

# Supporting Information

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**Table S1.** List of ancestor/descendant relationships where the range overlap falls within our proposed estimate of total error

Ancestor	Descendant	Ancestor LAD	Descendant FAD	Range overlap	Potential speciation mechanism
Paleogene					
<i>Hedbergella holmdelensis</i>	<i>Globanomalina archeocompressa</i> *	64.9	64.95	0.05	One potential anagenetic event
<i>Hedbergella monmouthensis</i>	<i>Praemurica taurica</i> *	64.85	64.9	0.05	
<i>H. monmouthensis</i>	<i>Eoglobigerina eobulloides</i> *	64.85	64.97	0.12	Two cladogenetic events and one potential anagenetic event <sup>†</sup>
<i>H. monmouthensis</i>	<i>Parasubbotina aff_pseudobulloides</i> *	64.85	65	0.15	
<i>Parasubbotina aff_pseudobulloides</i>	<i>Parasubbotina pseudobulloides</i> *	64.8	64.9	0.1	One potential anagenetic event
<i>Praemurica taurica</i>	<i>Praemurica psuedoinconstans</i>	63.8	64.8	1.0	One potential anagenetic event
<i>Eoglobigerina eobulloides</i>	<i>Subbotina trivialis</i>	63.6	64.92	1.32	One cladogenetic event <sup>†</sup> ( <i>E. eobulloides</i> – <i>S. trivialis</i> )
<i>E. eobulloides</i>	<i>Eoglobigerina edita</i>	63.6	64.9	1.3	(1) and one potential anagenetic event ( <i>E. eobulloides</i> – <i>E. edita</i> )
<i>Globanomalina archeocompressa</i>	<i>Globanomalina planocompressa</i>	62.8	64.9	2.1	One cladogenetic event, one potential anagenetic event <sup>†</sup>
<i>G. archeocompressa</i>	<i>Globanomalina compressa</i> *	62.8	62.9	0.1	
<i>Globanomalina planocompressa</i>	<i>Globanomalina imitata</i> *	62	62.05	0.05	One potential anagenetic event
<i>Praemurica psuedoinconstans</i>	<i>Praemurica inconstans</i>	61.15	62.8	1.65	One potential anagenetic event
<i>Eoglobigerina edita</i>	<i>Eoglobigerina spiralis</i> *	61.15	61.2	0.05	One potential anagenetic event
<i>Subbotina trivialis</i>	<i>Subbotina triangularis</i> *	61	61.1	0.1	One cladogenetic event and one potential anagenetic <sup>†</sup>
<i>S. trivialis</i>	<i>Subbotina cancellata</i> *	61	61.2	0.2	
<i>Praemurica inconstans</i>	<i>Praemurica uncinata</i> *	60.9	61.2	0.3	One potential anagenetic event
<i>Praemurica uncinata</i>	<i>Praemurica lozanoi</i> *	60.7	60.75	0.05	Two cladogenetic events <sup>†</sup>
<i>P. uncinata</i>	<i>Igorina pusilla</i> *	60.7	60.95	0.25	( <i>P. uncinata</i> – <i>P. lozanoi</i> ;
<i>P. uncinata</i>	<i>Morozovella praeangulata</i> *	60.7	61.15	0.45	<i>P. uncinata</i> – <i>P. pusilla</i> ) and one putative anagenetic event <sup>‡</sup> ( <i>P. uncinata</i> – <i>M. praeangulata</i> ) (2)
<i>Globanomalina compressa</i>	<i>Globanomalina ehrenbergi</i> *	60.7	61.1	0.4	One potential anagenetic event
<i>Morozovella praeangulata</i>	<i>Acarinina strabocella</i> *	60	60.8	0.8	One cladogenetic event <sup>†</sup> ( <i>M. praeangulata</i> – <i>A. strabocella</i> )
<i>M. praeangulata</i>	<i>Morozovella angulata</i>	60	60.95	0.95	and one putative anagenetic event <sup>‡</sup> ( <i>M. praeangulata</i> – <i>M. angulata</i> ) (2)
<i>Igorina pusilla</i>	<i>Igorina tadjikistanensis</i> *	59.5	60	0.5	One cladogenetic event and one potential anagenetic event <sup>†</sup>
<i>I. pusilla</i>	<i>Igorina albeari</i> *	59.5	60.05	0.55	
<i>Globanomalina ehrenbergi</i>	<i>Globanomalina pseudomenardii</i> *	59.38	59.4	0.02	One cladogenetic event and one potential anagenetic event <sup>†</sup>
<i>G. ehrenbergi</i>	<i>Globanomalina chapmani</i> *	59.38	59.45	0.07	

