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Coccidia (Apicomplexa: Eimeriidae) Infecting Cricetid Rodents from Alaska, U.S.A., and Northeastern Siberia, Russia, and Description of a New *Eimeria* Species from *Myodes rutilus*, the Northern Red-Backed Vole

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ABSTRACT: During the summers of 2000, 2001, and 2002, 1,950 fecal samples from 4 families, 10 genera, and 16 species of rodents in Alaska, U.S.A. ($N = 1,711$), and Siberia, Russia ($N = 239$) were examined for coccidia (Apicomplexa: Eimeriidae). The 4 families sampled were Dipodidae (jumping mice), Erethizontidae (New World porcupines), Muridae (mice, rats), and Cricetidae (voles, lemmings). Nineteen oocyst morphotypes were observed, of which 10 were consistent with descriptions of known coccidia species from murid hosts, 8 were similar to oocysts described previously from other genera than those in which they are found here (and are called *Eimeria* species 1–8), and 1 is described as new. In the Dipodidae, all from Alaska, 0/15 *Zapus hudsonius* had coccidian oocysts in their feces when examined. In the Erethizontidae, all from Alaska, 0/5 *Erethizon dorsatum* had oocysts when examined. In the Muridae, all from Russia, 0/5 *Apodemus peninsulae* had oocysts when examined. In the Cricetidae from Alaska, we found the following infections: 15/72 (21%) *Lemmus trimucronatus* (*Eimeria* spp. 3, 4, 5); 10/29 (34%) *Microtus longicaudus* (*Eimeria saxei*, *Eimeria wenrichi*); 41/88 (47%) *Microtus miurus* (*Eimeria coahiliensis*, *Eimeria ochrogasteri*, *Eimeria saxei*, *Eimeria wenrichi*); 278/405 (68%) *Microtus oeconomus* (*E. ochrogasteri*, *E. saxei*, *E. wenrichi*); 116/159 (73%) *Microtus pennsylvanicus* (*E. saxei*, *E. wenrichi*); 9/52 (17%) *Microtus xanthognathus* (*E. wenrichi*); 218/699 (31%) *Myodes rutilus* (*Eimeria cernae*, *Eimeria gallati*, *Eimeria marconii*, *Isospora clethrionomydis*, *Isospora clethrionomydis*, and a new *Eimeria* species); 34/187 (18%) *Synaptomys borealis* (*Eimeria* spp. 6, 7, 8, *Eimeria synaptomys*). In the Cricetidae from Siberia, we found the following infections: 5/24 (21%) *Alticola macrotis* (*Eimeria* spp. 1, 2); 0/5 *Dicrostonyx torquatus*; 1/11 (9%) *Lemmus lemmus* (*Eimeria* sp. 3); 30/48 (52%) *Mi.oeconomus* (*E. saxei*, *E. wenrichi*); 5/53 (9%) *Myodes rufocanus* (*E. cernae*, *E. gallati*, *I. clethrionomydis*, the new *Eimeria* sp.); 21/85 (25%) *Myodes rutilus* (*E. cernae*, *E. gallati*, *E. marconii*, the new *Eimeria* sp.); 0/8 *Myopus schisticolor*. Oocysts of the new species, found in both *My. rutilus* (Alaska, Siberia) and *My. rufocanus* (Siberia), are ellipsoidal with a striated outer wall and measured 30.6×20.5 ($27\text{--}33 \times 19\text{--}23$) μm ; micropyle and oocyst residuum absent, but a polar granule is present. Sporocysts are ellipsoidal, 14.5×9.1 ($13\text{--}16 \times 8\text{--}10$) μm ; Stieda body, sub-Stieda body and sporocyst residuum are present.

KEY WORDS: Apicomplexa, Eimeriidae, *Eimeria*, *Isospora*, coccidia, Rodentia, Cricetidae, *Alticola*, *Dicrostonyx*, *Lemmus*, *Microtus*, *Myodes*, *Myopus*, Erethizontidae, *Erethizon*, Dipodidae, *Zapus*, Muridae, *Apodemus*, Siberia, Russia, Alaska, U.S.A.

Over 3 summer field seasons (2000–2002) rodents were collected in Alaska, U.S.A., and northeastern Siberia, Russia, and examined for parasites as part of the Beringia Co-evolution Project (BCP) (see Hoberg et al., 2003). Rarely is there an opportunity to conduct a parasite survey of this magnitude. The main objective of our part in the overall study was to identify the coccidia found in all host animals collected and answer 2 simple questions: 1) Do the same host family, genera, and/or species on different continents share the same coccidia species? 2) How are the parasite assemblages similar or different among closely related hosts? The answers may help us understand the distributions of each parasite species across this large geographic area that was once continuous but is now separated by the Bering Strait.

MATERIALS AND METHODS

Rodents were collected (Cook et al., 2005) using a variety of traps or firearms (University Alaska–Fairbanks animal protocol 99-012 and 01-016 and the Idaho State University animal protocol 03-02-442). Rodents were collected from 10 sites in Alaska, U.S.A., and 4 sites in northeastern Siberia, Russia. The Alaskan sites included 2 regional sites in western Alaska near Nome and Kotzebue, 3 national preserves: Bering Land Bridge, Yukon-Charley Rivers, and Noatak, 4 national parks: Denali, Gates of the Arctic, Kobuk Valley, and Wrangell-Saint Elias, and 1 national monument: Cape Krusenstern. The 4 regions sampled in northeastern Siberia were the Omolon, Anadyr, and Kolyma River basins and the Provideniya Oblast. Animals were dissected 10 min or less after death, their intestinal tract was removed, and feces were preserved in 2.5% (w/v) aqueous potassium dichromate ($\text{K}_2\text{Cr}_2\text{O}_7$) solution. Fecal-dichromate solutions were brought to the University of New Mexico within 1 mo after collection and stored at 4°C. Within 6 mo after collection, oocysts were isolated, measured, and photographed as described by Duszynski and Wilber (1997). Photosyntypes (Duszynski, 1999) of sporulated oocysts of the new eimerian were submitted to the United

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Table 1. Prevalence of coccidia from rodents collected in the summers of 2000-2002, from Siberia, Russia, and Alaska, U.S.A.

Locality	Family/genus/species	No. infected/no. collected (%)	<i>Eimeria</i> and <i>Isoospora</i> species found
Russia	Cricetidae (<i>N</i> = 239)		
	<i>Alticola macrotis</i>	5/24 (21)	<i>E. sp. 1, E. sp. 2</i>
	<i>Dicrostonyx torquatus</i>	0/5	—
	<i>Lemmus lemmus</i>	1/11 (9)	<i>E. sp. 3</i>
	<i>Microtus oeconomus</i>	30/48 (52)	<i>E. saxei, E. wenrichi</i>
	<i>Myodes rufocanus</i>	5/53 (9)	<i>E. cernae, E. gallatii, E. rutilus, I. clethrionomydis</i>
	<i>My. rutilus</i>	21/85 (25)	<i>E. cernae, E. gallatii, E. marconii, E. rutilus</i>
	<i>Myopus schisticolor</i>	0/8	—
	Muridae (<i>N</i> = 5)		
	<i>Apodemus peninsulae</i>	0/5	—
U.S.A.	Erethizontidae (<i>N</i> = 5)		
	<i>Erethizon dorsatum</i>	0/5	—
	Cricetidae (<i>N</i> = 1,691)		
	<i>L. trimucronatus</i>	15/72 (21)	<i>E. sp. 3, E. sp. 4, E. sp. 5</i>
	<i>Mi. longicaudus</i>	10/29 (34)	<i>E. saxei, E. wenrichi</i>
	<i>Mi. miurus</i>	41/88 (47)	<i>E. coahuiliensis, E. ochrogasteri, E. saxei, E. wenrichi</i>
	<i>Mi. oeconomus</i>	278/405 (68)	<i>E. ochrogasteri, E. saxei, E. wenrichi</i>
	<i>Mi. pennsylvanicus</i>	116/159 (73)	<i>E. saxei, E. wenrichi</i>
	<i>Mi. xanthognathus</i>	9/52 (17)	<i>E. wenrichi</i>
	<i>My. rutilus</i>	218/699 (31)	<i>E. cernae, E. gallatii, E. marconii, E. rutilus, I. clethrionomydis, I. clethrionomydis</i>
	<i>Synaptomys borealis</i>	34/187 (18)	<i>E. sp. 6, E. sp. 7, E. synaptomys, E. sp. 8</i>
	Dipodidae (<i>N</i> = 15)		
	<i>Zapus hudsonius</i>	0/15	—
4 families, 16 species	783/1,950 (40)	17 <i>Eimeria</i> , 2 <i>Isoospora</i> species	

States National Parasite Collection (USPNC) in Beltsville, Maryland, U.S.A. (see Bandoni and Duszynski, 1988) as were photomicrographs of other forms (e.g., photoneotypes) when appropriate. Symbiotype host specimens (Frey et al., 1992; Brooks, 1993), in which all oocyst species/forms were seen and identified here, are maintained in the University of Alaska Museum of the North (UAM). All measurements are presented in μm , and we used the abbreviations of Wilber et al. (1998) as follows: length (L), width (W), micropyle (M), oocyst residuum (OR), polar granule (PG), Stieda body (SB), sub-Stieda body (SSB), sporocyst residuum (SR), sporozoite (SZ), and refractile body (RB).

