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HISTORIA NATURAL Y DEMOGRAFÍA DEL MARSUPIAL ARBÓREO *DROMICIOPS GLIROIDES* EN BOSQUES NATIVOS PRIMARIOS Y DEGRADADOS DEL NORTE DE LA ISLA GRANDE DE CHILOÉ

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A Chiloé y su gente

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RESUMEN

En esta tesis se describen aspectos ecológicos y conductuales del marsupial endémico del bosque templado Monito del monte (*Dromiciops gliroides*) en hábitat de bosques y matorrales en un paisaje rural del norte de la Isla de Chiloé (42 S), Chile. Estudiamos los cambios y las consecuencias de la degradación antrópica del bosque sobre la abundancia poblacional, la demografía y rasgos de historia de vida de *D. gliroides* en parches de bosque antiguo, bosques expuestos a tala selectiva y matorrales como elementos más frecuentes de este paisaje rural.

Durante cuatro años (2005–2009), en la época reproductiva (noviembre) y no reproductiva (febrero-marzo), se estimó la densidad poblacional y parámetros demográficos utilizando la metodología de captura-recaptura en dos fragmentos de bosque antiguo y dos fragmentos de bosques con tala selectiva. Junto con ello, durante los dos primeros años, se monitoreó la presencia de individuos en dos hábitat de matorrales. Tanto en bosques antiguos como expuestos a tala selectiva, se caracterizó la vegetación y recursos alimenticios para *D. gliroides*. Además se comparó cambios en las estimaciones del ámbito de hogar, dieta (contenido fecal), tamaño corporal y carga parasitaria. Considerando todos los individuos capturados, se estudiaron aspectos reproductivos mediante análisis de hormonas, y a través del uso de cajas-nidos se describió la conducta de hibernación.

En bosques antiguos, la conducta de sopor grupal fue más frecuente (64.1%), constituyendo el primer registro de sopor comunal, siendo más frecuente en animales juveniles que en adultos. El ámbito de hogar de machos fue significativamente mayor

que el de hembras independiente del tipo de manejo del bosque. Su dieta es omnívora, consumiendo principalmente artrópodos, sin embargo también con un alto contenido de frutos en individuos adultos en verano y en bosques antiguos. La proporción de insectos y frutos en el contenido fecal varió en forma proporcional a la disponibilidad en el ambiente. Ni el tipo de manejo del bosque ni el sexo afectaron la prevalencia de ectoparásitos, sin embargo se registró una mayor incidencia de ectoparásitos en individuos juveniles provenientes de bosques expuestos a tala selectiva. La carga parasitaria no afectó el peso ni la sobrevivencia de los individuos.

La tala selectiva modifica la estructura de la vegetación de los bosques. La cobertura del sotobosque y la densidad de troncos y árboles grandes fue mayor en bosques antiguos, mientras que la densidad del bambú nativo o quila (*Chusquea* spp.) y la abundancia artrópodos del sotobosque fue mayor en bosques con tala selectiva.

Se documentó la presencia de individuos de *Dromiciops gliroides* en todos los hábitat estudiados, pero su abundancia en matorrales fue considerablemente menor que en fragmentos de bosque. No se observaron diferencias en la abundancia poblacional entre bosques antiguos y con tala selectiva, pero se observó un patrón con mayores abundancias en el período de verano. Tanto en bosques antiguos como en bosques expuestos a tala selectiva no se encontraron diferencias en la sobrevivencia y reclutamiento poblacional, por ende se registraron tasas de crecimiento poblacional similares. Variaciones interanuales en la sobrevivencia y en el crecimiento poblacional podrían estar asociadas a variaciones en los recursos tróficos y condiciones ambientales. Los resultados permiten concluir que junto con la importancia de mantener remanentes de bosques antiguos, la tala selectiva como manejo tradicional

de bosques a pequeña escala, permite mantener poblaciones viables de *D. gliroides* en un paisaje antropogénico.

ABSTRACT

The ecology and behavior of the endemic marsupial Monito del monte (*Dromiciops gliroides*) was described in remnant temperate rain forest in a rural landscape of northern Chiloé Island (42 S), Chile. *D. gliroides* were studied in two remnant old-growth forest fragments, two selectively-logged forest fragments and two shrublands during four years (2005-2009).

We estimated and compared among habitat types: i) lifetime home range, ii) diet (through fecal content), iii) body size and iv) ectoparasitic load. We also characterized reproductive season, through reproductive hormones, and hibernation behavior using artificial nest boxes.

Dromiciops gliroides presence was found in all studied habitat types, but its abundance in shrublands was significantly lower than in forested habitat. Lifetime home range of males was significantly higher than females regardless of habitat types. *D. gliroides* is omnivorous, consuming mainly arthropods, but also with a high content of fruit in adult individuals in summer and in old-growth forest fragments. The proportion of insects and fruit in diet content varied according to its availability. Neither habitat types nor sex affected ectoparasites prevalence, however, there was a higher incidence in juveniles from selective-logged forest. The parasitic load did not affect animal weight of individuals.

Communal nesting was more frequently (64.1%) than solitary individuals, being the first record of communal hibernation, and being more common in juvenile than adults individuals.

Differences in vegetation structures and food resources as a result of forest degradation was described and its consequences on population density and demography of *D. gliroides* were assessed through capture-recapture procedures for four years (2005-2009), during the reproductive (November) and non-breeding (February-March) season between old growth and selective-logged forest patches as common elements of this rural landscape.

Selective logging altered the structure of forest vegetation. Canopy cover and density of understory and large trees was higher in old forests, while the density of native bamboo or quila (*Chusquea* spp.) and understory arthropod abundance was greater in selective-logged forest.

Population density did not differ between habitats however, there was a trend with higher densities in summer period. Demographic parameters did not differ between both habitats; hence there were similar rates of population growth. Interannual variation in survival and population growth could be linked to changes in trophic resources and environmental conditions. The results strongly suggest that along with the importance of remaining old growth forests, selective logged-forest such as a traditional small-scale logging, allows to maintain viable populations of *D. gliroides* in an anthropogenic landscape of northern Chiloé.

INTRODUCCIÓN

Actualmente, la pérdida mundial de la superficie de bosques, debido principalmente al cambio en el uso del suelo (e.g., expansión de la frontera agrícola y explotación forestal; Sala et al. 2000; FAO 2009), ha implicado una disminución del hábitat para muchas especies (Andren, 1994). El área de los bosques templados del sur de Sudamérica fue originalmente limitada por la presencia de la Cordillera de los Andes y el desierto del norte de Chile, quedando restringidos en gran medida a una estrecha faja en el margen oeste del continente al sur de los 35 grados (Hinojosa & Villagran 1997). La superficie cubierta por bosques templados entre los 35 y 42° S ha disminuido en aproximadamente un 51%, quedando sólo el 38% de la superficie remanente de estas regiones cubierta por bosques nativos y el resto cubierto por plantaciones (Lara & Veblen 1993). Esta reducción, sumado a la ausencia de una adecuada legislación y manejo de los bosques nativos, ha llevado a que junto con la actividad forestal industrial de gran escala, fragmentos remanentes de bosque, sean explotados por los propietarios para extracción local de manera y leña, por lo que el paisaje original se ha transformado en un mosaico de remanentes de bosques antiguos y jóvenes, junto con grandes áreas de plantaciones de especies exóticas, además de extensos matorrales, cultivos y praderas (Willson & Armesto 1996; CONAF–CONAMA–BIRF 1999; Echeverría et al. 2007). Este escenario podría representar una amenaza para muchas poblaciones de especies de vertebrados restringidos a los hábitats de bosques nativos (Acosta-Jamett & Simonetti 2004), lo que se manifiesta en que muchos de ellos han

sido incluidos en alguna categoría de amenaza tanto a nivel nacional como internacional (Glade 1993; www.redlist.org).

El Monito del Monte (*Dromiciops gliroides*), una especie restringida a los bosques templados del sur de Sudamérica (Hershkovitz 1999) es una de las especies más carismáticas y, sin embargo, menos estudiadas en su hábitat natural (Redford & Eisenberg 1992; Kelt 2000). Este marsupial es considerado un fósil viviente, ya que es el único representante vivo del Orden Microbiotheria, un Orden monotípico de una línea filogenéticamente primitiva de los mamíferos a nivel mundial (Reig, 1955; Palma & Spotorno 1999). *Dromiciops gliroides* posee un hábito arborícola, frecuentemente asociado a fragmentos de bosque nativo con denso sotobosque, mayoritariamente constituido por quila o bambú (*Chusquea* spp.) de 2-10 m de alto, y con un dosel ocupado por gran diversidad de elementos epífitos y enredaderas (Mann 1955; Patterson et al. 1990; Hershkovitz 1999). *D. gliroides* nidifica en cavidades de árboles que fueron abandonadas por otras especies, o construye nidos inmersos en el denso sotobosque de quila, para hibernar y reproducirse (Mann 1955). Por esta razón, la presencia de árboles con cavidades y de sotobosque denso podrían ser elementos claves del hábitat para asegurar el éxito de la reproducción y sobrevivencia invernal de esta especie.

Dromiciops gliroides posee un hábito trófico insectívoro-omnívoro, ya que consume una gran variedad de artrópodos, especialmente coleópteros (Mann 1955; Meserve et al. 1988). También consume frutos, por lo que algunos autores lo consideran un importante dispersante de las semillas de la planta hemiparásita *Tristerix corymbosus* (Loranthaceae) y de otras especies epífitas de las familias Gesneriaceae,

Philesiaceae y arbustos del genero *Gaultheria* (Ericaceae) (Armesto et al. 1987; Salinas 2008; Rodríguez-Cabal et al. 2007; Amico et al. 2009). Ocasionalmente, depreda huevos y polluelos de *Elaenia albiceps*, *Aphrastura spinicauda* y *Troglodytes musculus* (Jiménez & Rageot 1979; Cornelius 2007).

Estos atributos de la historia de vida de *Dromiciops gliroides*, los que están fuertemente asociados al hábitat de bosques nativos (Mann 1978; Jiménez & Rageot, 1979, Kelt & Martínez 1989), hacen de esta especie un interesante sujeto de estudio para evaluar las consecuencias de la pérdida de cobertura y degradación de bosques por acción antrópica.

En el paisaje rural, sus poblaciones podrían quedar limitadas exclusivamente a los remanentes de vegetación nativa (Patterson et al. 1990; Saavedra & Simonetti 2005). Sin embargo, estos remanentes de bosque son comúnmente degradados a través de la tala selectiva o "floreo", que modifica la estructura y composición del bosque posiblemente afectando la viabilidad de las poblaciones de *Dromiciops* en los paisajes fragmentados. Además, *Dromiciops gliroides*, está categorizado por la legislación chilena como "inadecuadamente conocido", debido a la falta de información ecológica y por las actuales tasas de pérdida de su hábitat lo que implica que se considera en la categoría vulnerable o en peligro de extinción, (D.S. N° 151, 2007). A nivel mundial, *D. gliroides* integra la lista de especies con problemas de conservación, clasificada como "casi amenazado" (Díaz & Teta 2008; véase en www.redlist.org).

La degradación de los bosques remanentes, en paisajes rurales del sur de Chile, actúa como un proceso invisible a las estimaciones de cobertura forestal debido a las tradicionales prácticas de tala selectiva, floreo, tala, roce o quemas (Armesto et al. 2009) y la remoción del sotobosque para la protección del ganado (Willson et al. 1994). En el caso de los bosques templados de Chile, se estima que el 62% de la superficie remanente de bosques nativos (entre Maule y la región de Los Lagos, 35 - 42° S) se encuentra en distintos estados de degradación (i.e., en las condiciones denominadas renoval, adulto-renoval y achaparrado; según CONAF–CONAMA–BIRF 1999). Una de las formas de degradación más común es la tala selectiva, la que involucra la remoción selectiva de los árboles maduros, alteración de la estructura de la vegetación por apertura de claros, aumentos de la cobertura de quila, desecación del hábitat y pérdida de diversidad y biomasa de epifitas (Laurance 1996; Lindenmayer et al. 1997; Díaz et al. 2005; Pérez et al. 2009; Díaz et al. in press). Algunos elementos estructurales considerados “críticos” para especies de vertebrados están relacionados con sitios de alimentación, refugio y nidificación (Stenseth 1980; Tews et al. 2004). La pérdida de estas estructuras (e.g., árboles de gran tamaño, troncos muertos, presencia de sotobosque, y elementos epifitos) sumada a la modificación de los recursos tróficos (i.e., flores, frutos e invertebrados) contribuyen a cambiar la calidad del hábitat, modificando así los procesos demográficos y las abundancias poblacionales de muchas especies que dependen de la presencia de estos elementos estructurales y tróficos (Raphael & White 1984; Saunders et al. 1991; Cornelius 2007). Mamíferos arborícolas, tales como *D. gliroides*, pueden ser particularmente vulnerables a la pérdida de estructuras y recursos tróficos del bosque nativo ya que, como propondré, la

densidad del sotobosque, la presencia de grandes árboles e indirectamente de cavidades y también la abundancia de frutos carnosos e insectos serían determinantes para la persistencia de esta especie en los fragmentos de bosques remanentes del paisaje rural. En consecuencia, postulo que la degradación de los remanentes de bosque por efectos de la tala selectiva podría afectar negativamente las poblaciones de *Dromiciops gliroides* en estos ambientes.

Se ha documentado que la tala de árboles y la remoción del sotobosque en bosques fragmentados del norte de la Isla de Chiloé han afectado la riqueza y abundancia de aves, alterando la abundancia de sus presas (Reid et al. 2004), y que la presencia de una matriz de matorrales y praderas restringe su capacidad de dispersión y colonización de nuevos hábitats (Willson 2004), confinando los individuos a fragmentos de bosque de reducido tamaño y de menor calidad de hábitat (Willson et al. 1994; Sieving et al. 1996; Willson 2004; Díaz et al. 2005; Cornelius 2007; Castellon & Sieving 2006 a, b). En bosques tropicales de Madagascar, Australia y sudeste de Brasil, la pérdida de árboles de gran tamaño y cambios en cobertura del dosel, lianas y sotobosque a consecuencia de la explotación maderera ha afectado negativamente las abundancias de poblacionales de aves y mamíferos, incidiendo directamente en su densidad y éxito reproductivo (Laurance 1990; Laurance 1996; Aleixo 1999; Gentile & Fernández 1999; Quental et al. 2001; Rendigs et al. 2003; Pardini 2004; Viveiros de Castro & Fernández 2004; Watson et al. 2004; Lambert et al. 2005; Bragagnolo et al. 2007).

Varios autores argumentan que las actuales superficies de áreas silvestres protegidas en Chile y otros países no serían suficientes para asegurar la protección de

toda la biodiversidad, fallando especialmente en el caso de los vertebrados (Simonetti & Mella 1997; Tognelli et al. 2008; Ranganathan et al. 2008). Sobre esta base, es interesante conocer la capacidad que tienen los remanentes de vegetación nativa que se encuentran en el paisaje antropogénico, para albergar poblaciones de especies especialistas de hábitat como *Dromiciops gliroides*, y de esta manera complementar los esfuerzos de conservación hoy limitados a las áreas protegidas formales (Armesto et al. 1998).

El tema central de esta tesis es determinar cuál es el efecto de la tala selectiva de los fragmentos de bosque nativo, remanentes en el paisaje rural del norte de la isla de Chiloé, sobre la densidad poblacional y demografía del marsupial arbóreo *Dromiciops gliroides*. Debido al escaso conocimiento ecológico de *D. gliroides*, el primer objetivo de este trabajo se centrará en investigar aspectos no conocidos de la historia natural y ecología de *Dromiciops gliroides* en paisajes de bosques remanentes, para luego analizar las consecuencias de la degradación del bosque por tala selectiva sobre sus poblaciones locales. Específicamente, responderé las siguientes preguntas: ¿cuál es el tamaño de las poblaciones de *Dromiciops* en fragmentos de bosque antiguo (sin tala), bosques degradados por tala selectiva y en matorrales secundarios, como principales ambientes que caracterizan el paisaje rural actual del norte de la Isla de Chiloé? Considerando el hábito estrictamente arborícola de *D. gliroides* y su reconocida afinidad por hábitats de bosques, también me pregunto ¿cuáles serán las consecuencias de la degradación de los fragmentos de bosque para la demografía de la especie y, por ende, el futuro de las poblaciones de este marsupial en los fragmentos de bosque del actual paisaje rural? Debido a que el proceso de degradación del hábitat

actuaría en conjunto con la pérdida de superficie de bosques y acentuaría sus efectos, espero que *D. gliroides* muestre una gran sensibilidad a los cambios de estructura de los parches de hábitat, lo cual debería reflejarse en una menor densidad poblacional y éxito reproductivo, limitando su capacidad de persistir en los actuales paisajes rurales de la isla de Chiloé.

Marco teórico. — El conocimiento de los factores que regulan la dinámica de las poblaciones en su ambiente ha sido uno de los temas centrales de la ecología (Andrewartha & Birch 1954). Los modelos de selección de hábitat predicen que hábitats de mejor calidad, serían preferidos por sobre hábitats de inferior calidad resultando en una distribución "libre-ideal" (Fretwell & Lucas 1970). Bajo este escenario, parches de bosques antiguos serían ocupados primeramente por sobre áreas con bosques expuestos a tala selectiva (supuestamente hábitats de "calidad inferior"), hasta que su oferta de recursos se reduzca debido a la alta densidad poblacional, igualando al hábitat sub-óptimo. Sobre la base de este modelo teórico, se esperaría que la densidad poblacional de *Dromiciops gliroides* sea menor en bosques modificados por la tala selectiva en comparación con bosques antiguos no intervenidos. Este modelo de selección de hábitats supone que los individuos se mueven libremente entre los hábitats, pero en paisajes antropogénicos este puede no ser el caso. Si la conectividad entre los hábitats remanentes en el paisaje antropogénico es reducida, a medida que se incrementa la densidad, procesos denso-dependientes pueden aumentar su importancia y reducir la sobrevivencia poblacional (Turchin 1999).

La estructura del hábitat tiene una considerable influencia en la distribución y abundancia de los vertebrados (MacArthur & MacArthur 1961; Rosenzweig 1991). Estudios recientes realizados en marsupiales arborícolas, que respecto a la utilización de recursos tróficos se comportan como generalistas, han documentado un alto grado de especialización en el uso de ciertas estructuras del hábitat (Laurance 1990; Laurance 1996; Lindenmayer et al. 1994; Lindenmayer et al. 1997). Teóricamente, en términos de tiempo y energía es más "fácil" ser especialista de hábitat que especialista trófico (Schoener 1974; Kozakiewicz 1995; Morris 1996; Bowman et al. 2000). Esto sugiere que la pérdida o modificación de estructuras del hábitat de los bosques, producto de la degradación por tala selectiva o quema, tendría un mayor impacto sobre las poblaciones de vertebrados que la disminución de algunos recursos.

La densidad de una población es resultado de la interacción entre las características de historia de vida de los organismos y la disponibilidad de hábitat y recursos tróficos. En el caso particular del marsupial arbóreo *Dromiciops gliroides* en bosques de Chiloé, la degradación de un parche de bosque puede reducir marcadamente la heterogeneidad vertical (Aravena et al. 2002; Pérez et al. 2009) y aumentar la varianza espacial y temporal en la disponibilidad de recursos (Foley et al. 2007). De acuerdo a este marco teórico, estas diferencias alterarían los procesos demográficos locales, con consecuencias negativas para la densidad poblacional de *D. gliroides*. Frecuentemente, la densidad poblacional constituye un indicador de calidad del hábitat, ya que un mayor número de individuos en un área respondería a una mayor cantidad de recursos disponibles. Sin embargo, a veces es posible encontrar altas densidades poblacionales en hábitat subóptimos producto efectos de dominancia,

territorialidad o estacionalidad (Van Horne 1983; Pulliam 1988; Loeb 1999; Brawn & Robinson 1996; Manning & Edge 2004), o debido a que la densidad responde a la denominada "deuda de extinción" en hábitats fragmentados (Tilman et al. 1994). En consecuencia, parámetros demográficos de la población, tales como las tasas de sobrevivencia y fecundidad son variables más apropiadas para estimar la probabilidad de persistencia de las poblaciones en parches de hábitats remanentes (Loeb 1999; Brawn & Robinson 1996; Manning & Edge 2004).

Estudios previos han documentado grandes diferencias en las densidades de poblaciones de *D. gliroides* en bosques del sur de Chile. Murúa (1996) clasificó a esta especie como escasa, con rangos de densidad entre 10-19 individuos/ha. Por otra parte, densidades mucho menores se registran en hábitats fragmentados (Saavedra & Simonetti 2005; Rodríguez-Cabal et al. 2007), mientras que Patterson et al. (1990) registraron altas densidades en claros y bordes del bosque, lo que atribuyeron a una mayor abundancia de artrópodos asociados al bambú nativo, *Chusquea* sp. Sin embargo, Fortúrbel (2008) registró densidades de hasta 21 individuos/ha en un fragmento de bosque primario en Las Cascadas en Osorno, argumentando que las menores densidades estimadas en estudios previos se debían a problemas de detectabilidad de *D. gliroides* (Kelt & Martínez 1989). En este sentido, metodologías más robustas para estimar la densidad de poblaciones (Parmenter et al. 2003) de *D. gliroides* en hábitats de bosques antiguos y bosques perturbados por tala selectiva nos ayudarían a comprender las consecuencias poblacionales de la degradación antropogénica del hábitat, proceso que ha ido en incremento en las últimas décadas (Echeverría et al. 2008).

Finalmente, los parámetros demográficos están relacionados con atributos de historia de vida de las especies. Por ello, un conocimiento más profundo de estos atributos en el caso de *Dromiciops gliroides*, nos ayudaría a comprender su susceptibilidad a la modificación del hábitat y a explicar o predecir las tendencias demográficas en parches de bosques remanentes del ambiente rural. Por lo tanto, el conocimiento de la historia natural de esta especie, y otras especies amenazadas por la pérdida y degradación de sus hábitats, constituye una herramienta fundamental para estimar la probabilidad de persistencia de las especies en distintos escenarios de paisajes de bosques fragmentados.

Tal como se señaló al principio, el objetivo general de esta tesis es determinar cómo los cambios antropogénicos de la vegetación (bosque antiguo vs. bosque con tala selectiva y matorrales) -- que involucran cambios en cobertura de sotobosque y productividad del hábitat -- afectan la ecología y demografía de las poblaciones locales de *Dromiciops gliroides*, y determinan así el valor del paisaje rural del norte de la Isla de Chiloé como reservorio para las poblaciones de esta especie y complemento del Sistema Nacional de Áreas Protegidas del Estado.

Estructura de la Tesis. — La presentación de esta tesis está dividida en dos capítulos y dos anexos. El primer capítulo, describe los aspectos menos conocidos de historia natural, atributos de historia de vida y ecología de *Dromiciops gliroides* en fragmentos de bosques nativos antiguos, bosques afectados por tala selectiva y matorrales secundarios, como principales tipos de comunidades en el paisaje rural del norte de la

Isla de Chiloé. En este capítulo se estima su densidad poblacional en cada tipo de comunidad y analizan las consecuencias de la degradación de los parches de bosque y cambio potencial del paisaje (relación bosque/matorral) sobre la ecología de las poblaciones.

Los objetivos específicos del capítulo fueron: (i) Estimar la densidad poblacional de *D. gliroides* en hábitats de bosque antiguo, expuestos a tala selectiva y matorrales del paisaje rural del norte de la isla de Chiloé, (ii) estimar el ámbito de hogar y dieta, (iii) describir su ciclo reproductivo y conducta de hibernación y (iv) describir su carga parasitaria, caracterizada con mayor detalle en ambos anexos. Esta información ecológica nos permitiría comprender la respuestas a los cambios del hábitat de *Dromiciops gliroides* a nivel del paisaje.

En el segundo capítulo se caracteriza la demografía de las poblaciones de *D. gliroides* en remanentes de bosques antiguos y bosques sujetos a tala selectiva en el paisaje rural del norte de la Isla de Chiloé. Los objetivos específicos de este capítulo fueron: (i) caracterizar las diferencias en la estructura de la vegetación y recursos alimenticios (e.g., frutos e insectos) entre bosques antiguos y sujetos a tala selectiva, (ii) estimar los tamaños poblacionales en los dos tipos de bosques y (iii) estimar y comparar los parámetros demográficos (sobrevivencia, reclutamiento y tasa de crecimiento poblacional) de *D. gliroides* en bosques antiguos y sujetos a tala selectiva.

