

Resolving the phylogenetic position of Darwin's extinct ground sloth (*Mylodon darwini*) using mitogenomic and nuclear exon data

Frédéric Delsuc, Melanie Kuch, Gillian C. Gibb, Jonathan Hughes, Paul Szpak, John Southon, Jacob Enk, Ana T. Duggan' and Hendrik N. Poinar

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Table S1. Detailed taxonomic sampling and corresponding sequence accession numbers of the mitochondrial and nuclear datasets. Newly acquired sequences from this study are indicated in bold.

Species	Common name	Mitogenomes	ADORA3	APOB	BCHE	BRCA1	BRCA2	RAG2	TTN
<i>*Bradypus pygmaeus</i>	Pygmy sloth	KT818523	LT852562	LT852583	LT852604	LT852627	LT852645	LT852662	LT852684
<i>*Bradypus torquatus</i>	Maned sloth	KT818524	LT852563	LT852584	LT852605	LT852628	LT852646	LT852663	LT852685
<i>*Bradypus tridactylus</i>	Pale-throated three-toed sloth	KT818525	LT852564	LT852585	LT852606	LT852629	LT852647	LT852664	LT852686
	Brown-throated three-toed sloth	NC_006923	LT852565	LT852586	LT852607	LT852630	LT852648	LT852665	LT852687
<i>*Bradypus variegatus</i>	Southern two-toed sloth	NC_006924	LT852570	LT852591	LT852613	LT852634	LT852651	LT852671	LT852693
<i>*Choloepus didactylus</i>	Hoffmann's two-toed sloth	KT818538	LT852571	LT852592	LT852614	LT852635	LT852652	LT852672	LT852694
<i>*Choloepus hoffmanni</i>	Darwin's ground sloth	MF061314	LT852575	LT852596	LT852619	LT852639	LT852656	LT852676	LT852698
<i>*Mylodon darwini</i>	Pygmy anteater	KT818539	LT852572	LT852593	LT852615	LT852636	LT852653	LT852673	LT852695
<i>*Cyclopes didactylus</i>	Giant anteater	KT818549	LT852576	LT852597	LT852620	LT852640	LT852657	LT852677	LT852699
<i>*Myrmecophaga tridactyla</i>									
<i>*Tamandua mexicana</i>	Northern tamandua	KT818551	LT852578	LT852599	LT852622	LT852641	LT852659	LT852679	LT852701
<i>*Tamandua tetradactyla</i>	Southern tamandua	NC_004032	LT852579	LT852600	LT852623	LT852642	LT852660	LT852680	LT852702
<i>*Dasypus kappleri</i>	Greater long-nosed armadillo	KT818541	LT852574	LT852595	LT852617	LT852638	LT852655	LT852675	LT852697
<i>*Dasypus hybridus</i>	Southern long-nosed armadillo	KT818540	LT852573	LT852594	LT852616	LT852637	LT852654	LT852674	LT852696
<i>Dasypus septemcinctus</i>	Seven-banded armadillo	KT818546	-	-	-	-	-	-	-
<i>Dasypus novemcinctus</i>	Nine-banded armadillo	NC_001821	XM_004473	JN413957	XM_0125197	AF484222	XM_012519	XM_004483	XM_0125
US	armadillo		603		96		803	174	18926
<i>Dasypus novemcinctus</i>	Nine-banded armadillo	KT818542	-	-	-	-	-	-	-
FG	armadillo								
<i>Dasypus pilosus</i>	Hairy long-nosed armadillo	KT818544	-	-	-	-	-	-	-

<i>Dasybus sabanicola</i>	Northern long-nosed armadillo	KT818545	-	-	-	-	-	-	-
<i>Dasybus yepesi</i>	Yunga's long-nosed armadillo	KT818547	-	-	-	-	-	-	-
* <i>Chaetophractus vellerosus</i>	Screaming hairy armadillo	KT818533	LT852568	LT852589	LT852611	LT852632	NA	LT852669	LT852691
* <i>Chaetophractus villosus</i>	Large hairy armadillo	KT818535	LT852569	LT852590	LT852612	LT852633	NA	LT852670	LT852692
* <i>Euphractus sexcinctus</i>	Six-banded armadillo	KT818548	AY011193	NA	LT852618	AF484224	NA	AY011922	NA
* <i>Zaedyus pichiy</i>	Pichi	KT818555	LT852582	LT852603	LT852626	LT852644	NA	LT852683	LT852705
* <i>Calyptophractus retusus</i>	Greater fairy armadillo	KT818532	LT852567	LT852588	LT852610	FR821709	LT852650	LT852668	LT852690
<i>Chlamyphorus truncatus</i>	Pink fairy armadillo	KT818536	-	-	-	FR821708	-	-	-
* <i>Priodontes maximus</i>	Giant armadillo	KT818550	LT852577	LT852598	LT852621	AF484225	LT852658	LT852678	LT852700
* <i>Tolypeutes matacus</i>	Southern three-banded armadillo	KT818553	LT852580	LT852601	LT852624	AF484227	LT852661	LT852681	LT852703
* <i>Tolypeutes tricinctus</i>	Brazilian three-banded armadillo	KT818554	LT852581	LT852602	LT852625	LT852643	NA	LT852682	LT852704
* <i>Cabassous chacoensis</i>	Chacoan naked-tailed armadillo	KT818528	NA	NA	LT852608	LT852631	NA	LT852666	LT852688
* <i>Cabassous unicinctus</i>	Southern naked-tailed armadillo	KT818531	LT852566	LT852587	LT852609	AF484228	LT852649	LT852667	LT852689
<i>Cabassous centralis</i>	Northern naked-tailed armadillo	KT818527	-	-	-	-	-	-	-
<i>Cabassous tatouay</i>	Greater naked-tailed armadillo	KT818529	-	-	-	-	-	-	-
<i>Dugong dugon</i>	Dugong	NC_003314	JN633415	KR827337	KR827307 XM_0105960	AF284019	JN414242	JN633302	JN632896 XM_0105
<i>Loxodonta africana</i>	African elephant	NC_000934	AY011204	JN413956	53 XM_0079487	AF284021	JN414245	AY011932	86379
<i>Orycteropus afer</i>	Aardvark	NC_002078	AY011206	JN413950	19	AF284030	JN414239	AY011935	JN632894

Notes -- *: Libraries selected for nuclear capture; NA: Not available because capture failed to provide enough coverage.

Table S2: Detailed results of the PartitionFinder analysis for the mitogenomic dataset.

Subset	Best Model	Subset Partitions
1	GTR+I+G	ATP6_p1, ATP8_p1, ATP8_p2, ATP8_p3, CYTB_p1, ND1_p1, ND2_p1, ND3_p1, ND4L_p1, ND4_p1, ND5_p1, ND6_p2, ND6_p3, rRNA_12S, rRNA_16S, tRNAs
2	GTR+I+G	ATP6_p2, COX1_p1, COX1_p2, COX2_p1, COX2_p2, COX3_p1, COX3_p2, CYTB_p2, ND1_p2, ND2_p2, ND3_p2, ND4L_p2, ND4_p2, ND5_p2
3	GTR+I+G	ATP6_p3, COX1_p3, COX2_p3, COX3_p3, CYTB_p3, ND1_p3, ND2_p3, ND3_p3, ND4L_p3, ND4_p3, ND5_p3, ND6_p1

Settings used

alignment : ./mylo_35mitos_concat_seqCat.phylip
branchlengths : unlinked
models : GTR+I+G, GTR+G
model_selection : bic
search : greedy

Best partitioning scheme

Scheme Name : step_39
Scheme lnL : -169349.5486
Scheme BIC : 340923.742014
Number of params : 231
Number of sites : 15222
Number of subsets : 3

RaxML-style partition definitions

DNA, p1 = 1-681\3, 682-876\3, 683-876\3, 684-876\3, 3883-5022\3, 5023-5976\3, 5977-7020\3, 7021-7365\3, 7366-7662\3, 7663-9039\3, 9040-10842\3, 10844-11367\3, 10845-11367\3, 11368-12305, 12306-13766, 13767-15222

DNA, p2 = 2-681\3, 877-2415\3, 878-2415\3, 2416-3099\3, 2417-3099\3, 3100-3882\3, 3101-3882\3, 3884-5022\3, 5024-5976\3, 5978-7020\3, 7022-7365\3, 7367-7662\3, 7664-9039\3, 9041-10842\3

DNA, p3 = 3-681\3, 879-2415\3, 2418-3099\3, 3102-3882\3, 3885-5022\3, 5025-5976\3, 5979-7020\3, 7023-7365\3, 7368-7662\3, 7665-9039\3, 9042-10842\3, 10843-11367\3

Table S3: Detailed results of the PartitionFinder analysis for the seven nuclear exons dataset.

Subset	Best Model	Subset Partitions
1	GTR+G	ADORA3_pos1, ADORA3_pos2, APOB_pos2
2	GTR+G	ADORA3_pos3, APOB_pos3, RAG2_pos3
3	GTR+G	APOB_pos1, BRCA1_pos1
4	GTR+G	BCHE_pos1, RAG2_pos1, TTN_pos1
5	GTR+I+G	BCHE_pos2, RAG2_pos2, TTN_pos2
6	GTR+G	BCHE_pos3, BRCA2_pos3
7	GTR+G	BRCA1_pos2, BRCA2_pos2
8	GTR+G	BRCA1_pos3, TTN_pos3
9	GTR+G	BRCA2_pos1

Settings used

alignment : ./Mylo_7exons_seqCat.phy
branchlengths : linked
models : GTR+I+G, GTR+G
model_selection : bic
search : greedy

Best partitioning scheme

Scheme Name : step_12
Scheme lnL : -61830.71542
Scheme BIC : 125038.535538
Number of params : 143
Number of sites : 15216
Number of subsets : 9

RaxML-style partition definitions

DNA, p1 = 1-312\3, 2-312\3, 314-2724\3
DNA, p2 = 3-312\3, 315-2724\3, 10374-10797\3
DNA, p3 = 313-2724\3, 3706-6483\3
DNA, p4 = 2725-3705\3, 10372-10797\3, 10798-15216\3
DNA, p5 = 2726-3705\3, 10373-10797\3, 10799-15216\3
DNA, p6 = 2727-3705\3, 6486-10371\3
DNA, p7 = 3707-6483\3, 6485-10371\3
DNA, p8 = 3708-6483\3, 10800-15216\3
DNA, p9 = 6484-10371\3

Table S4: Detailed comparisons between our *Mylodon* mitogenome and the one produced by Slater et al. (2016).

Annotated regions	Length (bp)	Identity (%)	Problems in Slater et al. (2016) sequence (KR336794)
tRNA-Phe	70	100	None
12S rRNA	963	99.2	1 divergent region (12 bp) including a 4 bp insertion
tRNA-Val	68	100	None
16S rRNA	1575	98.7	1 substitution + 1 divergent region (26 bp) + 3 Ns
tRNA-Leu	75	100	None
ND1	957	88.7	4 substitutions + 3 divergent regions (26 bp, 14bp, 19bp) including missing start codon + 60 Ns (7 stop codons)
tRNA-Ile	69	100	None
tRNA-Gln	73	100	None
tRNA-Met	69	100	None
ND2	1044	91.0	2 substitutions + 4 divergent regions (88 bp, 41 bp, 51 bp, 6 bp) + 3 deletions + 1 insertion + 4 Ns (8 stop codons)
tRNA-Trp	69	100	None
tRNA-Ala	69	100	None
tRNA-Asn	73	100	None
Origin of L-strand replication	34	100	None
tRNA-Cys	67	100	None
tRNA-Tyr	66	100	None
COX1	1542	90.5	3 substitutions + 43 Ns
tRNA-Ser	69	100	None
tRNA-Asp	68	97.1	1 substitution + 1 insertion
COX2	684	94.2	21 substitutions + 2 divergent regions (10 bp, 19 bp)
tRNA-Lys	69	100	None
ATP8	204	52.5	3 substitutions + 94 gaps (1 stop codon)
ATP6	681	41.8	13 substitutions + 3 divergent regions (39 bp, 29 bp, 34 bp) + 175 Ns + 259 gaps (1 stop codon)
COX3	784	96.4	2 substitutions + 1 divergent region (23 bp) + 1 insertion (10 stop codons)
tRNA-Gly	70	100	None
ND3	346	54.3	7 substitutions + 151 Ns
tRNA-Arg	70	100	None
ND4L	297	59.1	1 divergent region (14 bp) + 110 Ns
ND4	1378	85.7	187 Ns + 2 insertions (29 stop codons)
tRNA-His	69	100	None
tRNA-Ser	60	100	None

tRNA-Leu	70	100	None
ND5	1824	53.5	13 substitutions + 2 divergent regions (10 bp, 3 bp) + 809 gaps + 16 Ns (9 stop codons)
ND6	528	72.2	6 substitutions + 1 divergent region (13 bp) + 4 gaps + 126 Ns (2 stop codons)
tRNA-Glu	69	100	None
CYTB	1140	98.9	11 substitutions + 2 insertions (27 stop codons)
tRNA-Thr	67	29.4	22 gaps + 26 Ns
tRNA-Pro	70	0	70 Ns
Control Region	993	60.0	48 gaps + 310 Ns

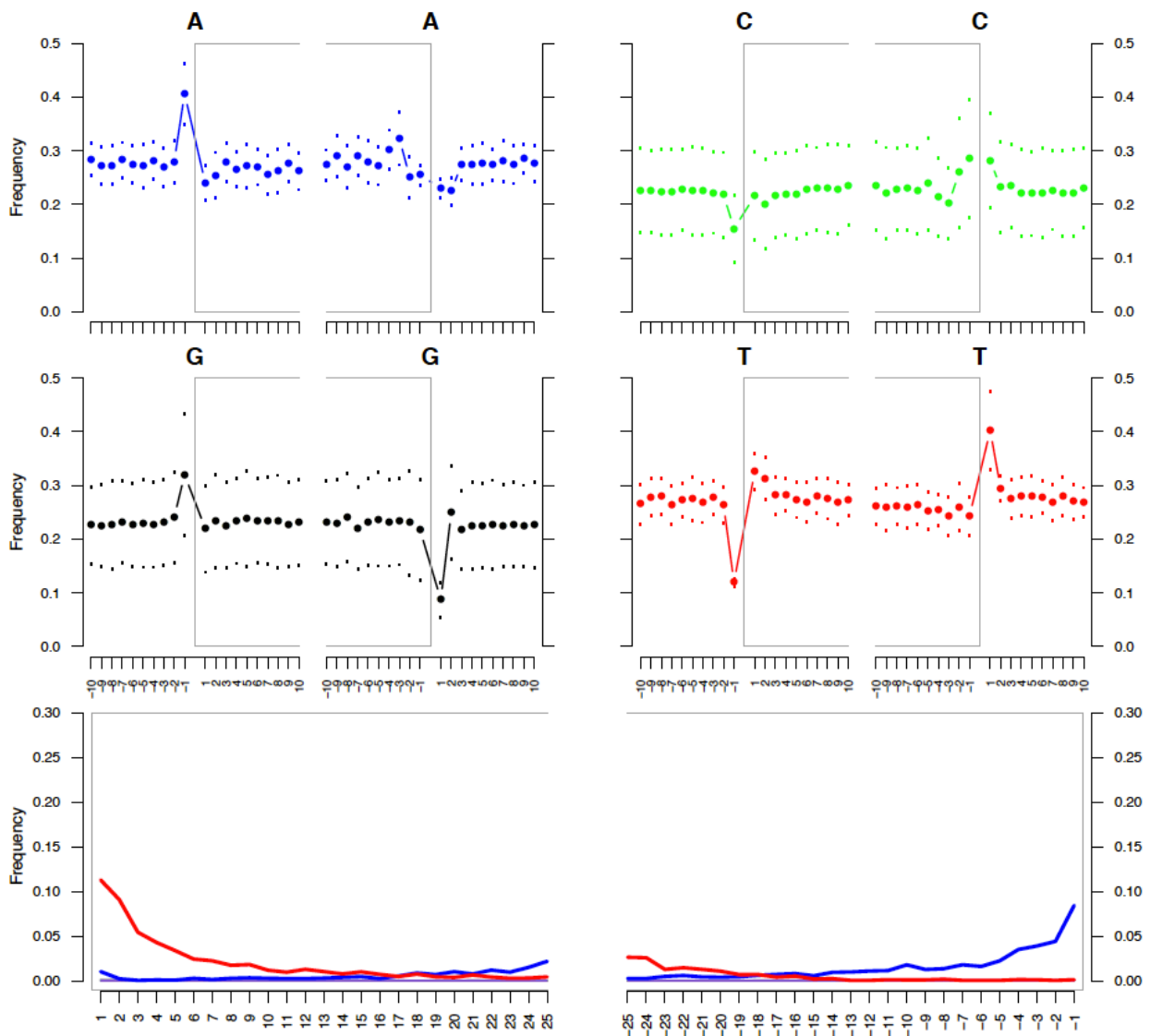


Figure S1. mapDamage DNA fragmentation and nucleotide mis-incorporation plots of the 53,550 cleaned shotgun Illumina reads mapping to the *Myiodon darwinii* SP57 mitogenome. 1st and 2nd rows: DNA fragmentation patterns for the 10 bp preceding read starts (positions -1 to -10 on the left panels of each base composition profile) and the 10 bp following read ends (positions 1 to 10 on the right panels of each base composition profile). 3rd row: Nucleotide mis-incorporation patterns for the first 25 nucleotides and the last 25 sequenced, respectively, with C=>T substitution frequencies (red) and G=>A substitution frequencies (blue).

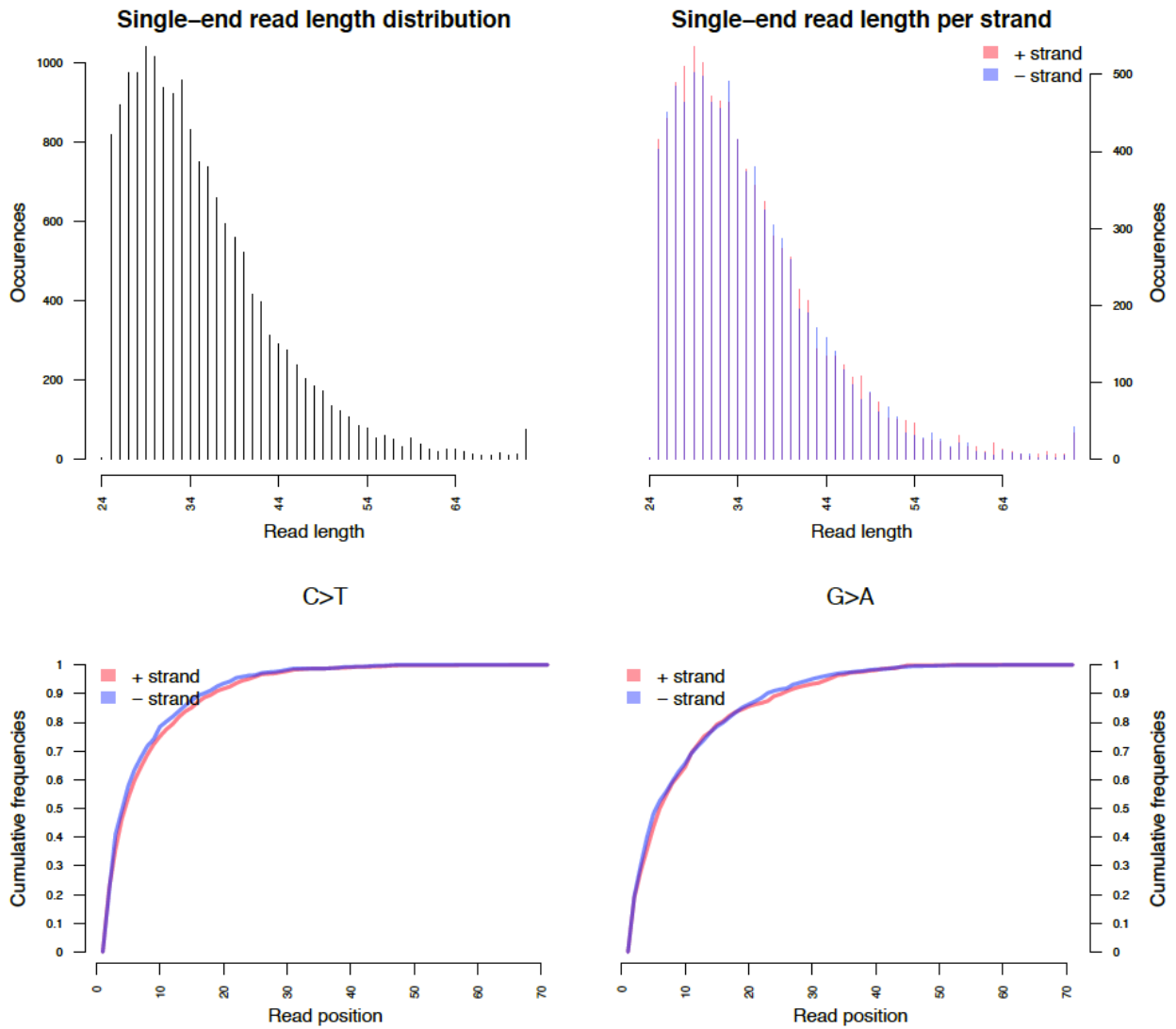


Figure S2. mapDamage fragment length distribution of the 53,550 shotgun cleaned Illumina reads mapping to the *Myiodon darwinii* SP57 mitogenome. Mean fragment length was 30 base pairs.

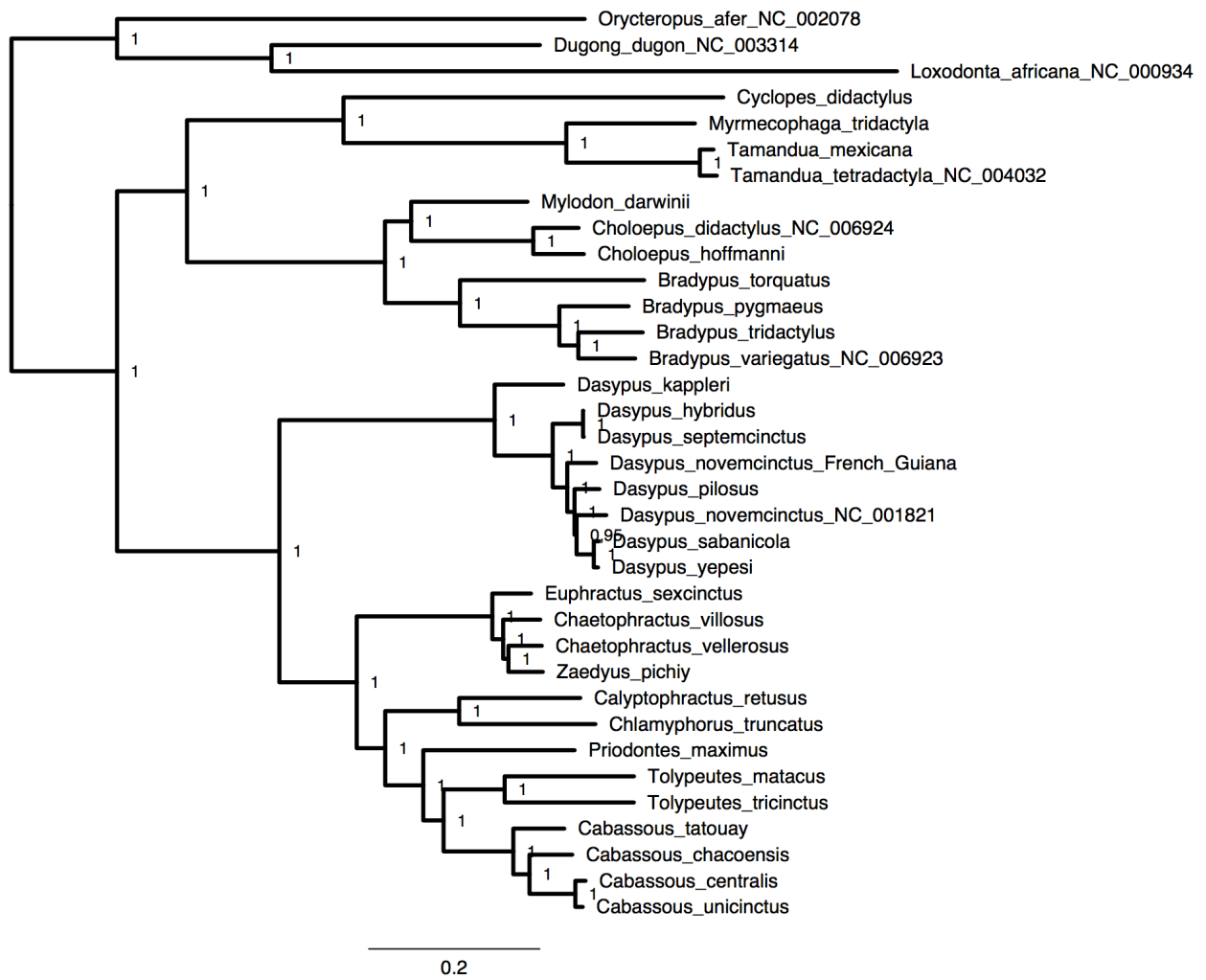


Figure S4: Bayesian consensus mitogenomic tree obtained with MrBayes under a mixed model for each best-fitting partition as estimated by PartitionFinder. Bayesian posterior probabilities are indicated at nodes. Branch lengths represent mean number of substitutions per site.

