Partial and clear-cut harvesting of high-elevation spruce-fir forests: implications for small mammal communities

Walt Klenner and Thomas P. Sullivan

Abstract: Subalpine spruce (Picea) - fir (Abies) forests occur throughout the Cordillera of western North America. A variety of alternative silvicultural systems to clear-cutting are being investigated in these high-elevation forests in terms of their impacts on ecosystem components. We tested the hypotheses that abundance, reproduction, and survival of populations of (i) southern red-backed voles (Clethrionomys gapperi Vigors) will decline, (ii) long-tailed voles (Microtus longicaudus Merriam) and northwestern chipmunks (Tamias amoenus J.A. Allen) will increase, and (iii) deer mice (Peromyscus maniculatus Wagner) will be similar, with respect to decreasing levels of tree retention. Small mammal populations were live-trapped from 1994 to 1998 in replicated sites of uncut forest, single tree selection, 0.1-ha patch cuts. 1.0-ha patch cuts, a 10.0-ha clearcut, and edges in an Engelmann spruce (Picea engelmannii Parry ex Engelm.) subalpine fir (Abies lasiocarpa (Hook.) Nutt.) forest near Sicamous, British Columbia, Canada. Clethrionomys gapperi dominated the small mammal community, reaching a peak abundance of 80 animals/ha with mean values ranging from 31 to 50 animals/ha. Over the 4 postharvest years, abundance, reproduction, and survival of C. gapperi populations were consistently similar among uncut forest, single tree selection, and patch cut treatments compared with clear-cut sites where this species declined to extirpation. Microtus longicaudus and T. amoenus occurred predominantly in clearcut sites, whereas P. maniculatus was present in low numbers on all sites. With respect to small mammals, partial harvesting systems appear to provide a means for combining timber extraction with maintenance of mature forest habitat in these subalpine ecosystems.

Résumé : Des forêts subalpines d'épinette (Picea) et de sapin (Abies) croissent un peu partout dans la Cordillère de l'ouest de l'Amérique du Nord. Les impacts écosystémiques de plusieurs systèmes sylvicoles sont actuellement étudiés comme solution de rechange à la coupe à blanc dans ces forêts de haute altitude. Nous avons testé les hypothèses voulant que l'abondance, la reproduction et la survie de populations vont, en relation avec une diminution du niveau de rétention d'arbres : (i) diminuer chez le campagnol à dos roux de Gapper (Clethrionomys gapperi Vigors); (ii) augmenter chez le campagnol à longue queue (Microtus longicaudus Merriam) et le tamia du Nord-Ouest (Tamias amoenus J.A. Allen); et (iii) rester stables chez la souris sylvestre (Peromyscus maniculatus Wagner). Les populations de petits mammifères ont fait l'objet de piégeage non mortel de 1994 à 1998 dans des sites répétés de forêt intacte, de jardinage par pied d'arbre, de coupes par trouées de 0,1 ha, de coupes par trouées de 1,0 ha, de coupes à blanc de 10,0 ha et de bordures, dans une forêt d'épinettes d'Engelmann (Picea engelmannii Parry ex Engelm.) et de sapins subalpins (Abies lasiocarpa (Hook.) Nutt.) près de Sicamous, en Colombie-Britannique, au Canada. Clethrionomys gapperi dominait la communauté de petits mammifères, avec une abondance maximale atteignant 80 individus/ha et des valeurs moyennes de 31 à 50 individus/ha. Durant les 4 années après coupe, l'abondance, la reproduction et la survie des populations de C. gapperi étaient de façon soutenue semblables dans la forêt intacte, le jardinage par pied d'arbre et les coupes par trouées en comparaison avec les coupes à blanc où l'espèce a diminué jusqu'à l'extirpation. Microtus Longicaudus et T. amoenus occupaient surtout les coupes à blanc tandis que P. maniculatus était présent partout mais peu abondant. Les systèmes de coupe partielle semblent donc, quant aux petits mammifères, un moyen de combiner la récolte ligneuse et le maintien d'habitats de forêt mature dans ces écosystèmes subalpins.

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Introduction

Green-tree retention systems have become more common across a range of forested ecosystems in North America where clear-cutting has been the dominant method of forest harvesting. Variable retention harvesting systems can maintain or increase structural diversity of the regenerating stand and provide some mature forest habitat (McComb et al. 1993; Franklin et al. 1997). Retention of structural characteristics includes not only "green trees" but also snags, logs, system structure and function and enhance connectivity across the forested landscape.

The sustainability of clear-cutting has been questioned (Franklin 1995; Wagner et al. 1998) and the practice is particularly under scrutiny with respect to subalpine forests in the southern interior of British Columbia (Vyse 1999). These high-elevation forests are projected to be a major source of timber over the next several decades. In addition, clear-cut harvesting near the highest elevations, close to the tree line, has prompted concerns about forest regeneration (Vyse 1999). These forests may be at the "physiological edge" where low temperatures, high winds, and deep snowfalls combine to reduce biological activity above and below ground and hence tree regeneration (Vyse 1997). Although much work on regeneration at high elevations has been done in North America (see review by Alexander 1986), there is little information on the responses of other ecosystem components to forest harvesting and stand regeneration.

A major component of the fauna in these high-elevation Engelmann spruce (Picea engelmannii Parry ex Engelm.) subalpine fir (Abies lasiocarpa (Hook.) Nutt.) forests is the small mammal community. Forest floor small mammals provide a measure of ecosystem function and may be indicators of sustainability of forests in the Pacific Northwest and elsewhere (Carey and Harrington 2001; Sullivan and Sullivan 2001). Ecosystem functions include serving as prey for a wide variety of reptiles, carnivores, hawks, and owls (Verts and Carraway 1998), consumption of invertebrates (Buckner 1966; Gunther et al. 1983) and various plant products (Sullivan et al. 1990; Carey et al. 1999), and dispersal of fungal spores (Maser et al. 1978). Thus, the abundance and dynamics of small mammal populations may serve as a measure of the impact of alternative harvesting treatments on ecosystem function in subalpine forests.