Recently Wilson and Reeder (2005) made substantial changes in the nomenclature of the mammalian order Rodentia. Principally, the long-standing vole genus, *Clethrionomys*, became a junior synonym of *Myodes*. Since we have 2 infected host genera that we refer to repeatedly, and that both begin with the same letter, we use the following genus abbreviations to avoid confusion; *Mi.* for *Microtus* and *My.* for *Myodes*.

RESULTS

In all, 1,950 rodents were collected representing 4 families, 10 genera, and 16 species in Alaska, U.S.A. (*N* = 1,711) and Russia (*N* = 239). The families sampled were Dipodidae (jumping mice), Erethizontidae (New World porcupines), Muridae (mice, rats), and Cricetidae (voles and lemmings).

Wilson and Reeder (2005) list 6 subfamilies in the

Dipodidae. In our study only 1 subfamily, Zapodinae, was sampled with all 15 specimens of *Zapus hudsonius* Zimmerman, 1780, the meadow jumping mouse, negative for coccidian oocysts.

There are 5 genera and 16 species in the Erethizontidae. In this study only 5 specimens of *Erethizon dorsatum*, L., 1758, the North American porcupine, were sampled; all 5 were negative.

We collected only 5 *Apodemus peninsulae* Thomas, 1907, the Korean field mouse, from the Muridae (subfamily Murinae), and all were negative.

Wilson and Reeder (2005) list 6 subfamilies of cricetids. We collected 13 host species in 7 genera within the subfamily Arvicolinae, which included 8 species of "voles" and 5 species of "lemmings." Lemmings and voles comprise an enigmatic group of arvicolid rodents that, as yet, have unresolved phylogenetic relationships (see Discussion). Here we list the 19 putative coccidia species we found in vole genera first, followed by the coccidia found in lemming hosts (see Table 1). For the new eimerian, we give a complete description following the format of Wilber et al. (1998). For the coccidia that either have been recorded from a host genus or species before, or those that resemble strongly species described from other host genera, we provide only

a taxonomic summary and any discrepancies from, or changes to, the original description.

Voles *Eimeria* sp. 1

Host: *Alticola macrotis* (Radde, 1862), no common name recorded.

Other hosts: None, see *Remarks*, below.

Collected: V. B. Fedorov, K. E. Galbreath, 26 July 2002.

Locality: Asia: Russia, northern Siberia, Provideniya Oblast, Chukotka, Ulhum River, 15 km W of Chaplino Village (64°25'N, 172°32'W).

Geographic distribution: Asia: Russia.

Prevalence: 1/24 (4%) (Table 1).

Material deposited: Symbiotype host, UAM 83818 (IF 7551).

Remarks: Svanbaev (1956) was the first to examine a member of this host genus, *Alticola strelzowi* (Kastchenko, 1899), for coccidia and reported finding *Eimeria arvicolae* Galli-Valerio, 1905, originally described from the snow vole, *Microtus nivalis* (Martins, 1842), in western Kazakhstan. Later Svanbaev (1958) reported *E. arvicolae* in 4/43 (9.3%) *A. strelzowi* in the Karaganda Region of central Kazakhstan. The oocysts we found in *A. macrotis* (29 × 21 [26–31 × 19–23]) differed from the spherical oocysts of *E. arvicolae* (14–18 in diameter). In 1980 Dzerzhinskii and Svanbaev (1980) reported finding oocysts they called *Eimeria argentata* in a species of *Alticola* from the Altai Mountains in Kazakhstan, but did not describe the oocysts or provide any images. Thus, this name becomes a *nomen nudum* (fails to conform to Article 12 of the *International Code of Zoological Nomenclature*). Finally, Levine and Ivens (1990) provided the name *Eimeria bassagensis* to the form earlier called *E. arvicolae* by Svanbaev (1958, 1969). The sporulated oocysts we saw in *A. macrotis* differ in shape (broadly ellipsoidal vs. ovoidal), size (29 × 21 vs. 25 × 22), and L/W ratio (1.4 vs. 1.1) from those of *E. bassagensis* and have sporocysts with a SB and SSB, which those of *E. bassagensis* lack. Oocysts of our species closely resemble those of *Eimeria subsimi* Vance and Duszynski, 1985, in *Microtus mexicanus* Goldman, 1938, from Mexico (Vance and Duszynski, 1985). Oocyst and sporocyst size means are slightly larger than those of *E. subsimi* (25 × 19

and 14 × 7), but the oocysts of both lack a M and OR, possess a PG, and their sporocysts contain both SB and SSB and a membrane-bound, faceted SR. Rather than suggesting the form we saw can cross generic boundaries and call it *E. subsimi*, or assuming that it can't and call it a new species, we believe it best at this time to document its presence in *A. macrotis* and do neither.

A line drawing and photomicrographs of sporulated oocysts of *E. subsimi* already have been published (Vance and Duszynski, 1985), and photosyntypes are in the USNPC as #88512. We did not feel it necessary to submit phototypes of species with uncertain taxonomic status to the USNPC (*Eimeria* spp. 1–8).

Eimeria sp. 2

Host: *Alticola macrotis* (Radde, 1862), no common name recorded.

Other hosts: None, see *Remarks* under *Eimeria* sp. 1 and below.

Collected: V. B. Fedorov, K. E. Galbreath, 16 August 2002.

Locality: Asia: Russia, northern Siberia, Provideniya Oblast, Chukotka, 10 km SW of Yanrakynnot Village (64°52'N, 172°40'W).

Geographic distribution: Asia: Russia, Provideniya Oblast and Kolyma River Basin.

Prevalence: 4/24 (17%) (Table 1).

Material deposited: Symbiotype host, UAM 84096 (IF 7831).

Remarks: The oocysts we saw differed in size and other qualitative features from both *Eimeria* sp. 1 and *E. bassagensis* noted above. They closely resemble sporulated oocysts first described as "*Eimeria wenrichi* species A" by Saxe et al. (1960) (see Vance and Duszynski, 1985), from *Microtus pennsylvanicus* Ord, 1815, in Pennsylvania. Our oocyst (22 × 17 [19.5–23 × 16–19]) and sporocyst (12 × 7 [11–12 × 7–8]) sizes were slightly larger than those in the original description (19 × 14 and 10 × 6), but all unique qualitative features (e.g., small dark SB, faceted SR) were consistent with the original description. Because of rapid host speciation documented by Conroy and Cook (1999), this eimerian may be able to infect multiple host genera; however, as noted in reasons given for *Eimeria* sp. 1 (above), we believe it best at this time only to document its presence in *A. macrotis*.

A line drawing (Saxe et al., 1960) and photomicrograph of sporulated oocysts (Vance and Duszynski, 1985) of *E. wenrichi* have already been published, and photosyntypes are in the USNPC as #88517.

Eimeria cernae
Levine and Ivens, 1965
(Fig. 1)

Type host: *Myodes (Clethrionomys) glareolus* (Shreber, 1780), common red-backed vole.

Other hosts: *Myodes (Clethrionomys) rufocanus* Sundevall, 1846 (Russia), *Myodes (Clethrionomys) rutilus* (Pallas, 1779) (Alaska, Russia), both in this study (Table 1).

Collected (this study): A. M. Runck, 13 July 2001.

Type locality: Europe: Czech Republic.

Locality (this study): North America, U.S.A., Alaska, Noatak National Preserve, west bank of Situkuyok River (67°12.3'N, 163°9.7'W).

Geographic distribution: Europe: Czech Republic; Asia: Russia, Siberia; North America: U.S.A., Alaska.

Prevalence: Unknown in the type host; 1/53 (<2%) in *My. rufocanus* from Russia; 19/85 (22%) in *My. rutilus* from Russia; 181/699 (26%) in *My. rutilus* from Alaska (Table 1).

Sporulation: 2–4 days (Lewis and Ball, 1982).

Prepatent period: 6 days (Lewis and Ball, 1982).

Patent period: 4–6 days.

Site of infection: Epithelial cells of the cecum, colon, and rectum.

Endogenous stages: Lewis and Ball (1982) described 3 meront generations and the macro- and microgamonts and gametes from *My. glareolus*.

Material deposited: We deposited a photoneotype (see Duszynski, 1999) of a sporulated oocyst from *My. rutilus* (Alaska) in the USNPC as #99497. Symbiotype host (this study), UAM 56126 (AF 46214).

