed Clock model (BEAST v2.4.4). Maximum clade credibility tree depicting branch lengths equal to the median ages calculated from 9,000 post burn-in chronograms (10%). Bars show 95% Highest Posterior Density intervals of age nodes and the numbers in the brackets show dates on million years ago (Ma). Calibration points are shown above the branches in red, these refer to the fossils of Appendix B – Priors for divergence time estimation. The values shown below the tree are dates on million years. Sister group of the long-faced (LF) and short-faced (SF) bumblebees were shown.

Table S9. Basic information on the two data sets used in analyses about the impacts of missing data.

Data Set	Character Data	Number Taxa	Number Characters	Number of Missing Data Characters	Missing Data (%) Per Matrix
1	4 mitochondrial, 4 nuclear genes	279	1,726,731	746,360	43.2
2	2 mitochondrial, 4 nuclear genes	248	943,888	96,090	10.2

Table S10. Data of full-length concatenated alignment data used on phylogenetic analysis. Introns of the genes Argk, Opsin and PECK belonging to outgroup taxa and the 12S were removed of analyses. TL – Total length; N – Number of charsets; Numbers between parenthesis refer to indels and when followed by 'N' the nucleotide base is not known.

Taxon	TL (pb)	N	Argk				EF1 α					
			16S	Exon	Argk Intron	COI	EF1 α Exon	Intron	Opsin Exon	Opsin Intron	PEPCK Exon	PEPCKI
<i>Apis cerana</i>	2496	6	412 (3)	530	0	449	529	85	491	0	0	0
<i>Apis dorsata</i>	2795	7	384 (6)	468	0	480	529	85	501	0	348	0
<i>Apis mellifera</i>	2893	7	384 (4)	599	0	480	529	85	462	0	354	0
<i>Bombus affinis</i> voucher SC167	3805	10	441	599	248	480	529	85	501	101	381	440 (40)
<i>Bombus alagestanus</i> voucher SC085	3736	10	441	539	248	471	529	85	501	101	381	440
<i>Bombus alboanalis</i> voucher SC257	3325	9	441	599	248	0	529	85	501	101	381	440
<i>Bombus alpinus</i> voucher SC029	3805	10	441	599	248	480	529	85	501	101	381	440
<i>Bombus apollineus</i> voucher SC084	3285	9	441	599	248 (32)	0	529	85	501	101	357	424
<i>Bombus applanatus</i> voucher EF209	1810	5	0	488	248 (1)	480	509	85	0	0	0	0
<i>Bombus appositus</i> voucher SC145	3709	10	441	599	248 (1)	384	529	85	501	101	381	440 (11)
<i>Bombus ardens</i> voucher SC131	3736	10	441	530	248	480	529	85	501	101	381	440
<i>Bombus argillaceus</i> voucher SC058	3805	10	441	599	248 (1)	480	529	85	501	101	381	440
<i>Bombus armeniacus</i> voucher SC080	3325	9	441	599	248 (1)	0	529	85	501	101	381	440
<i>Bombus ashtoni</i> voucher SC164	3736	10	441	530	248 (2)	480	529	85	501	101	381	440 (37)
<i>Bombus asiaticus</i> voucher SC249	3167	9	441	539	248	0	529	85	501	92	381	351
<i>Bombus atripes</i> voucher SC066	3311	9	441	588	248 (57)	0	529	85	501	101	378	440

