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DATA REPOSITORY

Appendix 1. Taxonomic list of nannofossils and foraminifera used to define the biozones of the BS.

Table S1. Table S1 reports the values of the principal component analysis (*Kirschvink*, 1980), of the ChRM of every sample. Samples are divided in classes of quality as described in the paragraph 3.1.2 of the main text.

Table S2. The table represents the quantitative data of calcareous nannofossils.

Supporting Information for

High-resolution integrated magnetobiostratigraphy of a new middle Eocene section from the western branch of the Neo-Tethys (Elazığ Basin, central Turkey) and its tectonic significance

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The supporting information includes datasets of the characteristic remanent magnetization (ChRM), quantitative analysis of calcareous nannofossils, and a taxonomic list of nannofossils and

foraminifera. The supplementary material is meant to complete the information provided by the main text.

Appendix 1

Taxonomic list of nannofossils and foraminifera used to define the biozones of the BS.

Taxonomic list.

Blackites
B. gladius (Locker 1972) Varol 1989
Calcidiscus Kamptner 1950
C. protoannulus (Gartner 1971) Loeblich & Tappan 1978
Chiasmolithus Hay Mohler and Wade 1966
C. gigas (Bramlette and Sullivan 1961) Radomski 1968
C. grandis (Bramlette and Riedel 1954) Radomski 1968
C. oamaruensis (Deflandre 1954) Hay Mohler and Wade 1966
C. solitus (Bramlette and Sullivan 1961) Locker 1968
C. titus Gartner 1970
Cribrocentrum Perch-Nielsen 1971
C. erbae Fornaciari in Fornaciari et al. (2010)
C. reticulatum (Gartner and Smith 1967) Perch-Nielsen 1971
Coccolithus Schwarz 1894
C. eopelagicus (Bramlette and Riedel 1954) Bramlette and Sullivan 1961
C. pelagicus Schiller 1930
Cyclicargolithus Bukry 1971
C. floridanus (Roth and Hay in Hay et al. 1967) Bukry 1971
Dictyococcites Black 1967
D. bisectus (Hay Mohler and Wade 1966) Bramlette and Wilcoxon 1967
D. scrippsae Bukry and Percival 1971
Discoaster Tan Sin Hok 1927
D. barbadiensis Tan Sin Hok 1927
D. bifax Bukry 1971
D. deflandrei Bramlette and Riedel 1954
D. nodifer (Bramlette and Riedel 1954) Bukry 1973
D. saipanensis Bramlette and Riedel 1954

D. tanii Bramlette and Riedel 1954
Ericsonia Black 1964
E. formosa (Kamptner 1963) Haq 1971
E. obruta Perch-Nielsen 1971
Helicosphaera Kamptner 1954
H. bramlettei Muller 1970
H. compacta Bramlette and Wilcoxon 1967
H. euphratis Haq 1966
H. lophota Bramlette and Sullivan 1961
H. papillata Bukry and Bramlette 1969
H. reticulata Bramlette and Wilcoxon 1967
H. salebrosa Perch-Nielsen 1971
H. seminulum Bramlette and Sullivan 1961
H. wilcoxonii Gartner 1971
Lanternithus Stradner 1962
L. minutus Stradner 1962
Nannotetrina Achuthan and Stradner 1969
N. cristata (Martini 1958) Perch-Nielsen 1971
Pontosphaera Lohmann 1902
Pseudotriquetrorhabdulus Wise in Wise and Constans 1976
P. inversus (Bukry and Bramlette 1969) Wise in Wise and Constans 1976
Reticulofenestra Hay Mohler and Wade 1966
R. daviesii ((Haq, 1968) Haq (1971)
R. umbilicus (Levin 1965) Martini and Ritzkowski 1968
Sphenolithus Deflandre in Grassé 1952
S. furcatolithoides Locker 1967
S. obtusus Bukry1971
S. predistentus Bramlette and Wilcoxon 1967
S. radians Deflandre in Grassé 1952
S. richteri Bown and Dunkley Jones, 2012
S. runus Bown & Dunkley Jones, 2006
S. spiniger Bukry 1971
Zygrhablithus Deflandre 1959
Z. bijugatus (Deflandre in Deflandre and Fert 1954) Deflandre 1959

Discocyclina pratti (Michelin 1846)
D. trabayensis (Neumann 1955)
*D. augustae (*van der Weijden 1940)
D. radians (d'Archiac 1850)
D. dispansa (Sowerby 1840)
D. pulcra (Checchia-Rispoli 1909)
Orbitoclypeus douvillei (Schlumberger 1903)

O. haynesi (Samanta & Lahiri 1985)
O. varians (Kaufmann 1867)
O. zitteli (Checchia-Rispoli 1908)
O. baskilensis sp. nov.
Asterocyclina stellata (d'Archiac 1846)
A. stella (Gümbel 1861)
A. sireli Özcan & Less 2006)
A. kecskemetii (Less 1987)
A. alticostata (Nuttall 1926)
Nummulites garganicus (Tellini 1890)
*N. hormoensis (*Nuttall & Brighton 1931)
N. perforatus (de Montfort 1808) -*biedai* Schaub 1962)
N. maximus (d'Archiac 1850)
*N. alponensis (*Schaub 1981)
N. gizehensis (Forskâl 1775) -*lyelli* (d'Archiac & Haime 1853)
N. ptukhiani (Kacharava 1969)
N. striatus (Bruguière 1792)
N. biarritzensis (d'Archiac & Haime 1853)
N. anomalus (de la Harpe 1877)
Heterostegina armenica (Grigoryan 1986)
Operculina ex. gr. *gomezi (*Colom et Bauzá 1950)
Assilina schwageri (Silvestri 1928)
A. exponens (Sowerby 1840)
Chapmanina gassinensis (Silvestri 1905)
Silvestriella tetrahedra (Gümbel 1868)

Table S1

Table S1 reports the values of the principal component analysis (*Kirschvink*, 1980), of the ChRM of every sample. Samples are divided in classes of quality as described in the paragraph 3.1.2 of the main text.

msl	NRM intensity (A/m)	dec. (°)	inc. (°)	MAD (°)
357.00	3.37E-04	160.80	42.20	3.40
354.00	8.36E-04	135.80	40.30	2.40
352.00	6.46E-04	190.60	15.00	1.20
348.00	8.56E-04	190.90	19.30	1.80
345.00	1.03E-03	197.00	33.30	1.50
341.00	1.46E-03	129.00	20.90	1.90
339.00	4.75E-04	278.00	29.70	6.00
335.00	4.86E-04	265.60	39.00	2.50
332.00	5.24E-04	192.20	48.60	1.90
329.00	8.18E-04	32.20	43.80	3.70
326.00	5.78E-04	184.20	57.70	4.30
324.00	4.13E-04	213.10	32.40	2.50
322.00	3.24E-04	280.20	42.50	2.70
321.00	2.36E-04	193.30	-40.00	8.30
320.00	2.45E-04	205.30	-13.00	7.80
319.00	1.97E-04	197.50	-39.10	2.30
317.00	4.79E-04	201.30	-31.20	4.80
315.00	5.27E-04	177.20	-20.70	3.40
312.00	1.13E-03	221.70	26.90	1.80
311.00	1.64E-04	199.30	42.00	2.90
310.00	4.59E-04	237.80	30.70	1.70
308.00	3.52E-04	260.20	27.70	1.80
306.00	4.36E-04	210.90	47.70	1.70
304.00	4.75E-04	204.20	-17.80	7.20
303.00	2.46E-04	215.10	-58.50	6.70
302.00	4.61E-04	183.60	-36.00	3.50
300.00	2.88E-04	183.40	52.90	9.00
299.00	6.03E-04	187.50	39.80	2.70
298.00	3.19E-04	179.60	57.40	10.50
297.00	3.06E-04	159.60	20.90	5.60
296.00	2.57E-04	208.10	43.00	2.20
294.00	4.68E-04	195.30	41.90	4.00
292.00	2.96E-04	233.50	17.70	4.60
291.00	3.99E-04	236.90	32.50	1.90

