



# The Effects of Area and Habitat Heterogeneity on Bird Richness and Composition in High Elevation Wetlands (“Bofedales”) of the Central Andes of Peru

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**Abstract** We assessed area and habitat heterogeneity effects on avian richness and composition in bofedales that differed in size and microhabitat diversity. We analyzed data collected in 2 seasons and 24 bofedales using General Linear Models, Ordinary Least Square models to establish the relationship of predictor variables on richness and Akaike Information Criterion for model selection. We evaluate composition classifying species into groups using Bray Curtis ordination, followed by Multiple Response Permutation Procedure to test for differences among groups, and Indicator Species Analysis to identify species. Bofedales differed in richness ( $F = 5.1, p < 0.001$ ) and microhabitat diversity ( $F = 23.4, p < 0.001$ ), but no seasonal differences emerged ( $p > 0.05$ ). The best model indicates that 54% of variance in richness was explained by area and microhabitat diversity, however, a tendency to decrease in

microhabitat diversity as area increases, suggests a relatively more important role of area. Results are supported by composition, as microhabitats not only differed pairwise ( $T = -94.14, A = 0.601, p < 0.001$ ) and had significant indicator species ( $p < 0.05$ ), but because its differential contribution to richness, as some microhabitats were more speciose than others. As such, few species-rich microhabitats may contribute more to richness than many species-poor ones which is not predicted by the habitat heterogeneity hypothesis. Disentangling the influence of area and habitat heterogeneity on species richness is important to establish conservation priorities that ensure bofedales integrity under imminent climate change.

**Resumen** Evaluamos el efecto del área y la heterogeneidad del hábitat en la riqueza y composición de aves en bofedales que difieren en tamaño y diversidad de microhábitats. Los datos recopilados en 2 estaciones y 24 bofedales fueron analizados usando Modelos Generales Lineales, Modelos de Mínimos Cuadrados Ordinarios para establecer la relación entre las variables predictivas y la riqueza, y el Criterio de Información de Akaike para seleccionar los modelos. Evaluamos la composición de especies clasificándolas en grupos con el Ordenamiento de Bray Curtis, seguido por el Análisis de Permutación de Respuesta Múltiple para detectar diferencias entre los grupos, y el Análisis de Especies Indicadoras para identificar las especies. Los bofedales difieren en riqueza ( $F = 5.1, p < 0.001$ ) y diversidad de microhábitats ( $F = 23.4, p < 0.001$ ), pero no hallamos diferencias estacionales ( $p > 0.05$ ). El modelo seleccionado indica que el área y la diversidad de microhábitats explican 54% de la varianza en la riqueza, sin embargo, encontramos una tendencia inversa entre la diversidad de microhábitats y el área, la cual sugiere un papel relativamente más importante del área en la riqueza de especies. Nuestros resultados son respaldados por los datos de composición, ya que los

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microhábitats no sólo fueron diferentes en comparaciones pareadas ( $T = -94.14$ ,  $A = 0.601$ ,  $p < 0.001$ ) y estuvieron representados significativamente por especies indicadoras ( $p < 0.05$ ), sino que contribuyeron diferencialmente con la riqueza. Así, pocos microhábitats ricos en especies contribuirían más a la riqueza que varios microhábitats pobres en especies lo cual no concuerda con las predicciones de la hipótesis de heterogeneidad del hábitat. Determinar la influencia que el área y la heterogeneidad tienen en la riqueza de especies es importante para establecer prioridades de conservación que garanticen la integridad de los bofedales ante el inminente cambio climático.

**Keywords** Diversity · Microhabitats · Cover · Cushion bogs · Indicator species

**Palabras clave** Diversidad · Microhábitats · Cobertura · Bofedales · Especies indicadoras

## Introduction

The number and identity of species present at a site are influenced by the interaction of factors operating at different spatial and temporal scales (Ricklefs and Schluter 1993; Huston 1999). At a regional scale, species richness and community composition are likely influenced by speciation rates, biogeographic factors, and/or climate, while at a local scale, physical and ecological factors such as habitat structure, micro-environmental conditions, and interactions (e.g., predation, competition, mutualism, etc.) play a prominent role in determining species richness and composition (Vuilleumier 1970; Terborgh 1977; Angermeier and Winston 1998; Rahbek 2005).

The scale dependence of species richness is represented by the species–area relationship (SAR; Arrhenius 1921; Gleason 1925). SAR has been studied in the context of two not mutually exclusive hypotheses: the ‘*area per se*’ (Preston 1960, 1962), derived from the island biogeography equilibrium theory (MacArthur and Wilson 1967) and the ‘*habitat heterogeneity hypothesis*’ (Williams 1964), developed within the context of niche theory (Hutchinson 1957). In the former, as area and isolation increases, the probability of encountering more species decreases through their effects on colonization and extinction rates (even in a uniform environment); in the latter, as area increases, more habitat types are encountered, allowing more species to coexist due to niche partitioning (MacArthur and MacArthur 1961; Holt 2009; Stein et al. 2014). As such, the relationship between area and habitat heterogeneity is usually positive, however, it could also be negative when greater microhabitat diversity reduces the amount of suitable area available for each species and, after some threshold, richness decreases (*‘area-heterogeneity trade off’*,

Kadmon and Allouche 2007, Allouche et al. 2012, Bar-Massada and Wood 2014).

The notion that increasing environmental heterogeneity promotes a larger number of species with different ecological requirements has been widely acknowledged (e.g., Ricklefs and Lovette 1999; Allouche et al. 2012; Bar-Massada and Wood 2014; Stein et al. 2014), but its relative contribution has proven difficult to discern because area and habitat heterogeneity are strongly correlated at large spatial scales (Simberloff 1976; Kohn and Walsh 1994; Ricklefs and Lovette 1999).