Remarks: The morphology of sporulated oocysts we studied from *My. rufocanus* and *My. rutilus* fits well with the composite description of *E. cernae* provided in the studies by Levine and Ivens (1990) and Lewis

and Ball (1982, 1983). Our oocyst and sporocyst size ranges ($17.5\text{--}22 \times 14\text{--}18.5$ and $9\text{--}11 \times 5\text{--}7$) overlap those ($13\text{--}23 \times 11\text{--}17$ and $9\text{--}15 \times 4\text{--}7$) given by Levine and Ivens (1990), but they did not give mean size measurements. Our quantitative data also are similar to the means provided by Lewis and Ball (1982) from *My. glareolus* in England (20×17 and 10×6 vs. 20×16 and 12×7). However, we observed 2 oocyst walls while all other descriptions of this coccidium only observed 1. Duszynski (2002) suggested that reports of oocysts having only 1 wall likely are in error and should be viewed cautiously when using older descriptions. *Eimeria cernae* was reported previously in *My. rutilus* from the Taimyr Peninsula (Arnastauskiene, 1977). Its recovery from *My. rutilus* in Alaska is a new geographic record, while its recovery from *My. rufocanus* in Russia is a new host record.

Line drawings (Levine and Ivens, 1965; Lewis and Ball, 1982, 1983) and a photomicrograph of a sporulated oocyst (Lewis and Ball, 1983) of *E. cernae* already have been published, and these can be compared to our Figure 1, which we have deposited as a photoneotype, as noted above.

Eimeria coahuiliensis
Vance and Duszynski, 1985

Type host: *Microtus mexicanus subsimus* Goldman, 1938, meadow vole.

Other hosts: *Mi. miurus* (Table 1).

Collected (this study): V. B. Fedorov, A. Fedorov, E. C. Waltari, B. Wagner, 25 July 2001.

Type locality: North America: Mexico, Coahuila, 14.8 km E of San Antonio de las Alazanas.

Locality (this study): North America: U.S.A., Alaska, Noatak National Preserve, Misheguk Mountain (68°28'N, 161°28'W).

Geographic distribution: North America: Mexico, Coahuila; U.S.A.: Alaska.

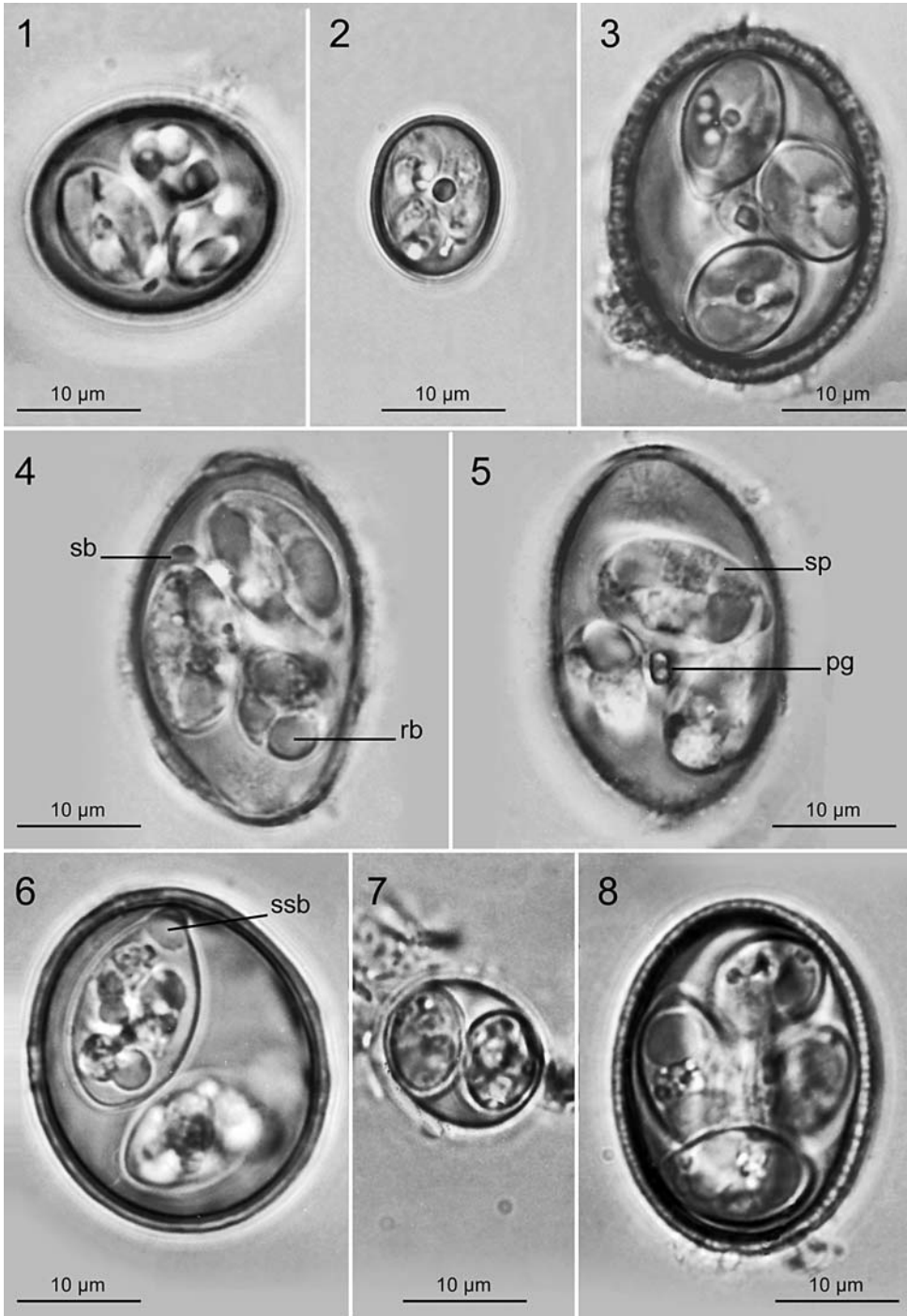
Prevalence: 2/22 (9%) *Mi. m. subsimus* (type host); 2/88 (2%) *Mi. miurus* from Alaska (Table 1).

Sporulation: Unknown.

Prepatent and patent periods: Unknown.

Site of infection: Unknown, oocysts recovered from feces.

Endogenous development: Unknown.



Figures 1–8. Photomicrographs of sporulated oocysts of *Eimeria* and *Isospora* spp. from rodents from Alaska and Russia. **1.** *Eimeria cernae* from *Myodes rutilus* from Alaska. **2.** *Eimeria marconii* from *My. rutilus* from Alaska. **3.** *Eimeria ochrogasteri* from *Microtus miurus* from Alaska. **4, 5.** *Eimeria rutilus* from *My. rutilus* from Alaska. **6.** *Isospora clethrionomydis* from *My. rutilus* from Alaska. **7.** *Isospora clethrionomydis* from *My. rutilus* from Alaska. **8.** *Eimeria synaptomys* from *S. borealis* from Alaska (sb, Stieda body at tip of sporocyst; rb, refractile in sporozoite; sp, sporozoite; pg, polar granule in oocyst; ssb, sub-Stieda body below sb).

Material deposited: A photosyntype is deposited in the USNPC as #88509. Symbiotype host (this study): UAM 56682 (AF 48384).

Remarks: Sporulated oocysts ($N = 25$) from *Mi. miurus* were 32.3×22.0 ($29\text{--}35 \times 19.5\text{--}26$), somewhat larger than those in the original description, 29.6×19.6 ($27\text{--}34 \times 18\text{--}22$), but the L/W ratios, 1.5, were the same, and all other features of the oocysts and sporocysts were the same. Line drawings and photomicrographs of oocysts are in the original description. This report represents new host and locality records for *E. coahuiliensis*.

Eimeria gallatii
Straneva and Kelly, 1979

Type host: *Myodes (C.) gapperi* (Vigors, 1830), red-backed vole.

Other hosts: *My. rufocanus* (Russia), *My. rutilus* (Alaska, Russia), Table 1.

Collected (this study): S. Kutz, A. A. Tsvetkova, A. A. Eddingsaas, M. McCain, 19 July 2001.

Type locality: North America: U.S.A., Pennsylvania, Indiana County, Brush Valley.

Locality (this study): North America: U.S.A., Alaska, Wrangell–St. Elias National Preserve, Nabesna, Chisana ($62^{\circ}3.9'N$, $142^{\circ}2.8'W$).

Geographic distribution: North America: U.S.A., Pennsylvania, Alaska; Asia: Russia, Siberia.

Prevalence: 1/15 (7%) in *My. gapperi* (type host); 4/53 (7.5%) in *My. rufocanus* from Russia; 1/85 (1%) in *My. rutilus* from Russia; 16/699 (2%) in *My. rutilus* from Alaska (Table 1).

Sporulation: 7 days at $22\text{--}25^{\circ}C$ (Straneva and Kelley, 1979).

Prepatent and patent periods: Unknown.

Site of infection: Unknown, oocysts recovered from fecal material.

Endogenous stages: Unknown.

Material deposited: Symbiotype host (this study), UAM 57764 (AF 54520).