Hipótesis de trabajo.

1. La tala selectiva modifica la estructura de la vegetación y los recursos tróficos (e.g., frutos e insectos), a través de: a) la remoción de grandes árboles del dosel e

indirectamente de especies epífitas, b) la disminución del sotobosque, y c) la pérdida de hábitats para las poblaciones de artrópodos que sirven de alimento a *Dromiciops*. Como consecuencia, se espera que la densidad poblacional de *Dromiciops gliroides* en bosques antiguos sea mayor que en los bosques sujetos a tala selectiva.

2. Debido a la remoción de árboles de gran tamaño por tala selectiva, se espera una reducción de la densidad de potenciales sitios de hibernación y nidificación, lo que afectaría negativamente la tasa de crecimiento poblacional de *Dromiciops gliroides* a través de la modificación de parámetros demográficos (i.e., sobrevivencia y reclutamiento) en comparación con los bosques antiguos sin tala.

3. La extracción de grandes árboles del dosel produce claros que conllevan aumentos en la densidad de quila (*Chusquea* spp.) en el sotobosque, la que es un hábitat para una alta densidad de artrópodos (Reid et al. 2004) y, a la vez, constituye un refugio contra los depredadores de *Dromiciops gliroides*. El aumento de la quila podría compensar la pérdida de grandes árboles con cavidades y especies epífitas con frutos carnosos. Por lo tanto, se espera como hipótesis alternativa, que la densidad de *Dromiciops gliroides* en bosques con tala selectiva y con una alta densidad de *Chusquea* spp., sea igual o mayor a la densidad en los bosques antiguos.

1 CAPITULO I

2

3

4

5

6 **POPULATION DENSITIES, FOOD HABITS, AND BEHAVIOR OF THE THREATENED**
7 **ARBOREAL MARSUPIAL *DROMICIOPS GLIROIDES* IN A RURAL MOSAIC OF**
8 **FORESTS AND SHRUBLANDS IN CHILOE ISLAND, CHILE**

9

9 **ABSTRACT**

10 The ecology and behavior of the endemic and threatened Monito del monte (*Dromiciops*
11 *gliroides*) were studied in different habitats in a rural area of northern Chiloé Island (42
12 S), Chile. Individuals were live-trapped for four years (2005–2009) during their breeding
13 (November) and non-breeding (February) season. MNKA was used as an estimation of
14 population density, and home range, diet (through fecal content) and health status
15 (ectoparasite load) were assessed. The breeding season was estimated through
16 reproductive hormones and winter torpor behavior was documented using artificial nest
17 boxes. Population varied seasonally in accordance with breeding, with higher densities
18 recorded during summer. Population densities were similar in old growth than in
19 selective logged forest, but were considerable higher in comparison with shrublands.
20 Social torpor was documented for the first time and was highly frequent (64.1%),
21 especially among juveniles. Home ranges were two times larger for males than for
22 females. *Dromiciops gliroides* is omnivorous, with a diet based mainly on arthropods,
23 and a higher consumption of fleshy fruits in summer. Habitat and age had significant
24 effects on ectoparasite prevalence with higher incidences in juveniles at selective
25 logged forest. Deeper knowledge on the ecology of *D. gliroides* is crucial for the
26 assessment of its conservation status in a changing forest landscape.

27

28 Key words: countryside, selective-logged forest, old-growth forest, communal nesting,
29 natural history, marsupial, temperate rainforest.

30

30 INTRODUCTION

31 *Dromiciops gliroides* (Microbiotheria: Microbiotheriidae), locally known as “Monito del
32 monte”, is the only extant member of the order Microbiotheria (Reig, 1955), with strong
33 phylogenetic affinity with Australasian marsupials (Springer et al., 1998; Palma and
34 Spotorno, 1999; Phillips et al., 2006). It is considered a living fossil or relict in the South
35 American mammal fauna (Springer et al., 1998; Palma and Spotorno, 1999). The
36 species is endemic to temperate rainforests of South America (Mann, 1978;
37 Hershkowitz, 1999) with a distribution that extends along the Chilean Pacific rim from
38 Talca (ca. 35° 30') (Lobos et al., 2005) to Chiloé Island (ca. 44° 00') and adjacent
39 forested habitats of Neuquén and Río Negro provinces in southwestern Argentina
40 (Hershkowitz, 1999).

41 From late 1800's onwards, deforestation has been intensive in South American
42 temperate rainforests (Armesto et al., 1998; Echeverría et al., 2006), including northern
43 Chiloé Island (Willson and Armesto, 1996; Echeverría et al., 2007; Jaña-Prado et al.,
44 2006). In addition, fragmentation and degradation of forested habitats through logging,
45 grazing by cattle and fire, alter forest structure and tree species composition affecting
46 the populations of forest vertebrates (Díaz et al., 2005). The current rural landscape of
47 northern Chiloé Island comprises a mosaic of remnant forest fragments (including old
48 growth and secondary forest), wet scrublands, pastures and agricultural fields (Willson
49 and Armesto, 1996; Aravena et al., 2002; Armesto et al., 2009).

50 The massive loss of forested habitat, together with the limited geographical range
51 of *D. gliroides*, are possibly leading to its decline over recent decades, as shown by its
52 inclusion in the IUCN red list first as “vulnerable,” recently changed to “near threatened”

53 (Díaz and Tetas, 2008). *Dromiciops gliroides* has been recently classified in the Chilean
54 legislation as “inadequately known,” which means that it could be either endangered or
55 vulnerable, but it remains to be assessed (Chilean law Supreme Decree N° 151, March,
56 2007).

57 Despite its near threatened condition, the natural history, food habits, and
58 behavior of this arboreal marsupial remains poorly known in southern South American
59 forests (Jiménez and Rageot, 1979; Kelt and Martínez, 1989; Redford and Eisenberg,
60 1992; Muñoz-Pedreros and Palma, 2000) and virtually nothing is known about its
61 habitat use in rural landscape mosaics. *D. gliroides* has been described as an
62 scansorial nocturnal marsupial, most commonly found or seen in evergreen-
63 broadleaved temperate rainforests (Mann, 1978; Jiménez and Rageot, 1979;
64 Hershkowitz, 1999), often dominated by southern beeches (*Nothofagus*:
65 *Nothofagaceae*), with dense understory cover of native bamboos of the genus
66 *Chusquea* (Mann, 1978; Patterson et al., 1990). During winter *D. gliroides* nests in snag
67 cavities or inside the dense bamboo understory (Mann, 1978; Kelt and Martínez, 1989)
68 and the animals hibernate under cold-temperate winter conditions and/or during food
69 shortages (Mann, 1978; Bozinovic et al., 2004). According to Mann (1978), *D. gliroides*
70 reaches sexual maturity in its second year of life, and it has been reported as a
71 monoestrous, seasonal breeder, producing one litter per year, with births occurring in
72 the early austral spring, October to November (Muñoz-Pedreros et al., 2005). Based on
73 summer analyzes of stomach contents of an Andean population, *D. gliroides* has been
74 classified as an insectivore (Meserve et al., 1988), but recent studies highlight its trophic

75 role as a seed disperser of several fleshy-fruited temperate forest trees and vines
76 (Armesto et al., 1987; Amico and Aizen, 2000; Aizen, 2003; Rodríguez-Cabal et al.,
77 2007; Amico et al., 2009), and also as a nest predator of the forest insectivorous bird
78 *Aphrastura spinicauda* (Cornelius 2007).

79 Considering that most current knowledge of the species derives from Andean
80 forests, despite its widespread range in lowland and coastal Chilean forests, and that
81 lowland areas are changing rapidly due to human land use (Echeverría et al., 2007),
82 this study seeks to advance our knowledge of the ecology of the relict *D. gliroides* in a
83 rural landscape of northern Chiloé Island and its propensity to become endangered due
84 to the loss of forested habitats.

85 We concentrate our study in the two main habitats within the rural landscape
86 where this arboreal species could dwell, including forested patches (both old growth and
87 selectively logged patches) and shrublands. We did not sample the other two main rural
88 habitats, pastures and agricultural fields, because previous intensive sampling in
89 northern Chiloé (E. Palma and M. Bustamante, personal communication, and J.L.C.
90 personal observations), and data from mainland sites (Fontúrbel et al., 2009) showed
91 no evidence of presence of *D. gliroides* in these habitats.

92 Our specific goals were: (1) to assess the population densities of *D. gliroides* in
93 forest patches (i.e., old growth and selectively-logged) and shrublands that are part of
94 the rural landscape mosaic in Chiloé Island; (2) to assess lifetime home range and diet
95 of the species; (3) to establish the seasonality of breeding and torpor behaviors; and (4)
96 to describe its ectoparasites. We use this information to discuss the conservation status
97 of this relict marsupial in the rural mosaic of northern Chiloé Island. Such knowledge is

98 critical for proposing science-based guidelines for the conservation of this unique
99 marsupial, and accompanying biodiversity, especially where old growth forests are
100 becoming scarce (Gutiérrez et al., 2009) and limited land can be allocated to new
101 reserves.

102

103 MATERIAL AND METHODS

104 *Study site.* — The study was conducted in a rural area of northeastern Chiloé Island,
105 Ancud County, Chiloé province (Fig. 1). Within the northern Chiloé rural landscape, we
106 studied two types of forest habitat (Fig. 2; Aravena et al., 2002; Díaz et al., 2005;
107 Cornelius 2007): two old growth forests (OGF; one within Senda Darwin Biological
108 Station and the other in Caulín) and two selective-logged forests (SLF; named
109 Llanquihue and Quilar), where valuable timber trees had been removed by landowners
110 with no defined silvicultural system (Cornelius 2007). We also looked for *Dromiciops*
111 *gliroides* in dense shrublands (ESS) surrounding forested areas within Senda Darwin
112 Biological Station (ESS1 and ESS2). OGF habitats had an age of at least 300 years
113 based on tree-ring counts of dominant canopy trees, and had no visible signs of logging
114 or fire (Aravena et al., 2002; Gutiérrez et al., 2004). These OGF patches were
115 characterized by a multi-layered canopy, with large emergent trees of *Nothofagus nitida*
116 (Nothofagaceae), *Drimys winteri* (Winteraceae), *Podocarpus nubigena* and
117 *Saxegothaea conspicua* (both Podocarpaceae), plus few individuals of *Eucryphia*
118 *cordifolia* (Eucryphiaceae) and *Caldcluvia paniculata* (Cunoniaceae) in the lower
119 canopy. Trees were often covered with vines and epiphytes such as ferns, mosses and

120 bromeliads (Muñoz et al., 2003). Large amounts of dead biomass (snags and logs) are
121 frequent within OG stands. The understory had a high density of tree saplings, few
122 shrubs, and dense thickets of 2-3 m tall native bamboo (*Chusquea* spp.) within tree-fall
123 gaps (Gutiérrez et al., 2004). SL patches are frequent in the rural landscape due to the
124 widespread practice of small-scale logging, and had areas with a closed canopy of
125 dense juvenile trees, 10–20 m tall, predominantly shade-intolerant *D. winteri*, and *N.*
126 *nitida*. There are also few large trees, legacies from the pre-disturbance forest. In SLFs,
127 open areas with a dense cover of native bamboos were frequent, with a sparse
128 presence of tree seedlings and saplings due to shading (Aravena et al., 2002).
129 Shrublands developed after anthropogenic forest fires, and were dominated by a 1-2 m
130 tall layer of *Baccharis magellanica* (Asteraceae) and a <1 m tall layer of *Blechnum*
131 *chilensis* (Dicksoniaceae), with sparse cover of *Berberis darwinii* and *B. microphylla*
132 (both Berberidaceae) (Díaz et al., 2005). These shrublands have sparse regeneration of
133 shade-intolerant tree species such as *D. winteri*, and *N. nitida* (1–3 m tall) forming small
134 patches.

135 The prevailing climate in northern Chiloé is wet-temperate with a strong oceanic
136 influence (Di Castri and Hajek 1976). Climatic records (i.e., air temperature and
137 precipitation) to characterize the studied area were obtained from Senda Darwin
138 Biological Station (45° 53' S, 73° 40' W) (see appendix 1).

139

140 *Animal trapping and field data.* — Live captures of *D. gliroides* were carried out in forest
141 habitats from spring 2005 to spring 2009 and only until summer of 2007 for both

142 shrubland sites. Trapping was conducted over 10 consecutive nights, twice each year:
143 (1) in mid-spring (November-December), after torpor and before the onset of the
144 breeding season, and (2) late austral summer (February-March), after the breeding
145 season had ended. We did not trap animals in autumn or winter because the harsh
146 environmental conditions increases mortality risk and animals are largely inactive. We
147 used modified Tomahawk-live traps (baited with banana) because Sherman traps were
148 not effective (see Kelt and Martínez, 1989; Patterson et al., 1989; Fontúrbel and
149 Jiménez, 2009). Traps were placed on tree branches 1-2 m high, because of dense
150 understory and structural complexity of both forest fragments and shrublands, and
151 arranged spatially in a web design, which consisted of 12 radial lines of 100 m each.
152 Each trapping line had 12 stations; the first four were at 5 m intervals from the center (5,
153 10, 15, and 20 m), and the following eight were at 10 m intervals (30 m up to 100 m).
154 Thus, 144 traps were set up in each site per trapping night (Parmenter et al., 2003). The
155 area covered by each web of traps was 3.46 ha. Each trap was checked daily between
156 0600-0800 h.

157 For all captured animals we recorded sex, age, breeding status (based on
158 scrotum size, pouch morphology and mammary gland development), and body mass,
159 using a Pesola ® (Baar, Switzerland) spring balance, 0.1 g precision. Animals were
160 marked with numerated 2 mm long aluminum ear tags (Tag no. 1005-1, National Band
161 and Tag Co., Newport, KY). Each individual was released at the capture site. To avoid
162 pseudo replication in statistical analyses, we considered only new unmarked individuals.
163 ESS had to be excluded due to low sample sizes and low capture probabilities. Capture

164 procedures followed guidelines approved by the American Society of Mammalogists
165 (Gannon et al., 2007) and complied with current Chilean laws (SAG permit No. 2605).

166 The effect of habitat (OG forest, SL forest and shrubland), season and sex on
167 adult body mass were analyzed using three-way ANOVA (STATISTICA; Statsoft Inc.
168 2001), with log-transformed mass as the dependent variable and habitat, season and
169 sex as a grouping factor.

170

171 *Population abundance.* — The minimum number of individuals known alive (MNKA) was
172 used as an index of population abundance for all habitat types and sites (Lima et al.,
173 2001; Rodríguez-Cabal et al., 2007). Population abundances of *D. gliroides* from each
174 forest fragment were associated with monthly temperature and precipitation obtained
175 from Senda Darwin Biological Station using a nonparametric Spearman rank correlation
176 (STATISTICA; Statsoft Inc. 2001).

177

178 *Home range.* — Lifetime home ranges (i.e., four years) were estimated in both forested
179 habitat for marked-recaptured individuals from the entire study period (2005 to 2009),
180 using capture positions (XY-coordinates from trap data) and analyzed using Animal
181 Movement SA (v2.04 beta -ArcView GIS 3.2). Despite some limitations of mark-
182 recapture data, this method still is widely used to estimate home range for small
183 mammals (Ribble et al., 2002). For adult males and females trapped more than three
184 times, the 95% minimum convex polygon (MCP) was used as an estimate of home
185 range size (Agüero and Simonetti, 1988). The MCP method is more robust than other

186 home-range estimators (e.g. radio-tracking), when the number of fixes is low (Agüero
187 and Simonetti, 1988).

188 The effect of habitat and sex on home range areas were analyzed using two-way
189 ANOVA (STATISTICA; Statsoft Inc. 2001), with log transformed MCP areas as the
190 dependent variable and habitat and sex as a grouping factor. In addition, we assessed
191 the relationship between home-range area and adult body mass (for both sexes) using
192 a nonparametric Spearman rank correlation (STATISTICA; Statsoft Inc. 2001).

193

194 *Diet analysis.* — The diet of *D. gliroides* in all sites was studied by examining prey and
195 vegetal remains in fecal samples during the first three years of trapping periods. Feces
196 were collected in the field from inside or under traps in both forest fragments. Feces
197 were placed in labeled 1.5-ml Eppendorf vials with 70% alcohol (Pinheiro et al., 2002)
198 and dried in the laboratory. They were later teased apart under a 40X binocular
199 dissecting scope and remains of arthropods were determined to at least ordinal level
200 with keys for exoskeleton fragments (Pinheiro et al., 2002). Seeds were identified using
201 reference samples for plant species in the forest and surrounding areas. Frequencies of
202 occurrence (FO%) were estimated by counting the number of each prey or seed type in
203 feces and then dividing by the total number of prey or seeds identified without correction
204 for differential digestibility (Jiménez, 2007).

205 We compared variation in the proportions of items in the diet of *D. gliroides*
206 among both forested habitats (i.e., OGF and SLF) and seasons (i.e., spring and
207 summer) using the-log likelihood ratio (G Test: Zar 1999). For this comparison, we

208 pooled data from juveniles and adults individuals, due to inadequate sample size ($n <$
209 30) at each category for a simultaneous analysis. In addition, we explored differences in
210 prey content between juveniles and adults individuals. This comparison was done only
211 for summer, due to the larger number of juveniles. Finally, we described the overall
212 composition of the diet of *D. gliroides* in terms of items in fecal content using all
213 samples.

214 For graphical comparisons, dietary results by habitat and season were simplified
215 further by combining categories into three broad classes (following Meserve et al.,
216 1988): arthropods and vertebrates, vegetative plant material and fungi, and fruits and
217 seeds. We used De Finetti diagrams following procedures of Meserve et al. (1988) for
218 illustration purposes, where the trophic position of a group is represented as a single
219 point whose distance to each endpoint of an equilateral triangle is proportional to the
220 dietary composition along three axes categories.

221

222 *Ectoparasites and health status.* — Ectoparasites were collected from unmarked
223 individuals captured by live trapping from spring 2005 until summer 2007. Ectoparasites
224 were collected after visual examination following Marín-Vial et al. (2007). Samples were
225 placed in labeled 1.5 ml Eppendorf vials with 70% alcohol and identified taxonomically
226 in the laboratory.

227 Ectoparasites were identified using descriptive keys, and their prevalence and
228 abundance were measured according to definitions given by Bush et al. (1997). To
229 analyze the effect of habitat, season, sex, and age of *D. gliroides* in ectoparasite

230 prevalence, a GLIM model selection with binomial distribution was conducted in R
231 project software (version 2.7.1, 2008). Mass of parasitized adults *versus* non-parasitized
232 individuals was analyzed using one-way ANOVA (STATISTICA; Statsoft Inc., 2001).
233 Proportions of recaptured individual, at least once, as a proxy of survival, were
234 compared between infected and non-infected individuals using a proportion test in R
235 project software (version 2.7.1 2008).

236

237 *Nest box occupancy and torpor.* — In winter 2002 and 2004, artificial nest boxes were
238 installed in two OG forests (i.e., Senda Darwin and Caulín) as part of a parallel bird
239 survey (see Moreno et al., 2005). All nest boxes (N=194) were placed more than 500 m
240 away from the trapping area and were fastened to tree trunks or branches ca. 150 cm
241 above the ground. Nest box dimensions were described in Moreno et al. (2005). To
242 describe torpor behavior of *D. gliroides* we monitored the nest boxes in the austral
243 winter period (July-August) during 2005 and 2006, when birds were not using them. All
244 animals found in boxes were measured and marked as those captured with web live
245 trapping method.

246 To assess the effect of age and sex on the probability of communal *versus*
247 solitary nests found in the occupied boxes, data were analyzed using GLIM model
248 selection with binomial distribution at R project software (version 2.7.1, 2008).

249

250 *Reproduction.* — At all sites sexual maturity and breeding season were assessed by
251 measuring the levels of plasma testosterone in males and progesterone in females from

252 blood samples, as signs of reproductive activity (i.e., follicular development and corpora
253 lutea). Blood samples (100 μ l) were obtained from adults or juveniles that were
254 anesthetized in a closed chamber for 20 – 30 sec with isofluran in 0.5 – 1 ml volume
255 and blood extracted from the infraorbital sinus with heparinized hematocrit capillary
256 tubes (Van Herck et al., 2000). Samples were centrifuged and the plasma analyzed. All
257 individuals were released after recovery from anesthesia at the site of capture. Due to
258 the small amount of plasma, individuals captured at different sites but with similar ages
259 were pooled. Plasma testosterone and serum progesterone concentrations were
260 analyzed with solid phase I^{125} radioimmunoassay, using a total testosterone or
261 progesterone Siemens kit.

262 The relationship between reproductive hormones and secondary sexual traits,
263 such as testicular size, was assessed using a nonparametric Spearman rank correlation
264 (STATISTICA; Statsoft Inc., 2001).

265 We quantified the proportion of reproductive females and counted in-pouch
266 young for all females at all sites throughout study period. To confirm reproductive data,
267 which suggest that *D. gliroides* becomes sexually active during the second year of life
268 (Mann, 1978), we recorded the age at maturity based on morphological changes (i.e.,
269 testicular size, and pouch development and coloration). For this analysis, we considered
270 only reproductive individuals with previously recorded body mass information at the
271 juvenile stage.

272

272 **RESULTS**

273 *Dromiciops population description.* — From spring 2005 to summer 2009, we live-
274 trapped 502 different individuals of *D. gliroides* with 1065 captures in all habitats and
275 sites. A total of 119 animals were trapped at Senda Darwin, 111 at Caulín, 181 at
276 Llanquihue, 83 at Quilar, 4 in ESS1 and 6 in ESS2 (Table 1). Only two juveniles
277 captured in shrubland (i.e., at ESS2) were trapped again as adults in the Senda Darwin
278 OG forest patch. On average, each animal was recaptured 2.12 times (3.4 times for
279 recaptured individuals). Accidental trapping death was 1.8% (19 out of 1065 captures),
280 and was marginally significantly higher during the spring months (13) than in summer
281 (6, $\chi^2 = 3.31$, *d.f.* = 1, *P* = 0.06; Table 1). On two occasions we observed owls (*Strix*
282 *rufipes* and *Glaucidium nanum*) killing trapped individuals at SL patches.

283 Population varied seasonally in all habitats (Fig. 3). In both forested habitats (i.e.,
284 OGF and SLF), we observed a trend towards higher abundances in the austral summer
285 (February), which corresponded to the juvenile recruitment period, than in spring
286 (November), when mating and breeding takes place (Fig. 3). In contrast, even though
287 abundances were extremely low, we observed an inverse trend in shrublands (Fig. 3).
288 Abundances were significantly correlated with mean temperature, but only for Senda
289 Darwin OG forest (Spearman rank correlation $R = 0.87$, $n = 8$, $P = 0.004$). Population
290 abundances were higher than 45 individuals/ha, and occasionally lower than 6
291 individuals/ha in spring or in shrublands. Over all seasons, years, and sites, OGF had a
292 mean of 4.91 individuals/ha (SE = 0.7); SLF a mean of 5.36 individuals/ha (SE = 0.8)
293 and shrublands a mean of 0.40 individuals/ha (SE = 0.14) (Table 1).

294 We found no significant differences in body mass of *Dromiciops* adults captured
295 between habitats, OG forest, SL forest and shrublands ($F_{1, 268} = 3.75$, $P > 0.06$; Table
296 2). However, adult mass differed significantly between seasons within a year ($F_{1, 268} =$
297 28.71 , $P < 0.01$; Table 2) and between sexes ($F_{1, 268} = 5.43$, $P = 0.02$; Table 2), with
298 lower body mass during spring when individuals were recovering from torpor ($n = 197$,
299 mean \pm SE = 26.87 g \pm 0.4), and higher body mass at the end of summer period ($n =$
300 79 , 30.54 g \pm 0.56). On the other hand, females ($n = 95$, mean \pm SE = 29.5 g \pm 0.5)
301 were heavier than males ($n = 181$, mean \pm SE = 27.9 g \pm 0.46). In addition, the
302 interaction between seasons and sex of the animals was statistically significant ($F_{1, 268} =$
303 5.52 , $P < 0.02$), because summer body mass of females was higher than females in
304 spring and for males in both seasons.

