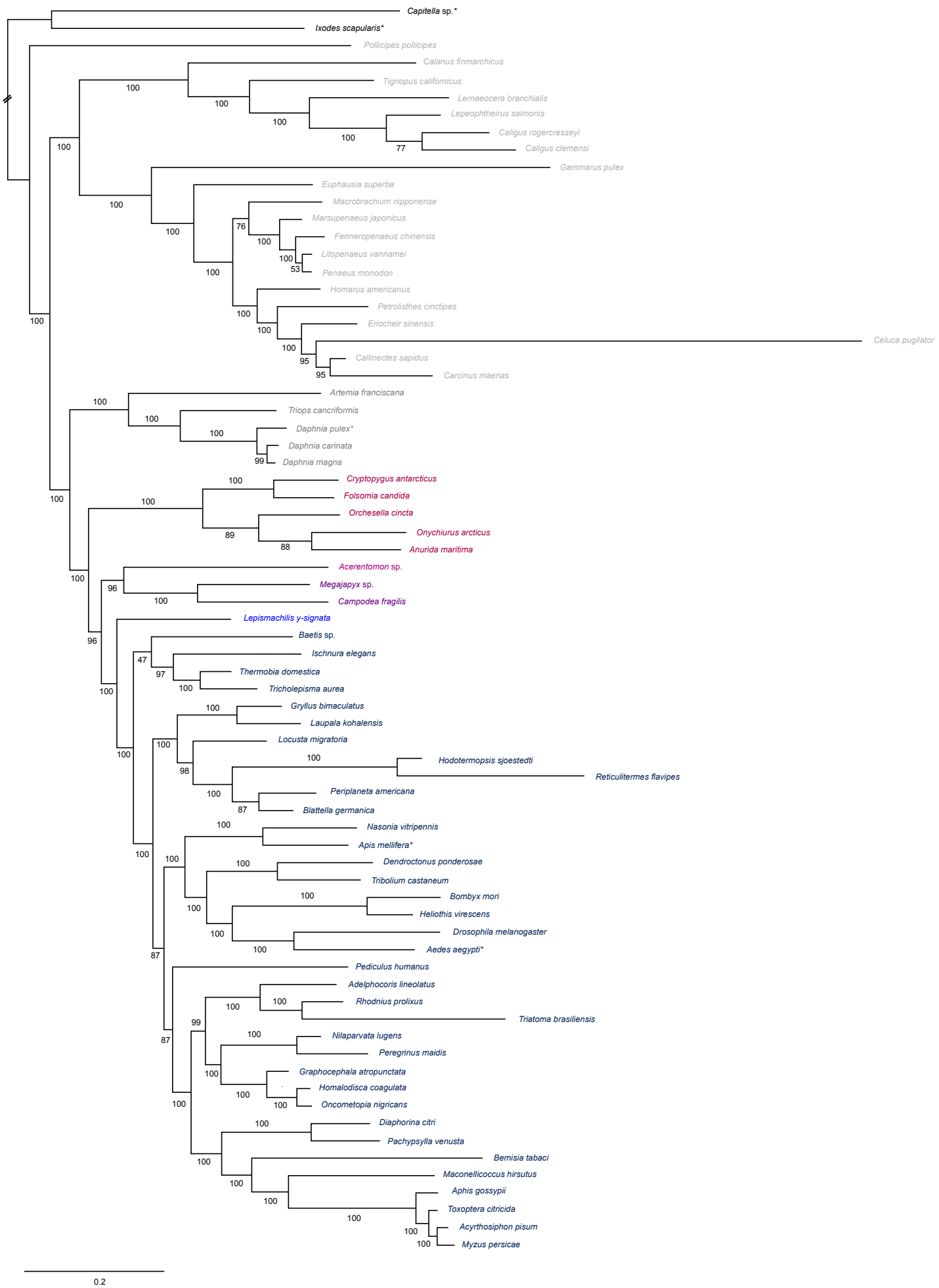
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Caligus clemensi
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Euphausia superba
Macrobrachium nipponense
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Fenneropenaeus chinensis
Litopenaeus vannamei
Penaeus monodon
Homarus americanus
Petrolisthes cinctipes
Eriocheir sinensis
Celuca pugilator
Callinectes sapidus
Carcinus maenas
Artemia franciscana
Triops cancriformis
Daphnia pulex*
Daphnia carinata
Daphnia magna
Cryptopygus antarcticus
Folsomia candida
Orchesella cincta
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Anurida maritima
Acerentomon sp.