289.00	6.27E-04	138.30	56.70	6.40
288.00	8.70E-04	134.80	32.80	7.50
287.00	5.58E-04	146.10	-48.00	5.90
286.00	9.59E-04	37.80	-20.20	2.90
285.00	1.47E-03	216.20	-42.70	4.80
284.00	5.87E-04	254.30	-41.20	7.30
283.00	1.03E-03	248.40	-69.50	5.20
281.00	1.74E-04	256.20	-73.40	9.90
280.00	4.79E-04	219.50	9.80	7.00
279.00	4.08E-04	172.00	18.40	9.90
277.00	3.60E-04	162.10	-28.00	4.40
276.00	3.28E-04	116.80	-53.40	1.70
275.20	1.92E-04	228.80	-26.00	4.00
272.00	7.00E-04	160.30	-37.20	1.20
270.00	2.72E-04	215.00	-12.20	6.80
267.00	9.93E-04	152.40	41.70	2.60
263.00	6.04E-04	226.60	50.90	6.70
262.00	4.35E-04	249.50	24.00	6.90
261.00	3.35E-04	238.50	10.30	7.70
259.00	7.71E-04	179.90	48.50	2.50
258.00	2.11E-04	139.40	20.20	8.40
257.00	5.15E-04	206.20	43.10	9.40
256.00	5.78E-04	153.20	20.70	5.60
254.00	5.13E-04	181.00	53.70	8.50
253.00	1.81E-04	132.60	-30.40	8.20
252.00	3.59E-04	283.50	-26.20	2.30
250.50	4.64E-04	308.80	16.10	5.90
250.00	6.85E-04	187.40	41.70	3.00
249.00	6.26E-04	224.10	18.00	2.20
248.00	3.82E-04	160.10	-41.30	4.50
247.00	4.75E-04	172.80	-10.00	6.80
246.00	8.27E-04	223.60	44.00	6.10
245.00	7.75E-04	121.40	22.70	2.40
243.00	4.31E-04	254.50	19.10	6.30
242.00	1.75E-04	113.00	12.90	22.60
241.00	3.11E-04	188.60	33.40	4.80
240.00	2.36E-04	168.00	38.30	8.90
239.00	3.71E-04	164.50	58.80	4.70
237.00	1.63E-04	203.30	32.80	2.70
236.00	1.22E-04	145.00	59.40	8.70
226.00	4.16E-04	149.70	30.40	4.70
225.00	4.35E-04	185.10	31.60	4.30
222.00	3.00E-04	217.10	41.50	6.80

221.00	6.27E-04	341.00	42.60	6.90
220.00	1.14E-03	219.60	28.40	3.70
219.00	5.62E-04	237.50	19.60	8.20
218.00	4.47E-04	181.30	-12.40	6.80
217.00	1.01E-03	223.90	-14.70	8.30
216.00	4.65E-04	224.30	61.00	5.10
215.00	3.93E-04	135.90	33.60	9.60
214.00	1.00E-03	222.40	14.50	7.00
213.00	6.70E-04	107.00	31.70	6.10
212.00	1.14E-03	255.50	39.10	3.40
211.00	6.48E-04	242.40	58.20	2.50
210.00	2.94E-03	136.00	14.60	4.50
209.00	1.71E-03	135.70	18.40	3.40
207.00	4.63E-04	177.40	8.90	4.90
205.00	4.62E-04	200.60	44.60	2.20
204.50	1.55E-03	141.70	41.60	3.60
204.00	3.82E-04	243.30	22.80	4.40
203.50	2.62E-03	149.40	39.40	2.30
201.00	4.13E-04	202.80	48.80	3.40
200.00	3.90E-04	191.90	12.20	9.10
199.00	2.54E-04	303.10	-41.30	3.50
197.00	3.10E-04	156.40	-32.50	10.90
196.00	2.45E-04	227.00	-27.90	2.70
194.00	1.66E-04	157.70	-19.70	4.60
193.00	2.70E-04	251.70	-22.10	9.80
192.00	5.26E-04	247.10	67.30	2.80
189.50	3.17E-04	238.00	11.70	3.40
188.00	2.49E-04	204.70	46.00	7.50
186.00	6.23E-04	200.60	27.70	8.80
183.00	3.29E-04	191.40	39.10	9.80
182.00	2.94E-04	309.30	73.70	5.40
179.00	3.60E-04	212.10	41.30	9.70
177.00	2.44E-04	203.30	20.60	8.80
175.00	1.95E-04	189.60	34.20	9.00
173.00	2.80E-04	195.90	32.00	8.50
172.00	1.20E-04	214.10	41.20	10.40
167.00	2.58E-04	41.20	44.20	5.00
164.00	2.79E-04	126.40	35.80	5.80
161.00	4.90E-04	159.00	8.70	2.10
157.00	1.11E-03	177.50	30.30	10.50
156.00	2.23E-04	107.40	46.60	9.80
154.00	1.79E-04	117.20	14.00	7.70
153.00	1.53E-03	185.20	17.00	7.80

152.00	4.19E-04	215.20	-30.50	4.70
151.00	2.55E-04	51.60	-49.40	1.70
148.00	1.55E-04	230.10	-43.70	8.30
147.00	4.08E-04	173.90	-16.20	9.80
138.00	2.84E-04	258.90	-44.60	4.90
133.00	1.51E-04	147.90	-49.40	8.80
123.00	2.73E-04	226.80	-50.60	9.20
112.00	5.61E-04	320.60	-40.20	8.30
103.00	3.08E-04	326.70	-17.00	2.50
93.00	3.48E-04	189.30	-51.20	9.70
85.00	2.42E-04	217.60	-26.33	6.23
85.00	2.42E-04	217.60	-26.33	6.23
70.00	2.76E-04	213.80	-46.70	6.60
69.50	5.44E-04	148.30	-40.60	9.10
69.00	7.35E-05	50.30	-31.80	10.40
68.00	1.38E-03	223.00	-45.00	8.00
66.00	1.42E-03	211.20	-40.30	10.70
65.00	4.68E-04	238.70	-43.50	4.90
65.00	8.02E-04	188.10	-47.60	7.70
63.50	6.62E-04	209.20	-58.80	7.40
61.00	5.10E-04	247.20	-45.20	7.50
54.00	6.02E-04	355.70	59.50	5.80
53.00	2.83E-04	277.20	42.00	8.90
52.00	1.64E-04	120.50	43.00	11.50
51.00	5.65E-04	247.40	19.70	10.40
50.90	1.61E-03	154.80	38.60	1.50
50.00	6.57E-04	321.70	50.60	11.40
47.00	5.35E-04	161.50	49.30	10.20
43.50	4.89E-04	144.00	42.00	8.00
42.00	1.25E-03	200.70	-36.80	8.80
39.00	1.11E-03	189.60	-53.50	9.60
38.00	6.66E-04	95.30	-49.40	10.90
36.00	7.72E-04	236.90	-30.60	8.90
34.00	8.56E-04	174.10	-18.40	9.90
33.00	5.75E-04	174.10	-21.10	7.10
31.00	5.76E-04	212.50	-34.90	8.60
27.00	2.09E-03	86.30	-40.40	9.70
26.00	7.39E-04	179.30	-59.80	6.80
25.00	4.76E-04	209.90	-29.90	6.80
24.00	6.07E-04	175.40	-47.80	10.80
23.00	2.92E-04	199.90	-39.90	2.10
21.00	3.84E-03	136.30	-17.00	7.60
19.00	6.38E-04	123.50	-20.40	11.20

15.00	2.26E-03	206.10	-51.90	11.30
13.00	1.23E-03	211.20	-61.50	11.10
12.00	6.50E-04	154.30	-30.20	10.50
9.00	2.67E-03	227.20	-17.30	5.80
8.00	2.79E-03	214.10	-23.30	4.00
7.00	8.89E-04	205.50	-21.20	7.90
5.00	3.37E-03	151.80	-23.70	7.20
4.00	4.07E-03	238.10	-57.90	3.90
3.00	2.36E-02	201.40	-49.30	3.90
2.00	2.42E-03	220.00	-45.40	6.40

bad quality				
311.00	2.05E-04	215.10	-75.50	25.30
295.00	3.19E-04	179.60	57.40	10.50
287.00	3.01E-04	236.40	51.10	12.60
279.00	1.64E-04	278.30	40.00	18.10
278.00	2.82E-04	251.10	49.10	16.30
275.00	2.82E-04	251.10	49.10	10.30
272.00	2.47E-04	122.30	40.90	16.90
252.00	4.57E-04	162.20	33.50	19.50
239.00	1.75E-04	113.00	12.90	22.60
231.00	2.78E-04	227.10	44.40	10.40
230.00	1.46E-04	210.80	35.70	15.30
224.00	5.40E-04	219.60	32.40	11.50
221.00	4.00E-04	100.90	57.20	12.50
194.00	3.10E-04	156.40	58.50	10.90
185.50	1.52E-04	210.20	-17.20	16.10
178.00	2.04E-04	216.50	41.80	10.10
177.00	2.66E-04	225.50	50.20	19.80
169.00	1.20E-04	214.10	41.20	10.40
168.10	1.45E-04	135.50	4.50	17.60
168.00	3.39E-04	219.20	53.50	14.70
167.00	2.01E-04	169.30	44.40	11.60
166.00	2.12E-04	180.60	54.00	13.10
163.00	5.00E-04	173.40	65.30	12.60
162.00	4.87E-04	152.60	-46.80	12.20
159.00	1.75E-03	223.20	-38.00	12.80
157.00	3.78E-04	170.10	38.40	15.40
155.00	3.52E-04	215.10	-29.00	14.70
154.00	1.11E-03	177.50	30.30	10.50