The southern red-backed vole (Clethrionomys gapperi Vigors) is a common inhabitant of late successional coniferous and deciduous forests across temperate North America (Merritt 1981) and may be an important indicator species of "old-forest conditions". This vole occurs in substrates with abundant organic debris composed of stumps, logs, and exposed roots (Martell and Radvanyi 1977; Yahner 1986) that maintain hypogeous ectomycorrhizal fungi as a major food supply (Maser et al. 1978). Other prominent forest floor species include the deer mouse (Peromyscus maniculatus Wagner), long-tailed vole (Microtus longicaudus Merriam), meadow vole (Microtus pennsylvanicus Ord), and northwestern chipmunk (Tamias amoenus J.A. Allen). The deer mouse is a habitat generalist occupying a wide variety (type and age) of forested habitats (Baker 1968; Martell 1983; Sullivan et al. 1999). Tamias amoenus is also a habitat generalist found primarily in early successional harvested sites (Sutton 1992; Sullivan and Sullivan 2001). The preference of M. longicaudus and M. pennsylvanicus for early successional habitats after natural or forest harvesting disturbances may be related to the abundance of herbs and grasses that provide food and cover (Reich 1981; Getz 1985).

Clethrionomys gapperi is often the most abundant species in montane and old-growth subalpine spruce-fir forests (RaMedin and Booth 1989; Steventon et al. 1998; Von Trebra al. 1998; Sullivan and Sullivan 2001). In subalpine spruc fir forests, patch cutting appeared to have little effect on a abundance of *C. gapperi* (Hayward et al. 1999). However comprehensive evaluation of the responses of *C. gapp* and other small mammal species to a range of partial cutt; treatments in subalpine spruce-fir forests has not been dor

We tested the hypotheses that abundance, reproduction and survival of (i) C. gapperi, a species associated we closed canopy forests, will decline with decreasing levels tree retention, (ii) M. longicaudus and T. amoenus, spec associated with early successional habitats, will increwith decreasing levels of tree retention, and (iii) P. ma culatus, a habitat generalist species, will be similar acrthe levels of tree retention.

Materials and methods

Study area

As part of the Sicamous Creek Silvicultural Systems P: ject, this study was located in the Salmon Arm Forest D trict, south of the north fork of Sicamous Creek and north Mount Mara, near Sicamous, British Columbia, Can. (50°50'N, 119°50'W). The forest corresponds to the Eng mann Spruce Subalpine Fir (ESSF) biogeoclimatic ze (Meidinger and Pojar 1991) and is representative of t ESSF wet cold subzone (ESSFwc2) described by Lloyd al. (1990). This subzone is the largest of the seven ES subzones in the Kamloops Forest Region and is very simi to the cold, wet ESSF subzones in the neighbouring Nels and Cariboo forest regions. The site elevation ranges free 1530 to 1830 m and the tree line is found between 1850 a 1900 m immediately to the south. The forest is a matu stand of subalpine fir and Engelmann spruce that original approximately 300 years ago and has experienced individu tree or group mortality since the mid-1800s (Parish et 1999). The standing volume of live timber on the area w estimated to be 264 m3/ha, of which approximately 35% w spruce and 65% was subalpine fir. Thirty percent of standing trees were snags. The understory vegetation w primarily white rhododendron (Rhododendron albiflor Hook.), black huckleberry (Vaccinium membranace. Dougl.), and oval-leaved blueberry (Vaccinium ovalifoli. Smith). The project area forms part of the Sicamous Cru Community Watershed (Vyse 1999).

The project area was divided into three blocks based elevation (mean values 1615 m for block A, 1690 m block B, and 1750 m for block C), and five harvesting truments representative of a range of tree retention patte were applied to each block (Fig. 1). All treatments were h vested in the winter of 1994–1995. The largest opening s of 10 ha (320 m \times 320 m) was planned with a 100-m-w buffer strip of mature forest, and other buffer strips were as needed to separate the treatment units. Each unit was proximately 30 ha. After harvesting, most treatment ut were site prepared in 1995 using excavators to create sn mounds (<1 m² and raised 30 cm) and then planted in 1⁶ with Engelmann spruce seedlings. Some small portions Location of the Sicamous Creek site in south-central British Columbia and the arrangement of treatment units at the site. d areas indicate harvested units and dark grey areas are the single tree selection units.



atment areas were neither site prepared nor planted 1999).

five treatments were as follows. (i) Control: no reof trees; (ii) single tree cut: 33% volume removal by on over 30 ha by cutting every fifth tree using fallers' (no tree marking), with marked skid trails, but with empt to "improve the stand"; (iii) 0.1-ha patch cut: moval of volume over a 30-ha area by cutting approxy sixty 0.1-ha square openings with 60 m between and skid trails linking all groups, (iv) 1.0-ha patch % removal of volume over a 30-ha area by cutting roups of 1 ha each with 200 m between centres and nking skid trails; (v) 10-ha clearcut: 33% removal of over a 30-ha area by cutting one 10-ha opening apnately 330 m square.

mental design

study had a randomized-block design with three replies of each treatment. Within each block, there were es: uncut forest (UF), single tree selection (STS), patch cut (0.1 PC), 1.0-ha patch cut (1.0 PC), 10.0-ha t (10.0 CC), and an edge habitat. On the patch cut nts, the trapping grids were positioned to represent pitat types following harvest. For example, on the 0.1-ha sites, the trapping grids extended across several openings, with about 60% of the sampling area in uncut forest. The edge sites were located on the forest-opening interface of the 10.0-ha clearcut. These sites were established to determine the responses of small mammals to the edges and leave strips created during harvesting.