Megajapyx sp.
Campodea fragilis
Lepismachilis y-signata
Baetis sp.
Ischnura elegans
Thermobia domestica
Tricholepisma aurea
Gryllus bimaculatus
Laupala kohalensis
Locusta migratoria
Hodotermopsis sjoestedti
Reticulitermes flavipes
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Blattella germanica
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Apis mellifera*
Dendroctonus ponderosae
Tribolium castaneum
Bombyx mori
Heliothis virescens
Drosophila melanogaster
Aedes aegypti*
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Adelphocoris lineolatus
Rhodnius prolixus
Triatoma brasiliensis
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Peregrinus maidis
Graphocephala atropunctata
Homalodisca coagulata
Oncometopia nigricans
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Pachypsylla venusta
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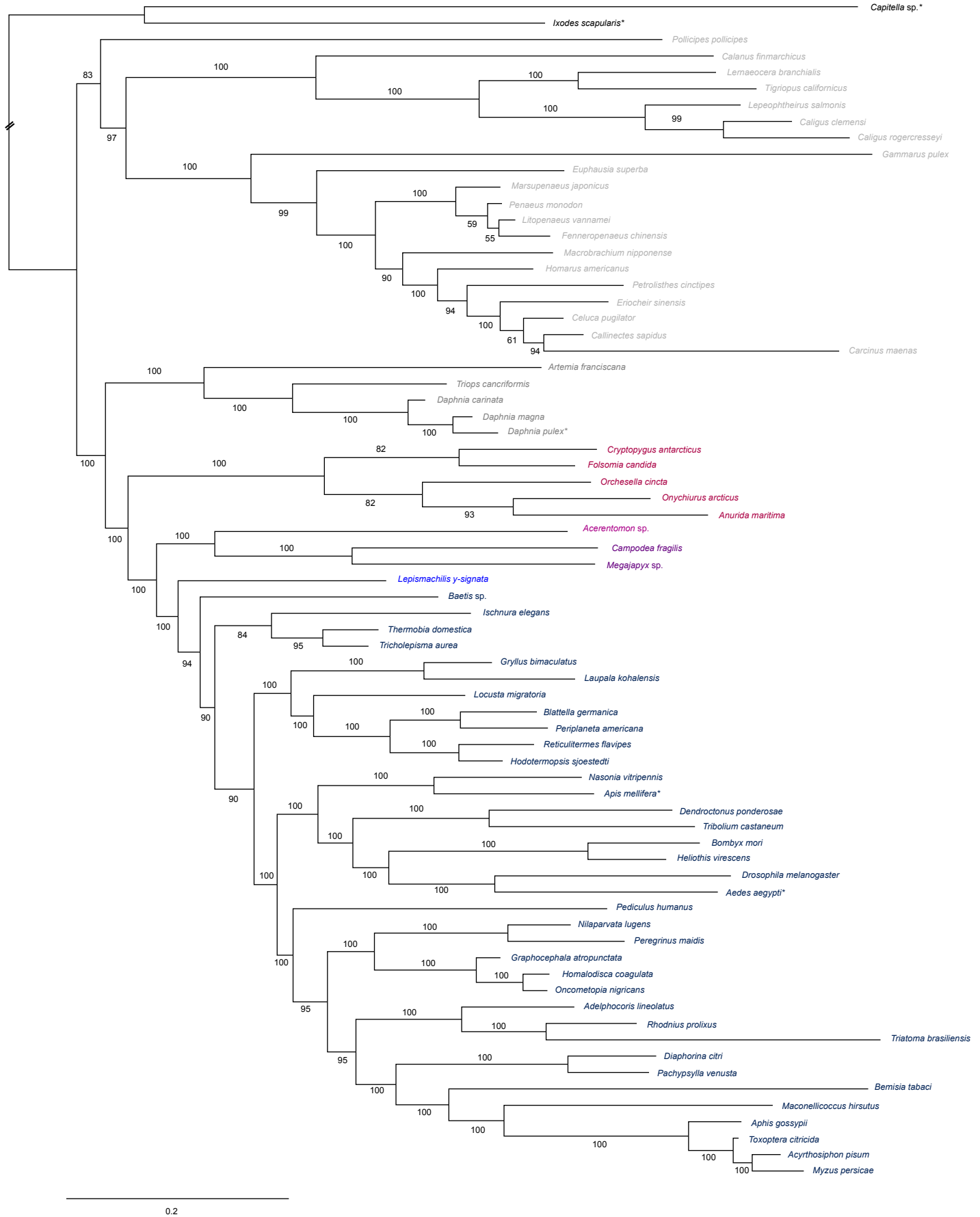
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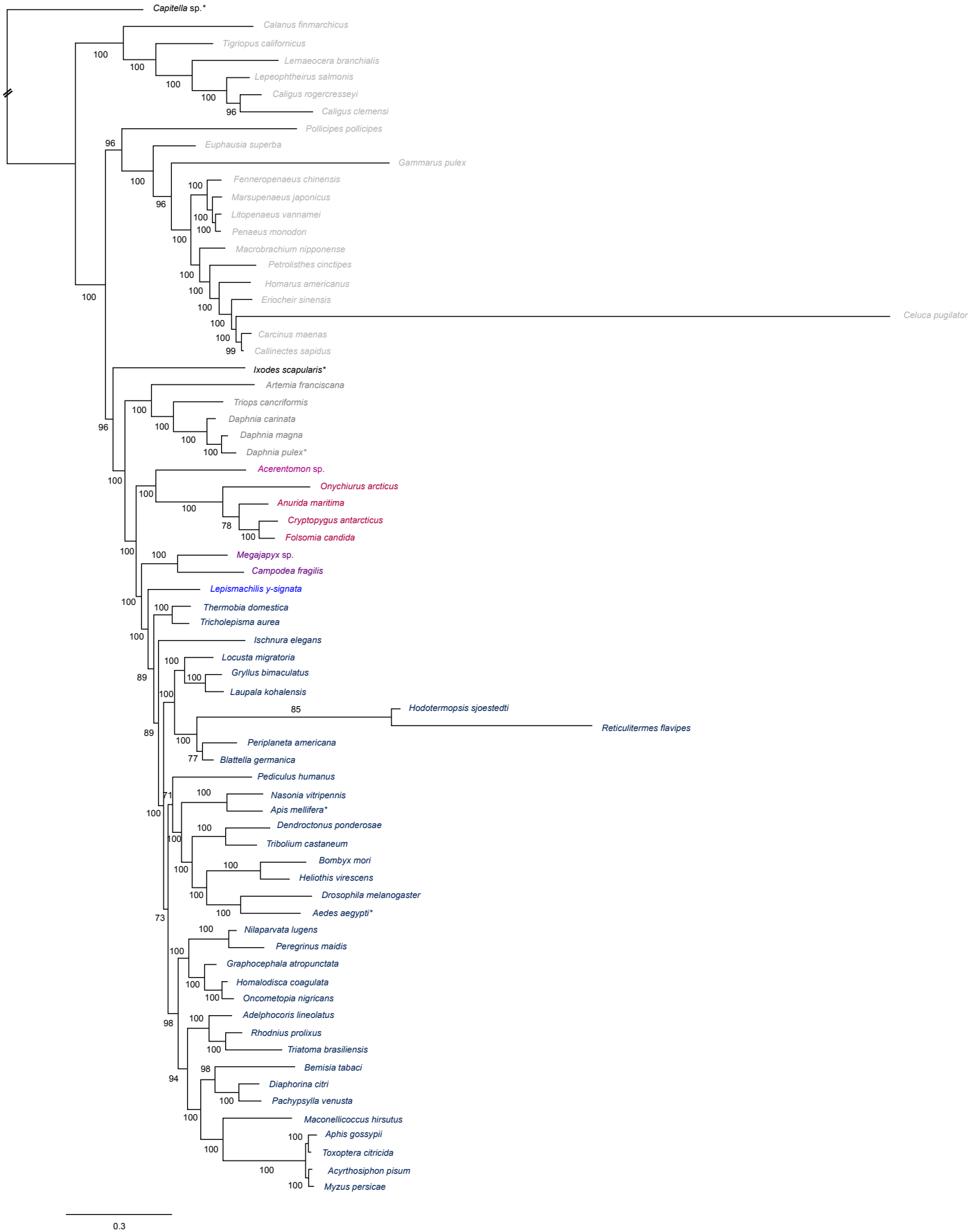
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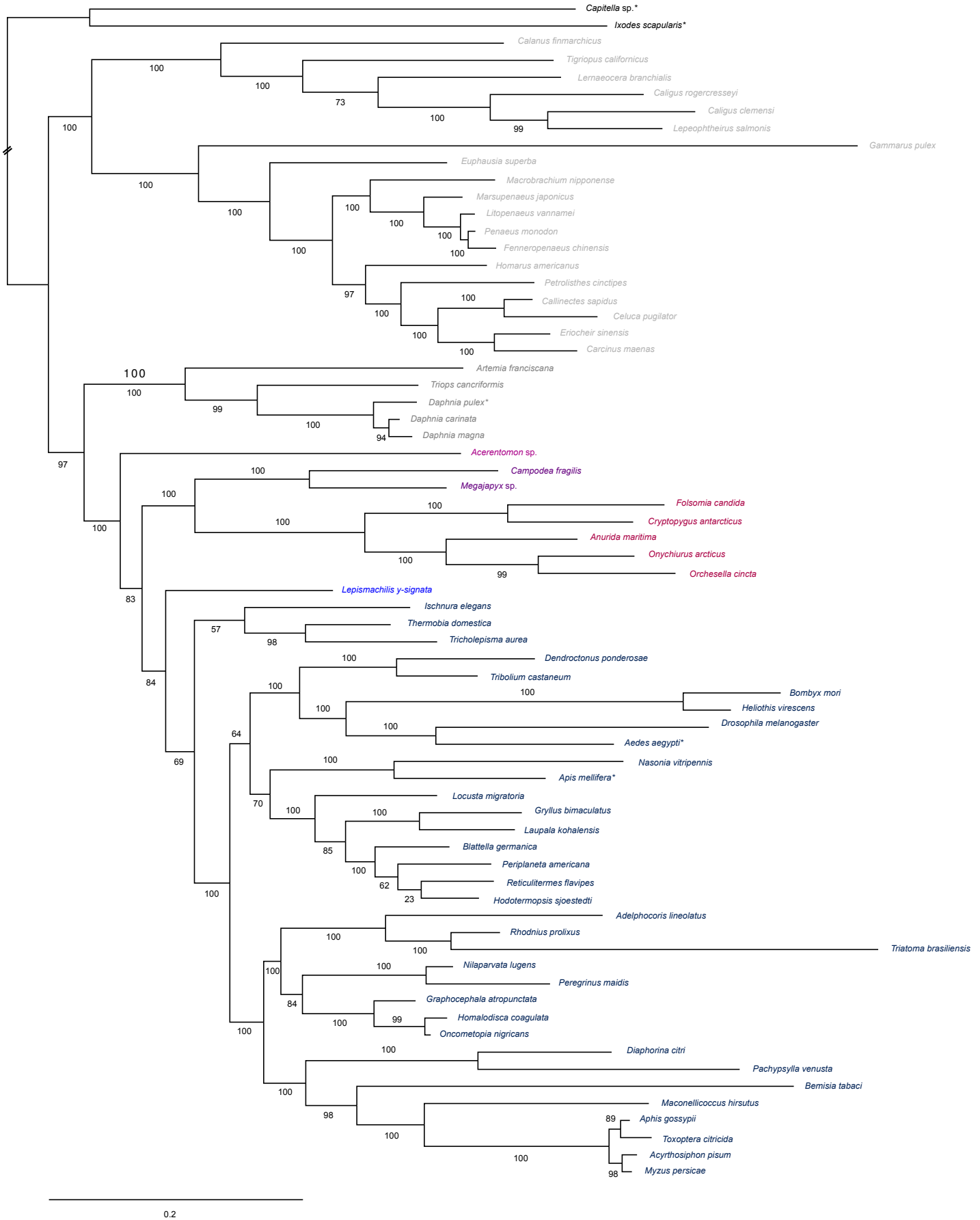
