${ }^{1}$ Heidi Swanson, Martin Lysy, Michael Power, Ashley Stasko, Jim Johnson, and James Reist.
2 2014. A new probabilistic method for quantifying n-dimensional ecological niches and niche overlap. Ecology

4 Appendices

5 Appendix A. Computation of niche regions.

## Appendix A: Computation of Niche Regions.

Suppose that $X=\left(X_{1}, \ldots, X_{n}\right)$ is an $n$-dimensional isotope distribution with joint pdf $f(x)$. We have defined the niche region to be

$$
\begin{equation*}
\mathrm{N}_{\mathrm{R}}=\{x: f(x)>C\}, \tag{A.1}
\end{equation*}
$$

where $C$ is chosen such that $P\left(X \in \mathrm{~N}_{\mathrm{R}}\right)=95 \%$. This is in fact the smallest region occupying $95 \%$ of the probability space (e.g., Box and Tiao, 1973; Wei and Tanner, 1990; Chen and Shao, 1999). Figure A1 illustrates this for a one-dimensional isotope $X$.

For a symmetric distribution (left panel), $\mathrm{N}_{\mathrm{R}}$ simply consists of the interval between the $2.5 \%$ and $97.5 \%$ quantiles of the distribution. For skewed distributions (middle panel), however, this is not the case. In fact, for bimodal distributions (right panel) the niche region can even be disjoint.


Fig. A1: Probabilistic $95 \%$ niche region for various univariate distributions. These are the intervals or collection of intervals highlighted in red. For symmetric distributions, these coincide with the $2.5 \%$ and $97.5 \%$ quantiles which are depicted with crosses. In each case the red interval(s) contain(s) the smallest region covering $95 \%$ of the probability distribution.

Of particular interest here is the case where $X=\left(X_{1}, \ldots, X_{n}\right) \sim \mathcal{N}(\mu, \Sigma)$ is an $n$-dimensional

2 multivariate Normal isotope distribution. The joint pdf of $X$ is

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$$
f(x)=(2 \pi)^{-n / 2}|\Sigma|^{-1 / 2} \exp \left\{-\frac{1}{2}(x-\mu)^{\prime} \Sigma^{-1}(x-\mu)\right\},
$$

which is a constant in $x$ for fixed $Q(x)=(x-\mu)^{\prime} \Sigma^{-1}(x-\mu)=q$. Since increasing $Q(x)$ results in smaller values of $f(x)$, the niche region for Normal data is

$$
\begin{equation*}
\mathrm{N}_{\mathrm{R}}=\left\{x:(x-\mu)^{\prime} \Sigma^{-1}(x-\mu)<C\right\} . \tag{A.2}
\end{equation*}
$$

Upon rearranging terms, $\mathrm{N}_{\mathrm{R}}$ can be written as the quadratic inequality

$$
\begin{equation*}
\mathrm{N}_{\mathrm{R}}=\left\{x=\left(x_{1}, \ldots, x_{n}\right): \sum_{i=1}^{n} \sum_{j=1}^{n} a_{i j} x_{i} x_{j}+\sum_{i=1}^{n} b_{i} x_{i}+d<C\right\} \tag{A.3}
\end{equation*}
$$

although the canonical form in A. 2 is computationally simpler to use for the niche overlap calculation described in Appendix B.

In order to determine the value of $C$ in A.2, we may utilize the fact that

$$
(X-\mu)^{\prime} \Sigma^{-1}(X-\mu)=Z^{\prime} Z=\sum_{i=1}^{n} Z_{i}^{2}
$$

where $Z=\left(Z_{1}, \ldots, Z_{n}\right)^{\prime}=\Sigma^{-1 / 2}(X-\mu) \sim \mathcal{N}(0, I)$. In other words, $Q(X)$ is the sum of the squares of $n \operatorname{iid} \mathfrak{N}(0,1)$ random variables, such that

$$
Q(X)=\sum_{i=1}^{n} Z_{i}^{2} \sim \chi_{(n)}^{2}
$$

Thus, for a $95 \%$ niche region we require that $P(Q(x)<C)=.95$, a value which can be

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1 OMalley, A.J. and Zaslavsky, A.M. (2008). Domain-level covariance analysis for multilevel
${ }_{4}$ Wei, G.C.G. and Tanner, M.A. (1990). Calculating the content and boundary of the highest posterior density region via data augmentation. Biometrika, 77(3): 649-652.

