# **Using bear rub data and spatial capture-recapture models to estimate trend in a brown bear population**

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## **SUPPLEMENTARY INFORMATION**

## **Introduction**

Although bear populations are challenging to monitor, bears possess an attribute that facilitates sampling them. Bears routinely rub on trees and other objects vigorously, often leaving hair behind. The most likely explanation for this ubiquitous activity is that it is a form of marking behaviour and functions as chemical signaling to other bears<sup>1</sup>. This tendency to rub results in the deposition of hair from individual bears at multiple sites. Because bears often use trails to travel, rub objects along trail networks can be located visually and therefore represent entirely passive detection; the sampling process has no effect on bear behaviour. While bears with a previous history of capture and handling have lower detection rates at baited hair corrals than bears that have not been trapped<sup>2</sup>, no such effect has been found at bear rubs<sup>3</sup>.

Grizzly bear and black bear (*U. americanus*) marking behaviour is well documented, if imperfectly understood<sup>4,5,6,7</sup>. Bears of all sex and age classes rub throughout the active season with variable frequency<sup>8</sup>. Evidence of rubbing is found throughout occupied habitat off-trail as well as along maintained and wildlife trails and by roads. Bears do not appear to discriminate much in selecting objects to mark. Bear hair has been found on a wide variety of natural (e.g. trees of all sizes and taxa, stumps, rocks) and man-made objects (e.g. sign and fence posts, utility poles, bridges, cabins)<sup>8,9,10</sup>. Female and subadult grizzlies rarely rub during the breeding season (May—June), however, their detection probability increases after low rates in spring and approaches male rates by late summer<sup>3,8,11,12</sup>.

Many studies have demonstrated that genetic detection through hair collection is a highly effective way to sample bear populations; hair collection from a grid of baited hair corrals has been used extensively to study bear population abundance throughout the world (e.g. 13,14,15,16,). Kendall *et al.*<sup>3,8</sup> added hair collection at natural bear rub sites as a secondary sampling approach to a primary method using an array of baited hair corrals. Because there was only partial overlap in individuals detected at hair corrals and rubs, the addition of this second sampling approach increased sample coverage and population abundance estimate precision while abundance estimates themselves were not significantly changed<sup>17</sup>. Results of subsequent studies that used concurrent sampling at hair corrals and bear rubs also found that the addition of rub site sampling was a cost-effective way to increase sample size and estimate precision<sup>11,15,18</sup>. While detection rates per sample site are higher at baited hair corrals than at unbaited rub sites<sup>3,8,17,19,20</sup> it is possible to sample a similar or larger portion of the population at rub sites than at baited corrals<sup>11,18</sup>. Sampling at rubs can also be more cost effective than at

baited hair corrals. In most studies using both methods, more field crew time was put into sampling at baited corrals than at rubs even when rubs greatly outnumbered corrals. For example, the ratio of field crews assigned to hair corral vs. rub sampling was over 4:1, while the ratio of individuals detected at corrals and rubs ranged from 1:1 to 2.5:1 for the studies conducted by Kendall *et al.*<sup>3,8,18</sup>. However, the overall efficiency of rub tree sampling depends on monitoring higher numbers of rub trees to offset the lower per site sampling efficiency of rub trees<sup>19,20</sup>. Therefore, rub tree sampling is most viable in areas with greater access, such as maintained trails.

## **Methods**

## **Study area**

Approximately 85% of the Northern Continental Divide Ecosystem (NCDE) is in public ownership, where grizzly bear conservation is a management objective. It includes Glacier National Park (NP), parts of 4 National Forests (Flathead, Kootenai, Helena-Lewis and Clark, and Lolo), five Wilderness areas (Bob Marshall, Great Bear, Mission Mountain, Rattlesnake, and Scapegoat), three Wilderness study areas (Mount Hefty-Tuchuck, Ten Lakes, and Thompson-Seton), Bureau of Land Management, U.S. Fish and Wildlife Service, and State land, and approximately half of the Blackfeet and one-third of the Flathead Indian Reservations. Private lands comprising the remaining 15% of the NCDE occur chiefly along the eastern, southern and western peripheries, but the Clearwater and Swan River Drainages, which are interior to the NCDE, contain both extensive diffuse housing and several small towns. Waterton Lakes National Park, Alberta and Akamina-Kishinena Provincial Park and provincial Crown lands in British Columbia, Canada border the NCDE to the north.