305

306 *Home range.* — A total of 80 adults out of 287 individuals were captured three or more
307 times during the study period (Table 3), enabling a reliable estimation of lifetime home
308 ranges. Because no individuals were captured more than three times in ESS, we did not
309 estimate home range in this habitat. Estimated home range sizes of the 80 individuals
310 from forest habitats ranged from 0.12 to 0.35 ha (Table 3). Home range was not
311 statistically different between OG and SL forest habitats, but it was significantly larger
312 for males than for females, with considerable spatial overlap between sexes (Table 3
313 and Fig. 4). Body mass and home range area were uncorrelated ($r_s = 0.12$, $P = 0.29$, n
314 = 80).

315

316 *Diet.* — A total of 146 fecal samples of *D. gliroides* were obtained from forested habitats
317 (i.e., OGF and SLF) in two seasons of the year (i.e., spring and summer) from 2005 to
318 2008. Only one fecal sample was obtained from a shrubland animal, and hence we
319 excluded this habitat from further analyses. Arthropods were the most frequent item in
320 *D. gliroides* feces, over 40% in all habitats and seasons. However, significant
321 differences were found in the proportion of food items between forest habitats and
322 seasons ($G = 13.95$, $d.f. = 6$, $P < 0.03$, Fig. 5). The proportion of arthropods was higher
323 in SLD than in OGF in both spring and summer (Fig. 5). On the contrary, the proportion
324 of seeds in feces was higher in OGF than in SLF, and this difference increased in
325 summer (Fig. 5). No differences were found between age classes in the proportion of
326 food items from summer samples ($G = 4.74$, $d.f. = 2$, $P < 0.09$). In particular, adults from
327 OGF showed a higher proportion of seeds in summer, when most of shrubs and trees
328 were in fruiting (Smith-Ramírez and Armesto, 1994), in comparison with spring. Hence,
329 in summer samples, fruits were more frequent in feces than were arthropods.

330 Among arthropods, Coleoptera and Orthoptera were the most frequent prey item
331 in feces of this marsupial, with a frequency of occurrence $FO\% = 42.9$ and 24.6 ,
332 respectively (Table 4). Lepidoptera (mostly larvae and pupae), Hymenoptera and
333 Hemiptera were also found at lower frequencies (Table 4). Two fecal samples contained
334 bird bones and feathers (only from spring samples). Seeds of twelve plant species were
335 identified, including forest epiphytes, vines and shrubs. *Tristerix corymbosus*, *Gaultheria*
336 *insana* and *Luzuriaga polyphylla* were the most frequent seeds found in feces,
337 representing 45, 19 and 8% of the total frequency of occurrence (Table 5); however,

338 considering total seed numbers, *Tristerix corymbosus*, *Myrteola numularia* and
339 *Gaultheria* sp. were more abundant in *D. gliroides* feces (Table 5).

340

341 *Ectoparasites and health status.* — Four different taxonomic groups of ectoparasites
342 were identified infesting *D. gliroides*: ticks, fleas, mites and leaches. Ectoparasites were
343 identified to species level for ticks and fleas. For ticks, we detected *Ixodes*
344 *neuquenensis*, the only tick previously reported from *D. gliroides* (recently recorded in
345 Chiloé; Marín-Vial et al., 2007) and for fleas, we detected *Chiliopsylla allophyla* and
346 *Plocopsylla diana*, both species shared with other temperate rainforest rodents
347 (Alarcón, 2003).

348 A total of 102 (55%) of 185 individuals *D. gliroides* were infested with some or all
349 of these ectoparasites in all sites studied (Table 6). Excluding shrublands, due to small
350 sample size ($n = 7$), a GLIM model including forest habitat, season (spring versus
351 summer) and animal age, had the best adjustment to explain the prevalence of
352 ectoparasites (AIC = 236.91, res. dev. = 220.91). Prevalence was statistically different
353 between forest habitats (GLM $W = 5.33$, $P < 0.039$), with higher values in SL forest
354 (62.73%) than in OG forest (44.78%), and was higher in juveniles (62.1%) than in adults
355 (50.9%). Flea prevalence had the same pattern (GLM $W = 5.33$, $P < 0.02$), been higher
356 in juveniles (52.2%) than in adults (31.5%). In addition, tick prevalence was statistically
357 different between summer and spring (GLM $W = 5.33$, $P < 0.02$), with higher prevalence
358 in spring (33.3%) than in summer (15.7%). No significant differences were found

359 between sexes (GLM $W = 5.33$, $P > 0.05$), unlike the findings of Guglielmone et al.
360 (2004).

361 Body mass of infested adults was not significantly different from that of non-
362 infested individuals (ANOVA $F_{1, 115} = 2.24$, $P = 0.14$). Moreover, the proportion of
363 recapture rate of infested adults was not significantly different from that of non-infested
364 individuals ($\chi^2 = 0.04$, $d.f. = 1$, $P = 0.8$).

365

366 *Nest box occupancy and torpor.* — In winter 2005, 28 out of 194 nest boxes were used
367 by *D. gliroides* as den sites in two OG forests. Twenty-three (15.4%) nest boxes were
368 occupied in Senda Darwin and five (11.1%) in Caulín. Similar level of occupancy was
369 observed in winter 2006 with twenty-six (17.5%) and six (13.3%) boxes used by
370 *Dromiciops* in Senda Darwin and Caulín respectively. Nests inside boxes were
371 spherical, with a single entrance, built with materials from epiphytes, vines, and
372 understory vegetation (*Chusquea* leaves, *Hymenophyllum* ferns), and lined with
373 epiphytes, ferns, or mosses.

374 During winter 2005 and 2006, 77 and 62 torpid animals, respectively, were found
375 nesting in these boxes. Overall nest occupancy rate was 16.5 % ($n = 388$ boxes). Mean
376 number of animals in nests was 2.3 ± 0.2 SE (range = 1-9). Forty-one of 64 nest boxes
377 (i.e., 64.1%) were communal nests, significantly greater than expected by random ($\chi^2 =$
378 9.03 , $d.f. = 1$, $P = 0.002$). Animal age had a significant effect on communal nest
379 behavior (Fisher's exact test, $d.f. = 1$, $P = 0.05$; Table 7); juveniles were found solitary in
380 17.4% and communally in 38.3% of the nests, whereas adults were solitary in all the

381 remaining nests, 82.6%. Contrary to expectations, neither sex (Fisher's exact test, *d.f.* =
382 1, $P = 0.34$) nor body mass (three way ANOVA, $F_{1, 139} = 0.0049$, $P = 0.94$), had a
383 significant effect on communal nesting.

384

385 *Reproductive parameters.* — A total of 44 blood samples were obtained from adult
386 males captured throughout the study period. Plasma testosterone levels had a single
387 annual peak during austral spring (November), followed by a steady decline from spring
388 to winter (July), when males entered torpor (Fig. 6). Levels of plasmatic testosterone
389 were significantly correlated with testicular volume (Heideman et al., 2000; $r_s = 0.12$, $n =$
390 36, $P < 0.001$), and related to mating season. Interestingly, like other marsupials (Scott,
391 1979), we observed that most adult males individuals had different scrotal colorations.

392 A total of 21 blood samples were obtained from 19 adult and 2 subadult females,
393 captured in all sites. Plasma progesterone concentration in adult females showed an
394 increase during spring (November) lasting until summer (February), followed by a
395 reduction in winter (August), when females entered torpor (Fig. 6). During spring,
396 subadult females had lower progesterone concentrations than mature females (22.3
397 ng/ml *versus* 73.0 ng/ml). Overall, considering all forest sites and years, we recorded 27
398 adult females with their young still attached to the marsupium, with a mean litter size of
399 2.26 ± 0.17 (1 SE). Two of these females were also trapped pregnant in the
400 reproductive season (i.e., November) of a previous year. In addition, in the summer (i.e.,
401 post reproduction) of 2009, six females were trapped with young still attached to the
402 marsupium (mean \pm SE: 2.17 ± 0.17), from which three were found breeding in the

403 previous (i.e., reproductive) trapping session, suggesting a second reproductive event.
404 Anecdotally, in spring of 2005, we observed parturition and migration of two immature
405 embryos into the pouch. Three days later, we trapped the same female bearing young
406 inside the pouch.

407 Based on hormone profiles and/or primary sexual characters, field observations
408 suggest that subadult individuals become sexually mature when they reach a body
409 mass of $25.2 \text{ g} \pm 0.47$ ($n = 21$) for males and $27.0 \text{ g} \pm 0.98$ ($n = 8$) for females from birth
410 to the next reproductive season (i.e., 10 month approximately)

411

412 **DISCUSSION**

413 In all remnant forest patches in the rural landscape of northern Chiloé Island, regardless
414 of whether some logging of large trees had taken place, we recorded abundant
415 presence of the arboreal marsupial *D. gliroides*. Population densities were similar in OG
416 and SL forest, and there were no differences in adult body mass between habitats.
417 Previous knowledge suggested that *D. gliroides* populations were strongly associated
418 with old-growth forest patches, because of its arboreal habit and nesting behavior,
419 occupying tree holes (Mann, 1978; Jiménez and Rageot, 1979, Kelt and Martínez,
420 1989; Patterson et al., 1989; Hershkovitz, 1999). Other studies have found similar
421 densities of *D. gliroides* in habitats that differed in successional stage such as old
422 growth and secondary forest patches (e.g., Fundo San Martín Experimental Forest
423 Reserve in Valdivia, and Las Cascadas in Osorno), which are characterized by dense
424 bamboo cover in the forest understory (Patterson et al., 1990; Fontúrbel et al., 2009).

425 *Dromiciops gliroides* abundances in forested habitats remaining in northern
426 Chiloé Island ranged from 5 to 48 individuals/ha, a density comparable to estimates
427 available for continuous forests in the Andes (Rodríguez-Cabal et al., 2007) and slightly
428 higher than estimates for an arboreal rodent of southern temperate rainforests,
429 *Irenomys tarsalis* (see Murúa 1996).

430 Considering an annual forest loss rate of 1.1 % (Echeverría et al., 2007), and an
431 increase in the proportion of SL in remnant forest patches in northern Chiloé
432 (Echeverría et al., 2007; Jaña-Prado et al., 2006; Armesto et al., 2009), the finding of
433 comparable densities of *D. gliroides* in OG and SL forest patches of the rural landscape
434 constitutes an important result. If *D. gliroides* is negatively affected by habitat
435 fragmentation, as shown for Andean forests in Argentina (Rodríguez-Cabal et al., 2008),
436 and pastures constitute powerful barriers for movement between forest fragments
437 (Rodríguez-Cabal et al., 2007; Fontúrbel et al., 2009), shrublands and logged forests
438 could facilitate the movement of animals across the rural landscape. SL forests could
439 become relevant by providing suitable habitat for *D. gliroides* in areas where land cover
440 of OG forests has been greatly reduced. According to our results, shrublands that are
441 contiguous to forest habitats could also mitigate restrictions on the movement of
442 animals between forest fragments in the rural landscape. In this sense, recent studies in
443 forest mammals communities, have shown a crucial role of the matrix in maintaining
444 diversity in fragmented landscapes (Umetsu and Pardin, 2007). Consequently,
445 assessing the capacity of remnant forest to support regional biodiversity and inclusion of
446 matrix quality parameters in managing rural landscapes is of critical importance to

447 enhance conservation, especially where agriculture frontiers are still expanding and
448 limited land can be allocated to new reserves (Chan and Daily, 2008).

449 Populations phenology of *D. gliroides* varied between spring and summer each
450 year, with a trend in both forested habitats toward higher densities in summer,
451 associated to favorable weather. In spring 2008, we observed a drastic reduction in *D.*
452 *gliroides* population (Fig. 2). We suggest that a severe drought of spring-summer 2007-
453 2008 could explain this population pattern, but we did not have enough data to support
454 this hypothesis.

455 Reproduction in marsupials is usually cued to environment temperature and
456 resource abundance (Tyndale-Biscoe and Renfree, 1987; Quental et al., 2001).
457 Accordingly, favorable conditions of food and climate during the austral spring-summer,
458 result in synchronous reproductive patterns and maximum sex hormone levels during
459 this period, associated with the development of secondary sexual traits (Muñoz-
460 Pedreros et al., 2005). Based on the general climate of Chiloé Island (di Castri and
461 Hajek, 1979), we suggest a slight delay in the female reproductive cycle in comparison
462 to mainland populations (e.g. developmental morphology; see Muñoz-Pedreros et al.,
463 2005), because mating, gestation and parturition in Chiloé Island generally took place
464 from late October to early December. Higher and constant progesterone levels during
465 gestation and lactation, in contrast to male testosterone, might be attributed to the luteal
466 phase duration (Tyndale-Biscoe and Renfree, 1987). However, a second reproductive
467 event or a delayed one could sustain higher hormones levels during summer periods,
468 contrary to previous reports that *D. gliroides* breeds exclusively in spring (Mann, 1978;
469 Muñoz-Pedreros et al., 2005). In males, melanistic scrotal coloration, such as found in

470 males from Chiloé forest, has been described in species exposed to harsh
471 environments, and thought to aid in regulating sperm temperature (Scott, 1979) and
472 also suggested previously for *D. gliroides* (Grant and Temple-Smith, 1987).

473 In most marsupials, the culmination of lactation and weaning are critical periods
474 for offspring survival (Lee and Cockburn, 1985), and in rural Chiloé forests, these two
475 phases always occurred in summer, during the peak of arthropod and fruit abundance
476 (Smith-Ramírez and Armesto, 1994; Díaz, 2008). Similar population densities in OGF
477 and SLF in rural Chiloé Island, suggest that modification of forest resources by limited
478 logging of some tree species does not translate into demographical consequences (see
479 chapter II).

480 Mann (1978) indicated that *D. gliroides* reached its sexual maturity during its
481 second year. However, our data on sex hormone levels and secondary sexual traits
482 (e.g. testicular size in males and pouch development and coloration in females),
483 suggest that individuals born in mid-spring (November) could reach sexual maturity in
484 the following breeding season (i.e., next November), if they have reached a minimum
485 mass of 25 g and 27 g for males and females, respectively.

486 Adult males often have larger home ranges during mating periods (see
487 Kraaijeveld et al., 2002; Belcher and Darrant, 2004). Considering that our home range
488 estimates from captured animals in Chiloé forests were smaller than radio-tracking
489 estimates (Fontúrbel et al., 2009; unpublished data), larger home ranges were
490 documented in the spring, when mating occurs (Fontúrbel et al., 2009). Smaller home
491 ranges based on mark-recapture methods, in comparison to radio-tracking estimates,
492 can be equivalent to the core areas of radio-tracking estimates (Pavey et al., 2003). In

493 this case, mean adult home range (0.2 ha) from Chiloé forests was similar to the core
494 area defined in an Andean population (0.2 ha) (Fontúrbel et al., 2009).

495 Artificial nest boxes are widely used as ecological research tools and wildlife
496 management resources, especially in field studies of arboreal marsupials (Lindenmayer
497 et al., 2003). Although of controversial application in forest conservation (Lindenmayer
498 et al., 1991; Gibbons and Lindenmayer, 2002), their use may reduce the negative
499 effects of declines in nesting sites due to habitat loss (Lindenmayer et al., 1997;
500 Cornelius 2007). In our case, nest boxes provided useful tools as a proxy to estimate
501 animal abundances and to study nesting and reproductive behavior of a cryptic
502 nocturnal and arboreal marsupial.

503 Mann (1978) reported solitary torpor of *D. gliroides*, and its physiological bases
504 were analyzed by Bozinovic et al. (2004), who noted the first record of hibernation in a
505 South American mammal. Nesting and torpor in tree hollows are common in
506 Australasian marsupials, and appears to be a response to harsh environmental
507 conditions to increase winter survival (Geiser 1994; Canals et al., 1997; Bozinovic et al.,
508 2004). In marsupials, the number of animals nesting in the same place shows wide
509 variation (Lee and Cockburn, 1985; Lindenmayer, 1997). We report the first record of
510 communal nesting for a South American marsupial in the case of *Dromiciops* in Chiloé
511 forests. Communal torpor or huddling is energetically advantageous for social
512 thermoregulation (Lee and Cockburn, 1985; Bozinovic et al., 1988; Rhind, 2003). It is
513 facilitated by reproductive cessation and low relatedness (Rhind, 2003; Ebensperger et
514 al., 2004). Ontogeny and group size may influence the effectiveness of huddling,
515 because early developmental stages and the number of animals reduce the

516 surface/volume ratio, and hence individual metabolic rates (Canals et al., 1998).
517 Accordingly, in rural populations of *D. gliroides*, communal nests are often composed of
518 juveniles with lower body mass, as opposed to solitary nests, which are mostly occupied
519 by adults. Communal nesting also reduces time needed for nest construction, so that
520 one individual may use more than one nest within its home range (Lee and Cockburn,
521 1985). In this sense, in our studied area, we observed winter movement of torpid
522 animals between nests; however, further analysis is needed to determine the relevance
523 of communal nesting to sociality. Some authors suggest that the decline of large and old
524 trees, as a result of logging may affect population abundance and winter survival of
525 many arboreal mammals (Laurance, 1996; Lindenmayer et al., 1997) and forest
526 specialist birds (Diaz et al., 2005; Cornelius, 2007), due to resource limitation (as is the
527 case of the arboreal marsupial *Hemibeldeus lemuroids* in Australian tropical forest and
528 *Aphrastura spinicauda*, a secondary-cavity forest bird in Chilean temperate rain forest;
529 Laurance, 1996; Cornelius, 2007). Nevertheless, we found no evidence of a reduction in
530 population abundance of *Dromiciops* residents in SL forests in Chiloé Island. This result
531 suggests that the type of logging practiced in rural forest patches has not negatively
532 affected nesting sites of *D. gliroides*; however, demographic consequences need to be
533 assessed (see Chapter 2).

534 As other small marsupials, *D. gliroides* is omnivorous. Arthropods, particularly
535 Coleoptera and Orthoptera, were common items in the diet, but fruits were also an
536 important component of the animals' diet (Armesto et al., 1987; Amico et al., 2009). *D.*
537 *gliroides* showed high consumption of a diversity of fleshy fruits, particularly in OG
538 forest, some of which had no dispersal agent known (Table 5). Hence, *D. gliroides* may

539 have a key role in forest patches, especially when dispersing seeds of epiphytes and
540 vines (Rodríguez-Cabal et al., 2007; Amico et al., 2009). Among mammals, *D. gliroides*
541 is responsible for the dispersal of up to 24% of all fleshy-fruited species in Chiloé Island
542 forests (J. L. Celis, unpublished data). Considering that in temperate rainforests of
543 Chiloé Island, about 70% of the woody flora bears fleshy fruits (Armesto and Rozzi,
544 1989), animal frugivores such as *D. gliroides* may facilitate the processes leading to
545 seed dispersal and forest regeneration. The fruit-eating behavior of *D. gliroides* is
546 exacerbated in summer, coinciding with the fruiting peak of many fleshy-fruited forest
547 species (Smith-Ramírez and Armesto, 1994; Riveros and Smith-Ramírez, 1996). High
548 arthropod abundance in Chiloé forests during spring-summer (Díaz, 2008) becomes
549 particularly important for *D. gliroides* during the first months following hibernation, both
550 in SL and OG forest patches. We propose that a higher proportion of arthropods in the
551 diet of *D. gliroides* detected in SL forests can be related to greater bamboo cover in the
552 understory of open canopy stands (Pérez et al., 2009). Bamboos sustain a great
553 abundance of some arthropod groups (Reid et al., 2004). Alternatively, our data support
554 the hypothesis of Amico et al. (2009), which suggests that the higher incidence of
555 arthropods in *Dromiciops* diet in logged forests could be related to a reduction in the
556 local abundances of epiphytes and vines after logging.

557 Although higher parasitism has been generally attributed to greater population
558 stress and low habitat quality (Daszak et al., 1999), higher prevalence of ectoparasites
559 in *D. gliroides* populations (55%) at SL forests compared to OG forests did not have
560 noticeable effects on the animal's health condition, as there were no differences in body
561 mass, recapture rate, or population density between habitats. This relict marsupial is the

562 main host for *I. neuquenensis*, the only tick ever recorded on *D. gliroides* (Marin-Vial et
563 al., 2007) and likely, the biological vector of a recently documented blood parasite of
564 this species (Merino et al., 2009). Health consequences of these vector-parasite-host
565 interactions in different forested habitats of northern Chiloé Island are still unknown.

566 In short, knowledge of the population density, behavior, and food habits of *D.*
567 *gliroides* and other forest specialist species that use remnant forest patches in rural
568 landscapes and lowland areas of the Chilean Lake District (Castellon and Sieving,
569 2006; Fontúrbel et al., 2009; Smith-Ramírez et al., ms), is urgently needed in the
570 context of regional conservation of biodiversity outside protected areas, and also
571 considering a rapidly changing rural scenario, due to unabated rates of forest cover
572 loss, fragmentation and reduction of old-growth forest habitats, and climate change
573 (Armesto et al., 1998; Echeverría et al., 2007; Jaña et al., 2007; Armesto et al., 2009).
574 On the positive side, maintenance of *D. gliroides* populations in rural forest patches has
575 been documented under current selective logging practices, which means that under
576 regulated management of rural forest patches, conservation of forest dwelling species
577 with strict habitat requirements is possible. Consequently, our report provides valuable
578 guidance for *D. gliroides* conservation and for planning the sustainable management of
579 rural areas.

580

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594

594 **FIGURE LEGENDS**

595

596 **Figure 1.** Satellite image (gray scale) of the rural landscape of northern Chiloé Island
597 (Landsat 2001). Darker areas indicate wooded habitats, and lighter areas indicate
598 shrublands and agricultural lands. Circles indicate studied sites where Old-growth forest
599 (OGF), Selective-logged forest (SLD) and ESS are found.

600

601 **Figure 2.** Different habitats where *Dromiciops* was studied in a rural landscape of
602 northern Chiloé. A and B = Old Growth forest, C and D = Selective Logged forest and D
603 and E = shrubland.

604

605 **Figure 3.** Population density (based on mnka) of *Dromiciops gliroides* in the studied
606 sites in northern Chiloe Island. (A) Densities in different sites SD = Senda Darwin OG
607 forest; Caulín = Caulin OG forest; LL = Llanquihue SL forest; QU = Quilar SL forest;
608 ESS1 = Senda Darwin shrubland 1; ESS2 = Senda Darwin shrubland 2, and (B)
609 Densities in different habitat types. OFG = Old-growth forest; SLF = Selective logged
610 forest and ESS= Shrublands.

611

612 **Figure 4.** Example of home range overlaps among two adult males (horizontal lines)
613 and four adult females (vertical lines) at Senda Darwin OGF in Chiloé Island.

614

615 **Figure 5.** Characterization of the trophic niche of *Dromiciops gliroides* in forest habitats
616 and during spring and summer, from the three-year of data in northern Chiloé Island.

617

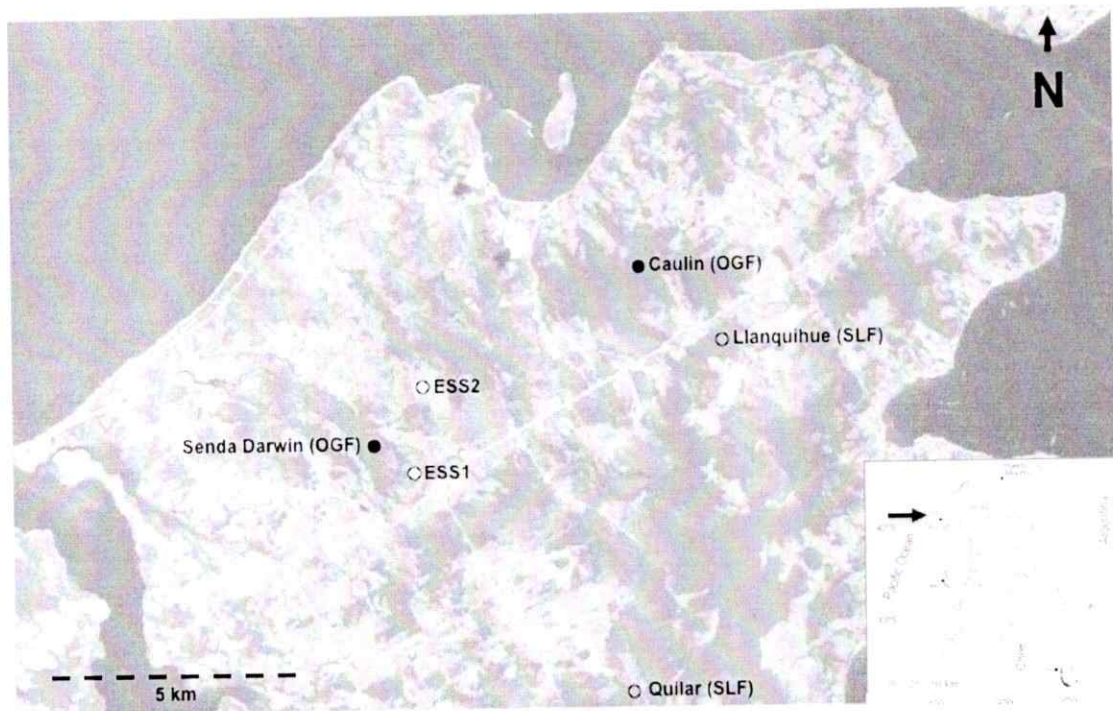
618 **Figure 6.** Mean (± 1 SE) plasma testosterone and progesterone concentration (ng/ml)
619 in blood samples from 44 males (black) and 19 females (white) of *D. gliroides* from
620 pooled data for all rural forest sites in northern Chiloé Island.