Taxon	TL (pb)	N	16S	Argk			EF1 α					
				Exon	Argk Intron	COI	EF1 α Exon	Intron	Opsin Exon	Opsin Intron	PEPCK Exon	PEPCKI
<i>Bombus auricomus</i> voucher SC062	3650	10	441	444	248 (49)	480	529	85	501	101 (11)	381	440 (3)
<i>Bombus avanus</i> voucher SC272	3264	9	441	599	248	0	529	85	501	101	357	403 (4)
<i>Bombus avinoviellus</i> voucher SC242	3237	9	441	539	248 (12)	0	529	85	473	101 (1 'N', 2)	381	440
<i>Bombus baeri</i> voucher SC174	3265	9	441	539	248 (17)	0	529	85	501	101	381	440
<i>Bombus bahiensis</i> 52893	3572	10	441	539	248 (1)	480	513	85	473	101	348	344 (1)
<i>Bombus balteatus</i> voucher SC039	3796	10	441	599	248	471	529	85	501	101	381	440
<i>Bombus barbutellus</i> voucher SC073	3718	10	441	599	248 (1)	393 (4 'N')	529	85	501	101	381	440 (37)
<i>Bombus beaticola</i> J5	2463	7	0	530	248	480	518	85	501	101	0	0
<i>Bombus bellicosus</i> voucher SC221	3656	10	441	539	248 (1)	419	529	85	473	101 (1 'N', 2)	381	440
<i>Bombus bicoloratus</i> voucher SC225	3639	10	441	530	248 (1)	480	529	85	501	101	348	376
<i>Bombus bifarius</i> voucher SC208	3460	10	441	530	248 (78)	204	529	85	501	101	381	440
<i>Bombus bimaculatus</i> voucher SC218	3805	10	441	599	248 (78)	480	529	85	501	101	381	440
<i>Bombus biroi</i> voucher SC210	3265	9	441	539	248	0	529	85	501	101	381	440
<i>Bombus bohemicus</i> voucher SC055	3649	10	441	530	248 (2)	393 (4 'N')	529	85	501	101	381	440 (37)
<i>Bombus borealis</i> voucher SC250	3688	10	441	530	248 (1)	480	529	85	501	101	351	422 (11)
<i>Bombus brachycephalus</i> voucher SC230	3325	9	441	599	248 (17)	0	529	85	501	101	381	440
<i>Bombus brasiliensis</i> 1218400	3585	10	441	539	248 (1)	480	529	85	473	101	348	341 (1)
<i>Bombus brasiliensis</i> voucher SC219	2796	8	441	0	0	435	529	85	473 (3 'N')	101 (7 'N', 2)	381	351
<i>Bombus breviceps</i> voucher SC190	3739	10	441	599	248	414	529	85	501	101	381	440
<i>Bombus brevivillus</i> 1200916	3412	10	441	414	248 (1)	435	521	85	473	101	348	346 (1)
<i>Bombus brevivillus</i> 1202754	3501	10	441	508	248 (1)	435	520	85	473	101	348	342 (1)
<i>Bombus brevivillus</i> 1207210	3407	10	441	414	248 (1)	461	497	85	473	101	348	339 (1)
<i>Bombus brevivillus</i> 1207320	3324	10	441	414	248 (1)	466	529	85	473	101	348	219
<i>Bombus brevivillus</i> 1207321	3444	10	441	414	248 (1)	466	529	85	473	101	348	339 (1)
<i>Bombus brodmannicus</i> voucher SC077	3773	10	441	599	248	448	529	85	501	101	381	440
<i>Bombus californicus</i> voucher SC306	3297	9	441	599	248 (1)	0	529	85	473	101 (2)	381	440

Taxon	TL (pb)	N	16S	Argk			EF1 α					
				Exon	Argk Intron	COI	EF1 α Exon	Intron	Opsin Exon	Opsin Intron	PEPCK Exon	PEPCKI
<i>Bombus caliginosus</i> voucher SC150	3256	9	441	530	248	0	529	85	501	101	381	440
<i>Bombus campestris</i> voucher SC040	3727	10	441	530	248 (2)	480	520	85	501	101	381	440 (37)
<i>Bombus centralis</i> voucher SC146	3691	10	441	599	248	366	529	85	501	101	381	440
<i>Bombus chinensis</i> voucher SC152	3325	9	441	599	248 (1)	0	529	85	501	101	381	440 (37)
<i>Bombus cingulatus</i> voucher SC212	3647	10	441	530	248	480	529	85	501	101	381	351
<i>Bombus citrinus</i> voucher SC170	3736	10	441	530	248 (53)	480	529	85	501	101	381	440 (37)
<i>Bombus coccineus</i> voucher SC137	3325	9	441	599	248 (17)	0	529	85	501	101	381	440
<i>Bombus confusus</i> voucher SC083	3805	10	441	599	248 (173)	480	529	85	501	101	381	440
<i>Bombus consobrinus</i> voucher SC261	3736	10	441	530	248 (1)	480	529	85	501	101	381	440
<i>Bombus convexus</i> voucher SC109	3189	9	441	539	248 (12)	0	529	85	501	101	348	397
<i>Bombus cornutus</i> voucher SC271	2930	9	441	539	248 (1)	0	529	85	234	33	381	440 (37)
<i>Bombus crotchii</i> voucher SC071	3256	9	441	530	248 (17)	0	529	85	501	101	381	440
<i>Bombus cryptarum</i> voucher SC127	3805	10	441	599	248	480	529	85	501	101	381	440 (40)
<i>Bombus dahlbomii</i> voucher SC016	3460	10	441	530	248 (1)	204	529	85	501	101	381	440
<i>Bombus defector</i> J3	3729	10	441	530	248 (12)	480	529	85	501	101	381	433
<i>Bombus deuteronymus</i> voucher SC147	3655	10	441	530	248 (3)	480	529	83	501	101	381	361 (4)
<i>Bombus difficillimus</i> voucher SC154	3325	9	441	599	248 (1)	0	529	85	501	101	381	440
<i>Bombus digressus</i> voucher SC307	3148	9	441	539	248 (26)	0	529	85	473	101 (4 'N', 2)	381	351 (1)
<i>Bombus diligens</i> voucher SC171	3777	10	441	599	248 (24)	480	529	85	473	101 (1 'N', 2)	381	440
<i>Bombus distinguendus</i> voucher SC197	3748	10	441	599	248 (11)	423	529	85	501	101	381	440 (11)
<i>Bombus diversus</i> voucher SC120	3736	10	441 (2)	530	248 (1)	480	529	85	501	101	381	440
<i>Bombus ecuadorius</i> voucher SC135	3325	9	441	599	248 (17)	0	529	85	501	101	381	440 (5)
<i>Bombus ephippiatus</i> voucher SC198	3649	10	441 (1)	599	248 (78)	324	529	85	501	101	381	440
<i>Bombus erzurumensis</i> voucher SC126	3796	10	441	599	248	471	529	85	501	101	381	440
<i>Bombus excellens</i> voucher SC308	3717	10	441	539	248 (1)	480	529	85	473	101 (2)	381	440
<i>Bombus exil</i> voucher SC232	3245	9	441	539	248 (49)	0	529	85	501	101	361	440

