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Primer	Sequence	Strand	Gene / Region	3'-End (Ref. Seq.)	Author (ref.)
MVZ 01	AAACTGGGATTAGATACCCAC	L	12s RNA	506 (Mus);1088 (Hu)	T. White
MVZ 02	GAGGGTGACGGGCGGTGTG	H	12s RNA	902 (Mus);1479 (Hu)	T. White
MVZ 03	GCTTCCATCCAACATCTCAGCATGATG	L	cyt b	14230 (Mus)	T. White
MVZ 03'	CCATCCAACATCTC[A/T]GC[A/T]TGATG	L	cyt b	14230 (Mus);14838 (Hu); 14987 (Chi)	N.K. Johnson/C. Cicero
MVZ 04	GCAGCCCCCTAGAATGATATTTGTCTC	H	cyt b	14542 (Mus)	T. White
MVZ 04'	GTAGCACCTCAGAA[C/G/T]GATATTTG	H	cyt b	(Hu);15304 (Chi)	N.K. Johnson/C. Cicero
MVZ 05	CGAAGCTTGATATGAAAAACCATCGTTG	L	cytb/tRNA-Glu	14115 (Mus)	S. Pääbo, L14724
MVZ 05-Bt	Bt-CGAAGCTTGATATGAAAAACCATCGTTG	L	cytb/tRNA-Glu	14115 (Mus)	C. Orrego
MVZ 05-GC	CGCCCGCCGCGCCCCGCGCCCGGCCCGCCCGCCCCGCCCC- CGAAGCTTGATATGAAAAACCATCGTTG	L	cytb/tRNA-Glu	14115 (Mus)	C. Orrego
MVZ 05-M13RSP	GTGAGCGGATAACAATTTTCACACAGGA- CGAAGCTTGATATGAAAAACCATCGTTG	L	cytb/tRNA-Glu	14115 (Mus)	C. Orrego
MVZ 06	GCTGTGTCTGATGTGTAGTGTAT	H	cyt b	14295 (Mus)	M. F. Smith
MVZ 07	AACCCCATCTAACATTTC[A/T]TC[C/T]TGATG	L	cyt b	14230 (Mus)	M. F. Smith
MVZ 08	ATGTGAAG[G/A]TAGATGCA[G/A]AT[G/A]AA	H	cyt b	14409 (Mus)	C. Orrego
MVZ 09	GTACAATTCGG[A/C]TGACT[A/C]AT[C/T]CG	L	cyt b	14377 (Mus)	C. Orrego
MVZ 10	TATGAGCCGTAGTA[A/G]A[T/G][T/G]CCTC	H	cyt b	14437 (Mus)	C. Orrego
MVZ 11	CCTCCGTAGCCCACA[T/C][A/T]TG[C/T]CG	L	cyt b	14350 (Mus)	C. Orrego
MVZ 11-Bt	Bt-CCTCCGTAGCCCACA[T/C][A/T]TG[C/T]CG	L	cyt b	14350 (Mus)	C. Orrego
MVZ 12	TTATTACTCCAAT[A/G]TTTCA[T/G/A]GT[C/T]TC	H	cyt b	14469 (Mus)	D. Good
MVZ 13	TTTTTAGCTATACATTATACA[G/T/C]C[A/C/T/G]GA	L	cyt b	14311 (Mus)	D. Good
MVZ 14	GGTCTTCATCT[C/T][C/T/A]GG[T/C]TTACAAGAC	H	cytb/tRNA-Thr	15309 (Mus);15915 (Hu); 16065 (Chi)	C. Orrego
MVZ 14-Bt	Bt-GGTCTTCATCT[C/T][C/T/A]GG[T/C]TTACAAGAC	H	cyt b	15309(Mus);15915(Hu);	
MVZ 14-M13SP	CGCCAGGGTTTTCCAGTCACGAC- GGTCTTCATCT[C/T][C/T/A]GG[T/C]TTACAAGAC	H	cyt b	15309(Mus);15915(Hu);	C. Orrego
MVZ 15	GAACTAATGGCCACAC[A/T][A/T]TACG[G/A/T/C]AA	L	cyt b	14155 (Mus)	D. Good / C. Orrego