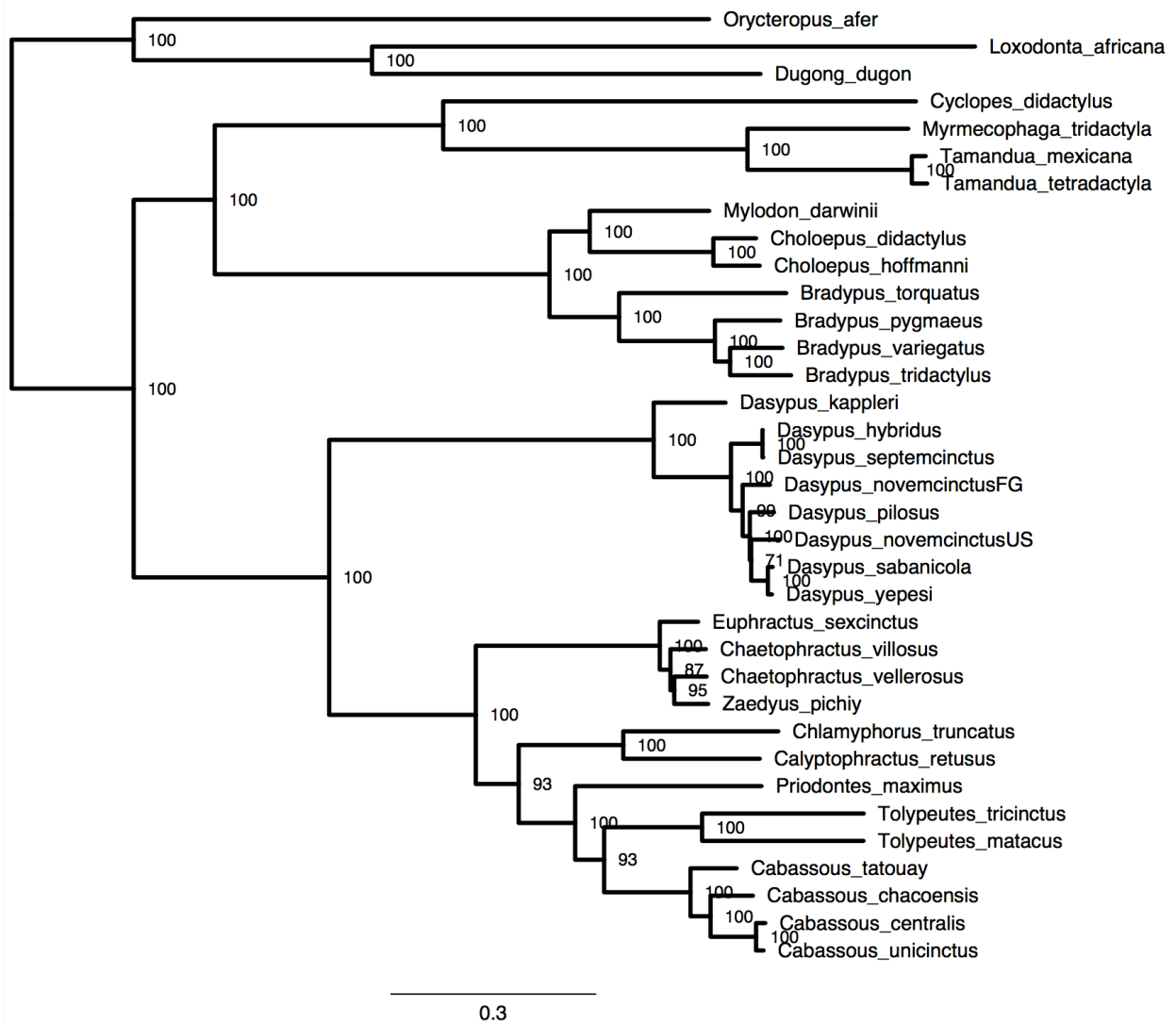


Figure S5: Maximum likelihood mitogenomic tree obtained with RAxML under a GTR+G+I mixed model for each best-fitting partition as estimated by PartitionFinder. Bootstrap percentages are indicated at nodes. Branch lengths represent mean number of substitutions per site.

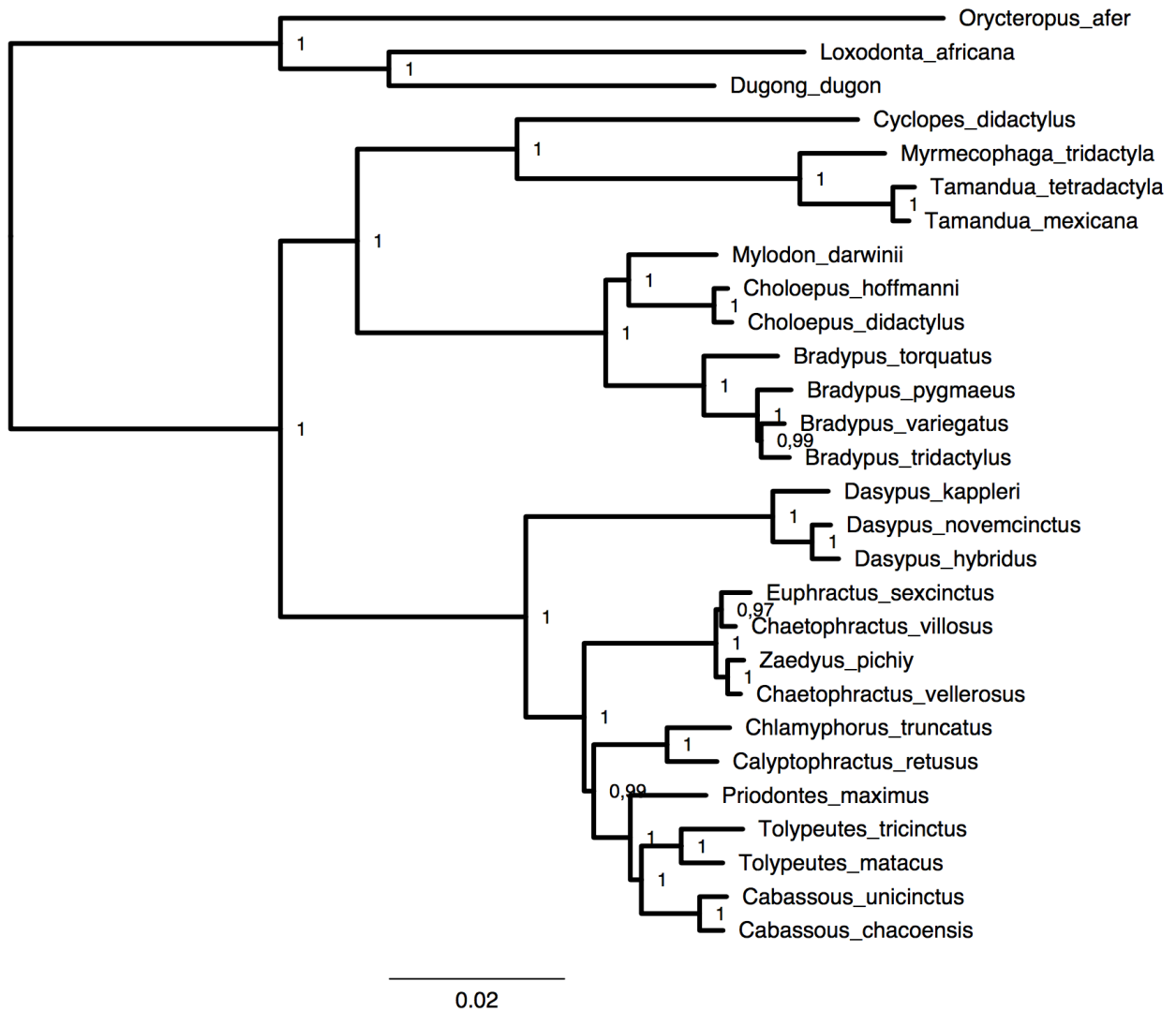


Figure S6: Bayesian consensus nuclear tree (concatenation of 7 exons) obtained with PhyloBayes under the CAT-GTR+G₄ mixture model. Bayesian posterior probabilities are indicated at nodes. Branch lengths represent mean number of substitutions per site.

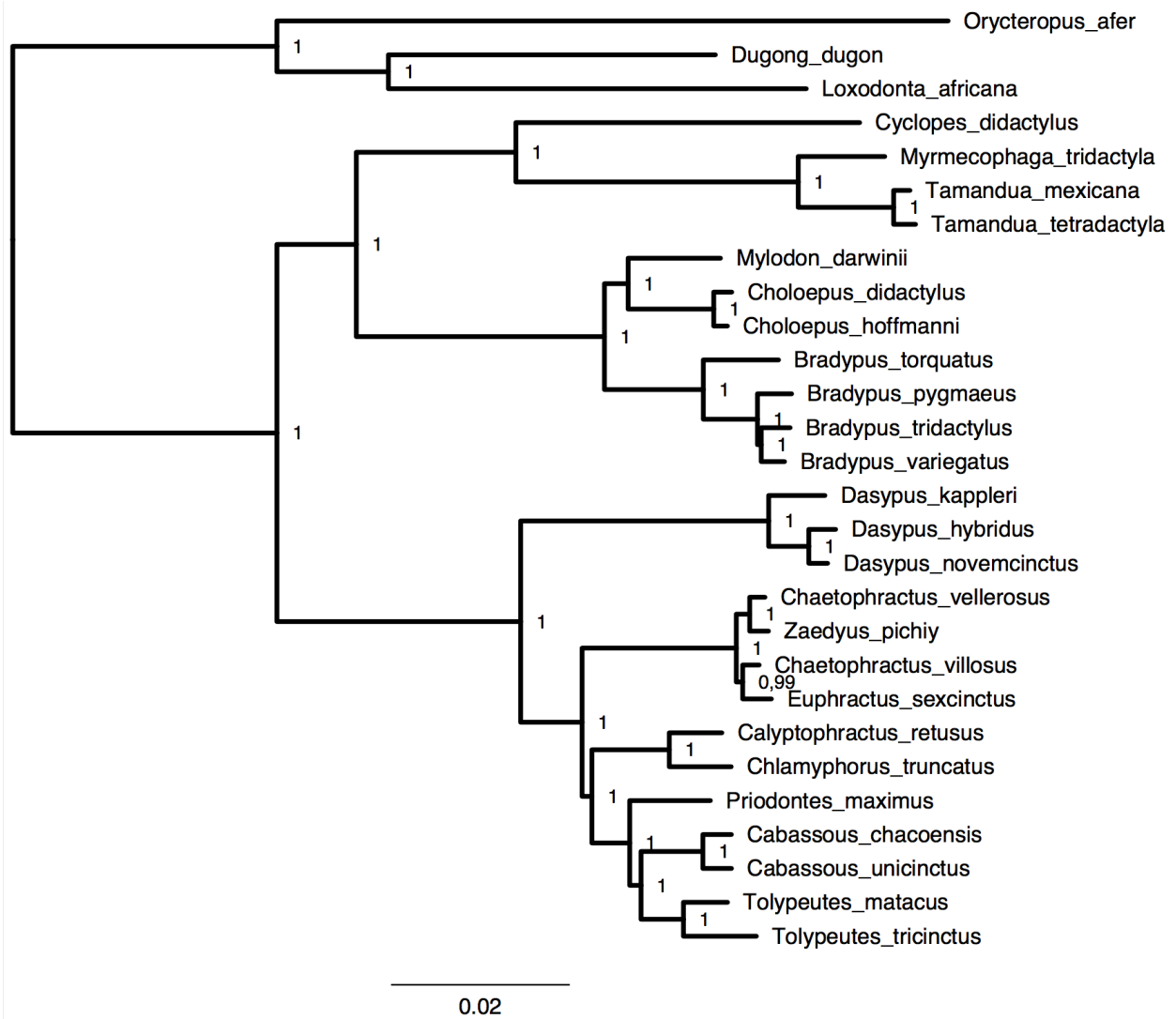


Figure S7: Bayesian consensus nuclear tree (concatenation of 7 exons) obtained with MrBayes under a mixed model for each best-fitting partition as estimated by PartitionFinder. Bayesian posterior probabilities are indicated at nodes. Branch lengths represent mean number of substitutions per site.

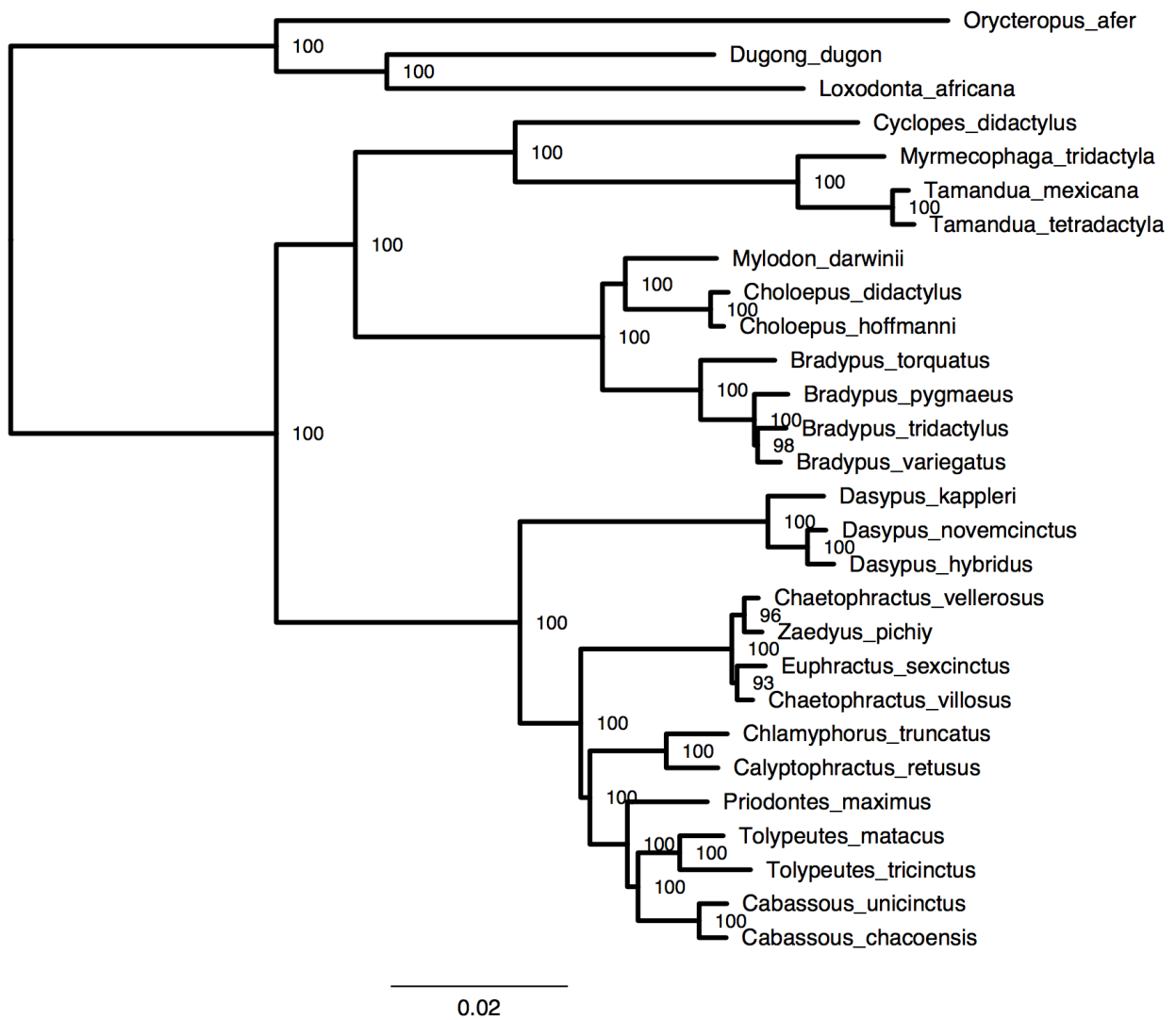


Figure S8: Maximum likelihood nuclear tree (concatenation of 7 exons) obtained with RAxML under a GTR+G mixed model for each best-fitting partition as estimated by PartitionFinder. Bootstrap percentages are indicated at nodes. Branch lengths represent mean number of substitutions per site.

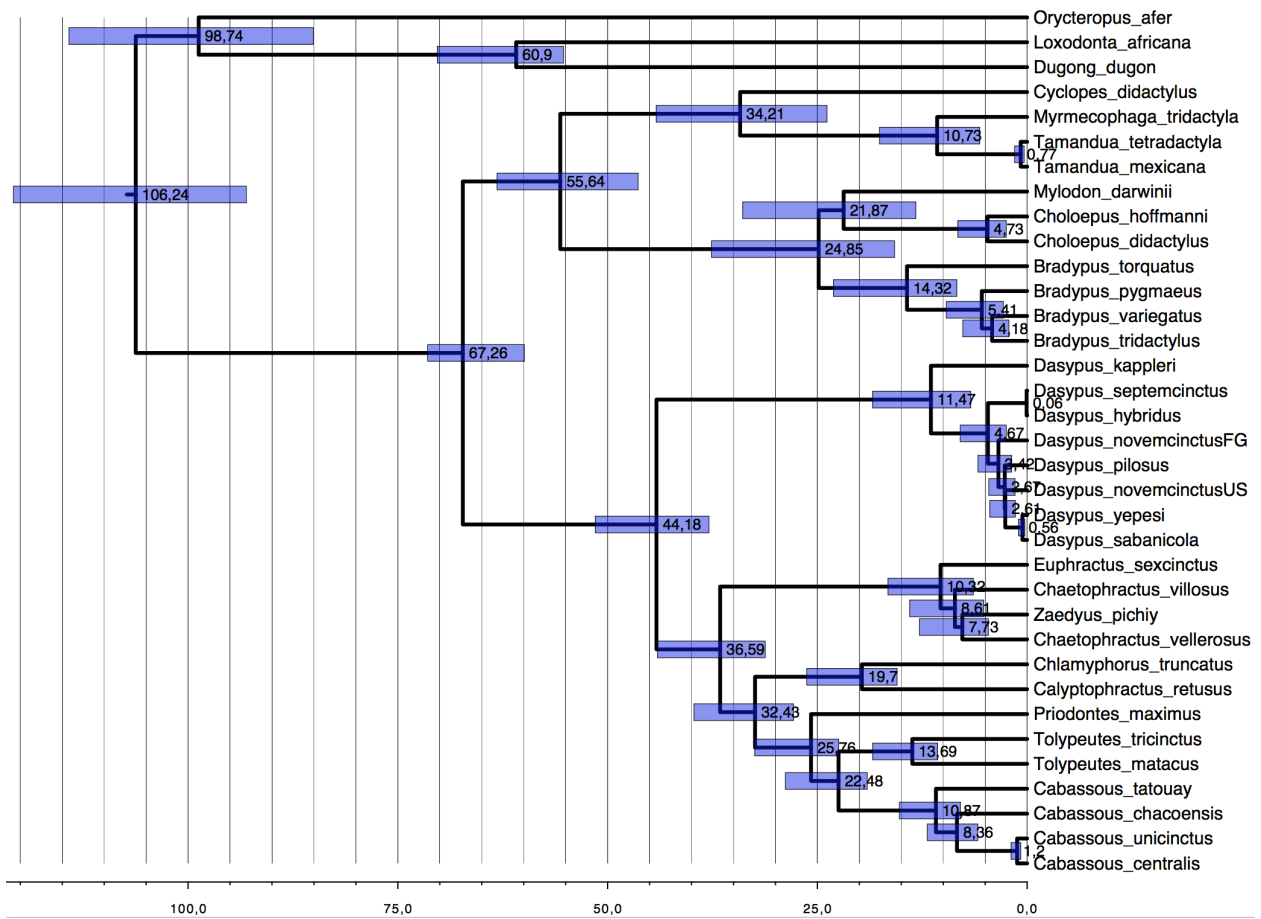


Figure S9: Bayesian mitogenomic chronogram obtained with PhyloBayes under the CAT-GTR+G₄ mixture model and an autocorrelated lognormal model of clock relaxation. Timescale is in million years ago.