Remarks: The oocysts and sporocysts we studied and measured resembled *E. gallatii*, first described from a red-backed vole in Pennsylvania, in most ways

except that those in the original description are slightly smaller (27.7×19.3 and 13.5×8.8) than the ones we measured ($N = 50$; 30.6×20.5 [$29\text{--}33 \times 19\text{--}23$] and 14.5×9.1 [$13\text{--}16 \times 8\text{--}10$]). All qualitative features are the same. The only uncertainty is whether or not a M really exists in the oocyst wall or whether the wall thins so much at the flattened pole of the oocyst to give the impression of a M when the osmotic concentration of the flotation fluid causes it to indent slightly. Unfortunately all our infected samples were discarded during a laboratory renovation after measurements were made, but before we were able to make photomicrographs of this species to deposit in the USNPC. Only a line drawing exists in the original description by Straneva and Kelley (1979).

Eimeria marconii
Straneva and Kelley, 1979

Fig. 2

Type host: *Myodes (C.) gapperi* (Vigors, 1830), southern red-backed vole.

Other hosts: *My. rutilus* (Russia, Alaska), Table 1.

Collected (this study): A. M. Runck, 21 July 2001.

Type locality: North America: U.S.A., Pennsylvania, Indiana County, Brush Valley.

Locality (this study): North America: U.S.A., Alaska, Cape Krusenstern National Monument, Red Dog Mine Road ($67^{\circ}44.97'N$, $163^{\circ}36.56'W$).

Geographic distribution: North America: U.S.A., Pennsylvania, Alaska; Asia, Russia, Siberia.

Prevalence: 1/15 (7%) in *My. gapperi* (type host); 1/85 (1%) in *My. rutilus* from Russia, and 40/699 (6%) in *My. rutilus* from Alaska (Table 1).

Sporulation: Unknown.

Prepatent and patent periods: Unknown.

Site of infection: Unknown, oocyst recovered from feces.

Endogenous stages: Unknown.

Material deposited: A photoneotype (see Duszynski, 1999) of a sporulated oocyst from *My. rutilus* (Alaska) is deposited in the USNPC as #99498. Symbiotype host (this study), UAM 55963 (AF 46245).

Remarks: The morphology of oocysts from *My. rutilus* in Alaska is similar to the description

provided by Straneva and Kelley (1979) of *E. marconii* described from *My. gapperi* in Pennsylvania. Oocyst and sporocyst sizes were similar to the original description (13×10 and 7×4 vs. 13×11 and 8×4). Unique qualitative features (e.g., single PG, small, dark SB, membrane-bound SR) were consistent with the original description. The recovery of *E. marconii* from *My. rutilus* in Russia and Alaska is a new host and geographic record for this parasite. A line drawing of the sporulated oocyst appears in the description by Straneva and Kelley (1979).

Eimeria ochrogasteri
Ballard, 1970
Fig. 3

Type host: *Microtus ochrogasteri* Wagner, 1842, prairie vole.

Other hosts: *Mi. mexicanus fulviventer*, *Mi. m. mexicanus*, *Mi. p. pennsylvanicus* (Vance and Duszynski, 1985), *Mi. miurus*, *Mi. oeconomus* (Table 1).

Collected (this study): V. B. Fedorov, 26 July 2001.

Type locality: North America: U.S.A., Colorado, Weld County, Black Hollow west of Ault.

Locality (this study): North America: U.S.A., Alaska, Noatak National Preserve, 8 km w of Copter Peak ($68^{\circ}28'N$, $161^{\circ}28'W$).

Geographic distribution: North America: U.S.A., Alaska, Colorado, Massachusetts; Mexico: Mexico, Oaxaca.

Prevalence: 1/71 (1%) *Mi. ochrogasteri* (type host); 3/26 (11.5%) in *Mi. m. fulviventer* from Oaxaca; 1/14 (14%) in *Mi. m. mexicanus* from Mexico; 1/11 (9%) in *Mi. p. pennsylvanicus* from Massachusetts (Vance and Duszynski, 1985); 1/88 (1%) in *Mi. miurus* and 2/405 (0.5%) in *Mi. oeconomus*, both from Alaska (Table 1).

Sporulation: Up to 9 days at 20°C.

Prepatent and patent periods: Unknown.

Site of infection: Unknown, oocysts recovered from feces.

Endogenous development: Unknown.

Material deposited: We deposited a photoneotype (see Duszynski, 1999) of a sporulated oocyst from *Mi. miurus* (Alaska) in the USNPC as #99499.

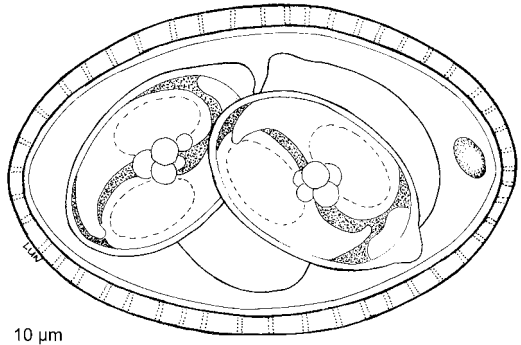


Figure 9. Composite line drawing of a sporulated oocyst of *Eimeria rutilus*.

Symbiotype host (this study), UAM 56698 (AF 48417).

Remarks: The morphology of oocysts from *Mi. oeconomus* and *Mi. miurus* in Alaska is similar to the description of *E. ochrogasteri* provided by Ballard (1970) from *Mi. ochrogasteri* in Colorado. Oocysts ($N = 11$) from *Mi. miurus* were 28.6×23.4 ($26\text{--}32 \times 22\text{--}25$), and those from *Mi. oeconomus* ($N = 25$) were 29.0×23.4 ($26\text{--}32 \times 22\text{--}25$), while those from *Mi. ochrogasteri* ($N = 100$) were 24.0×20.5 ($18\text{--}29 \times 16.5\text{--}24$); similarly, sporocysts ($N = 11$) from *Mi. miurus* (13.1×8.7 [$12\text{--}14 \times 8\text{--}9$]) and from *Mi. oeconomus* ($N = 25$: 13.2×8.6 [$12\text{--}14 \times 8\text{--}9$]) were slightly larger than those in the original description ($N = 50$: 12.3×8.2 [$11\text{--}14 \times 7\text{--}9$]). Ballard (1970) said that the average L/W ratio of his oocysts was 1.7, but this is incorrect; his data show a L/W of 1.2 ($24/20.5$), as do ours. The qualitative features (e.g., thick oocyst wall, variable OR, “capped” SB, and dispersed SR) we saw in oocysts from Alaskan microtines were similar to those in the original description. The recovery of *E. ochrogasteri* here represent new host records and a new geographic record. A line drawing of the sporulated oocyst appears in the description by Ballard (1970).

***Eimeria rutilus* n. sp.**
Figs. 4, 5, 9

Description of sporulated oocyst: Oocyst shape: ellipsoidal; number of walls: 2; wall thickness: ~ 1.75 ; wall characteristics: thinner at ends, outer slightly striated, 3/4 of total width, inner, membranous; $L \times W$ ($N = 50$): 30.6×20.5 ($27\text{--}33 \times 19\text{--}23$); L/W ratio: 1.5 (1.4–1.7); M: absent; OR: absent; PG:

present, single elliptical body, $\sim 3 \times 2$. Distinctive features of oocyst: large, ellipsoid shape with walls that become thinner at the 2 ends.

Description of sporocyst and sporozoites: Sporocyst shape ($N = 50$): ellipsoidal; $L \times W$: 14.5×9.1 ($13\text{--}16 \times 8\text{--}10$); L/W ratio: 1.6 (1.4–1.9); SB: present, dark, 3 times wider than high; SSB: present, colorless, twice as wide as SB, but just as high; PSB: absent; SR: present; SR characteristics: large refractile granules congregated in the center of the sporocyst; SZ: comma-shaped with one large, long RB, $\sim 7\text{--}3$. Distinctive features of sporocyst: presence of SB/SSB complex, SR with large granules, large RB.

Taxonomic summary

Type host: *Myodes (C.) rutilus* (Pallas, 1779), northern red-backed vole.

Other host: *My. rufocanus* (Russia), Table 1.

Collected: V. B. Fedorov, 15 July 2001.

Type locality: North America: U.S.A., Alaska, Noatak National Preserve, near Sidik Lake ($68^{\circ}08'N$, $158^{\circ}59'W$).

Geographic distribution: Asia: Russia, northeastern Siberia; North America: U.S.A., Alaska.

Prevalence: 1/85 (1%) in *My. rutilus* (Russia); 16/699 (2%) in *My. rutilus* (Alaska); 3/53 (6%) in *My. rufocanus* (Russia).

Sporulation: Unknown.

Prepatent and patent periods: Unknown.

Site of infection: Unknown, oocysts recovered from feces.

Endogenous stages: Unknown.

Material deposited: A photosyntype of a sporulated oocyst from the symbiotype host is deposited in the USNPC as #99500. Symbiotype host, UAM 56279 (AF 48238, male).