MVZ 16	AAATAGGAA[A/G]TATCA[T/C]TCTGGTTT[A/G]AT	H	cyt b	14940 (Mus)	C. Orrego
MVZ 16-Bt	Bt-AAATAGGAA[A/G]TATCA[T/C]TCTGGTTT[A/G]AT	H	cyt b	14940 (Mus)	C. Orrego
MVZ 17	ACCTCCTAGGAGA[C/T]CCAGA[C/A/T]AA[C/T]T	L	cyt b	14904 (Mus)	C. Orrego
MVZ 18	GTCTTTGTATGAGAAGTATG	H	cyt b	14803(Mus)	C. Orrego/C. Schneider
MVZ 19	TGCCTATATATTCA[T/C]AT[T/C]G[G/C]A[C/G]G	L	cyt b	14437 (Mus)	C. Orrego/L. Apfelbaum
MVZ 20	TATGCGAATAGGAA[G/A]TA[T/C]CA[T/C]TC	H	cyt b	14949(Mus);15557 (Hu); 15706 (Chi)	N.K. Johnson/C. Cicero
MVZ 21	GCAATCCCATATATTGGCACAA[A/C][C/T/A][C/T]T	L	cyt b	14617 (Mus)	C. Orrego/L. Apfelbaum
MVZ 22	CGTTAGCATGTAT[A/G]TATCG[G/A]AT	H	cyt b	14373 (Mus)	C. Orrego/L. Apfelbaum

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MVZ 23	TACTCTTCCTCCACGAAAC[A/C]GG[A/C/T/G]TC	L	cyt b	14752 (Mus)	C. Orrego/L. Apfelbaum
MVZ 24	TGTAGTT[A/G]TC[A/T]GGGTC[A/T/C]CCTA	H	cyt b	14887 (Mus)	M. DaSilva/C.Orrego
MVZ 25	TAAAGAAACTTGAAA[T/C]AT[T/C]GG[A/T/C]GT	L	cyt b	14488 (Mus);16601 (Xen)	C. Orrego/C. Moritz
MVZ 26	AGATCTTTGATTGTGTAGTAGGGGT	H	cyt b	14800 (Mus)	M. F. Smith
MVZ 27	TACCTAAACAAAGAAAC[G/C/T]TG[G/A]AA	L	cyt b	14479(Mus);15087(Hu); 15236(Chi)	N.K. Johnson/C. Cicero
MVZ 28	CGAGGC[G/C]CC[T/C]GCAAT[A/G]ATAA	H	cyt b	14698 (Mus)	C. Orrego/A. A. Graybeal
MVZ 29	GACTGTGACAAAATCCC[G/A/T/C]TTCCA	L	cyt b	14800(Mus);15408(Hu); 15557(Chi)	N.K. Johnson/C. Cicero
MVZ 30	GGAGCCAAA[A/G]TTTCATCAGGC	H	cyt b	14223 (Mus);16339 (Xen)	C.Schneider/C.Orrego
MVZ 31	TCTCAGGATTATGAAA[T/C]TT[T/C]GG	L	cyt b	14239 (Mus)	C. Orrego/L. Apfelbaum
MVZ 32	TCCTA[A/G]TAA[T/A/G]GATCC[A/G]AA[A/G]TTTCA	H	cyt b	14229 (Mus)	G. Applebaum/C. Orrego
MVZ 33	GGTCTCTACTTGGAAT[A/T]TG[T/C]TT	L	cyt b	14260 (Mus)	C. Orrego/L. Apfelbaum
MVZ 34	CCTA[C/T]AAA[G/T]GCTGTGGCTATGA	H	cyt b	?15506? (Mus)	C. Orrego/P. Barber
MVZ 35	TCTCAGAAACCTG[A/G]AA[C/T]AT[T/C]GG	L	cyt b	14485 (Mus)	M.F. Smith
MVZ 36	TCTTCTACTGGTTGTCTCCGATTCA	H	cyt b	15114 (Mus)	P. Barber/C. Orrego