Small mammal populations

Populations of forest floor small mammals were livetrapped at 3-week intervals from July to September 1994, 1995, and 1996, from June to September 1997, and from June to August 1998. One trapping grid (1 ha), with 49 (7 × 7) trap stations at 15.0-m intervals and one or two Longworth-type live-traps at each station, was located on each site (Ritchie and Sullivan 1989). The 1.0-ha patch cut and edge sites were not trapped in 1994. Traps were baited with a mixture of whole oats, sunflower seeds, and a slice of apple. The nest chamber of each trap was filled with coarse brown cotton for insulation. During each trapping session, traps were set for two consecutive nights and were locked open during the day. Between trapping sessions, traps were locked open to allow animals to enter and maintain familiarity with the traps. When capture success exceeded 80% of the 49 traps on a sampling grid on both days of the trapping

Forest floor small mammal species sampled by this protocol included C. gapperi, M. longicaudus, P. maniculatus, T. amoenus, heather vole (Phenacomys intermedius Merriam), northern bog lemming (Synaptomys borealis Richardson), water vole (Microtus richardsoni DeKay), jumping mouse (Zapus sp.), and shrews (Sorex spp.). Captured animals were marked with unique numeric ear tags, and the species, mass (Pesola[®] spring balance), sex, and reproductive condition recorded before release at the point of capture. Palpation of male testes and mammaries of females was used to measure reproductive state (Krebs et al. 1969). Unfortunately, there was a high mortality rate for shrews in the traps overnight, but this was unavoidable in practice.

Demographic analysis

Estimates of abundance for the major species, C. gapperi, P. maniculatus, M. longicaudus, and T. amoenus, were derived from the Jolly-Seber stochastic model (Seber 1982). The minimum number of animals known to be alive (Krebs 1966) was used as the population estimate for the first and last sampling periods of the study when the Jolly-Seber estimate was not calculated. Because the reliability of the Jolly-Seber model declines when population sizes are consistently very low and (or) no marked animals are captured (Krebs et al. 1986), the minimum number of animals known to be alive was used to estimate abundance of P. intermedius, M. pennsylvanicus, S. borealis, and M. richardsoni. Number of captures was calculated for Sorex spp. These uncommon species all occurred in very low abundance in all treatment sites.

We evaluated reproduction, recruitment, and survival of the four major species. Reproductive performance was based on the number of successful pregnancies calculated for each population in 1994-1998. A pregnancy was considered successful if a female was lactating during the period following the estimated time of birth of a litter (Sullivan 1990). Each record of a lactating female was considered an independent litter based on a 3-week nursing period within our 3-week trapping intervals. Early juvenile survival is an index relating recruitment of young into the trappable population to the number of lactating females (Krebs 1966). A modified version of this index is the number of juvenile animals at week t divided by the number of lactating females caught in week t - 3. Mean Jolly-Seber 28-day survival rates were calculated for each population during the summer (June-September) and winter (October-May) periods each year.

We used age at sexual maturity to determine age-classes of voles. Body mass was used as an index of age. Animals were classified as juvenile (juvenile and subadult classes pooled) or adult by body mass: C. gapperi, juvenile 1–18 g, adult \geq 19 g; M. longicaudus, juvenile 1–26 g, adult \geq 27 g; P. maniculatus, juvenile 1–15 g, adult \geq 16 g. Juveniles were considered to be young animals recruited during the study. Recruits were defined as new animals that entered the population through reproduction and immigration.

Statistical analyses

A repeated-measures analysis of variance (RM-ANOVA) was used to determine the effect of harvest treatments on

and mean index of juvenile survival and mean Jolly-Seber summer and winter survival for *C. gapperi* only, during the post-treatment period (1995–1998). A one-way ANOVA was conducted on these same variables during the pretreament year (1994). Data not conforming to properties of normality and equal variance were subjected to various transformations (Zar 1999). Sphericity (independence of data among repeated measures) was not detected in any of the data sets, which were subjected to Mauchly's *W* test statistic (Littel 1989; Kuehl 1994).

Proportional data were arcsine transformed prior to analy. sis (Zar 1999). Duncan's multiple range test (DMRT) was used to compare mean values based on ANOVA results. If there was a significant treatment effect and significant treatment x time interaction, a one-way ANOVA (and DMRT; was conducted for each post-treatment year. If there was a significant treatment effect but not a significant treatment x time interaction, DMRT was conducted on the overall set of post-treatment years. In all cases, the level of significance was at least P = 0.05.

Results

Abundance

Clethrionomys gapperi was the most abundant species with a total of 3998 individuals captured (66.9% of total small mammals). Total numbers of individuals of the other major species were 713 for *M. longicaudas*, 646 for *P. maniculatus*, and 278 for *T. amoenus*.

Estimates of trappability (susceptibility to capture) tended to be variable among species, with overall mean values of 77.5% for *C. gapperi*, 57.8% for *M. longicaudus*, 57.2% for *P. maniculatus*, to 69.0% for *T. amoenus*. Thus, Jolly-Seber population estimates were used for this study, for the reasons given in Jolly and Dickson (1983).