Remarks: Sporulated oocysts of *E. rutilus* resemble those of *Eimeria* species identified by Straneva and Kelley (1979) and Vance and Duszynski (1985): *E. gallatii* from *My. gapperi* and *E. coahuiliensis* from *Mi. mexicanus*, respectively. The oocyst and sporocyst sizes are similar (31×21 and 15×9 vs. 28×19 and 14×9 [Straneva and Kelley, 1979] vs. 30×20 and 14×9 [Vance and Duszynski, 1985]). All 3 species are characterized by having a SB and a SR

composed of loose granules aggregated in the center of the sporocyst, although at times the SR in *E. rutilus* appears to be membrane bound. There are 2 structures that differentiate these 3 species. Both *E. gallatii* and *E. coahuiliensis* have a M, while *E. rutilus* does not. We also identified a SSB in *E. rutilus*, while the other 2 lack this structure.

Etymology: The *nomen triviale* is derived from the specific name of the type host, *rutilus*.

Eimeria saxei

Vance and Duszynski, 1985

Type host: *Microtus pennsylvanicus* Ord, 1815, meadow vole.

Other hosts: *Mi. californicus sactidiegi*, *Mi. longicaudus*, *Mi. mexicanus fulviventer*, *Mi. m. mexicanus*, *Mi. m. subsimus*, *Mi. oregoni oregoni* (Vance and Duszynski, 1985), *Mi. miurus*, *Mi. oeconomus* (Table 1).

Collected (this study): S. O. MacDonald, A. N. Lazutkin, A. J. Lynch, S. Runck, K. Gamblin, L. B. Barrelli, N. MacDonald, N. I. MacDonald, 9 August, 2001.

Type locality: North America: U.S.A., Pennsylvania.

Locality (this study): North America: U.S.A., Alaska, Yukon–Charley Rivers National Preserve, Charley River ($65^{\circ}21.3'N$, $142^{\circ}57.5'W$).

Geographic distribution: North America: U.S.A., Alaska, California, Pennsylvania, Washington; Mexico: Coahuila, Mexico, Oaxaca; Asia: Russia, Siberia.

Prevalence: 1/1 (100%) in *Mi. pennsylvanicus* from Pennsylvania (type host); 1/3 (33%) in *Mi. c. sactidiegi* from California; 3/29 (10%) in *Mi. longicaudus* from Alaska (Table 1); 1/26 (4%) in *Mi. m. fulviventer* from Oaxaca; 1/14 (7%) in *Mi. m. mexicanus* from Mexico; 3/48 (6%) in *Mi. m. subsimus* from Coahuila; 1/4 (25%) in *Mi. o. oregoni* from Washington; 2/88 (2%) in *Mi. miurus* from Alaska (Table 1); 2/48 (4%) in *Mi. oeconomus* from Siberia (Table 1); 18/405 (4%) in *Mi. oeconomus* from Alaska (Table 1).

Sporulation: 2–3 days at $24\text{--}27^{\circ}C$.

Prepatent and patent periods: Unknown.

Site of infection: Unknown, oocysts recovered from feces.

Endogenous development: Unknown.

Material deposited: A photosyntype is deposited in the USNPC as #88510. Symbiotype host (this study), UAM 57278 (AF 52605).

Remarks: Originally described as “*E. wenrichi* species B” by Saxe et al. (1960), Vance and Duszynski (1985) renamed this species as *E. saxei*. The morphology of oocysts from *Microtus* species in Alaska and Russia is similar to both the original description for *E. saxei* from *Mi. pennsylvanicus* and the redescription by Vance and Duszynski (1985) from *Mi. mexicanus*, *Mi. oregoni*, and *Mi. californicus*. Oocysts we measured ($N = 25$) from *Mi. longicaudus* in Alaska were moderately larger (15×12 [$12-17.5 \times 10-13$]) than those measured by Saxe et al. (1960) (13×10 [$11-15 \times 8-11$]) and by Vance and Duszynski (1985) (13×11 [$11-14 \times 10-12$]), but all other quantitative and qualitative features were the same. Likewise, those we measured from *Mi. pennsylvanicus* in Alaska were also slightly larger (14×10 [$13-14 \times 9-11$]). This study represents new host and/or locality records for *Mi. longicaudus*, *Mi. miurus*, and *Mi. oeconomus* in Alaska and *Mi. oeconomus* in Siberia. A line drawing (Saxe et al., 1960) and photomicrograph of sporulated oocysts (Vance and Duszynski, 1985) of *E. saxei* have already been published.

Eimeria wenrichi
Saxe, Levine, and Ivens, 1960

Type host: *Microtus pennsylvanicus* Ord, 1815, meadow vole.

Other hosts: *Mi. mexicanus fulviventer*, *Mi. m. mexicanus*, *Mi. m. mogollonensis*, *Mi. m. subsimus*, *Mi. montanus arizonensis*, *Mi. o. oregoni*, *Mi. p. pennsylvanicus*, *Mi. oeconomus*, *Mi. longicaudus*, *Mi. miurus*, *Mi. xanthognathus*.

Collected (this study): H. Henttonen, J. Niemimeä, K. Gamblin, L. B. Barrelli, 26 July 2001.

Type locality: North America: U.S.A., Pennsylvania.

Locality (this study): North America: U.S.A., Alaska, Yukon-Charley Rivers National Preserve, Eagle ($64^{\circ}58'N$, $143^{\circ}3'W$).

Geographic distribution: North America: U.S.A., Alaska, Arizona, Massachusetts, Washington; Mexico, Coahuila, Oaxaca, Veracruz; Asia: Russia, Siberia.

Prevalence: 1/1 (100%) in *Mi. pennsylvanicus* (type species); 1/26 (4%) in *Mi. mexicanus fulviventer*

from Oaxaca, 4/15 (3%) in *Mi. m. mexicanus* from Veracruz, 1/1 (100%) in *Mi. m. mogollonensis* from Arizona, 8/48 (8%) in *Mi. m. subsimus* from Coahuila, 4/8 (50%) in *Mi. montanus arizonensis* from Arizona, 2/4 (50%) in *Mi. o. oregoni* from Washington, and 5/11 (45%) in *Mi. p. pennsylvanicus* from Massachusetts (all from Vance and Duszynski, 1985); 3/29 (10%) in *Mi. longicaudus*, 39/88 (44%) in *Mi. miurus*, 265/405 (65%) in *Mi. oeconomus*, 123/159 (77%) in *Mi. pennsylvanicus*, and 9/52 (17%) in *Mi. xanthognathus*, all from Alaska (Table 1); 29/48 (60%) in *Mi. oeconomus* from Russia (Table 1).

Sporulation: 2–3 days at 24–27°C.

Prepatent and patent periods: Unknown.

Site of infection: Unknown, oocysts recovered from feces.

Endogenous development: Unknown.

Material deposited: A photosyntype is deposited in the USNPC as #88517. Symbiotype host (this study), UAM 58313 (AF 49611).

Remarks: In the original description by Saxe et al. (1960) they referred to this species as “*E. wenrichi* species A.” The morphology of oocysts from *Microtus* species in Alaska and Russia is similar to those reported in the original description for *E. wenrichi* from *Mi. pennsylvanicus*. Oocysts we measured ($N = 25$) from *Mi. longicaudus* in Alaska were slightly larger (20.8×15.8 [$19-23 \times 14-16.5$]) than those measured by Saxe et al. (1960) (18.9×14.3 [$16-22 \times 12-16$]), but all other quantitative and qualitative features were the same. This study represents new host and/or locality records for *Mi. longicaudus* in Alaska. The oocysts we saw in the 5 *Microtus* species in this study (Table 1) resemble the form we call *E. sp. 2* from *A. macrotis* (from Russia), above. A line drawing (Saxe et al., 1960) and photomicrograph of sporulated oocysts (Vance and Duszynski, 1985) of *E. wenrichi* have already been published.

Isospora clethrionomydis
Golemansky and Yankova, 1973
Fig. 6

Type host: *Myodes (C.) glareolus* (Shreber, 1780), bank vole.

Other hosts: *My. rufocanus* (Russia), *My. rutilus* (Alaska), Table 1.

Collected (this study): H. Henttonen, 25 July 2001.

Type locality: Europe: Bulgaria.

Locality (this study): North America: U.S.A., Alaska, Yukon–Charley Rivers National Preserve, near base of Mount Sorenson, north of Seventy-Mile headwaters (64°58'N, 143°3'W).

Geographic distribution: Europe: Bulgaria; North America: U.S.A., Alaska; Asia: Russia, Siberia.

Prevalence: 39/109 (36%) *My. glareolus* (type host) from 5 localities in Bulgaria were infected with coccidia, but it was not stated how many of the 39 were infected with *I. clethrionomydis*; 1/53 (<2%) *My. rufocanus* (Russia); 4/699 (<1%) *My. rutilus* (Alaska) (Table 1).

Sporulation: Unknown.

Prepatent and patent periods: Unknown.

Site of infection: Small intestine.

Endogenous stages: Unknown.

Material deposited: A photoneotype (see Duszynski, 1999) of a sporulated oocyst from *My. rutilus* (Alaska) is deposited in the USNPC as #99502. Symbiotype host (this study), UAM 58263 (AF 49559).