MVZ 37	TACACCTT[T/C]ATAGA[A/G]AC[A/C/T]TG[A/G]A	L	cyt b	14478 (Mus)	M. DaSilva/C.Orrego
MVZ 38	TTCTTAGGGCTTTGAAGGCTC	H	ND2/tRNA-Trp	4975 (Mus)	M. F. Smith
MVZ 39	CA[C/T]TTACTATTT[T/C]T[A/T/C]CA[C/T]GA[G/A]AC	L	cyt b	14746 (Mus)	M. DaSilva/C.Orrego
MVZ 40	GCTAG[G/A]AA[G/A]AG[T/A]CC[G/A]GTTA	H	cyt b	14275 (Mus)	M. F. Smith
MVZ 41	CACCCTTA[C/T]TA[C/T]AC[A/C]A[C/T][C/T]AAAGA	L	cyt b	14821 (Mus)	M. DaSilva/C.Orrego
MVZ 42	TCGT[A/C]TAACCGCGG[T/C]GGCTG	H	12 s	310 (Mus); 2307 (Xen)	M. Garcia-Paris/Cristian
MVZ 43	GAGTCTGCCT[A/T]AT[T/C]GC[C/T]CA[A/G]AT	L	cyt b	14272 (Mus)	C. Orrego/A. Graybeal
MVZ 44	TT[A/C][C/T]AGAACAGGCTCCTCTAG	H	12 s	613(Mus); 2605(Xen)	M. Garcia-Paris/C. Orrego
MVZ 45	AC[A/C]AC[C/T/A]ATAGC[A/C]ACAGCATTCGTAGG	L	cyt b	14527 (Mus)	M.F. Smith
MVZ 46	TTTAGAAAATGTAGCCATTCTT	H	12 s	776 (Mus); 2768 (Xen)	M. Garcia-Paris/Cristian
MVZ 47	CCAGACAATT[A/T][C/T]AC[A/C/T]CCAGCAAA[C/T]C	L	cyt b	14919 (Mus)	M.F. Smith
MVZ 48	GTGACGGGCGGTGTGT[G/A]CG	H	12 s	898 (Mus); 2894 (Xen)	M. Garcia-Paris/C. Orrego
MVZ 49	ATAA[A/G]AACAATGACAAT[C/T]ATACGAA	L	cyt b	14154 (Mus);16267 (Xen)	C.Schneider/C.Orrego
MVZ 50	T[T/C]TCGGTGTAAG[T/C]GA[G/A]A[G/T]GCTT	H	12S/tRNA-Val	1045 (Mus); 3042 (Xen)	M. Garcia-Paris/C. Orrego
MVZ 51	CTAGATCCCCAAGT[C/T]TCAG[A/G]ATA	L	cyt b	13833 (Mus);15949 (Xen)	C.Schneider/C.Orrego
MVZ 51 CM	CCAATGACCAACCTACGTAA	L	cyt b	14155 (Mus)	C. Orrego
MVZ 52	ARA TRA ARA ATA TTG ATG CT	H	cyt b	14396 (Mus)	M. DaSilva/C. Orrego
MVZ 53	GGTTCTTATATATTTAAAGA[G/A]AC[T/C]TG	L	cyt b	14476 (Mus);16598 (Xen)	C. Schneider/C. Orrego
MVZ 54	ATGTARYKGAWRTKWTWTAATGGT	H	D-loop	2198 (Xen)	A. Graybeal/ C. Orrego
MVZ 55	TGAGGACAAATATCATTCTGAGGGGCTGCAG	L	cyt b	14571 (Mus)	P. Barber/C. Orrego

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MVZ 56	CTA AGC ATA GTG GGG TAT CTA ATC C	H	XL PCR	491 (Mus)	C. Schneider
MVZ 57	GCTATCGGGCCCATACCCCGA	L	ND2/tRNA-Met	3884 (Mus)	M. F. Smith
MVZ 58	CTA AGC ATA GTG GGG TAT CTA ATC CCA G	H			C. Schneider
MVZ 59	ATAGCACTGAAAA[T/C]GCT[T/A/G]AGATG	L	12S/tRNA-Phe	47 (Mus); 2180 (Xen)	M. Garcia-Paris/C. Orrego