The abundance of *C. gapperi* was significantly $(F_{[5,10]} = 132.94, P < 0.01)$ different among sites, with the highest mean numbers occurring in the uncut, single tree selection, and patch cut sites in 1995, ranging from 31.1 to 49.7 voles/ha (Table 1; Fig. 2). Numbers of *C. gapperi* were lowest in the clear-cut sites during all postharvest years. Overall, *C. gapperi* populations generally declined with time $(F_{[3,36]} = 26.24, P < 0.01)$. By 4 years postharvest, abundance of *C. gapperi* was highest in the uncut forest and single tree selection, with no differences among any of the partial harvesting methods and edge habitats (Table 1). This microtine had all but disappeared from the clear-cut sites by the third year postharvest.

Abundance of *M. longicaudus* was significantly $(F_{15,10} = 4.30, P = 0.02)$ different among sites and over time $(F_{13,36} = 10.43, P < 0.01)$ with the highest relative numbers in 1994 and 1998 (Table 1; Fig. 3). Clear-cut sites tended to have the highest mean abundance at 16.2 voles/ha in 1998. *Microhus longicaudus* disappeared from the uncut forest after 1995 (Table 1).

Numbers of *P. maniculatus* were also similar $(F_{15,10} = 0.11, P = 0.99)$ among sites but with relatively high numbers in 1994 and 1998 as indicated by the significant $(F_{13,30} = 6.90, P < 0.01)$ difference among years (Table 1).

fable 1. Mean	 (n = 3 replicat pretreatment ye Uncut forest 	te sites) alcund ar (1994) with Single tree selection	unce per hec results of a 0.1-ha patch	tare for c.ii one-way AN 1.0-ha patch	OVA and 4 10.0-ha	pperi, Miero post-treatmet Edue	ius longicaudus, Peromysc at years (1995-1998) with Treatment	us municulatur, and Tami results of RM-ANOVA. Time	is amoenus for the Treatment x time
lethrionomys gapperi						allow			and a month of the
1994	32.5±1.8	43.5±8.1	32,4±2.4	Ę	34.3±3.7	I	$F_{(1,0)} = 1.27, P = 0.35$.		
1995	31.1±3.6ub	44,0±8.3a	49.7±3.0a	38.1±6.0ab	10.8±0.4c	24.7±5.1b	$F_{[5,10]} = 132.94, P < 0.01$	$F_{13,361} = 26.24, P < 0.01$	Fi15 161 = 6.81, P <
1996	26.9±1.2a	24.2±1.7a	21.1±7.7a	20.7±3.4a	2.7±0.3b	18.9±3.9a			
1997	33.2±6.0a	20.8±2.2a	27,8±4.3a	20.0±2.1a	0.2±0.2b	19.5±4.8a			
1998	30.0±2.2a	22.8±0.7ab	18.7±3.8b	18.0±0.8b	0.1±0.1c	19.0±3.8b			
Aicrotus longicandus									
1994	7.7±1.8	3,6±2.5	6.8±4.3	j.	7.1±3.4	Î	$F_{[3,8]} = 0.35, P = 0.79$		
	В	В	В	в	V	в			
1995	8.5±2.9	1.2±0.2	2.6±0.6	1.4 ± 0.7	6.4±2.3	3.5±1.8	$F_{15,101} = 4.30, P = 0.02$	$F_{\rm FA, 461} = 10.43, P < 0.01$	FILENSI = 1.73, P =
9661	0.0	0.0	0.4±0.3	0.2±0.2	2.6±0.6	0.4 ± 0.3			Include
1997	0.0	1.3±0.6	1.9 ± 0.9	1.5±1.1	9.6±2.3	2.5±1.3			
1998	0.0	3.4±1.5	4.5±1.9	2.6±1.6	16.2±3.5	3.9±2.0			
peromyscus maniculan									
1994	11.0±2.1	7.3±3.8	8.6±1.0	J,	4.4±1.0	i	$F_{[1,8]} = 1.45, P = 0.30$		
1995	3.1±2.2	0.4 ± 0.2	0.4 ± 0.2	2.0±1.9	0.8±0.2	2.2±1.4	$F_{[5,10]} = 0.11, P = 0.99$	$F_{[3,36]} = 6.90, P < 0.01$	$F_{115,361} = 0.69, P =$
1996	1.2±0.7	0.4±0.3	1.3±1.3	3.3±2.1	2.0±1.5	0.8±0.7			
1997	1.5±1.3	4.0±3.4	1.9±1.7	4.5±7.2	0.8 ± 0.4	0.2±0.2			
1998	3.8±1.0	3.8±1.7	5,1±1.4	3.2±1.9	5.0±2.0	4.4±1.9			
lamias amoenus									
1994	0.1±0.1	0.2±0.2	1.0±0.4	Į,	0.3±0.2	Î,	$F_{10.01} = 3.21, P = 0.08$		
	U	BC	A	AB	~	BC			
1995	0.2±0.1	1.1±0.5	3,4±0.3	2.2 ± 0.8	1.9 ± 1.0	0.5±0.4	$F_{15,101} = 6.46, P < 0.01$	$F_{[1,06]} = 1.47, P = 0.24$	$F_{[15,36]} = 0.90, P =$
1996	0.8 ± 0.8	0.8 ± 0.4	3.4±0.7	2.4±1.1	4.8 ± 0.4	0.7±0.4			
1997	0.2±0.0	1.1±0.6	3.3±0.3	2.4±1.0	6.0±0.3	1.1±0.9			
1998	0.640.6	14+0.2	P.0+0 C	0.040.0	17+07	1.6+0.9			

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Fig. 3. Mean (n = 3 replicate sites) population density of long-tailed voles (Jolly-Seber) for the six treatments, 1994–1998.



Abundance of *T. amoenus* was significantly ($F_{[5,10]} = 6.46$; P < 0.01) different among sites, with the highest numbers recorded in the clearcut and 0.1-ha patch cuts during 1995–1997 (Table 1); mean abundance ranged from 0.1 to 6.0/ha.