The effects of area and habitat heterogeneity on local species richness and composition are particularly relevant for models that predict shifts in the geographic range and distribution of species. Shifts in distribution of species may affect overall richness and ecosystem function particularly under different scenarios of climate change, (e.g., Thomas and Lennon 1999; Graham et al. 2011; Young 2012; Herzog et al. 2012). Predicting patterns of species richness in models projections linking conservation efforts that maximize biodiversity (Margules and Pressey 2000; Brooks et al. 2006) and ecosystem function (Tilman et al. 1997; Chapin et al. 1998; Cadotte et al. 2011; but see Schwartz et al. 2000) have become a fundamental aspect in conservation. However, accuracy of models requires basic knowledge of spatial patterns of species richness and composition, which for most tropical ecosystems is far from being accomplished, even for some well-known groups of organisms such as birds (Herzog and Kattan 2011).

In the high Andes, the complex topography and elevational gradients provide a variety of local conditions likely influencing species components (Benham and Witt 2016). Cushion bogs (locally known as *bofedales*) are a high Andean wetland system characterized by water flowing pools and rivulets that depend on glacier melt and precipitation for constant water flow, surrounded by cushion vegetation (Weberbauer 1936; Squeo et al. 2006; Ruthsatz 2012). Cushions are thick layers of vegetation growing over slowly accumulating organic matter in various stages of decomposition due to low temperature and oxygen available at high elevations. Through time, cushions have the capacity for insulation and water retention providing important ecosystem services such as carbon and nitrogen sinks, sources of methane, regulators of permafrost and hydrological cycles (Keddy 2010; Vuille 2013). Bofedales are one of the most productive life-support systems, characterized by a distinct and diverse flora and fauna, but at the same time they constitute a fragile and threatened ecosystem markedly sensitive to human disturbances. Major threats to bofedales include unsustainable land use (e.g., overgrazing, peat mining; Foote and Krogman 2006), urban expansion (Mitsch and Gosselink 1993), and climate change (Bury et al. 2013; IPCC 2014). Since studies in the bofedal ecosystem are poorly documented (but see Telleria et al. 2006; Gibbons 2012) our main goal was to document bird species

richness and composition in relation to bofedal area and habitat heterogeneity (microhabitat diversity). We basically ask the following two questions: (1) How do the number of species and the composition of bird assemblages vary in bofedales that differ in size, distance from other bofedales (isolation), and microhabitat diversity? and (2) what variables (i.e., area, microhabitat diversity) best predict avian richness in bofedales?

## Methods

We studied 24 bofedales located in the Huancavelica and Ayacucho Departments (Fig. 1, Table 1) in the Central Andean Puna region (Olson et al. 2001). In the high Andes, climatic conditions vary depending on elevation, sun exposure and other local factors, but in general, daily temperatures are extremely oligothermic, while yearly variation in mean temperature is small (Vuille et al. 2008; Vuille 2013). On the other hand, the amount of precipitation defines a wet season, with 60% to 95% of total annual rainfall occurring from December to March, and a dry season with 0% to 40% rainfall from April to November. Data were collected during four 12-day visits in November–December 2010 and 2012; and April–May 2013 and 2014. Criteria for bofedal selection included sites above 4300 m (to avoid species from montane or non-Puna ecosystems), isolated by mountain ridges (to reduce connectivity), groundwater or glacial origin (i.e., microtrophic), and water presence throughout the year (i.e., hydromorphic). We avoided potential regional effects by restricting the study to the central part of a precipitation gradient that diminishes from east to west and corresponds to changes in bofedales plant species composition (Valencia et al. 2013). All bofedales in the region are under the influence of linear infrastructure projects (two underground gas pipelines run from east to west), numerous access roads, grazing pressure from sheep, alpaca, and llamas since pastoralism has dominated the region over millennia (Browman 1974; Villarroel et al. 2014).

On each visit, we conducted bird censuses in two non-consecutive days, from 7:45 AM to approximately 10:45 AM, along a line transect that surrounded each bofedal. Two trained surveyors walked in opposite directions at constant speed (~ 0.40 km/min, Servat et al. 2013). We identified and counted all individuals in the water-saturated and humid and sub-humid zones of the bofedal (sensu Valencia et al. 2013) and recorded the microhabitat(s) used by foraging birds (e.g., cushion vegetation, pools/shores, bare areas, etc.) Birds flying (or inactive) were not assigned to any microhabitat. We reduced variation in detectability due to (1) weather conditions, by not sampling during inclement weather, (2) temporal

differences, by visiting bofedales in different days within and among visits, and (3) observer differences, by having the same team of observers trained together to standardize census ability, and by randomly assigning observers to sites.

We estimated area by georeferencing 20–30 points (Garmin GPSmap 60CSx) along the perimeter of each bofedal using Google Earth Pro ([www.google.com/earth](http://www.google.com/earth)). We averaged the elevation data taken at georeferenced points to obtain a single measure for each bofedal. Isolation was estimated by measuring the linear distance from the border of each bofedal to the border of its nearest neighbor.

Bofedales have aquatic and terrestrial microhabitats with micro-invertebrates and plants suitable as food resources for foraging birds that included: (1) Cushion vegetation, composed mainly of *Distichia muscoides* and *Plantago tubulosa*; (2) Water pools, with plant species such as *Potamogeton strictus* and a few species of *Ranunculus* (Valencia et al. 2013), offering large concentrations of plankton for aquatic birds. Water pools are surrounded by shores that vary in substrate type from sieve mud, silt, or gravel, used by shorebirds and waders; (3) Bare expanses of soil, some of which may be dry vernal pools that are searched by ground-feeding birds; (4) Rocky outcrops from landslides that absorb and radiate heat (Weberbauer 1936) providing microsites for lithophilous plants, ferns, and tall-stalk herbs that attract pollinators, frugivorous and seed eating birds; (5) Tussock-forming grasses or tall perennial grasses growing in bunches or clumps (e.g., species of *Festuca*, *Poa*, *Calamagrostis*, etc.); (6) Short lawn-like vegetation, composed of graminoids (grasses and grass-like forms) and forbs (non-graminoid and broad leaved herbaceous plants) that may include degraded vegetation; and (7) Shrubs, scattered plants up to 30 cm height (mostly species of Asteraceae and Ericaceae) found in marginal areas of the bofedal.