Remarks: The morphology of sporulated oocysts we studied from *My. rutilus* in Alaska is similar to the description by Golemansky and Yankova (1973) of *I. clethrionomydis* from *My. glareolus* in Europe. Our sporulated oocysts were slightly larger than those in the original description (27 × 26 vs. 25 × 25), but this difference seems insignificant. Golemansky and Yankova (1973) described the sporocysts they saw as “elongated,” measuring 21–23 × 11–12. However, both their line drawing and photomicrograph of a sporulated oocyst show sporocysts that are broadly ovoidal. The sporocysts we measured were 17 × 11 (15–18 × 10–11), L/W ratio: 1.5, and seem more parsimonious with their original line drawing and (poor quality) photomicrograph. In the original they describe a “light-refracting Stieda body,” and the drawing shows a SB much wider than high. The SB in the sporocysts we studied was approximately 4 times wider than high. Additionally we report a SSB, which Golemansky and Yankova (1973) did not mention, although there is a hint of this structure in their line drawing. The SSB we observed was clearly visible and slightly wider than the SB. The recovery of *I. clethrionomydis* from *My. rutilus* in Alaska and

My. rufocanus from Russia are new host and geographic records for this parasite.

Isospora clethrionomysis
Arnastauskiene and Maldzhiunaite, 1981
Fig. 7

Type host: *Myodes* (*C.*) *glareolus* (Shreber, 1780), bank vole.

Other hosts: *My. rutilus* (Alaska), Table 1.

Collected (this study): V. B. Fedorov, 2 August 2001.

Type locality: Eastern Europe: Lithuania.

Locality (this study): North America: U.S.A., Alaska, Noatak National Preserve, near Aniralik Lake (68°12'N, 159°49'W).

Geographic distribution: Eastern Europe: Lithuania; North America: U.S.A., Alaska.

Prevalence: 5/946 (<1%) in *My. glareolus* (type host); 4/699 (<1%) in *My. rutilus* from Alaska (this study).

Sporulation: Unknown.

Prepatent and patent periods: Unknown.

Site of infection: Unknown, oocysts recovered from feces.

Endogenous development: Unknown.

Material deposited: A photoneotype (see Duszynski, 1999) of a sporulated oocyst from *My. rutilus* (Alaska) is deposited in the USNPC as #99503. Symbiotype host (this study), UAM 56434 (AF 48526).

Remarks: The morphology of oocysts from *My. rutilus* in Alaska is similar to the original description by Arnastauskiene and Maldzhiunaite (1981) of *I. clethrionomysis* from *My. glareolus* in 6 districts in Lithuania. In their description only oocyst size ranges were given (9–11 × 7–10). The oocysts we measured ($N = 25$) (10 × 10 [9–12.5 × 9–12]) are slightly larger. Sporocyst size was not reported in the original description, but the line drawing shows them to be spindle shaped (slightly pointed at both ends) and almost as long as the oocyst is wide. The sporocysts we measured ($N = 25$) are of comparable size (8 × 5 [7.5–8 × 4.5–5]), but we also saw that they possess a small, but distinct, Stieda body not reported by Arnastauskiene and Maldzhiunaite (1981). Finally, the sporozoites we saw were sausage-shaped rather

than the pointed, spindle-shaped ones described and drawn by Arnastauskiene and Maldzhiunaite (1981). Other qualitative features (e.g., small, membrane-bound SR) are similar to the original. The recovery of *I. clethrionomysis* from *My. rutilus* is a new host and geographic record in Alaska.

Lemmings *Eimeria* sp. 3

Host: *Lemmus trimucronatus* Richardson, 1825, arctic or brown lemming.

Other hosts: *Lemmus lemmus* L., 1758 (Table 1).

Collected: A. M. Runck, N. E. Dokuchaev, R. Foster, K. Fisher, E. Tomasik, 28 July 2001.

Locality: North America: U.S.A., Alaska, Bering Land Bridge National Preserve, Bendeleben (65°23.4'N, 163°15.9'W).

Geographic distribution: Asia: Russia, Siberia; North America: U.S.A., Alaska.

Prevalence: 1/11 (9%) in *L. lemmus*; 2/72 (3%) in *L. trimucronatus* (Table 1).

Material deposited: Symbiotype host, UAM 55562 (AF 46286).

Remarks: There are no coccidia yet described from this host species. Arnastauskie (1980) described oocysts of *Eimeria chatangae* from *Lemmus sibiricus* that were similar in size (13–14 × 11–13) to those we found in this study (12 × 11 [10–13 × 9–11.5]), but his sporocysts (8–9 × 5–6) were larger than those we measured (6 × 4 [4.5–6 × 3–4]), and there are qualitative differences between the species. Oocysts of *E. chatangae* lack a PG while the sporocysts are drawn as pointed, long structures, and do not have a SB. The sporulated oocysts we measured from *L. lemmus* were similar to, but smaller than, those of *E. saxei* first described from *Mi. pennsylvanicus* in Pennsylvania and later found in several other *Microtus* species/subspecies in 2 states in the U.S.A. and 3 states in Mexico (Vance and Duszynski, 1985). We also found similar looking sporulated oocysts in *Lemmus trimucronatus* (14 × 13 [12–15 × 12–14] and 7 × 4.5, [6–8 × 4–5]), which had slightly larger oocysts, but not sporocysts, than in the first description of *E. saxei*. In both *Lemmus* species, only the oocysts varied in size while the sporocysts and all structural and qualitative features were the same as in *E. saxei*. However, for reasons stated earlier and in our Discussion, we believe it best at this time to

document its presence in *L. lemmus* and *L. trimucronatus* but not name it.

Eimeria sp. 4

Host: *Lemmus trimucronatus* Richardson, 1825, arctic or brown lemming.

Other hosts: None.

Collected: V. B. Fedorov, A. Fedorov, B. Wagner, E. C. Waltari, 19 July 2001.

Locality: North America: U.S.A., Alaska, Noatak National Preserve, Ambler River, Laluch Creek Upland (67°39'N, 158°11'W).

Geographic distribution: North America: U.S.A., Alaska.

Prevalence: 5/72 (7%) (Table 1).

Material deposited: Symbiotype host, UAM 56355 (AF 48310).

Remarks: The oocysts we measured were most similar to those *E. subsimi* first described from *Mi. mexicanus*; they are also quite similar to those we call *E. sp. 1* from *A. macrotis*. However, as already noted, we believe it best only to document its presence in *L. trimucronatus*. A line drawing and photomicrographs of sporulated oocysts of *E. subsimi* already have been published (Vance and Duszynski, 1985), and photo-syntypes are in the USNPC as #88512.

Eimeria sp. 5

Host: *Lemmus trimucronatus* Richardson, 1825, arctic or brown lemming.

Other hosts: None.

Collected: V. B. Fedorov, A. Fedorov, B. Wagner, E. C. Waltari, 20 July 2001.

Locality: North America: U.S.A.: Alaska, Noatak National Preserve, Ambler River, Kaluich Creek Upland (67°39'N, 158°11'W).

Geographic distribution: North America: U.S.A., Alaska.

Prevalence: 8/72 (11%) (Table 1).

Material deposited: Symbiotype host, UAM 56378 (AF 48335).

Remarks: The morphology of sporulated oocysts from *L. trimucronatus* in Alaska is very similar to the

description provided by Upton and Pitts (1993) of *Eimeria synaptomys* described from *Synaptomys cooperi* in Missouri with oocyst (27×22 [$23\text{--}31 \times 19\text{--}24$]) and sporocyst (12×8 [$10.5\text{--}13.5 \times 7\text{--}9$]) sizes, as well as all qualitative features nearly identical to those in the original description except that their OR was a single globule ~ 6 , while ours was smaller, ~ 3 . Both quantitative and qualitative features of the sporulated oocysts of *Eimeria* sp. 5 and of *Eimeria synaptomys* are similar to those of *E. ochrogasteri* from *Microtus* species (Ballard, 1970; Vance and Duszynski, 1985) with only subtle differences. Given that *Synaptomys* and *Lemmus* are sister taxa (Conroy and Cook, 1999; Jarrell and Fredga, 1993), the most parsimonious conclusion might be that these oocysts in *L. trimucronatus* also represent *E. synaptomys*. It's known from cross-infection experiments, for example, that sister taxa of other rodents, for instance, *Peromyscus* and *Reithrodontomys* species, are capable of sharing coccidia species (Upton et al., 1992; Hnida and Duszynski, 1999a). However, we believe it best only to document its presence in *L. trimucronatus* until molecular evidence can clear up this picture to set the record straight (see *Remarks*, *Eimeria* sp. 1, and *Discussion*, below). A line drawing and photomicrographs of sporulated oocysts of *E. synaptomys* already have been published (Upton and Pitts, 1993).

***Eimeria* sp. 6**

Host: *Synaptomys borealis* Richardson, 1828, bog lemming.

Other hosts: None, see *Remarks*, below.

Collected: H. Henttonen, J. Niemimeä, A. V. Goropashnaya, J. Burch, 8 August 2001.

Locality: North America: U.S.A., Alaska, Yukon–Charley Rivers National Preserve, across Yukon River from Glenn Creek cabin ($65^{\circ}18'N$, $142^{\circ}14'W$).

Geographic distribution: North America: U.S.A., Alaska.