Taxon	TL (pb)	N	16S	Argk			EF1 α					
				Exon	Argk Intron	COI	EF1 α Exon	Intron	Opsin Exon	Opsin Intron	PEPCK Exon	PEPCKI
<i>Bombus eximius</i> voucher SC049	3805	10	441	599	248	480	529	85	501	101	381	440
<i>Bombus fernaldae</i> voucher SC088	3733	10	441	530	248 (2)	477	529	85	501	101	381	440 (37)
<i>Bombus fervidus</i> voucher SC309	3736	10	441	530	248 (1 'N', 1)	480	529	85	501	101	381	440
<i>Bombus festivus</i> voucher SC104	3325	9	441 (1)	599	248	0	529	85	501	101	381	440
<i>Bombus filchnerae</i> voucher SC206	3325	9	441	599	248 (1)	0	529	85	501	101	381	440 (4)
<i>Bombus flavescens</i> voucher SC181	3736	10	441	530	248	480	529	85	501	101	381	440
<i>Bombus flavidus</i> voucher SC041	3796	10	441	599	248 (2)	471	529	85	501	101	381	440 (37)
<i>Bombus flavifrons</i> voucher SC095	3736	10	441	530	248	480	529	85	501	101	381	440
<i>Bombus formosellus</i> J2	2474	7	0	530	248	480	529	85	501	101	0	0
<i>Bombus fragrans</i> voucher SC061	3325	9	441	599	248 (1)	0	529	85	501	101	381	440
<i>Bombus franklini</i> voucher SC256	3781	10	417	599	248	480	529	85	501	101	381	440 (40)
<i>Bombus fraternus</i> voucher SC183	3736	10	441 (1)	530	248 (17)	480	529	85	501	101	381	440
<i>Bombus frigidus</i> voucher SC185	3803	10	441	597 (2 'N')	248	480	529	85	501	101	381	440
<i>Bombus friseanus</i> voucher SC105	3325	9	441	599	248	0	529	85	501	101	381	440
						480 (30						
<i>Bombus funebris</i> voucher SC128	3795	10	431	599	248 (17)	'N')	529	85	501	101	381	440 (1)
<i>Bombus funerarius</i> voucher SC270	3744	10	441	599	248 (1)	419	529	85	501	101	381	440 (13)
<i>Bombus gerstaeckeri</i> voucher SC065	3802	10	441	599	248 (1)	477	529	85	501	101	381	440
<i>Bombus grahami</i> voucher SC273	3499	10	441	519	248	372	529	85	501	101	348	355 (4 'N')
<i>Bombus griseocollis</i> voucher SC082	3256	9	441	530	248 (17)	0	529	85	501	101	381	440
<i>Bombus haematurus</i> voucher SC211	1373	5	441	0	0	0	529	85	0	0	201	117
<i>Bombus haemorrhoidalis</i> voucher SC191	3697	10	441	491	248 (1 'N', 1)	480	529	85	501	101	381	440
<i>Bombus handlirschi</i> voucher SC132	3211	9	441	539	248 (17)	0	529	85	501	101	351	416 (4)
<i>Bombus handlirschianus</i> voucher SC087	3265	9	441	539	248 (16)	0	529	85	501	101	381	440
<i>Bombus hedinii</i> voucher SC129	3325	9	441	599	248 (6)	0	529	85	501	101	381	440 (4)
<i>Bombus honshuensis</i> J7	2474	7	0	530	248 (6)	480	529	85	501	101	0	0