152.00	1.08E-04	166.60	23.60	11.50
147.00	1.40E-04	120.20	-38.60	12.90
143.00	3.81E-04	169.20	54.70	10.50
142.00	7.83E-04	21.20	-68.60	25.50
141.00	1.80E-04	192.10	-28.80	16.80
140.00	2.83E-04	130.90	41.70	17.20
125.00	1.64E-04	135.30	-31.80	11.30
95.00	2.81E-04	157.60	-17.40	10.50
85.00	2.42E-04	217.60	-26.33	6.23
80.00	3.70E-04	165.00	23.40	12.10
77.00	2.01E-04	188.20	-39.20	15.30

outliers

314.00	5.98E-04	201.30	28.80	7.90
298.00	2.67E-04	31.80	-67.50	4.70
292.00	4.45E-04	83.20	-38.80	0.90
271.00	2.53E-04	235.30	38.40	7.00
263.00	4.95E-04	304.50	-49.90	5.30
260.50	8.81E-04	142.30	-70.40	1.90
257.00	5.08E-04	188.90	-46.40	8.70
254.00	4.92E-04	194.30	-60.80	8.00
232.00	1.65E-04	265.90	-36.70	5.10
220.00	4.89E-04	285.00	-14.70	8.70
195.00	3.85E-04	284.00	53.00	6.40
188.00	2.21E-04	239.40	-7.10	2.00
184.00	4.03E-04	218.70	-26.80	6.00
181.00	3.07E-04	230.40	-3.70	6.10
175.00	2.55E-04	214.70	-4.50	5.30
173.00	3.20E-04	149.20	-51.90	9.60
171.00	1.93E-04	143.70	-7.50	6.00
165.00	3.27E-04	171.40	-3.60	2.50
160.00	1.46E-03	340.50	-32.30	5.30
146.00	2.23E-04	149.30	39.00	6.21
146.00	3.81E-04	169.20	54.70	10.50
115.00	2.93E-04	207.00	22.42	8.31
75.00	5.62E-04	125.80	39.60	8.00
32.00	8.71E-04	246.00	34.00	9.20
19.00	6.38E-04	123.50	-20.40	11.20
10.00	4.83E-03	332.70	61.80	4.80
0.00	1.28E-03	142.80	44.80	7.50

low INC

348.00	4.96E-04	261.50	5.30	3.00
270.00	2.79E-04	284.50	-5.60	4.60
265.00	2.99E-04	230.60	-4.90	2.00
262.00	3.64E-04	242.00	0.90	7.00
248.00	4.80E-04	135.00	-5.50	6.30
241.00	5.66E-04	136.80	1.40	9.30
192.00	2.51E-04	221.80	-1.60	7.90
156.00	5.42E-04	153.20	4.20	7.80
302.00	7.11E-04	147.90	3.40	10.60
45.50	3.70E-04	170.10	9.00	3.90
56.00	4.54E-04	140.70	7.10	8.40
55.00	4.74E-04	19.50	7.40	8.40

Table S2

The table represents the quantitative data of calcareous nannofossils.

