## SUPPLEMENTARY MATERIALS

**Table S1** Springtail collection information showing the location, coordinates and BOLD sample IDs of sequences for each unique haplotype

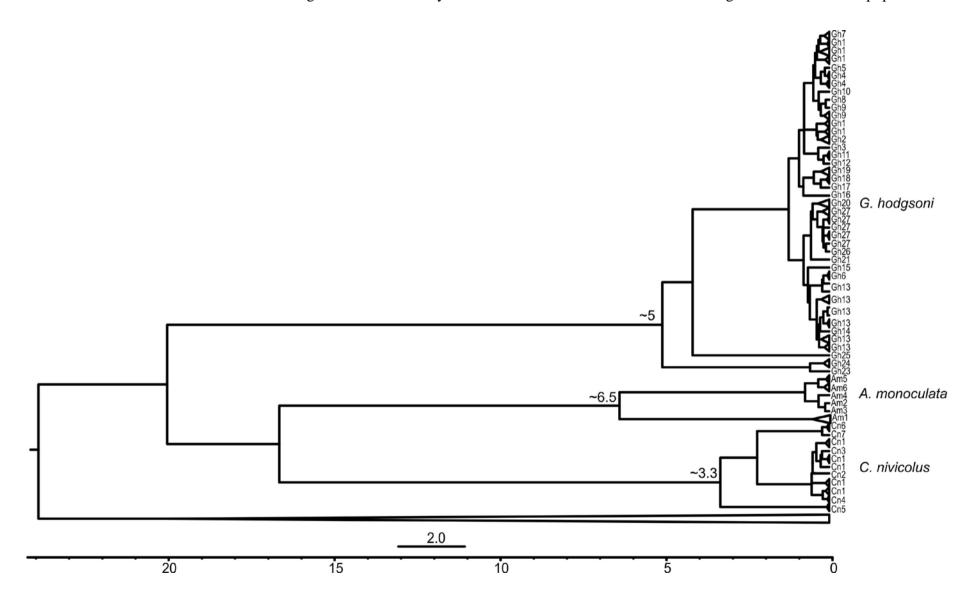
Haplotype #	Location	Coordinates	<b>BOLD Sample IDs</b>
G. hodgsoni			
Gh1	Tiger Island	-76.784 162.452	ANTSP607
	Flatiron	-77.005 162.408	ANTSP585, ANTSP587,
			ANTSP589, ANTSP605,
			ANTSP606, ANTSP607,
			ANTSP628, ANTSP629
	Benson Glacier	-76.822 162.107	ANTSP596, ANTSP597
	St Johns Range	-77.2801 161.731	ANTSP129, ANTSP131,
			ANTSP134, ANTSP136,
			ANTSP137, ANTSP138,
			ANTSP140, ANTSP141,
			ANTSP143, ANTSP151,
			ANTSP216
Gh2	St Johns Range	-77.2801 161.731	ANTSP128, ANTSP144,
			ANTSP145, ANTSP147,
			ANTSP148, ANTSP149,
			ANTSP214, ANTSP218
Gh3	St Johns Range	-77.208 161.7	ANTSP210
Gh4	Flatiron	-77.005 162.408	ANTSP584, ANTSP586,
			ANTSP604
	St Johns Range	-77.208 161.7	ANTSP213, ANTSP215
Gh5	Flatiron	-77.005 162.408	ANTSP631
Gh6	Benson Glacier	-76.87 161.754	ANTSP564, ANTSP565
Gh7	Mount Seuss	-77.015 161.75	ANTSP153, ANTSP166,
			ANTSP167, ANTSP534
Gh8	St Johns Range	-77.208 161.7	ANTSP209