MVZ 60	CTA AGC ATA GTG GGG TAT CTA ATC CCA GT	H			C. Schneider
MVZ 61	CAGCC[A/G]CCGCGGT[C/T]A[T/G]ACGA	L	12 s	329 (Mus); 2326 (Xen)	M. Garcia-Paris/C. Orrego
MVZ 62	ATG AAG GGA TGT TCT ACT GGT TG	H			C. Mojica
MVZ 63	A[A/G][G/C]TGG[G/A]AAGAAATGGGCTAC	L	12 s	789 (Mus); 2781 (Xen)	M. Garcia-Paris/C. Orrego
MVZ 64	TGT GGG TAA CAG ATG AGA ATG CTG	H			R. Robertson
MVZ 65	TAY AAA GAA ACH TGA AAY AT	L	cyt b	14482 (Mus)	S. dos Reis/C. Orrego
MVZ 66	GAT GAA TGG GTG TTC TAC TGG TT	H	cyt b	15160(Mus);15768(Hu); 15917(Chi)	C. Cicero
MVZ 67	CAY ATY AAA CCM GAA TGR TA	L	cyt b	14956 (Mus)	S. dos Reis
MVZ 68	GAT GTC CTT TGT GGA GAA GTA GG	H	cyt b	14803 (Mus);15560 (Chi)	C. Cicero
MVZ 69	AAA ATT GTA AAY CAY GCC	L	cyt b	14189 (Mus)	M.F. Smith/W.K. Thomas
MVZ 70	CAT CGT GAT GCC TTA TTT AA	H	D-loop	15836 (Mus)	L. F. Garcia
MVZ 71	TGA GGA CAA ATA TCV TTC TGA GG	L	cyt b	14563 (Mus);15321 (Chi)	C. Orrego/C. Cicero
MVZ 72	GGA GTA TGA GCC GTA GTA GAT TC	H			R. Robertson
MVZ 73	CGA TAT GCA CAY GCY AAY GGA	L	cyt b	14396 (Mus)	M. da Silva/C. Orrego
MVZ 74	GGG ATG GCT GAT AAG AGG TTT GTG A	H			R. Robertson
MVZ 75	GGW CAR CCN GTN GAR GAY CC	L		1318 (Xen)	A. Graybeal/C. Orrego
MVZ 76	GGT TAA GGA TAG TCA TGT ACT G	H			L. F. Garcia
MVZ 77	CGC ACA CAC CGC CCG TCA CCC TC	L	XL PCR	920 (Mus)	C. Schneider
MVZ 78	AAA CTG CTG CTT CTG CTG	H	SRY (see nuclear)		L. F. Garcia
MVZ 79	CAC TTC TAG GAM TAT GCC TAA TCA T	L			R. Robertson
MVZ 80	TCA TGG AAC TGC TGC TTC T	H	SRY (see nuclear)		L. F. Garcia
MVZ 81	GAT GAC TCA TYC GAT ACA TWC ATG C	L			R. Robertson
MVZ 82	CCA GAG ATT AGA GGG AAT CAG TG	H	CO I	6510(Mus);7110(Hu); 7827(Chi)	L. F. Garcia/S.R. Palumbi; identical to CO1e of Palumbi
MVZ 83	AAA GAA ACC TGA AAC ACA GGC GT	L	cyt b	14488(Mus);15245(Chi)	C. Cicero
MVZ 84	CGG CAG GTA TGT GTT ACT GAT G	H	cyt b	14329 (Mus)	M. Ruedi
MVZ 85	ACC TTC CTC CAT GAA ACA GGA TC	L	cyt b	14752(Mus);15509(Chi)	C. Cicero
MVZ 86	TTT GTR ATT ACT GTG GCC CCT CA	H	cyt b	14559(Mus);15316(Chi)	N. Allen