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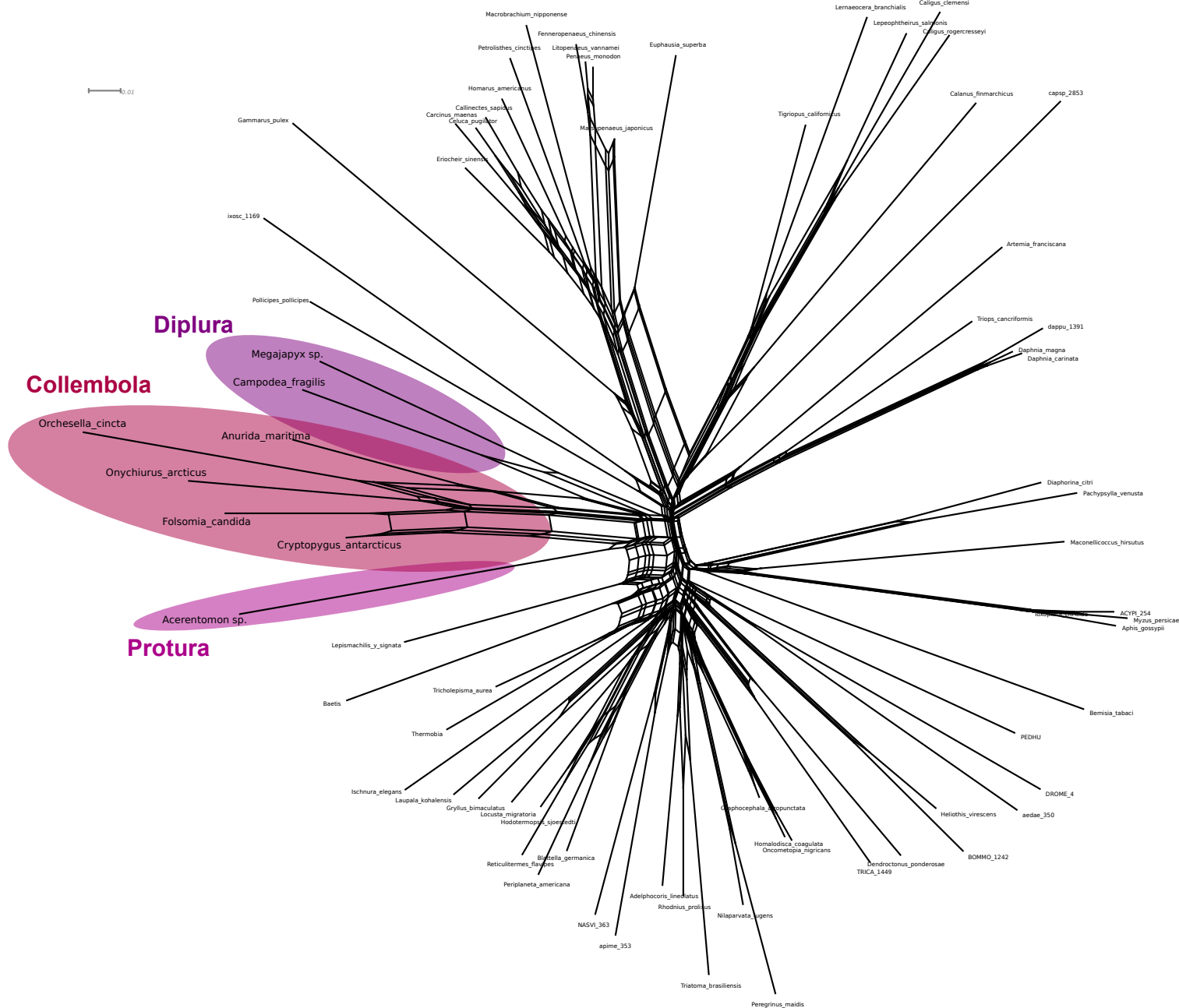
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DROME_4
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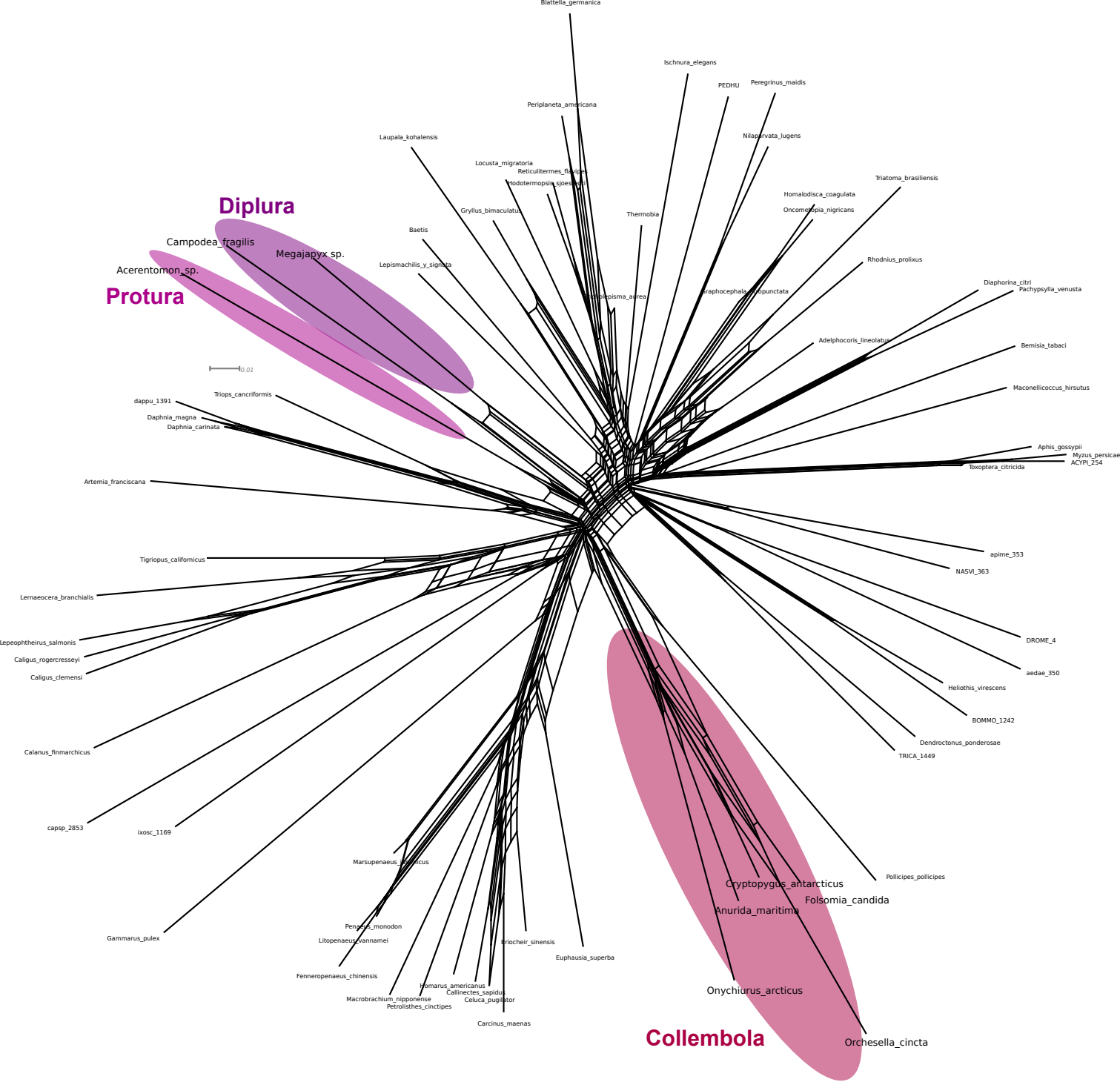