621

622 **Appendix 1.** Monthly climatic records from Senda Darwin Biological Station for the
623 studied periods (2005 – 2009).

624

624 **Figure 1.**

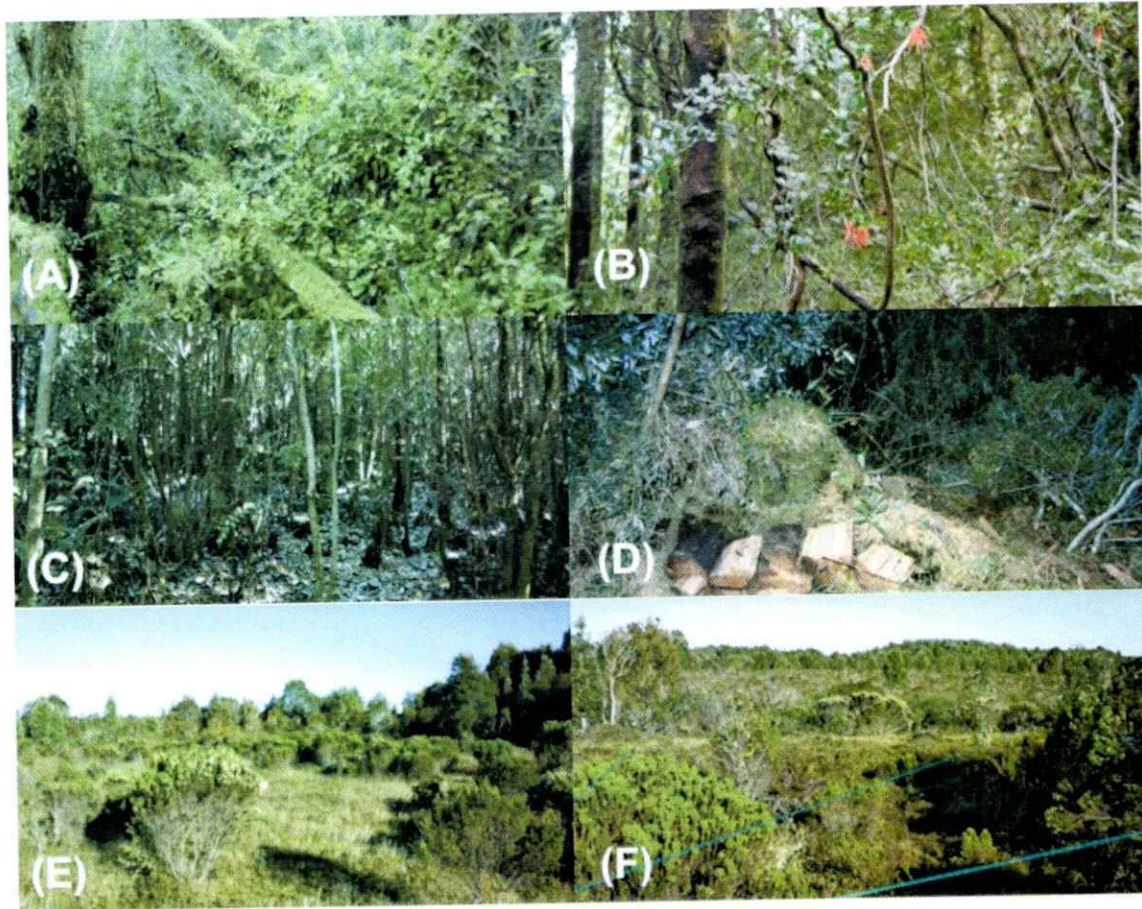


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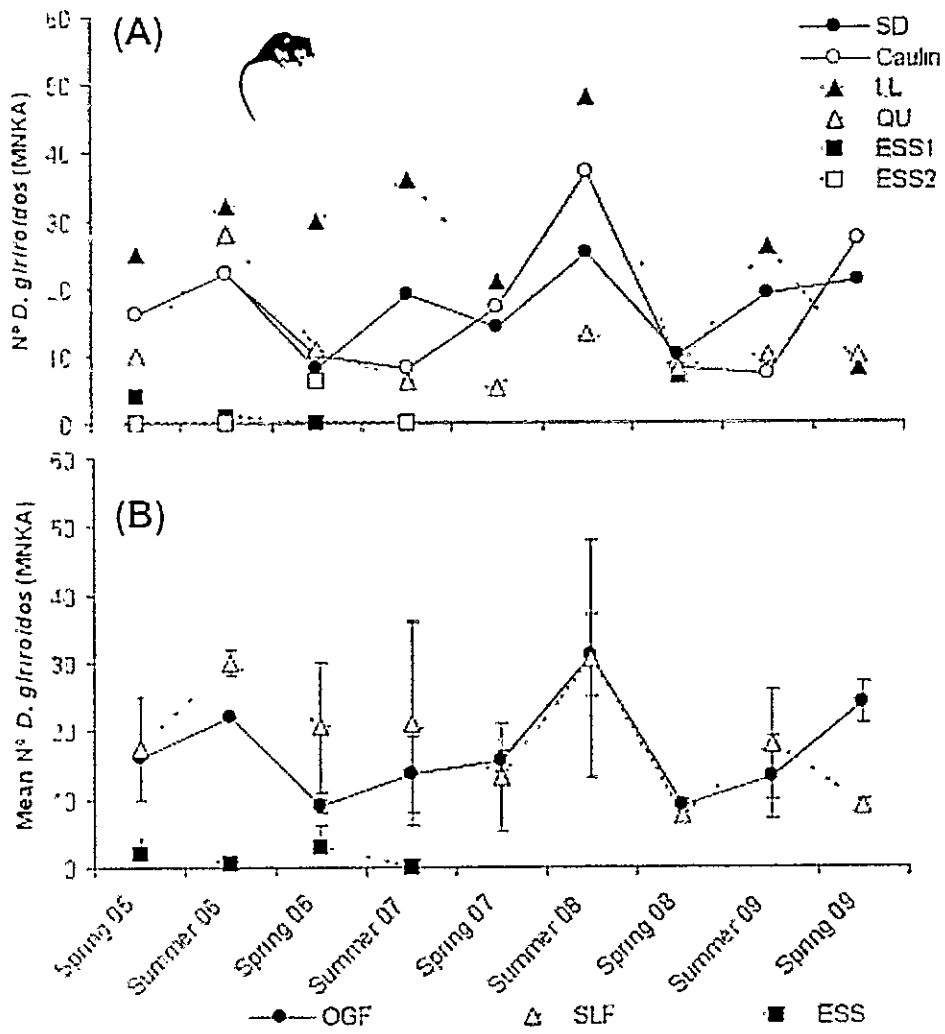
627 **Figure 2.**



628

629

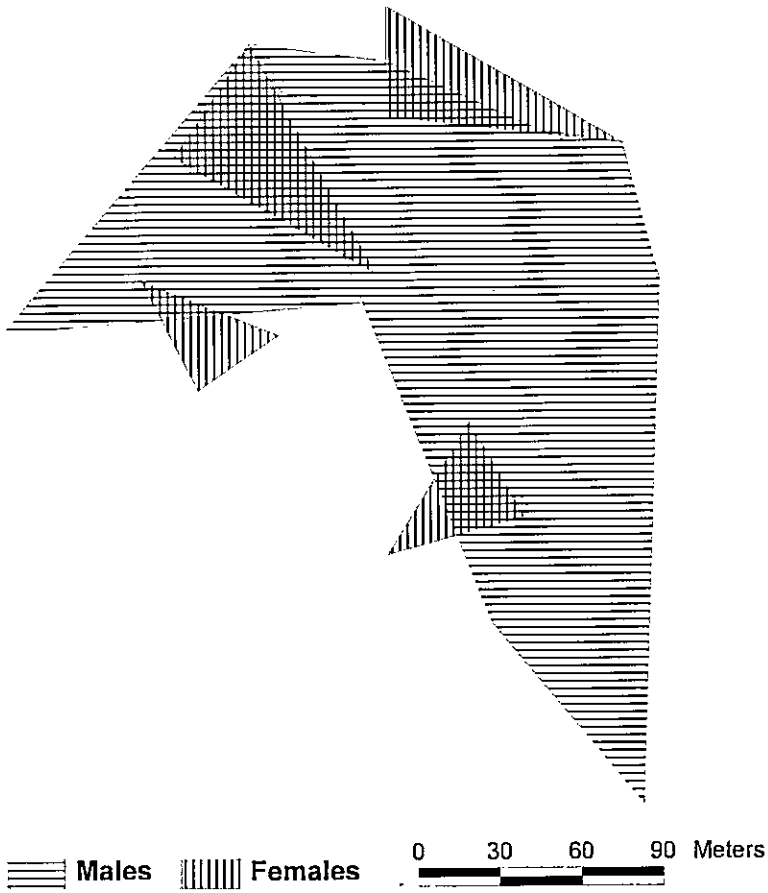
629 **Figure 3.**



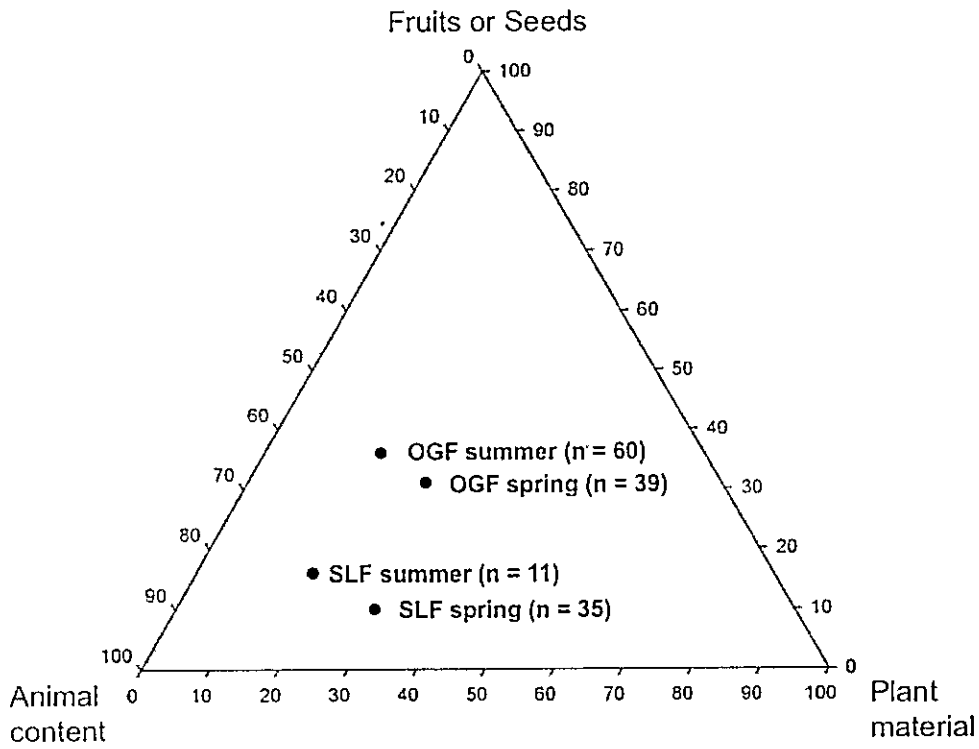
630

631

631 **Figure 4.**
632



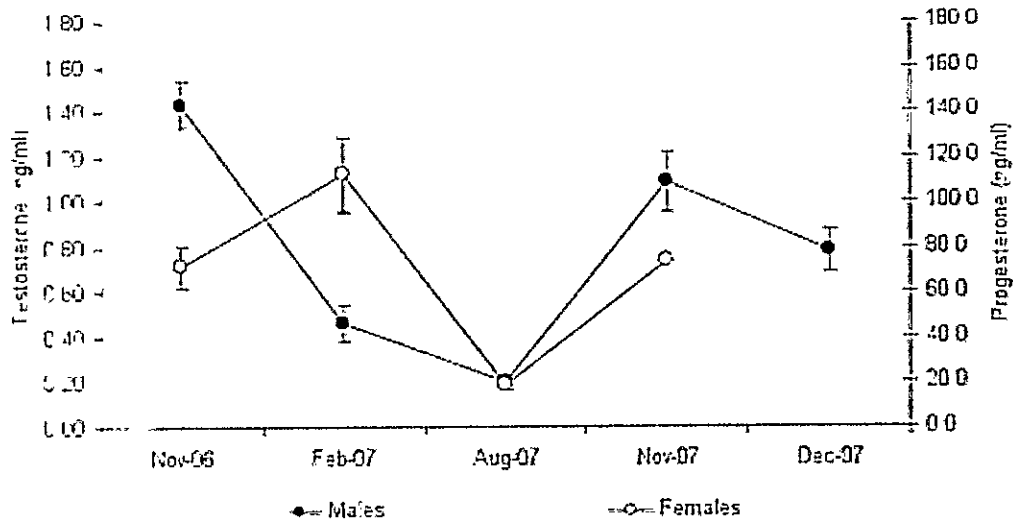
632 **Figure 5.**



633

634

634 **Figure 6.**



635

636 **Table 1.** Number of captured (MNKA) and recaptured individuals of *D. gliroides* from spring 2005 to spring 2009 according
 637 to sex and age categories in all study sites (forests and shrublands) in the rural environment of northern Chiloé Island.

Season	OG forest													
	Senda Darwin						Caulín							
	mnka	recaptures	dead	Males	Females	Pregnant females	Juvenile s	mnka	recaptures	dead	Males	Females	Pregnant females	juveniles
Spring 05	16	12	1	12	5	4	0	16	2	0	12	3	1	1
Summer 06	22	27	1	8	3	0	12	22	7	0	6	6	0	10
Spring 06	8	2	0	5	3	0	0	10	7	2	6	2	0	4
Summer 07	19	19	1	7	2	0	11	8	2	0	2	4	0	2
Spring 07	14	13	1	11	3	2	0	17	12	0	12	5	3	0
Summer 08	25	33	1	7	4	0	15	37	41	2	11	10	0	17
Spring 08	10	5	0	9	1	0	0	8	0	0	5	3	0	0
Summer 09	19	6	0	5	2	2	12	7	1	0	1	1	0	5
Spring 09	21	27	0	15	7	0	0	27	13	0	20	6	0	1

638

639

639 Table 1. cont.

Season	SL forest										Quillar			
	Llanquihue					Adult					Adult			
	minka	recaptures	dead	Males	Females	Pregnant females	juveniles	minka	recaptures	dead	Males	Females	Pregnant females	Juvenile s
Spring 05	25	11	2	15	12	0	0	10	5	1	5	6	5	0
Summer 06	32	16	3	4	6	0	25	28	8	0	7	5	0	16
Spring 06	30	24	3	20	8	0	5	11	10	3	9	1	0	3
Summer 07	36	27	1	10	13	0	13	6	9	1	0	3	0	4
Spring 07	21	12	1	14	8	8	0	5	2	0	4	1	1	0
Summer 08	48	23	0	7	6	0	35	13	20	0	3	2	0	8
Spring 08	7	1	0	4	3	2	0	8	4	0	7	1	0	0
Summer 09	26	2	0	6	6	4	14	10	1	0	3	1	0	6
Spring 09	8	5	2	6	4	0	0	10	5	0	7	3	0	0

640

641 Table 1. cont.

Season	ESS										ESS2			
	ESS1					Adult					Adult			
	minka	recaptures	dead	Males	Females	Pregnant females	juveniles	minka	recaptures	dead	Males	Females	Pregnant females	Juvenile s
Spring 05	4	0	0	1	2	1	1	0	0	0	0	0	0	0
Summer 06	1	0	0	0	0	0	1	0	0	0	0	0	0	0
Spring 06	0	0	0	0	0	0	0	6	0	0	3	3	0	0
Summer 07	0	0	0	0	0	0	0	0	0	0	0	0	0	0

642

642 **Table 2.** Mean (\pm 1 SE) body mass (g) of *D. gliroides* during spring and summer, and differences between sexes and ages
 643 classes from 2005-2009 in different forest and shrubland habitats of northern Chiloe Island. nd = no data.

Habitat	Forest	Season	Sex											
			Male						Female					
			adult	SE	n	juvenile	SE	n	adult	SE	n	juvenile	SE	n
OGF	Senda Darwin	Spring	26.5	0.7	35	n/d	n/d	n/d	25.8	1.1	13	n/d	n/d	n/d
		Summer	29.1	1.2	12	18.0	0.7	34	31.1	1.4	9	18.9	1.0	16
	Caulin	Spring	26.3	0.6	46	17.5	4.1	1	24.9	1.2	11	20.3	2.4	3
		Summer	29.6	1.6	7	17.0	0.9	20	31.2	1.4	9	17.3	1.1	13
SLF	Llanquihue	Spring	27.0	0.6	42	27	4.1	1	28.4	0.9	19	20.3	2.1	4
		Summer	29.0	1.3	10	18.8	0.6	53	31.4	1.0	17	19.5	0.7	31
	Quilar	Spring	28.1	0.9	21	n/d	n/d	n/d	28.3	1.3	10	20.3	2.4	3
		Summer	28.1	1.5	8	19.9	0.8	27	37.4	1.6	7	18.4	1.7	6
ESS	ESS1	Spring	28.0	4.1	1	n/d	n/d	n/d	27.8	2.9	2	n/d	n/d	n/d
		Summer	n/d	n/d	n/d	22.5	4.1	1	n/d	n/d	n/d	n/d	n/d	
	ESS2	Spring	28.8	2.4	3	n/d	n/d	n/d	36.3	2.4	3	n/d	n/d	n/d
		Summer	n/d	n/d	n/d	n/d	n/d	n/d	n/d	n/d	n/d	n/d	n/d	n/d

644

645 **Table 3.** Mean home range areas (ha) for adult male and female *D. gliroides* in
 646 forested habitats of northern Chiloé Island. OGF = Old growth forest patches and
 647 SLF = Selective logged forest patches.

Habitat	Forest	Sex					
		Male	SE	n	Female	SE	N
OGF	Senda Darwin	0.35	0.05	15	0.13	0.06	9
	Caulin	0.12	0.06	11	0.15	0.06	9
	<i>Mean</i>	<i>0.26</i>	<i>0.04</i>	<i>26</i>	<i>0.14</i>	<i>0.05</i>	<i>18</i>
SLF	Quilar	0.18	0.07	8	0.23	0.09	5
	Llanquihue	0.27	0.06	12	0.12	0.06	11
	<i>Mean</i>	<i>0.23</i>	<i>0.05</i>	<i>20</i>	<i>0.15</i>	<i>0.05</i>	<i>16</i>
	<i>Total mean</i>	<i>0.26</i>	<i>0.03</i>	<i>46</i>	<i>0.14</i>	<i>0.04</i>	<i>80</i>

649 **Table 4.** Relative frequency of occurrence (FO%) of arthropods in feces of
650 *Dromiciops gliroides* in rural forest habitats of northern Chiloé Island.

651

Arthropod order	FO % (n = 126)
<i>Coleoptera</i>	42.9
<i>Orthoptera</i>	24.6
<i>Lepidoptera</i>	11.9
<i>Hymenoptera</i>	2.4
<i>Hemiptera</i>	1.6
<i>Aranea</i>	0.8
<i>Acari</i>	1.6
Unidentified	14.3

652

653

653 **Table 5.** Relative frequency of occurrence (FO%), and relative abundance (%) of
 654 seeds over the total sample, and total number of seeds of 12 different fleshy-fruited
 655 forest species found in feces (n = 146) of *Dromiciops gliroides* in rural forest
 656 habitats of northern Chiloé Island. Number of fruits eaten (in parentheses) is
 657 estimated from number of seeds per fruit (Armesto et al., 1987) and personal
 658 observations*. na = not applicable. Known dispersal agent extracted from Armesto
 659 et al. (1987).

660

	Growth form	FO %	Abundance % (n = 359)	No. seeds (fruits)	Known dispersal agent
<i>Tristerix corymbosus</i>	Hemiparasite	43.08	25.07	90 (90)	<i>D. gliroides</i> – Birds
<i>Gaultheria insane</i>	Shrub	18.46	13.93	50 (na)	Unknown
<i>Gaultheria mucronata.</i>	Shrub	10.77	15.32	55 (6)	Birds
<i>Luzuriaga polyphylla</i>	Vine	7.69	4.46	16 (16)	Unknown
<i>Sarmienta repens</i>	Hollo-epiphyte	6.15	5.85	21 (na)	Unknown
<i>Mitraria coccinea</i>	Vine-Epiphyte	3.08	3.34	12 (1)	Unknown
<i>Griselinia racemosa</i>	Vine	3.08	5.29	19 (19)	Unknown
<i>Asteranthera ovata</i>	Vine-Epiphyte	1.54	2.23	8 (1)	Unknown
<i>Raukaua laetevirens</i>	Hemi-epiphyte	1.54	1.67	6 (2)	Unknown
<i>Myrteola numularia</i>	Shrub	1.54	22.28	80 (6)	Birds
<i>Philesia magellanica</i>	Shrub, vine	1.54	0.28	1 (1)	Birds
<i>Amomyrtus luma</i>	Tree	1.54	0.28	1 (1)	Birds

661

661 **Table 6.** Number of adult and juvenile individuals with and without ectoparasites in
 662 forest sites of rural environments in northern Chiloé Island.

663

Habitat	Forest	With		Without ectoparasites		Total
		Adults	Juveniles	Adults	Juveniles	
OGF	Senda Darwin	10	6	10	7	67
	Caulin	10	4	15	5	
	<i>Total OGF</i>	20	10	25	12	
SLF	Quillar	11	11	7	4	110
	Llanquihue	27	20	21	9	
	<i>Total SLF</i>	38	31	28	13	
ESS	ESS1	0	1	2	0	8
	ESS2	2	0	3	0	
	<i>Total ESS</i>	2	1	5	0	
<i>Total</i>		60	42	58	25	

664

665

665 **Table 7.** Mean mass (\pm 1 SE) and number of individuals of *D. gliroides* found
 666 nesting in artificial nest boxes in rural forest patches of northern Chiloé Island.

