

Supplementary Table S1. List of characters used in the phylogenetic analysis of the primitive syringophilid genera (genera with a full complement of body and leg setae).

No.	Character
1	Basal part of gnathosoma: not submerged (0); deeply submerged into idiosoma (1)
2	Stylophore apodeme: indistinct (0); distinctly developed (1)
3	Palpal tibia and tarsus: separated (0); fused (1)
4	Claw-like seta of palps: present (0); absent (1)
5	Posterior part of stylophore: constricted (0); rounded (1)
6	Projection on posterior part of stylophore: absent (0); present (1)
7	Hypostomal apex: unornamented (0); ornamented (1)
8	Hypostomal protuberances: one pair small and blunt-ended (0); one pair small and sharp-ended (1); two pairs (2); one pair large (3)
9	Size of hypostomal lips: small (0); large (1)
10	Hypostomal lips: equal (0); unequal (1)
11	Lateral branch of peritreme: normal i.e., 6-10 chambers (0); reduced i.e., 1-5 chambers (1); supernumerary i.e., 11-20 chambers (2)
12	Distal tip of chelicerae: edentate (0); dentate (1)
13	Propodonotal shield: strongly sclerotized (0); weakly sclerotized (1)
14	Hysteronotal shield: present (0); absent (1)
15	Hysteronotal shield: entire (0); divided (1)
16	Hysteronotal shield: reach level of setae <i>d2</i> (0); not reach <i>d2</i> (1)
17	Pygidial shield: present (0); absent (1)
18	Hysteronotal and pygidial shields: fused (0); not fused (1)
19	Position of setae <i>vi</i> and <i>ve</i> : at same level (0); <i>vi</i> anterior to <i>ve</i> (1)
20	Position of setae <i>c1</i> and <i>se</i> : at same level (0); not at same level (1)
21	Position of setae <i>se</i> and <i>c2</i> : <i>se</i> posterior to <i>c2</i> (0); <i>se</i> and <i>c2</i> at same level (1)
22	Position of setae <i>c2</i> : out of shield (0); on shield (1)
23	Ornamentation of idiosomal setae: present (0); absent (1)
24	Setae <i>e1</i> : present (0); absent (1)
25	Position of setae <i>f1</i> and <i>f2</i> : <i>f1</i> close to <i>f2</i> (0); <i>f2</i> far from <i>f1</i> (1)
26	Setae <i>ps3</i> : present (0); absent (1)
27	Genital setae: short (0); long (1); medium (2)
28	Position of setae <i>ag1</i> and <i>ag2</i> : <i>ag2</i> postero lateral to <i>ag1</i> (0); <i>ag2</i> posterior to <i>ag1</i> (1); <i>ag1</i> and <i>ag2</i> at same level (2)
29	Neotrichious aggenital setae: absent (0); present (1)
30	Seta <i>a'</i> on tarsus I: present (0); absent (1)
31	Proral setae <i>p'</i> and <i>p''</i> : rod-like (0); fan-like (1)
32	Number of tines in proral setae: reduced i.e., 4-10 tines (0); normal i.e., 11-20 tines (1); supernumerary 21-35 tines (2)
33	Setae <i>4a</i> : present (0); absent (1)
34	Seta <i>vs</i> on tarsus III: present (0); absent (1)
35	Seta <i>vs</i> on tarsus IV: present (0); absent (1)
36	Seta <i>dG</i> on genu IV: present (0); absent (1)
37	Seta <i>vF</i> on femur III: present (0); absent (1)
38	Seta <i>v</i> on trochanter III: present (0); absent (1)
39	Legs I and II: subequal in thickness (0); I thicker than II (1)
40	Legs I and II: I longer than II (0); I and II subequal in length (1)
41	Coxal fields I-II and III-IV: grouped together (0); widely separated (1)
42	Apodemes I and II: similar in shape (0); different in shape (1)
43	Apodemes I: divergent (0); parallel (1)
44	Divergence of apodemes I: strongly divergent (0); slightly divergent (1)
45	Apodemes I and II: fused (0); not fused (1)
46	Apodemes III and IV: present (0); absent (1)
47	Claw basal angle: absent (0); present (1)
48	Position of setae <i>3a</i> and <i>3b</i> : at same level (0); <i>3a</i> anterior to <i>3b</i> (1)
49	Body size: large i.e., more than 1000 (0); medium i.e., 700-990 (1); small i.e., less than 690 (2)

Supplementary Table S2. Data matrix

	1	2	3	4
	1234567890123456789012345678901234567890123456789			
<i>Cheyletus eruditus</i>	0000--0-00000000010000100000010-00000000000010000			
<i>Cheletopsis norneri</i>	0000--0-000001--1-0101100000010-00000000000010000			
<i>Aulobia dendroicae</i>	1111100-11200001011000010100001011111101111-11011			
<i>Aulobia cardueli</i>	1111100-11200001011000010100001011111101111-11011			
<i>Blaszakia rossae</i>	1111100-0000000001120011110100101111111110111012			
<i>Bubophilus aegolius</i>	1111000-00110001011000010101001111111101110111011			
<i>Charadriphilus ludmilae</i>	1111000-00200000011200111101001011111101110111012			
<i>Charadriphilus ralli</i>	1111000-01000000011210111101001011111101110111012			
<i>Colinophilus wilsoni</i>	1111000-0021000001120001010100111111111110111010			
<i>Corvitorotrogus alpha</i>	111100110001000101110101010-101111111101110001010			
<i>Creagonycha totani</i>	1111000-000111--0-1101010101001111111101100011000			
<i>Crotophagisyringophilus io</i>	11110011002101--0-1102010101001111111101110101010			
<i>Ixobrychiphilus wallacei</i>	1111000-00111001001100010101001011111101111-11012			
<i>Kalamotrypates cracidus</i>	1111110-00000000011200010100001011111111110111012			
<i>Megasyringophilus eos</i>	11110012001101--1-1101011100001111111101110010100			
<i>Megasyringophilus trichoglossus</i>	1111001?00010001011102011110001111111101110010100			
<i>Megasyringophilus aquilus</i>	1111001100010001011102011100001211111101110010100			
<i>Pteroclidisyringophilus otididus</i>	1111000-00100001011300010100001111111101110111012			
<i>Pteroclidisyringophilus re</i>	1111000-00100000011210010100001111111101110111012			
<i>Selenonycha charadriiformicus</i>	1111101-00010001011101011101001111111101100010000			
<i>Syringophilopsis turdi</i>	11110012002101--0-1101010102001011111101110001010			
<i>Syringophilopsis muscicapicus</i>	1111001200210011001101010101001011111101110001010			
<i>Syringophilus bipectinatus</i>	1111000-00000000011310010100001111111101110101011			
<i>Tinamiphilopsis ariconte</i>	1111010-00010000011300010101001111111101110111012			
<i>Tinamiphilopsis elegans</i>	1111010-00010000011300010101001111111101110111012			
<i>Tinamiphilopsis temmincki</i>	1111010-00010000011300010101001111111101110111012			
<i>Torotrogla meulae</i>	111100130001001100110101010-101111111101100011011			
<i>Torotrogla lusciniae</i>	11110013000101--0-110101010-101111111101100010011			
<i>Trypetoptila casmerodia</i>	1111011???0101--0-100001011-101210111101110001010			