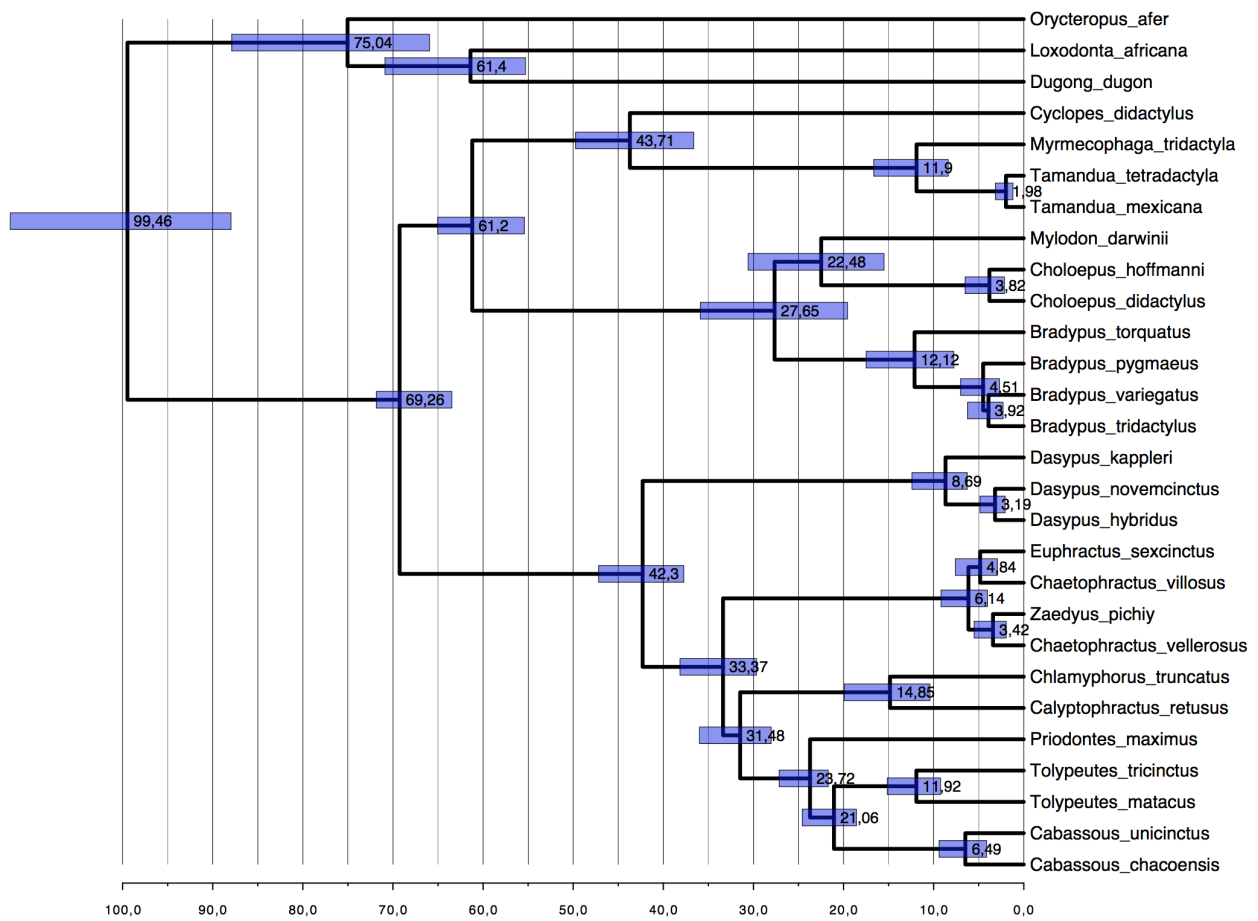


Figure S10: Bayesian nuclear chronogram (concatenation of 7 exons) obtained with PhyloBayes under the CAT-GTR+G₄ mixture model and an autocorrelated lognormal model of clock relaxation. Timescale is in million years ago.

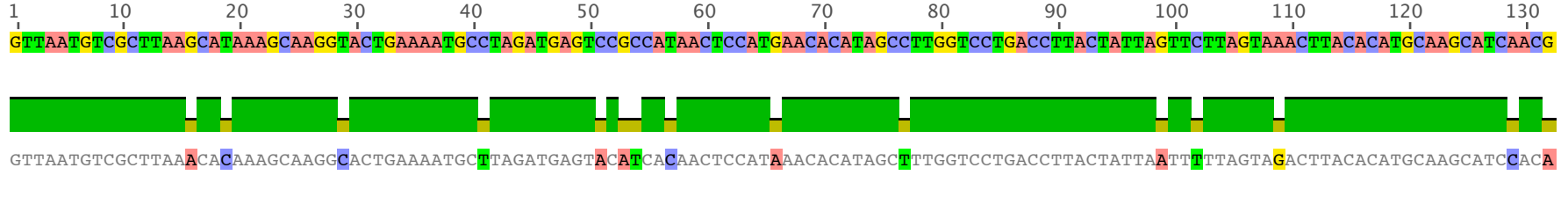
Figure S11: Comparison of available *Mylodon* mitogenomes with the *Choloepus didactylus* reference mitogenome.

1 10 20 30 40 50 60 70 80 90 100 110 120 130

Consensus
Translation

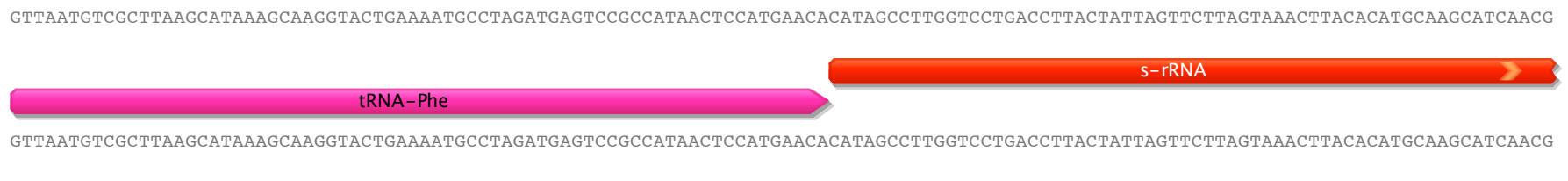
Identity

Choloepus_didactylus_NC
Translation



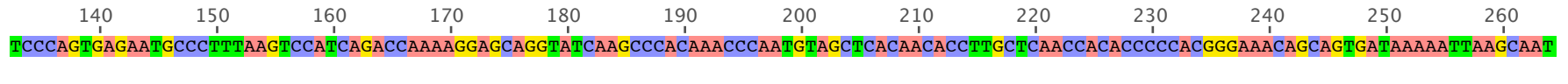
Myiodon_darwinii_Delsuc
Translation

Myiodon_darwinii_Slater
Translation



140 150 160 170 180 190 200 210 220 230 240 250 260

Consensus
Translation



Identity

Choloepus_didactylus_NC
Translation

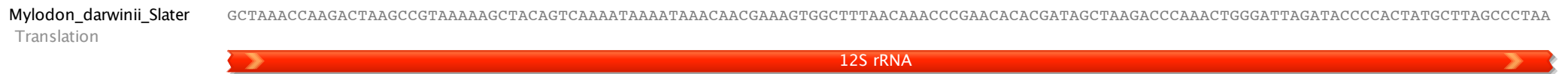
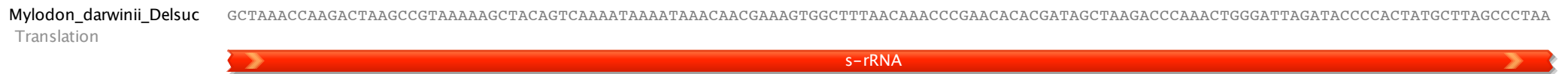
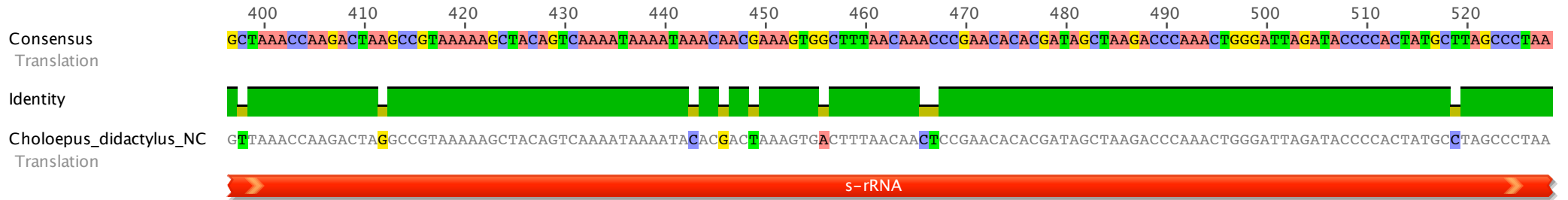
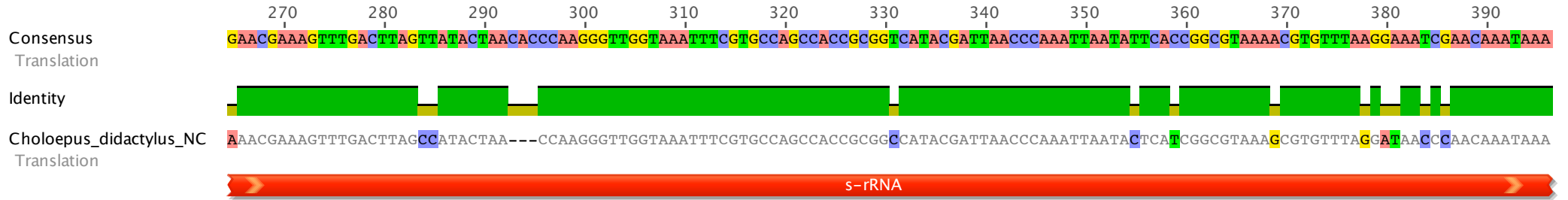


Myiodon_darwinii_Delsuc
Translation

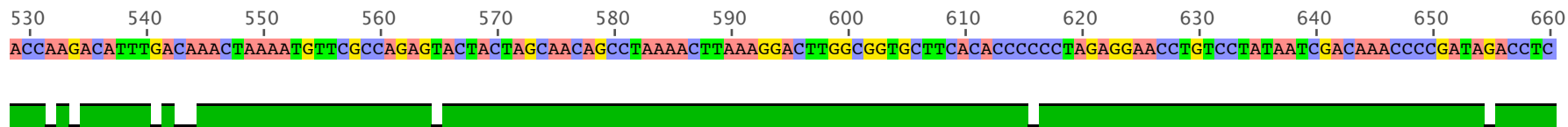


Myiodon_darwinii_Slater
Translation





530 540 550 560 570 580 590 600 610 620 630 640 650 660
Consensus
Translation
Identity



Choloepus didactylus_NC
Translation

ACCCAAACATTTTATCAACTAAAATGTTCCGCCAGAGCACTACTAGCAACAGCCTAAAACCTAAAGGACTTGGCGGTGCTTCACACCCCTAGAGGAACCTGTCTATAATCGACAAAACCCCGATAGACCTC



Myiodon darwinii_Delsuc
Translation

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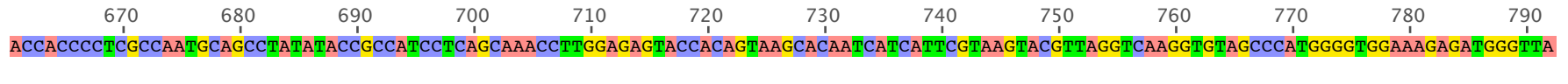


Myiodon darwinii_Slater
Translation

ACCAAGACATTTGACAAACTAAAATGTTCCGCCAGAGTACTACTAGCAACAGCCTAAAACCTAAAGGACTTGGCGGTGCTTCACACCCCCCTAGAGGAACCTGTCTATAATCGACAAAACCCCGATAGACCTC



670 680 690 700 710 720 730 740 750 760 770 780 790
Consensus
Translation
Identity



Choloepus didactylus_NC
Translation

ACCACCCTAGCCAAACACAGCCTATATACCGCCATCCAGCAGCAACCTTGGAGAGTACCACAGTAAGCACAATCATCATTCGTAAGTACGTTAGGTCAAGGTGTAGCCCATGAGGTGGAAAGAGATGGGTTA



Myiodon darwinii_Delsuc
Translation

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Myiodon darwinii_Slater
Translation

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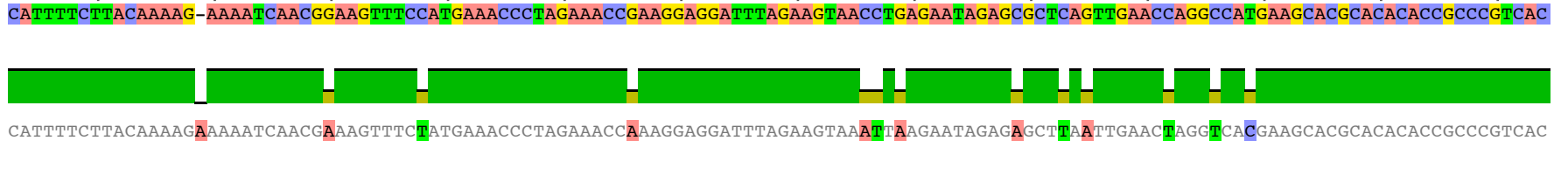


800 810 820 830 840 850 860 870 880 890 900 910 920

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation



Myiodon_darwinii_Delsuc
Translation



Myiodon_darwinii_Slater
Translation

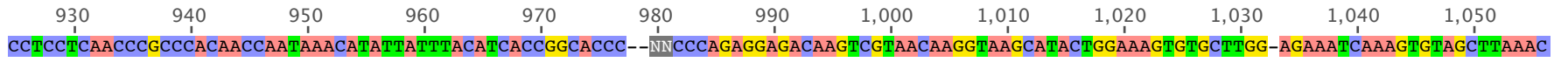


930 940 950 960 970 980 990 1,000 1,010 1,020 1,030 1,040 1,050

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

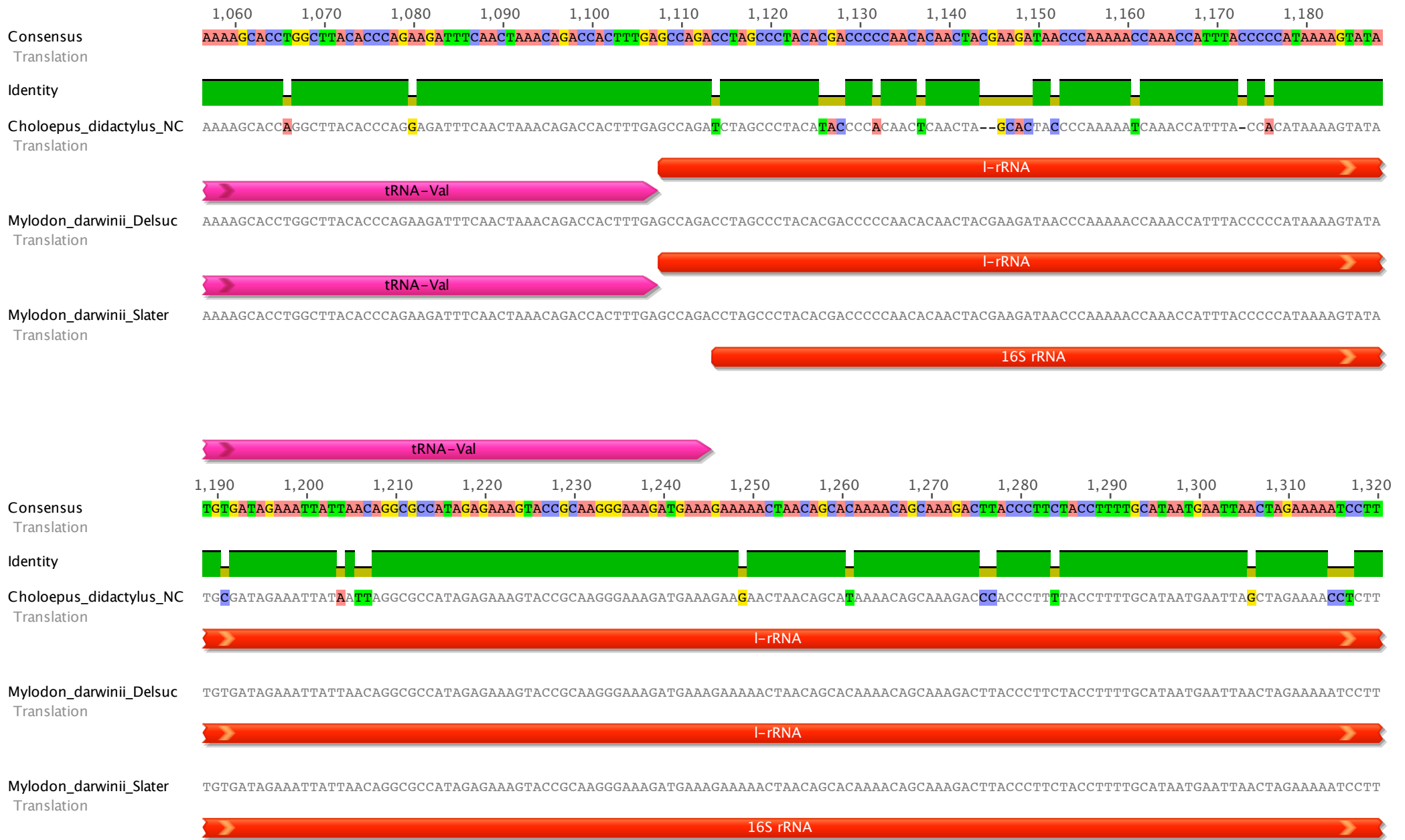


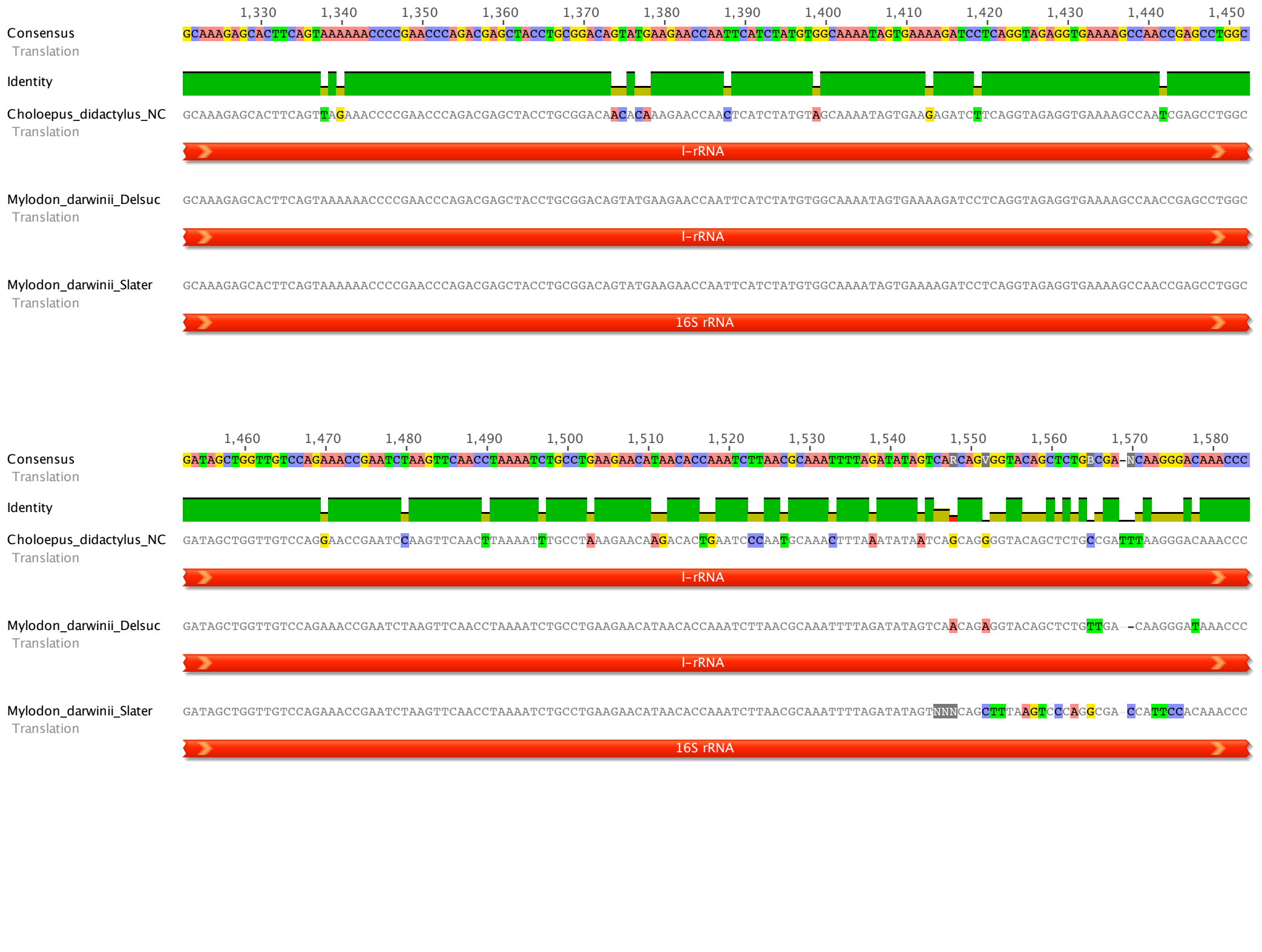
Myiodon_darwinii_Delsuc
Translation



Myiodon_darwinii_Slater
Translation







1,590 1,600 1,610 1,620 1,630 1,640 1,650 1,660 1,670 1,680 1,690 1,700 1,710
Consensus
Translation
Identity

Choloepus_didactylus_NC
Translation
I-rRNA

Myiodon_darwinii_Delsuc
Translation
I-rRNA

Myiodon_darwinii_Slater
Translation
16S rRNA

1,720 1,730 1,740 1,750 1,760 1,770 1,780 1,790 1,800 1,810 1,820 1,830 1,840
Consensus
Translation
Identity

Choloepus_didactylus_NC
Translation
I-rRNA

Myiodon_darwinii_Delsuc
Translation
I-rRNA

Myiodon_darwinii_Slater
Translation
16S rRNA

1,850 1,860 1,870 1,880 1,890 1,900 1,910 1,920 1,930 1,940 1,950 1,960 1,970 1,980

Consensus
Translation
 ACCACTAAACATATTCAAGTCTNCCCCACTGTTAACCCAACACAGGTGTGCATACTGGAAAGATTAAAAGAAGCAGAAGGAACTCGGCAAATATAAATCCCGCCTGTTTACCAAAAACATCACCTCTAGCAT

Identity

Choloepus_didactylus_NC
Translation
 ACCACCAAATACCACCGAGTCTTACCCATTGTTAACCCAACACAGGCRGTGGCACTGGAAAGATTAAAAGAAGTAAAGGAACTCGGCAAAACACAAATCCCGCCTGTTTACCAAAAACATCACCTCTAGCAT



Myiodon_darwinii_Delsuc
Translation
 ACCACTAAACATATTCAAGTCTCCCCACTGTTAACCCAACACAGGTGTGCATACTGGAAAGATTAAAAGAAGCAGAAGGAACTCGGCAAAATATAAATCCCGCCTGTTTACCAAAAACATCACCTCTAGCAT



Myiodon_darwinii_Slater
Translation
 ACCACTAAACATATTCAAGTCT-CCCCACTGTTAACCCAACACAGGTGTGCATACTGGAAAGATTAAAAGAAGCAGAAGGAACTCGGCAAAATATAAATCCCGCCTGTTTACCAAAAACATCACCTCTAGCAT



1,990 2,000 2,010 2,020 2,030 2,040 2,050 2,060 2,070 2,080 2,090 2,100 2,110

Consensus
Translation
 AACAACTATTAGAGGCACCTGCCTGCCAGTGACATATGTTAAACGGCCGCGGTACCCTGACCGTGCAAAGGTAGCATAATCATTGTTCCTTAATTAAGGACTTGTATGAAAGGCCCAACGAGGATTTTACT

Identity

Choloepus_didactylus_NC
Translation
 AACAACTATTAGAGGCACCTGCCTGCCAGTGACATATGTTAAACGGCCGCGGTACCCTGACCGTGCAAAGGTAGCATAATCATTGTTCCTTAATTAAGGACTTGTATGAAAGGCCCAACGAGGATTTTACT

