

Character partitions of subset 1 and 2

Partitions subset 1

COX3, first maximum clique = 1-528

COX2, first maximum clique = 529-916

16S, second maximum clique = 917-1413

ND2, first maximum clique = 1414-2103

12S, second maximum clique = 2104-2543

ND1, second maximum clique = 2544-2997

ITS2, first maximum clique = 2998-4099

COX2, second maximum clique = 4100-4487

ND6, first maximum clique = 4488-4873

ITS2, second group of taxa with compositional homogeneity = 4874-5975

tRNAs concatenated, first maximum clique = 5976-6522

18S, first maximum clique = 6523-7333

28S, first maximum clique = 7334-8067

ND3, first maximum clique = 8068-8301

COX1, first maximum clique = 8302-9267

ND5, first maximum clique = 9268-10403

18S, second group of taxa with compositional homogeneity = 10404-11214

ATP8, first maximum clique = 11215-11322

ATP6, second maximum clique = 11323-11778

12S, first maximum clique = 11779-12218

COX2, second group of taxa with compositional homogeneity = 12219-12606

16S, second group of taxa with compositional homogeneity = 12607-13103

ND4, first maximum clique = 13104-14001

ATP6, first maximum clique = 14002-14457

CytoB, first maximum clique = 14458-14839

28S, second group of taxa with compositional homogeneity = 14840-15573

28S, second maximum clique = 15574-16307

16S, first maximum clique = 16308-16804

ND4L, first maximum clique = 16805-17000

ND1, first maximum clique = 17001-17454

ATP8, second group of taxa with compositional homogeneity = 17455-17562

nuclear protein-coding seqs = 17563-88626

Partitions subset 2

COX2, first maximum clique = 1-388
18S, first maximum clique = 389-1199
ND2, first maximum clique = 1200-1889
CytoB, first maximum clique = 1890-2271
COX2, second maximum clique = 2272-2659
ND6, first maximum clique = 2660-3045
ITS2, second group of taxa with compositional homogeneity = 3046-4147
tRNAs concatenated, first maximum clique = 4148-4694
12S, second maximum clique = 4695-5134
COX1, first maximum clique = 5135-6100
ND5, first maximum clique = 6101-7236
18S, second group of taxa with compositional homogeneity = 7237-8047
ND3, first maximum clique = 8048-8281
16S, first maximum clique = 8282-8778
16S, second maximum clique = 8779-9275
ATP8, first maximum clique = 9276-9383
COX3, first maximum clique = 9384-9911
12S, first maximum clique = 9912-10351
COX2, second group of taxa with compositional homogeneity = 10352-10739
16S, second group of taxa with compositional homogeneity = 10740-11236
ND4, first maximum clique, 11237-12134
ATP6, first maximum clique = 12135-12590
ND1, second maximum clique = 12591-13044
28S, second group of taxa with compositional homogeneity = 13045-13778
ATP6, second maximum clique = 13779-14234
28S, second maximum clique = 14235-14968
ND4L, first maximum clique = 14969-15164
ND1, first maximum clique = 15165-15618
ITS2, first maximum clique = 15619-16720
ATP8, second group of taxa with compositional homogeneity = 16721-16828
28S, first maximum clique = 16829-17743
nuclear protein-coding seqs = 17744-88807