The climate and vegetation of the NCDE are shaped by the uplifted fault-blocks which formed the Rocky Mountains and by Pleistocene era ice ages<sup>21</sup>. Higher elevations receive more precipitation than lower slopes and valleys with annual precipitation ranging from over 170 cm in the higher elevations on the west side of Glacier NP to less than 38 cm on the eastern border. The Pacific maritime-influenced climate west of the Continental Divide is moister and more temperate than the colder and drier continental climate on the eastern side. The rugged terrain creates an array of habitats from alpine tundra and subalpine forests to moist mountain valleys, foothill groves, and semiarid grasslands in the rain shadow of large mountain masses<sup>22</sup>. Vegetation is characterized by coniferous forest, shrub fields and alpine tundra in the mountains, mixed deciduous–coniferous woodlands and herbaceous meadows in the valleys, and aspen (*Populus tremuloides*) stands, prairie grasslands, and agricultural fields along the eastern boundary<sup>3</sup>.

## **Field methods**

To identify bear rubs, we closely examined objects that included one or more of the following characteristics; smoothed and/or discoloured bark and branch stobs, bare ground at the base of trees/poles, entrenched bear trails leading to rubs from maintained trails, and bear bite or claw marks. We attempted to get broad and even geographic distribution of areas searched for rubs and did not prioritize areas thought to have higher densities of bears. In lower elevation areas

where there were fewer trees for bears to rub on, utility poles and fence lines provided sampling opportunities. To prevent damage to pack stock and their cargo, we used doublestranded smooth wire on trees that appeared to have been bumped by horse packs. We separated the twisted ends of smooth wire slightly to make them more effective in snagging hair. We visited rubs at least twice annually. All hair deposited on the wire was collected during each visit. The hair found on rubs during the initial visit each year was not analyzed and did not contribute to detection histories because it could have been deposited the previous year. A flame was passed under each barb and wire after sample collection, and hair between wires was removed to prevent contamination when the next bear rubbed. All hairs from a 4 point barb or wire end were considered one sample.

Rubs were initially identified and set up in 2003 for sampling in 2004. Because we removed tags and wire on monitored rubs at the end of the 2004 field season, the rub survey network was reestablished in 2009. The time required for initial set up meant that rubs in some remote areas were not set up until late summer 2009 or spring 2010, leading to lower rub sampling effort and extent in 2009 than in all other years.

#### **Genetic analysis**

In 2004, we attempted genetic analysis on all hair samples collected. During 2009—2012, we subsampled hair for analysis to limit genotyping costs. When we collected more than one hair sample during a visit to a rub, we genotyped the two samples with the most and/or largest follicles to maximize the amount of DNA in the sample. If sample quality was equal, we selected non-adjacent samples to increase the probability of detecting more than one individual per site. If there were distinct differences in hair colour between samples, we selected the best 2 samples from each, if available.

## **Spatial capture-recapture modeling**

We included several trap covariates to describe detection. We calculated security level based on the location of a trap, where National Park Service =10, Forest Service =7, State and private forest lands =3, and all other =1. We used the standard deviation of curvature within a 250-m buffer of the trap covariate based on Ironside *et al*. 23 , which found it to be the most robust measure of terrain ruggedness. We also evaluated the effects of percent canopy cover based on the cover layer from Landfire, using the most recent data for that year. We updated the 2004 layer to reflect reductions in canopy cover due to some logging and extensive fires between the 2000/2001 imagery and the 2004 dataset.

#### **Results**

## **Rub sampling effort**

Sampling primarily occurred in forested habitats, however, bears' predilection to rub on fence and utility poles ensured sampling opportunities in prairie and other treeless habitats and agricultural areas. In 2004, search effort to identify rubs was lower along the eastern edge of our study area because there were few trails or little bear sign to guide our efforts. Sampling in this area was primarily in riparian corridors. Anecdotal information also indicated that few

bears ranged east of the Rocky Mountain foothills at that time. After 2004, we expanded the number of monitored rub objects along the eastern foothills and plains as bears began to reoccupy former range. Starting in 2009, additional rub objects on the eastern edge were, therefore, added to the monitoring network. Rubs continued to be identified and added as sampling sites during early 2010 and then they generally remained in our monitoring network for the duration through 2012. A small number (<1%) of trees blew down or could not be relocated during our study; however, other monitored rubs were typically < 1 km away and in some cases, nearby new rubs were added to the monitoring network to replace them. As a result, sampling distribution and effort was nearly identical 2010—2012.

## **Discussion**

Our point estimate for the annual rate of change, 1.043, was fairly high for an interior grizzly/brown bear population. Stable growth rates ( $\lambda \approx 1.0$ ) have been reported in the past for a number of interior populations in the Rocky Mountains of the United States and Canada (e.g. <sup>24,25</sup>). More recently, however, several populations have grown at rates comparable to those we found in the NCDE, likely a reflection of demographic recovery in response to conservation measures<sup>26,27,28</sup>. A study examining demographic mechanisms underpinning genetic assimilation of remnant groups of grizzly bears in the NCDE found that the increases in genetic diversity they identified resulted primarily from immigration<sup>29</sup>. It is likely that a combination of growth from within the remnant population coupled with bears moving in from higher density areas and the mathematics of small numbers in a local area (a change from 1 to 2 bears is a 100% increase) was responsible for the highest rates of pixel-specific growth we found in the NCDE (Figure 3). Quantifying local change rates using alternate measures such as gain or loss in density per pixel may be useful in smoothing this effect.