Taxon	TL (pb)	N	16S	Argk			EF1 α					
				Exon	Argk Intron	COI	EF1 α Exon	Intron	Opsin Exon	Opsin Intron	PEPCK Exon	PEPCKI
<i>Bombus hortorum</i> voucher SC005	3736	10	441	530	248 (1)	480	529	85	501	101	381	440
<i>Bombus hortulanus</i> voucher SC200	3325	9	441	599	248 (17)	0	529	85	501	101	381	440 (5)
<i>Bombus humilis</i> voucher SC056	3724	10	441	599	248 (3)	480	529	83	501	101	381	361 (4)
<i>Bombus huntii</i> voucher SC151	3580	10	441	530	248 (78)	324	529	85	501	101	381	440
<i>Bombus hyperboreus</i> voucher SC070	3796	10	441	599	248 (6)	471	529	85	501	101	381	440
<i>Bombus hypnorum</i> voucher SC078	3701	10	441	530	248	471	529	85	501	101	355	440
<i>Bombus hypnorum</i> voucher SC207	3796	10	432	599	248	480	529	85	501	101	381	440
<i>Bombus hypocrita</i> voucher SC123	3736	10	441	530	248	480	529	85	501	101	381	440 (40)
<i>Bombus ignitus</i> voucher SC096	3736	10	441	530	248	480	529	85	501	101	381	440 (49)
<i>Bombus imitator</i> voucher SC028	3325	9	441	599	248 (1)	0	529	85	501	101	381	440 (30)
<i>Bombus impatiens</i> voucher SC060	3714	10	441	530	248 (78)	480	529	85	501	101	359	440
<i>Bombus impetuus</i> voucher SC284	3304	9	441	578	248 (6)	0	529	85	501	101	381	440 (46)
<i>Bombus incertus</i> voucher SC086	3736	10	441	539	248	471	529	85	501	101	381	440
<i>Bombus inexpectatus</i> B05	3042	9	411	509	248 (1)	0	529	85	459	101	348	352 (46)
<i>Bombus infirmus</i> voucher SC157	3315	9	441	599	248 (26)	0	529	85	501	101	381	430
<i>Bombus infrequens</i> voucher SC140	3325	9	441	599	248	0	529	85	501	101	381	440
<i>Bombus insularis</i> voucher SC162	3736	10	441	530	248 (53)	480 (1 'N')	529	85	501	101	381	440 (37)
<i>Bombus jonellus</i> voucher SC079	3805	10	441	599	248	480	529	85	501	101	381	440
<i>Bombus kashmirensis</i> voucher SC121	3324	9	441	598	248	0	529	85	501	101	381	440
<i>Bombus keriensis</i> voucher SC114	3325	9	441	599	248	0	529	85	501	101	381	440
<i>Bombus koreanus</i> voucher SC277	3736	10	441	530	248 (1)	480	529	85	501	101	381	440
<i>Bombus kulingensis</i> voucher SC097	3691	10	441	539	248 (1)	434	521	85	501	101	381	440
<i>Bombus ladakhensis</i> voucher SC158	3319	9	441	599	248	0	523	85	501	101	381	440
<i>Bombus laesus</i> voucher SC052	3792	10	441	599	248 (1)	480	529	85	501	101	369	439 (77)
<i>Bombus lapidarius caucasicus</i> voucher L084TR	2508	7	0	0	0	471	529	85	501	101	381	440
<i>Bombus lapidarius</i> voucher SC006	3736	10	441	530	248	480	529	85	501	101	381	440