MVZ 87	GAA GGC TAT ACC CTT CCT AGG ACA	L	D-loop		L. F. Garcia
MVZ 88	GGT GTA TGC ATC GGG GTA GTC	H	CO I	6651 (Mus);7227 (Hu)	L. F. Garcia; modified CO1a Palumbi
MVZ 89	ACA GGN TCA AAY AAC CCA ACA GG	L			E. Jackusch/K. Le

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MVZ 90	CCT GAC AAT AGG GGA AAT CAG TG	H	CO I		L. F. Garcia; modified COIe Palumbi
MVZ 91	TCC TAC ACA GAG AGA AAT ACC	L	SRY (see nuclear)		L. F. Garcia
MVZ 92	ATT TGK CCA ATK GTT GTG AAT G				
MVZ 93	CCT GCA GGA GGA GGA GAY CC	L	CO I	6010 (Mus);6569 (Hu); 7326 (Chi)	L. F. Garcia/S.R. Palumbi; identical to CO1f of Palumbi
MVZ 94	AAA TGA TCN CCM GCY ATA CTA TGA	H	CO I	6493 (Mus)	CO1f+ L Garcia
MVZ 95	GAC AAT TAT ACG CAA GTC TCA CCC	L	cyt b	14164 (Mus)	M. Ruedi
MVZ 96	GGATTGCGCTGTTATCCCTA	H	16S	2371 (Mus)	16Sbr+ L. Garcia
MVZ 97	CCT ACT TAG GAT CAT TCG CCC T	L	ND5/cyt b	14851 (Chi)	C. Cicero/J. G. Groth
MVZ 98	CCGGTCTGAACTCAGATCACGT	H	16S	2510 (Mus)	16Sbr of Palumbi
MVZ 99	CTA TAC AAA GAA ACC TGA AAC ACA GG	L	cyt b	14485(Mus);15242(Chi)	N. Allen
MVZ 100	ATGTTTTTGATAAACAGGCG	H	16S	1928 (Mus)	16Sa of Palumbi
MVZ 101	TCC TAC TTA GGA TCA TTC GCC CTA TC	L	cyt b	?14061(Mus);14818(Chi)	N. Allen
MVZ 102	TAAGTGCAGAAGGCTAGGACCAAACCT	H	D-loop/12S	72 (Mus); 00651 (Hu)	H00651 of Kocher et al 1989
MVZ 103	CAC CCT AAC MCG CTT CTT CGC	L	cyt b	14677 (Mus)	M. F. Smith
MVZ 104	CCTGAAGTARSAACCAGWWG	H	D-loop	15782 (Mus)	FUMA3 E. Lacey; mod. H16498M Fumagalli et al 97
MVZ 105	TCA ACC ACA CTC CAC ACA GG	L	ND5/cyt b	14818 (Chi)	C. Cicero/J. G. Groth
MVZ 106	CCATCTAAGCATTTTCAGTGCT	H	D-loop	27 (Mus)	FUMA4 E. Lacey; identical to H605N Fumagalli et al 1997
MVZ 107	CCC GCC GGA GGA GGA GAY CC	L	CO I	6010 (Mus);6569 (Hu)	L. F. Garcia; modified CO1f of Palumbi
MVZ 108	CCA ATG TAA TTT TTA TAC	H	cyt b/tRNA-Glu	15292 (Mus)	TUCO14A E. Lessa/J. Cook
MVZ 110	TTTCATCTTTTCCTTGCGGTAC	H	12S	1210 (Mus)	12G of Springer et al '95
MVZ 111	CCY CTG TST TTA GAT TTA CA	L	CO I	5299 (Mus)	L. F. Garcia