667

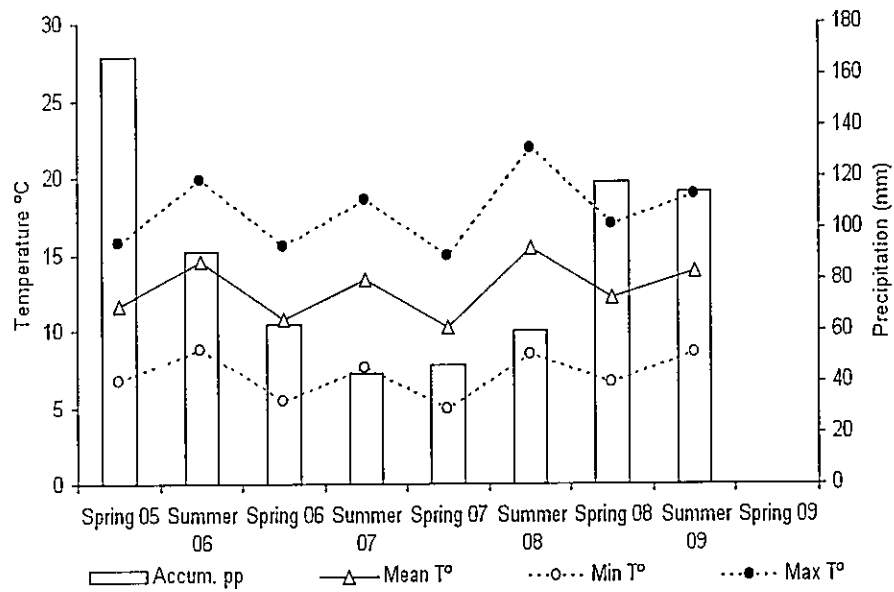
Weight	Solitary				Communal			
	Males		Females		Males		Females	
	Adults	juveniles	adults	juveniles	adults	juveniles	adults	juveniles
Mean	34.67	28	32.2	27.67	32.44	26.88	34.56	26.98
n	12	1	5	3	51	16	24	27
SE	1.82	Na	1.55	0.44	0.85	0.75	1.04	0.54

668

669

669 **Appendix 1. Monthly climatic records from Senda Darwin Biological Station for the**
 670 **studied periods (2005 – 2009).**

671



672

673

1 **CAPITULO 2**

2

3

4

5

6

7 **DEMOGRAPHY OF THE ARBOREAL MARSUPIAL *DROMICIOPS GLIROIDES***
8 **IN OLD GROWTH AND SELECTIVELY LOGGED FORESTS OF CHILOÉ**
9 **ISLAND, CHILE**

10

11

12

13

13 **ABSTRACT**

14 Demographic consequences of selective logging on the endemic arboreal
15 marsupial Monito del monte (*Dromiciops gliroides*) were assessed in forest patches
16 in a rural landscape of northeastern Chiloé Island (42 S), Chile. Two old growth
17 (OG) and two selectively logged (SL) temperate rainforest fragments were studied
18 as representative of remnant, forested areas, where populations of the relict and
19 threatened *D. gliroides* occur. We expected that habitat changes due to logging
20 could have negative effects on *D. gliroides* demography. Individuals were live-
21 trapped for four consecutive years (2005–2009) during the mating (November) and
22 non-mating (February) seasons. Differences in forest structure and food resources,
23 such as arthropods and fruits, were measured between OG and SL forests.
24 Demographic parameters were modeled using mark-recapture procedures. OG
25 forests had a higher density of larger canopy trees and snags, and more fleshy-
26 fruited epiphytes than SL forests. Cover of the understory native bamboo
27 (*Chusquea* sp.) was higher in SL than in OG forests and higher arthropod
28 abundances were associated with bamboo cover. Population sizes of *Dromiciops*
29 varied seasonally but did not differ between the two forest habitats. Demographic
30 parameters (e.g., monthly survival and recruitment) were not different between
31 habitat types, but survival varied across years, and recruitment between seasons
32 each year. Yearly population growth rate of *Dromiciops* was similar between
33 logged and unlogged forests but showed inter-annual and forest fragment
34 variation. In conclusion, small-scale, selective logging of remnant OG forest
35 patches in northern Chiloé does not translate into demographical consequences for
36 this arboreal marsupial, despite some changes in habitat structure. The

37 maintenance of understory bamboo cover is a key element for the persistence of
38 the population, by providing nesting materials, protection from predators, and
39 arthropod food resources. This result constitutes an important finding to support
40 the conservation of this forest-dependent relict species in anthropogenic rural
41 landscapes.

42

43

43 INTRODUCTION

44 Forestry practices such as logging can have profound implications for forest
45 structure and wildlife habitat (Lindenmayer et al., 1990; Newton, 1994;
46 Lindenmayer et al., 1997), depending on the degree of impact, along a gradient
47 from clear-cutting to selective cutting of single trees (Emmingham et al., 1992;
48 Laurance, 1996; Sullivan et al., 1999; Gagne et al., 1999). In anthropogenic
49 landscapes, such as rural environments, forest-dependent species are especially
50 susceptible to the often-invisible process of forest degradation, where forest area
51 remains unchanged but there is a significant loss of forest structures and habitat
52 quality due to firewood extraction, cattle grazing, local fires, and logging of remnant
53 forest patches (Lambert et al., 2005; Sekercioglu et al., 2007). Depending on
54 logging intensity, selective logging scenarios can substantially alter forest structure
55 and resources for wildlife. However, most damage to vegetation is incidental to the
56 forestry practices; thus the net effect is not "selective" removal of large canopy
57 trees of a few timber species, but a broader impact across a nearly random cross-
58 section of trees, understory, shrubs, epiphytes, vines (Johns 1988) as well as
59 terrestrial and flying arthropods (Lambert et al., 2005). Arboreal mammal
60 populations may be strongly influenced by changes in structure and composition of
61 forests (Laurance, 1996; Wilson and Friend, 1999; Ecke et al., 2002; Simard and
62 Fryxell, 2003; Lee, 2004; Renner et al., 2006; Isaac et al., 2008). For example,
63 dead or over-mature trees which are characteristic of primary forest are a crucial
64 source of den cavities for many arboreal vertebrates (Lindenmayer et al., 1997;
65 Gibbs et al., 1999, Gibson and Lindenmayer, 2002). Alternatively, openings

66 created within logged stands may provide new niches that could potentially
67 increase biodiversity.

68 Habitat quality is often defined in relation to population density and resource
69 availability. Moreover, population fluctuations are the result of demographic
70 processes (Julliard et al., 1999) and the loss of habitat quality such as forest
71 heterogeneity and resources due to logging could have long-term consequences,
72 which are translated into demographic parameters rather than current population
73 abundances (Mahan et al., 1996; Martin and Handasyde, 2006; Isaac et al., 2008).

74 In some cases, population density estimates are not adequate indicators of
75 habitat quality (Van Horne, 1983; Pulliam, 1988; Brawn & Robinson, 1996).
76 Consequently, habitat quality should be assessed directly by examining
77 relationships between population survival or growth rate and variation in habitat
78 characteristics (Loeb, 1999; Manning and Edge, 2004). However, few studies have
79 investigated demographic responses to habitat degradation as a way of
80 understanding the consequences of logging on forest-dwelling species.

81 Forest cover loss and fragmentation has been intense in South American
82 temperate rain forests during the past 100 years (Willson and Armesto, 1996;
83 Armesto et al., 1998; Jaña-Prado et al., 2006; Echeverria et al., 2007). Old-growth
84 forests in lowland areas of Chiloé Island occupy today less than one third of its
85 original distribution (Armesto et al., 2009) and consequently, Chile belongs to the
86 group of countries with biodiversity hotspots that have increased overall
87 deforestation rate from 1.02% during the 1980s to 1.76% during the 1990s (Jha
88 and Bawa, 2006). Deforestation has continued to drive the loss of old-growth forest
89 cover and greatly altered the landscape of this temperate forest region (Willson

90 and Armesto, 1996; Echeverria et al., 2007; Armesto et al., 2009). It has been
91 estimated that only 5% of logging of native forests is based on controlled
92 silvicultural practices (Lara, 1996). In Chile, most forests have been clear cut or
93 burned and replaced by exotic forestry plantations. However, in many rural areas
94 remnant forests are usually subjected to selective logging, where individual trees of
95 high timber value or high quality for firewood are removed, leaving large clearings
96 in the canopy. Given the frequency of traditional selective logging by rural
97 inhabitants and by small forestry operations, it is relevant to assess the impacts of
98 such practices on habitat quality for forest dwelling species (Nystrand and
99 Granstrom, 2000).

100 *Dromiciops gliroides* (Microbiotheria), a threatened arboreal marsupial
101 (IUCN 2008) is endemic to South American temperate rainforests (Herskhovitz,
102 1999), and has been frequently cited as an inhabitant of extensive old-growth
103 forest patches (see chapter 1; Mann, 1978; Kelt and Martinez, 1989; Rodriguez-
104 Cabal et al., 2007, 2008; Fontúrbel and Jiménez, 2009). Recently, this arboreal
105 species has been recorded with similar densities in secondary-growth forest as in
106 old-growth forest patches (see chapter 1; Patterson et al., 1990; Fonturbel et al.,
107 2009), and therefore it is interesting to know the consequences of habitat
108 degradation due to logging on the long-term demography of *D. gliroides*. This
109 should help assess the possible trends in *D. gliroides* populations of remnant forest
110 patches in rural landscapes, as habitat quality may deteriorate over time leading to
111 population declines. Here, we report a four-year study of *D. gliroides* populations in
112 remnant forests of rural northern Chiloé Island, which differ in structure due to
113 logging.

114 Our specific objectives were: (1) To characterize and compare the structure
115 of old-growth (OG) and selectively logged (SL) forest stands, as habitats for *D.*
116 *gliroides*, (2) to estimate *D. gliroides* population size in each habitat, and (3) to
117 estimate the main demographic parameters (*i.e.* survival, recruitment and
118 population growth rates) reflecting the status of *D. gliroides* populations in SL and
119 unlogged forest patches in northern Chiloé. On the basis of previous reports, we
120 expected to find similar densities in both forest types, but higher survival and
121 recruitment rates in OG than in SL forest habitats, because the former should
122 provide the best-quality habitat in terms of forest structure and availability of food
123 resources for *D. gliroides*. Considering *D. gliroides* hibernation behavior, the
124 greater abundance of hollowed trees (Diaz et al., 2005) and presumably the
125 greater abundance of fleshy-fruited epiphytes and vines in OG forests, results in
126 better habitat quality for *D. gliroides* than logged forests. Alternatively, SL forests
127 may provide opportunities for higher survival of *D. gliroides* due to the expansion of
128 native bamboo cover and associated arthropods that can serve as protective
129 habitat and resources for *D. gliroides*.

130

131 MATERIAL AND METHODS

132 *Study sites.* — The study was conducted in four medium size (> 50 ha) forest
133 fragments, representing two different levels of logging in northeastern Chiloé
134 Island, Ancud County, Chiloe province (41° 50' S, 73° 36' W; Fig. 1). The rural
135 landscape mosaic has about 40% of forest fragments, including both old growth
136 and secondary forest patches, embedded in an anthropogenic matrix of grazing
137 pastures, agricultural fields, and dense humid shrublands (Willson and Armesto

138 1996; Aravena et al., 2002; Díaz et al., 2005). We sampled two selectively logged
139 and two un-harvested forest patches, because of they represented two contrasting
140 habitat for *D. gliroides*, due to their different vegetation structures. The patches
141 selected were separated from each other by at least 5 km. Old-growth (OG) forest
142 fragments were within Senda Darwin Biological Station and in Fundo Cisnes at
143 Caulín (Fig.1), had no signs of human impact, and had large emergent trees in
144 their canopies. Selectively-logged (SL) forests were at Llanquihue and El Quilar,
145 where landowners had harvested valuable and relatively large timber trees with no
146 defined silvicultural system (see Cornelius 2007). Thus, we had two replicates for
147 logged and two for unlogged patches.

148 The two OG forest remnants were about 300 years old, based on tree-ring
149 counts of dominant canopy trees, and had no visible signs of recent logging or fire
150 (Aravena et al., 2002; Gutiérrez et al., 2004). These OG patches were
151 characterized by a multi-layered canopy, with large emergent trees of *Nothofagus*
152 *nitida* (Nothofagaceae), *Drimys winteri* (Winteraceae), *Podocarpus nubigena* and
153 *Saxegothaea conspicua* (both Podocarpaceae), plus few individuals of *Eucryphia*
154 *cordifolia* (Eucryphiaceae) and *Caldcluvia paniculata* (Cunoniaceae) in the lower
155 canopy. Trees were often covered with vines and epiphytes such as ferns, mosses
156 and bromeliads (Muñoz et al., 2003). Large amounts of dead biomass (snags and
157 logs) were frequent within OG stands. The understory had a high density of tree
158 saplings, few shrubs, and dense thickets of 2-3 m tall native bamboo (*Chusquea*
159 spp.) within tree-fall gaps (Gutiérrez et al., 2004).

160 SL forest patches are frequent in the rural landscape, due to the widespread
161 practice of small-scale harvesting, which generally results in an open canopy forest

162 with remnant large *Nothofagus nitida*, which are left standing. The patches
163 sampled had areas with a closed canopy of dense juvenile trees, 10–20 m tall,
164 predominantly shade-intolerant *D. winteri*, and *Amomyrtus luma*. In SL forests,
165 open areas with a dense cover of native bamboos, and a sparse presence of tree
166 seedlings and saplings due to shading, were frequent (Aravena et al., 2002). Both
167 SL patches endured limited logging at the end of summer during the study period
168 (Quillar in 2007 and Llanquihue in 2008); nevertheless, the two sites showed no
169 evidence of cattle impact or fire.

170

171 *Habitat structure and food resources.* — At each forest fragment thirty 10-m radius
172 circular plots were established. Each plot was assigned randomly to a trapping
173 station within each 100 m radial line (see below). We divided habitat variables into
174 three groups: habitat structures, vines, and resources. At each plot we counted or
175 measured the following: (a) The number of live trees, snags and fallen logs with a
176 DBH (diameter at breast height) >30 cm as a proxy for availability of potential
177 hollowed-trees or potential nest sites for secondary-cavity nesters (Newton, 1994;
178 Escobar et al., 2002; Cornelius 2007). This arbitrary cut-off diameter was defined
179 considering the smallest diameter of holes used by *D. gliroides* to built nests
180 (unpublished data), and was consistent with dimensions of nest-boxes used by *D.*
181 *gliroides*. (b) Within a 2-m radius circle at the center of each vegetation plot, we
182 measured native bamboo cover (*Chusquea* spp.), canopy, and non-bamboo
183 understory cover (i.e., shrubs and sapling). These different covers were estimated
184 visually as the percentage of ground covered by vegetation above 10 m (canopy)
185 and between 0,5 and 3 m (understory), assigning each point to one of five cover

186 classes: 0%, 1–25%, 26–50%, 51–75% and 75–100%. (c) As plant resource
187 variables we measured: presence of fleshy-fruited vines and epiphytes consumed
188 and probably dispersed by *D. gliroides* (i.e., *Tristerix corymbosus*, *Luzuriaga*
189 *poliphylla*, *Mitraria coccinea*, *Asteranthera ovata*, *Sarmienta repens*, *Gaultheria*
190 *insana*; see chapter 1; Amico et al., 2009). (d) Abundance of arthropods (>5 mm
191 long), was estimated as the number trapped by with 2 m height interception traps,
192 in one half of the circular plot at each forest fragment. Variables (a) and (b) are
193 descriptors of habitat structures while (c) and (d) are rough estimators of food
194 resources used by this arboreal marsupial. Differences between logged and OG
195 forests in habitat and resource variables were analyzed using a permutational
196 multivariate analysis of variance through PERMANOVA software (version 1.6
197 2005; Anderson, 2005). Data were previously transformed using $\log(x+1)$ for
198 habitat variables and presence/absence distance matrix for occurrence of fleshy-
199 fruited epiphytes (Anderson, 2005).

200

201 *Animal trapping.* — *Dromiciops gliroides* at each forest patch were studied from
202 spring 2005 to spring 2009 (four consecutive years). To estimate demographic
203 parameters, trapping was conducted over 10 consecutive nights in two seasons
204 each year: (a) in mid-spring (November-December), after animals became active
205 following winter torpor and before the onset of the breeding season, and (b) late
206 austral summer (February-March), after the reproductive season ended. Trapping
207 webs were established in both OG and SL forests, consisting of twelve radial lines
208 of 100 m long. Each trapping line had 12 stations; the first four were at 5 m
209 intervals from the center (5, 10, 15, and 20 m), and the following eight were at 10

210 m intervals (30, 40, 50, 60, 70, 80, 90, and 100 m). With this web design, 144 traps
211 were set in each forest patch covering 3.46 ha (Parmenter et al., 2003). We used
212 Tomahawk-like traps (baited with banana) fixed on tree branches between a
213 minimum height of 1 m and maximum of 3 m. Traps were checked daily between
214 06:00 and 08:00 hours. For all captured animals we recorded sex, age, breeding
215 status (based on scrotum size, pouch morphology and mammary gland
216 development), and body mass, using a Pesola ® (Baar, Switzerland) spring
217 balance, 0.1 g precision. In late summer, animal weight from adult individuals was
218 compared between sexes and both forest patches, because before entering in
219 winter torpor, weight can be a good proxy of habitat quality through health status.
220 To avoid pseudo replication we considered body mass for only new unmarked
221 adult individuals. Statistical analyses were performed using two-way ANOVA in
222 Statistica software (version 6.0, 2001, Statsoft Inc.).

223

224 *Model selection and demographic parameters.* — Population parameters were
225 estimated for OG and SL forests by capture-mark-recapture (CMR) procedures
226 (Lebreton et al., 1992). Our data consisted of captured-recaptured history from
227 nine primary occasions. We combined closed population models to estimate
228 population size within each trapping session (10 days per trapping period as
229 secondary sessions), with open population models to estimate survival between
230 primary sessions (Quental et al., 2001). Time between primary sessions was three
231 months in summer and nine months in winter. Survival estimation was
232 standardized to one month following Yoccoz et al. (1998). Population abundance
233 was estimated by software DENSITY (version 4.1, 2007; Efford et al., 2004), using

234 a Jackknife estimator that does not assume homogeneity of capture probabilities
235 among individuals (Burnham and Overton, 1979). This method was adopted
236 because, including heterogeneity in model selection we obtain a better adjustment
237 ($\Delta AIC > 2$) because many individuals were caught just once ($n = 260$ from a total
238 of 494) and few were caught 5 times or more ($n = 43$). Relative population
239 densities were estimated as the ratio of population abundance to the effective area
240 estimated as the trapping web area plus a buffer area of a width equal to half of the
241 largest recapture distance (i.e., MDA; Otis et al., 1978; Parmenter et al., 2003;
242 Fontúrbel et al., 2009). Because we expected differences in population densities
243 between seasons (see chapter 1), we performed a two-way ANOVA in Statistica
244 software (version 6.0, 2001, Statsoft Inc.), to test differences in population
245 densities between habitat types and seasons.

246 A Goodness of fit test for the general model was applied with U-Care
247 software (version 2.02; Choquet et al., 2005), searching for transients, as an
248 estimate of migration, and trap shy-happy individuals.

249 For all demographic parameters, we constructed a model set to test the
250 hypothesis of difference among forest patch sites (i.e., Senda Darwin, Caulin,
251 Llanquihue, Quilar), between OG and SL forests, between seasons (summer vs.
252 spring), and among years (from 2005 to 2009). Capture probabilities were also
253 tested for spatial and temporal variation. Survival (ϕ) and recruitment rates ($1-\gamma$)
254 were analyzed by CMR statistical modeling based on Akaike's information criterion
255 (AIC; Lebreton et al., 1992) using M-SURGE software (version 1.8.5, 2008;
256 Choquet et al., 2004).

257 We selected models with the lowest AIC ($\Delta AIC > 2$); otherwise models were
258 considered statistically identical (Burnham and Anderson, 1998), in which case we
259 selected the model with the smallest number of parameters, assuming a
260 parsimonious criterion (Crespin et al., 2006).

261 Recruitment rate was estimated with the method developed by Pradel
262 (1996), "reading" the capture histories backwards (reverse-time capture-recapture
263 methods). This makes it possible to estimate the probability that an individual
264 caught at time t_{+1} was present in the population at time t . These probabilities are
265 called seniority probabilities (γ ; Pradel, 1996) and represent a fraction of the
266 population. Seniority probabilities can be used to estimate other related
267 demographic parameters, such as population growth rate (λ) (Lima et al., 2001).

268 Reproductive rate was estimated from adult females as a fraction from
269 reproductively active individuals (pregnant or lactating) at both habitat types.

270 Finally, population growth rate (Lambda: λ) was estimated using Pradel's
271 estimator through MARK software (version 5.1; White and Burnham, 1999). For
272 this analysis we considered only one primary capture session per year in mid-
273 spring (i.e., in November).

274

275 RESULTS

276 *Habitat structures.* — OG and SL forests differed markedly in: (a) number of large
277 tree, snags, canopy cover, woody understory cover, and bamboo cover (ANOVA
278 $F_{1, 119} = 54,44$, $P < 0.001$), (b) food resources such as the presence of fleshy-
279 fruited vines and epiphytes (ANOVA $F_{1, 119} = 14,99$, $P < 0.001$), and (c) arthropods

280 per unit of understory vegetation (ANOVA $F_{1, 59} = 4,22$, $P = 0.015$). The number of
281 large trees and snags was greater in OG than in SL forests (Table 1). Understory
282 cover of shrubs and tree saplings was also higher in OG than in SL forests, while
283 bamboo cover was larger in the more open SL than in OG forest patches (Table 1).
284 Forest canopy cover also differed between forests, being denser in SL than in OG
285 patches (Table 1).

286 The frequency and species composition of fleshy-fruited epiphytes and
287 vines also differed between logged and unharvested forests. More fleshy-fruited
288 vines were found in OG forests. Finally, we recorded higher abundances of
289 arthropods at SL forests (Table 1). Arthropod abundances were positively and
290 significantly correlated with volume of understory bamboo in both logged and
291 unlogged stands (Spearman rank correlation $R = 0.20$, $n = 60$, $P = 0.025$).

292

293 *Abundance and population parameters of Dromiciops gliroides*— From spring
294 2005 to summer 2009, we recorded a total of 1054 captures, representing 494
295 different individuals in OG and SL forests combined. A total of 119 animals were
296 trapped at Senda Darwin, 111 at Caulín (both being OG forests), while 181 were
297 trapped at Llanquihue and 83 at Quilar (SL forests). On average, each animal was
298 captured 2.13 times. In mid-late summer, there was no significant difference in
299 adult body mass between-habitat types (ANOVA $F_{1, 75} = 0.16$, $P = 0.69$), but we did
300 detect marginal differences between both sexes (ANOVA $F_{1, 75} = 3.81$, $P = 0.054$).

301 Overall, mean population densities were not statistically different between
302 OG and SL forests (i.e., OGF mean = $6.18 \pm 1 \text{ SE} = 1.26$, $n = 9$; SLF mean = $8.8 \pm$

303 1 SE = 1.26, n = 9; GLM W = 3.15, P = 0.07; Fig. 2), however, we observed a slight
304 trend towards higher values in SL forests. Population density varied seasonally in
305 all habitats (Fig. 2). In both forested habitats (i.e., OGF and SLF), we observed
306 higher *Dromiciops* densities in the austral summer (February), which corresponded
307 to the juvenile recruitment period, than in the previous spring (November), when
308 mating and breeding took place (summer mean = $9.73 \pm \text{SE} = 2.09$, n = 8; SLF
309 mean = $5.3 \pm \text{SE} = 1.33$; GLM W = 10.6, P = 0.01; Fig. 2).

310 The highest and lowest *Dromiciops* densities were observed at SL forests
311 with a maximum of 20.9 individuals/ha in Quilar (e.g., summer of 2006) and a
312 minimum of 1.5 individuals/ha in the same forest patch (e.g., summer and spring of
313 2007; Fig. 2). In spring 2008, we recorded a drastic reduction in population in all
314 forest patches.

315

316 *Model selection and demographic parameter estimation.* — Based on the AIC
317 criterion, neither logging nor season had significant effects on survival rates of
318 *Dromiciops* populations ($\Delta \text{AICc} < 2$; Table 2). However, we did detect an inter-
319 annual variation in survival (Table 2). With the exception of 2007, we observed a
320 trend towards higher survival rates during summer than in winter, yielding no
321 overall difference between seasons. We detect a small proportion of transient
322 individuals in Senda Darwin OG forest. Testing model selection assumptions, no
323 effect on survival was detected for the general model. Capture probabilities were
324 significantly different between each studied sessions, but neither selective logging
325 nor season had significant effects (Table 2).

326 As expected, recruitment rates ($1 - \gamma$) showed strong seasonal variability with
327 higher values in summer (monthly recruitment = 0.33) than in spring (monthly
328 recruitment = 0.02; Table 3). Neither timber harvesting nor year had significant
329 effects on recruitment (Table 3). Reproductive rate, as the proportion of
330 reproductive females, was not different between both habitat types (0.17 in OG vs.
331 0.22 in SL forests; $\chi^2 = 0.4$, $df = 1$, $P = 0.53$).