We measured habitat heterogeneity as microhabitat diversity, a measure that combines two components: the number of microhabitat types and its cover (%) at each bofedal. To estimate components we took panoramic photographs from the same locations (two high elevation sites) in each bofedal per season, using a Cannon Powershot SX30IS camera with regular lens (i.e., 360° horizontal and 45° vertical fields of view). We drew contour lines around water bodies, tussock, bare areas, rocky outcrops, and cushion vegetation in the photographs; counted all microhabitat types and estimated the mean cover (%) of each microhabitat per bofedal per season. Data was used to calculate microhabitat diversity using Shannon-Wiener diversity index ( $H'$ ) (Shannon 1948 in Morris et al. 2014).

## Data Analysis

We estimated sampling effort, defined as the cumulative number of species observed at each bofedal by season, to assess the completeness and representativeness of bird surveys. We used the *species accumulation curves* function (*specaccum*) in the ‘vegan’ Package (Oksanen et al. 2012) within the CRAN platform (R Development Core Team 2006). We obtained the mean and variance of bird species richness per bofedal per season after randomization of the original matrix (999 times), with the number of species randomly taken for each sample (Colwell et al. 2012).

To test for differences in species richness, area, isolation, microhabitat diversity and cover, between seasons (2), and bofedales (24) we used General Linear Models (GLM). We explored significant results with Tukey’s Honest Significant Difference (HSD) post hoc tests (Sokal and Rohlf 2012). All data used in the analyses were checked for normality (Wilk-Shapiro test) and homogeneity of variances (Levene’s test). Data were transformed to  $\log_n$  (bird richness and microhabitat diversity) or arcsine (microhabitat cover) to meet normality assumptions.

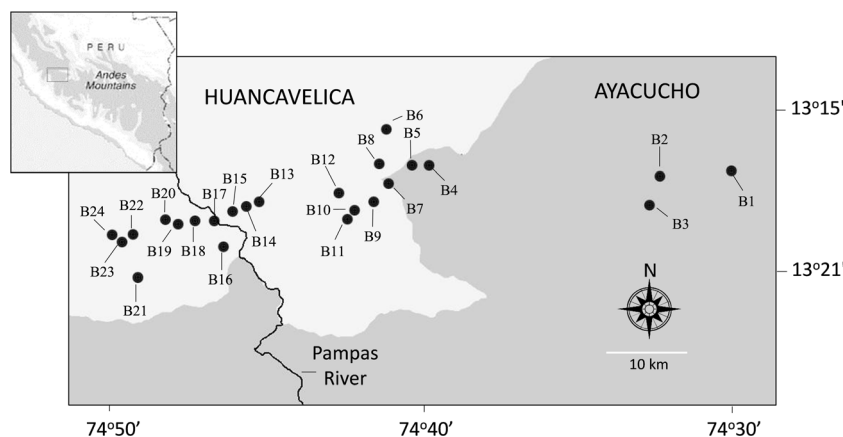
Descriptions of the species-area relationship have been mathematically represented by power and logarithmic functions (Connor and McCoy 1979, Rosenzweig 1995, Lennon et al. 2001). However, since power functions imply a constant increase in species richness with increasing area (whereas logarithmic functions imply a constant absolute increase) (White et al. 2006), we considered a linearized power function ( $\text{LogS} = C + \text{LogA}$ ) for multimodel inference. To assess collinearity among predictor variables (i.e., bofedal area, isolation, and microhabitat diversity), we estimate the Variance Inflation Factor (VIF), which measures how much the variance of an estimated regression coefficient is increased due to collinearity. We used as threshold for variable removal VIF values  $>3$  (Zuur et al. 2010) in the Companion to Applied Regression Package (CAR, Fox and Weisberg 2011) within the CRAN platform

(R Development Core Team 2006). To estimate the relationship between bird species richness and predictor variables, we used Ordinary Least Square (OLS) models. All candidate models were ranked using Akaike Information Criterion corrected for small sample sizes (AICc, Bolker et al. 2009) in the Multimodel Inference Package (MuMIn, Bartoń 2016) within the CRAN platform (R Development Core Team 2006). We chose the *dredge* function to explore the differences between the AICc values of the best model (i.e., model with the lowest AICc) and other candidate models ( $\Delta\text{AICc}$ ). Models with AICc differences  $<2$  were considered supported by the data (Bolker et al. 2009; Richards 2015). All analyses were run in R Software (R Development Core Team 2006), except where indicated.

Species composition was explored using Bray Curtis ordination (PC ORD Version 6, McCunne and Mefford 2011) to obtain a spatial configuration of species in different groups (microhabitats) within and across bofedales. To do so, we built a matrix of bird species (rows) by microhabitats (columns) for each bofedal, with the relative use of each microhabitat filling the cells of the matrix. In all matrices, we relativized data by row (species), and selected Sørensen (Bray Curtis) distance (i.e., the average of pairwise dissimilarities in microhabitat use between bofedales), and used the original Bray Curtis method for end-point selection.

To determine whether bird species composition in microhabitats within and among bofedales differed significantly, we used the species groups obtained from Bray Curtis ordination of each bofedal and perform *Multiple Response Permutation Procedure* (MRPP). MRPP uses classified bird groups to test the null hypothesis of no differences in use within and among microhabitats across bofedales providing measures of the degree of separation among microhabitats (T), within-microhabitat agreement (A), and *P* values (after 900 runs of the original matrix, using a random starting configuration) (McCunne and Grace 2002). To identify the species that best

**Fig. 1** Geographical location of bofedales (B1–B24) in Ayacucho (dark shaded) and Huancavelica (light shaded) in the Central Andes of Peru. All bofedales were above 4300 m, surrounded by mountain ridges, and with water present throughout the year (See Table 1 for specifics)



**Table 1** Geographic coordinates, bofedal area (ha), elevation (m ASL), isolation, mean microhabitat diversity ( $H'$  = Shannon Wiener index) by season (WS = wet season November – December 2010 and 2012, DS = dry season April – May 2013 and 2014) and species richness (LN) by season at each bofedal included in the study