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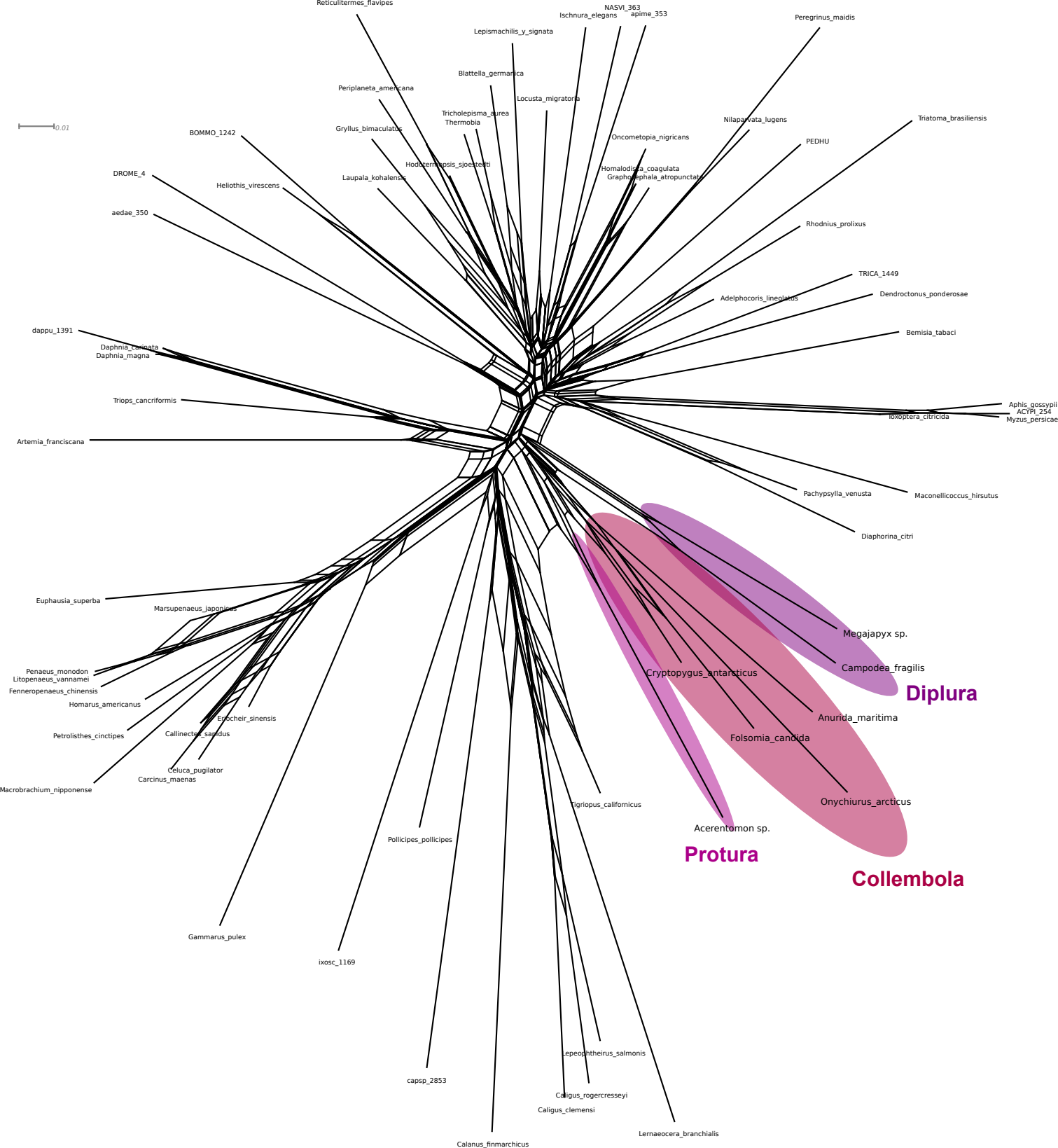
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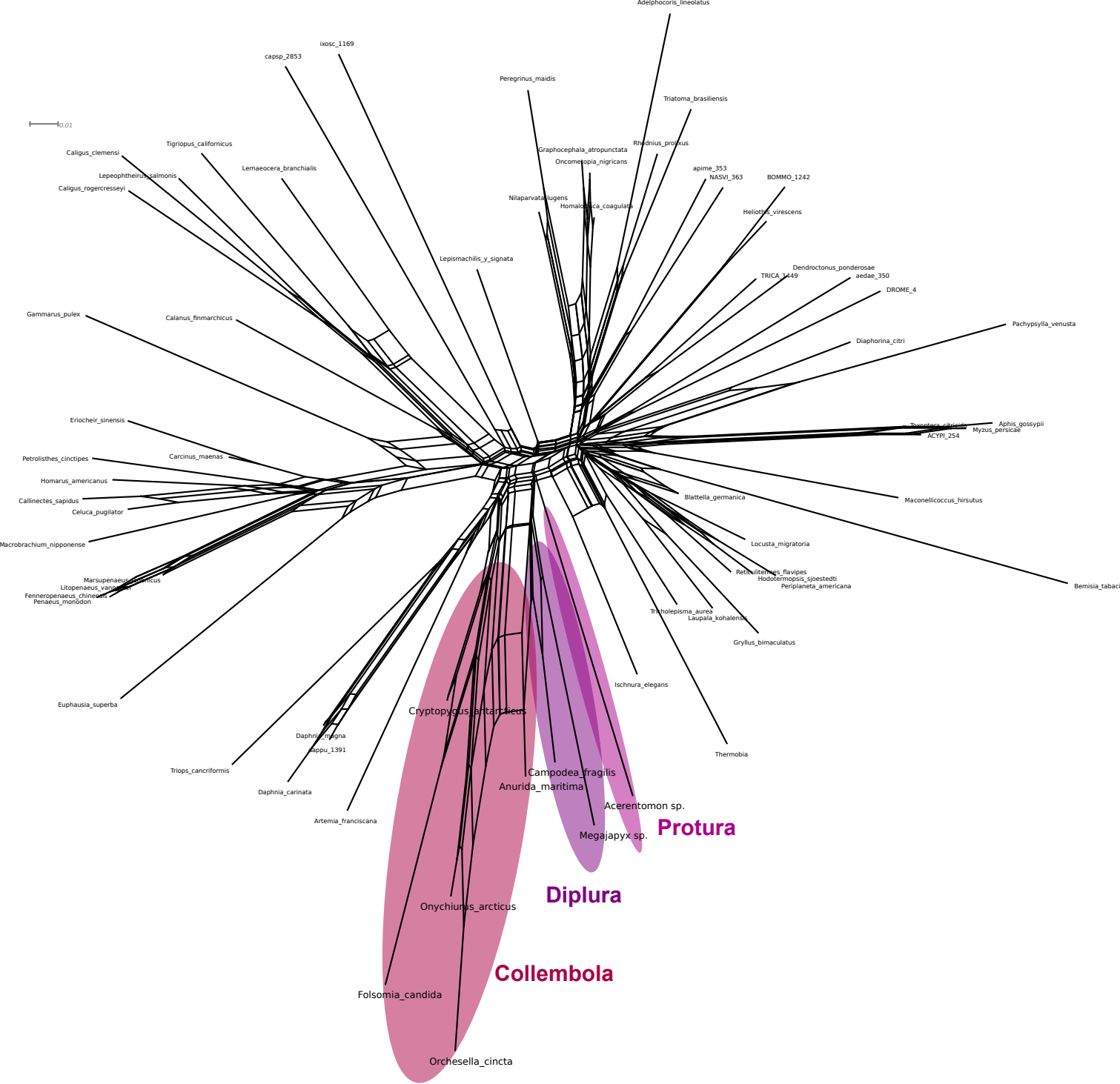
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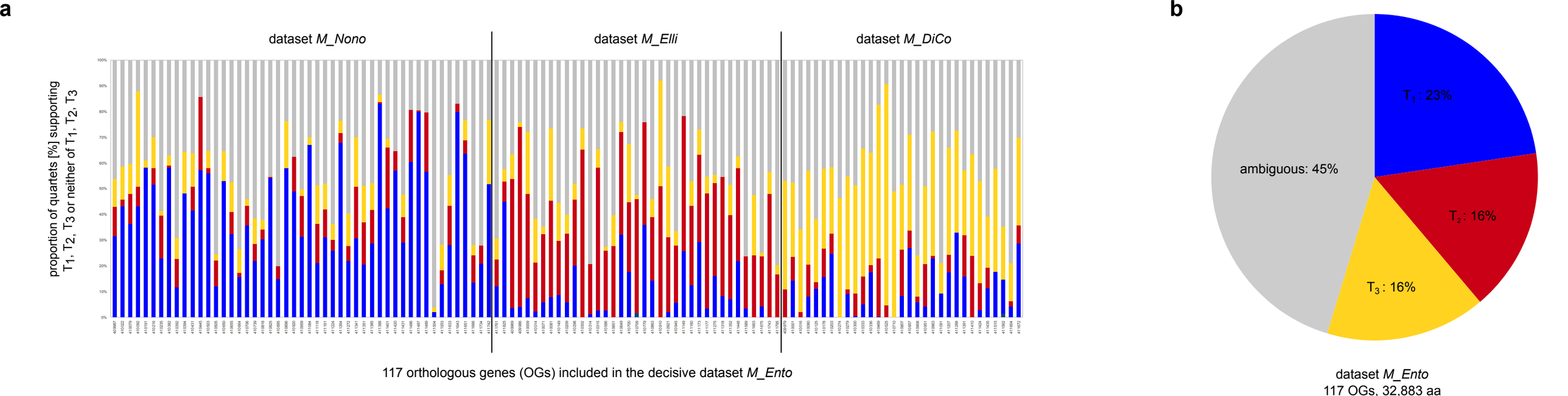
a
dataset M_Nono
dataset M_Elli
dataset M_DiCo
proportion of quartets [%] supporting T1, T2, T3 or neither of T1, T2, T3
100%
90%
80%
70%
60%
50%
40%
30%
20%
10%
0%
117 orthologous genes (OGs) included in the decisive dataset M_Ento
b
T1 : 23%
T2 : 16%
T3 : 16%
ambiguous: 45%
dataset M_Ento
117 OGs, 32,883 aa