332 The estimation of annual population growth rate (λ) ranged from 0.8 to 1.11
333 in a given forest fragment with a mean of 1.15 and 0.97 at OG and SL forest,
334 respectively. Annual population growth rate did not differ between OGF and SLF
335 (Table 3). However it did differ between each forest stand with the lowest value at
336 Llanquihue SL forest (Table 3). Overall annual population growth rates were $1.11 \pm$
337 0.09 , 1.19 ± 0.09 , 1.10 ± 0.12 and 0.83 ± 0.06 for Senda Darwin, Caulin, Quilar
338 and Llanquihue, respectively (Table 3), indicating slow population growth rate
339 during the four years of study.

340

341 **DISCUSSION**

342 Most studies on habitat loss have focused on landscape-level processes, such as
343 patch size, the spatial configuration of patches, and dispersal patterns of
344 organisms, paying less attention to habitat degradation within fragments (Harrison
345 and Bruna, 1999, Jaña-Prado et al., 2006).

346 Selective logging usually modifies food-resource availability, and shape life
347 history traits of forest-dwelling species (see chapter 1; Wheatley et al., 2002; Isaac
348 2005; Herbers & Klenner 2007) through changes in vegetation structure of remnant

349 forest patches (Pérez et al., 2009). This traditional logging practice by landowners
350 typically target large trees of selected timber species (Aravena et al., 2002;
351 Lambert et al., 2005; Diaz et al., 2005; Pérez et al., 2009), thus reducing the
352 abundance of tree holes, snags and understory trees. As a consequence of the
353 opening of the canopy, SL forests present higher cover of the understory bamboo,
354 *Chusquea* sp. (Pérez et al., 2009), especially under canopy openings created
355 within logged stands. These canopy gaps provide new opportunities for bird
356 species that can take advantage of the protection of bamboo cover, such as
357 tapaculos (Reid et al., 2004) and the Des Murs' Wiretail (Díaz et al., 2006). In this
358 regard, higher arthropods abundances documented for SL forests, associated with
359 denser understory bamboo cover or woody debris (Niemelä et al., 1993; Reid et
360 al., 2004), translate into increases in the frequency of arthropods in the diet of *D.*
361 *gliroides* at the expense of fleshy fruits, in comparison with individual from OG
362 forest (see chapter 1). Increased arthropod abundance represents an important
363 food resource for Chilean marsupials because of their higher energy content (Lima
364 et al., 2001) (see chapter 1; Meserve et al. 1988), as well as for specialized
365 understory birds (Reid et al., 2004; Diaz et al., 2005; Díaz 2008). In addition
366 bamboo cover also provides effective protection against natural predators for both
367 birds and mammals (Manning and Edge, 2004; Reid et al., 2004).

368 This dietary shift at SL forest could have consequences for seed dispersal
369 by *D. gliroides* of many vines and epiphytes, with potential cascading effects
370 throughout the rest of the plant community (see chapter 1; Armesto et al., 1987;
371 Rodriguez-Cabal et al., 2007; Salinas, 2008; Amico et al., 2009). In summary, our
372 results, documented that the consumption of fleshy fruits vs. arthropods (see

373 chapter 1), was related to the relative abundances of these food resources in each
374 forest type, SL vs. OG.

375 Considering that *D. gliroides* like other arboreal mammals, depends on
376 natural cavities in large canopy trees for shelter, hibernation and reproduction, and
377 that selective logging reduces the availability of cavities in the stand (Lindenmayer
378 et al., 1997; Cornelius, 2007), we expected a reduction in winter survival rate that
379 could translate into overall population survival, as has been documented for many
380 other arboreal mammal species (Lindenmayer et al., 1997; Pilastro et al., 2003),
381 and for the forest bird *Aphrastura spinicauda* in the same study area (Cornelius,
382 2007). However, despite a reduction in the density of old trees in SL forests
383 compared to OG forests, winter survival probabilities remained the same in both
384 types of forest over four years. One possible explanation is that the denser
385 bamboo cover developed after tree harvesting in the understory of SL stands may
386 compensate for the lack of tree holes. It is known that bamboo foliage and culms
387 are frequently used by *D. gliroides* to construct woven spherical nests, which
388 protect the animals against unfavorable weather conditions (Mann, 1978, Patterson
389 et al., 1990). In this regard, more than 80% of nests found in artificial nest boxes
390 (see chapter 1) had bamboo foliage as construction material. Moreover, although
391 hibernation is energetically challenging, we did not found differences in survival
392 between winter and summer seasons (Schaub & Vaterlaus-Schlegel, 2001;
393 Pilastro et al., 2003).

394 *Dromiciops gliroides* populations have often been thought to depend greatly
395 on the amount of land covered by OG forest patches along its geographical
396 distribution (see chapter 1, Mann, 1978; Kelt and Martínez, 1989; Hershkowitz,

397 1999; Rodriguez-Cabal et al., 2008). Most published evidence indicates that
398 arboreal mammals, and especially nocturnal marsupials in Australia, are highly
399 susceptible to logging damage (McNab, 1978; Johns, 1985, 1986; Laurance, 1990;
400 Lindenmayer et al., 1997) or logging intensity (Herbers & Klenner 2007) due to
401 reductions in population density and/or survival. Nevertheless, our results show
402 that OG and SL forests have similar population densities, and there is even a trend
403 toward higher densities at SL stands considering all sampling dates. Other works
404 have also shown no differences in *Dromiciops* densities between forest patches
405 with different vegetation structures such as OG and secondary forests or tree-fall
406 gaps (see chapter 1; Patterson et al., 1989; Fontúrbel et al., 2009; Smith-Ramirez
407 et al., ms). Variable densities in SL forests could be attributable to differences in
408 logging intensity (Fig. 2). For example, Laurance (1996) recorded higher
409 abundance of green ringtail possums in logged forests, while others arboreal
410 marsupials did not differ significantly between logged and unlogged forests.
411 Moreover, harvesting intensity was the more important determinant of squirrel
412 density (Herbers & Klenner 2007).

413 *Dromiciops* densities in logged and unlogged forest patches in the rural
414 landscape of Chiloé were considerable lower (overall mean density = 7.3
415 individual/ha) than mean densities from Andean populations in OG forests of
416 mainland Chile (ca. 19 individual/ha) (Fontúrbel et al., 2009). However, it is
417 important to note serious time-restrictions (one season or one year of sampling)
418 and methodological shortcomings (see Parmenter et al., 2003; small sample size
419 and less powerful trap spatial arrangements) of previous studies (see Rodriguez-
420 Cabal et al., 2008; Fontúrbel et al., 2009).

421 Habitat selection models predict that higher quality habitats will be preferred
422 over those of poor quality (Fretwell and Lucas, 1970). We initially hypothesized that
423 OG forests had higher habitat quality than logged forests. However, our results
424 show that *D. gliroides* densities did not differ between logged and unlogged stands,
425 which did not translate into differences in demographic parameters and population
426 growth rate. We propose that bamboo cover and arthropod abundances are key
427 structural elements and food resources respectively for the maintenance of viable
428 *D. gliroides* populations and for some understory-specialist birds (Willson and
429 Armesto, 2003), making SL forests relatively high quality habitats. Telleria et al.
430 (1991), Sullivan and Sullivan (2001) and Ecke et al. (2002) have also documented
431 higher abundance of small mammals in remnant forests in anthropogenic
432 landscapes. Moreover, in some temperate rainforests of South America, rodent
433 species can be more abundant in forest remnants compared large protected areas
434 due to increased availability of food resources and reduced predator abundance
435 (Saavedra and Simonetti, 2005). Coincidentally, we also found that the abundance
436 of *Irenomys tarsalis*, a nocturnal arboreal rodent, was higher in SL forest (JL Celis-
437 Diez, personal observation). Therefore, forests subjected to limited logging may
438 also serve as relatively high quality habitats for some small mammal species, in
439 contrast to large vertebrate predators (Sergio et al., 2005).

440 Because top predators are particularly sensitive to habitat disruption (Sergio
441 et al., 2005), we expect that they exert stronger negative pressure on prey survival
442 in OG stands. The main predators of *D. gliroides* in temperate rainforests are
443 Rufous-legged owls (*Strix rufipes*) and Kodkod cats (*Leopardus guigna*). Both
444 species are especially vulnerable to habitat degradation and are less commonly

445 found in open SL forests, small forest fragments and exotic pine plantations
446 (Martinez and Jaksic, 1996; Sanderson et al., 2002; Acosta-Jamett and Smonetti
447 2004). Lower predator densities in SL forests should result in higher survival of
448 *Dromiciops* in these stands, compared to OG stands, where predators remain
449 present, as a possible explanation of higher population densities at least in one SL
450 forest patches. However, a predator release hypothesis (Ferguson and Fox, 1984)
451 to explain the higher survival of *Dromiciops* in SL stands has to be rejected,
452 because we did not observe differences in survival rates of between OG and SL
453 forests.

454 This work is the longest demographic study of *D. gliroides* in South America
455 (four years; see Rodriguez-Cabal et al., 2008 for a previous record), but it is not yet
456 sufficient to provide reliable documentation of population dynamic and to predict
457 the fate of *D. gliroides* in remnant forests in anthropogenic landscapes. Annual
458 variation in survival has been documented to depend on food resources (Cezilly et
459 al., 1996; Lahti et al., 1998; Gauthier et al., 2001) and in temperate rain forests of
460 Chiloé, arthropods abundances vary according to season and years (Díaz 2008).

461 *Dromiciops* populations both in logged and unlogged forests did not seem to
462 be declining, and because of the limited time series, we could not ascertain
463 whether their population dynamics is regulated by denso-dependent or denso-
464 independent factors (Lima et al., 2001). Indeed, we suggest that the drastic
465 population drop recorded in all forest patches in the spring 2008 was possibly a
466 consequence of the severe drought of the summer 2007-2008. This highlights the

467 need to expand long-term demographic studies of this and other forest species
468 (Carmona et al., ms)

469 With our results, we can be certain that populations of the threatened
470 arboreal marsupial are not so rare as previously thought (Kelt, 2000), and that the
471 species can be found in both logged and unlogged forest remnants in the rural
472 landscape, despite the severe loss of forest cover in the recent decades
473 (Echeverria et al. 2007). Further, the impact of non-industrial logging, when limited
474 to the removal of few harvestable large trees from the forest, and the ensuing
475 proliferation of bamboo, does not harm the populations of this arboreal marsupial,
476 and enhances the potential for the conservation of biodiversity in rural areas under
477 productive management of remnant forests and OG forest patches.

478

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492 Chiloé, Chile.

493

493 **FIGURE LEGENDS**

494

495 **Figure 1.** Satellite image (gray scale) of the rural landscape of northern Chiloé
496 Island (Landsat 2001). Darker areas indicate wooded habitats, and lighter areas
497 indicate shrublands and agricultural land use. Circles indicate studied sites where
498 OGF and SLD are found.

499

500 **Figure 2.** Population density of the arboreal *Dromiciops gliroides* in the studied
501 area in northern Chiloe Island. (A) Mean densities (± 1 SE) of animals in both
502 habitat types. Circles OGF = Old-growth forest and triangles SLF = Selective
503 logged forest, and (B) Mean densities (± 1 SE) of animals in all studied sites, SD =
504 Senda Darwin OG forest; Caulín = Caulin OG forest; LL = Llanquihue SL forest;
505 QU = Quilar SL forest.

506

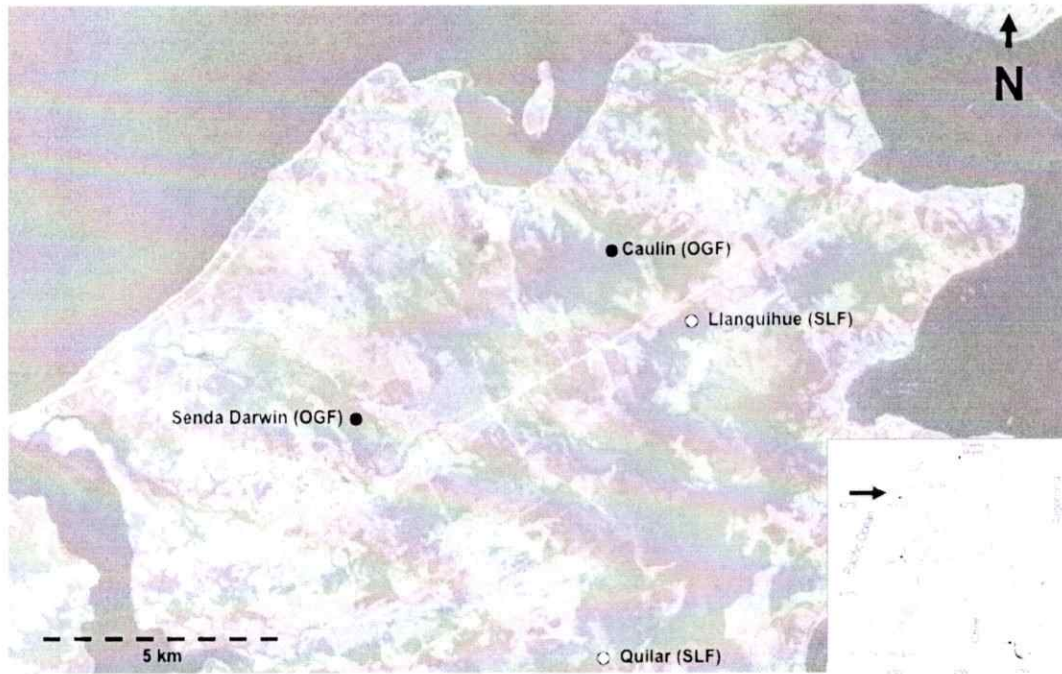
507 **Appendix 2.** Number of live captures (line, not including recaptures) and
508 proportion of adult males, females and juveniles (bars) in forest fragments in a rural
509 landscape of northern Chiloé Island. OGF sampled at Senda Darwin (a) and
510 Caulín; (b) SLF sampled at Quilar (c) and Llanquihue (d).

511

512

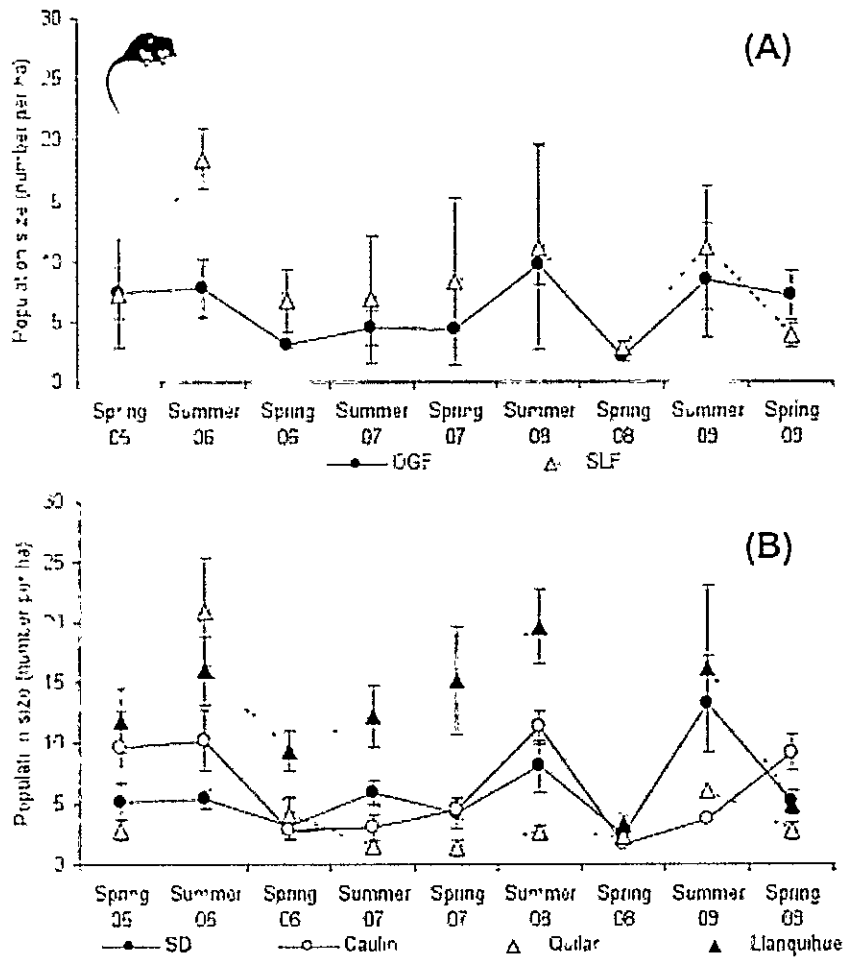
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513 **Figure 1.**



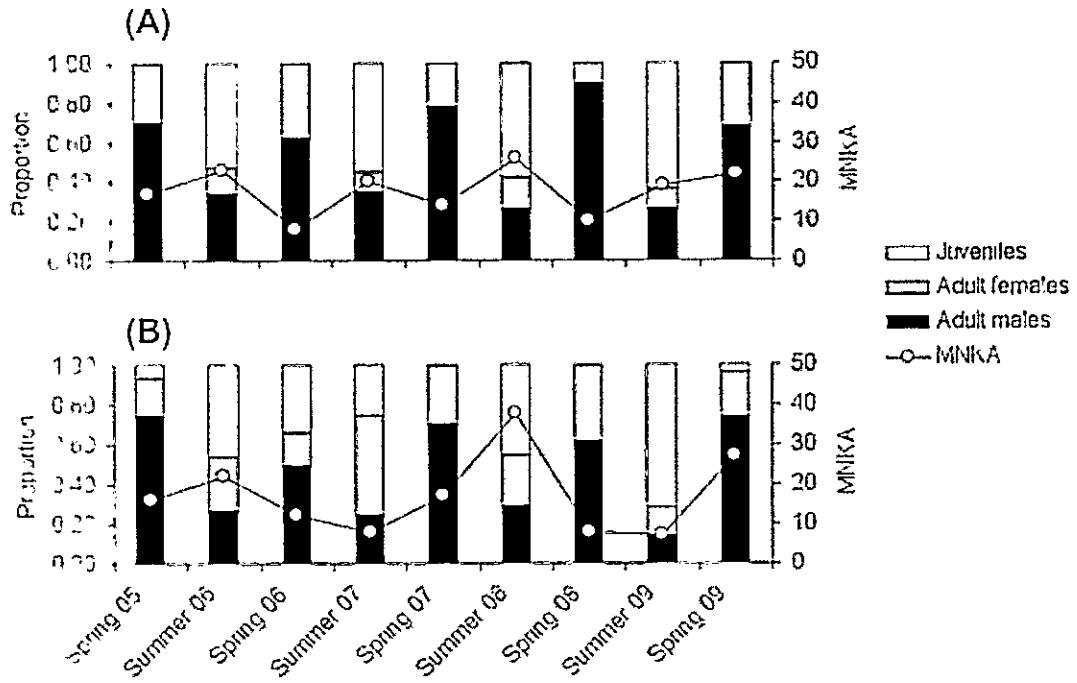
514

515 **Figure 2.**
516



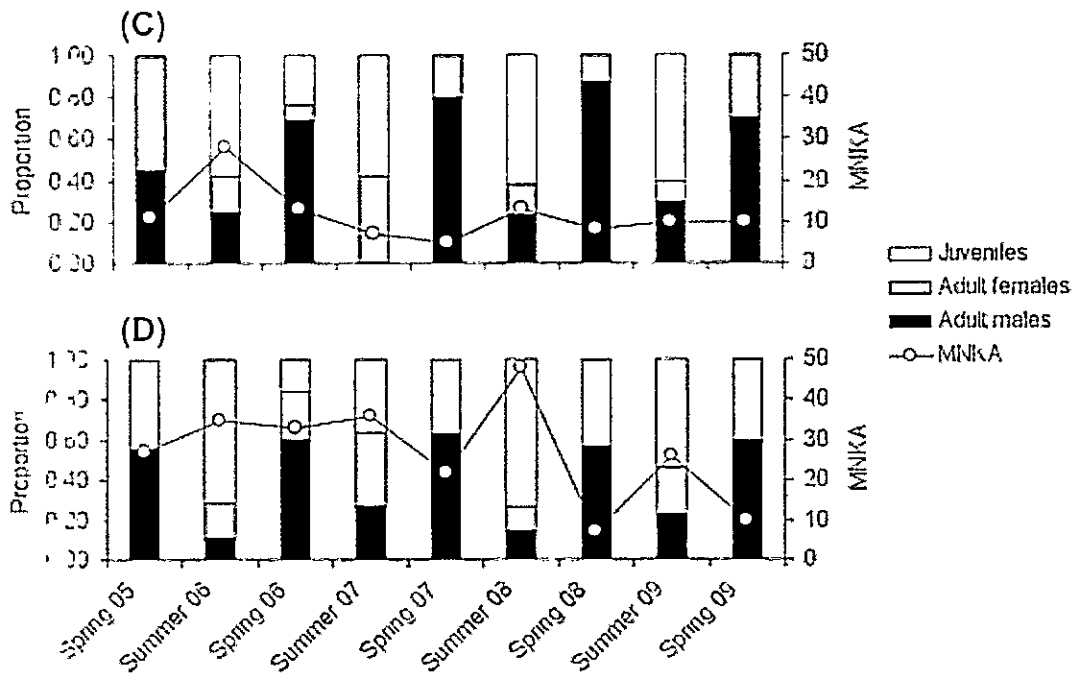
517

518 Appendix 2a.
519



520 **Appendix 2b.**

521



522 **Table 1.** Vegetation structure and food resources characteristics for both studied habitat types. OGF: Old growth
 523 forest; SLF: Selective logged forest; dbh: diameter at the breast height; Tc: *Tristerix corymbosus*; Lp: *Luzuriaga*
 524 *poliphylla*; Gi: *Gaultheria insana*; Mc: *Mitraria coccinea*; Sr: *Sarmienta repens*; Ao: *Astheranthera ovata*; (*) P <
 525 0.05.

526

Habitat	Mean number (per plot)			Frequency of occurrence (n = 60)							Median category		
	No. trees (dbh > 50 cm (*)	No. snags (*)	No. logs arthropods/tra p (*)	Mean number	Tc (*)	Lp	Gi (*)	Mc (*)	Sr (*)	Ao (*)	% Canopy cover	% Understor y cover (*)	% bamboo cover (*)
OGF	3.80	0.50	2.08	2.43	0.42	1	0.32	0.13	0.27	0.47	25	75	50
SLF	1.45	0.25	1.70	5.40	0.07	0.98	0.07	0.45	0.07	0.25	50	25	75

527

527 **Table 2.** The statistical model tested denoted according to each model-specific variation
 528 in survival (Φ) and capture probability (p). i denotes constant parameter; time denotes fulltime
 529 specific variation; forest denote habitat type (e.g., OGF vs. SLF); site denotes all studied forest sites (e.g.,
 530 Senda Darwin, Caulin, Llanquihue and Quilar); season denotes seasonal time-specific variation (e.g.,
 531 summer vs. winter) and year denotes between year variation (e.g., 2005-2009). AICc is the estimated
 532 Akaike's information criterion: models with lower values and marked with asterisk indicate more
 533 parsimonious models. *ne* indicate not estimated parameter.