Reproduction and recruitment

Reproductive performance, as measured by the number of

successful pregnancies, was significantly ($F_{[5,10]} = 65.52$, $P_{(5,10)} = 65.52$,

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= 65.52 /* no difference sites come nancies, set i5-1998 (he e appeared to 0.01).

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Table 2. Mean \pm SE (n = 3 replicate sites) number of successful pregnancies for *Clethrionomys gapperi*, *Microtus longicaudus*, and *Peromyscus maniculatus* for the six treaments during 1 pretreatment year (1994) with results of one-way ANOVA and 4 post-treatment years (1995–1998) with results of RM-ANOVA.

	ncut forest	selection	0.1-ha patch	1.0-ha patch	clearcut	Edge	Treatment	Time	
Clethrionomys gapperi	0 8 9					0		100	Ireatment × time
1994 18	3±1.3	11.3±1.7	18.0±4.0	1	15.3±3.3	1	$F_{11} = 1.34 \ P = 0.33$		
V		AB	AB	В	c	в	toret		
1995 16	1.7±5.8	22.7±4.7	19.0±2.1	14.0±2.5	4.7±1.5	14 0+3 2	F = 65.53 P > 0.01	E - 164 D - 001	
1996 18	.7±0.3	13.0±2.3	13.0±3.5	10.0±2.5	2.3±0.7	9.7+4.3	teen < a taccom = [01'cl -	10'0 > 1 'borb = [96'6] 1	r[15,36] = 1.80, P =
1997 23	.0±4.0	13.3±0.9	13.7±3,4	12.7±2.3	0.0	8.7±1.7			
1998 15	7±0.3	12.0±0.6	8.3±2.0	10.7±0.3	0.0	11 3+2 8			
Microtus longicandes									
1994 4	.0±1.0	2.7±1.5	3.7±3.7	1	6.7±3.3	1	$F_{1,,} = 0.43$ $P = 0.74$		
8		В	В	B	A	н	Nim - mount [girl] -		
1995 5	0±3.6	0.7±0.3	1.7±0.9	0.3±0.3	43±1.2	3.7±1.9	F_{ii} = 4.18 $P = 0.03$	F A 11 B - 0.01	r 0,40 r
0 9661	0	0.0	0.0	0.0	2.7±1.2	0.0	come + tores = [of'c] .	13.96 = 4.11, $t = 10.01$	r [15,36] = 0.08, P =
0 1997 0	0	2.3±1.5	1.0±0.6	0.7±0.7	7.3±1.5	2.3+1.5			
1998 0.	0	2.0±1.5	2.3±1.9	1.0±1.0	12.7±2.0	3.7±1.9			
Peromyscus maniculatus									
1994 0.	7±0.3	1.0±1.0	1.3±0.3	i	0.0	1	$F_{1111} = 1.06$ $P = 0.47$		
1995 0.	7±0.7	0.3±0.3	0.3±0.3	0.7±0.7	0.0	0.7+0.3	For a 10.0 P = 0.04	F - 0.14 D - 0.01	
1996 0.	0	0.0	0.0	0.7±0.7	0.7±0.7	0.0	Lora = 1 town = [01'c] .	r (1.36) = 0.14, r < 0.01	$r_{[15,36]} = 1.07, P =$
1997 0.	3±0.3	0.3±0.3	0.3±0.3	1.3±1.3	0.3±0.3	0.0			
1998 1.	3±0.9	1.7±0.9	1.0±0.6	1.0±1.0	3.7±1.7	0.7±0.3			

P

cauaus also differed ($F_{[5,10]} = 4.18$, P = 0.05) alloing sites. At 2 years postharvest, *M. longicaudus* on clear-cut sites had the most successful pregnancies, and this trend continued in 1997 and 1998 as well, with the edge sites in both years and the single tree selection sites in 1997 having numbers comparable with the clear-cut sites (Table 2).

This measure of reproductive effort was similar ($F_{[5,10]} = 0.23$, P = 0.94) among sites for *P. maniculatus* and was considerably lower than that for *C. gapperi* or *M. longicaudus* throughout the study (Table 2).

Recruitment of new C. gapperi was significantly ($F_{[5,10]} = 40.80$, P < 0.01) different among sites, with lower numbers of first captures in the clearcut than in the other sites (except edge sites in 1995) in all postharvest years (Table 3). Mean number of recruits ranged from 66.0 to 82.0 among sites in the preharvest year of 1994. Thereafter, mean numbers of recruits were similar among the uncut and partially harvested sites.

Recruitment of *M. longicaudus* was also significantly $(F_{[5,10]} = 9.74, P < 0.01)$ different among sites, with the clear-cut sites at high numbers of first captures in 1996–1998 (Table 3). Edge sites in 1997–1998 and the 0.1-ha patch cut site in 1998 had numbers of vole recruits similar to those of the clear-cut sites.

Numbers of *P. maniculatus* recruits were similar ($F_{[5,10]} = 0.17$, P = 0.97) among sites (Table 3). However, recruitment did appear to change significantly ($F_{[3,36]} = 5.07$, P < 0.01) with time, being high in 1994 and again in 1998. Recruitment of *T. amoenus* was also variable but was significantly ($F_{[5,10]} = 5.68$, P < 0.01) different among sites (Table 3). The harvested sites and edges tended to have somewhat similar levels of chipmunk recruitment in the postharvest years.

Survival

Mean estimates of Jolly-Seber survival for *C. gapperi* were similar among sites during the summer and winter periods but declined dramatically on clear-cut sites (Table 4). Mean survival for *M. longicaudus* and *P. maniculatus* was variable across seasons and sites. An index of early juvenile survival was similar among sites for *C. gapperi* but again was negligible on clear-cut sites in 1997 and 1998 (Table 5). Early juvenile survival was variable across sites for both *M. longicaudus* and *P. maniculatus*.