Taxon	TL (pb)	N	16S	Argk			EF1 α					
				Exon	Argk Intron	COI	EF1 α Exon	Intron	Opsin Exon	Opsin Intron	PEPCK Exon	PEPCKI
<i>Bombus lapponicus</i> voucher SC103	3804	10	441	599	248 (78)	480	528	85	501	101	381	440
<i>Bombus lemnicatus</i> voucher SC161	3288	9	441	599	248 (22)	0	529	85	501	101	358	426
<i>Bombus lepidus</i> voucher SC155	3325	9	441	599	248 (22)	0	529	85	501	101	381	440
<i>Bombus longipes</i> voucher SC194	3805	10	441 (2)	599	248 (1)	480	529	85	501	101	381	440 (6)
<i>Bombus lucorum</i> voucher SC184	3805	10	441	599	248	480	529	85	501	101	381	440 (40)
<i>Bombus lucorum</i> voucher SC217	3729	10	441	530	248	473	529	85	501	101	381	440 (40)
<i>Bombus macgregori</i> voucher SC231	3325	9	441	599	248 (17)	0	529	85	501	101	381	440
<i>Bombus maxillosus</i> voucher SC074	3325	9	441	599	248 (1)	0	529	85	501	101	381	440 (37)
						480 (24						
<i>Bombus medius</i> voucher SC222	3717	10	441	539	248 (6)	'N')	529	85	473	101 (3 'N', 2)	381	440
<i>Bombus melaleucus</i> voucher SC173	3265	9	441	539	248 (18)	0	529	85	501	101	381	440 (5)
<i>Bombus melanopygus</i> voucher SC215	3736	10	441	530	248 (78)	480	529	85	501	101	381	440
<i>Bombus melanurus</i> voucher SC022	3647	10	441	530	248 (1)	480	529	85	501	101	381	351
<i>Bombus mendax</i> voucher SC019	3730	10	441	530	248 (12)	480	523	85	501	101	381	440
<i>Bombus mesomelas</i> voucher SC037	3805	10	441	599	248 (1)	480	529	85	501	101	381	440
<i>Bombus mexicanus</i> voucher SC220	3432	10	441	527	248 (6)	207	529	85	473	101 (3 'N', 2)	381	440
<i>Bombus miniatus</i> voucher SC244	3265	9	441	539	248	0	529	85	501	101	381	440
<i>Bombus mixtus</i> voucher SC024	3715	10	441	530	248	459	529	85	501	101	381	440
<i>Bombus mlokosievitzii</i> voucher SC081	3325	9	441	599	248 (1)	0	529	85	501	101	381	440 (46)
<i>Bombus moderatus</i> voucher SC163	3805	10	441	599	248	480 (1 'N')	529	85	501	101	381	440 (40)
<i>Bombus modestus</i> voucher SC160	3805	10	441	599	248	480	529	85	501	101	381	440
<i>Bombus modestus</i> voucher SC238	3805	10	441 (1)	599	248	480	529	85	501	101	381	440
<i>Bombus monticola</i> voucher SC176	3805	10	441	599	248 (78)	480	529	85	501	101	381	440
<i>Bombus morio</i> B13	3548	10	441	537	248 (1)	435	529	85	473	101	348	351 (1)
<i>Bombus morio</i> voucher SC310	3732	10	441	599	248 (1)	435	529	85	473 (1 'N')	101 (2)	381	440
<i>Bombus morrisoni</i> voucher SC196	3325	9	441	599	248 (17)	0	529	85	501	101	381	440

Taxon	TL (pb)	N	16S	Argk			EF1 α					
				Exon	Argk Intron	COI	EF1 α Exon	Intron	Opsin Exon	Opsin Intron	PEPCK Exon	PEPCKI
<i>Bombus mucidus</i> voucher SC059	3641	10	441	530	248 (1)	395	519	85	501	101	381	440 (4)
<i>Bombus muscorum</i> voucher SC033	3796	10	441	599	248 (1)	471	529	85	501	101	381	440 (4)
<i>Bombus neoboreus</i> voucher SC188	3325	9	441	599	248 (6)	0	529	85	501	101	381	440
<i>Bombus nevadensis</i> voucher SC139	3324	10	441	428	248	204	529	85	501	101 (11)	357	430 (6)
<i>Bombus niveatus</i> voucher SC093	3224	9	441	599	248	0	529	84	501	90	381	351
<i>Bombus nobilis</i> voucher SC098	3325	9	441	599	248	0	529	85	501	101	381	440
<i>Bombus norvegicus</i> voucher SC089	3778	10	441	599	248 (2)	453	529	85	501	101	381	440 (37)
<i>Bombus oberti</i> voucher SC234	3316	9	441	599	248	0	529	85	501	92	381	440 (2)
<i>Bombus occidentalis</i> voucher SC026	3778	10	441	599	248	453	529	85	501	101	381	440 (40)
<i>Bombus opifex</i> voucher SC175	3805	10	441	599	248 (1)	480	529	85	501	101	381	440
<i>Bombus parthenius</i> voucher SC241	3265	9	441	539	248	0	529	85	501	101	381	440
<i>Bombus pascuorum</i> voucher SC023	3736	10	441	530	248 (5)	480	529	85	501	101	381	440 (4)
<i>Bombus patagiatus</i> voucher SC111	3736	10	441	530	248	480	529	85	501	101	381	440 (40)
<i>Bombus pauloensis</i> 77202	2767	8	441	0	0	448	529	85	473	101	348	342 (1)
<i>Bombus pauloensis</i> voucher SC305	3670	10	441	539	248 (6)	435	527	85	473	101 (1 'N', 2)	381	440
<i>Bombus pennsylvanicus</i> voucher SC311	3322	9	441	599	248 (1)	0	529	85	501	101	381	437
<i>Bombus pennsylvanicus</i> voucher SC317	3805	10	441	599	248 (1)	480	529	85	501	101	381	440
<i>Bombus perplexus</i> voucher SC166	3736	10	441	530	248	480	529	85	501	101	381	440
<i>Bombus persicus</i> voucher SC054	3273	9	441	573	248 (15)	0	527	85	501	101	359	438
<i>Bombus personatus</i> voucher SC138	3325	9	441	599	248 (1)	0	529	85	501	101	381	440
<i>Bombus picipes</i> voucher SC180	3325	9	441	599	248	0	529	85	501	101	381	440
<i>Bombus polaris</i> voucher SC223	3638	10	441	530	248	471	529	85	501	101	381	351
<i>Bombus pomorum</i> voucher SC053	3325	9	441	599	248 (1)	0	529	85	501	101	381	440
<i>Bombus portchinsky</i> voucher SC072	3767	10	441	599	248 (1)	480	529	85	501	101	353	430 (12 'N')
<i>Bombus potanini</i> voucher SC113	3317	9	441	599	248 (6)	0	521	85	501	101	381	440 (46)
<i>Bombus pratorum</i> voucher SC075	3686	10	441	530	248	480	529	85	501	101	351	420

