

Table S1. Curation notes regarding the chironomid database: comments, corrections, additional literature and omitted sequences.

Branch Number	Consensus Identification	# of sequences removed	Name and/or sequence removed from database as a likely error	Curated Branch Annotation (A= adult; L=larva)
11	<i>Dicrotendipes lucifer</i> agg.	-	<i>Dicrotendipes lucifer</i>	Specimens were identified as <i>D. lucifer</i> (1 A) and <i>D. simpsoni</i> (3L). Both species are part of a complex known as <i>D. lucifer</i> agg. (Epler, 1987) and therefore combined under this single name.
26	<i>Cryptotendipes pseudotener</i>	1	<i>Cladopelma</i> sp.	Specimens were identified as <i>C. pseudotener</i> (1A) and <i>Cladopelma</i> sp. (1A). Closest genus level sequences in GenBank include <i>Cryptotendipes</i> , (10.5% distance), not <i>Cladopelma</i> . <i>Cladopelma</i> genera barcodes, including those in GenBank and this database, are >13% different from the removed sequence
35	<i>Harnischia curtilamellata</i>	1	<i>Cryptochironomus</i> sp.	Specimens were identified as <i>Harnischia</i> sp. (3L), <i>Harnischia curtilamellata</i> (1A), and <i>Cryptochironomus</i> sp. (1L). <i>Cryptochironomus</i> sp. did not agree with the consensus genus and was removed.
36	<i>Chironomus entis/plumosus</i>	2	<i>Chironomus crassicaudatus</i> <i>Chironomus magnus</i>	Specimens were identified as <i>Chironomus</i> sp. (3L6A), <i>Chironomus plumosus</i> gr. (2A), <i>Chironomus plumosus</i> (5A), <i>Chironomus entis/plumosus</i> (1A) <i>Chironomus crassicaudatus</i> (1A), and <i>Chironomus magnus</i> (1A). The last two did not agree with the consensus species and were removed.[1,2].
40	<i>Chironomus bifurcatus</i>	1	<i>Chironomus decorus</i>	Specimens were identified as 5A <i>C. bifurcatus</i> , 1A <i>Chironomus</i> sp., and 1A <i>C. decorus</i> . The one <i>C. decorus</i> did not agree with the consensus species and was removed.
42	<i>Chironomus crassicaudatus</i>	1	<i>Goeldichironomus</i> sp.	Specimens were identified as 4L <i>C. crassicaudatus</i> , 60L <i>Chironomus</i> sp., 5A <i>C. crassicaudatus</i> , 1A <i>Chironomus</i> sp., and 1L <i>Goeldichironomus</i> sp. The <i>Goeldichironomus</i> sp. did not agree with the consensus genus and was removed.[3,4].
43	<i>Paratanytarsus</i> sp.	1	<i>Micropsectra</i> sp.	Specimens were identified as 1A <i>Paratanytarsus</i> sp. and 1A <i>Micropsectra</i> sp. BLAST of the <i>Micropsectra</i> sequence gave more than 40 >96.5% matches—to <i>Paratanytarsus</i> ! Thus, <i>Micropsectra</i> did not agree with the consensus identification and was removed.[5].
53	<i>Pseudochironomus</i> sp.	1	<i>Polypedilum halterale</i> gr.	Specimens were identified as 2L <i>Pseudochironomus</i> sp. and 1L <i>Polypedilum halterale</i> gr. Sequences were within 2.6% of another set of specimens (branch 52) that were 3L <i>Pseudochironomus</i> sp. and 1A <i>Pseudochironomus fulviventris</i> . <i>Polypedilum halterale</i> gr. did not agree with the consensus genus identification and was removed.[6].
54	<i>Polypedilum halterale</i> gr.	1	<i>Polypedilum scalaenum</i> gr.	Specimens were identified as 26L and 2A <i>Polypedilum halterale</i> gr., and 1L <i>Polypedilum scalaenum</i> gr. <i>Polypedilum scalaenum</i> gr. did not agree with the consensus species and was removed.
56	<i>Polypedilum halterale</i> gr.	1	<i>Polypedilum scalaenum</i> gr.	Specimens were identified as 17L <i>Polypedilum halterale</i> gr. and 1L <i>Polypedilum scalaenum</i> gr. . <i>Polypedilum scalaenum</i> gr. did not agree with the consensus species and was removed
58	<i>Cricotopus sylvestris</i>	1	<i>Cricotopus festivellus</i>	Specimens were identified as 7L and 1A <i>Cricotopus</i> sp., 1A <i>Cricotopus sylvestris</i> , and 1A <i>Cricotopus festivellus</i> . <i>C. festivellus</i> did not agree with the consensus GenBank sequences within 3.5% of the sequence and was >5.9% different from other specimens identified as <i>C. festivellus</i> . Therefore, <i>C. festivellus</i> was removed from this branch.[7-9].
78	<i>Procladius sublettei</i>	1	<i>Cladopelma viridulum</i>	Specimens were identified as 10L <i>Procladius</i> sp., 2A <i>Procladius sublettei</i> , and 1A <i>Cladopelma viridulum</i> . <i>Cladopelma viridulum</i> did not agree with the consensus genus and was removed.