msl	Total counted	C. protoannula	C. pelagicus C.eopelagicus	C. erbae	C. reticulatum	C. floridanus	D. biesectus	D. scrippsae	Dictyococcites sp.	E. formosa	E. obruta	L. minutus	Reticulofenestra	R. daviesii	R. umbilicus	Z. bijugatus	Sphenolithus sp.	S. furcatolithoides	S. obtusus	S. predistentus	S. radians	S. richteri	S. runus	S. spiniger	S. strigosus	D. barbadiensis	D. bifax	D. deflandrei	Star shaped Discoaster with terminal nodes	D. nodifer	D. saipanensis	D. tanii	Discoaster sp.	Total Discoaster	Ch. gigas	Ch. grandis	Ch. oamaruensis	Ch. solitus	Ch. titus	Ciasmolithus sp.	Total Chiasmolithus	H. bramlettei	H. compacta	H. euphratis	H. lophota	H. papillata	H. reticulata	H. selebrosa	H. seminulum	H. wilcoxonii	Total Helicosphaera	N. cristata	Pontosphaera sp.	P. inversus	B. gladius	Fornaciari et al. 2010	Agnini et al. 2014	Martini 1971	Okada & Bukry 1980	
357.0	300	0.0	18.0	4.6	12.0	16.0	1.0	7.6	0.0	5.0	0.0	21.0	0.0	1.6	0.0	1.6	11.6	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	9	0	0	1	0	2	0	4	16	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	1	2	0	0	0	0					
355.0	300	0.0	8.0	9.0	7.6	19.6	0.3	10.0	0.0	1.6	0.0	19.0	0.0	3.3	0.3	1.3	20.0	0.0	0.0	0.0	1.0	0.0	0.0	0.0	0.0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	1	0	1	1	0	0	0	0	0	0	2	0	1	0	0	MNP18	CNE18			
353.0	300	0.0	16.6	1.6	13.0	17.3	0.3	0.0	0.0	7.0	0.0	26.6	0.0	5.6	0.0	2.0	10.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	18	0	0	0	0	0	0	2	20	0	0	0	0	0	1	1	0	1	0	0	0	0	0	0	0	1	0	0	0	0			NP18	CP15a	
348.0	300	0.0	16.0	17.3	10.6	9.6	2.6	3.0	0.0	9.0	0.0	17.0	0.3	0.3	2.0	2.0	10.3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	26	0	0	0	2	2	3	2	35	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	1	0	1	0	0					C. erbae AE
345.0	300	0.3	17.0	19.0	16.0	12.6	0.0	0.0	0.0	9.6	0.0	6.6	0.0	0.6	3.0	2.3	13.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	16	0	3	1	8	3	14	6	51	0	0	7	0	0	2	9	0	1	3	0	0	0	1	0	2	7	0	7	0	0					
342.0	300	0.3	20.3	18.6	11.6	13.0	1.0	2.3	0.0	3.6	0.0	13.6	0.0	0.0	1.6	0.0	14.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	3	0	0	0	0	3	0	2	8	0	0	2	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	MNP18A	CNE17			Ch. oamarue
339.0	300	0.3	13.6	25.0	15.6	19.0	0.3	1.6	0.0	4.0	0.0	14.0	0.0	0.0	1.0	1.0	4.6	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	31	0	4	0	0	9	2	6	52	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	1	0	3	0	0					
336.0	300	0.0	9.0	35.0	16.6	5.6	0.3	1.6	0.0	4.6	0.0	16.6	0.0	0.0	1.0	2.6	7.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	10	0	2	0	0	3	2	3	20	0	5	0	0	0	0	5	0	0	0	0	0	0	1	0	0	1	0	3	0	0					Ch. grandis H
332.0	300	0.0	13.0	31.0	12.6	11.0	2.0	2.0	0.0	7.6	0.0	8.6	0.0	0.0	1.6	0.3	10.3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	5	0	2	1	5	3	2	3	21	0	4	0	0	0	0	4	0	1	1	0	0	0	0	0	0	2	0	0	0	0					C. erbae AB
329.0	300	0.0	11.6	3.6	24.0	17.0	3.0	9.6	0.0	8.6	0.0	13.0	0.0	0.0	1.0	1.6	7.0	0.0	0.0	0.0	1.0	0.0	0.0	0.0	0.0	8	0	0	0	3	2	0	3	16	0	0	0	0	0	0	0	0	0	1	0	0	0	2	0	0	3	0	1	0	0					
323.0	300	0.6	10.3	0.6	11.3	16.3	9.0	16.0	0.0	3.0	0.0	24.6	0.0	0.0	1.0	2.0	5.0	0.0	0.0	0.0	1.0	0.0	0.0	0.0	0.0	6	0	1	0	2	4	2	3	18	0	1	0	0	0	2	3	0	1	1	0	0	0	1	0	0	3	0	2	0	0					
319.0	300	0.0	5.0	0.3	9.0	21.0	3.0	7.6	0.0	2.6	0.0	34.0	0.0	0.0	0.0	2.0	15.3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0	0	0	0	0	0	0	0	0	0	2	0	0	0	0	2	0	0	1	0	0	0	0	0	1	2	0	0	0	0					
312.0	300	0.0	9.0	1.0	10.0	12.6	3.0	10.0	0.0	6.0	0.0	32.0	0.0	0.0	0.0	3.3	13.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	8	0	0	1	0	2	0	6	17	0	0	0	0	0	0	0	0	0	0	0	0	0	2	0	0	2	0	1	0	0					
307.0	300	0.0	8.6	1.0	9.6	26.6	6.6	13.6	0.0	2.0	0.0	12.3	0.0	0.0	0.0	0.6	19.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0					
303.0	300	1.6	11.0	0.3	15.0	12.0	3.0	7.6	0.0	7.0	0.0	21.0	0.3	0.0	2.6	7.0	11.6	0.0	0.0	1.0	0.0	0.0	0.0	0.0	0.0	38	0	7	1	2	8	0	5	61	0	5	0	0	0	2	7	0	0	1	0	0	0	0	0	2	3	0	0	0	0					
299.0	300	1.0	7.6	0.3	7.6	16.0	3.0	17.6	0.0	3.0	0.0	22.0	0.0	0.0	1.6	1.3	19.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	7	0	2	0	0	2	0	3	14	0	1	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	MNP17B	CNE16	NP17	CP14b	
295.0	300	0.0	8.0	0.3	14.0	26.6	5.6	13.3	0.0	3.6	0.0	13.6	0.0	0.0	2.6	2.3	10.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	7	0	2	0	4	2	0	8	23	0	2	0	0	0	1	3	0	0	0	0	0	0	0	0	0	0	0	1	0	0					
291.0	300	2.6	12.0	0.0	15.0	20.3	3.6	8.6	0.0	7.0	0.0	14.0	0.0	0.0	1.6	5.0	10.3	0.0	0.0	0.0	1.0	0.0	0.0	0.0	0.0	16	0	4	2	1	1	1	1	26	0	2	0	0	0	1	3	0	0	0	0	0	0	0	0	0	0	0	1	0	0					
287.0	300	0.0	10.3	0.0	23.6	15.6	2.6	8.3	0.0	5.6	0.0	23.6	0.0	0.0	0.6	1.6	8.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	12	0	0	0	0	4	0	5	21	0	3	0	0	0	0	3	0	0	0	0	0	0	0	0	0	0	0	1	0	0					
286.0	300	0.0	11.0	0.0	19.0	21.0	3.0	5.0	0.0	3.0	0.0	18.0	0.0	0.0	2.0	4.0	14.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	18	0	4	0	0	11	0	12	45	0	3	0	0	0	0	3	0	1	0	0	0	0	0	0	0	1	0	0	0	0					
284.0	300	0.0	5.0	0.0	18.0	22.3	3.0	10.0	0.0	2.0	0.0	18.0	0.0	0.0	0.0	2.6	19.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	2	0	0	0	0	0	0	5	7	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	1	0	0	0	0					
283.0	300	0.3	10.0	0.0	20.0	23.0	2.0	12.3	0.0	3.0	0.0	18.6	0.0	0.0	2.0	1.6	7.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	24	0	2	1	9	17	3	18	74	0	1	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0					
282.0	100	0.0	8.0	0.0	30.0	31.0	1.0	5.0	0.0	3.0	0.0	5.0	0.0	0.0	1.0	1.0	15.0	0.0	1.0	0.0	0.0	0.0	0.0	0.0	0.0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0					S. obtusus
278.0	300	0.0	6.0	0.0	9.3	38.0	5.0	14.3	0.0	3.5	0.0	9.3	0.0	0.0	2.0	0.0	12.6	0.0	4.0	0.0	0.0	0.0	0.0	0.0	0.0	0	0	0	0	0	0	0	2	2	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	0	2	0	0					
273.0	300	0.0	3.6	0.0	12.6	35.0	6.3	15.6	0.0	1.6	0.0	18.0	0.0	0.0	1.0	1.0	5.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	5	0	2	0	1	3	0	4	15	0	2	0	0	0	0	2	0	0	1	0	0	0	1	0	0	2	0	0	0	0					