MVZ 112	ACC TCG GTA GCA GTG GTT AGC	H	ND4/tRNA-Ser	11626 (Mus)	G Parra
MVZ 113	TCA TAG TAT RGC DGG NGA TCA TTT	L	CO I	6457 (Mus)	CO1a+ L. Garcia
MVZ 114	TGGCTGGCACGARATTTACCA	H	D-loop/12S	295 (Mus)	E. Lacey (tucu 12S)
MVZ 115	ATAAGACGAGAAGACCCTATG	L	16S	2169 (Mus)	16Sar+ L. Garcia
MVZ 116	GTT GAA CTT TCA TTT CTA GGG YCA C	H	ND4	11569 (Mus);13701 (Xen)	M. Mahoney (NDHis)
MVZ 117	CGCCTGTTTATCAAAAACAT	L	16S	1947 (Mus)	16Sar of Palumbi
MVZ 118	YAT GDD DTT GGC TTG AAA CC	H	CO I	6902 (Mus)	L. F. Garcia
MVZ 119	ACGTGATCTGAGTTCAGACCGG	L	16S	2522 (Mus)	16Sb of Palumbi
MVZ 121	ACCAYYARCACCCAAAGCTRA	L	D-loop	15393 (Mus)	FUMA1 E. Lacey; mod. L15999M Fumagalli et al 1997
MVZ 123	CWTCTGGTTBYTACTTCAGG	L	D-loop	15801 (Mus)	FUMA2 E. Lacey; mod. L16517M Fumagalli et al 1997

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MVZ 125	TTC TAA TTA AAC TAT TTC TTG	L	D-loop/tRNA-Pro	15416 (Mus)	TUCO-PRO E. Lessa/J. Cook
MVZ 127	TRY TAC CAT GAG GAC AAA TAT C	L	cyt b	14554 (Mus)	TUCO1 E. Lacey
MVZ 129	CCY CAY ATY AAA CCA GAA TG	L	cyt b	14953 (Mus)	TUCO2 E. Lacey
MVZ 131	AAAGCAAARCACTGAAAATG	L	12S	25 (Mus)	12C of Springer et al '95
MVZ 133	GGC TAT GGG ATA ATC CGA GTT ATA TTT A	L	ND4	10905 (Mus);13008 (Xen)	M. Mahoney (ND4-M)
Cyt b1	AAAAAGCTTCCATCCAACATCTCAGCATGATGAAA	L	cyt b	14233(Mus);14841 (Hu)	T. Kocher
Cyt b2	AAACTGCAGCCCCCTCAGAATGATATTTGTCCTCA	H	cyt b	14541(Mus);15149(Hu)	T. Kocher
H15768	ATGAAGGGATGTTCTACTGGTTG	H	cyt b	15159(Mus);15768(Hu); 15916(Chi)	C. Cicero/S. V. Edwards
Dipotas	CTTGTCATGTTTCATCGTGCATT	L			F.X. Villablanca
Dpd 4	TACATTAGACATTAATCCACATAA		D-loop		F.X. Villablanca
Dpd 6	GTCCTTGTCATATGACTATC		D-loop		F.X. Villablanca
Dpd 7	TGGTTTCACGGAGGATGGTA		D-loop		F.X. Villablanca
Tb-1	TAATTCCTCCTCCATTAACCCA	L	D-loop		K. Thomas
Tb-2	GGTTGCTGTTTCACGGAGG	H	D-loop		K. Thomas
Tb-2 + GC	CGCCCGCCGCGCCCCGCGCCCGCCGCGCCCCCGCCCC- GGTTGCTGTTTCACGGAGG		D-loop		E. Lessa
Tb-3	AAACTCTTCAAACCAACGTA		D-loop		E. Lessa
Tb-5	TCCACCACATAACTCAAC		D-loop		E. Lessa
TDKD	CCTGAAGTAGGAACCAGATG	H	D-loop	15782(Mus);16498 (Hu)	T. Kocher