Gh9	St Johns Range	-77.2801 161.731	ANTSP132, ANTSP133,
			ANTSP135, ANTSP139,
			ANTSP211, ANTSP212
Gh10	St Johns Range	-77.285 161.726	ANTSP217
Gh11	Flatiron	-77.005 162.408	ANTSP634, ANTSP635
Gh12	St Johns Range	-77.2849 161.726	ANTSP150
Gh13	Mount Seuss	-77.023 161.738	ANTSP152, ANTSP154,
			ANTSP157, ANTSP158,
			ANTSP159, ANTSP160,
			ANTSP161, ANTSP163,
			ANTSP164, ANTSP165,
			ANTSP168, ANTSP169,
			ANTSP172, ANTSP174,
			ANTSP175, ANTSP225,
			ANTSP518, ANTSP520,
			ANTSP521, ANTSP522,
			ANTSP524, ANTSP532,
			ANTSP533, ANTSP540,
			ANTSP541
	Benson Glacier	-76.87 161.754	ANTSP563
Gh14	Mount Seuss	-77.021 161.737	ANTSP521
Gh15	Mount Seuss	-77.015 161.75	ANTSP535
Gh16	Flatiron	-77.005 162.408	ANTSP627
Gh17	Tiger Island	-76.783 162.452	ANTSP639
Gh18	Tiger Island	-76.783 162.452	ANTSP590, ANTSP612
Gh19	Tiger Island	-76.783 162.452	ANTSP592, ANTSP637,
			ANTSP638, ANTSP640,
			ANTSP642, ANTSP643,
			ANTSP644, ANTSP646
	Benson Glacier	-76.822 162.107	ANTSP623
Gh20	Mount Seuss	-77.011 161.768	ANTSP142, ANTSP173,
			ANTSP519, ANTSP525,
			ANTSP531

Gh21	Mount Seuss	-77.023 161.738	ANTSP539
Gh22	Mount Seuss	-77.2849 161.726	ANTSP142
Gh23	Towle Glacier	-76.655 161.093	ANTSP561
Gh24	Towle Glacier	-76.729 161.011	ANTSP556, ANTSP557,
			ANTSP558, ANTSP559
Gh25	Mount Gran	-76.982 161.16	ANTSP202
Gh26	St Johns Range	-77.2849 161.726	ANTSP146
Gh27	Benson Glacier	-76.822 162.107	ANTSP594, ANTSP595,
			ANTSP598, ANTSP599,
			ANTSP600, ANTSP601,
			ANTSP624, ANTSP649,
			ANTSP651, ANTSP652,
			ANTSP653, ANTSP654,
			ANTSP655, ANTSP656
A. monoculato	ı		
Am1	Cliff Nunatak	-76.11 162.015	ANTSP552, ANTSP553,
			ANTSP554, ANTSP555,
			ANTSP566, ANTSP567,
			ANTSP568, ANTSP569,
			ANTSP579, ANTSP580,
			ANTSP581, ANTSP582
	Mount Murray	-76.11 162.014	ANTSP570,
			ANTSP571,ANTSP572,
			ANTSP573, ANTSP574,
			ANTSP575, ANTSP576,
			ANTSP577, ANTSP578,
			ANTSP613, ANTSP615
Am2	Pegtop	-77.046 161.362	ANTSP618
	Mountain		
Am3	Pegtop	-77.046 161.362	ANTSP616
	Mountain		
Am4	Benson Glacier	-76.822 162.107	ANTSP622
Am5	Springtail Point	-77.168 160.71	ANTSP196, ANTSP235

Am6	Springtail Point	-77.168 160.71	ANTSP194, ANTSP195, ANTSP203, ANTSP204, ANTSP205
C. nivicolus			
Cn1	Tiger Island	-76.784 162.452	ANTSP609
	Mount Seuss	-77.023 161.738	ANTSP124, ANTSP155,
			ANTSP156, ANTSP536,
			ANTSP543, ANTSP544,
			ANTSP548, ANTSP611
	Mount Gran	-76.972 161.148	ANTSP619, ANTSP620,
			ANTSP621
Cn2	Mount Seuss		ANTSP170
Cn3	Mount Seuss	-77.023 161.738	ANTSP547
Cn4	Mount Seuss	-77.023 161.746	ANTSP537, ANTSP550
Cn5	Towle Glacier	-76.655 161.093	ANTSP602, ANTSP603
Cn6	Springtail Point	-77.1676 160.71	ANTSP119, ANTSP121
Cn7	Springtail Point	-77.1676 160.71	ANTSP118

**Fig. S1** Bayesian Inference tree showing estimated divergence times for *Gomphiocephalus hodgsoni*, *Cryptopygus nivicolus* and *Antarcticinella monoculata*. Timescale along the bottom is in My. Numbers on nodes refer to estimated divergence times for those populations.



**Fig. S2** Phylogenetic trees (Bayesian followed by Neighbour Joining) of sequences representing 40 unique haplotypes for three springtail species including three Symphypleona outgroups (*Sminthurides aquaticus* IHCO046-03, *S. malmgreni* IHCO047-03 and *Sminthurinus elegans* COONT067-08). Support values over 0.95/95 are displayed with trees coloured according to the seven BINs present.