269.0	300	0.3	8.0	0.0	12.0	30.0	6.0	13.3	0.0	4.6	0.0	10.0	0.0	0.3	0.0	2.3	13.6	0.0	8.0	1.0	0.0	0.0	0.0	0.0	0.0	16	0	2	0	3	10	0	9	40	0	1	0	1	0	0	2	0	0	0	2	0	0	0	0	1	3	0	0	0	0					Ch. solitus H
264.0	300	0.0	1.3	0.0	7.6	26.3	6.0	19.6	0.0	1.6	0.0	13.0	0.0	0.0	0.0	0.6	24.0	0.0	cf	0.0	0.0	0.0	0.0	0.0	0.0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	1	0	1	0	0					
259.0	300	0.0	4.3	0.0	4.0	29.3	5.6	20.6	0.0	1.6	0.0	6.0	0.0	0.0	0.0	0.6	28.0	0.0	3.0	2.0	1.0	0.0	0.0	0.0	0.0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	2	0	0	0	0	0	0	0	2	0	0	0	0					
254.0	300	0.0	4.6	0.0	4.3	40.6	2.6	16.0	0.0	2.6	0.0	10.3	0.0	0.0	0.3	0.0	19.0	0.0	1.0	0.0	1.0	0.0	0.0	0.0	0.0	4	0	0	0	0	0	0	3	7	0	0	0	0	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0					
249.0	300	0.0	4.6	0.0	13.3	30.3	5.0	12.0	0.0	3.0	0.0	6.3	0.0	0.0	0.0	1.0	24.3	0.0	2.0	1.0	0.0	0.0	0.0	0.0	0.0	1	0	0	0	0	0	0	2	3	0	0	0	0	0	2	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0					
245.0	300	0.0	5.3	0.0	10.6	42.3	4.6	14.3	0.0	3.6	0.0	2.6	0.0	0.0	0.6	1.6	14.3	0.0	2.0	0.0	0.0	0.0	0.0	0.0	0.0	13	0	4	0	0	6	0	5	28	0	6	0	0	0	0	6	0	0	0	0	1	0	0	0	0	1	0	1	0	0	MNP17A				
240.0	300	0.0	4.3	0.0	8.6	40.0	5.0	17.3	0.0	2.6	0.0	3.3	0.0	0.0	0.6	0.3	18.0	0.0	2.0	0.0	0.0	0.0	0.0	0.0	0.0	5	0	1	0	0	1	0	4	11	0	3	0	0	0	0	3	0	0	0	0	0	0	0	0	0	0	0	0	0	0					
230.0	300	0.0	5.0	0.0	18.6	25.0	6.0	14.6	0.0	5.6	0.0	2.3	0.0	0.0	1.0	0.3	21.6	0.0	4.0	0.0	0.0	0.0	0.0	0.0	0.0	8	0	2	0	0	1	1	12	24	0	3	0	0	0	0	3	0	2	0	0	0	0	0	0	0	2	0	2	1	0					
220.0	300	0.0	6.0	0.0	10.6	33.0	6.6	19.3	0.0	4.6	0.0	3.3	0.0	0.0	1.6	0.0	15.0	0.0	3.0	1.0	0.0	0.0	0.0	0.0	0.0	0	0	0	0	0	1	1	5	7	0	5	0	0	0	0	5	0	0	0	0	0	0	1	0	0	1	0	1	0	0					
215.0	300	0.0	4.6	0.0	7.0	44.0	4.6	13.6	0.0	4.3	0.0	2.6	0.0	0.0	1.0	0.6	17.6	0.0	1.0	1.0	0.0	0.0	0.0	0.0	0.0	2	0	1	0	0	0	0	1	4	0	1	0	0	0	0	1	0	1	0	0	0	0	0	0	0	1	0	0	1	0					
210.0	300	0.0	7.0	0.0	8.6	35.0	6.6	15.0	0.0	2.6	0.0	3.3	0.3	0.0	0.6	1.0	20.0	0.0	4.0	1.0	0.0	0.0	0.0	0.0	0.0	1	0	3	0	1	5	0	5	15	0	2	0	0	0	0	2	0	1	0	0	0	0	1	0	0	2	0	1	2	0					
206.0	300	0.0	5.0	0.0	15.3	40.0	3.0	10.6	0.0	1.3	0.0	5.0	0.3	0.0	1.0	0.3	18.0	1.0	1.0	2.0	0.0	0.0	0.0	0.0	0.0	5	0	1	0	0	0	0	5	11	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	1	0	2	0	0					
202.0	300	0.0	6.0	0.0	14.6	39.0	4.6	13.6	0.0	1.6	0.0	2.3	1.0	0.0	1.0	0.3	16.0	0.0	0.0	1.0	0.0	0.0	0.0	0.0	0.0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0					
200.5	300	0.0	14.3	0.0	19.0	38.0	1.0	11.6	0.0	6.0	0.0	1.3	0.0	0.0	2.0	1.6	5.0	0.0	3.0	1.0	1.0	0.0	0.0	0.0	0.0	8	0	0	0	1	3	1	2	15	0	2	0	0	0	0	2	0	0	0	0	1	0	0	0	0	1	0	1	0	0		CNE15			S. obtusus
198.0	300	0.0	18.0	0.0	13.0	46.0	1.0	4.0	0.0	6.0	0.0	0.0	1.0	0.0	2.0	0.0	9.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0					
197.0	300	0.3	13.0	0.0	7.0	52.0	1.0	3.0	0.0	4.6	0.0	1.0	1.0	0.0	5.0	1.0	11.0	0.0	0.0	1.0	0.0	0.0	0.0	0.0	0.0	0	0	0	0	0	0	0	1	1	0	1	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0					
195.0	300	0.0	22.0	0.0	10.3	16.0	2.6	15.0	0.0	18.3	0.0	0.0	2.6	0.0	2.0	0.3	11.0	0.0	0.0	1.0	2.0	0.0	0.0	0.0	0.0	3	0	3	1	7	1	0	17	32	0	10	0	0	0	1	11	0	0	0	0	1	0	0	0	0	1	0	2	0	0					
192.0	300	0.0	13.0	0.0	9.0	21.0	6.0	14.6	0.0	17.0	0.0	1.0	2.0	0.3	1.0	0.0	15.0	0.0	0.0	6.0	2.0	0.0	0.0	0.0	0.0	5	0	7	3	2	0	0	5	22	0	3	0	0	0	0	3	0	0	0	0	1	0	0	0	0	1	0	2	0	0	MNP16Bc				
191.0	300	0.0	14.0	0.0	6.0	37.0	3.0	9.0	0.0	3.0	0.0	3.0	1.0	0.0	1.0	0.0	23.0	0.0	0.0	2.0	0.0	0.0	0.0	0.0	0.0	4	0	0	0	0	2	1	5	12	0	3	0	0	0	0	3	0	0	0	0	0	0	0	0	0	0	0	0	0	0					
190.0	300	0.0	26.6	0.0	8.6	23.6	1.6	7.0	0.0	13.0	0.0	1.0	1.6	0.0	3.0	0.0	14.0	0.0	0.0	2.0	2.0	0.0	0.0	1.0	0.0	2	0	4	1	2	1	1	5	16	0	3	0	1	0	0	4	0	1	0	0	1	0	0	0	0	2	0	1	0	0					S. spiniger H
188.0	300	0.3	24.0	0.0	4.3	19.3	3.6	20.0	0.0	15.0	0.0	0.0	1.6	0.6	2.3	0.0	9.0	0.0	0.0	0.0	0.0	0.0	0.0	4.0	0.0	0	0	0	0	0	0	2	3	5	0	1	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	1	0					
186.5	300	0.0	24.0	0.0	4.0	19.0	0.6	24.0	0.0	12.0	0.0	0.0	3.0	0.3	4.0	0.0	9.0	1.0	0.0	3.0	1.0	0.0	0.0	4.0	0.0	8	0	3	0	5	2	0	3	21	0	3	0	1	0	0	4	0	3	0	0	0	0	0	2	0	5	0	0	0	0					
185.0	300	0.0	21.3	0.0	14.3	26.3	0.6	9.0	0.0	6.0	0.0	2.6	1.6	0.0	1.3	0.0	17.0	0.0	0.0	5.0	0.0	0.0	0.0	3.0	0.0	1	0	3	1	6	4	1	12	28	0	0	0	1	0	0	1	0	1	0	0	0	0	1	0	0	2	0	2	0	0					
180.0	300	0.0	13.6	0.0	15.6	24.6	0.6	9.3	0.0	14.3	0.0	0.0	2.6	0.0	3.0	0.0	16.3	0.0	0.0	3.0	2.0	0.0	0.0	5.0	0.0	6	0	12	3	15	2	3	32	73	0	8	0	1	0	0	9	0	3	0	0	0	0	0	1	4	8	0	3	0	0					
175.0	300	0.0	9.3	0.0	9.0	36.3	2.0	8.0	0.0	6.3	0.0	6.0	3.6	0.0	0.0	1.0	18.3	0.0	0.0	0.0	2.0	0.0	0.0	2.0	0.0	0	0	0	0	2	0	0	3	5	0	8	0	1	0	0	9	0	4	0	0	0	0	0	0	1	5	0	0	0	0	MNP16Bb				
170.0	300	0.0	18.0	0.0	11.6	22.0	0.3	7.0	0.0	14.3	0.0	0.0	4.0	0.0	1.3	0.3	21.0	1.0	0.0	3.0	1.0	0.0	0.0	13.0	0.0	6	0	13	3	15	7	5	26	75	0	7	0	0	0	1	8	0	5	1	0	0	0	2	0	4	12	0	3	0	0					
160.0	300	0.0	12.0	0.0	8.0	32.0	2.6	16.0	0.0	8.0	0.0	6.0	0.6	0.0	0.6	1.0	13.0	0.0	0.0	0.0	1.0	0.0	0.0	3.0	0.0	11	0	9	1	2	8	1	5	37	0	2	0	0	0	1	3	0	0	0	0	0	0	1	0	0	1	0	1	1	0					
157.0	300	0.0	8.6	0.0	22.0	34.0	0.6	10.6	0.0	4.6	0.0	0.0	1.6	0.6	1.0	1.3	15.0	0.0	0.0	1.0	4.0	0.0	0.0	6.0	0.0	8	0	3	0	3	5	0	8	27	0	0	0	1	0	0	1	0	0	0	0	0	0	1	0	0	1	0	0	0	0					
155.0	300	0.0	9.6	0.0	8.6	37.0	2.6	18.0	0.0	4.6	0.0	1.6	3.0	0.0	2.0	1.0	12.0	0.0	0.0	0.0	3.0	0.0	0.0	9.0	0.0	0	0	0	0	1	0	0	0	1	0	1	0	1	0	0	2	0	1	1	0	0	0	2	2	0	6	0	1	0	0			NP16		
154.0	300	0.0	8.0	0.0	6.6	34.6	4.0	19.0	0.0	10.0	0.0	1.6	1.0	0.0	3.0	0.3	12.0	0.0	0.0	0.0	1.0	0.0	0.0	4.0	0.0	17	0	0	1	2	5	0	5	30	0	3	0	0	0	0	3	0	0	0	0	0	0	0	0	0	0	0	0	0	0					D. bisectus L
153.0	300	0.0	12.0	0.0	2.0	24.6	0.0	15.0	0.0	9.3	0.0	3.6	7.3	0.0	1.6	1.0	23.6	0.0	0.0	2.0	7.0	0.0	0.0	13.0	0.0	5	0	2	0	2	2	0	6	17	0	4	0	1	0	0	5	0	1	0	0	0	0	0	1	0	2	0	0	1	0					
150.0	300	0.0	11.0	0.0	5.0	25.0	0.0	19.0	0.0	5.0	0.0	6.0	6.0	0.0	2.0	0.3	20.0	1.0	0.0	0.0	3.0	0.0	0.0	13.0	0.0	12	0	3	0	10	9	0	9	43	0	1	0	0	0	0	1	0	0	0	2	0	0	3	0	0	5	0	0	1	0					