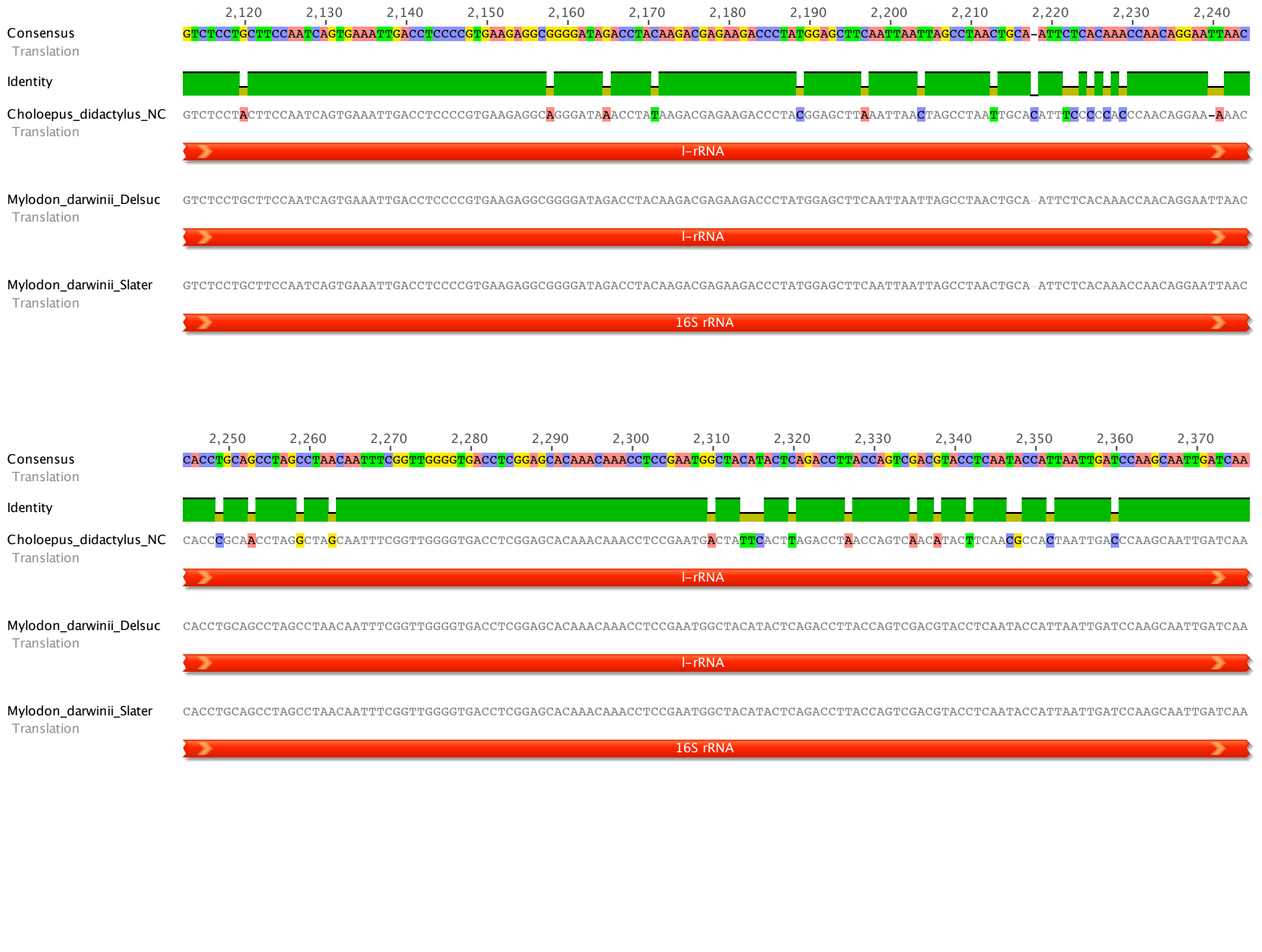


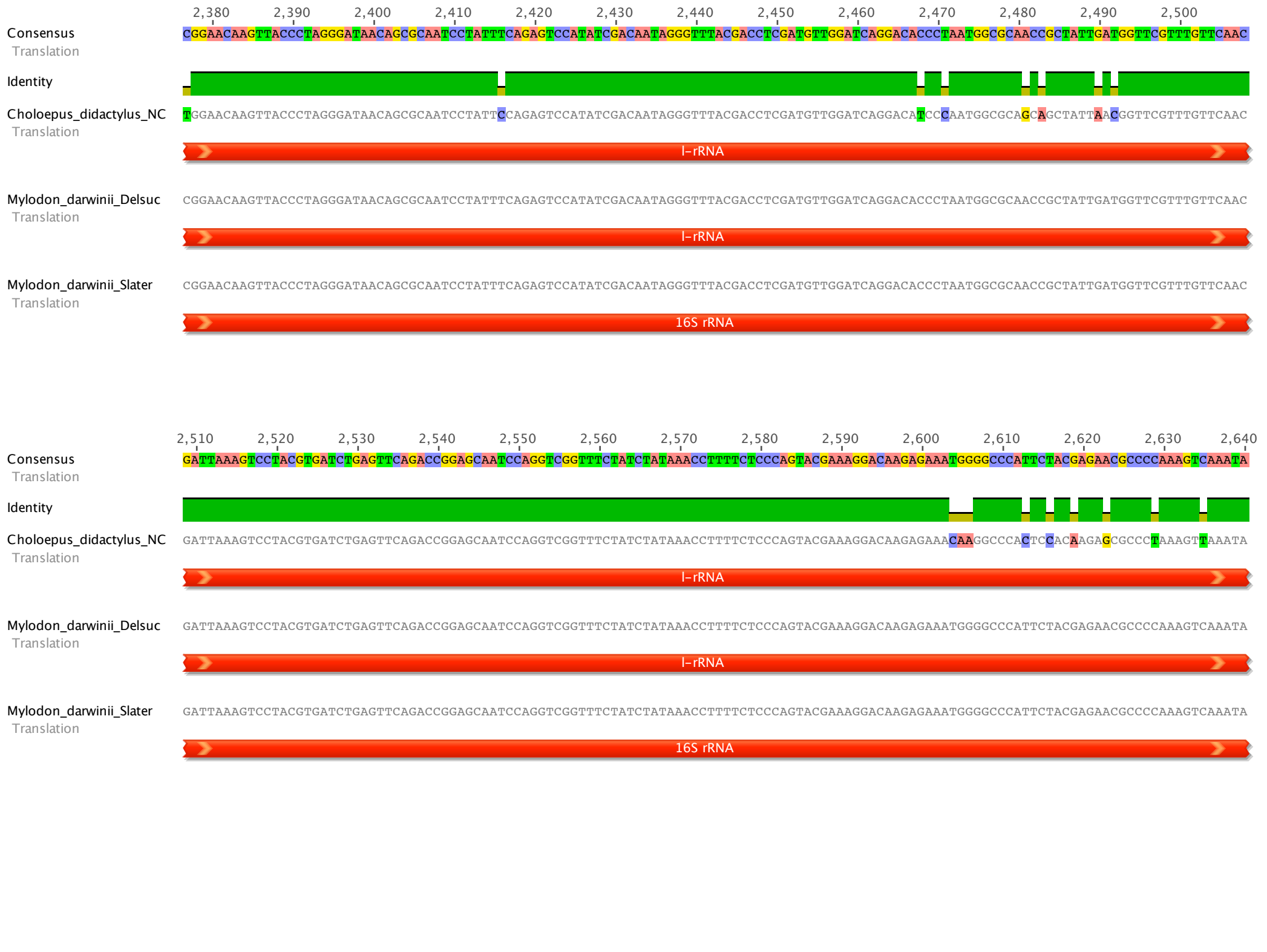
Myiodon_darwinii_Delsuc
Translation
 AACAACTATTAGAGGCACCTGCCTGCCAGTGACATATGTTAAACGGCCGCGGTACCCTGACCGTGCAAAGGTAGCATAATCATTGTTCCTTAATTAAGGACTTGTATGAAAGGCCCAACGAGGATTTTACT

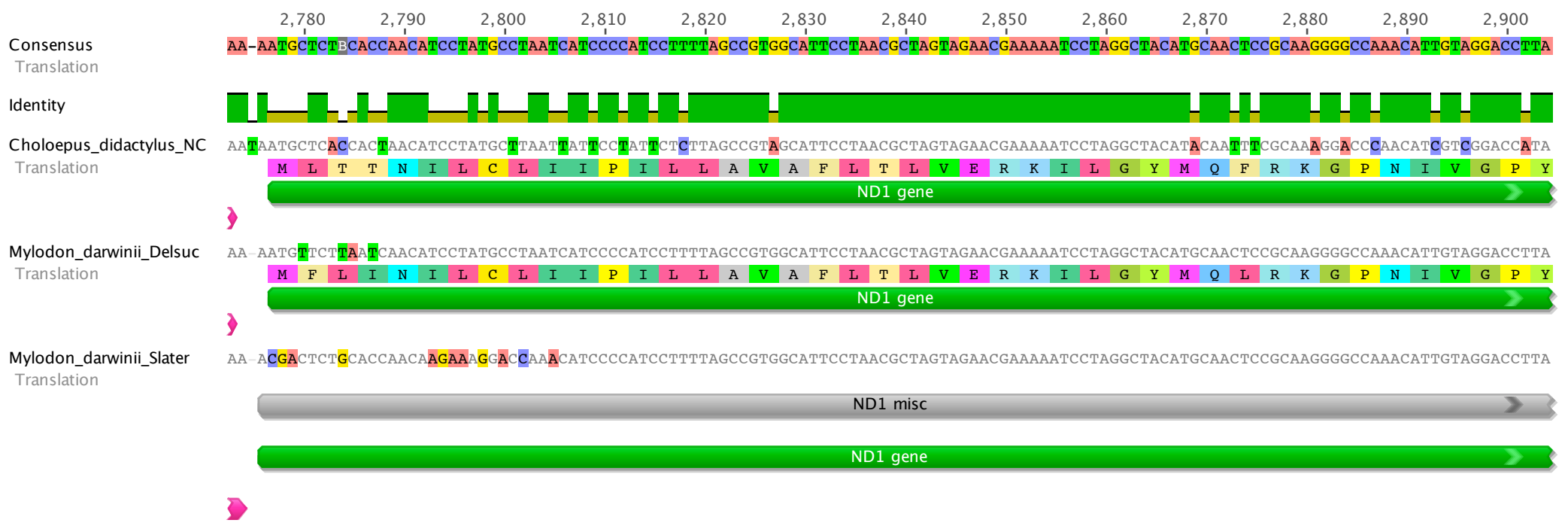


Myiodon_darwinii_Slater
Translation
 AACAACTATTAGAGGCACCTGCCTGCCAGTGACATATGTTAAACGGCCGCGGTACCCTGACCGTGCAAAGGTAGCATAATCATTGTTCCTTAATTAAGGACTTGTATGAAAGGCCCAACGAGGATTTTACT









2,910 2,920 2,930 2,940 2,950 2,960 2,970 2,980 2,990 3,000 3,010 3,020 3,030

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

ND1 gene

Myiodon_darwinii_Delsuc
Translation

ND1 gene

Myiodon_darwinii_Slater
Translation

ND1 misc

ND1 gene

3,040 3,050 3,060 3,070 3,080 3,090 3,100 3,110 3,120 3,130 3,140 3,150 3,160

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

ND1 gene

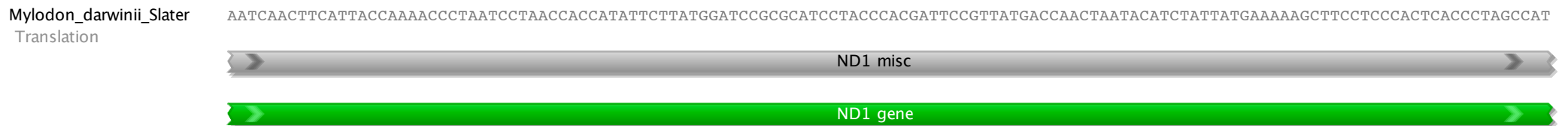
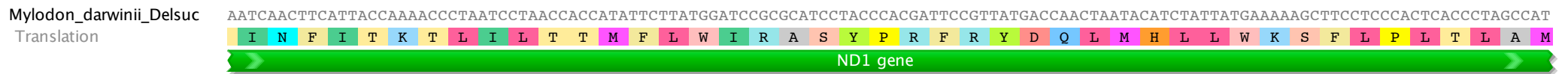
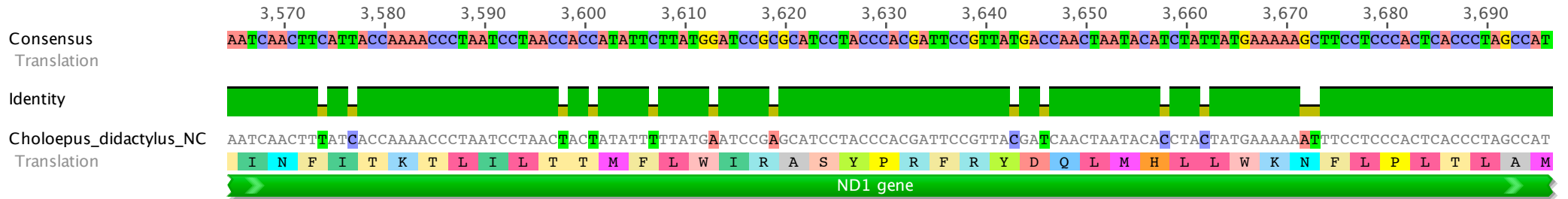
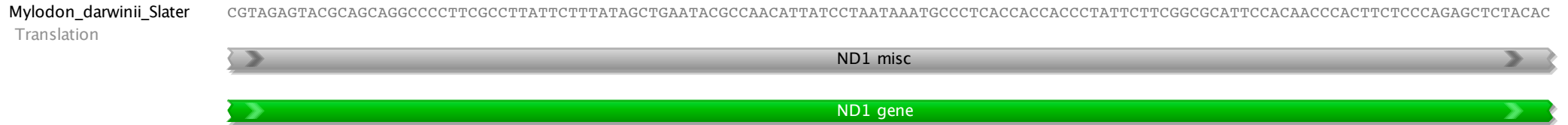
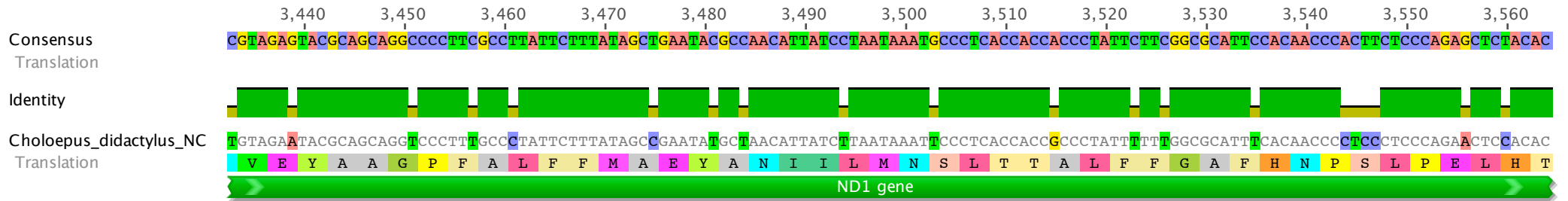
Myiodon_darwinii_Delsuc
Translation

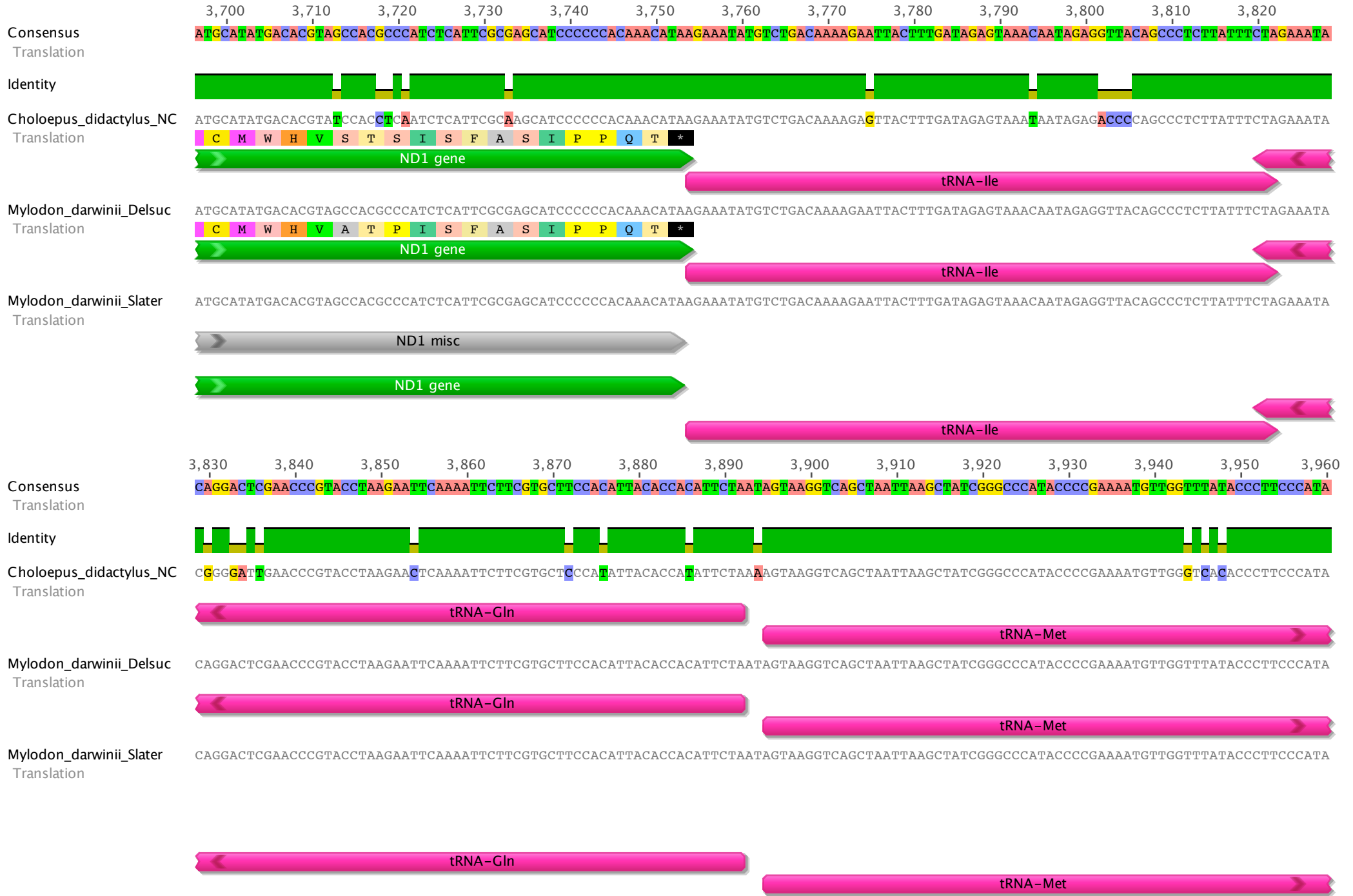
ND1 gene

Myiodon_darwinii_Slater
Translation

ND1 misc

ND1 gene





3,970 3,980 3,990 4,000 4,010 4,020 4,030 4,040 4,050 4,060 4,070 4,080 4,090

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

Myiodon_darwinii_Delsuc
Translation

Myiodon_darwinii_Slater
Translation

4,100 4,110 4,120 4,130 4,140 4,150 4,160 4,170 4,180 4,190 4,200 4,210 4,220

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

Myiodon_darwinii_Delsuc
Translation

Myiodon_darwinii_Slater
Translation

4,230 4,240 4,250 4,260 4,270 4,280 4,290 4,300 4,310 4,320 4,330 4,340 4,350

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

ND2 gene

Myiodon_darwinii_Delsuc
Translation

ND2 gene

Myiodon_darwinii_Slater
Translation

ND2 misc

ND2 gene

4,360 4,370 4,380 4,390 4,400 4,410 4,420 4,430 4,440 4,450 4,460 4,470 4,480

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

ND2 gene

Myiodon_darwinii_Delsuc
Translation

ND2 gene

Myiodon_darwinii_Slater
Translation

ND2 misc

ND2 gene

4,490 4,500 4,510 4,520 4,530 4,540 4,550 4,560 4,570 4,580 4,590 4,600 4,610 4,620

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

ND2 gene

Myiodon_darwinii_Delsuc
Translation

ND2 gene

Myiodon_darwinii_Slater
Translation

ND2 misc

ND2 gene

4,630 4,640 4,650 4,660 4,670 4,680 4,690 4,700 4,710 4,720 4,730 4,740 4,750

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

ND2 gene

Myiodon_darwinii_Delsuc
Translation

ND2 gene

Myiodon_darwinii_Slater
Translation

ND2 misc

ND2 gene

4,760 4,770 4,780 4,790 4,800 4,810 4,820 4,830 4,840 4,850 4,860 4,870 4,880

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

L P K W M I I Q E M T K N N L I V M P T M M A I L A L L N L Y F Y M R L I Y S T S L T L

ND2 gene

Myiodon_darwini_Delsuc
Translation

L P K W M I I Q E M T K N G L I I M P T V M A I L A L L N L Y F Y M R L I Y S T S L T L

ND2 gene

Myiodon_darwini_Slater
Translation

ND2 misc

ND2 gene

4,890 4,900 4,910 4,920 4,930 4,940 4,950 4,960 4,970 4,980 4,990 5,000 5,010

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

F P T T N N T K I K W Q F K P T T P P P L T P T L I I L S T T I I P L S P M L C I L E *

ND2 gene

Myiodon_darwini_Delsuc
Translation

F P S T N N T K I K W Q F K S T T P S P I T P T L I A L S T M I I P L S P M L C T L E *

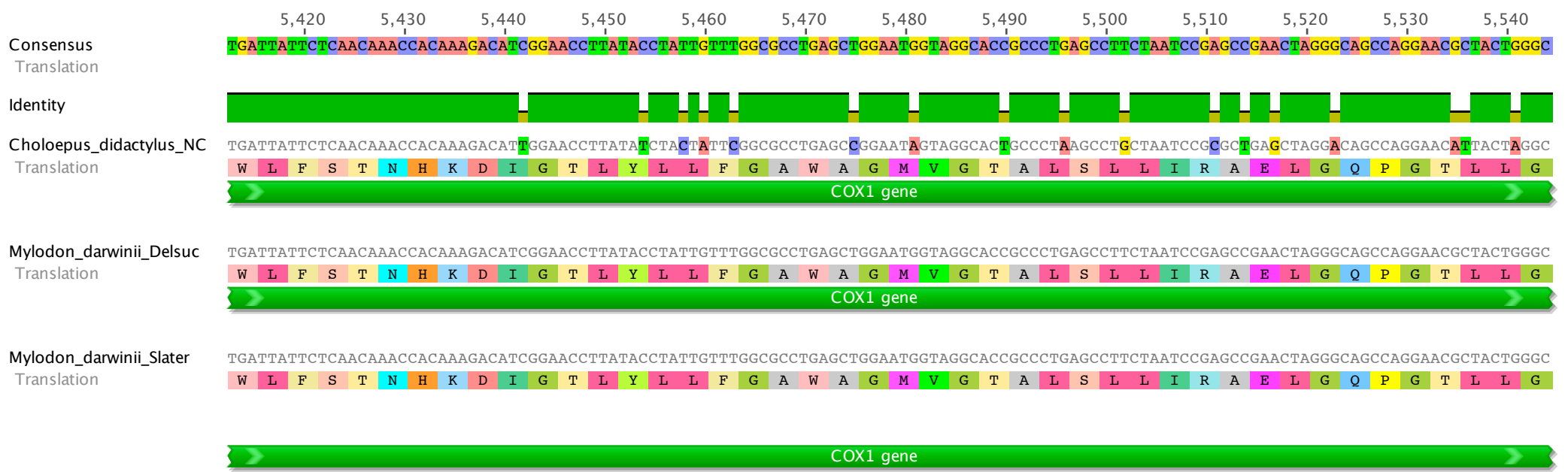
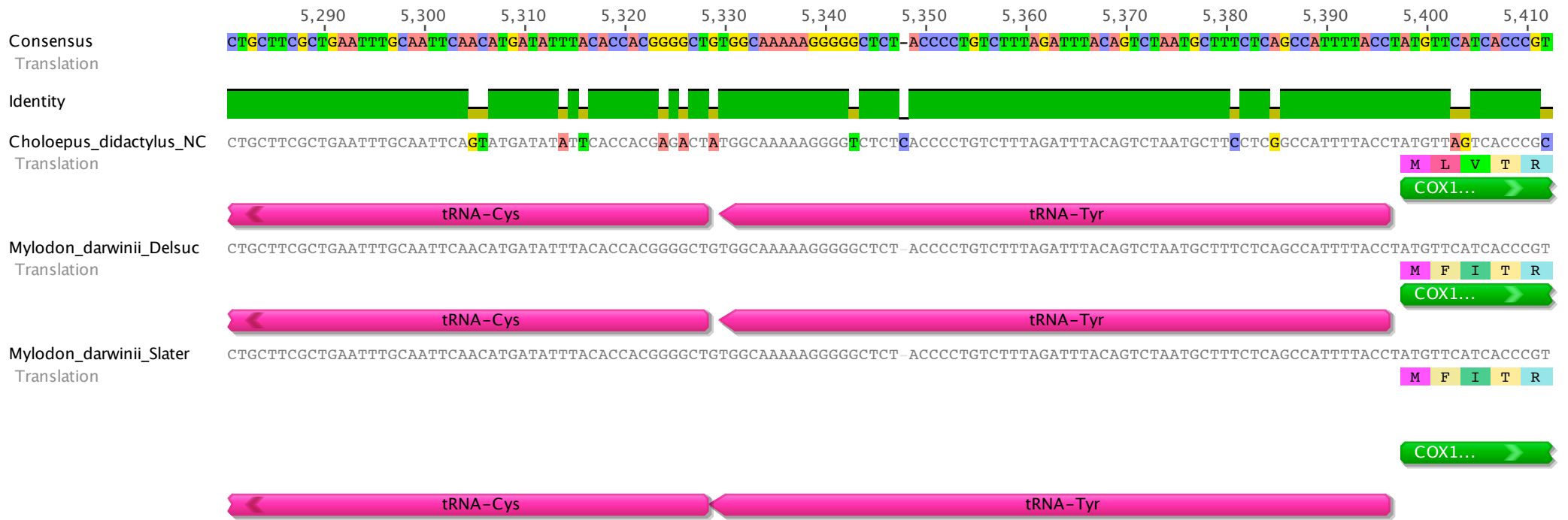
ND2 gene