Table S1. Cont.

Ancestor	Descendant	Ancestor LAD	Descendant FAD	Range overlap	Potential speciation mechanism
<i>Acarinina strabocella</i>	<i>Acarinina nitida</i> *	59.3	59.5	0.2	One putative anagenetic event <sup>§</sup> (3)
<i>Morozovella conicotruncata</i>	<i>Morozovella velascoensis</i> *	59.3	59.9	0.6	One potential anagenetic event
<i>Morozovella angulata</i>	<i>Morozovella apantesma</i> *	59.3	60	0.7	One cladogenetic event <sup>†</sup>
<i>M. angulata</i>	<i>Morozovella conicotruncata</i>	59.3	60.8	1.5	( <i>M. angulata</i> – <i>M. apantesma</i> ) (4) and one putative anagenetic event <sup>§</sup> ( <i>M. angulata</i> – <i>M. conicotruncata</i> ) (2)
<i>Subbotina cancellata</i>	<i>Subbotina patagonica</i> *	56.8	57	0.2	One cladogenetic event <sup>†</sup>
<i>S. cancellata</i>	<i>Subbotina velascoensis</i>	56.8	59.2	2.4	( <i>S. cancellata</i> – <i>S. velascoensis</i> ) (1) and one potential anagenetic event ( <i>S. cancellata</i> – <i>S. patagonica</i> )
<i>Acarinina mckannai</i>	<i>Acarinina soldadoensis</i> *	56.3	56.4	0.1	One putative anagenetic event <sup>§</sup> (5)
<i>Acarinina nitida</i>	<i>Acarinina esnaensis</i> *	56.2	56.3	0.1	One cladogenetic event and one potential anagenetic event <sup>†</sup>
<i>A. nitida</i>	<i>Acarinina coalingensis</i> *	56.2	56.5	0.3	
<i>Globanomalina imitata</i>	<i>Globanomalina australiformis</i> *	55.8	55.88	0.08	One cladogenetic event and one potential anagenetic event <sup>†</sup>
<i>G. imitata</i>	<i>Globanomalina ovalis</i> *	55.8	56.5	0.7	
<i>Subbotina triangularis</i>	<i>Subbotina roesnaesensis</i> *	55.55	55.6	0.05	One potential anagenetic event
<i>Igorina tadjikistanensis</i>	<i>Igorina lodoensis</i> *	55.4	55.8	0.4	One cladogenetic event <sup>†</sup> (6)
<i>Acarinina sibaiyaensis</i>	<i>Acarinina africana</i> *	55.3	55.4	0.1	One potential anagenetic event
<i>Globanomalina ovalis</i>	<i>Globanomalina luxorensis</i> *	55.3	55.7	0.4	One potential anagenetic event
<i>Morozovella velascoensis</i>	<i>Morozovella edgari</i> *	54.45	54.9	0.45	One cladogenetic event <sup>†</sup> (2, 7)
<i>Morozovella apantesma</i>	<i>Morozovella aequa</i>	54.45	56.5	2.05	One cladogenetic event