Taxon	TL (pb)	N	16S	Argk			EF1 α					
				Exon	Argk Intron	COI	EF1 α Exon	Intron	Opsin Exon	Opsin Intron	PEPCK Exon	PEPCKI
<i>Bombus pressus</i> voucher SC239	1365	5	441	0	0	0	529	85	0	0	201	109
<i>Bombus pseudobaicalensis</i> voucher SC253	3736	10	441	530 (2 'N')	248 (6)	480	529	85	501	101	381	440 (4)
<i>Bombus pullatus</i> voucher SC312	3501	10	441	599	248 (1)	204	529	85	473 (1 'N')	101 (3 'N', 2)	381	440
<i>Bombus pyrenaicus</i> voucher SC035	3805	10	441	599	248	480	529	85	501	101	381	440
<i>Bombus quadricolor</i> voucher SC090	3805	10	441	599	248 (2)	480	529	85	501	101	381	440 (37)
<i>Bombus religiosus</i> voucher SC141	3600	10	441	599	248 (1)	480	529	85	296	101	381	440
<i>Bombus remotus</i> voucher SC192	3325	9	441	599	248 (6)	0	529	85	501	101	381	440 (46)
<i>Bombus robustus</i> voucher SC050	3469	10	441	539	248 (17)	204	529	85	501	101	381	440 (5)
<i>Bombus rubicundus</i> voucher SC202	3325	9	441	599	248 (17)	0	529	85	501	101 (2)	381	440
<i>Bombus ruderarius</i> voucher SC047	3726	10	441	530	248 (1)	480	519	85	501	101	381	440 (46)
<i>Bombus ruderatus</i> voucher SC018	3736	10	441	530	248 (1)	480	529	85	501	101	381	440
<i>Bombus rufocinctus</i> voucher SC186	3705	10	441	530	248 (8)	449	529	85	501	101 (1)	381	440
<i>Bombus rufofasciatus</i> voucher SC133	3296	9	441	599	248	0	529	85	501	101	352	440
<i>Bombus rupestris</i> voucher SC009	3736	10	441	530	248 (1)	480	529	85	501	101	381	440 (37)
<i>Bombus sandersoni</i> voucher SC255	3782	10	418	599	248	480 (4 'N')	529	85	501	101	381	440
<i>Bombus schrencki</i> voucher SC298	3691	10	412	530	248 (6)	480	513	85	501	101	381	440 (4)
<i>Bombus securus</i> voucher SC142	3805	10	441	599	248 (1)	480	529	85	501	101	381	440
<i>Bombus semenoviellus</i> voucher SC236	3805	10	441	599	248 (32)	480	529	85	501	101	381	440
<i>Bombus shaposhnikovii</i> voucher SC099	3265	9	441	539	248 (16)	0	529	85	501	101	381	440
<i>Bombus sibiricus</i> voucher SC274	3705	10	441	599	248	480	529	85	501	90	381	351
<i>Bombus sichelii</i> voucher SC034	3420	10	441	599	248	480	529	85	196	21	381	440
<i>Bombus simillimus</i> voucher SC243	3324	9	441	599	248	0	529	85	501	101	380	440
<i>Bombus sitkensis</i> voucher SC144	3325	9	441	599	248 (7)	0	529	85	501	101	381	440
<i>Bombus skorikovi</i> voucher SC159	3325	9	441	599	248 (2)	0	529	85	501	101	381	440 (37)
<i>Bombus sonani</i> J6	2474	7	0	530	248	480	529	85	501	101	0	0
<i>Bombus sonorus</i> voucher SC051	3315	9	431	599	248 (1)	0	529	85	501	101	381	440