Myiodon_darwini_Slater
Translation

ND2 misc

ND2 gene





5,550 5,560 5,570 5,580 5,590 5,600 5,610 5,620 5,630 5,640 5,650 5,660 5,670

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

D D Q I Y N V I V T A H A F V M I F F M V M P I M I G G F G N W L V P L M I G A P D M A

COX1 gene

Myiodon_darwinii_Delsuc
Translation

D D Q I Y N V I V T A H A F V M I F F M V M P I M I G G F G N W L V P L M I G A P D M A

COX1 gene

Myiodon_darwinii_Slater
Translation

D D Q I Y N V I V T A H A F V M I F F M V M P I M I G G F G N W L V P L M I G A P N M A

COX1 gene

5,680 5,690 5,700 5,710 5,720 5,730 5,740 5,750 5,760 5,770 5,780 5,790 5,800

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

F P R M N N M S F W L L P P S F L L L L A S S M V E A G A G T G W T V Y P P L A G N L A

COX1 gene

Myiodon_darwinii_Delsuc
Translation

F P R M N N M S F W L L P P S F L L L L A S S M V E A G A G T G W T V Y P P L A G N L A

COX1 gene

Myiodon_darwinii_Slater
Translation

F P R M N N M S F W L L P P S F L L L L A S S M V E A G A G T G W T V Y P P L A G N L A

COX1 gene

5,810 5,820 5,830 5,840 5,850 5,860 5,870 5,880 5,890 5,900 5,910 5,920 5,930 5,940

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

COX1 gene

Myiodon_darwinii_Delsuc
Translation

COX1 gene

Myiodon_darwinii_Slater
Translation

COX1 gene

5,950 5,960 5,970 5,980 5,990 6,000 6,010 6,020 6,030 6,040 6,050 6,060 6,070

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

COX1 gene

Myiodon_darwinii_Delsuc
Translation

COX1 gene

Myiodon_darwinii_Slater
Translation

COX1 gene

6,080 6,090 6,100 6,110 6,120 6,130 6,140 6,150 6,160 6,170 6,180 6,190 6,200

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

GGTGACCCCATCCTATAACCAACATCTATTCTGATTCTTCGGACACCCCTGAGGTCTACATTCTTATCCTCCCAGGCTTCGGCATGATTTCTCACATTGTACATATTACTCTGGAAAAAAGGAACCATTTCGGC

G D P I L Y Q H L F W F F G H P E V Y I L I L P G F G M I S H I V T Y Y S G K K E P F G

COX1 gene

Myiodon_darwinii_Delsuc
Translation

GGTGACCCCATCCTATAACCAACATCTATTCTGATTCTTCGGACACCCCTGAGGTCTACATTCTTATCCTCCCAGGCTTCGGCATGATTTCTCACATTGTACATATTACTCTGGAAAAAAGGAACCATTTCGGC

G D P I L Y Q H L F W F F G H P E V Y I L I L P G F G M I S H I V T Y Y S G K K E P F G

COX1 gene

Myiodon_darwinii_Slater
Translation

GGTGACCCCATCCTATAACCAACATCTATTCTGATTCTTCGGACACCCCTGAGGTCTACATTCTTATCCTCCCAGGCTTCGGCATGATTTCTCACATTGTACATATTACTCTGGAAAAAAGGAACCATTTCGGC

G D P I L Y Q H L F W F F G H P E V Y I L I L P G F G M I S H I V T Y Y S G K K E P F G

COX1 gene

6,210 6,220 6,230 6,240 6,250 6,260 6,270 6,280 6,290 6,300 6,310 6,320 6,330

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

TATATGGGTATAGTAGTGAGCTATAAATGTCGATTGGATTTCCTAGGGTTCATTGTCTGAGCTCACCACATGTTACAGTAGGAATGGACGTGGACACACGAGCATACTTCACATCAGCGACCATAATTATTGCT

Y M G M V W A M M S I G F L G F I V W A H H M F T V G M D V D T R A Y F T S A T M I I A

COX1 gene

Myiodon_darwinii_Delsuc
Translation

TACATAGGGATAGTGTGAGCTATAAATGTCGATTGGATTTCCTAGGGTTCATTGTCTGAGCTCACCACATGTTACAGTAGGAATGGACGTGGACACACGAGCATACTTCACATCAGCGACCATAATTATTGCT

Y M G M V W A M M S I G F L G F I V W A H H M F T V G M D V D T R A Y F T S A T M I I A

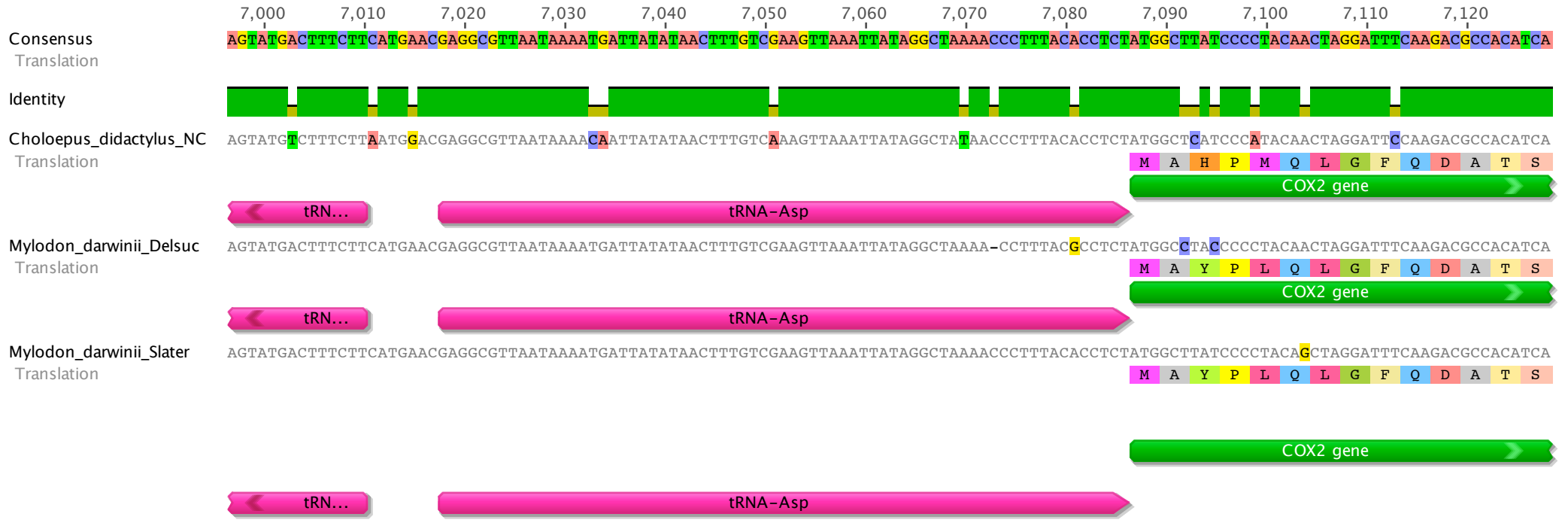
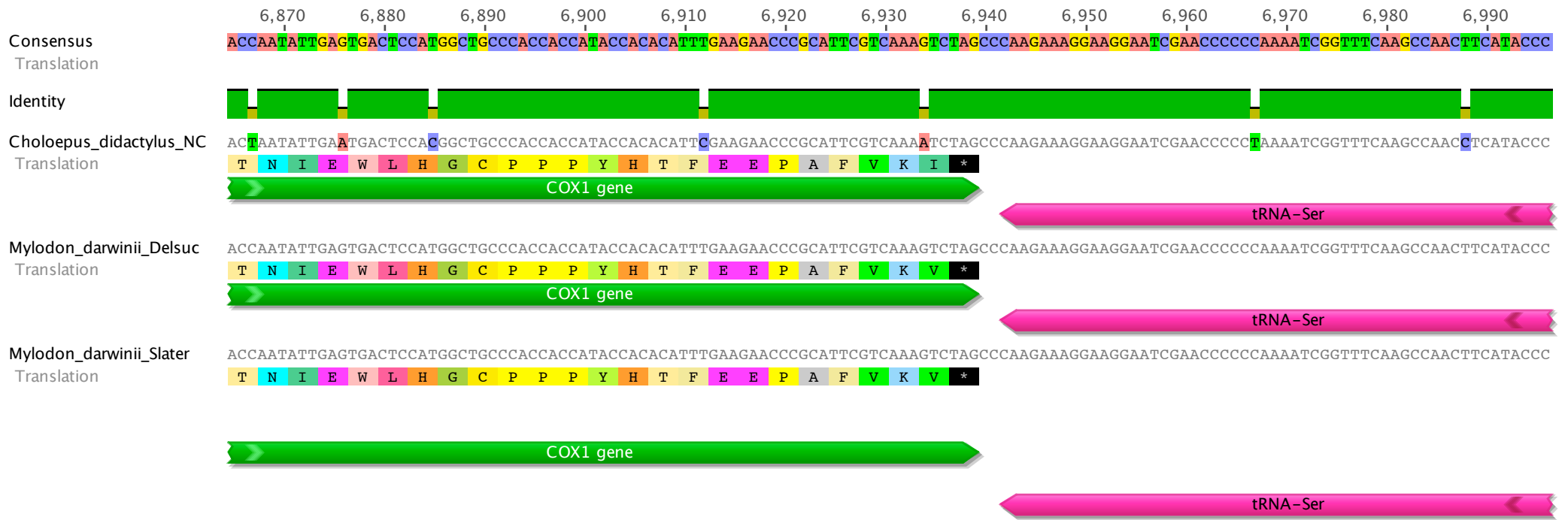
COX1 gene

Myiodon_darwinii_Slater
Translation

TACATAGGGATAGTGTGAGCTATAAATGTCGATTGGATTTCCTAGGGTTCATTGTCTGAGCTCACCACATGTTACAGTAGGAATGGACGTGGACACACGAGCATACTTCACATCAGCGACCATAATTATTGCT

Y M G M V W A M M S I G F L G F I V W A H H M F T V G M D V D T R A Y F T S A T M I I A

COX1 gene



7,130 7,140 7,150 7,160 7,170 7,180 7,190 7,200 7,210 7,220 7,230 7,240 7,250 7,260

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

COX2 gene

Myiodon_darwinii_Delsuc
Translation

COX2 gene

Myiodon_darwinii_Slater
Translation

COX2 gene

7,270 7,280 7,290 7,300 7,310 7,320 7,330 7,340 7,350 7,360 7,370 7,380 7,390

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

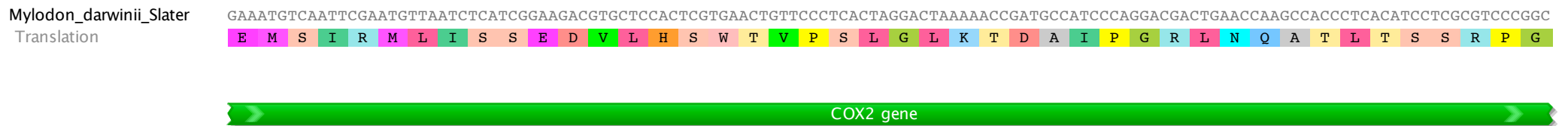
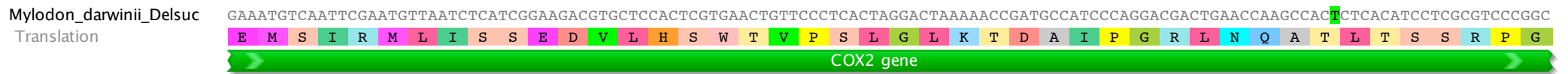
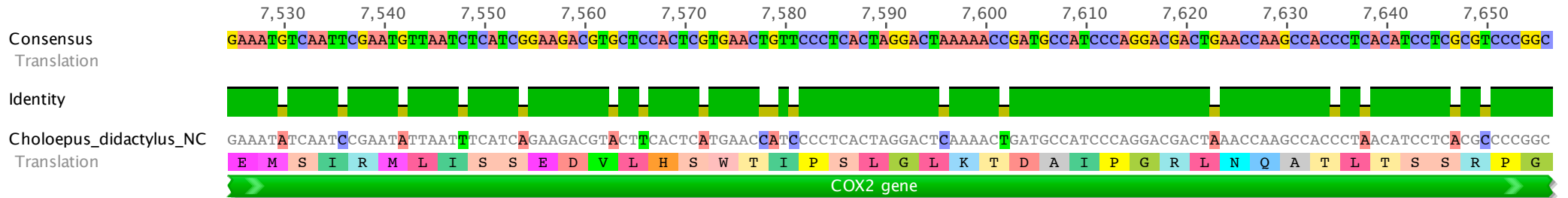
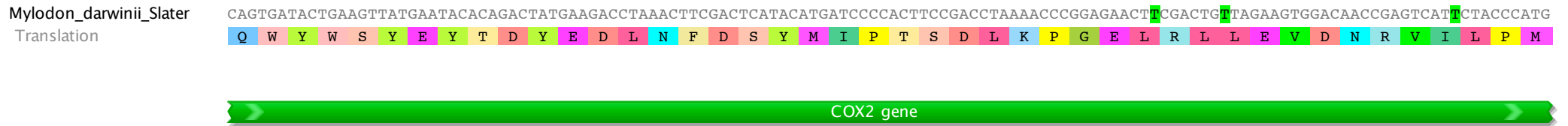
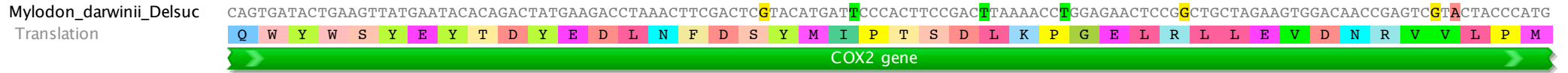
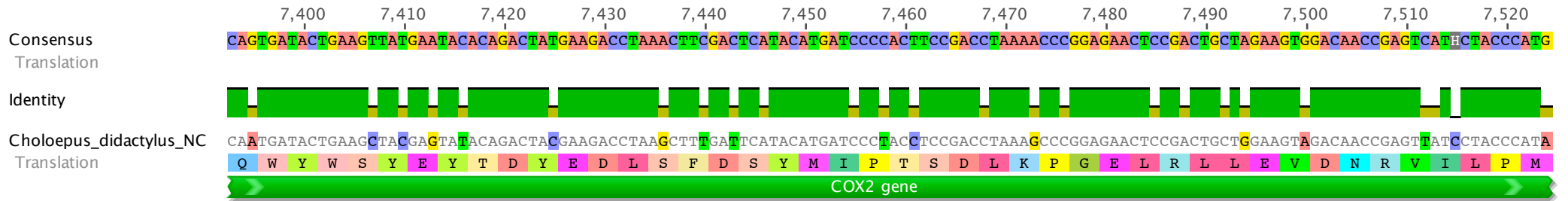
COX2 gene

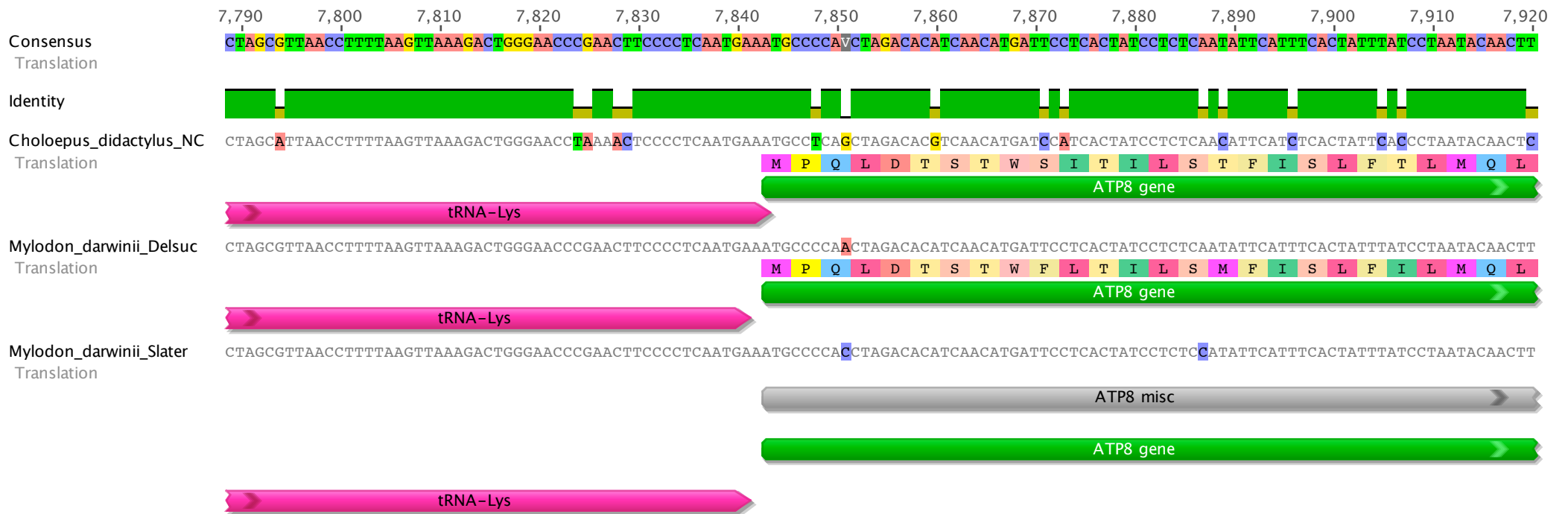
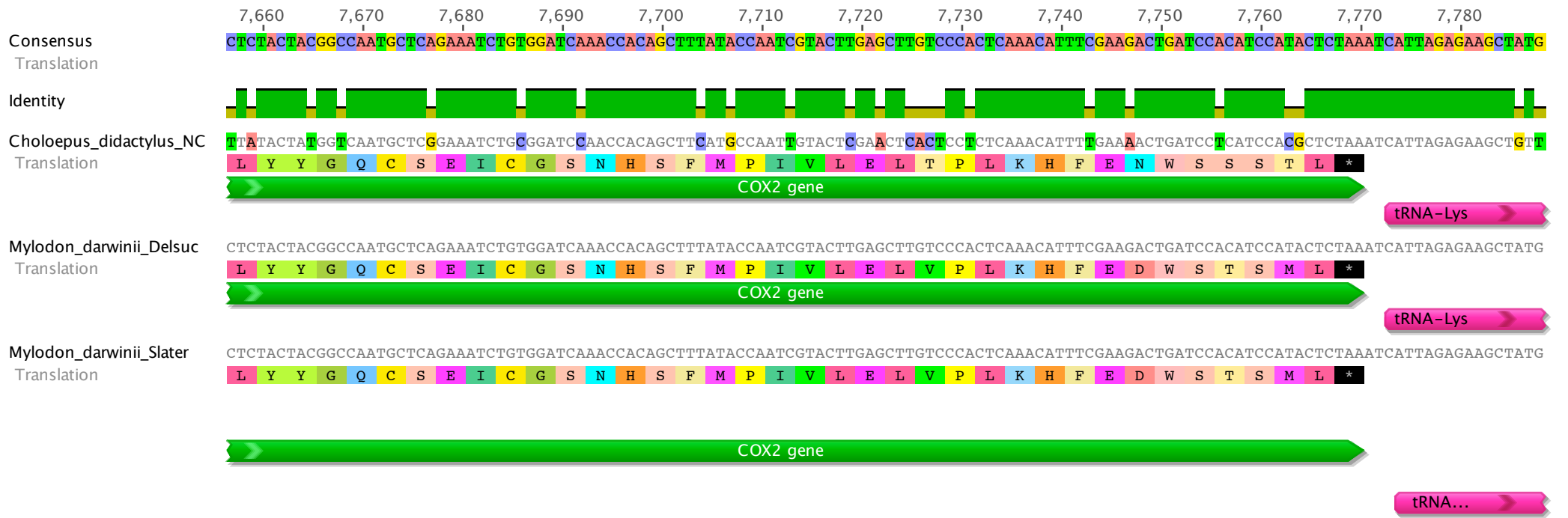
Myiodon_darwinii_Delsuc
Translation