Bofedal	Latitude (S)	Longitude (W)	Elevation (m asl)	Area (ha)	Isolation (m)	Microhabitat diversity ( $H'$ )		Species Richness	
						WS	DS	WS	DS
B1	13°16'53"	74°30'34"	4346	10.7	1604.0	1.04	1.16	0.110	0.030
B2	13°17'09"	74°32'45"	4400	8.0	539.0	1.15	1.29	0.560	0.490
B3	13°17'21"	74°33'10"	4500	7.3	710.2	0.80	1.04	0.490	0.400
B4	13°16'48"	74°39'50"	4546	10.0	547.0	1.03	1.05	0.590	0.640
B5	13°16'47"	74°40'17"	4636	10.0	1186.0	1.43	1.39	0.260	0.100
B7	13°17'19"	74°41'02"	4965	5.1	866.0	1.48	1.41	1.260	1.260
B6	13°15'42"	74°41'05"	4620	4.9	665.5	1.53	1.43	1.110	1.040
B8	13°16'45"	74°41'18"	4716	3.7	813.0	1.20	1.19	1.160	1.160
B9	13°17'50"	74°41'28"	4620	2.1	722.0	1.35	1.37	1.550	1.550
B10	13°18'05"	74°42'01"	4670	5.1	630.0	1.53	1.52	0.670	1.320
B11	13°18'21"	74°42'14"	4541	5.5	671.0	1.31	1.40	1.130	0.930
B12	13°17'33"	74°42'30"	4825	3.6	417.9	1.23	1.31	0.500	0.660
B13	13°17'48"	74°44'54"	4877	5.9	487.1	1.21	1.26	0.700	0.920
B14	13°18'02"	74°44'55"	4706	5.6	557.0	1.46	1.48	1.110	0.680
B15	13°18'06"	74°45'43"	4752	7.5	707.0	1.42	1.45	0.280	0.280
B17	13°19'05"	74°45'58"	4661	5.9	411.9	1.20	1.32	0.620	0.530
B16	13°18'27"	74°46'21"	4849	3.9	763.0	1.53	1.56	1.260	1.180
B18	13°18'31"	74°46'52"	4816	10.0	200.0	1.19	1.24	0.640	1.550
B19	13°18'31"	74°47'26"	4756	4.3	531.6	1.57	1.60	1.550	0.620
B20	13°18'27"	74°47'49"	4661	5.44	868.5	1.49	1.61	1.080	1.010
B21	13°18'47"	74°48'43"	4770	8.06	756.9	1.42	1.47	0.960	0.910
B22	13°20'04"	74°48'32"	4638	3.16	645.0	1.45	1.46	1.560	1.560
B23	13°18'02"	74°48'26"	4717	7.03	875.7	1.59	1.57	0.530	0.610
B24	13°19'30"	74°50'30"	4772	3.06	676.7	1.44	1.44	0.670	1.550

characterize each microhabitat, we used *Indicator Species Analysis* (ISA) (PC ORD Version 6, McCunne and Mefford 2011). ISA provides an indicator value (IV) for each species based on its frequency of occurrence in each microhabitat across bofedales (IV reaches 100% when the species uses the same microhabitat in all bofedales where it is present), and a  $P$  value after comparing the observed with the predicted association (after a randomization procedure with 900 runs of the original matrix) between species and microhabitats to determine its statistical significance.

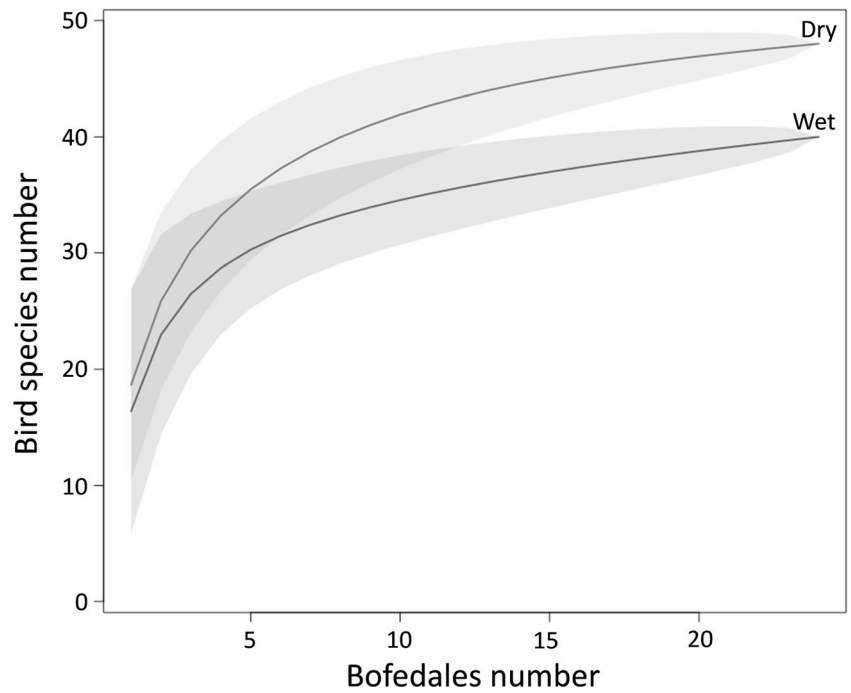
## Results

We found 52 bird species in 24 bofedales that varied in area from 2.1 to 10.7 ha (Mean =  $8.0 \pm 2.43$ ) (Table 1, Appendix 1). Most birds in bofedales (86%) had narrow to medium size geographic distribution ( $< 500,000$  to  $1' 500,000$  km<sup>2</sup>) and 78% had an elevational range of distribution of 1000 to 5500 m. (Appendix 1).

The cumulative number of bird species in bofedales tends to reach an asymptote in both the dry and wet seasons (Fig. 2). Seasonal variation in species richness was not significant ( $p > 0.05$ ), despite more species being detected during the dry season (49) than during the wet season (39) (Fig. 2, Appendix 1). Spatial differences emerged when comparing richness across bofedales (GLM  $F_{\text{bofedales}} = 5.1$ ,  $df = 23, 3$ ;  $p < 0.001$ ) (Fig. 3), with significant variation due to the “species-rich” bofedales B9, B19 and B22; and the “species-poor” B1, B5 and B18 (Fig. 3), as indicated by post-hoc Tukey test.