83	<i>Coelotanypus scapularis</i>	2	<i>Chironomus</i> sp. <i>Procladius</i> sp.	Specimens were identified as 61L <i>Coelotanypus</i> sp., 1A <i>Coelotanypus scapularis</i> , 1L <i>Chironomus</i> sp., and 1L <i>Procladius</i> sp. <i>Chironomus</i> sp. and <i>Procladius</i> sp. did not agree with the consensus genus and was removed.
86	<i>Tanytarsus mendax</i>	1	<i>Procladius</i> sp.	Specimens were identified as. and 1A <i>Tanytarsus mendax</i> and 1A <i>Procladius</i> sp. BLAST of the <i>Procladius</i> sequence gave more than 100 >96.5% matches—to <i>Tanytarsus mendax</i> ! Also, this sequence clusters within the Tanytarsini (see tree Fig. 2). Thus, <i>Procladius</i> did not agree with the consensus identification and was removed.[10].
91	<i>Rheotanytarsus exiguus</i> gr.	1	<i>Paratanytarsus</i> sp.	Specimens were identified as 3L <i>Rheotanytarsus exiguus</i> gr. and 1L <i>Paratanytarsus</i> sp. <i>Paratanytarsus</i> sp. did not agree with the consensus genus and was removed.[11,12].
93	<i>Stictochironomus devinctus</i>	1	<i>Polypedilum halterale</i> gr.	Specimens were identified as 13L <i>Stictochironomus</i> sp., 3A <i>Stictochironomus devinctus</i> , and 1L <i>Polypedilum halterale</i> gr. <i>Polypedilum</i> did not agree with the consensus genus and was removed.[13].
95	<i>Cladotanytarsus</i> sp.	3	<i>Coelotanypus</i> sp. <i>Cryptochironomus</i> sp. <i>Stempellinella</i> sp.	Specimens were identified as 27L <i>Cladotanytarsus</i> sp., 1L <i>Coelotanypus</i> sp., 1L <i>Cryptochironomus</i> sp., and 1L <i>Stempellinella</i> sp. <i>Coelotanypus</i> sp., <i>Cryptochironomus</i> sp., and <i>Stempellinella</i> sp. did not agree with the consensus genus and were removed.
99	<i>Micropsectra insignilobus</i>	-	<i>Cladotanytarsus</i> sp.	Failla et al. (2016) determined that this specimen's sequence, originally identified as <i>Cladotanytarsus</i> sp., was most likely a misidentified <i>Micropsectra insignilobus</i> . The sequence matches 100% to a large number of <i>Micropsectra insignilobus</i> in BOLD and is therefore renamed here as <i>Micropsectra insignilobus</i> .
100	<i>Paratanytarsus natvigi</i>	1	<i>Micropsectra</i> sp.	Specimens were identified as 1L <i>Paratanytarsus</i> sp., 1A <i>Paratanytarsus</i> sp., 2A <i>Paratanytarsus natvigi</i> , and 1A <i>Micropsectra</i> sp. <i>Micropsectra</i> sp. did not agree with the consensus genus and was removed.
Total number of sequences removed from database as a likely error		21		

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