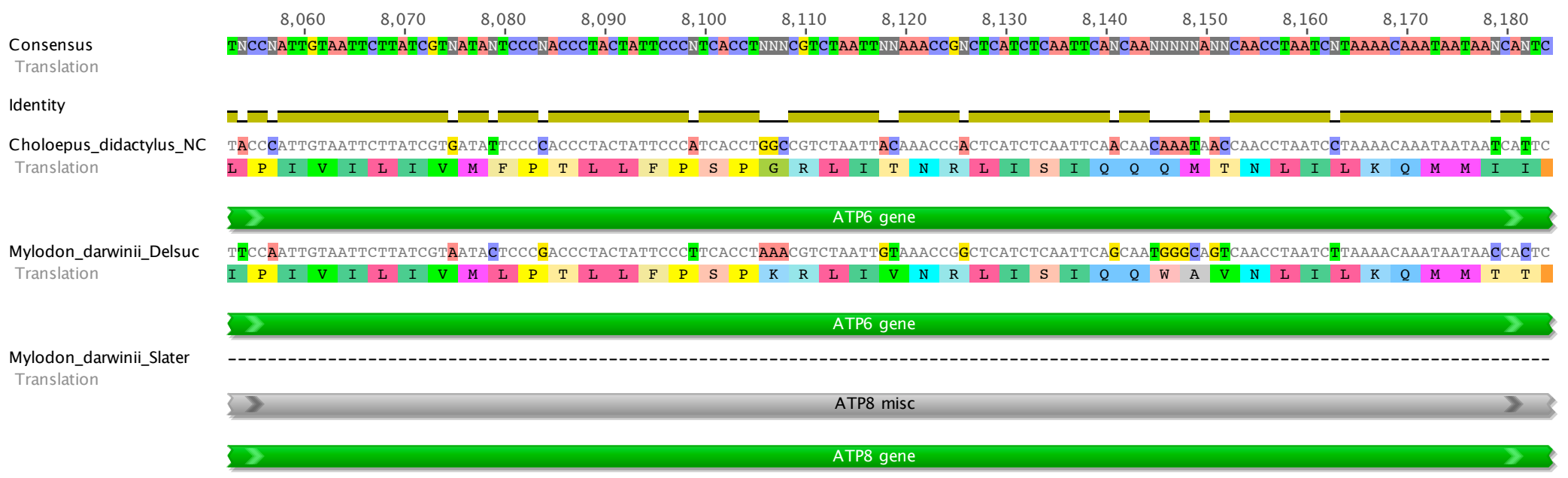
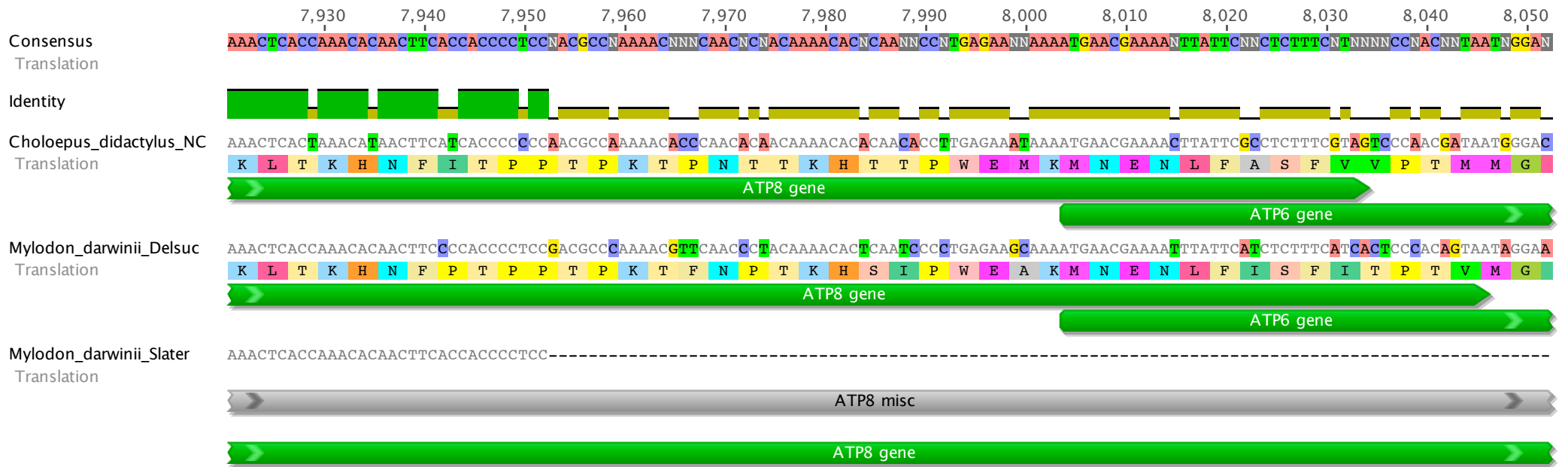
COX2 gene

Myiodon_darwinii_Slater
Translation

COX2 gene







8,190 8,200 8,210 8,220 8,230 8,240 8,250 8,260 8,270 8,280 8,290 8,300 8,310

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation



Myiodon_darwinii_Delsuc
Translation



Myiodon_darwinii_Slater
Translation



8,320 8,330 8,340 8,350 8,360 8,370 8,380 8,390 8,400 8,410 8,420 8,430 8,440

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation



Myiodon_darwinii_Delsuc
Translation



Myiodon_darwinii_Slater
Translation



8,450 8,460 8,470 8,480 8,490 8,500 8,510 8,520 8,530 8,540 8,550 8,560 8,570 8,580

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation



Myiodon_darwinii_Delsuc
Translation



Myiodon_darwinii_Slater
Translation



8,590 8,600 8,610 8,620 8,630 8,640 8,650 8,660 8,670 8,680 8,690 8,700 8,710

Consensus
Translation

Identity

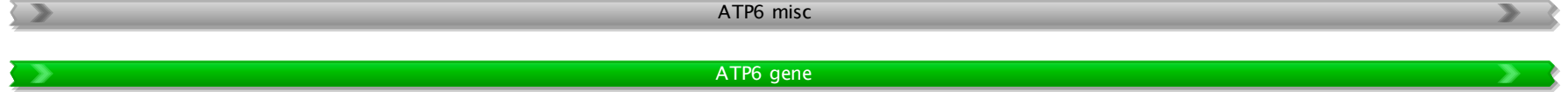
Choloepus_didactylus_NC
Translation

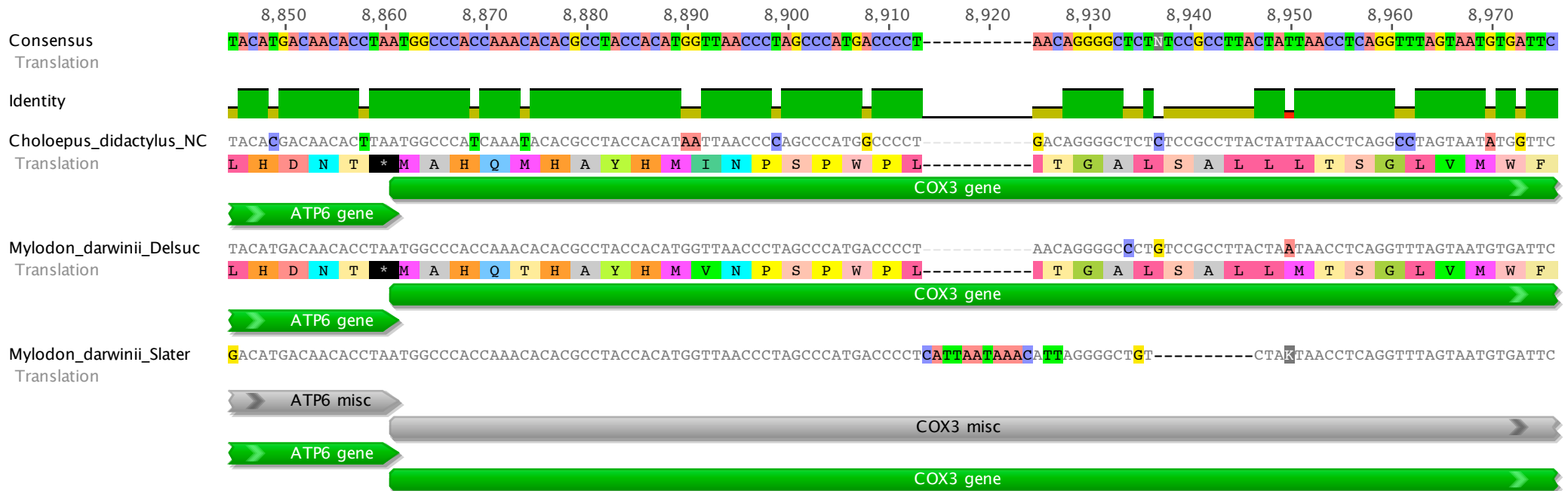
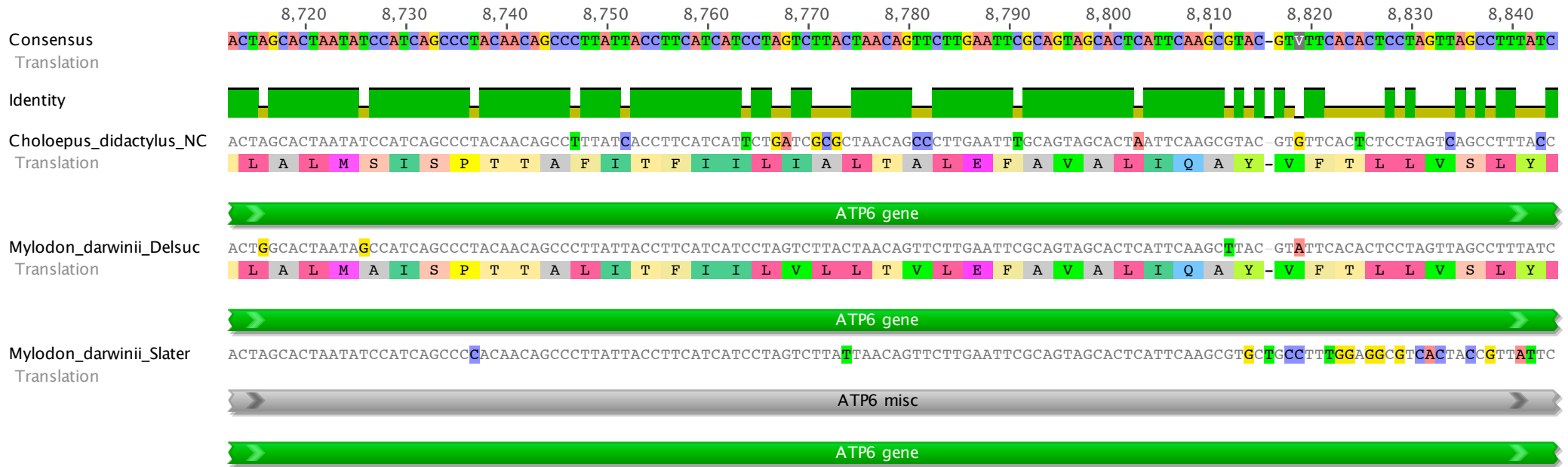


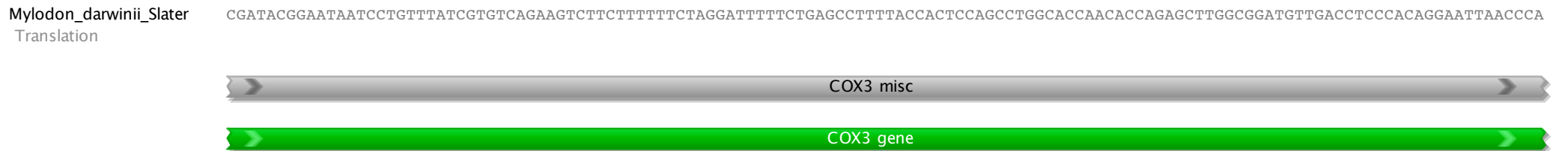
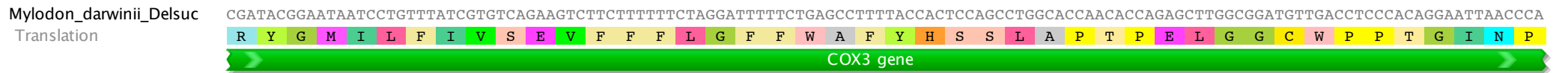
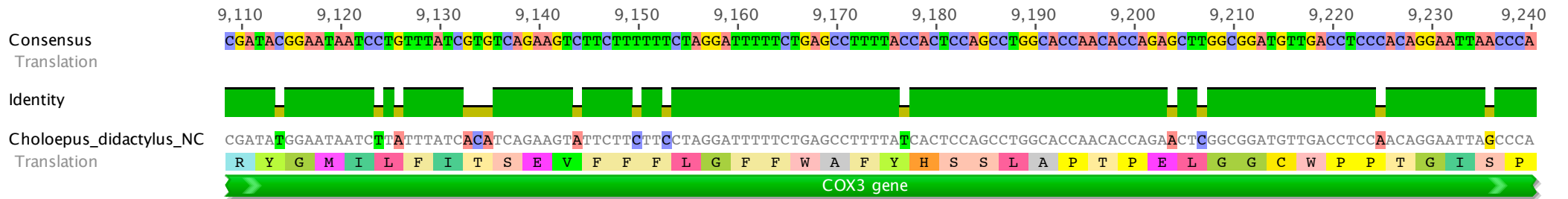
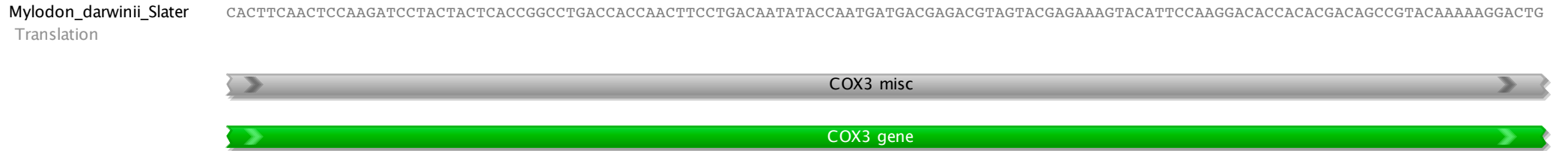
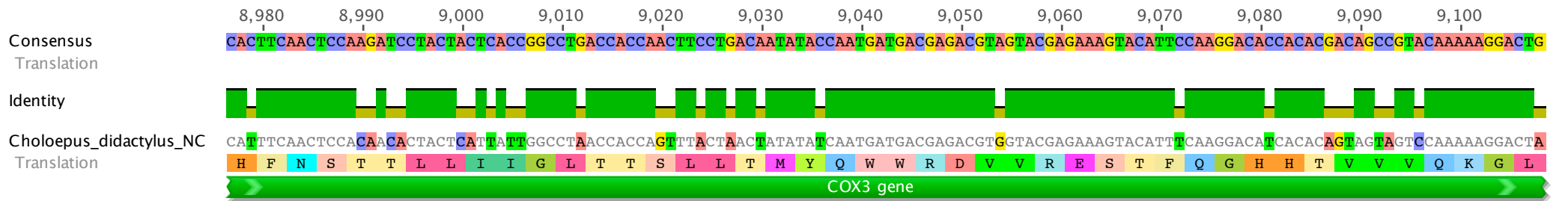
Myiodon_darwinii_Delsuc
Translation



Myiodon_darwinii_Slater
Translation







9,250 9,260 9,270 9,280 9,290 9,300 9,310 9,320 9,330 9,340 9,350 9,360 9,370

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

COX3 gene

Myiodon_darwinii_Delsuc
Translation

COX3 gene

Myiodon_darwinii_Slater
Translation

COX3 misc

COX3 gene

9,380 9,390 9,400 9,410 9,420 9,430 9,440 9,450 9,460 9,470 9,480 9,490 9,500

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

COX3 gene

Myiodon_darwinii_Delsuc
Translation

COX3 gene

Myiodon_darwinii_Slater
Translation

COX3 misc

COX3 gene

9,510 9,520 9,530 9,540 9,550 9,560 9,570 9,580 9,590 9,600 9,610 9,620 9,630

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

S S F L I I C F L R Q L K F H F T S N H H F G F E A A A W Y W H F V D V V W L L L Y V S

COX3 gene

Myiodon_darwinii_Delsuc
Translation

S S F L I V C F L R Q L K F H F T S N H H F G F E A A A W Y W H F V D V V W L F L Y V S

COX3 gene

Myiodon_darwinii_Slater
Translation

COX3 misc

COX3 gene

9,640 9,650 9,660 9,670 9,680 9,690 9,700 9,710 9,720 9,730 9,740 9,750 9,760

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

I Y W W G S M N L I L T L L I N S A L A

COX3 gene

tRNA-Gly

ND3 gene

Myiodon_darwinii_Delsuc
Translation

I Y W W G S M N L I L S L L I N S T L A

COX3 gene

tRNA-Gly

ND3 gene

Myiodon_darwinii_Slater
Translation

M N L I L S L L I N S T L A

COX3 misc

COX3 gene

tRNA-Gly

ND3 gene

9,770 9,780 9,790 9,800 9,810 9,820 9,830 9,840 9,850 9,860 9,870 9,880 9,890 9,900

Consensus
 Translation

Identity

Choloepus_didactylus_NC
 Translation

ND3 gene

Myiodon_darwinii_Delsuc
 Translation

ND3 gene

Myiodon_darwinii_Slater
 Translation

gap

ND3 gene

9,910 9,920 9,930 9,940 9,950 9,960 9,970 9,980 9,990 10,000 10,010 10,020 10,030

Consensus
 Translation

Identity

Choloepus_didactylus_NC
 Translation

ND3 gene

Myiodon_darwinii_Delsuc
 Translation

ND3 gene

Myiodon_darwinii_Slater
 Translation

gap

10,040 10,050 10,060 10,070 10,080 10,090 10,100 10,110 10,120 10,130 10,140 10,150 10,160

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

Myiodon_darwinii_Delsuc
Translation

Myiodon_darwinii_Slater
Translation

10,170 10,180 10,190 10,200 10,210 10,220 10,230 10,240 10,250 10,260 10,270 10,280 10,290

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

Myiodon_darwinii_Delsuc
Translation

Myiodon_darwinii_Slater
Translation

10,300 10,310 10,320 10,330 10,340 10,350 10,360 10,370 10,380 10,390 10,400 10,410 10,420

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

ND4L gene

Myiodon_darwinii_Delsuc
Translation

ND4L gene

Myiodon_darwinii_Slater
Translation

10,430 10,440 10,450 10,460 10,470 10,480 10,490 10,500 10,510 10,520 10,530 10,540 10,550 10,560

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

N...

ND4 gene

Myiodon_darwinii_Delsuc
Translation

N...

ND4 gene

Myiodon_darwinii_Slater
Translation

CTACAATGCTAAAAC TTATTATCCACAATTATACTAATTCCCTCACCTGACTATCAAAAAATAGCATAATCTGAATTAAC TCCACGCCACAGCCTCTTCATCAGTTTAATTAGCCTGTCAC TATTCC

10,830 10,840 10,850 10,860 10,870 10,880 10,890 10,900 10,910 10,920 10,930 10,940 10,950

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

Myiodon_darwinii_Delsuc
Translation

Myiodon_darwinii_Slater
Translation

ND4 gene

ND4 gene

ND4 misc

ND4 gene

10,960 10,970 10,980 10,990 11,000 11,010 11,020 11,030 11,040 11,050 11,060 11,070 11,080

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

Myiodon_darwinii_Delsuc
Translation

Myiodon_darwinii_Slater
Translation

ND4 gene

ND4 gene

ND4 misc

ND4 gene

11,090 11,100 11,110 11,120 11,130 11,140 11,150 11,160 11,170 11,180 11,190 11,200 11,210 11,220

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

Myiodon_darwinii_Delsuc
Translation

Myiodon_darwinii_Slater
Translation

ND4 gene

ND4 gene

ND4 misc

ND4 gene

11,230 11,240 11,250 11,260 11,270 11,280 11,290 11,300 11,310 11,320 11,330 11,340 11,350

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

Myiodon_darwinii_Delsuc
Translation

Myiodon_darwinii_Slater
Translation

ND4 gene

ND4 gene

ND4 misc

ND4 gene

11,360 11,370 11,380 11,390 11,400 11,410 11,420 11,430 11,440 11,450 11,460 11,470 11,480

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

W S F M G A S T L M I A H G L T S S L L F C L A N T N Y E R T H S R T M M L T R G L Q T



Myiodon_darwinii_Delsuc
Translation

W S F M G A S A L M I A H G L T S S L L F C L A N T N Y E R V H S R T M I L A R G L Q T



Myiodon_darwinii_Slater
Translation



11,490 11,500 11,510 11,520 11,530 11,540 11,550 11,560 11,570 11,580 11,590 11,600 11,610

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

L L P L M A L W W L L A N L S N L A L P P T I N L I G E L F V I M P T I S W S N I T I A



Myiodon_darwinii_Delsuc
Translation

L L P L M A L W W L L A S L A N L A L P P T I N L I G E L F I I L P T F S W S N I T I T



Myiodon_darwinii_Slater
Translation



11,620 11,630 11,640 11,650 11,660 11,670 11,680 11,690 11,700 11,710 11,720 11,730 11,740

Consensus
 Translation

Identity

Choloepus_didactylus_NC
 Translation



Myiodon_darwini_Delsuc
 Translation



Myiodon_darwini_Slater
 Translation



11,750 11,760 11,770 11,780 11,790 11,800 11,810 11,820 11,830 11,840 11,850 11,860 11,870 11,880

Consensus
 Translation

Identity

Choloepus_didactylus_NC
 Translation

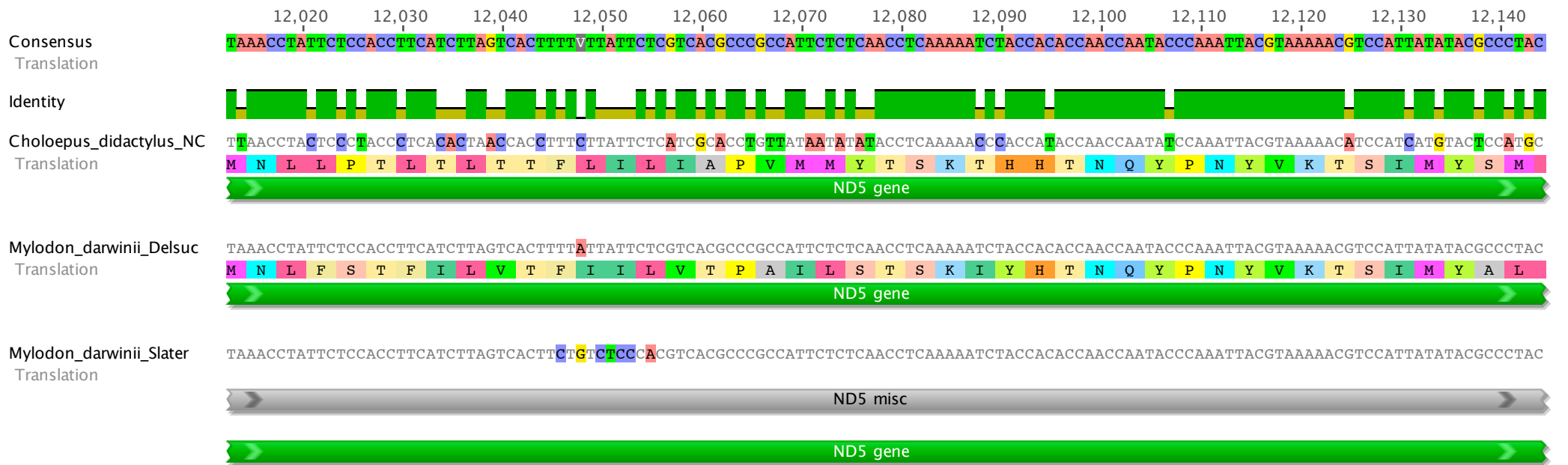
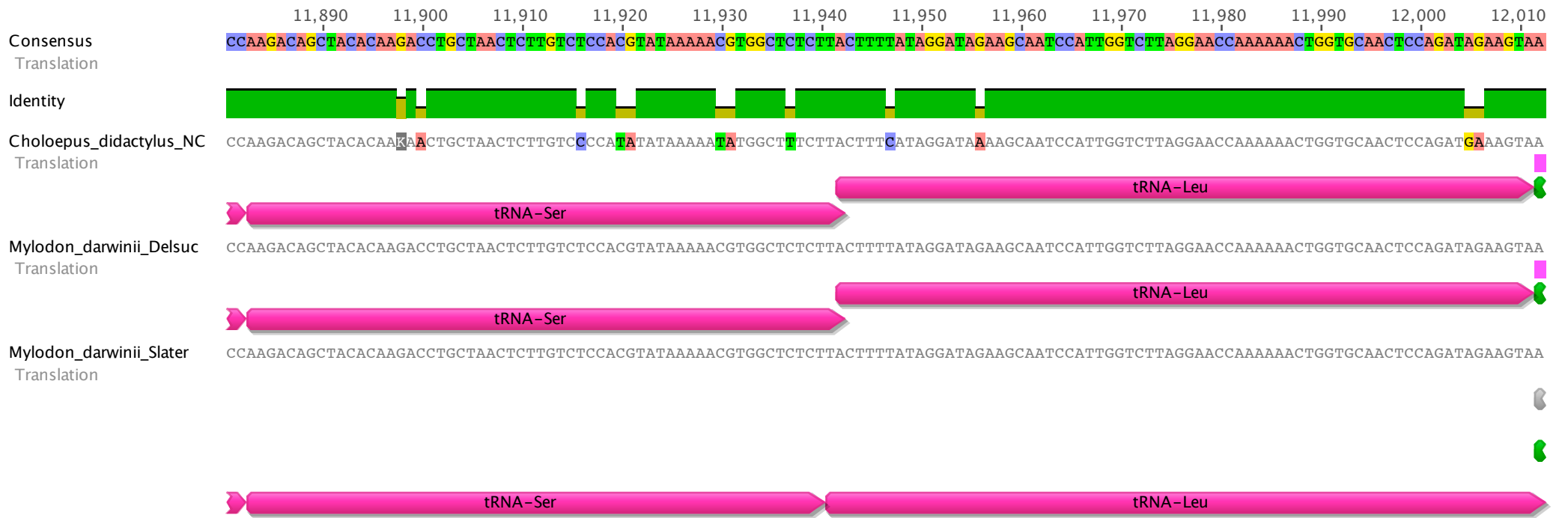