<i>M. apantesma</i>	<i>Planorotalites pseudoscitula</i>	54.45	55.9	1.45	( <i>M. apantesma</i> – <i>P. pseudoscitula</i> ) <sup>†</sup> (1) and one potential anagenetic event ( <i>M. apantesma</i> – <i>M. aequa</i> )
<i>Subbotina velascoensis</i>	<i>Subbotina hornibrooki</i>	54.45	55.6	1.15	One cladogenetic event <sup>†</sup> (1)
<i>Morozovella velascoensis</i>	<i>Morozovella allisonensis</i>	54.45	55.5	1.05	One cladogenetic event <sup>†</sup> (1)
<i>Globanomalina luxorensis</i>	<i>Pseudohastigerina wilcoxensis</i>	54.45	55.3	0.85	One potential anagenetic event
<i>Globanomalina planoconica</i>	<i>Planoglobanomalina pseudoalgeriana</i> *	50.6	50.8	0.2	One potential anagenetic event
<i>Morozovella lensiformis</i>	<i>Morozovella aragonensis</i>	50.6	52.3	1.7	One potential anagenetic event
<i>Acarinina interposita</i>	<i>Acarinina pentacamerata</i>	50.4	52.3	1.9	One potential anagenetic event
<i>Acarinina coalingensis</i>	<i>Acarinina primitiva</i> *	50.1	50.8	0.7	One potential anagenetic event
<i>Acarinina pseudotopilensis</i>	<i>Acarinina mcgowrani</i>	47.25	49.25	2.0	One potential anagenetic event
<i>Parasubbotina inaequispira</i>	<i>Guembeltrioides nuttalli</i>	45.4	46.4	1.0	One cladogenetic event <sup>†</sup> (1)
<i>Clavigerinella caucasica</i>	<i>Hantkenina singanoae</i> *	44.45	44.5	0.05	One cladogenetic event <sup>†</sup> (8)
<i>Hantkenina singanoae</i>	<i>Hantkenina mexicana</i> *	44.4	44.41	0.01	One potential anagenetic event
<i>Hantkenina mexicana</i>	<i>Hantkenina liebusi</i> *	43.8	44.2	0.4	One potential anagenetic event
<i>Igorina broedermanni</i>	<i>Igorina anapetes</i>	43.7	45.4	1.7	One putative anagenetic <sup>§</sup> (6)
<i>Subbotina roesnaesensis</i>	<i>Globigerina officinalis</i> *	43.2	43.5	0.3	One potential anagenetic event
<i>Globoturborotalita bassriverensis</i>	<i>Globoturborotalita ouachitaensis</i> *	42.4	43.1	0.7	Two cladogenetic events ( <i>G. bassriverensis</i> – <i>G. martini</i> ; <i>G. bassriverensis</i> – <i>Turborotalita carcoselleensis</i> ) <sup>†</sup>
<i>G. bassriverensis</i>	<i>Globoturborotalita martini</i>	42.4	44	1.6	
<i>G. bassriverensis</i>	<i>Turborotalita carcoselleensis</i>	42.4	44.4	2.0	(1) and one potential anagenetic event
<i>Acarinina pseudosphaerica</i>	<i>Acarinina echinata</i>	42.3	43.2	0.9	One cladogenetic event <sup>†</sup> (1)