149.0	300	0.0	7.0	0.0	5.0	38.6	0.0	12.0	0.0	4.0	0.0	4.0	4.0	0.3	3.0	0.0	22.0	0.0	0.0	1.0	8.0	0.0	0.0	9.0	0.0	25	0	2	4	2	10	0	16	59	0	1	0	1	0	1	3	0	0	0	1	0	0	1	0	0	2	0	0	0	0					
148.0	300	0.0	23.0	0.0	9.0	18.3	0.0	0.3	0.0	7.0	0.0	6.0	9.0	0.3	7.0	0.0	20.0	0.0	0.0	2.0	3.0	0.0	0.0	18.0	0.0	5	0	1	0	4	10	0	6	26	0	0	0	0	0	1	1	0	0	0	0	0	0	1	0	0	1	0	1	0	0	MNP16Ba				
147.0	300	0.0	17.0	0.0	5.6	17.3	0.0	0.0	0.0	3.0	0.3	17.0	9.6	0.0	6.6	8.3	15.3	0.0	0.0	0.0	2.0	0.0	0.0	10.0	0.0	7	0	9	1	3	20	0	26	66	0	5	0	0	0	0	5	0	0	0	0	0	0	1	0	0	1	0	0	0	0					
145.0	300	0.0	18.0	0.0	7.0	26.0	0.0	0.0	0.0	2.6	0.0	7.3	4.3	0.0	3.0	4.0	28.0	1.0	0.0	0.0	3.0	0.0	0.0	10.0	0.0	2	0	0	0	2	1	0	6	11	0	5	0	0	0	0	5	0	0	0	2	0	0	0	0	0	2	0	1	0	0					
144.0	300	0.0	11.0	0.0	2.0	48.0	0.0	0.0	0.0	3.0	0.0	8.0	4.0	0.0	2.0	3.0	19.0	0.0	0.0	0.0	1.0	0.0	0.0	11.0	0.0	0	0	0	0	1	7	0	2	10	0	1	0	0	0	1	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0					
143.0	300	0.0	8.0	0.0	4.0	31.0	0.0	0.0	0.0	5.0	0.0	12.0	5.0	0.0	1.0	5.0	29.0	1.0	0.0	0.0	2.0	0.0	0.0	12.0	0.0	9	0	0	1	2	10	0	14	36	0	2	0	0	0	0	2	0	0	0	1	0	0	0	0	0	1	0	1	0	0					
142.0	300	0.0	10.0	0.0	5.0	36.0	0.0	0.0	0.0	9.0	0.0	12.0	1.0	0.0	2.0	2.0	23.0	12.0	0.0	1.0	4.0	0.0	0.0	10.0	0.0	5	0	0	0	1	4	0	9	19	0	4	0	0	0	0	4	0	0	0	1	0	0	0	0	0	1	0	0	0	0					S. furcatolitho
141.0	300	0.0	12.0	0.0	4.0	40.0	0.0	0.0	0.0	4.0	0.0	7.0	2.0	0.0	3.0	4.0	24.0	18.0	0.0	0.0	2.0	0.0	0.0	7.0	0.0	7	0	3	1	4	10	0	29	54	0	8	0	0	0	0	8	0	0	0	0	0	0	0	0	0	0	0	1	0	0					HCO
140.0	300	0.0	7.6	0.0	4.0	32.0	0.0	0.0	0.0	2.0	0.0	10.6	1.6	0.0	3.0	4.0	35.0	18.0	0.0	0.0	0.0	0.0	0.0	16.0	0.0	24	0	2	0	15	4	0	20	65	0	6	0	0	0	0	6	0	0	0	0	0	0	0	0	0	0	0	0	0	0					
135.0	300	0.0	3.6	0.0	4.0	39.0	0.0	0.0	0.0	4.6	0.0	7.6	2.6	0.0	2.0	4.6	32.0	26.0	0.0	0.0	0.0	0.0	0.0	8.0	0.0	6	0	4	0	0	0	0	3	13	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0					
130.0	300	0.0	6.0	0.0	4.3	39.0	0.0	0.0	0.0	6.0	0.0	6.3	1.0	0.0	1.6	6.0	30.0	40.0	0.0	0.0	0.0	0.0	0.0	4.0	0.0	5	0	6	0	2	4	0	10	27	0	3	0	0	0	0	3	0	0	0	0	0	0	0	0	0	0	0	0	0	0					
125.0	300	0.0	9.6	0.0	4.3	52.0	0.0	0.3	0.3	1.6	0.0	3.0	1.6	0.0	1.0	1.3	25.3	14.0	0.0	0.0	2.0	0.0	0.0	8.0	0.0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0					
115.0	300	0.0	15.0	0.0	7.6	41.3	0.0	1.0	0.6	7.0	0.0	3.0	0.3	0.0	3.6	1.6	19.0	20.0	0.0	0.0	2.0	0.0	0.0	2.0	0.0	0	0	0	0	0	0	0	0	0	0	2	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	1	1	0					
109.0	300	0.6	17.0	0.0	4.6	24.6	0.0	0.0	0.0	14.3	1.3	2.6	2.6	0.0	8.0	4.6	19.6	25.0	0.0	1.0	1.0	0.0	0.0	13.0	0.0	57	0	31	8	6	1	0	46	149	0	9	0	0	0	1	10	0	2	3	0	0	0	1	0	0	6	0	0	1	0					
105.0	300	0.0	13.0	0.0	3.6	26.0	0.0	0.3	0.0	1.6	0.0	8.6	0.6	0.0	1.0	2.0	43.0	4.0	0.0	0.0	2.0	0.0	0.0	1.0	0.0	0	0	0	0	0	0	0	0	0	0	4	0	0	0	0	4	0	0	0	0	0	0	0	0	0	0	0	0	0	0					
100.0	300	0.0	11.0	0.0	4.0	32.0	0.0	0.0	0.0	5.6	0.0	4.6	3.6	0.0	2.3	2.0	35.0	2.0	0.0	0.0	2.0	0.0	0.0	4.0	0.0	1	0	2	0	0	0	0	2	5	0	2	0	0	0	0	2	0	0	0	0	0	0	1	0	0	1	0	0	0	0					
95.0	300	0.0	8.0	0.0	12.0	44.0	0.0	0.0	0.0	5.0	0.0	4.0	3.0	0.0	2.0	2.0	20.0	5.0	0.0	0.0	1.0	0.0	0.0	6.0	0.0	1	0	2	2	0	0	0	2	7	0	1	0	3	0	3	7	0	0	0	0	0	0	0	0	0	0	0	0	0	0					
90.0	300	0.0	13.0	0.0	5.6	37.0	0.0	0.0	0.0	6.3	0.0	1.0	4.0	0.0	1.6	1.6	30.0	8.0	0.0	0.0	6.0	0.0	0.0	6.0	0.0	2	0	4	0	1	0	0	1	8	0	2	0	1	0	1	4	0	0	0	0	0	0	1	0	0	1	0	1	0	0					
85.0	300	0.0	13.3	0.0	3.0	41.0	0.0	0.3	0.3	9.0	0.0	6.0	4.6	0.0	0.6	5.0	17.0	8.0	0.0	0.0	3.0	0.0	0.0	4.0	0.0	1	0	4	0	1	0	0	1	7	0	0	0	3	0	4	7	0	0	0	0	0	0	2	0	0	2	0	1	0	0		CNE14			
80.0	300	0.0	12.0	0.0	3.0	50.0	0.0	0.6	0.6	4.6	0.0	1.6	7.6	0.0	1.6	4.3	14.0	6.0	0.0	2.0	2.0	0.0	0.0	2.0	0.0	0	0	0	0	0	0	0	0	0	0	1	0	2	0	0	3	0	0	0	0	0	0	1	0	0	1	0	0	0	0					
77.5	300	0.0	17.0	0.0	6.0	33.0	0.0	0.3	1.3	3.6	0.0	2.6	10.6	0.0	0.0	0.6	25.0	8.0	0.0	1.0	1.0	0.0	0.0	6.0	0.0	0	0	0	0	0	0	0	0	0	0	1	0	1	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0					
70.0	300	0.0	17.0	0.0	5.0	32.0	0.0	0.6	2.0	10.6	0.0	5.6	8.0	0.0	2.0	3.3	14.0	6.0	0.0	0.0	0.0	0.0	0.0	4.0	0.0	0	0	0	0	0	0	0	0	0	0	2	0	0	0	0	2	0	0	0	0	0	2	3	0	0	5	0	0	0	0					
65.0	300	0.0	20.0	0.0	2.6	29.3	0.0	0.0	3.6	6.0	0.0	2.6	6.6	0.0	1.6	0.6	27.0	4.0	0.0	0.0	2.0	0.0	0.0	0.0	0.0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	0	0	2	0					
63.5	300	0.0	9.6	0.0	3.0	37.6	0.0	1.0	1.3	2.3	0.0	2.6	11.6	0.0	0.3	2.0	28.6	3.0	0.0	0.0	2.0	0.0	0.0	4.0	0.0	1	0	0	0	0	0	0	4	5	0	6	0	1	1	1	9	0	0	0	0	0	0	0	0	0	0	0	0	0	0				CP14a	
62.0	300	0.0	13.3	0.0	2.0	46.0	0.0	0.0	1.0	4.0	0.0	3.0	9.0	2.0	0.3	0.3	19.0	6.0	0.0	0.0	5.0	0.0	0.0	5.0	1.0	0	0	3	0	0	1	0	1	5	0	9	0	1	0	0	10	0	0	0	0	0	0	0	0	0	0	0	0	0	0					
61.0	300	0.0	9.6	0.0	1.0	33.0	0.0	0.6	0.6	10.0	0.0	0.0	23.3	0.6	0.6	0.6	31.0	5.0	0.0	0.0	1.0	0.0	0.0	11.0	4.0	2	0	5	0	0	0	0	2	9	0	6	0	1	1	0	8	0	0	0	0	0	1	0	0	0	0	0	0	0	1					B. gladius H
60.0	300	0.0	15.0	0.0	3.0	30.0	0.0	0.0	2.0	4.0	0.0	0.3	22.6	5.0	0.0	3.0	15.0	0.0	0.0	0.0	1.0	0.0	0.0	2.0	1.0	0	0	1	0	0	1	0	0	2	0	1	0	0	3	0	4	0	0	0	0	0	0	0	0	0	0	0	0	0	0					
59.0	SB	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0					
58.0	300	0.0	20.0	0.0	2.0	31.3	0.0	2.0	2.3	2.0	0.0	1.0	6.3	0.0	0.3	1.6	31.0	8.0	0.0	0.0	1.0	1.0	1.0	8.0	5.0	5	0	2	0	0	0	0	7	14	0	5	0	0	0	1	6	0	0	0	0	0	0	0	0	0	0	0	0	0	0					
57.0	300	0.0	16.3	0.0	3.0	27.6	0.0	0.6	0.0	6.3	0.0	3.0	8.6	0.0	0.3	4.0	30.3	2.0	0.0	0.0	2.0	1.0	0.0	11.0	1.0	2	0	3	0	0	0	0	1	9	0	2	0	1	0	0	3	0	0	0	0	0	0	0	0	0	0	0	1	0	0					
56.0	300	0.0	9.6	0.0	2.0	50.0	0.0	1.3	2.0	5.3	0.0	5.0	4.3	0.0	0.3	1.0	19.0	4.0	0.0	0.0	2.0	0.0	0.0	2.0	5.0	4	0	0	0	1	0	0	1	6	0	4	0	0	0	0	4	0	0	0	0	0	0	0	0	0	0	0	2	0	0					
55.0	300	0.0	14.6	0.0	0.6	50.0	0.0	0.3	1.0	4.0	0.0	0.0	4.0	0.6	0.0	1.3	23.6	2.0	0.0	0.0	2.0	0.0	0.0	2.0	2.0	1	0	3	0	0	0	0	1	5	0	2	0	0	0	0	2	0	0	0	1	0	0	0	0	0	1	0	1	1	0					
54.0	300	0.0	11.0	0.0	2.0	49.3	0.0	0.6	1.0	8.6	0.0	7.0	3.0	0.6	0.3	0.6	16.0	2.0	0.0	0.0	0.0	0.0	0.0	2.0	3.0	0	0	0	1	0	0	0	2	3	0	3	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0					