534

Model	AICc	# Parameters	Deviance	Monthly parameter values (for the significant model)
Capture (p)				
$\Phi i p time$	884.89 (*)	9	866.63	p1 = 0.32; p2 = 0.60; p3 = 0.41; p4 = 0.82; p5 = 0.65; p6 = 0.37; p7 = 0.38; p8 = 0.64
$\Phi i p season$	893.63	3	887.59	
$\Phi i p i$	896.68	2	892.66	
$\Phi i p year$	901.03	5	890.94	
Survival (Φ)				
$\Phi year p time$	879.17 (*)	12	854.70	Φ 2005- 2006 = 0.85
$\Phi time p time$	881.05	15	850.33	Φ 2006- 2007 = 0.89
$\Phi forest+year p time$	881.24	13	854.70	Φ 2007- 2008 = 0.85
$\Phi season p time$	882.93	10	862.61	Φ 2008- 2009 = <i>ne</i>
$\Phi forest.year p time$	883.49	16	850.68	
$\Phi forest+season p time$	883.89	11	861.50	
$\Phi i p time$	884.89	9	866.63	
$\Phi forest p time$	885.76	10	865.44	
$\Phi site p time$	887.78	12	863.31	

535 **Table 3.** The statistical model tested denoted according to each model-specific variation
 536 in seniority (γ) (recruitment rate = $1-\gamma$) and population growth rate ($\text{Lambda} = \lambda$). *i* denotes
 537 constant parameter; *time* denotes fulltime specific variation; *forest* denote habitat type (e.g., OGF vs.
 538 SLF); *site* denotes all studied forest sites (e.g., Senda Darwin, Caulin, Llanquihue and Quilar); *season*
 539 denotes seasonal time-specific variation (e.g., summer vs. winter) and *year* denotes between year
 540 variation (e.g., 2005-2009). AICc is the estimated Akaike's information criterion: models with lower values
 541 and marked with asterisk indicate more parsimonious models. *ne* indicate not estimated parameter.
 542

Model	AICc	# Parameters	Deviance	Monthly parameter values (for the significant model)
Seniority (γ) (recruitment rate = $1-\gamma$)				
γ <i>season p time</i>	889.4	10	869.05	γ spring = 0.98
γ <i>fores+season p time</i>	891.4	11	869.02	γ summer = 0.67
γ <i>time p time</i>	896.2	15	865.5	
γ <i>year p time</i>	943.4	12	918.9	
γ <i>i p time</i>	944.9	9	926.60	
γ <i>forest p time</i>	946.9	10	926.52	
Growth rate (λ)				
λ <i>site</i>	989.59	9	58.6	λ SD = 1.11 ± 0.09
λ <i>forest</i>	992.68	8	63.8	λ Caulin = 1.19 ± 0.09
λ <i>forest+time</i>	995.9	13	56.3	λ Quilar = 1.10 ± 0.12
λ <i>constant</i>	998.7	7	71.9	λ Llanquihue = 0.83 ± 0.06
λ <i>time</i>	1002.3	9	71.3	

543

CONCLUSION

Los resultados de esta tesis aportan al conocimiento de la mastozoología de los bosques templados de nuestro país, en particular respecto de una especie de marsupial endémica de los bosque templados de Sudamérica, cuya historia natural, biología, estado de conservación y dinámica poblacional en paisajes antropogénicos habían sido escasamente estudiados. La especie en muchos catálogos nacionales e internacionales se considera aún inadecuadamente conocida (Redford & Eisenberg 1992; Glade 1993; Muñoz-Pedrerros & Palma 2000; www.redlist.org). La mayor parte de los datos poblacionales sobre pequeños mamíferos de los bosques templados del sur de Chile, incluido *Dromiciops gliroides*, ha sido generada prácticamente en tres localidades, La Picada (Puerto Montt), Las Cascadas (Osorno), y Fundo San Martín (Valdivia) (véase revisión en Kelt, 2006), dos de ellas andinas y la última costera, por lo que dado los actuales cambios en el uso del suelo, se hace urgente conocer el estado de las poblaciones en zonas antroponegizadas.

Datos de censos poblacionales de cuatro años (2005-2009), mediante métodos de captura-recaptura, del marsupial arbóreo *Dromiciops gliroides* en parches de bosque antiguos, expuestos a tala selectiva y en matorrales, que conforman los principales tipos de parches del paisaje rural del norte de Chiloé (Echeverría et al., 2007), registran la presencia de individuos en todos los hábitat, a pesar de las grandes diferencias en la estructura de la vegetación y en los tipos y abundancia de recursos tróficos. El principal resultado de este trabajo es que, en comparación con otros micromamíferos del bosque templado, *Dromiciops gliroides* fue más común en el norte de Chiloé de lo que había

sido previamente descrito en otras localidades del sur de Chile (Kelt & Martínez, 1989; Saavedra & Simonetti, 2005; Kelt, 2006). La densidad poblacional fue considerablemente mayor en hábitats de bosque antiguo y sujeto a tala selectiva que en matorrales secundarios; sin embargo no se observaron diferencias en densidad poblacional entre los dos tipos de bosques comparados en este trabajo.

En ambos bosques, independientemente del impacto e la tala, la densidad poblacional mostró una fuerte variación estacional con mayores densidades en verano que en primavera. También hubo variación entre los sitios de bosque sujetos a tala selectiva, atribuible a diferencias en intensidad de tala, lo que es muy difícil de controlar en experimentos naturales. Sin embargo, no se observaron tendencias de declinación del tamaño poblacional en ninguno de los bosques en el período de cuatro años de monitoreo (2005 a 2009). Se podría inferir que en el período de estudio las poblaciones han permanecido relativamente estables con variaciones atribuibles a la fenología de la población (Cezilly et al., 1996).

Por otra parte, a pesar de nuestras predicciones originales, la sobrevivencia y reclutamiento poblacional no difirieron entre las poblaciones de bosques antiguos y expuestos a tala selectiva, observándose también una fuerte variación interanual en la sobrevivencia, sumada a una variación estacional en el reclutamiento. En consecuencia, al combinar estos parámetros demográficos no se registraron diferencias en la tasa anual de crecimiento poblacional de *Dromiciops* en ninguno de los bosques comparados. Dada las actuales tasas de pérdida y degradación del hábitat de bosques remanentes en la zona rural de Chiloé (Echeverría et al., 2007), estos

resultados permiten sugerir que la conservación de parches de bosques remanentes en paisajes rurales pueden sostener poblaciones viables de *D. gliroides* por períodos largos de tiempo, asegurando su viabilidad futura y complementando el esfuerzo de conservación en áreas protegidas.

Considerando que frecuentemente existen diferencias metodológicas asociadas con las estimaciones de densidad poblacional en mamíferos pequeños (Van Horne, 1983; Efford, 1992), el uso de modelos demográficos, más que las estimaciones de densidad per se, es una importante herramienta para evaluar el estado de conservación y tendencias de largo plazo de las poblaciones de especies amenazadas. Debido a que esta metodología demanda censos sistemáticos durante varios años de estudio sucesivos, es necesario implementarlos como parte de un programa de estudios de largo plazo. Mediante este tipo de información es posible entender las consecuencias de las perturbaciones antrópicas del paisaje, incluyendo cambios de cobertura y degradación de los parches, sobre la calidad de hábitats para los taxa de interés.

La mantención de áreas con bosque antiguo, asegura la conservación de esta especie arbórea de marsupial en el paisaje rural, pero la tala selectiva, como explotación local de madera de los bosques, asociada a la remoción selectiva de árboles de mayor tamaño, al promover la presencia de quila, mantiene una adecuada cobertura vegetal en el sotobosque que favorece algunas especies restringidas a fragmentos de bosque tales como *D. gliroides* junto a otras aves del sotobosque (Reid et al., 2004; Castellon and Sieving, 2006a y b)

Nuestros datos demuestran que *Dromiciops gliroides*, al igual que algunas especies de aves del bosque, es un importante vector de semillas de árboles y epífitas de los bosques templados (Armesto et al., 1987; Amico & Aizen, 2000; Rodríguez-Cabal et al., 2007; Amico et al., 2009), de las cuales no existía registro previo de sus dispersantes bióticos (Salinas, 2008). Estos datos amplían el espectro de frutos y semillas consumidos por este marsupial. En el contexto de la trama de interacciones planta-frugívoro en bosques templados del sur de Chile, la disminución o aislamiento de las poblaciones de *Dromiciops gliroides* de los parches de bosque remanentes en la zona rural podría afectar negativamente la regeneración y sobrevivencia de una alta diversidad de especies de bosque (Rodríguez-Cabal et al., 2007).

Considerando el ámbito de hogar de *D. gliroides* se puede inferir los tamaños de parches mínimos o las dimensiones de corredores biológicos que permiten mantener poblaciones viables de esta especie en el ámbito rural. El registro en este estudio, de la presencia de *Dromiciops gliroides* en matorrales secundarios del norte de Chiloé, constituye un importante hallazgo, debido a que este tipo de matorrales no sería una barrera infranqueable al movimiento de individuos de *D. gliroides* entre fragmentos de bosque, favoreciendo la conectividad, a escala de paisaje, entre parches de bosques. También registramos la presencia en las fecas de este marsupial de semillas de *Myrteola numularia*, una especie frecuente en bordes y matorrales, lo que podría indicar que estos matorrales pueden proveer recursos para esta especie durante sus desplazamientos o incluso proponer dinámicas fuente-sumideros entre parches de bosques y matorrales. Al respecto, se ha documentado depredación de nidos de *Aphrastura spinicauda* en matorrales por *D. gliroides* (S. Ippi comunicación personal y

datos no publicados). Sin embargo, faltan estudios más detallados sobre el valor de la matriz como lugar de tránsito entre parches de bosque, para modelar la distribución de especies en distintos escenarios de paisajes rurales o dinámicas metapoblacionales (Umetsu & Pardini, 2007; Umetsu et al., 2008).

En base a la información ecológica generada en esta tesis, y en conjunto con recientes trabajos (Fontúrbel et al. 2009; Smith-Ramírez et al. ms), se propone revisar la clasificación actual de *Dromiciops gliroides*, incluido en la categoría “insuficientemente conocida” propuesta para los vertebrados de Chile (D.S. N° 151, 2007) a “vulnerable”, a nivel nacional, de manera de concordar con la clasificación internacional de “Casi Amenazada” (Díaz & Teta 2008) debido al actual conocimiento de sus densidades poblacionales en Chiloé y en el continente, y la rápida tasa de deforestación y degradación del bosque nativo (Echeverría et al. 2008), que constituye el principal hábitat de esta especie.

Finalmente, en base a los datos de esta tesis, se sugiere considerar las siguientes medidas en el manejo forestal para asegurar la viabilidad de las poblaciones de *D. gliroides* en zonas rurales: (a) mantener, cuando sea posible, los troncos muertos en pie o árboles grandes y viejos sin valor económico en los rodales, (b) evitar el ingreso de ganado al bosque, manteniendo el sotobosque denso y la cobertura de quila en los claros (*Chusquea* spp.), (c) favorecer la presencia de corredores de vegetación nativa arbórea como cercos vivos a través de las praderas, y cobertura arbustiva (matorrales) favoreciendo la conectividad entre fragmentos de bosque o con áreas protegidas (Willson, 2004; Díaz et al., 2005), (d) proteger y monitorear la presencia de

D. gliroides en las áreas con bosques primarios remanentes en ambientes rurales. Con las primeras tres medidas, incorporadas en los requisitos para la obtención de planes de manejo en la nueva legislación forestal (CONAF, 2008) o de producción y protección agrícola (SAG, 1998), se proveería de un paisaje más favorable para la sobrevivencia de especies como *Dromiciops gliroides* y otras restringidas a los fragmentos de bosque, que pueden ser indicadores del impacto del manejo sobre la calidad de los hábitats (Willson & Armesto, 2003; Díaz et al., 2005; Cornelius, 2007).

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SHORT COMMUNICATION

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Presence of *Ixodes neuquenensis* Ringuelet, 1947 (Acari: Ixodidae) on the endangered Neotropical marsupial Monito del monte (*Dromiciops gliroides* Thomas, 1894, Microbiotheria: Microbiotheriidae) at Chiloé Island, Chile

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Abstract The presence of *Ixodes neuquenensis* (Ringuelet, *Notas Mus la Plata* 12:207–216, 1947) (Acari: Ixodidae) parasitizing populations of *Dromiciops gliroides* Thomas, 1894 (Microbiotheria: Microbiotheriidae) at Chiloé Island confirms that this tick species is established in Chile. No preference of the ticks for sex or age of the host was observed.

Keywords Ticks · Marsupials · Ectoparasites · Chile

Introduction

The “Monito del monte”, *Dromiciops gliroides* (Microbiotheria: Microbiotheriidae), is the only living species of the Order Microbiotheria and considered a fossil of the

South American fauna (Hershkovitz 1999). This small and rare mouse opossum is an endemic marsupial of the temperate rainforests of South America which extends along the Pacific rim in Chile and adjacent parts of Argentina and in western Neuquén and southwestern Río Negro provinces. These rainforests hold endemic and endangered flora and fauna (Armesto et al. 1998). In addition, this marsupial has an enigmatic origin, being closely related to Australian marsupials rather than the highly diversified American ones (Kirsh et al. 1991). *D. gliroides* is confined to southern South America, from Talca province (region VII, ca. 35°30') to Chiloé Island (region X, ca. 44°00') in Chile and east to Argentina near Lago Nahuel Huapi (Hershkovitz 1999; Saavedra and Simonetti 2001; Lobos et al. 2005). *D. gliroides* is a tree climber species with nocturnal habits, associated to southern genus *Nothofagus* (Fagaceae) and *Chusquea* bamboo (Jiménez and Rageot 1979). According to IUCN, this species is threatened due to habitat loss, and its population has declined over the previous years (Nowak 1999). Its parasites are also consequently at risk of extinction (Guglielmone et al. 2004).

In continental Chile, *D. gliroides* was found infested with the fleas *Chiliopsylla allophyla* Rothschild 1908 and *Plocopsylla diana* Beaucournu et al. 1986 (Beaucournu and Gallardo 1991, 1992). In Argentina, Ringuelet (1947) presents a description of the holotype female of *Ixodes neuquenensis* on *D. gliroides*, and Guglielmone et al. (2004) recently described the nymph and larvae and re-described the female of *I. neuquenensis*, collected from *D. gliroides* in Llao-Llao Reserve (Río Negro, Argentina). González-Acuña and Guglielmone (2005) previously concluded that, if *I. neuquenensis* was endemic to *D. gliroides*, this parasite–host interaction could also be found in Chile. In this paper, we present information that supports such inference together with additional information on the *D. gliroides*–*I. neuquenensis* relationship at Chiloé Island.

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Materials and methods

The study area is located northeast of Chiloé Island, Ancud city (41°50' S, 73°36' W). This area is characterized by a wet temperate weather with a strong ocean influence. The annual mean temperature is 12°C, with monthly maximum and minimum means of 16°C (January) and 5°C (July), respectively (Aravena et al. 2002). The rainfall amount is 2,000–2,500 mm yearly, with a dry period during January–February (Di Castri and Hajek 1976). The present-day rural landscape is characterized by a mosaic of remnant forest fragments, woodlands, and grazing pastures in rolling hills with altitudes ranging from 50 to 100 m (Aravena et al. 2002).

The physiognomy of these temperate rainforests is characterized by a multi-layer canopy, with large emergent trees such as *Nothofagus nitida* (Nothofagaceae), *Eucryphia cordifolia* (Eucryphiaceae), *Saxegothaea conspicua* (Podocarpaceae), and *Aextoxicon punctatum* (Aextoxicaceae) and large amounts of dead logs biomass and stump covered with epiphytes and vines. The understory is quite abundant, composed of trees, shrubs, and bamboo (Aravena et al. 2002).

Captures of *D. gliroides* were carried out during February to March 2005 at two old-growth forest fragments located in the northeastern area of the island. One of the sites was located inside the private protected area of the "Fundación Senda Darwin" (41° 52' S, 73° 40' W) and the second one was in Fundo "Los Cisnes" located in Bahía Caulín (41° 50' S, 73° 36' W). To capture *D. gliroides* individuals, Tomahawk-like traps (baited with banana) were used as *D. gliroides* are not prone to be caught with Sherman-like traps (Kelt and Martínez 1989). The traps were set up on tree branches at a minimum height of 1 m and disposed in 10×10 grid, with ten lines of 100 m each and 10 m distance between traps. The trapping period was performed in two occasions in both study sites for 15 and 10 nights, respectively (trapping effort=1,500 and 1,000 traps/night, respectively). Each trap was checked daily between 06:00 and 08:00 A.M. To avoid re-sampling, the animals captured were tagged with a 2-mm-long aluminum tag on the right ear, identified according to sex (i.e., female or male) and age (i.e., juvenile or adult), weighed, and finally tick specimens collected. After 10 min of handling, the animals were released at the same capture site.

The site of attachment on the body of the animal was recorded for each tick. All specimens were immediately preserved in 70% alcohol. The description and redescription

of *I. neuquenensis* by Guglielmone et al. (2004) were used to confirm the specific status of the ticks.

Exact Fisher's test was applied to test differences in tick parasitism on *D. gliroides* in relation to sex and age. Tick abundance according to sex and age was also evaluated with a non-parametric ANOVA for ranking data (Sokal and Rohlf 1995).

Results and discussion

Twenty seven (40.3%) from a total of 67 captured *D. gliroides* were infested with *I. neuquenensis* (Table 1). *I. neuquenensis* is the only tick ever reported from *D. gliroides*. A total of 56 larvae, 10 nymphs, and 3 female ticks were collected; 87% of them were observed attached to the ears or nearby. These results confirm the hypothesis proposed by González-Acuña and Guglielmone (2005); *I. neuquenensis* is a tick species present in Chile.

No significant differences were found in tick prevalence according to the sex or age of the hosts (Fisher $P=0.66$). The abundance of *I. neuquenensis* in relation to sex (males vs females $F_{1,63}=0.85$, $P=0.39$) and age (juveniles vs adults $F_{1,63}=0.57$, $P=0.45$) were also non-significant, as well as the interaction between the two mentioned variables ($F_{1,63}=44.25$, $P=0.70$).

These results show that *I. neuquenensis* is quite abundant at the study site but independent of the age and sex of its hosts in contrast to observations in Argentina where ticks were only observed on male hosts (Guglielmone et al. 2004). The male of *I. neuquenensis* remains still unknown. This probably means that this tick species is nidicolous with a non-parasitic male which mates female(s) of the host. An alternative explanation could be that this tick species is parthenogenetic. Further sampling of *D. gliroides*' nests may provide data about the existence of *I. neuquenensis* males. Larvae, nymphs, and female ticks were found at the same time of the year, indicating that this tick needs at least 3 years to complete its life cycle.

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Table 1 Prevalence of infestation of 67 *D. gliroides* by the tick *I. neuquenensis*

Sex	Age								
	Juvenile			Adult			Total		
	Total	Positive	%	Total	Positive	%	Total	Positive	%
Female	17	5	29.4	10	4	40	27	9	33.3
Male	24	10	41.6	16	8	50	40	18	45.0
Total	41	15	36.6	26	12	46.2	67	27	40.3

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ANEXO 2. Artículo publicado en Biological Journal of the Linnean Society (2009)

Biological Journal of the Linnean Society, 2009, **98**, 568–576. With 3 figures

Molecular characterization of an ancient *Hepatozoon* species parasitizing the ‘living fossil’ marsupial ‘Monito del Monte’ *Dromiciops gliroides* from Chile

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The Microbiotheriid *Dromiciops gliroides*, also known as ‘Monito del Monte’, is considered to be a threatened species and the only living representative of this group of South American marsupials. During the last few years, several blood samples from specimens of ‘Monito del Monte’ captured at Chiloé island in Chile have been investigated for blood parasites. Inspection of blood smears detected a *Hepatozoon* species infecting red blood cells. The sequences of DNA fragments corresponding to small subunit ribosomal RNA gene revealed two parasitic lineages belonging to *Hepatozoon* genus. These parasite lineages showed a basal position with respect to *Hepatozoon* species infecting rodents, reptiles, and amphibians but are phylogenetically distinct from *Hepatozoon* species infecting the order Carnivora. In addition, the *Hepatozoon* lineages infecting *D. gliroides* are also different from those infecting other micro-mammals living in sympatry, as well as from some that have been described to infect an Australian species of bandicoot. The potential vector of this parasite appears to be the host-specific tick *Ixodes neuquenensis* because the sequencing of a long amplicon determined the presence of one of the two lineages found in the marsupial. © 2009 The Linnean Society of London, *Biological Journal of the Linnean Society*, 2009, **98**, 568–576.

ADDITIONAL KEYWORDS: hemoparasites – host-parasite coevolution – phylogeny – rodents – South America – tick.

INTRODUCTION

Among the American marsupials, there exists a species, the so-called ‘Monito del Monte’, *Dromiciops gliroides* Thomas, 1894, which is the last living representative of the Order Microbiotheria (Redford & Eisenberg, 1992; Hershkovitz, 1999). This small mammal is a tree climber species with nocturnal activity, which lives in southern forest of *Nothofagus*

(Fagaceae) and *Chusquea* Bamboo in southern Chile and Argentina (Hershkovitz, 1999; Lobos *et al.*, 2005). The oldest fossil record from microbiotherids comes from the Early Paleocene of Bolivia (Gayet, Marshall & Sempere, 1991), and phylogenetic analyses of marsupials situated *Dromiciops* as being more closely related to Australidelphian than to Ameridelphian marsupials (Palma & Spotorno, 1999). It is clear that ‘Monito del Monte’ is a ‘living fossil’, being the last representative of a point of union between Australian and American marsupials.

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The protozoan blood parasites under the genera *Hepatozoon* are known to infect a variety of vertebrates, from anurans to mammals (Smith, 1996). These parasites are found in peripheral blood, where they infect white blood cells, except in marsupials and some rodents, where red blood cell infections are common (Smith, 1996). These parasites are transmitted to vertebrate hosts by ingestion of infected haematophagous ectoparasites, where the parasite is developed in the haemocoel. The variety of ectoparasites acting as definitive hosts of *Hepatozoon* parasites is also very diverse, varying from mosquitoes to mites, bugs, flies, lice, ticks, and fleas (Smith, 1996).

In the present study, we characterized, at the molecular and morphological level, *Hepatozoon* infections from a wild population of 'Monito del Monte' in Southern Chile. We also explored the phylogenetic position of these parasite lineages with respect to other previously described *Hepatozoon* species. In addition, we looked for the presence of these lineages infecting other micro-mammals cohabiting in the same area. Finally, we searched for potential parasite vectors using molecular tools.

MATERIAL AND METHODS

The study was conducted at Senda Darwin Biological Station and surrounding forest patches in northern Chiloé Island, Southern Chile. *Dromiciops gliroides* individuals were captured inside bird nest-boxes or by using Tomahawk-live traps (baited with banana), and blood samples were collected in August 2005 (for details of the study area, see Moreno *et al.*, 2005). Each animal was individually marked with numbered or coloured ear-tags to avoid resampling. Approximately 50 µL of blood were obtained with a heparinized hematocrit capillary tube from the tail vein or the infraorbital sinus (van Herck *et al.*, 2000). One part of the blood was immediately smeared on a slide, air dried, and later fixed with 96 ° ethanol and stained with Giemsa (1 : 10, v/v) for 40 min. Blood smears were scanned in search of blood parasites in accordance with previously described methods (Merino, Potti & Fargallo, 1997). The rest of the blood was transferred with the aid of a capillary to a ster-

ilized tube containing a buffer composed of 100 mM Tris (pH 8.0), 100 mM ethylenediaminetetraacetic acid (EDTA), 2% sodium dodecyl sulphate (SDS) (Jarvi, Schultz & Atkinson, 2002) to preserve blood samples for DNA analyses, or were stored on FTA classic cards (Whatman International Ltd.; Gutiérrez-Corcherro *et al.*, 2002). To identify potential vectors of the *Hepatozoon* infections, animals were also captured three times a year in 2007 and 2008 (i.e. August, November, and February) and examined in search of ectoparasites. Ticks were the most common ectoparasite detected and collected (see Results). Ticks and two unidentified fleas were maintained alive for 24 h to allow the blood digestion and then conserved in 70% ethanol. Subsequently, ectoparasites were lyophilized to facilitate transport to the laboratory in Spain and DNA extraction. In addition, during the summer of 2007, in the same study area, we also captured and blood sampled the following rodents; two *Oligoryzomys longicaudatus* Bennett, 1832, 11 *Abrotrix (Chroeomys) olivaceus* Thomas, 1916, and four *Abrotrix sanborni* Thomas, 1916, to check whether the *Hepatozoon* species detected in *D. gliroides* were also infecting other micro-mammals in the area of study. Only blood samples for DNA analyses were obtained from these animals and were analysed as described below.

Parasite morphometric measurements were made with the aid of image analyser software (Scion Image) from pictures of parasites taken under a ×100 oil immersion objective using an Olympus BX41 optic microscope. The length and width of the parasite and parasite nucleus were measured for each parasite (Table 1). Infected and uninfected erythrocytes were also measured.