Microhabitat diversity differed across bofedales ( $F_{H'} = 23.4$ ,  $df = 23$ ,  $p < 0.001$ ,  $n = 48$ ), independent of season ( $p > 0.05$ ). Our results revealed that microhabitat diversity was significant and inversely related to bofedal area, although the relation had a low predictive value ( $R^2 = 0.13$ ,  $p = 0.01$ ,  $n = 48$ ). Comparisons of microhabitat cover (%) across bofedales differed significantly in shrubs ( $F = 185.5$ ,  $df = 23$ ,  $p < 0.001$ ), tussock ( $F = 381.8$ ,  $df = 23$ ,  $p < 0.001$ ), rocky outcrops ( $F = 35.3$ ,  $df = 23$ ,  $p < 0.001$ ), and bare areas ( $F = 16.8$ ,  $df = 23$ ,  $p < 0.001$ ), but not for cushion vegetation and water pools/

**Fig. 2** Cumulative mean number of bird species in 24 bofedales surveyed during the wet and dry seasons. The lines indicate the mean accumulation increase of detected species (after 999 randomizations of the original matrix). The *shadowed area* around each solid line indicates 95% confidence intervals

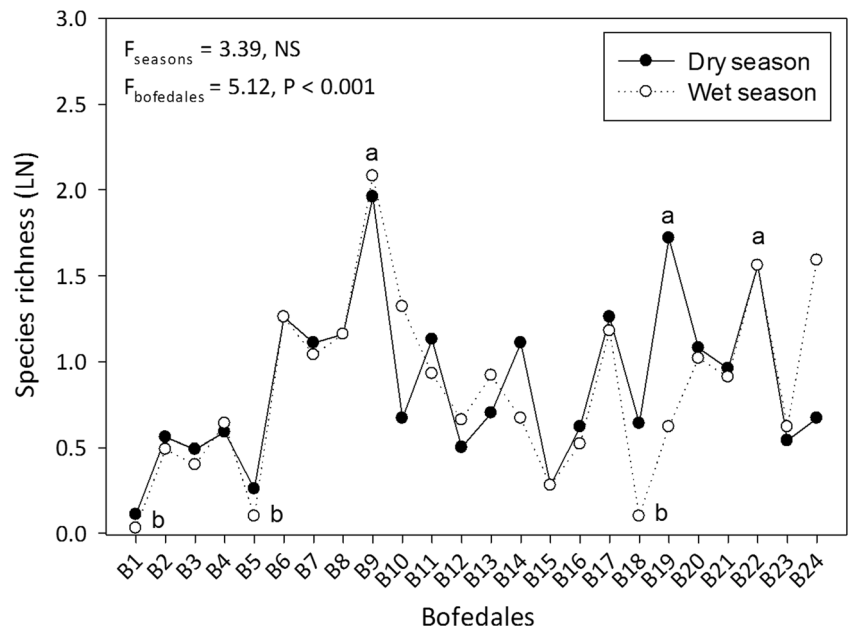


shores ( $p > 0.05$ ). Seasonal differences only emerged for cushion vegetation ( $F = 13.9, n = 47, p < 0.001$ ) and water pools ( $F = 28.5, n = 47, p < 0.001$ ), as during the wet season water increases at the expense of cushion vegetation (conversely, in the dry season cushion vegetation and shores increase).

Predictor variables (bofedal area, isolation, elevation and microhabitat diversity) were not collinear ( $VIF < 2$ ), so we included all in the OLS model to explain bird

species richness in bofedales. Model selection (from the 16 OLS models by season obtained using 4 variables, Appendix 2), reveals that the combined effect of area and microhabitat diversity explained 54% of the variation in species richness in the dry season, while in the wet season, 38% of the variation in species richness was predicted by bofedal area (Table 2). The effect of isolation when combined with area only explained 43% (Table 2).

**Fig. 3** Species richness/ha logarithmically transformed (LN) in 24 bofedales (B1-B24) surveyed in the dry (discontinuous line and white circles) and wet season (continuous line and black circles). General Linear Model (GLM) ANOVA (F) values for comparisons among seasons and bofedales are provided. Lowercase fonts (*a, b*) indicate bofedales that significantly contribute with GLM differences



**Table 2** Highest rank models and predictors of species richness (SpRich) in the dry (DS) and wet seasons (WS), based on the Akaike information criterion adjusted for small sample size (AICc). Differences in AICc between model *i* and the best supported model ( $\Delta i$  AICc), and

model weight (*w<sub>i</sub>* AICc) are provided for each model. Predictor variables are: Area (bofedal size), LN MD<sub>DS</sub> (Log transformed microhabitat diversity–Dry season), ISOL (Isolation, distance to nearest bofedal). Models with  $\Delta i > 2$  are not included in the table

Response variable	Intercept	Model and predictor variables	R <sup>2</sup> <sub>Adj</sub>	AICc	ΔAICc	AICcw
SpRich-DS	0.73	−1.45*Area + 0.87*LN MD <sub>DS</sub>	0.54	14.9	0.00	1.00
SpRich-WS	2.08	−1.6*Area	0.38	24.3	0.10	0.49
SpRich-WS	2.28	−1.45*Area−0.0004 ISOL	0.43	24.2	0.00	0.52

Nearly 45% of all bird species associated with bofedales were observed in 1 or 2 foraging microhabitats while 55% were found in more than three (Appendix 1). Ordination of species within and among microhabitats in the 24 bofedales studied followed five general patterns that differed in the amount of variation explained by the two first axes (Table 3). The first pattern shows bird species separated in groups associated to tussock

vegetation from those in bare areas along axis *x* (B5, B6, B7, B15, B20, B22; Table 2). Subsequent patterns show birds species associated with pools/shores separated from species in bare areas along axis *x* (B8, B12, B13; Table 2); shrubs along axis *y* (B1, B4; Table 3); rocky outcrops along axis *y* (B3, B9, B11, B14, B17, B19, B21, B24; Table 2); and in cushion vegetation along axis *y* (B2, B10, B16, B18, B23; Table 3).