Table S1. Cont.

Ancestor	Descendant	Ancestor LAD	Descendant FAD	Range overlap	Potential speciation mechanism
<i>Globigerinatheka curryi</i>	<i>Globigerinatheka euganea</i>	41.6	42.8	1.2	One potential anagenetic event
<i>Turborotalia frontosa</i>	<i>Turborotalia pomeroli</i>	40.8	43.2	2.4	One potential anagenetic event
<i>Globigerinatheka euganea</i>	<i>Globigerinatheka luterbacheri*</i>	40.2	40.3	0.1	One cladogenetic event and one potential anagenetic event <sup>†</sup>
<i>G. euganea</i>	<i>Orbulinoides beckmanni*</i>	40.2	40.5	0.3	
<i>Hantkenina dumblei</i>	<i>Hantkenina compressa</i>	39	41.2	2.2	One potential anagenetic event
<i>Acarinina primitiva</i>	<i>Dentoglobigerina galavisi*</i>	38.8	39	0.2	One potential anagenetic event
<i>Turborotalita carcoselleensis</i>	<i>Turborotalita praequinqueloba*</i>	35.9	36	0.1	One potential anagenetic event
<i>Turborotalia cocoaensis</i>	<i>Turborotalia cunialensis*</i>	33.8	34.1	0.3	One potential anagenetic event
<i>Subbotina linaperta</i>	<i>Subbotina utilizindex</i>	33.7	35.4	1.7	One potential anagenetic event
<i>Dentoglobigerina sp.</i>	<i>Dentoglobigerina prasaepis*</i>	32.9	33.3	0.4	One cladogenetic event, one potential anagenetic event <sup>†</sup>
<i>Dentoglobigerina sp.</i>	<i>Dentoglobigerina tapuriensis*</i>	32.9	33.7	0.8	
<i>Subbotina sp. 1</i>	<i>Subbotina sp. 2</i>	32.6	33.8	1.2	One cladogenetic event <sup>‡</sup> (1)
<i>Subbotina eocaena</i>	<i>Subbotina sp. 1</i>	32.2	34.1	1.9	One cladogenetic event <sup>‡</sup> (1)
<i>Globoturborotalita martini</i>	<i>Globoturborotalita labiacrassata*</i>	30.3	30.4	0.1	One cladogenetic event, one potential anagenetic event <sup>†</sup>
<i>G. martini</i>	<i>Globoturborotalita woodi*</i>	30.3	30.6	0.3	
<i>Turborotalia ampliapertura</i>	<i>Turborotalia euapertura</i>	30.3	32.65	2.35	One potential anagenetic event
<i>Dentoglobigerina galavisi</i>	<i>Dentoglobigerina larmeu*</i>	27.2	27.3	0.1	One potential anagenetic event
<i>Dentoglobigerina sellii</i>	<i>Dentoglobigerina binaiensis</i>	24.1	26	1.9	One cladogenetic event <sup>‡</sup> (1)
<i>Protentelloides primitiva</i>	<i>Protentelloides dalhousiei</i>	23.8	25.7	1.9	One potential anagenetic event
<i>Paragloborotalia pseudokugleri</i>	<i>Paragloborotalia kugleri*</i>	23.5	23.8	0.3	One potential anagenetic event
Neogene					
<i>Paragloborotalia kugleri</i>	<i>Fohsella peripheroronda</i>	21.5	22	0.5	One cladogenetic event <sup>‡</sup> (1)
<i>Catapsydrax parvulus</i>	<i>Catapsydrax stainforthi*</i>	16.9	17.1	0.2	One potential anagenetic event
<i>Praeorbulina circularis</i>	<i>Orbulina suturalis*</i>	14.9	15.1	0.2	One potential anagenetic event
<i>Hirsutella praescitula</i>	<i>Hirsutella scitula*</i>	14.8	15	0.2	One cladogenetic event, one potential anagenetic event <sup>†</sup>
<i>H. praescitula</i>	<i>Hirsutella challenger</i>	14.8	15.15	0.35	
<i>Menardella archeomenardii</i>	<i>Menardella praemenardii</i>	14.2	14.6	0.4	One cladogenetic event <sup>‡</sup> (1)
<i>Fohsella peripheroronda</i>	<i>Fohsella peripheroacuta</i>	13.8	14.3	0.5	One potential anagenetic event
<i>F. peripheroacuta</i>	<i>Fohsella praefohsi</i>	13.4	13.8	0.4	One cladogenetic event <sup>‡</sup>
<i>F. peripheroacuta</i>	<i>Fohsella linguaensis</i>	13.4	13.7	0.3	( <i>F. peripheroacuta</i> – <i>F. linguaensis</i> ) (1); one potential anagenetic event ( <i>F. peripheroacuta</i> – <i>F. praefohsi</i> )
<i>Fohsella praefohsi</i>	<i>Fohsella fohsi</i>	13.1	13.4	0.3	One cladogenetic event <sup>‡</sup> (9)
<i>Menardella praemenardii</i>	<i>Menardella menardii*</i>	12.3	12.31	0.01	One potential anagenetic event
<i>Hirsutella primitiva</i>	<i>Hirsutella praemargaritae*</i>	5.7	5.9	0.2	One potential anagenetic event
<i>Globorotalia merotumida</i>	<i>Globorotalia plesiotumida</i>	5.8	6.4	0.6	One potential anagenetic event
<i>Globoconella conoidea</i>	<i>Globoconella conomiozea</i>	5.7	6	0.3	One cladogenetic event <sup>‡</sup> (10)
<i>Globoconella conomiozea</i>	<i>Globoconella terminalis*</i>	5.5	5.6	0.1	One cladogenetic event <sup>‡</sup> (10)
<i>Globoconella terminalis</i>	<i>Globoconella pliozea*</i>	5	5.05	0.05	One cladogenetic event <sup>‡</sup>
<i>G. terminalis</i>	<i>Globoconella sphericomiozea</i>	5	5.4	0.4	( <i>G. terminalis</i> – <i>G. pliozea</i> ) (11) and one putative anagenetic event <sup>§</sup> ( <i>G. terminalis</i> – <i>G. sphericomiozea</i> ) (11)