Taxon	TL (pb)	N	16S	Argk			EF1 α					
				Exon	Argk Intron	COI	EF1 α Exon	Intron	Opsin Exon	Opsin Intron	PEPCK Exon	PEPCKI
<i>Bombus sonorus</i> voucher SC318	3297	9	441 (1)	599	248 (1)	0	529	85	473	101 (1 'N', 2)	381	440
<i>Bombus soroensis</i> voucher SC136	3736	10	441	530	248	480	529	85	501	101 (3)	381	440 (17)
<i>Bombus sporadicus</i> voucher SC193	3805	10	441	599	248 (61)	480	529	85	501	101	381	440 (40)
<i>Bombus steindachneri</i> voucher SC313	3297	9	441	599	248 (1 'N', 1)	0	529	85	473	101 (2)	381	440
<i>Bombus subterraneus</i> voucher SC046	3727	10	441	530	248 (1)	471	529	85	501	101	381	440
<i>Bombus suckleyi</i> voucher SC091	3325	9	441	599	248 (1)	0	529	85	501	101	381	440 (37)
<i>Bombus sulfureus</i> voucher SC064	3219	9	441	594	248	0	529	84	501	90	381	351
<i>Bombus superbis</i> B02	1857	5	441	0	0	0	0	0	501	101	381	433
<i>Bombus supremus</i> voucher SC101	3805	10	441	599	248 (1)	480	529	85	501	101	381	440
<i>Bombus sushkini</i> voucher SC143	3736	10	441	530	248 (1)	480	529	85	501	101	381	440
<i>Bombus sylvarum</i> voucher SC110	3769	10	441	599	248 (1)	480	529	85	501	101	348	437 (44)
<i>Bombus sylvestris</i> voucher SC020	3757	10	441 (1 'N')	599	248 (2)	480	516	85	466	101	381	440 (37)
<i>Bombus sylvicola</i> voucher SC108	3775	10	441	599	248 (78)	477	529	85	501	101	354	440
<i>Bombus ternarius</i> voucher SC116	3736	10	441	530	248 (78)	480	529	85	501	101	381	440
<i>Bombus terrestris</i> voucher SC003	3736	10	441	530	248	480	529	85	501	101	381	440 (47)
<i>Bombus terricola</i> voucher SC205	3736	10	441	530	248	480	529	85	501	101	381	440 (40)
<i>Bombus tibetanus</i> voucher SC134	3325	9	441	599	248 (1)	0	529	85	501	101	381	440 (37)
<i>Bombus tichenkoi</i> J1	2474	7	0	530	248 (1)	480	529	85	501	101	0	0
<i>Bombus transversalis</i> voucher SC314	3777	10	441	599	248 (6)	480	529	85	473	101 (3 'N', 2)	381	440
<i>Bombus tricornis</i> voucher SC148	3736	10	441 (1)	530	248	480	529	85	501	101	381	440
<i>Bombus trifasciatus</i> voucher SC015	3686	10	441 (2)	530	248 (1)	434	525	85	501	101	381	440
<i>Bombus trinominatus</i> voucher SC229	3325	9	441	599	248 (60)	0	529	85	501	101	381	440
<i>Bombus tucumanus</i> voucher SC276	3325	9	441	599	248 (17)	0	529	85	501	101	381	440 (5)
<i>Bombus turkestanicus</i> B03	1857	5	441	0	0	0	0	0	501	101	381	433
<i>Bombus ussurensis</i> voucher SC130	3656	10	412 (2)	530	248 (1)	480	529	85	501	101	350	420 (6)
<i>Bombus vagans</i> voucher SC044	3788	10	424	599	248	480	529	85	501	101	381	440