**Table 3** Percent of variation explained by primary axes of Bray-Curtis ordination of bird species by microhabitat in 24 bofedales

Bofedal	Variation explained (%)			Endpoint microhabitats <sup>a</sup>
	Axis <i>X</i>	Axis <i>Y</i>	Total	
B5	<b>38.0</b>	27.8	65.8	Tussock – Bare areas
B6	<b>41.9</b>	23.2	65.1	Tussock – Bare areas
B7	<b>40.5</b>	25.2	65.7	Tussock – Bare areas
B15	<b>54.4</b>	17.1	71.5	Tussock – Bare areas
B22	<b>42.0</b>	32.3	74.3	Tussock – Bare areas
B8	<b>40.9</b>	23.2	64.1	Pools and shores – Bare areas
B12	<b>42.7</b>	36.4	79.1	Pools and shores – Bare areas
B13	<b>58.7</b>	23.5	82.2	Pools and shores – Bare areas
B20	<b>50.5</b>	23.2	73.7	Pools and shores – Bare areas
B1	28.4	<b>36.6</b>	65.0	Shrubs – Pools and shores
B4	21.1	<b>51.7</b>	72.8	Shrubs – Pools and shores
B3	23.2	<b>59.4</b>	82.6	Pools and shores – Rocky outcrops
B9	23.1	<b>50.9</b>	74.0	Pools and shores – Rocky outcrops
B11	24.0	<b>58.8</b>	82.8	Pools and shores – Rocky outcrops
B14	19.6	<b>47.8</b>	67.4	Pools and shores – Rocky outcrops
B17	18.9	<b>51.6</b>	70.5	Pools and shores – Rocky outcrops
B19	20.0	<b>50.0</b>	70.0	Pools and shores – Rocky outcrops
B21	19.1	<b>49.8</b>	68.9	Pools and shores – Rocky outcrops
B24	23.2	<b>41.8</b>	65.0	Pools and shores – Rocky outcrops
B2	18.7	<b>39.7</b>	58.4	Cushion vegetation – Pools and shores
B10	24.7	<b>37.1</b>	61.7	Cushion vegetation – Pools and shores
B16	18.2	<b>39.5</b>	57.7	Cushion vegetation – Pools and shores
B18	23.2	<b>38.0</b>	61.2	Cushion vegetation – Pools and shores
B23	23.1	<b>35.1</b>	58.2	Cushion vegetation – Pools and shores

In bold is the axis that contributes most to ordination. We used Sørensen’s dissimilarity to measure the distance gradient among microhabitats; and the original Bray Curtis endpoint selection method to identify the microhabitats that polarize the ordination

<sup>a</sup> Microhabitats that polarize the ordination for the axis (in bold) that contributes most to the ordination

**Table 4** Pairwise comparisons in species composition between and within microhabitats obtained with Multiple Response Permutation Procedures (MRPP). T = Degree of separation between microhabitats (higher negative values indicate greater separation); A = Level of agreement within microhabitats (higher values indicate more similarity). Pair-wise comparisons were significantly different at  $\alpha = 0.05$

Microhabitat pairwise comparisons		T	A	P
Cushion vs.	Pools/shores	-30.21	0.40	0.000
	Bare	-31.83	0.52	0.000
	Tussock	-31.64	0.51	0.000
	Rocks	-27.81	0.31	0.000
	Short vegetation	-29.21	0.35	0.000
	Shrubs	-27.20	0.48	0.000
Pools/shores vs.	Bare	-31.98	0.54	0.000
	Tussock	-31.67	0.50	0.000
	Rocks	-31.37	0.47	0.000
	Short vegetation	-27.08	0.27	0.000
	Shrubs	-28.07	0.55	0.000
Bare vs.	Tussock	-32.16	0.58	0.000
	Rocks	-30.98	-0.44	0.000
	Short vegetation	-32.05	0.52	0.000
	Shrub	-27.59	0.53	0.000
Tussock vs.	Rocks	-31.33	0.48	0.000
	Short vegetation	-29.79	0.36	0.000
	Shrubs	-28.20	0.56	0.000
Rocks vs.	Short vegetation	-30.67	0.41	0.000
	Shrubs	-27.00	0.44	0.000
Short vegetation vs.	Shrubs	-28.08	0.54	0.000

Classification of bird species by microhabitat was supported by MRPP analysis, as significant differences in pairwise comparisons between microhabitats and random-corrected within-microhabitat agreement were found ( $T = -94.14$ ,  $A = 0.601$ ,  $p < 0.000$ ). Relative comparisons of significant A values shows that bird assemblages in cushion vegetation shared less species with rocky outcrops ( $A = 0.31$ , Table 4). In the same manner, assemblages associated with pools/shores shared less species with short vegetation ( $A = 0.27$ , Table 4), suggesting less homogeneity in species composition.

Microhabitats were also characterized by the presence of indicator species. ISA reveals that 37 bird species were significant indicators of microhabitats ( $p < 0.05$ , Table 5) from which 17 were present in the same microhabitat across bofedales more than 30% of the time (Table 5). As such, *Diuca speculifera* and *Lessonia oreas* were significant indicator species in cushion vegetation,

while *Chloephaga melanoptera*, *Phegornis mitchellii* and *Plegadis ridgwayii* were significant indicators of pools and shores of bofedales (Table 5). Other species such as *Upucerthia validirostris* and *Oreotrochilus melanogaster* in rocky outcrops, *Gallinago andina* in tussock and *Colaptes rupicola* and *G. cunicularia* in short vegetation, had high values as indicator species (the only exception being bare areas, with no indicators) (Table 5).