TDKD+GC	CGCCCGCCGCGCCCCGCGCCCGCCGCGCCCCCGCCCC- CCTGAAGTAGGAACCAGATG	H	D-loop	15782(Mus);16498 (Hu)	F.X. Villablanca/C. Orrego
Thr	TCAAAGCTTACACAGTCTTGTAACCC	L	D-loop/tRNA-Thr	15320(Mus);15926 (Hu)	L15926 of Kocher et al, 1989
Thr+GC	CGCCCGCCGCGCCCCGCGCCCGCCGCGCCCCCGCCCC- TCAAAGCTTACACAGTCTTGTAACCC	L	D-loop/tRNA-Thr	15320(Mus);15926 (Hu)	
Thr-Bt	Bt-TCAAAGCTTACACAGTCTTGTAACCC	L	D-loop/tRNA-Thr	15320(Mus);15926 (Hu)	
ND2-1	AAC TTC CTT CCA CTT ACM CTA GC	L	ND2	3646 (Mus); 5709 (Xen)	M. Mahoney/C. Schneider
ND2-2	AGT CGT ATG TAG AAG AAT AGG YT	H	ND2	4778 (Mus); 6843 (Xen)	M. Mahoney/C. Schneider
ND2-3	GCA TCT ACA AAA TAC TTC TTV AC	L	ND2	4098 (Mus); 6163 (Xen)	M. Mahoney/C. Schneider
ND2-4	GGT ATA AAT CCT GTT AGT GGC GG	H	ND2	4676 (Mus); 6741 (Xen)	M. Mahoney/C. Schneider
ND2-B1	GCT AAC AAA GCT ATC GGG CCC AT	L	ND2/tRNA-Met	3877 (Mus); 5204 (Chi)	C. Cicero/J. G. Groth
ND2-B2	CCT TGG AGT ACT TCT GGG AAT CAG A	H	ND2	4251 (Mus); 5578 (Chi)	C. Cicero/J. G. Groth
ND2-B3	AAT GCA TGA TCC ACC GGC CAA TGA GA	L	ND2	4167 (Mus); 5494 (Chi)	C. Cicero/J. G. Groth
ND2-B4	TTG GTT AGT TCT TGG ATA ATG AGT CA	H	ND2	4703 (Mus); 6030 (Chi)	C. Cicero/J. G. Groth
ND2-B5	GCC TTC TCA TCC ATC TCC CAC CTA GGA TGA AT	L	ND2	4482 (Mus); 5809 (Chi)	C. Cicero/J. G. Groth
ND2-B6	CTT ATT TAA GGC TTT GAA GGC C	H	ND2/tRNA-Trp	4986 (Mus); 6312 (Chi)	C. Cicero/J. G. Groth

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ND3-B1	CTC TAC ACA ACC ATC TAC TGA TGA GG	L	ND3/COIII	10701 (Chi)	C. Cicero/J. G. Groth
ND3-B2	GAT AGT ATT ATG CTT TCT AGG CA	H	ND3/ND4L	11289 (Chi)	C. Cicero/J. G. Groth
LEU	CAT TAC TTT TAC TTG GAT TTG CAC CA	H	ND4/tRNA-Leu	11713 (Mus)	Arevalo et al '94
ND4	CAC CTA TGA CTA CCA AAA GCT CAT GTA GAA GC	L	ND4	10828 (Mus)	Arevalo et al '94
His	AACCAAAACATCAGATTGTGAATCT	L	ND5/tRNA-His	11570 (Mus);12175 (Hu)	T. Kocher
ND5	AATGTTTGGGTCTGAGTTTATATATCA	H	ND5	12051 (Mus);12652 (Hu)	T. Kocher
NS7	GAG GCA ATA ACA GGT CTG TGA TGC				C. Orrego
TAS					F.X. Villablanca