Discussion

Small mammal communities

This investigation is the first evaluation of the population dynamics of small mammals over a range of harvesting treatments in high-elevation spruce-fir forests. Although 10 small mammal species were sampled during the study, the majority captured were *C. gapperi*. Abundance of this microtine peaked at 80 animals/ha with mean values of 31-50 voles/ha, indicating that these old-growth Engelmann spruce – subalpine fir forests provided prime habitat for *C. gapperi*. This finding was similar to reports elsewhere for *C. gapperi* in montane and old-growth forests dominated by spruce or true firs in western North America (Raphael 1988; Nordyke and Buskirk 1991; Hayward and Hayward 1995; Sullivan et al. 1999). Subalpine and boreal spruce-fir forests fungi, which are major determinants of putimum habital C. gapperi (Getz 1968; Maser et al. 1978). In define woody debris in these forests may provide protection durcritical periods of freezing and thawing Merritt and Merrit 1978; Merritt 1985).

Clethrionomys gapperi is considered an ndicator of a growth forest conditions (Nordyke and Buskirk 988). The definition is partly attributable to the above tabitat requirements and also because this microtine is an important species for several carnivores (Hayward and Verner Ruggiero et al. 1995). Therefore, maintenance of C. tabitation in forest harvesting treatments should also be potentially to vourable to predators (Garton et al. 1989).

Response to harvesting treatments

Because C. gapperi are closely associated with closed canopy forests, we expected them to respond negatively a terms of abundance, reproduction, and survival to decrease levels of tree retention, as per the prediction of hypothese (i). However, these demographic attributes were consistent similar among the uncut forest, single tree selection. The patch cut treatments compared with clear-cut sites where C. gapperi declined markedly. In most years, population harvesting treatments. Thus, our hypothesis is only partially supported, at least with respect to total removal of tree cour on clearcuts and the partial harvesting treatments (in Fig. 4).

Medin (1986) similarly reported that C. gapperi disap peared from plots when the basal area was reduced from 20.7 to 5.1 m²/ha, but no difference in numbers of the microtine was recorded after one third of the stand volume was removed (Medin and Booth 1989). In contrast, Moether and Soutiere (1985) reported greater abundance of C. peri in partially cut stands (50% basal area reduction) regenerating clearcuts than in uncut stands. West d (1980) reported similar numbers of northern red-backed voles (Clethrionomys rutilus) in shelterwood and uncut in est 1 year after harvest. Von Trebra et al. (1998) and Serve ton et al. (1998) reported higher abundance of C. gappen light removal and shelterwood stands than in uncut fores the first 1 or 2 years postharvest in west-central British Co lumbia. Retention of 10% basal area and woody seemed to maintain C. gapperi in aspen mixedwood from (Moses and Boutin 2001).

Although not statistically significant, we also observed slightly higher numbers of *C. gapperi* in the single tree lection and 0.1-ha patch cut than in the uncut forest first year after harvesting (Table 1; Fig. 2). As discussed Von Trebra et al. (1998), some partial cutting treatment "light disturbance" of the forest may temporarily call local forage and cover by increasing and seeds that were dislodged from the canopy or expect the duff layer of the forest floor. Tree removal may hypogeous fungi and some cover requirements may imbecause of additional coarse woody debris on he site logging (Tallmon and Mills 1994).

Similar to the results reported by Hayward and Construction and Sullivan (2001), C. gapperi

ring 1 pret Uncut forest 66.0±3.1 62.0±4.0ab 70.7±2.4a 70.7±2.4a 72.3±12.9a 35.0±2.5a	Single tree selection 79.7±20.4ab 62.7±2.8a 50.3±3.7a 29.3±1.5a	0.1-ha patch 77.7±2.9 88.0±6.1a 64.0±4.6a 44.7±6.3a 23.0±5.1ab	1.0-ha patch 66.3±10.4ab 48.7±3.4a 38.7±7.4a 22.3±0.3ab) for Clethrin -way ANON -way ANON 10.0-ha clearcut 82.0±12.0 28.0±1.2c 6.0±2.1b 1.0±1.0b 0.3±0.3c 	A and 4 pos A and 4 pos Edge 43.3±2.4a 34.3±4.9a 13.3±4.9a	Treatment years (1995– Treatment $F_{[3,8]} = 0.59, P = 0.64$ $F_{[5,10]} = 40.80, P < 0.01$. Peromyscus maniculatus (998) with results of RM. Time $F_{[3,36]} = 44.50, P < 0.01$, and <i>Tamias</i> ANOVA. Treatment × F[15.36] = 5.9
b±3.1 b±4.6a bc bc	10.7±3.7 3.3±0.3ab 0.0c 4.7±1.8b	17.7±8.7 7.3±2.4ab 1.3±0.9bc 7.0±2.5b	- 3.7±2.0b 1.0±0.6bc 4.3±2.0b	18.7±8.7 20.0±7.5a 9.0±1.0a 22.3±2.6a	10.0±2.5ab 2.0±0.6b 10.3±1.8ab	$F_{[3,8]} = 0.31, P = 0.82$ $F_{[5,10]} = 9.74, P < 0.01$	$F_{[3,36]} = 22.56, P < 0.01$	$F_{[15,36]} = 3.5$
x H±5.3 H±1.5 H±1.5 H±1.5 H±1.5	6.3±3.8b 21.3±7.2 1.7±0.7 1.0±0.6 6.3±4.4 7.7±2.4	9.0±5.1ab 26.0±3.0 1.0±0.6 5.3±3.9 13.0±2.5 13.0±2.5	4.7±2.0b 4.7±4.2 8.3±5.6 8.7±6.7 6.3±3.5	26.3±4.3a 11.7±0.9 2.3±1.2 8.0±5.1 3.3±0.7 13.7±4.7	9.3±2.8µb 3.3±2.4 1.0±0.6 12.0±4.9	$F_{[3,03]} = 1.66, P = 0.25$ $F_{[5,10]} = 0.17, P = 0.97$	$F_{[3,36]} = 5.07, P < 0.01$	$F_{[15,36]} = 0.5$
±0.3 ±0.3 ±0.7 ±0.0	0.7±0.7 CD 3.0±0.6 1.7±0.9 2.3±1.5	2.3±0.9 AB 4.7±1.7 5.0±1.5 7.3±0.3		3 3±1 3 A 3 7±0 3 9 7±1 5 9 3±0 7	 BCD 2.0±1.0 3.0±1.0 3.0±0.6	$F_{[3,16]} = 2.23, P = 0.16$ $F_{[5,10]} = 5.68, P < 0.01$	$F_{[3,36]} = 0.30, P = 0.82$	$F_{[15,36]} = 0.70$