Bear species throughout the world exhibit rubbing behaviour; e.g., grizzly<sup>30</sup>, European brown bear<sup>31,32</sup>, Japanese brown bear<sup>33</sup>, American black bears<sup>34</sup>, and Asiatic black bears<sup>15</sup>. A growing number of studies have taken advantage of this behaviour to study bear demography (e.g.,  $^{8,11,35}$ ), dispersal<sup>29</sup>, and behaviour<sup>7,36</sup>. Developing a better understanding of which bears rub, why they rub, and where and when they do it will make this sampling method more powerful. The data obtained from genetic detection can inform multiple aspects of population status and mechanisms. For example, studies of habitat effects on abundance<sup>38</sup>, connectivity and dispersal<sup>38,39</sup>, the role of social learning<sup>36</sup>, genetic assimilation in remnant subpopulations<sup>29</sup>, and the relationship between effective and demographic population size in continuously distributed populations<sup>40</sup>, were based on the genetic samples collected for this and earlier genetic detection studies in the NCDE<sup>3,8</sup>.

Efficient sampling methods increase sample coverage while decreasing monitoring costs. The number of individual bears we sampled through hair collection at rub sites was an order of magnitude larger and more evenly spatially distributed, e.g. in the expansion area, than would typically be possible through live capture methods. Thus, we expect that genetic monitoring will be more sensitive to spatial and temporal changes in population demography than approaches that involve fewer individuals. Our genetic detection network yielded large sample sizes and precise estimates of population growth rate with 5 years of rub detection data (2004, 2009—2012). Other approaches to obtaining similar spatial data, such as a grid of bear hair corrals, cost millions of dollars to implement across a region of this size<sup>8</sup>; repeated sampling to obtain trend estimates using these methods across areas as broad as the NCDE has to date proved unsustainable. It may be possible to monitor populations such as ours with adequate precision with less sampling effort and at lower cost than in this study. The use of multiple detection methods<sup>17</sup>, clustered and stratified sampling designs<sup>19,35</sup>, adding telemetry data, or integrated population models that combine SCR data with lower resolution information<sup>41,42</sup> could further reduce the sampling and, thus, expense of SCR-based studies in the future.

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Table S1. Description of full set of candidate models developed to estimate grizzly bear population density using spatial capture recapture data from northwestern Montana, USA. Spatial detection histories were developed through genetic identification of individual bears from hair collected at natural bear rub sites. Model 17 (= Model 8 + a term for translocated bears) and Model 18 (= Model 16 + translocations) did not converge because few individuals were identified as having been translocated in our dataset.



a Model notation:



Table S2. Female and male parameter estimates (untransformed) for the most supported models (Table 3) developed to estimate trend 2004—2012 in a grizzly bear population using spatial capture-recapture data from northwestern Montana, USA.



a Parameter notation:



Table S3. Model selection results for females including all detections, i.e. **including one additional detection 190 km from the other detections of that bear not included in the models shown in Table 3**. Spatial covariates (spatial covs= cover, cover2, curv) were calculated within a 250-m radius of each rub sampling site. Only models with cumulative weights up to 1 are presented.



Table S4. Female parameter estimates (untransformed) for the most supported models (Table S3) using all detections, i.e. **including one additional detection 190 km from the other detections of that bear not included in the models shown in Tables 3 and 4**. Estimates for model 4 were nearly identical to model 3 and estimates for model 2 were nearly identical to model 1, so are not shown.



a Parameter notation:





Table S5. Point estimates (transformed) of female density (bears/1,000 km<sup>2</sup>), spatial scale of movement (σ), and baseline detection probability (*p*0) parameters for models developed to estimate grizzly bear population trend using spatial capture-recapture data from the Northern Continental Divide Ecosystem in northwestern Montana, USA. a. **Estimates that include one outlier detection of a single female in 2010 that was 190 km distant from other detections of that individual.** b. Estimates that exclude the outlier detection point. The baseline detection probability reported here is for b=0, initial behavior covariate (prior to the first detection), and all other covariates at their mean level. When the outlier was included, the best model had yearspecific density (model #3) versus a linear yearly trend in density (model #1) when the outlier point was excluded.





#### **b. Females – excluding outlier point. Best model is #1**



Figure S1. Locations of grizzly bear detections 2004, 2009—2012 (N=1,539) derived from sampling at natural bear rub sites in the Northern Continental Divide Ecosystem in northwestern Montana, USA. Maps were created from the USGS National Elevation Dataset<sup>43</sup> using ArcMap 10.2.