53.0	300	0.0	10.3	0.0	1.6	38.6	0.0	0.0	1.0	9.0	0.0	1.6	10.0	0.3	0.3	3.3	24.0	4.0	0.0	0.0	2.0	1.0	0.0	13.0	5.0	1	0	0	0	0	1	0	1	3	0	1	0	1	3	0	5	1	0	0	0	0	0	2	3	0	5	0	0	0	0		
52.0	300	0.0	17.0	0.0	1.6	20.0	0.0	0.3	1.0	9.0	0.0	2.0	16.0	2.0	0.0	2.0	28.0	3.0	0.0	0.0	1.0	1.0	0.0	12.0	3.0	0	0	1	0	0	0	0	2	3	0	3	0	4	2	1	13	0	0	0	0	0	0	0	1	0	1	0	0	0	0		
51.0	300	0.0	16.0	0.0	0.6	27.3	0.0	0.3	1.0	10.6	0.6	0.6	15.3	0.0	0.3	1.0	26.3	3.0	0.0	0.0	1.0	1.0	0.0	10.0	5.0	2	0	1	0	0	0	0	4	7	0	1	0	8	1	2	13	0	0	0	0	0	0	0	2	0	2	0	0	0	0		
50.9	300	0.0	19.0	0.0	4.0	30.0	0.0	4.0	1.6	3.0	?	1.0	6.0	0.0	0.3	6.0	25.0	3.0	0.0	0.0	5.0	1.0	0.0	4.0	3.0	0	0	0	0	0	0	0	2	2	0	2	0	2	0	0	4	0	0	0	0	0	0	0	1	0	1	0	0	0	0		
50.0	300	0.0	20.3	0.0	1.3	34.0	0.0	0.0	1.0	5.0	?	2.0	7.6	0.0	0.3	0.3	28.0	3.0	0.0	0.0	1.0	0.0	1.0	9.0	3.0	0	0	1	0	0	0	0	0	1	0	7	0	0	2	1	12	0	0	0	0	0	0	0	0	0	0	0	0	0	0		C. reticulatun
45.5	300	0.0	18.6	0.0	0.0	24.6	0.0	0.0	0.6	8.0	?	0.0	12.0	0.6	0.0	5.0	26.6	4.0	0.0	0.0	1.0	0.0	0.0	10.0	0.0	3	0	5	0	1	0	0	5	14	cf	1	0	4	0	2	10	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
43.5	300	0.0	13.6	0.0	0.0	16.6	0.0	0.6	2.0	3.0	?	7.0	19.0	0.0	0.0	6.0	32.0	3.0	0.0	0.0	5.0	0.0	0.0	13.0	2.0	0	0	4	0	0	2	0	1	7	0	8	0	0	1	2	15	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
42.0	300	0.0	11.6	0.0	0.0	26.6	0.0	0.6	0.6	7.3	?	4.0	12.0	2.0	0.0	5.0	30.3	3.0	0.0	0.0	1.0	0.0	0.0	10.0	1.0	1	1	4	0	0	0	0	4	10	0	6	0	2	1	1	11	0	0	0	0	0	0	1	0	0	1	0	0	0	0		
40.0	300	0.0	16.0	0.0	0.0	27.0	0.0	0.6	4.0	3.3	?	1.0	12.0	4.0	0.3	1.0	30.6	2.0	0.0	0.0	2.0	0.0	0.0	8.0	2.0	0	0	0	0	0	0	0	1	1	0	1	0	2	0	0	5	0	0	0	1	0	0	3	0	0	4	1	0	0	0		N. cristata H
39.0	300	0.0	11.3	0.0	0.0	27.3	0.0	0.0	1.0	3.0	?	2.0	25.0	2.0	0.3	1.0	27.0	5.0	0.0	0.0	4.0	0.0	0.0	13.0	3.0	1	0	0	0	0	0	0	1	2	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	1	1	0	0	0	MNP16A	
38.0	300	0.0	12.3	0.0	0.0	33.0	0.0	0.0	0.3	8.6	?	2.0	17.0	1.6	0.0	0.6	24.6	4.0	0.0	0.0	2.0	1.0	1.0	6.0	0.0	1	0	0	0	0	1	0	1	3	0	0	0	1	2	1	4	0	0	0	0	0	0	0	1	0	1	0	0	0	0		
37.0	300	0.0	16.0	0.0	0.0	27.0	0.0	0.0	0.0	7.0	?	4.0	19.0	0.0	0.0	3.0	24.0	12.0	0.0	0.0	4.0	2.0	1.0	24.0	5.0	15	0	14	0	3	2	0	10	44	0	8	0	8	4	2	22	0	0	0	0	0	0	0	0	0	0	0	0	1	0		
36.0	100	0.0	17.0	0.0	0.0	27.0	0.0	0.0	0.3	4.0	?	4.0	23.0	1.3	0.0	0.0	23.3	4.0	0.0	0.0	1.0	3.0	0.0	4.0	0.0	1	0	0	0	0	0	0	0	1	0	0	0	3	0	1	6	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
35.1	300	0.0	22.6	0.0	0.3	22.6	0.0	0.0	0.3	7.0	?	0.6	20.6	0.6	0.3	0.6	24.0	1.0	0.0	0.0	2.0	1.0	0.0	16.0	2.0	4	0	2	0	0	0	0	8	14	0	2	0	3	3	2	14	0	0	0	1	0	0	0	0	0	0	0	0	0	0		
35.0	300	0.0	18.0	0.0	0.0	19.3	0.0	0.0	2.0	5.0	?	5.0	18.3	1.0	0.3	7.0	24.0	6.0	0.0	0.0	1.0	0.0	0.0	13.0	1.0	4	0	3	0	0	0	0	5	12	0	3	0	3	1	7	17	0	0	0	0	0	0	0	0	0	0	1	0	0	0		
34.0	300	0.0	12.0	0.0	0.0	29.0	0.0	0.0	1.0	3.0	?	3.0	10.0	0.0	0.0	4.0	38.0	8.0	0.0	0.0	7.0	0.0	0.0	13.0	7.0	5	0	5	0	0	0	0	5	15	0	4	0	0	3	0	8	1	0	0	0	0	0	1	1	0	2	0	0	0	0		
33.0	SB	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	?	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
32.0	300	0.0	8.0	0.0	0.3	24.6	0.0	0.0	1.0	4.0	?	10.0	13.0	2.0	0.0	11.0	26.0	7.0	0.0	0.0	3.0	1.0	0.0	11.0	1.0	4	0	2	0	0	0	0	1	7	0	5	0	2	0	0	8	0	0	0	0	0	0	0	0	0	0	2	0	0	0		
31.0	300	0.0	15.0	0.0	0.3	30.3	0.0	0.0	1.0	5.0	?	6.0	10.0	1.0	0.3	6.0	25.0	5.0	0.0	0.0	2.0	1.0	0.0	15.0	5.0	2	0	3	0	0	1	0	2	8	0	9	0	4	2	2	21	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
29.0	300	0.0	20.0	0.0	0.0	23.6	0.0	0.0	1.0	4.3	?	4.3	19.0	4.0	0.0	3.0	20.6	5.0	0.0	0.0	1.0	1.0	0.0	10.0	1.0	2	0	2	0	0	0	0	1	5	0	3	0	2	1	0	8	0	0	0	0	0	0	0	0	0	0	1	0	0	0	NP15	
28.0	300	0.0	18.0	0.0	0.0	14.0	0.0	0.0	1.0	3.0	?	1.0	35.0	2.0	0.0	3.0	23.0	5.0	0.0	0.0	7.0	1.0	0.0	8.0	7.0	2	0	2	0	0	3	0	3	10	0	3	0	1	6	0	12	0	0	0	1	0	0	0	3	0	4	1	0	0	0		
27.0	300	0.3	15.0	0.0	0.0	19.0	0.0	0.0	2.0	4.0	?	1.0	19.0	5.6	0.0	2.0	32.0	4.0	0.0	0.0	3.0	0.0	0.0	12.0	2.0	1	0	1	0	0	0	0	5	7	0	0	0	3	4	2	11	1	0	0	0	0	0	0	3	0	0	0	0	0	0		
26.0	300	0.0	17.0	0.0	0.0	24.6	0.0	0.6	1.0	12.3	?	1.0	23.0	0.0	0.0	3.0	17.3	6.0	0.0	0.0	2.0	1.0	0.0	7.0	0.0	0	0	0	0	0	1	0	3	4	0	0	0	3	2	1	7	0	0	0	0	0	0	0	1	0	0	0	0	0	0		