## Discussion

Avian richness in bofedales varied with some of them being consistently rich or consistently poor independent of season. Bofedal area and microhabitat diversity were the best predictors of species richness, although elucidating the direct and indirect effects of area on species richness is still debated (Tews et al. 2004 and references therein). Species richness could be influenced by area directly, as large areas may have higher colonization rates and support larger populations providing less vulnerability to stochastic extinction (area per se hypothesis); or indirectly, through its effect on habitat heterogeneity, often presumed to increase in direct relation to area (habitat heterogeneity hypothesis, e.g., Kohn and Walsh 1994, Allouche et al. 2012; Bar-Massada and Wood 2014, Stein et al. 2014). Our results however, show a tendency for a decrease in microhabitat diversity as area increases, suggesting a relatively more important role of area per se on species richness in the bofedal system

The overriding effect of area on richness of bofedales is further supported by our results on bird species composition. Our findings revealed that sets of species consistently (and significantly) characterized cushion vegetation, pools and shores, rocky outcrops, shrubs, tussock and short vegetation (the exception being bare areas, used opportunistically by different species) (Table 2). Nonetheless, microhabitats contributed differently to species richness as some had fewer associated species (e.g., tussock, scrubs) than others (e.g., cushion vegetation, pools/shores) (Table 3), suggesting that few species-rich microhabitats (i.e., “homogenous”) may contribute more to increase bird richness (e.g., key habitat types, Davidar 2001), than many species-poor microhabitats (i.e., “heterogeneous”), a result not supported by the habitat heterogeneity hypothesis. Moreover, bird species associated with bofedales may differ in their foraging requirements, and thus species may respond to the presence of suitable foraging habitats. For instance, some species that



**Table 5** Indicator species analysis for each microhabitat

Microhabitat	Indicator species <sup>a</sup>	IV	Montecarlo simulations mean ± SD
Cushion vegetation	<i>Geositta saxicolina</i>	30.6	10.5 ± 2.11
	<i>Muscisaxicola albifrons</i>	33.3	7.3 ± 2.23
	<i>Muscisaxicola flavinucha</i>	33.3	11.0 ± 2.03
	<i>Asthenes humilis</i>	47.9	8.7 ± 2.15
	<b><i>Diuca speculifera</i></b>	<b>83.3</b>	5.8 ± 2.18
	<b><i>Lessonia oreas</i></b>	<b>91.7</b>	6.1 ± 2.19
Pools/shores	<i>Anas flavirostris</i>	50.0	4.8 ± 2.26
	<i>Lophonetta specularioides</i>	50.0	4.8 ± 2.2
	<b><i>Chloephaga melanoptera</i></b>	<b>91.7</b>	6.1 ± 2.19
	<b><i>Phegornis mitchellii</i></b>	<b>100</b>	6.3 ± 2.18
Scattered shrubs	<b><i>Plegadis ridgwayi</i></b>	<b>100</b>	6.3 ± 2.18
	<b><i>Upucerthia validirostris</i></b>	<b>70.8</b>	5.4 ± 2.2
Rocky outcrops	<i>Muscisaxicola juninensis</i>	47.9	8.6 ± 2.17
	<b><i>Oreotrochilus melanogaster</i></b>	<b>50.0</b>	8.8 ± 2.15
Short vegetation	<i>Geositta cunicularia</i>	31.9	10.7 ± 2.09
	<b><i>Colaptes rupicola</i></b>	<b>33.3</b>	11.0 ± 2.07
Tussock	<b><i>Gallinago andina</i></b>	<b>34.8</b>	7.1 ± 2.23

Indicator values (IV) range from 0 (no indication) to 100 (perfect indication). The statistical significance of the maximum IV recorded for each species was generated by a Monte Carlo test. Species with significant IV values ( $p < 0.001$ ) are shown in bold

<sup>a</sup> Only species with IV > 30 are shown

are restricted to forage in one or two microhabitats (e.g., *Phegornis mitchellii*, *Diuca speculifera*) could favor homogeneous bofedales (with less microhabitats) as their preferred microhabitat may be better represented, while other species that use different foraging microhabitats (e.g., *Muscisaxicola cinereus*) may prefer heterogeneous bofedales as these provide more suitable microhabitats to forage (e.g., Riffell et al. 2001).

Our results do not provide support for isolation effects, as the distances between bofedales may not be large enough to drive a response on species richness (Watling and Donnelly 2006). However, our measures of isolation, the linear distance to nearest bofedal, may not necessarily reflect isolation, as bofedales in this study were surrounded by mountains and a complex topography that may effectively isolate them (Table 1). High dispersal capabilities of birds could also override area effects (Ricklefs and Lovette 1999) although data on dispersal is lacking, a possible indication of its role could be determined by the lack of endemics associated with bofedales in the region of study. Further studies should focus on species with different dispersal capabilities and levels of restriction to the system including bofedales from other Andean regions.

Field studies provide key information to identify gaps in existing knowledge for accurate projections that maximize biodiversity and avoid misleading conservation efforts. Disentangling the role of area and habitat heterogeneity in promoting species richness is important to understand which conditions help explain the coexistence of species in threatened ecosystems such as bofedales. In the face of a rapidly changing climate, the ability to make informed decisions is extremely urgent to maintain the integrity of the bofedal system.

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## Appendix 1

**Table 6** Altitudinal (m) and geographic (km<sup>2</sup>) ranges of distribution, and foraging microhabitats used by bird species associated with bofedales. Microhabitats: Ba = bare areas, Cu = cushion vegetation, Sv = short vegetation, Sr = scattered shrubs, Ro = rock outcrops, Tu = tussock, and Ws = water pools and shores. DS and WS = species reported in the dry or wet season exclusively. The list excludes species observed during flight (*Geranoaetus melanoleucus*, *G. polyosoma*, and *Haplochelidon andecola*)