Prevalence: 4/187 (2%) (Table 1).

Material deposited: Symbiotype host, UAM 60299 (AF 49479).

Remarks: The oocysts we measured (13.5×12 [$11\text{--}16 \times 10\text{--}13.5$]) most closely resemble those of *E. saxei* described from *Microtus* species in Pennsylvania, California, and Washington, U.S.A., and from Oaxaca, Mexico, and Coahuila, Mexico (Saxe et al., 1960;

Vance and Duszynski, 1985); they also resemble the oocysts we found in *Mi. longicaudus* and *Mi. pennsylvanicus* and *Eimeria* sp. 3 from *L. lemmus* in this study (Table 1). It may be possible that this species is actually *E. saxei*, capable of infecting multiple host genera that are closely related, but we believe it best at this time only to document its presence in *S. borealis* and wait until molecular data can provide evidence that this is either one or multiple species.

***Eimeria* sp. 7**

Host: *Synaptomys borealis* Richardson, 1828, bog lemming.

Other hosts: None, see *Remarks*, below.

Collected: S. O. MacDonald, A. N. Lazutkin, A. J. Lynch, S. Runck, 24 July 2001.

Locality: North America: U.S.A., Alaska, Yukon–Charley Rivers National Preserve, Charley River, 12.5 km east-northeast of Kathul Mountain ($65^{\circ}22'N$, $142^{\circ}01'W$).

Geographic distribution: North America: U.S.A., Alaska.

Prevalence: 17/187 (9%) (Table 1).

Material deposited: Symbiotype host, UAM 57419 (AF 52076).

Remarks: Oocysts we measured (30×20 [$28\text{--}32 \times 17.5\text{--}22$]) seemed most similar to those of *E. subsimi* described from *Mi. mexicanus* from Mexico (Vance and Duszynski, 1985) and were quite similar to *Eimeria* sp. 1 from *A. macrotis* from Russia (this study) and *Eimeria* sp. 4 from *L. trimucronatus* from Alaska (this study). Because of rapid host speciation (Conroy and Cook, 1999), we could suggest that this species may be able to infect multiple host genera; however, we believe it best only to document its presence in *S. borealis* and wait until molecular data can provide evidence that this is either one or multiple species. A line drawing and photomicrographs of sporulated oocysts of *E. subsimi* already have been published (Vance and Duszynski, 1985), and photosyntypes are in the USNPC as #88512.

***Eimeria synaptomys* Upton and Pitts, 1993 Fig. 8**

Type host: *Synaptomys cooperi* Baird, 1858, bog lemming.

Other hosts: *Synaptomys borealis* Richardson, 1828.

Collected (this study): H. Henttonen, J. Niemimeä, K. Gamblin, L. B. Barrelli, 27 July 2001.

Type locality: North America: U.S.A., Missouri, St. Charles County, 3.4 km West Weldon Spring.

Locality (this study): North America: U.S.A., Alaska, Yukon–Charley Rivers National Preserve, at base of Mount Sorenson, north of Seventy-Mile headwaters (64°58'N, 143°3'W).

Geographic distribution: North America: U.S.A., Alaska, Missouri.

Prevalence: 2/22 (9%) in *S. cooperi* (type host); 9/187 (5%) in *S. borealis* (Table 1).

Sporulation: Unknown.

Prepatent and patent periods: Unknown.

Site of infection: Unknown, oocysts recovered from feces.

Endogenous development: Unknown.

Material deposited: A photoneotype (see Duszynski, 1999) of a sporulated oocyst is deposited in the USNPC as #99501. Symbiotype host (this study), UAM 58343 (AF 49647).

Remarks: The oocysts and sporocysts we measured were similar to those seen in the original description by Upton and Pitts (1993). These oocysts also were similar to those of *Eimeria* sp. 5 that we found in *L. trimucronatus*. A line drawing and photomicrographs of sporulated oocysts of *E. synaptomys* already have been published (Upton and Pitts, 1993).

***Eimeria* sp. 8**

Host: *Synaptomys borealis* Richardson, 1828, bog lemming.

Other hosts: None, see *Remarks*, below.

Collected: S. Kutz, A. A. Tsvetkova, A. A. Eddingsaas, M. McCain, 4 August 2001.

Locality: North America: U.S.A., Alaska, Wrangell-St. Elias National Park and Preserve, Harry's Gulch (61°3.8'N, 143°54'W).

Geographic distribution: North America: U.S.A., Alaska.

Prevalence: 4/187 (2%) (Table 1).

Material deposited: Symbiotype host, UAM 56004 (AF 53552).

Remarks: The oocysts (21 × 15 [20–24 × 13–16]) and sporocysts (11 × 6 [10–12 × 5–7.5]) we measured were similar to those seen in the original description of *E. wenrichi* (species "A") from 1 *Mi. pennsylvanicus* collected in Pennsylvania, U.S.A. (Saxe et al., 1960; Vance and Duszynski, 1985). They also are similar to oocysts we identify as *E. wenrichi* from *Microtus* species in this study and to *Eimeria* sp. 2 in *A. macrotis* from Siberia in this study. As stated, we believe it best at this time to record its presence in *S. borealis* and wait until further study can provide evidence that this is either one or multiple species. A line drawing (Saxe et al., 1960) and photomicrograph of sporulated oocysts (Vance and Duszynski, 1985) of *E. wenrichi* have already been published, and photosyntypes are in the USNPC as #88517.

DISCUSSION

This study compares the coccidia of closely related rodent taxa separated by a geographic barrier, the Bering Strait. Rodentia is the most speciose order of mammals and occupies a wide variety of habitats. Because of their virtual ubiquity on land, we know more about the coccidia of rodents than we do about those of any other mammalian order (Duszynski and Upton, 2001). Nonetheless, <15% of the 2,277 rodent species (Wilson and Reeder, 2005) have been examined for coccidia.

In spite of the paucity of information, the picture that emerges is that the *degree* of host specificity seems to vary from host group to host group. For example, in murid rodents of the subfamily Sigmodontinae, Reduker et al. (1987) recovered a similar eimerian morphotype, "type A," across multiple host genera (*Peromyscus*, *Neotoma*, and later *Baiomys*, *Onychomys*, and *Reithrodontomys* [Upton et al., 1992]). However, when the "type A" morphotypes were subjected to cross-infection experiments (Hnida and Duszynski, 1999a, Upton et al., 1992) and DNA sequencing (Hnida and Duszynski, 1999b), each host was found to have a different species of *Eimeria*.

Conversely, in sciurids (squirrels and their relatives), some experimental evidence (Todd and Hammond, 1968a, b; Thomas and Stanton, 1994) suggests eimerians may easily switch between host species in closely related genera, and an eimerian from the chinchilla is known to infect 7 genera of mammals in 2 families (DeVos, 1970). Work by Hafner (1984) suggested that sciurids differentiated in

the Pleistocene and that many may be able to harbor similar parasites. Based on these efforts, Wilber et al. (1998) decided to use the “morphological species concept” to revise and summarize the coccidia of the Marmotini, the largest tribe within the Sciuridae. Recently this approach also has been used by others (e.g., Seville et al., 2005).

Phylogenetic work by Conroy and Cook (1999) suggested rapid pulses of speciation in arvicoline rodents due to a series of unresolved polytomies found at several periods in the arvicoline radiation. Many of the same rodent taxa in their analyses were surveyed in this study, and the coccidia we identified by their sporulated oocysts are consistent with a rapid speciation process in these rodents because similar oocyst morphotypes were found across both genera and species in geographically sympatric host taxa. Thus, the question becomes what should we do with the oocyst morphotypes we find in these genera and species, most of which closely resemble “species” described from other host genera. On one hand, our results may suggest that these high-latitude arvicoline rodents have *Eimeria* that are able to cross generic boundaries. Hoberg et al. (2003) suggested that drivers for speciation for the rodents collected in this study might be different than those for other rodents, which, in turn, might influence the speciation patterns among parasitic protists and other parasites. On the other hand, Tenter et al. (2002) said that without multiple lines of evidence, for instance, mensural and molecular data, coccidia that are morphologically indistinguishable and identified from closely related hosts should not be described as new species. Thus, we believe the most prudent choice is to name new species (e.g., *E. rutilus*) only when we feel the morphological mensural data are sufficiently different from all other eimerians described from related host species to warrant new taxon status. In all other instances, such as in the other morphotypes we saw (*Eimeria* spp. 1–8), we only mention their presence and point out the similarities to their closest structural relatives.