Taxon	TL (pb)	N	16S	Argk			EF1 α					
				Exon	Argk Intron	COI	EF1 α Exon	Intron	Opsin Exon	Opsin Intron	PEPCK Exon	PEPCKI
<i>Bombus vandykei</i> voucher SC149	3675	10	441	530	248	419	529	85	501	101 (1 'N')	381	440
<i>Bombus variabilis</i> voucher SC316	3297	9	441	599	248 (1)	0	529	85	473	101 (3 'N', 2)	381	440 (37)
<i>Bombus velox</i> voucher SC094	3252	9	441	526	248 (1)	0	529	85	501	101	381	440 (46)
<i>Bombus vestalis</i> voucher SC169	3805	10	441	599	248 (1)	480	529	85	501	101	381	440 (37)
<i>Bombus veteranus</i> voucher SC187	3805	10	441	599	248 (6)	480	529	85	501	101	381	440 (46)
<i>Bombus vogti</i> voucher SC172	3325	9	441	599	248 (17)	0	529	85	501	101	381	440 (5)
<i>Bombus volucelloides</i> voucher SC122	2435	7	441	530	248 (18)	0	529	85	501	101	0	0
<i>Bombus vorticosus</i> voucher SC124	3158	9	441	533	248	0	529	84	501	90	381	351
<i>Bombus vosnesenskii</i> voucher SC112	3580	10	441	530	248 (78)	324	529	85	501	101	381	440
<i>Bombus waltoni</i> voucher SC102	3716	10	441	539	248 (12)	480	529	85	501	101	352	440
<i>Bombus weisi</i> voucher SC315	3500	10	441	599	248 (89)	203	529	85	473 (1 'N')	101 (3 'N', 2)	381	440
<i>Bombus wilmattae</i> voucher SC199	3649	10	441	599	248 (78)	324	529	85	501	101	381	440
<i>Bombus wurflenii</i> voucher SC001	3670	10	441	530	248	414	529	85 (1 'N')	501	101	381	440
<i>Bombus zonatus</i> voucher SC063	3325	9	441	599	248 (17)	0	529	85	501	101	381	440 (46)
<i>Eufriesea pulchra</i>	1634	4	0	530	0	0	529	85	490	0	0	0
<i>Eufriesea surinamensis</i>	2175	5	0	583	0	480	526	85	501	0	0	0
<i>Euglossa championi</i>	2579	6	416 (10)	582	0	480	529	85	487	0	0	0
<i>Euglossa imperialis</i>	2425	6	383 (1)	467	0	460	529	85	501	0	0	0
<i>Eulaema boliviensis</i>	2983	7	441 (2)	599	0	480	529	85	501	0	348	0
<i>Eulaema nigrita</i>	2440	6	416 (5)	474 (2 'N')	0	480 (1 'N')	498	85	487	0	0	0
<i>Geniotrigona thoracica</i>	2423	6	441 (1)	556	0	0	494 (5 'N')	85	475	0	372	0
<i>Heterotrigona itama</i>	2408	6	441 (2)	539	0	0	506	85	486	0	351	0
							528 (1 'N',					
<i>Hypotrigona gribodoi</i>	2396	6	441	539	0	0	3)	85 (1)	442	0	361	0
<i>Liotrigona mahafalya</i>	2516	6	441	539	0	480	529 (1 'N')	85 (1)	442 (1 'N')	0	0	0
<i>Plebeia frontalis</i>	2413	6	441	539	0	0	529	85 (1)	442	0	377	0

Taxon	TL (pb)	N	16S	Argk			EF1 α					
				Exon	Argk Intron	COI	EF1 α Exon	Intron	Opsin Exon	Opsin Intron	PEPCK Exon	PEPCKI
<i>Trigona amazonensis</i>	2463	6	441 (2)	530	0	0	529	85 (1)	501	0	377 (3)	0

Table S11. Best partitioning schemes and substitution models.

Subset	Best Model	Subset Partitions	Subset Sites
1	GTR+I+G	16S	1-441
2	GTR+I+G	Argk exon 1, EF1 α exon 1	442-1040\3, 1769-2297\3
3	GTR+I+G	EF1 α exon2, Opsin exon2, Argk exon2	1770-2297\3, 2384-2883\3, 443-1040\3
4	SYM+G	Argk exon3, Opsin exon3, PEPCK exon	444-1040\3, 2385-2883\3, 2987-3365\3
5	GTR+G	Opsin intron, PEPCK intron, Argk intron	2884-2984, 3366-3805, 1041-1288
6	GTR+I+G	COI1	1289-1768\3
7	HKY+I+G	COI2	1290-1768\3
8	GTR+I+G	COI3	1291-1768\3
9	TVM+G	EF1 α exon3	1771-2297\3
10	TVM+G	EF1 α intron	2298-2382
11	K81UF+I+G	PEPCK exon1, Opsin exon1	2985-3365\3, 2383-2883\3
12	TRNEF+I+G	PEPCK exon2	2986-3365\3

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