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		Single tree			10.0-ha	1			
Year	Uncut forest	selection	0.1-ha patch	1.0-ha patch	clearcut	Edge	Treatment	Time	Treatment × time
Summer	145)	1 1000 12 11		200 - 100 - 100 - 100 - 100 - 100 - 100 - 100 - 100 - 100 - 100 - 100 - 100 - 100 - 100 - 100 - 100 - 100 - 100					
1994	0.70±0.17	0.80±0.07	0.62±0.11		0.67±0.03		$F_{[3,8]} = 0.53, P = 0.68$		
1995	0.60±0.11	0.63±0.09	0.52±0.07	0.54±0.04	0.39±0.11	0.66±0.03	$F_{[5,8]} = 0.94, P = 0.50$	$F_{[2,20]} = 11.96, P < 0.01$	$F_{[10,20]} = 5.90, P < 0$
1996	0.68±0.02	0.61±0.04	0.70±0.04	0.64±0.07	0.66±0.17	0.58±0.03			
1997	0.81±0.03	0.70±0.04	0.81±0.04	0.80±0.07	0.30	0.64±0.03			
Winter									
1995–1996	0.76±0.01	0.78±0.04	0.80±0.02	0.78±0.02	0.73±0.04	0.76±0.07	$F_{[4,8]} = 0.32, P = 0.85$	$F_{[2,20]} = 5.07, P = 0.02$	$F_{[8,20]} = 0.52, P = 0.52$
1996-1997	0.80±0.01	0.73±0.05	0.83±0.01	0.82±0.03	0.24±0.24	0.81±0.03			
1997-1998	0.84±0.03	0.86±0.01	0.83±0.02	0.84±0.02		0.85±0.02			

Table 4. Mean \pm SE (n = 3 replicate sites) estimates of Jolly-Seber 28-day survival for *Clethrionomys gapperi* for the six treatments during 1994 (summer pretreatment) with results of a one-way ANOVA and 1995-1998 (summer and winter post-treatment) with results of RM-ANOVA.

Table 5. Mean \pm SE (n = 3 replicate sites) index of early juvenile survival (numbers of juvenile recruits per successful pregnancy) for *Clethrionomys gapperi* for the six treat ments during 1 pretreatment year (1994) with results of a one-way ANOVA and 4 post-treatment years (1995–1998) with results of RM-ANOVA.

Year	Uncut forest	Single tree selection	0.1-ha patch	1.0-ha patch	10.0-h clearcut	Edge	Treatment	Time	Treatment × time
	2	2.23±0.23			1.47±0.21		$F_{[3,8]} = 4.59, P = 0.04$		
	1	l.89±0.09			4.55±1.99	2.02±0.44	$F_{[4,8]} = 0.73, P = 0.60$	$F_{[3,30]} = 19.41, P < 0.01$	$F_{12,30} = 0.35, P = 0.5$
	3	3.18±0.36			1.56±0.87	4.27±1.44			
	1	1.44±0.12				2.20±0.66			
	(0.61±0.26				0.27±0.09			

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confidence intervals for deer mice and northwestern chipmunks the six treatments in the 4 postharvest years, 1995-1998.



The maintained over the range of patch cuts up to, and initing, 1.0 ha in area. Hayward et al. (1999) also reported C. gapperi used edge habitats in a manner similar to C. gapperi used edge habitats. Although overall mean mance of *C.* gapperi was lower in the edge than in the forest, numbers of this microtine were similar in edge partially harvested sites (Fig. 4). Conversely, Mills C. gapperi was lower in the edge than in the partially harvested sites (Fig. 4). Conversely, Mills C. avoided edges in a highly fragmented and scape.

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Other studies from western North America also reported bundance of C. gapperi declined on clearcuts wiler 1970; Martell and Radvanyi 1977; Sullivan et al. Moses and Boutin 2001; Sullivan and Sullivan 2001). Wer, Kirkland's (1990) review of 21 published studies ded that the species principally responsible for the abundance of microtine rodents on recent clearcuts C. gapperi, which increased in abundance in 15 of 21 This apparent contradiction may relate to differences of C. gapperi in some parts of eastern North America forest harvesting by clear-cutting has occurred for at 300 years. It is possible that C. gapperi may have to conditions in human-managed forests whereby and physiological changes allow substitution of lichens in their diet by other items such as seeds, leaves, and berries (Schloyer 1977; Gliwicz and Glowack: 2000).