Myiodon_darwini_Delsuc
 Translation



Myiodon_darwini_Slater
 Translation





12,150 12,160 12,170 12,180 12,190 12,200 12,210 12,220 12,230 12,240 12,250 12,260 12,270

Consensus
 Translation

Identity

Choloepus_didactylus_NC
 Translation

ND5 gene

Myiodon_darwinii_Delsuc
 Translation

ND5 gene

Myiodon_darwinii_Slater
 Translation

ND5 misc

ND5 gene

12,280 12,290 12,300 12,310 12,320 12,330 12,340 12,350 12,360 12,370 12,380 12,390 12,400

Consensus
 Translation

Identity

Choloepus_didactylus_NC
 Translation

ND5 gene

Myiodon_darwinii_Delsuc
 Translation

ND5 gene

Myiodon_darwinii_Slater
 Translation

ND5 misc

ND5 gene

12,410 12,420 12,430 12,440 12,450 12,460 12,470 12,480 12,490 12,500 12,510 12,520 12,530 12,540

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

ND5 gene

Myiodon_darwinii_Delsuc
Translation

ND5 gene

Myiodon_darwinii_Slater
Translation

ND5 misc

ND5 gene

12,550 12,560 12,570 12,580 12,590 12,600 12,610 12,620 12,630 12,640 12,650 12,660 12,670

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

ND5 gene

Myiodon_darwinii_Delsuc
Translation

ND5 gene

Myiodon_darwinii_Slater
Translation

ND5 misc

ND5 gene

12,680 12,690 12,700 12,710 12,720 12,730 12,740 12,750 12,760 12,770 12,780 12,790 12,800

Consensus
 Translation

Identity

Choloepus_didactylus_NC
 Translation

TAGGAAAAATCCGCCCAATTCGGCCCTACACCCATGACTACCCCTCAGCCATGGAAGGACCTACTCCAGTATCCGCATTATTGCATTCCAGTACTATAGTAGTGGCAGGCATCTTCCTACTCGTACGATTCCACC

T G K S A Q F G L H P W L P S A M E G P T P V S A L L H S S T M V V A G I F L L V R F H

ND5 gene

Myiodon_darwinii_Delsuc
 Translation

TAGGAAAAATCCGCCCAATTCGGCCCTACACCCATGACTACCCCTCAGCCATGGAAGGACCTCACCCTCGTATCCGCATCTATTCGATTCCAGTACTATAGTAGTGGCAGGCATCTTCCTACTCGTACGATTCCACC

M G K S A Q F G L H P W L P S A M E G P T P V S A L L H S S T M V V A G I F L L V R F H

ND5 gene

Myiodon_darwinii_Slater
 Translation

TAGGAAAAATCCGCCCAATTCGGCCCTACACCCATGACTACCCCTCAGCCCTGGTAAAGACCTACTCCAGTATCCGCATTATTGCATTCCAGTACTATAGTAGTGGCAGGCATCTTCCTACTCGTACGATTCCACC

ND5 misc

ND5 gene

12,810 12,820 12,830 12,840 12,850 12,860 12,870 12,880 12,890 12,900 12,910 12,920 12,930

Consensus
 Translation

Identity

Choloepus_didactylus_NC
 Translation

CCCTCCTACAAAAAATAAACTAATCCTCACACTAGCCATATGTCTAGGGGCCATCACCACGCTATTCACCAGCAATCTGTGCCTTAAACACAGAATGATATTAATAAAAAATCGTAGCCTTTTCTACATCCAGCC

P M L Q N N S P I L T L T M C L G A I T T L F T A I C A L T Q N D I K K I I A F S T S S

ND5 gene

Myiodon_darwinii_Delsuc
 Translation

CCCTCCTACAAAAAATAAACTAATCCTCACACTAGCCATATGTCTAGGGGCCATCACCACGCTATTCACCAGCAATCTGTGCCTTAAACACAGAATGATATTAATAAAAAATCGTAGCCTTTTCTACATCCAGCC

P L L Q N N K L I L T L A M C L G A I T T L F T A I C A L T Q N D I K K I V A F S T S S

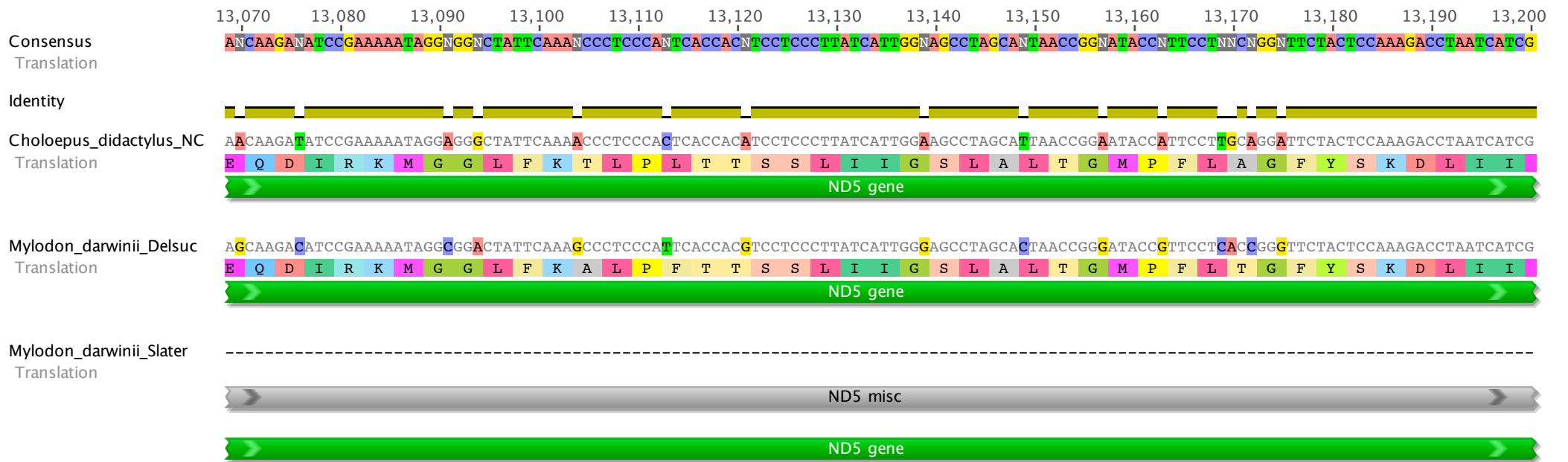
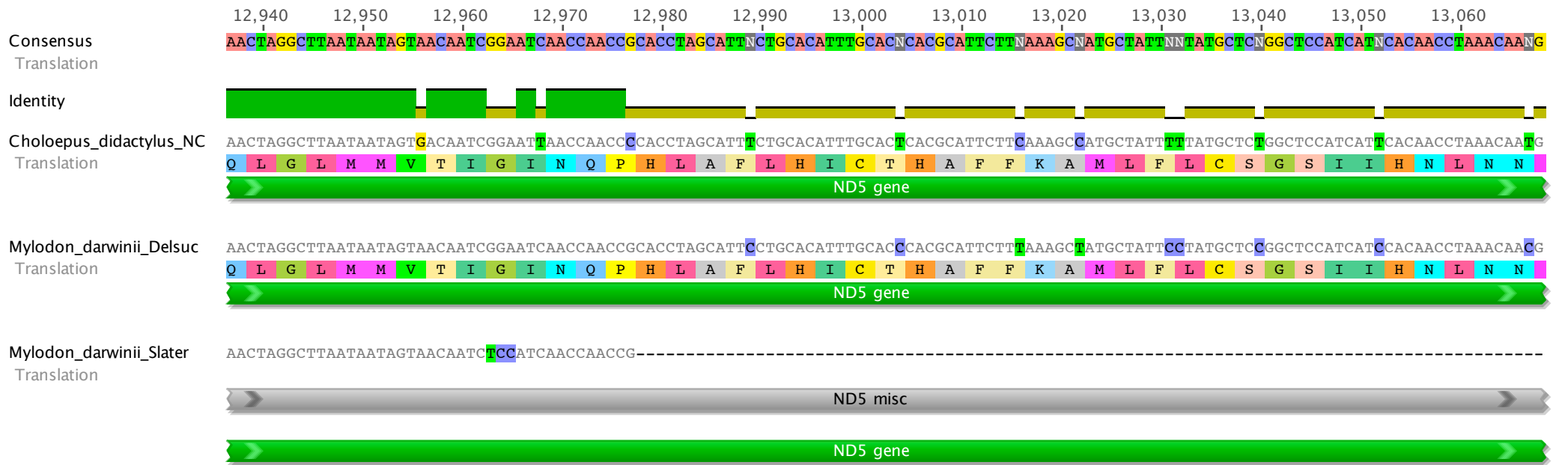
ND5 gene

Myiodon_darwinii_Slater
 Translation

CCCTCCTACAAAAAATAAACTAATCCTCACACTAGCCATATGTCTAGGGGCCATCACCACGCTATTCACCAGCAATCTGTGCCTTAAACACAGAATGATATTAATAAAAAATCGTAGCCTTTTCTACATCCAGCC

ND5 misc

ND5 gene



13,210 13,220 13,230 13,240 13,250 13,260 13,270 13,280 13,290 13,300 13,310 13,320 13,330
Consensus
Translation

Identity
Choloepus_didactylus_NC
Translation

Myiodon_darwinii_Delsuc
Translation

Myiodon_darwinii_Slater
Translation

13,340 13,350 13,360 13,370 13,380 13,390 13,400 13,410 13,420 13,430 13,440 13,450 13,460
Consensus
Translation

Identity
Choloepus_didactylus_NC
Translation

Myiodon_darwinii_Delsuc
Translation

Myiodon_darwinii_Slater
Translation

13,470 13,480 13,490 13,500 13,510 13,520 13,530 13,540 13,550 13,560 13,570 13,580 13,590

Consensus
 Translation

Identity

Choloepus_didactylus_NC
 Translation

ND5 gene

Myiodon_darwinii_Delsuc
 Translation

ND5 gene

Myiodon_darwinii_Slater
 Translation

ND5 misc

ND5 gene

13,600 13,610 13,620 13,630 13,640 13,650 13,660 13,670 13,680 13,690 13,700 13,710 13,720

Consensus
 Translation

Identity

Choloepus_didactylus_NC
 Translation

ND5 gene

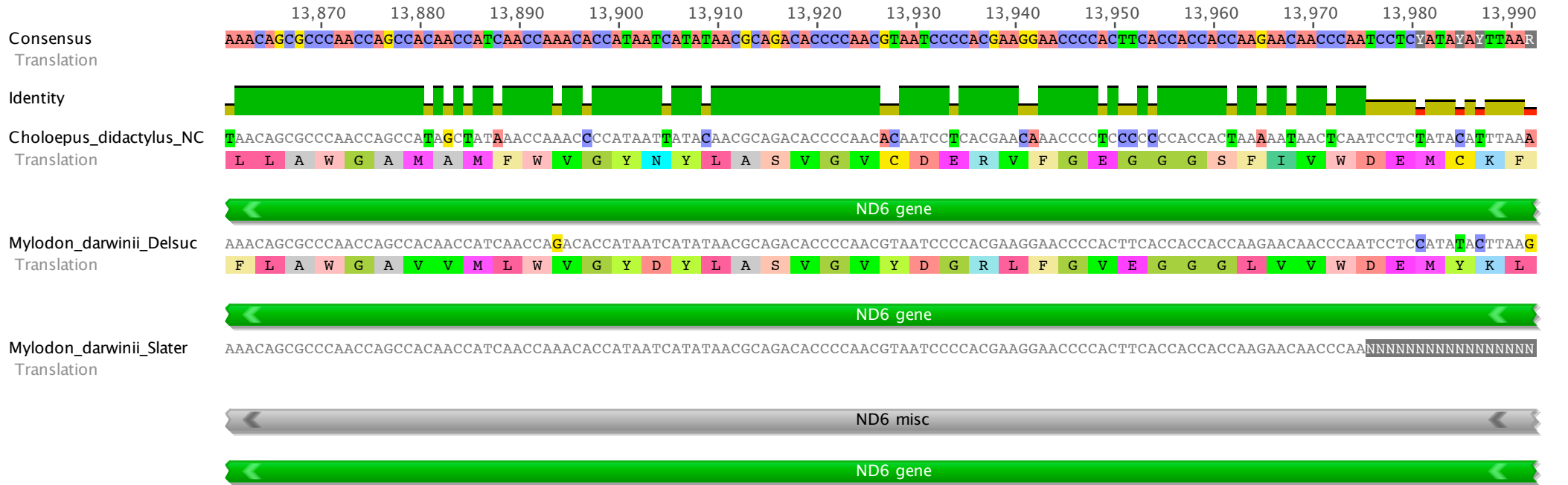
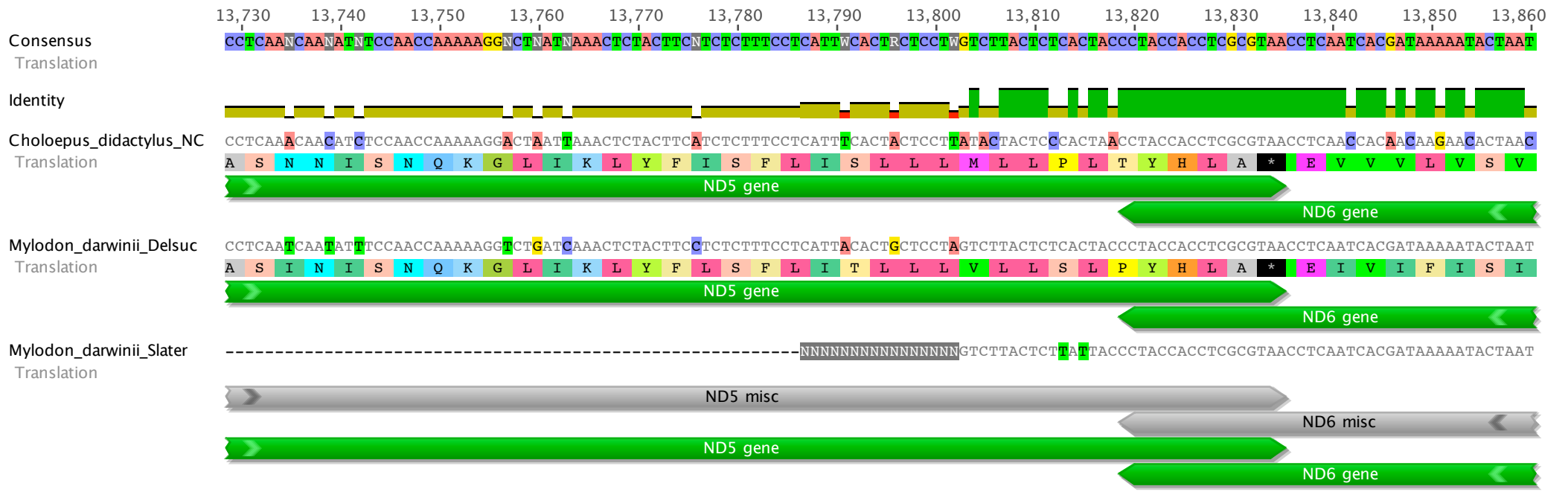
Myiodon_darwinii_Delsuc
 Translation

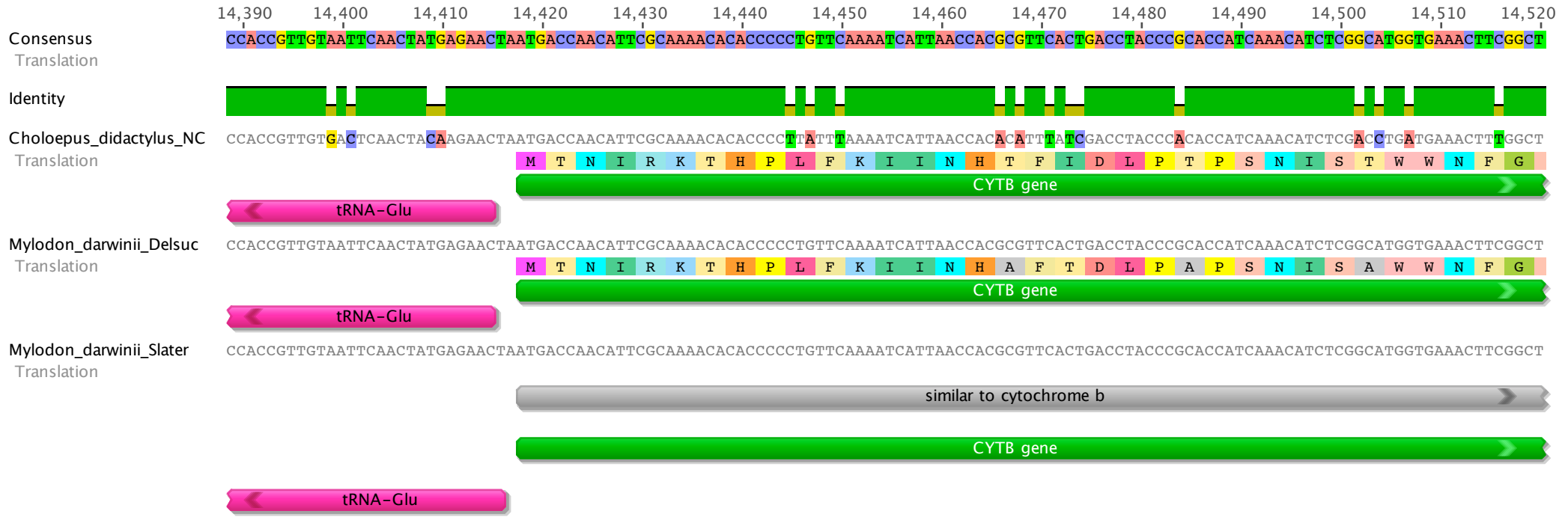
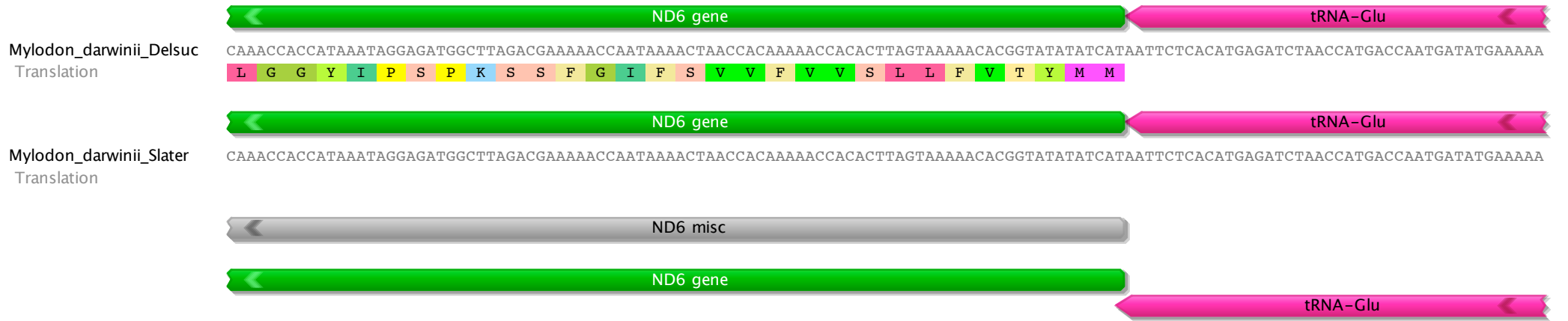
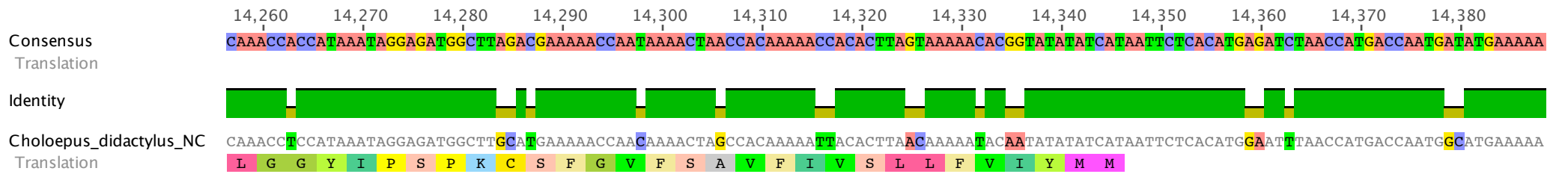
ND5 gene

Myiodon_darwinii_Slater
 Translation

ND5 misc

ND5 gene





14,530 14,540 14,550 14,560 14,570 14,580 14,590 14,600 14,610 14,620 14,630 14,640 14,650

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

S L L G M C L I M Q I L T G L F L A M H Y -- T P D T T T A F S S V T H I C R D V N Y G W I

CYTB gene

Myiodon_darwinii_Delsuc
Translation

S L L G I C L I M Q I L T G L F L A M H Y -- T S D T I T A F S S V T H I C R D V N Y G W I

CYTB gene

Myiodon_darwinii_Slater
Translation

CCCGCATCCGACACAATCACCGCCTTCTCGTCCGTACCCACATCTGCCGAGACGTAAACTACGGCTGAAT

similar to cytochrome b

CYTB gene

14,660 14,670 14,680 14,690 14,700 14,710 14,720 14,730 14,740 14,750 14,760 14,770 14,780