Myodes is a Holarctic genus composed of 9 Palearctic, 2 Nearctic, and 1 Holarctic species (Wilson and Reeder, 2005). Prior to this study, 3 species, *My. gapperi*, *My. glareolus*, and *My. rutilus*, had been surveyed for coccidia (Levine and Ivens, 1990). Levine and Ivens (1990) recognized *E. cernae* and *E. rysavyi* from the bank vole, *My. glareolus*, in Czechoslovakia, and Arnastauskiene (1977) reported *E. cernae* and described *Eimeria schiwicki* from *My. rutilus* in Russia. In North America Straneva and Kelley (1979) described and named 4 coccidia from

the southern red-backed vole, *My. gapperi*: *E. clethrionomyis*, *E. gallattii*, *E. marconii*, and *E. pileata*. Here we found coccidia from *My. rufocanus* and *My. rutilus* that have been previously reported from Old World *My. glareolus* and New World *My. gapperi*. We also discovered a new coccidium, *E. rutilus*, in *My. rufocanus* and *My. rutilus*. A recent phylogenetic tree created with the cytochrome b gene suggests that *My. gapperi* is more closely related to *My. glareolus* than to *My. rutilus*, and all 3 are more closely related to one another than they are to *My. rufocanus* (Cook et al., 2004). It is interesting to note that 2 coccidia, *E. clethrionomyis* and *E. pileata*, identified by Straneva and Kelley (1979) from *My. gapperi*, were not recovered in this study.

The genus *Microtus* has 65 species; 44 are Palearctic, 20 Nearctic, and 1 (*Mi. oeconomus*) is Holarctic in distribution (Wilson and Reeder, 2005). To date, 36 species of *Eimeria* and 3 species of *Isospora* have been described from *Microtus*. The majority of these species (29 *Eimeria*, 1 *Isospora*) are described from Eurasian *Microtus* spp., while all the coccidia reported from the 5 species of *Microtus* in this study were similar to coccidia reported from North America. Conroy and Cook (2000) suggested endemic species of North American *Microtus* are monophyletic and that *Mi. oeconomus* is a recent immigrant (>55,000 years) to North America via the BLB (Galbreath and Cook, 2004). Of the 5 *Microtus* spp. we examined, only *Mi. pennsylvanicus* had been surveyed previously for coccidia. We expected to find coccidia in the 4 North American *Microtus* spp. (Table 1) that were similar to those in other *Microtus* spp. in North America (see Vance and Duszynski, 1985; Ballard, 1970) and coccidia in *Mi. oeconomus* (a Eurasian species) that would be similar to coccidia described from Eurasian *Microtus* spp. Instead, we found *Mi. oeconomus* on both continents infected by the same species of coccidia that infect the North American endemic species (Table 1).

Other genera included in this study have had few coccidia reported from them: *Alticola*, *E. bassagensis*; *Dicrostonyx*, *E. dicrostonicis*; *Lemmus*, *E. chatangae* and *E. nativa*; and *Synaptomys*, *E. synaptomys* (Levine and Ivens, 1990). Our results suggest that 3 of these 4 host genera were infected with coccidia that discharged oocysts quite similar in their morphology to those seen previously to infect the genus *Microtus*.

The data from arvicoline rodents may be interpreted at least 2 ways. It is possible that several of their eimeriid coccidia (*E. saxei*, *E. subsimi*, *E. synaptomys*, *E. wenrichi*) are temporally, ecologically,

and genetically the same species of generalist coccidia found across multiple host taxa. For example, oocysts of *E. saxei* from *Microtus* are similar to the oocysts of *Eimeria* spp. 3 and 6 from *Lemmus* and *Synaptomys*, respectively. Similarly oocysts of *E. subsimi* (*Microtus*) resemble *Eimeria* spp. 1 (*Alticola*), 4 (*Lemmus*), and 7 (*Synaptomys*); those of *E. synaptomys* (*Synaptomys*) resemble *Eimeria* sp. 5 (*Lemmus*); and those of *E. wenrichi* (*Microtus*) resemble those of *Eimeria* sp. 2 (*Alticola*) and 8 (*Synaptomys*). This interpretation suggests there have been insufficient selective pressure and/or time for these coccidia to speciate. While each of these hosts is not sympatric with all of the other hosts and there is fine-scale niche partitioning (see Getz [1985] for *Microtus* and Linzey [1984] for data on *Synaptomys* and *Microtus*), their combined ranges are contiguous (see trap line data sheets for Beringia Co-evolution Project stored at the University of Alaska–Fairbanks and the University of New Mexico). It is possible, then, that there is flow of coccidia across host genera. The other interpretation is that morphologically similar oocysts are indeed different species in different hosts. The oocysts identified here with similar morphology in multiple host genera may instead each be different species representing distinct lineages that reflect host relationships. The idea of cryptic speciation (i.e., speciation without obvious morphological divergence) has been suggested as characterizing the parasites of Arctic arvicoline rodents because long-term climatic oscillations have caused repeated fragmentation and isolation as well as subsequent divergence, especially in lemming and vole populations (Fedorov et al., 1999; Fedorov and Goropashnaya, 1999; Hoberg et al., 1999). In fact, Haukisalmi et al. (2004) documented multiple species of cestodes hidden by their apparent morphological similarity to *Paranoplocephala omphalodes*, a Holarctic parasite of *Microtus* voles in Alaska, Russia, and elsewhere, and Baverstock et al. (1985) showed that another cestode thought to be a host-generalist parasite of wombats, wallabies, and kangaroos includes at least 12 biological species that are strictly host specific when isozyme electrophoretic data were examined. DNA sequence data in the form of *Eimeria* gene phylogenies or population-level microsatellite work on *Eimeria* spp. from these host groups may provide us with answers to these enigmatic observations.

Since the advent of phylogenetic reconstructions, innumerable schemes have been proposed for various families and genera within the order Rodentia. Homoplasy among morphological characters has

made it difficult to discern a consistent interpretation of relationships among the 29 families. Recently Debry (2003) combined data from multiple genes to analyze familial relationships. His (2003) work suggests there are 2 distinct clades. One contains Myodonta (Muroidea plus sister taxa Dipodidae), Geomyoidea, Pedetidae, and Castoridae, while the other clade is composed of Sciuroidea, Gliridae, and Hystricognathi.

Here, based on morphology of the sporulated oocysts, one interpretation suggests that coccidia in arvicoline rodents (family Cricetidae, superfamily Muroidea) may be able to infect multiple genera much like the coccidia in sciurids. Based on Debry's (2003) work, these 2 groups of rodent hosts aren't closely related. However, the fossil record and molecular data support a rapid arvicoline radiation (Conroy and Cook, 1999), much like the sciurids (Hafner, 1984). This speciation alone may be enough to explain the similarity in the ability of some *Eimeria* spp. to infect multiple host genera. If this is the case, then other groups of rodents (and possibly other mammals) that have undergone rapid speciation also may have multiple genera infected by similar coccidia.

Additionally, host ecology certainly affects transmittance of coccidia between individuals sharing an environment. Many sciurid rodents are highly social and dwell in subterranean tunnels (Murie and Michener, 1984). Harsh weather conditions that may affect coccidia transmission may be mitigated by the microclimates created by burrows (Thomas et al., 1995). Also, sympatry of many ground-dwelling sciurids would help to create and maintain a guild of hosts sharing the same coccidia (Shults et al., 1990). In general, voles and lemmings aren't as social as sciurids; however, they utilize runways and create shallow burrows that multiple vole and lemming species use (Gromov and Polyakov, 1992). Hertel and Duszynski (1987) suggested that, at least for shrews, similar habitat conditions (runways and burrows) might be conducive to oocyst transmission. The similarity of habitat usage creates a flow of oocysts across multiple host species that might inhibit parasite speciation.

Of the 1,925 cricetid specimens, 68 (3.5%) harbored multiple infections with coccidia; 65 with 2 species and 3 were infected with 3 coccidia simultaneously. Of those, 23 were double infections and 2 were triple infections of coccidia in *My. rutilus*. The majority of double infections (17) were composed of *E. cernae* with *E. marconii*. The rest (6) were *E. cernae* with *E. rutilus*. One triple infection

was composed of *E. cernae*, *E. marconii*, and *E. rutilus*, while the other had *I. clethrionomysis* rather than *E. marconii*. There were 42 *Microtus* spp. that were host to infections of 2 coccidia: *Mi. longicaudus*, 1; *Mi. miurus*, 1; *Mi. pennsylvanicus*, 16; and *Mi. oeconomus*, 24. Both *Mi. longicaudus* and *Mi. miurus* had 1 infection of *E. saxei* with *E. wenrichi*. In *Mi. oeconomus*, infections were composed of *E. wenrichi* with *E. saxei* (13), *E. wenrichi* with *E. subsimi* (8), and *E. wenrichi* with *E. ochrogasteri* (3). In *Mi. pennsylvanicus* there were equal numbers of infections of *E. wenrichi* with *E. saxei* (8) and *E. wenrichi* with *E. subsimi* (8). Only 1 specimen of *Mi. pennsylvanicus* harbored 3 coccidia: *E. wenrichi*, *E. saxei*, and *E. subsimi*. No other murids were host to multiple infections.

Three previous studies on the coccidia of *Z. hudsonius* reported 2 species of *Eimeria* (Levine and Ivens, 1990). The prevalence of coccidia across these 3 studies varied considerably. Duszynski et al. (1982) reported a prevalence of 6% (2/35); Gerard et al. (1977), 80% (4/5); and Whitaker (1963), 23% (7/31). It seems unusual that no coccidia were found in the 15 samples we collected. Including this study, there have been no reports of coccidia from the Erethizontidae.

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