Microtus longicaudus occurred predominantly in the clear-cut sites (Fig. 4). This pattern was similar to that recorded for Microtus spp. in other studies comparing a range of harvesting treatments (Moses and Boutin 2001; Sullivan and Sullivan 2001). The persistence of M. longicaudus on a variety of sites in our study fits the range of habitats occupied by this vole (Van Horne 1982; Smolen and Keller 1987). Habitats with some open areas and shrub and sapling cover up to 10 years after clear-cutting were optimum for M. longicaudus in Alaska (Van Horne 1982). Lower densities of this microtine occur in later seral stages that have less understory vegetation and a greater degree of canopy closure. High numbers of M. longicaudus in 1998 on clear-cut sites was similar to the high populations of this species recorded in clear-cut and seed-tree sites in a Douglas-fir (Pseudotsuga menziesii (Mirb.) Franco) - lodgepole pine (Pinus contorta Dougl.ex Loud.) forest in 1998 (Sullivan and Sullivan 2001), thereby supporting the presumed synchronous nature of some vole population fluctuations (Taitt and Krebs 1985; Krebs 1996).

Tamias amoenus also was associated with early successional habitat on our clear-cut sites, albeit at overall mean numbers of <5 animals/ha (Fig. 5). Tamias amoenus

edge sites. Subalpine forests in the southern interior of British Columbia appear to provide less suitable habitat for *T. amoenus* than mixed Douglas-fir – lodgepole pine forests at lower elevations (Sullivan et al. 2000; Sullivan and Sullivan 2001). Thus, hypothesis (*ii*), that abundance and related demographic attributes of *M. longicaudus* and *T. amoenus* will increase with decreasing levels of tree retention, seems to be supported.

Overall mean abundance of *P. maniculatus* was low (<4 animals/ha) but similar among sites (Fig. 5). Our prediction, hypothesis (*iii*), that a habitat generalist such as *P. maniculatus* will have similar demographic responses across a gradient of tree retention, appears to be supported and fits the prediction of Lehmkuhl et al. (1999) that habitat generalists will increase or have no change in abundance with decreasing levels of tree retention.

Experimental design

Our study was a classical randomized block design with blocks assigned on the basis of elevation and treatments randomly assigned to sites within each block (Zar 1999). There were three true replicates of treatment sites; however, it could be argued that these replicate units were not spatially segregated because the three blocks were contiguous (Hurlbert 1984). Ideally the individual blocks should have been located in independent areas of Engelmann spruce – subalpine fir forest in the south-central interior of British Columbia. Such a design would have permitted extrapolation of results and conclusions to a wider geographical area. However, the logistics and costs of establishing such a design were prohibitive and may not have achieved the desired experimental rigour needed for some components of this multidisciplinary study.

Treatment sites were the size of typical forestry operations in southern British Columbia and perhaps for subalpine forests in cordilleran regions of western North America. The study design tested operational-scale treatments of varying levels and patterns of green-tree retention on various forest ecosystem components as per Franklin et al. (1997), Aubry et al. (1999), and Sullivan et al. (2001). Inferences from our study reflect responses in small mammal communities in the first 4 postharvest years only. Longer-term (>5 years) measurement of responses will require periodic repeated sampling over decades to provide a clear understanding of the impact of these treatments.

The pretreatment year provided an important benchmark for assessing responses of small mammals to harvesting treatments. Ideally, all treatments should have been represented, but there were no pretreatment data for the 1.0-ha patch cut and edge sites because the precise locations of the treatment boundaries were not established until 1995. Responses of the forest floor small mammal communities to treatments represented relatively short summer and early autumn seasons and may not have been the same during winter months. However, there were 4 years of postharvest data and changes in abundance from one year to the next, during the overwinter periods when data were not available, were not qualitatively different from those of other forest small mammal studies. The results of our study suggest that partial harvesting high-elevation spruce-fir forests, at least up to and includ 1.0-ha patch cuts or openings, did not negatively affect sr mammal populations. Clethrionomys gapperi accounted 66.9% of the small mammal community in these forests a except for the 10.0-ha clear-cut site, were not affected terms of abundance, reproduction, or survival attributes of a 4-year postharvest period. Early successional species su as *M. longicaudus* and *T. amoenus* preferred the clearsites but occurred at low numbers.

Our results suggest that the partial harvesting (single t selection to 1.0-ha patches) of these forests may be one; tential method for combining timber extraction with main nance of mature forest habitat (McComb et al. 1993; Hanet al. 1995). Clearly, not all forest harvesting activities the same, as is often misconstrued by researchers and en ronmentalists in their communications to managers. As edenced by this and other studies, harvesting systems : quite different from each other and clearly have different fects on forest-dwelling wildlife species. Thus, it may possible to conduct harvesting activities in these subalpforests without disrupting the small mammal communite

Our experimental harvesting treatments were designed create a range of postharvest conditions that would help cl. ify the influence of residual overstory condition on sm. mammal communities. The treatments were conceptual nature but serve to illustrate the role that residual trees a gap size play in influencing small mammal communities. a more complex operational environment, topography a stand conditions will necessitate modifications to the shap and size of openings, but the basic principles of small mar mal community response to opening size and forest structu should remain applicable.

It is clear from our results that no single treatment is : ther beneficial or detrimental to all species of small ma: mals monitored at the Sicamous Creek study area. addition to the stand-level habitat effects on small mammdescribed in this paper, the operational application of vharvesting treatments at large spatial scales should consid a broader set of implications to small mammals and v overall biotic community. For example, partial-cut harve ing systems can maintain important structural features in " postharvest stand but create a more extensive network roads that may have undesirable ecological consequence (Forman et al. 2003). Similarly, treatment effects are me gated over time through forest growth and succession. the long-term habitat suitability of different treatments new to be considered. Balancing the habitat requirements small mammals and other species of wildlife will require careful assessment of the costs and benefits associated •* treatments that are evaluated in a landscape context.

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