Table S1. Cont.

Ancestor	Descendant	Ancestor LAD	Descendant FAD	Range overlap	Potential speciation mechanism
<i>Globoconella sphericomiozea</i>	<i>Globoconella puncticulata</i> *	4.5	4.6	0.1	One cladogenetic event <sup>‡</sup> (10)
<i>Pulleniatina praespectabilis</i>	<i>Pulleniatina spectabilis</i> *	4.4	4.6	0.2	One potential anagenetic event
<i>Pulleniatina spectabilis</i>	<i>Pulleniatina finalis</i> *	4.2	4.25	0.05	One potential anagenetic event
<i>Hirsutella margaritae</i>	<i>Hirsutella hirsuta</i> *	2.95	3	0.05	One cladogenetic event, one potential anagenetic event <sup>†</sup>
<i>H. margaritae</i>	<i>Hirsutella theyeri</i>	2.95	3.5	0.55	
<i>Globoconella puncticulata</i>	<i>Globoconella inflata</i>	2.3	2.6	0.3	One cladogenetic event <sup>‡</sup> (10)
<i>Menardella menardii</i>	<i>Menardella fimbriata</i> *	0	0.2	0.2	One cladogenetic event <sup>¶</sup>
<i>Beella digitata</i>	<i>Beella megastoma</i>	0	0.3	0.3	One cladogenetic event <sup>¶</sup>

The listed events represent ancestor/descendant transitions where range overlap is within our estimate of total-error margins (2.4 Ma for the Paleogene; 0.6 Ma for the Neogene). Where the range of a morphospecies crosses the Paleogene/Neogene boundary, we apply our Paleogene estimate for total error to the FAD ( $\pm 1.2$  Ma) and our Neogene estimate for the error to the LAD ( $\pm 0.3$  Ma). Events where overlap is equal to or less than dating-error margins (0.8 Ma for the Paleogene; 0.2 Ma for the Neogene) are marked with an \*. For each listed event, we assume potential anagenesis, except where available evidence indicates putative anagenesis or cladogenesis. Evidence used includes:

<sup>†</sup>Two or more descendants originated from a single ancestor, indicating one potential anagenetic event and multiple cladogenetic events.

<sup>‡</sup>Frequency distributions of ancestor and descendant morphotypes are nonoverlapping in morphospace, or are overlapping in morphospace, but distinguishable, indicating cladogenesis (1, 11, 12). For events where ancestor/descendant overlap falls within dating error, we invoke ‡ only where co-occurring specimens can be demonstrated to constitute two or more populations with distinguishable frequency-distributions in morphospace (e.g. ref. 12). We use ancestor/descendant transitions designated speciation events by Aze et al. (1) to resolve typological error in light of their statement that empty morphospace between populations in multidimensional space is necessary for a speciation event to have occurred.

<sup>§</sup>Ancestor–descendant pairs cannot be differentiated in morphospace, indicating putative anagenesis.

<sup>¶</sup>Ancestor and descendant are both extant and therefore contemporaneous, indicating cladogenesis.

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