DNA ANALYSIS

Genomic DNA from *D. gliroides* samples conserved in lysis buffer was obtained using the UltraClean DNA BloodSpin kit (MO BIO Laboratories, Inc.). On the other hand, genomic DNA present in FTA cards was extracted to a soluble solution before polymerase chain reaction (PCR) using the following protocol: cored samples were transferred to collection vials

Table 1. Measurements of parasites and erythrocytes found in 35 *Dromiciops gliroides*

	Parasite	Parasite nucleus	Infected erythrocyte	Uninfected erythrocyte (σ)
Length (µm)	9.39 (0.64)	3.42 (0.43)	11.33 (2.07)	5.86 (0.50)
Width (µm)	2.59 (0.34)	1.95 (0.47)	4.57 (1.13)	–
Sample size	23	23	21	69

The diameter of uninfected erythrocytes is also presented. Standard deviations are shown in parenthesis.

Table 2. List of primers used in the present study

Primers	Sequences (5'- to 3')
NBA1b	GTT GAT CCT GCC AGT AGT
NBA2	GCC TGC TGC CTT CCT TA
HEP1	CGC GCA AAT TAC CCA ATT
HEP4	TAA GGT GCT GAA GGA GTC GTT TAT
HPF1	CTA TGC CGA CTA GAG ATT G
HPF2	GAC TTC TCC TTC GTC TAA G
NBA1	GGT TGA TCC TGC CAG TAG T

with 250 μ L of SET buffer (0.15 M NaCl, 0.05 M Tris, 0.001 M EDTA; pH 8) at 4 °C for 6 h. Subsequently, SDS 20% (7 μ L) and proteinase K (50 μ g) were added to the vials and incubated at 55 °C overnight. After incubation, ammonium acetate 4 M (250 μ L) was added to the vials at room temperature for 30 min. Subsequently, vials were centrifuged at 13 000 g for 10 min. After removing the pellet, DNA was precipitated with ethanol and resuspended in sterile water.

Amplification of 18S rDNA gene (1769 bp) was accomplished by PCR using the primers NBA1b/HPF2 (Table 2). PCR reactions consisted of 25- μ L reaction volumes containing 20 ng of template DNA, 50 mM KCl, 10 mM Tris-HCl, 1.5 MgCl₂, 0.2 mM of each dNTP, 1 μ M of each primer, and 1.25 U of AmpliTaq Gold (Applied Biosystems). The reactions were cycled at the following parameters using a thermal cycler (MasterCycler Personal, Eppendorf): 94 °C for 10 min (polymerase activation), 40 cycles at 95 °C for 40 s, 54 °C for 1 min, 72 °C for 2 min, and a final extension at 72 °C for 10 min. The amplicons obtained after PCR assays were recovered from agarose gels and subjected to direct sequencing using the primers NBA1b, NBA2, HEP1, HEP4, HPF1, and HPF2 (Table 2). The sequencing was carried out using an ABI 3130 (Applied Biosystems) automated sequencer. DNA sequences were aligned and edited using CLUSTALW (Thompson, Higgins & Gibson, 1994) and BIOEDIT software (Hall, 1999), respectively. The phylogenetic analyses were performed using three different methods of inference after removing all columns containing gaps or missing data. The Neighbour-joining method (maximum composite likelihood model), the maximum likelihood method (GTR model; the gamma distribution parameter, the proportion of invariable sites, and the transition/transversion ratio were estimated) and Bayesian estimation (Markov chain Monte Carlo algorithm) were carried out using MEGA, version 4.0 (Tamura *et al.*, 2007), PHYML, version 3.0 (Dereeper *et al.*, 2008) and MrBayes (Ronquist & Huelsenbeck, 2003), respectively. Tree consistency was estimated by bootstrap

analysis with 100 replications. All *Hepatozoon* sequences in GenBank with a similar size to those lineages found in *D. gliroides* were included in the phylogenies (Table 3). Although with some limitations, small subunit rDNA (SSU) is generally considered to be a good marker for addressing relationships of apicomplexan species and their history (Morrison & Ellis, 1997; Zhu, Keithly & Philippe, 2000; Šlapeta *et al.*, 2003). In addition, only SSU rDNA sequences are available for sufficient *Hepatozoon* species to make evolutionary inferences of parasites within this genus.

To detect the presence of the parasite in potential ectoparasite vectors, we followed the method of Schall & Smith (2006). This method aims to avoid the amplification of DNA from parasites digested with the blood meal. This allows the discrimination of whether a particular species acts as vector of a particular parasite because only parasites that survive to digestion are transmitted. Arthropods were maintained alive for 24 h, thus allowing for the blood meal to be digested, including the parasite DNA. Thus, it would be very difficult to amplify a long DNA fragment from the parasite, except if the arthropod is the vector. In this case, the parasites would be located in the haemocoel, which is a safe place from digestive endonucleases.

The ticks and two fleas achieved from 'Monito del Monte' were maintained alive for 24 h before storage in 70% ethanol. DNA from lyophilized ectoparasites was extracted using the method previously described for FTA cards. Amplification of 18S rDNA gene (1096 bp) was accomplished by PCR using the primers NBA1/HEP4 (Table 2). PCR reactions were carried out using the conditions described above.

STATISTICAL ANALYSIS

To identify potential differences in the prevalence of infection by *Hepatozoon* between individuals of different sexes and ages, we conducted statistical analyses using Yates corrected chi-square tests (STATISTICA, version 6.0; StatSoft, Inc.).

RESULTS

Overall, we captured 77 *D. gliroides*, comprising 44 males (39 adults and four juveniles; one male was not aged), 29 females (11 adults and 18 juveniles) and four individuals not sexed and aged. Excluding the last four individuals, blood smears were obtained from all animals. Also, with the exception of two juvenile females, blood for DNA analyses was obtained for a total of 75 individuals.

Thirty-five out of 73 (47.9%) blood smears from different individuals were found infected by a *Hepa-*

Table 3. Protozoan species used for phylogenetic analyses indicating their hosts, Genbank accession numbers, location, and reference when available

Parasite	Host	Genbank accession number	Location	Reference
<i>Hepatozoon catesbiana</i>	<i>Rana catesbiana</i>	AF 130361	USA	Carreno, Martin & Barta (1999)
<i>Hepatozoon boigae</i>	<i>Boiga fusca</i>	AF 297085	Australia	Unpublished
<i>Hepatozoon</i> sp.	<i>Isoodon obesulus</i>	EF152218 to EF152230	Australia	Wicks <i>et al.</i> (2006)
<i>Hepatozoon</i> sp.	<i>Abrotrix olivaceus</i>	FJ719815, FJ719817, FJ719818	Chile	Present study
<i>Hepatozoon</i> sp.	<i>Abrotrix sanborni</i>	FJ719816 FJ719819	Chile	Present study
<i>Hepatozoon</i> sp.	<i>Bandicota indica</i>	AB181504	Thailand	Unpublished
<i>Hepatozoon</i> sp.	<i>Clethrionomys glareolus</i>	AY 600625, AY 600626	Spain	Criado-Fornelio <i>et al.</i> (2006)
<i>Hepatozoon felis</i>	<i>Felis catus</i>	AY 620232, AY 628681	Spain	Criado-Fornelio <i>et al.</i> (2006)
<i>Hepatozoon americanum curupira</i>	<i>Dusycion thous</i>	AY 461377	Brazil	Criado-Fornelio <i>et al.</i> (2006)
<i>Hepatozoon canis</i>	<i>Canis familiaris</i>	AY 461378	Spain	Criado-Fornelio <i>et al.</i> (2006)
<i>Hepatozoon canis</i>	<i>Vulpes vulpes</i>	AY731062, AY 150067	Spain	Criado-Fornelio <i>et al.</i> (2006)
<i>Hepatozoon canis curupira</i>	<i>Dusycion thous</i>	AY 461375	Brazil	Criado-Fornelio <i>et al.</i> (2006)
<i>Hepatozoon canis curupira</i>	<i>Pseudalopex gymnocercus</i>	AY471615, AY 461376	Brazil	Criado-Fornelio <i>et al.</i> (2006)
<i>Hepatozoon</i> sp.	<i>Dromiciops gliroides</i>	FJ719813, FJ719814	Chile	Present study
<i>Adelina bambarooniae</i>	<i>Dermolepida albohirtum</i>	AF494059	Australia	Unpublished

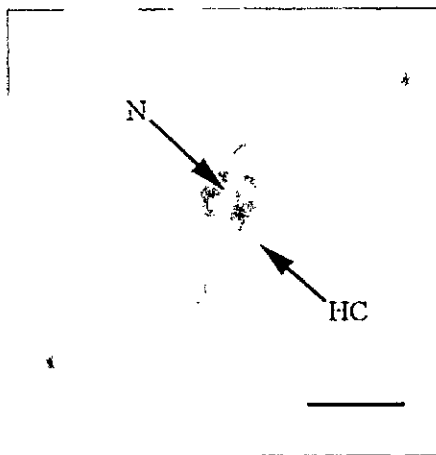


Figure 1. Photomicrograph of a blood smear from the 'Monito del Monte' (*Dromiciops gliroides*) showing an erythrocyte infected with *Hepatozoon*. N, parasite nucleus; HC, host cell. Scale bar = 5 µm.

tozoon species. Parasites infect red blood cells and show the typical banana shape of these parasite species (Fig. 1). Positive slides are deposited at the collection of Museo Nacional de Ciencias Naturales,

Madrid, Spain (accession numbers: MNCN 35.02/33, MNCN 35.02/34, and MNCN 35.02/35 blood smears from *D. gliroides*). Molecular methods increased the prevalence of infection detected up to 86.7% (i.e. 65 out of 75 individuals infected).

There were no significant differences between sexes in the proportion of infected individuals estimated either by molecular ($\chi^2 = 1.52, P = 0.22$) or microscopic methods ($\chi^2 = 0.08, P = 0.78$). Significant differences between age classes were only found for data based on blood smears ($\chi^2 = 14.20, P < 0.001$), indicating that more adults than juveniles were uninfected. The fact that this difference was not significant for molecular data on infection ($\chi^2 = 0.01, P = 0.96$) indicates that the intensity of infection is higher in juveniles because many infections were not detected in adults by microscopic as compared to molecular methods. Differences in the prevalence of infection between sexes by age classes were not significant (data not shown, $P > 0.05$).

The tick *Ixodes neuquenensis* Ringuelet 1847 was collected from *D. gliroides* (two adults and 15 nymphs in 2007 and 13 adults and 12 nymphs in 2008). Long sequences from the *Hepatozoon* lineages infecting 'Monito del Monte' were recovered from three adult *I. neuquenensis* (one from 2007 and two from 2008). We

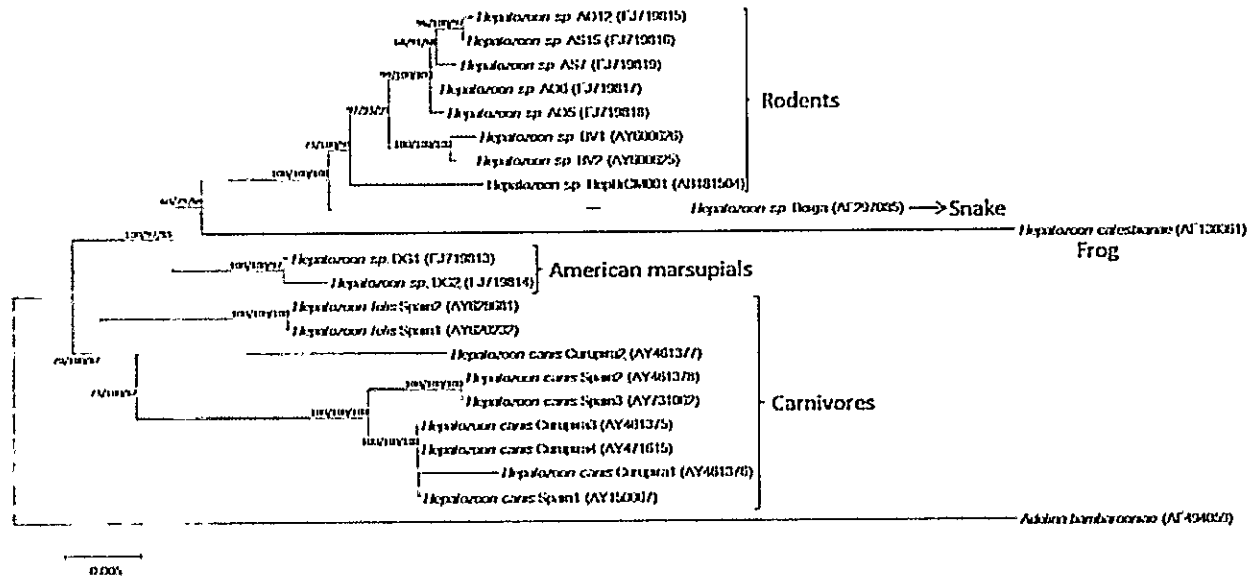


Figure 2. Phylogenetic tree obtained with the MEGA, version 4 software, using the Neighbour-joining method. Bootstrap values are shown at the corresponding nodes. When three values appear, they correspond to maximum composite likelihood, Bayesian, and maximum likelihood methods, respectively. Figures in parentheses after the species name indicate the GenBank accession number of the isolate. The hosts where *Hepatozoon* species were recovered are indicated. Samples from 'Monito del Monte' comprise those from American marsupials. This analysis is based on 1627 bp.

were unable to recover long segments of *Hepatozoon* from the two unidentified fleas.

Finally, we detected *Hepatozoon* infections in two of the three rodent species co-existing in the same area with 'Monito del Monte'. We found six *Abrotrix olivaceus* and two *A. sanborni* infected by five different *Hepatozoon* lineages. Three of them were present in *A. olivaceus* and two in *A. sanborni*. None of them were found infecting the 'Monito del Monte' (see sequences with accession numbers FJ719815 to FJ719819 in Fig. 2; Table 3).

PHYLOGENIES

The tree including the short sequences of *Hepatozoon* species from the Australian bandicoot are based on 743 bp after removing columns containing gaps or missing data. This tree is used to determine the phylogenetic relationships between *Hepatozoon* lineages isolated from 'Monito del Monte' and those from the bandicoot (Fig. 3), and we infer phylogenetic relationships among parasites infecting different host taxa from the tree based on 1627 bp (see below; Fig. 2) although conclusions did not vary considerably by using either of them except for bootstrap support. The *Hepatozoon* lineages found in both marsupial species are not closely related between them and the lineages found in Australian marsupials, American marsupials/anurans, rodents/reptiles, and carnivore species form clearly separated groups (Fig. 3). To infer

a sound phylogenetic relationship, we consider the more robust tree based on 1627 bp after removing columns containing gaps or missing data (Fig. 2). The topology of the tree indicates the grouping of the lineages in two clades supported with a high bootstrap value. One of them is formed by all *Hepatozoon* lineages found in carnivore species and the other by the lineages found in rodents, amphibians, reptiles, and the American marsupial. In this case, the lineages found in marsupials have a basal position in the clade.

DISCUSSION

The *Hepatozoon* species infecting 'Monito del Monte' appears to be clearly separated from lineages recovered from the Australian bandicoot. This may be interpreted as indicating that different *Hepatozoon* lineages developed independently in marsupials in both continents because the lineages recovered from *D. gliroides* appear to evolve from a common ancestor, previous to the separation of *Hepatozoon* species infecting rodents and *Isoodon obesulus*, although this tree topology lacks strong support (Fig. 3). Indeed, it has been recently suggested that parasite-host coevolution implying Apicomplexa follows the definitive hosts, namely those where sexual reproduction appear (Šlapeta *et al.*, 2003). Based on molecular and paleontological data, Palma & Spotorno (1999) pro-

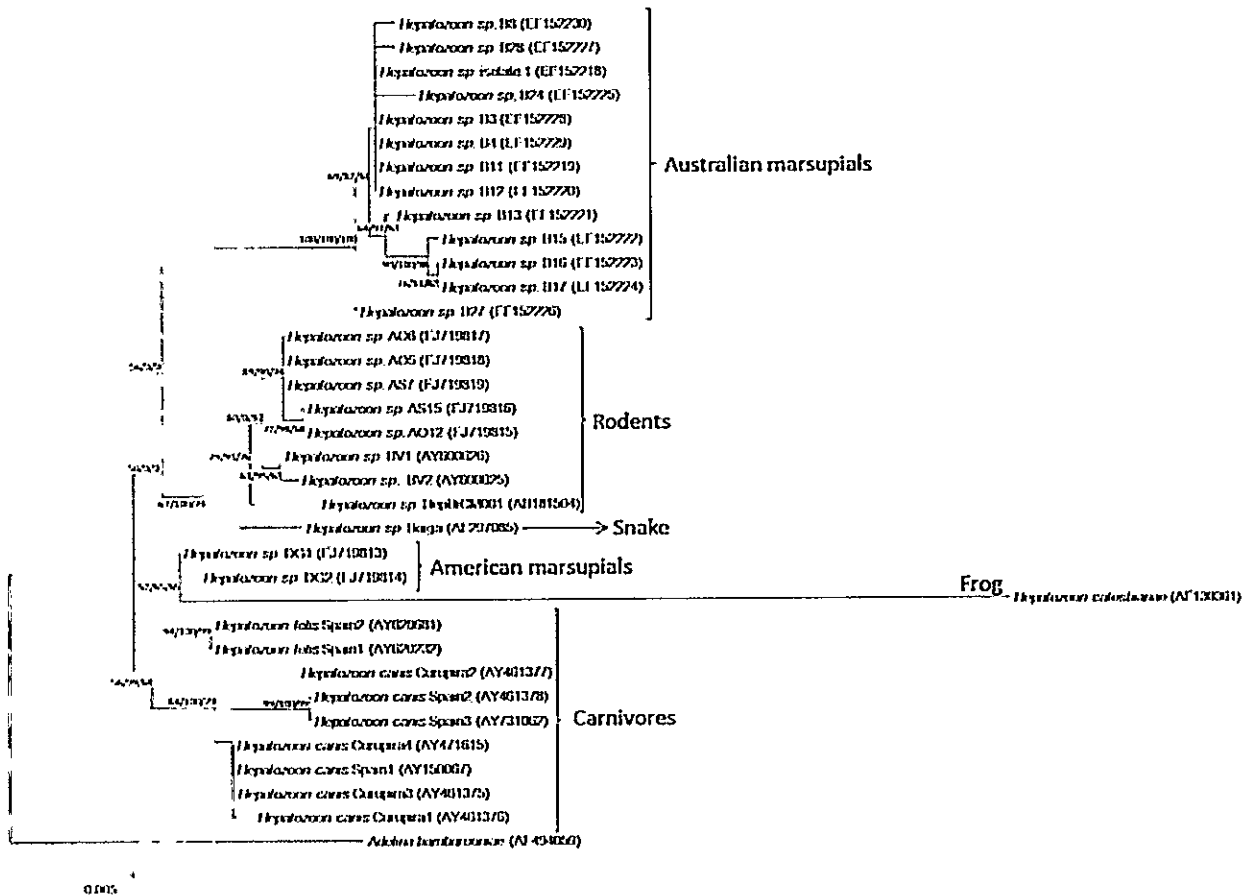


Figure 3. Phylogenetic tree obtained with the MEGA, version 4 software, using the Neighbour-joining method. Bootstrap values are shown at the corresponding nodes. When three values appear, they correspond to maximum composite likelihood, Bayesian, and maximum likelihood methods, respectively. Figures in parentheses after the species name indicate the GenBank Accession number of the isolate. The hosts where *Hepatozoon* species were recovered are indicated. Samples from 'Monito del Monte' are those from American marsupials. Samples from Australian marsupials are reported by Wicks *et al.* (2006). This analysis is based on 743 bp.

posed that Microbiotherids evolve in South America and later migrate to Australia across Antarctica prior to the separation of continents. If *Hepatozoon* species evolve with their definitive hosts and if *Hepatozoon* in *D. gliroides* separated from a common ancestor of other *Hepatozoon* species infecting Australian marsupials (Fig. 3), we can propose that the relationships *Hepatozoon*-*D. gliroides* is the result of a long coevolving relationship between host and parasite. That is, the relationship may have existed for a long time subsequent to the separation of continents. The complete sequence of 18S rDNA of *Hepatozoon* lineages from Australian marsupials may help to solve the position of the common ancestor of these parasites.

The possibility that *Hepatozoon* species infecting American marsupials were the ancestor to species infecting other mammals except carnivores obtains

some reinforcement from the tree based on a longer fragment (1627 bp) of the small subunit ribosomal RNA gene, because the *Hepatozoon* lineages infecting *D. gliroides* appear in a basal position with respect to species infecting anurans, ophidians, and rodents, including lineages isolated from sympatric rodents. The *Hepatozoon* species parasitizing carnivores appear to be clearly separated in another clade (Fig. 2). Thus, the lineages isolated from *D. gliroides* appear to be host-specific and to be an ancient species of *Hepatozoon* with respect to those infecting rodents and lower vertebrates. The basal position of *Hepatozoon* in 'Monito del Monte' with respect to those in anurans and ophidians may indicate that these lower vertebrates were infected secondarily from mammals. *Hepatozoon* species present in rodents in the area also form a robust clade; thus, lineages of this parasite in these mammals also appear to be endemic to the area.

The basal position of *Hepatozoon* infecting 'Monito del Monte' prompted us to consider that the relationship between the marsupial and the parasite should also be mediated by an ancient and host-specific ectoparasite. In this respect, we recovered long segments of *Hepatozoon* DNA corresponding to the species infecting *D. gliroides* from three ticks using the method described above (Schall & Smith, 2006). Therefore, *I. neuquenensis* appears to be a potential vector of the *Hepatozoon* species infecting 'Monito del Monte'. This is interesting because *I. neuquenensis* is considered to be a host-specific ectoparasite of this vertebrate (Guglielmone *et al.*, 2004; Guglielmone & Nava, 2005; Marín-Vial *et al.*, 2007). However, this tick did not appear as a very ancient species compared to tick species from other vertebrates (Guglielmone *et al.*, 2006), although these data are only based on phylogenetic analyses of sequences of 460 bp. This may imply that the *Hepatozoon* parasite has adapted to *I. neuquenensis* later than to *D. gliroides*, and that either we failed to sample the original vector or it has now disappeared. The possibility that *I. neuquenensis* may have had a broad specificity and only recently invaded *D. gliroides* is not supported by the lack of reports of this tick species infecting other hosts (Guglielmone & Nava, 2005). It is possible that other vectors of the parasite exist because the prevalence of these *Hepatozoon* lineages in *I. neuquenensis* is very low compared to the prevalence in the vertebrate host. However, this may be related to host susceptibility and not necessarily to a relationship between high prevalences of infection in vertebrate hosts and a high abundance of vectors (Sol, Jovani & Torres, 2000).

Several species of *Hepatozoon* have been described infecting marsupials both in Australia and America, and all of them infect erythrocytes. A common species infecting American marsupials and described as *Hepatozoon didelphydis* was recently found to be a Sarcocystidae (Merino *et al.*, 2008), and a similar case has been recently reported for a marsupial species in Australia (Zhu *et al.*, 2009). Criado-Fornelio *et al.* (2006) failed to find *Hepatozoon* infections in 15 opossums (*Didelphis albiventris*) from Brazil either by molecular methods or microscopy, in contrast to the 25% prevalence detected by microscopy in the same species from French Guyana (de Thoisy *et al.*, 2000). *Hepatozoon* infections have also been reported infecting other neotropical marsupials, including *Didelphys marsupialis*, *Philander opossum*, and *Metachirus nudicaudatus* (d'Utra e Silva & Arantes, 1916; Regendanz & Kikuth, 1928; Garnham & Lewis, 1958; Deane & Deane, 1961; Ayala *et al.*, 1973). However, these reports are only based in microscopy and some of them are attributed to *H. didelphydis*, and may correspond to infections by an erythrocytic sarcocystidae (Merino *et al.*, 2008).

The high prevalence of these parasites in 'Monito del Monte' implies that it is a common parasite in the area of study. The infection is more difficult to detect in blood smears from adult individuals, thus indicating that juveniles suffer from more intense infections. This implies that juveniles are more susceptible to the parasite and adults probably control the infection better. This pattern agrees with an increase in immune responses as animals age, although, in most cases, the disease is not completely cleared, as demonstrated by the molecular data. Alternatively, highly-infected young individuals could die as a result of infection or associated causes and only individuals that are able to control infection reach adulthood. Information on the effects of infection on *D. gliroides* and on variation in the prevalence between populations, as well as on other potential vectors, will help to reveal the real impact of these parasites on this threatened 'living fossil' species of marsupial because *Hepatozoon* infections have been shown to be pathogenic with respect to their hosts (Ewing & Panciera, 2003).

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