25.0	300	0.0	18.3	0.0	0.0	24.3	0.0	0.0	2.0	6.0	?	0.6	26.0	1.3	0.0	0.3	21.0	4.0	0.0	0.0	0.0	1.0	0.0	12.0	1.0	0	0	1	0	0	0	0	2	3	0	2	0	0	0	0	4	0	0	0	0	0	0	1	0	0	0	3	0	0	0		
24.0	300	0.0	19.3	0.0	0.0	26.0	0.0	0.0	4.0	3.0	?	1.0	22.6	1.0	0.0	3.0	20.0	2.0	0.0	0.0	0.0	0.0	0.0	2.0	1.0	0	0	0	0	0	0	0	0	0	0	0	0	2	1	2	6	0	0	0	1	0	0	0	0	0	0	0	0	0	0		
23.0	300	0.0	13.0	0.0	0.0	39.3	0.0	0.3	3.0	4.0	?	5.0	5.0	0.0	0.3	2.0	28.0	1.0	0.0	0.0	2.0	0.0	0.0	8.0	1.0	0	0	0	0	0	0	0	2	2	0	3	0	1	0	1	5	0	0	0	1	0	0	1	0	0	0	0	0	0	0	CNE13	
22.0	SB	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	?	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0	0	0	0	0	0	0	0		0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
20.0	300	0.0	12.3	0.0	0.3	12.6	0.0	0.0	0.0	5.0	?	7.3	20.0	0.0	0.0	8.3	34.0	5.0	0.0	0.0	3.0	2.0	0.0	4.0	5.0	3	0	2	0	0	0	0	3	8	0	5	0	7	1	4	20	0	0	0	0	0	0	0	0	0	0	0	1	0	0		
19.0	300	0.0	15.0	0.0	0.3	17.0	0.0	0.0	2.0	5.0	?	4.0	23.6	0.0	0.0	4.0	29.0	5.0	0.0	0.0	7.0	0.0	0.0	11.0	6.0	3	0	0	0	0	0	0	1	4	0	2	0	9	1	1	17	0	0	0	0	0	0	0	2	0	0	0	1	0	0		
18.0	300	0.0	22.0	0.0	0.0	20.6	0.0	0.0	2.0	3.0	?	12.0	22.0	0.0	0.0	5.0	13.3	6.0	0.0	0.0	4.0	0.0	2.0	3.0	3.0	2	0	2	0	0	0	0	3	8	0	3	0	4	3	0	14	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
16.0	300	0.0	20.3	0.0	0.0	25.0	0.0	0.6	2.0	3.0	?	1.0	30.0	2.0	0.0	1.0	15.0	2.0	0.0	0.0	0.0	2.0	0.0	6.0	5.0	1	0	0	0	0	0	0	2	3	0	4	0	6	2	1	15	1	0	0	3	0	0	2	2	0	7	0	0	0	0		
15.0	300	0.0	13.0	0.0	0.0	41.0	0.0	0.3	4.0	4.0	?	1.0	13.0	2.0	0.3	6.0	15.3	3.0	0.0	0.0	1.0	2.0	0.0	7.0	3.0	3	0	2	0	0	0	0	11	16	0	5	0	4	1	2	12	0	0	0	2	0	0	2	1	0	5	1	0	0	0		
14.0	300	0.0	13.3	0.0	1.0	53.0	0.0	1.0	5.0	4.0	?	0.0	11.6	0.3	0.0	0.6	10.0	4.0	0.0	0.0	0.0	1.0	0.0	4.0	3.0	5	0	0	0	1	4	0	1	11	0	0	0	0	3	0	3	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
13.9	300	0.0	14.3	0.0	0.0	30.6	0.0	0.0	5.0	7.0	?	1.0	19.3	1.0	0.0	3.0	18.6	7.0	0.0	0.0	6.0	2.0	0.0	9.0	3.0	1	0	1	1	0	0	0	1	4	0	0	0	2	3	2	11	2	0	0	0	0	0	3	0	0	0	0	0	0	0		
13.0	300	0.0	5.6	0.0	0.0	43.0	0.0	0.0	9.0	2.0	?	2.0	7.3	3.0	0.0	2.0	26.0	5.0	0.0	0.0	2.0	1.0	0.0	4.0	1.0	0	0	0	0	0	0	0	0	0	0	1		1	0	0	3	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
12.0	SB	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	?	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
11.0	300	0.0	20.0	0.0	0.0	14.0	0.0	0.0	2.0	6.0	?	2.0	25.0	2.0	0.0	7.0	22.0	6.0	0.0	0.0	1.0	1.0	0.0	6.0	3.0	5	2	4	0	0	1	0	7	19	cf	4	0	1	1	0	8	0	0	0	1	0	0	0	0	0	0	0	0	0	0		
10.0	300	0.0	21.3	0.0	0.3	27.6	0.0	0.0	1.3	4.0	?	3.0	10.0	3.0	0.0	5.0	24.3	2.0	0.0	0.0	0.0	0.0	0.0	6.0	1.0	2	0	0	0	0	0	0	1	3	0	0	0	4	1	0	6	0	0	0	0	0	0	0	1	0	0	0	0	0	0		
9.0	300	0.0	9.6	0.0	0.0	27.6	0.0	0.3	2.0	4.0	?	3.0	6.0	1.0	0.3	3.0	43.0	1.0	0.0	0.0	0.0	1.0	0.0	4.0	3.0	0	0	0	0	0	0	0	2	2	0	1	0	0	0	0	4	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
8.0	300	0.0	17.0	0.0	0.0	20.6	0.0	0.3	2.0	5.0	?	2.0	16.6	0.0	0.0	4.0	32.3	0.0	0.0	0.0	1.0	0.0	0.0	2.0	0.0	0	0	0	0	0	0	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
7.0	300	0.0	5.0	0.0	0.3	36.0	0.0	0.0	12.0	3.3	?	7.0	6.3	2.0	0.0	5.0	22.0	2.0	0.0	0.0	2.0	0.0	0.0	6.0	5.0	0	0	0	0	0	0	0	6	6	0	0	0	3	0	2	7	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
5.0	300	0.0	8.6	0.0	0.0	34.3	0.0	0.0	2.0	6.0	?	2.0	11.0	0.0	0.0	2.0	34.0	1.0	0.0	0.0	3.0	0.0	0.0	8.0	1.0	0	0	0	0	0	1	0	3	4	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
4.0	300	0.0	14.0	0.0	0.0	18.3	0.0	0.0	5.0	2.0	?	9.0	13.6	0.0	0.0	4.0	33.0	0.0	0.0	0.0	1.0	0.0	0.0	4.0	0.0	2	2	0	2	0	0	0	1	7	0	1	0	3	0	1	6	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
3.0	300	0.0	16.0	0.0	0.0	23.3	0.0	0.0	3.0	2.0	?	5.0	10.0	3.0	0.0	9.0	28.0	3.0	0.0	0.0	1.0	1.0	0.0	7.0	1.0	5	1	0	2	0	0	0	8	16	0	5	0	4	0	4	15	0	0	0	3	0	0	0	0	0	0	0	0	0	0		
2.0	300	0.0	10.6	0.0	0.0	36.0	0.0	0.0	3.0	3.0	?	3.0	20.0	3.0	0.0	3.3	18.0	2.0	0.0	0.0	2.0	1.0	1.0	9.0	1.0	3	2	3	0	0	1	0	4	13	0	5	0	5	0	0	12	0	0	0		0	0	0	0	0	0	1	1	0	0		
0.0	300	0.0	17.0	0.0	0.0	34.6	0.0	0.6	2.0	1.0	?	0.3	5.0	6.0	0.3	3.0	30.0	7.0	0.0	0.0	2.0	3.0	0.0	8.0	0.0	0	0	3	0	0	0	0	9	12	0	4	0	6	0	0	10	0	0	0		0	0	0	0	0	0	0	0	0	0		

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