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

I R Y L H T N G A S M F F I C L Y L H I G R S M Y Y G S Y T Y L K T W N I G I M L L F T

CYTB gene

Myiodon_darwinii_Delsuc
Translation

I R Y L H A N G A S M F F V C L Y M H I G R G L Y Y G S Y S Y L E T W N I G V L L L F T

CYTB gene

Myiodon_darwinii_Slater
Translation

ACTCCTACTCAATACTAGAAACCTGAAACATTGGAGTCATCTACTATTAC

similar to cytochrome b

CYTB gene

14,790 14,800 14,810 14,820 14,830 14,840 14,850 14,860 14,870 14,880 14,890 14,900 14,910

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

CGTAATAGCCACAGCATTTCATAGGCTATGTACTTCCATGAGGACAAAATATCATTCTGAGGAGCAACAGTAAATACAAACCTACTCTCAGCCATCCCATACATCGGAACAGATCTAGTAGAATGAATCTGAGG

V M A T A F M G Y V L P W G Q M S F W G A T V I T N L L S A I P Y I G T N L V E W I W G

CYTB gene

Myiodon_darwinii_Delsuc
Translation

TGTTATAGCCACAGCATTATATAGGCTATGTACTTCCATGAGGACAAAATATCATTCTGAGGAGCAACAGTAATTACAAACCTACTCTCAGCCATCCCATACATCGGAACAGATCTAGTAGAATGAATCTGAGG

V M A T A F M G Y V L P W G Q M S F W G A T V I T N L L S A I P Y I G T D L V E W I W G

CYTB gene

Myiodon_darwinii_Slater
Translation

TGTTATAGCCACAGCATTATATAGGCTATGTACTTCCATGAGGACAAAATATCATTCTGAGGAGCAACAGTAATTACAAACCTACTCTCAGCCATCCCATACATCGGAACAGATCTAGTAGAATGAATCTGAGG

similar to cytochrome b

CYTB gene

14,920 14,930 14,940 14,950 14,960 14,970 14,980 14,990 15,000 15,010 15,020 15,030 15,040

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

GGGATTTCCGTAGACAAAAGCCACATTAACCCGATTCTTTCCTTCCATTTCACTCCATTCATATCCCTGGCCCTCGTACTAACACACCTGCTTTTCCCTACACGAAACCGGATCAAACAACCCACTAGG

G F S V D K A T L T R F F A F H F I L P F I I L A L V L T H L L F L H E T G S N N P L G

CYTB gene

Myiodon_darwinii_Delsuc
Translation

CGGATTTCCGTAGACAAAAGCCACATTAACCCGCTTCTTTCCTTCCATTTTCATCCTTCCATTCATATCCTGGCCCTCGTACTAACACACCTGCTTTTCCCTACACGAAACCGGATCAAACAACCCACTAGG

G F S V D K A T L T R F F A F H F I L P F I I L A L V L T H L L F L H E T G S N N P L G

CYTB gene

Myiodon_darwinii_Slater
Translation

CGGATTTCCGTAGACAAAAGCCACATTAACCCGCTTCTTTCCTTCCATTTTCATCCTTCCATTCATATCCTGGCCCTCGTACTAACACACCTGCTTTTCCCTACACGAAACCGGATCAAACAACCCACTAGG

similar to cytochrome b

CYTB gene

15,050 15,060 15,070 15,080 15,090 15,100 15,110 15,120 15,130 15,140 15,150 15,160 15,170 15,180

Consensus
 Translation

Identity

Choloepus_didactylus_NC
 Translation

I S P D S D K I P F H P Y Y T M K D I L G L L P M I F I M M A L A L F Y P D L L G D P D

CYTB gene

Myiodon_darwinii_Delsuc
 Translation

I S S D S D K I P F H P Y Y T V K D I L G L F L M I L I M L T L V L F Y P D L L G D P D

CYTB gene

Myiodon_darwinii_Slater
 Translation

similar to cytochrome b

CYTB gene

15,190 15,200 15,210 15,220 15,230 15,240 15,250 15,260 15,270 15,280 15,290 15,300 15,310

Consensus
 Translation

Identity

Choloepus_didactylus_NC
 Translation

N Y T P A N P L N T P P H I K P E W Y F L F A Y A I L R S I P N K L G G V M A L V L S I

CYTB gene

Myiodon_darwinii_Delsuc
 Translation

N Y T P A N P L N T P P H I K P E W Y F L F A Y A I L R S I P N K L G G V L A L I C S I

CYTB gene

Myiodon_darwinii_Slater
 Translation

similar to cytochrome b

CYTB gene

15,320 15,330 15,340 15,350 15,360 15,370 15,380 15,390 15,400 15,410 15,420 15,430 15,440

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

L I L A I L P M L H T A K Q R S L M F R P I S Q C L F W M L M T N L L V L T W I G G Q P

CYTB gene

Myiodon_darwinii_Delsuc
Translation

L V L V I I P M L H T A K Q R S L M F R P I S Q C L F W M L V A N L L I L T W I G G Q P

CYTB gene

Myiodon_darwinii_Slater
Translation

similar to cytochrome b

CYTB gene

15,450 15,460 15,470 15,480 15,490 15,500 15,510 15,520 15,530 15,540 15,550 15,560 15,570

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

V E H P F T I I G Q L A S I S Y F S I I L I L M P M A G L I E N N L M K W

CYTB gene

Myiodon_darwinii_Delsuc
Translation

V E H P F T I I G Q L A S I S Y F S I I L V L M P V A G L I E N N L M K W *

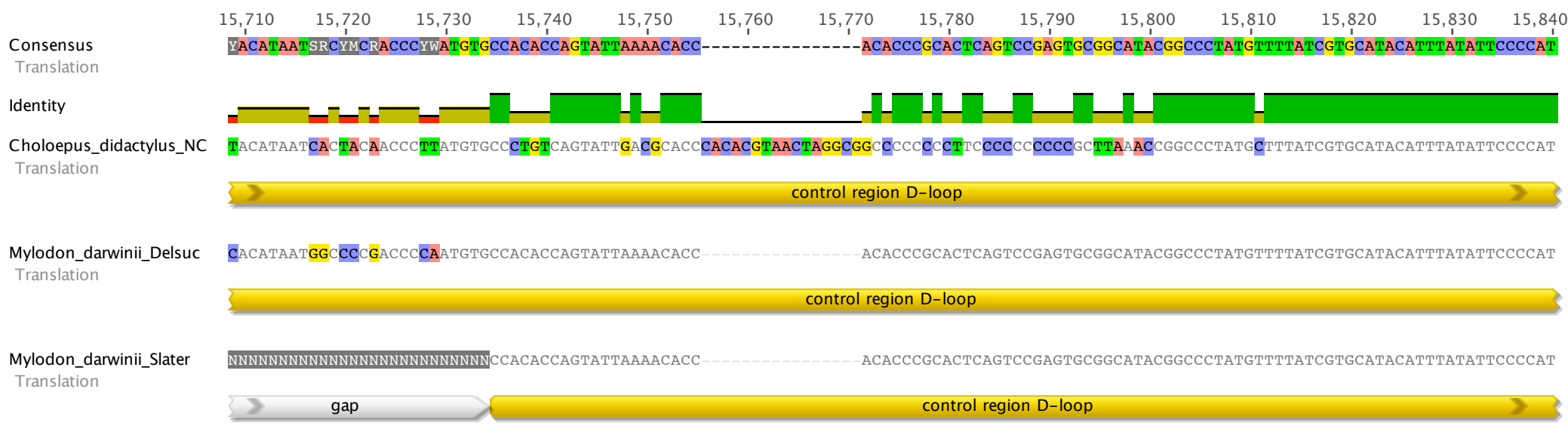
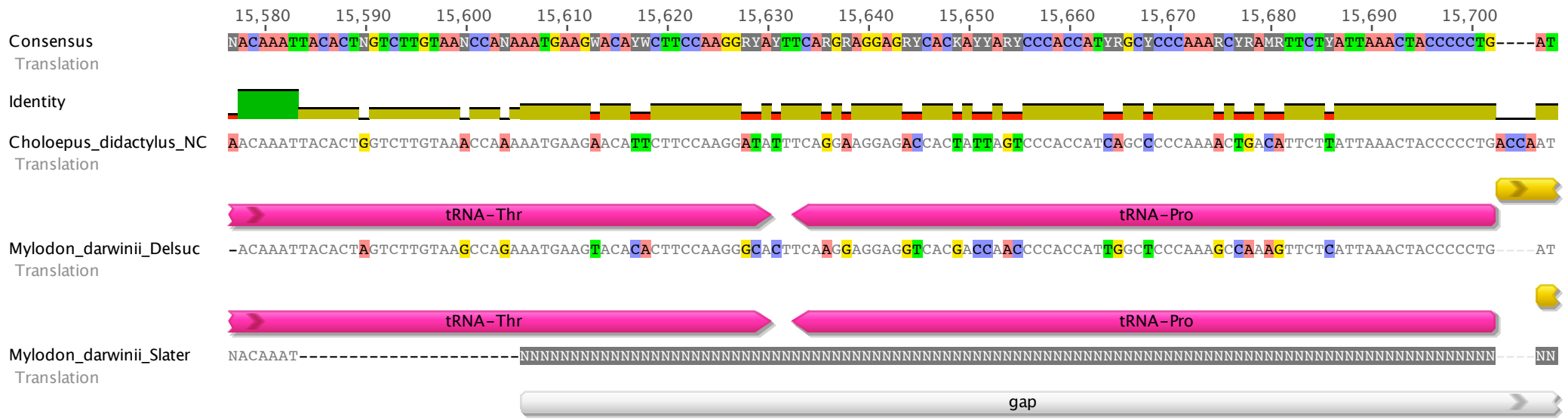
CYTB gene

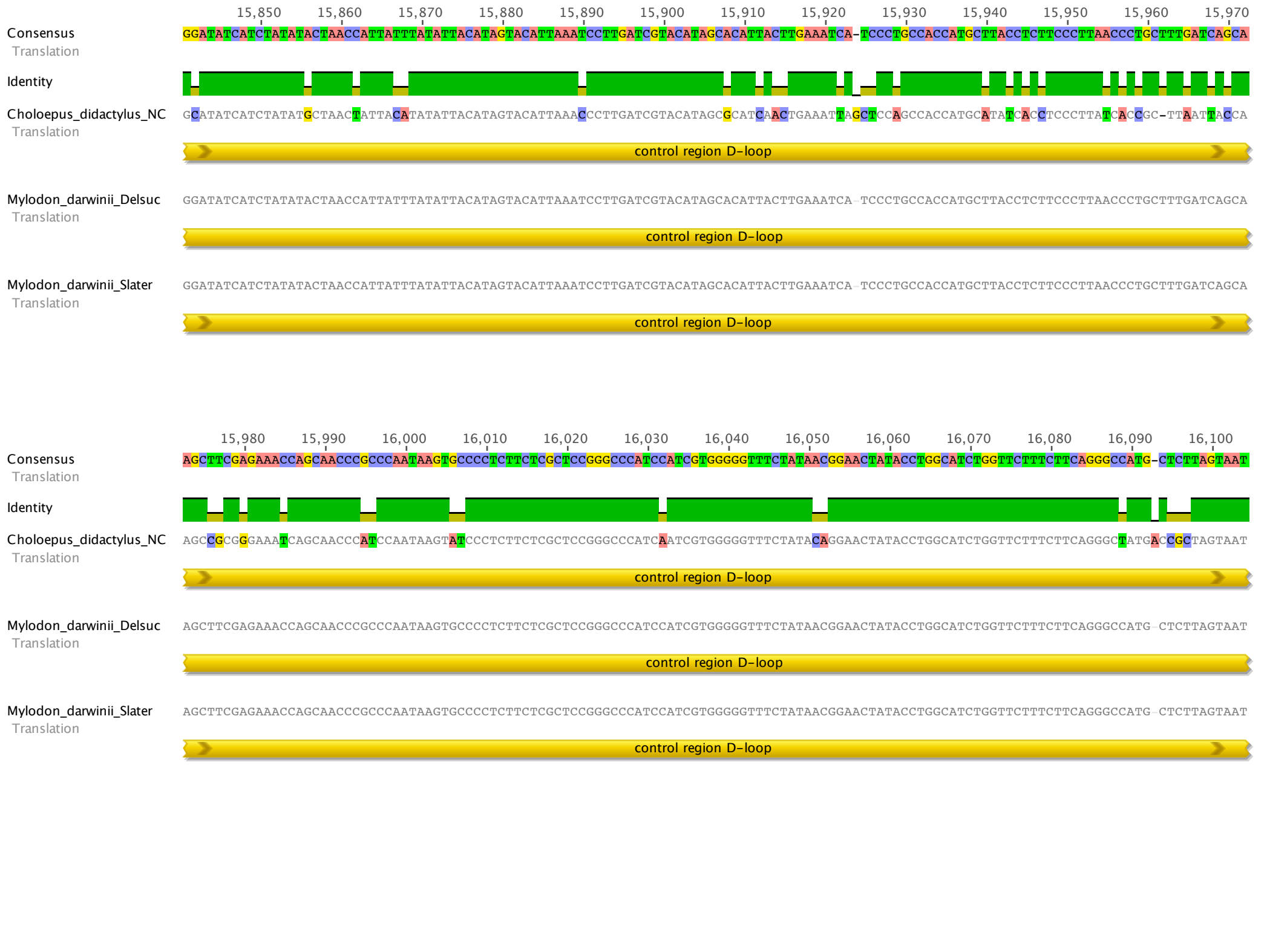
Myiodon_darwinii_Slater
Translation

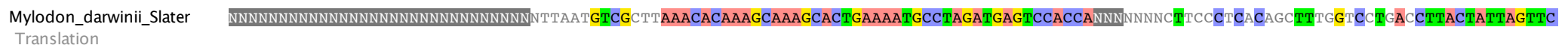
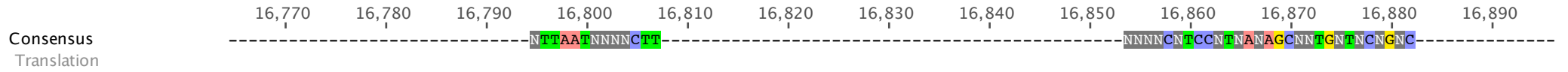
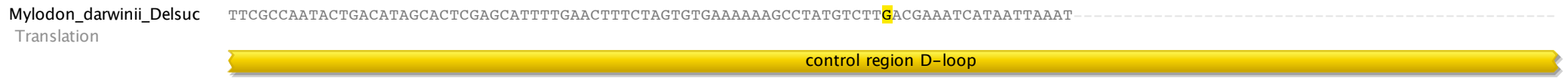
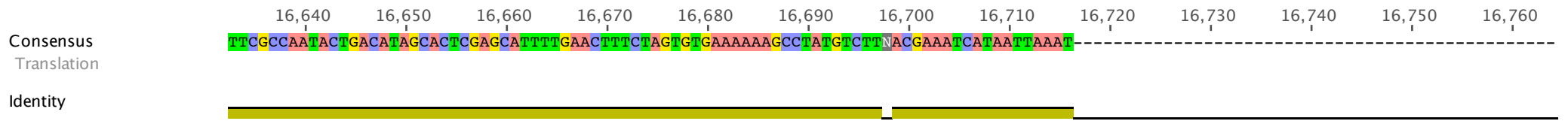
similar to cytochrome b

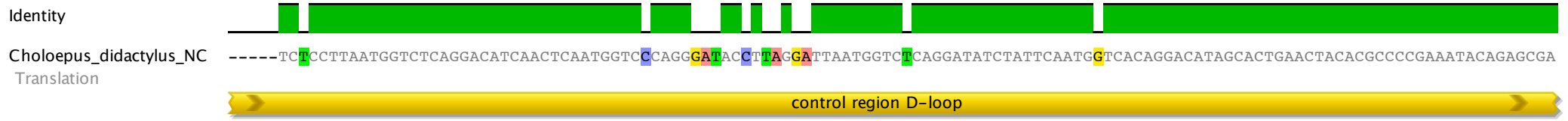
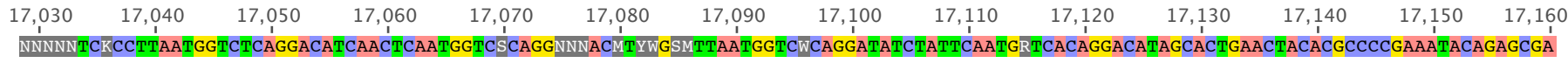
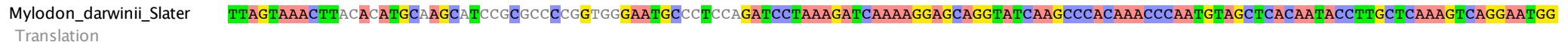
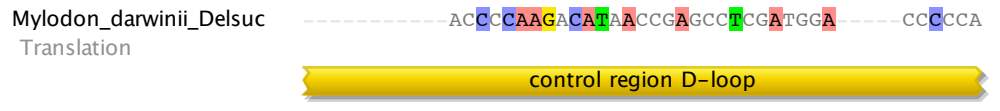
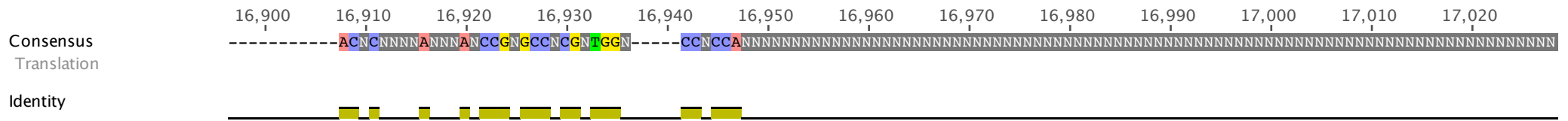
CYTB gene

tRN...

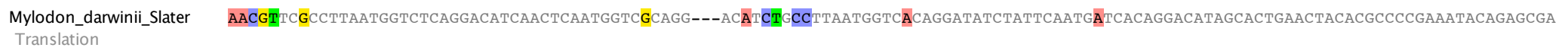








Myiodon_darwinii_Delsuc
Translation



17,170 17,180 17,190 17,200 17,210 17,220 17,230 17,240 17,250 17,260 17,270 17,280 17,290

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

TAAGCATAAGTGCCCAACAAATGACAATACTCCTATTTTTTTCTCATTATAATAGCAATATTTCCATTTGCGCCAATACTGACATAGCACTCAAGCATTTTGAATTTCTACTGTGAAAAAGCCATATGTCT

control region D-loop

Myiodon_darwinii_Delsuc
Translation

Myiodon_darwinii_Slater
Translation

TAAGCATAAGTGCCCAACAAATGACAATACTCCTATTTTTTTCTCATTATAATAGCGATATTTCCATTTGCGCCAATACTGACATAGTACTTAAGCATTTTGAATTTCTACTGTGAAAAAGCCATATGTCT

17,300 17,310 17,320 17,330 17,340 17,350 17,360 17,370 17,380 17,390 17,400 17,410 17,420

Consensus
Translation

Identity

Choloepus_didactylus_NC
Translation

TGATGAAACTATAATTAATTTCAATTTCTTTTTTTTT--CATTTTAAACACGTGCTATCAACACGCCAATGATACAACGCGGCCCCACACACAAGTTAATGTCGCTTAAACACAAAGCAAGGCACGAAA

control region D-loop

Myiodon_darwinii_Delsuc
Translation

Myiodon_darwinii_Slater
Translation

TGATGAAACTATAATTAATTTCAATTTCTTTTTTTTTTNCATTTTAAACACGTGCTATCAACACGCCAATGATACAAC-----CAGCCTCGATGG

17,430 17,444
Consensus
Translation

A Y S C Y Y A G A T G A G T A C A T C A

Identity



Choloepus_didactylus_NC

A T G C T T A G A T G A G T A C A T C A

Translation



Myiodon_darwinii_Delsuc

Translation

Myiodon_darwinii_Slater

A C C C C A

Translation

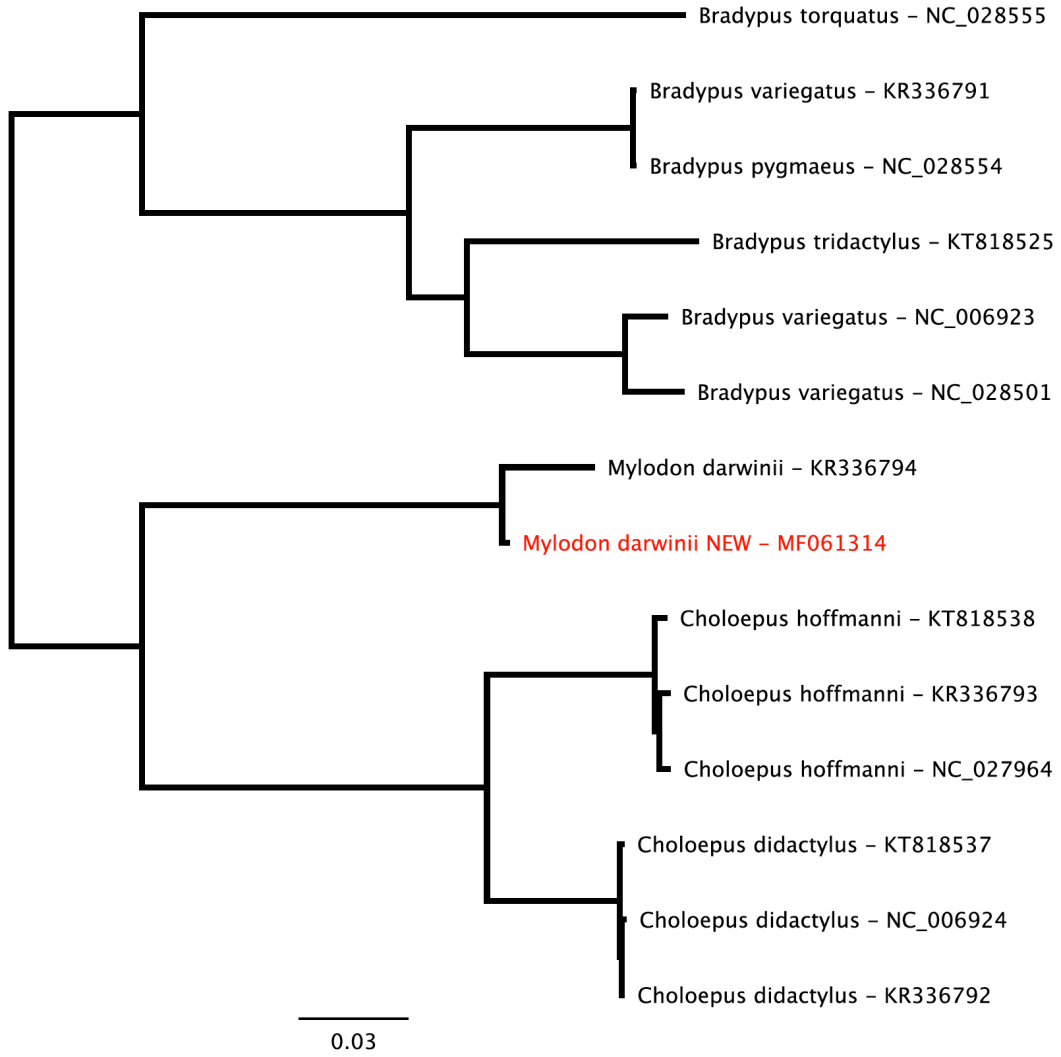


Figure S12: Maximum likelihood tree of available sloth mitogenomes obtained with RAxML under a single GTR+G model. Our new Mylodon mitogenome sequence is indicated in red to illustrate its much shorter branch length relative to the one previously obtained by Slater et al. (2017). Branch lengths represent mean number of substitutions per site.