

Table S2. ND2 sequence alignment for the hummingbird species registered in the Cloud Forest Sanctuary.

Species	GenBank accession number	ND2 sequence (1041 bp)
<i>Chaetura pelagica</i> (outgroup)	MH719133.1	ATGAACCCTTACGCTAAACTAATCTCACCATAGCCTACTCCTAGGAACAACCATAACAATCTTAGCAGCC ACTGAATCCTAGCCTGGACAGGACTAGAAATCAACACCCTAGCCATCATCCCCTACTCATCGAAAACACCACCA CCCACGAGCCATCGAACGCCAAATATTCCTAGTCCAAGCAGCCGCTAACACTACTACTATTCTCA AGCACAATTAATGCCTGGCTCCGGACAATGGGACATCACACAACTAAGCCACCCATCTCATGTGCCCTAC TAACAACAGCGATTGCAATAAAACTCGGGCTAGTCCCATTCTGATTCCCAGAAGTCCTCCAAGGATC CTCATTAACAACCGCAATACTCCTATCACAAATAAAATTCCCCCAATCACCATCCTCTCATAACAGCC CACTCACTAAACCCAGCCCTCCTTACCCATGGCAGTTCTCAGCCGCTTGAGGCTGAATAGGGCTTA ACCAAACACAAACTCGAAAAACTCTAGCTTCTCATCTCACCTCGGATGAATAGCTATCATCCTTAT TTACAACCCAAACTAACCTACTAACCTTCTACGTATATTGCCTATAACCATCACTGTATCCTCACCCCTA AACACAACAAAACTCTTAAACTATCTACAATAAACTTCCTGAACAAAACCCAAACACTAAATGCAGCCT TAATACTAATCTCCTATCCTTAGCCGGACTCCCACCTCTAACAGGATTCTACCTAAATGACTCATCACTCA AGAACTTACCAAACAAGAAATAACTCAACAGCCACAATCATCACACTCCTCTCCCTTAGGCCTATTCTC TACCTCCGCCCGCATACTACTCAACAATTACACTTCCTCCAAACCCACAAACACATAAAACAGTGACATG TCTACAAACTAACAAATACCCTATTCCATCCTCACACACTCTCACCTGCTCCTCCCCCTCCTCCCCAT AATCCTTGCCTATTAG
<i>Atthis heloisa</i>	KX855345.1	ATGAACCCTTACGCTAAACTAATTCTCTTTGAGCCTTATATTAGGGACAACCATTACCCATTCAAGTAGCC ATTGAATGTCGGCCTGGCCGGACTTGAGATCAACACCCTGGCCATCATCCGCTATCTCAAAGTCCCACCA CCCCCGAGCCATTGAAGCAGCGATCAAATATTCCTCGTCCAAGCTGCTGCCTTACCCACTGCTATTCTCA AGCATAATAATGCCTGACACACAGGACAATGAGACATCACTCAACTAAACCACCCACATCATCCCTATTAC TAACCACAGCGATTGCAATAAAACTGGCTGGTACCCCTTCACTCTGATTCCCCGAAGTCCTGCAAGGCTC CCCCTTAACAAACGCCACTCTTACAGTAATAAAATTCCCTCAATGACGATCCTTCACATCC CACTCCCTAAACCCAAACCTACTCACTACCATAGCAGTCTCAGCCGCTTAGGGGGTGAATAGGACTAA ATCAAACACAGACTCGAAAAACTCTAGCCTTCTCATCAATCTCATTTAGGGTGAATAACCACATCATCCTCAT CTACAACCCAAACTAACACTAGTAACCTTCTACCTTATTGCATCATAAACCGGTGCCATCTCCTAACTTTA AATACCACAAATCCCTTAAACTCTCAACGATAATAACCTCTGAACAAAACCTCCGTACTAACACCACAT TAATAATAACACTACTCACTAGCAGGATTACCCCACTAACAGGATTCTGCCAAATGACTCATCATCCA AGAACTCTCCAAGCAAGAAATAGCCACCTCCGCTACAATTATGCCATCCTATCCCTACTGGGCCTATTCTC TATCTCCGACTAGCCTACTACTCAACAATCACTCTCCGCTAACCCACAAACCATATAAAACAATGACACA CTAATAAACCGCACAAGCGCTTAATCGCTACACTCTCCTCTGTCAACCCCTTTACTCCCACATCCCCAAT AGTCCTGCCCTCCATCTAA
<i>Lampornis amethystinus</i>	KJ602255.1	ATGAACCCTTACGCTAAATTAAACATTCCCTATAAGCCTTATGCTAGGAACAACCATTCACTGCTCAAGTAATC ACTGAGTGTCAAGCCTGAGCCGGACTTGAAATCAACACCCTAGCCATCATCCCTAAATTCTAAGTCCCACCA CCCCCGAGCCATTGAAGCAGCAATCAAATATTCCTTGTCCAAGCTGCTGCCTCCACTCTTCTATTATTCTCA AGCATGATAATGCTGACACACAGGACAATGAGACATCACTCAACTAAACCACCCACATCATCCCTACTAC TAACCACAGCAATTGCAATAAAACTCGGCTTGTGCCCTTCCATTCTGATTCCCCAGAAGTCCTGCAAGGCTC CCCCCTAACACAGCTACTTCTACAGTAATAAAATTCCCCCAGTCACCATCTTCCACATCT CACTCCCTAAACCCAACTCTACTTACCATATAAGCCCTAGCCTCCGCGATTAGGGGGTGAATAGGACTAA ACCAAACACAAATTCTGAAAGTCTAGCCTTCTCATCAATCTCACACCTAGGATGAATGGCATTATCCTTAT CTACAACCCAAACTAACATTAATAACCTTCTACCTTACTGTGTCTAACACCAGCGCCATCTCCTAACCCCTA AACACCACAAATCCCTCAAACATCAACAATAATGACCTCTGGACAAAACCTCTGTACTCAACACCGCAT TAATAATAACACTACTCTCATGGCAGGCTTACCCCACTTACAGGATTCTACCCAAATGACTTGTATCCA AGAACTCTCCAAGAAATATCTTCTCTGCCACAGTTATGCCATCTGTCTTATTGGGTTATTTTC TACCTCCGCTAGCCTACTACTCAACCATTCACTCTCCACCAAACCCACAAACCATAAAACAATGATACA CTTATAAAACCCACAAGCAGCCTAATGCCACGCTCTCCTCCATCAGCCCTTTACTCCCACGTCCCCAAT AATCCTGCCCTCCATCTAA

*Pampa excellens** KJ602200.1

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TACTTACGCCCTAGCCTACTACTCAACAAATCACCTACCACCAAGCCCTACAAACCATAAAACAATGATACA
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AATCCTCGCCTCCATCTAA

Amazilia yucatanensis KJ602180.1

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Saucerottia cyanocephala KJ602167.1

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Chlorestes candida KJ602164.1

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Saucerottia beryllina KJ602161.1

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Anthracothonax prevostii EU983409.1

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Colibri thalassinus EU042544.1

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CTAATAAACCTGCAAGCAGCCTAATGCCATGCTCTCCCTGTCAGCCCTCCTACTCCACTATCCCCAAT
AATCCTCGCCTCCATTTAA

* Due the available sequences on GenBank for Wedge-tailed Sabrewing (*Pampa curvipennis*) are partial (KC858426 393 bp and KC858427 393 bp), we used a complete sequence of gene ND2 of closely related species Long-tailed Sabrewing (*Pampa excellens*).