Bird species <sup>a</sup>	Elevation range (m ASL) <sup>b</sup>	Geographic range (km <sup>2</sup> ) <sup>c</sup>	Foraging microhabitats
<i>Agriornis albicauda</i> <sup>DS</sup>	3360–4300	582,000	Sr, Ba, Sv
<i>Agriornis montanus</i>	2000–4500	2,170,000	Sr, Ba, Sv
<i>Anas flavirostris</i>	0–4400	3,940,000	Ws
<i>Anthus bogotensis</i>	2100–4500	336,000	Cu, Sv
<i>Asthenes humilis</i>	2750–4800	146,000	Ba, Cu, Sv
<i>Asthenes modesta</i>	0–4200	2,020,000	Ws, Ro, Sv
<i>Attagis gayi</i>	3390–4800	1,030,000	Cu, Ro, Sr, Sv
<i>Calidris bairdi</i>	2500–4700	2,810,000	Ws
<i>Chalcostigma olivaceum</i> <sup>DS</sup>	3600–4600	92,100	Cu, Ba, Sv
<i>Chloephaga melanoptera</i>	3300–4700	1,160,000	Ws
<i>Chroicocephalus serranus</i>	4000–5300	980,000	Ba, Sv
<i>Cinclodes albiventris</i>	0–5000	705,000	Ws, Cu, Sv
<i>Cinclodes atacamensis</i>	2200–4500	871,000	Ws, Cu, Ro
<i>Colaptes rupicola</i>	2000–5000	894,000	Cu, Ba, Sv
<i>Diuca specularifera</i>	4245–5500	208,000	Cu
<i>Falco femoralis</i>	0–4600	11,700,000	Sr, Ba, Sv, Ro
<i>Gallinago andina</i>	3100–4400	777,000	Cu, Ws, Tu
<i>Geositta cunicularia</i>	0–5000	2,910,000	Cu, Ba, Sv
<i>Geositta saxicolina</i>	3700–4900	44,800	Ro, Sr, Sv
<i>Geositta tenuirostris</i>	2500–4600	622,000	Ba, Sv, Sv
<i>Lessonia oreas</i>	3100–4300	832,000	Cu, Sv
<i>Lophonetta specularioides</i>	0–4700	1,860,000	Ws
<i>Metriopelia aymara</i> <sup>DS</sup>	2800–5000	832,000	Sr, Ba, Sv
<i>Metriopelia melanoptera</i> <sup>DS</sup>	2000–4400	1,470,000	Sr, Ba, Sv
<i>Muscisaxicola albifrons</i>	4000–5600	286,000	Cu, Sv
<i>Muscisaxicola cinereus</i> <sup>ds</sup>	2500–5000	650,000	Cu, Ro, Sr, Sv, Ba
<i>Muscisaxicola flavinucha</i> <sup>DS</sup>	500–4500	693,000	Cu, Ro, Sr
<i>Muscisaxicola frontalis</i> <sup>DS</sup>	2500–4300	223,000	Ro, Ba, Sv
<i>Muscisaxicola griseus</i>	2700–4550	313,000	Ro, Ba, Sv
<i>Muscisaxicola juninensis</i>	3200–4600	514,000	Ro, Sv
<i>Muscisaxicola rufivertex</i>	3000–4500	1,040,000	Ro, Sr, Ba, Sv
<i>Nycticorax nycticorax</i> <sup>WS</sup>	0–4800	41,200,000	Ws
<i>Ochthoeca oenanthoides</i>	2000–4500	624,000	Cu, Sr, Sv
<i>Oreotrochilus estella</i> <sup>DS</sup>	2400–5000	596,000	Ro, Sv
<i>Oreotrochilus melanogaster</i> <sup>DS</sup>	3500–4200	55,300	Ro, Sv
<i>Phalacrocorax megalopterus</i>	2000–5000	1,010,000	Tu, Ba, Sv
<i>Phegornis mitchellii</i>	3000–5000	762,000	Ws
<i>Phrygilus plebejus</i>	2500–4500	951,000	Cu, Sr, Ba, Sv
<i>Phrygilus punensis</i>	2000–4300	310,000	Sr, Ba, Sv
<i>Phrygilus unicolor</i>	2700–4500	1,360,000	Cu, Sr
<i>Plegadis ridgwayi</i>	3080–4800	457,000	Ws
<i>Sicalis uropygialis</i>	3500–4800	706,000	Ba, Sv
<i>Sporagra atrata</i>	1800–4800	840,000	Sr, Ba, Sv, Ro
<i>Sporagra magellanica</i>	0–5000	6,080,000	Ba, Sv
<i>Theristicus melanopis</i>	0–3000	1,330,000	Cu, Ba, Sv

**Table 6** (continued)

Bird species <sup>a</sup>	Elevation range (m ASL) <sup>b</sup>	Geographic range (km <sup>2</sup> ) <sup>c</sup>	Foraging microhabitats
<i>Thinocorus orbignyianus</i>	1000–5000	1,570,000	Cu, Ba, Sv
<i>Tringa melanoleuca</i> <sup>DS</sup>	0–4000	4,410,000	Ws
<i>Upucerthia validirostris</i>	3000–5000	522,000	Ba, Sv
<i>Vanellus resplendens</i>	1500–4600	735,000	Ba, Sv

<sup>a</sup> Nomenclature follows Remsen et al. (2016)<sup>b</sup> Lowest and highest elevational records from Birdlife International (2016), e-bird (2016) and Del Hoyo et al. (2016)<sup>c</sup> Geographic distribution records obtained from Birdlife International (2016) and e-bird (2016)

## Appendix 2

**Table 7** Excel spreadsheet with all 16 OLS models by season obtained using 4 variables: isolation, should say: elevation, area and microhabitat heterogeneity. (See attachment)

Season	Rank selection	Intercept	Isolation	Elevation	Area (log)	Microhabitat heterogeneity (log)	df	logLik	AICc	Delta	Weight
Wet	6	2.282	0.000	NA	-1.451	NA	4	-7.032	24.169	0.000	0.327
	5	2.075	NA	NA	-1.600	NA	3	-8.543	24.286	0.117	0.308
	7	-0.069	NA	0.000	-1.419	NA	4	-8.268	26.641	2.471	0.095
	13	2.112	NA	NA	-1.602	-0.026	4	-8.542	27.188	3.019	0.072
	8	1.033	0.000	0.000	-1.354	NA	5	-6.932	27.198	3.029	0.072
	14	2.057	0.000	NA	-1.430	0.166	5	-6.966	27.266	3.096	0.069
	15	-0.243	NA	0.001	-1.398	-0.187	5	-8.199	29.732	5.562	0.020
	16	1.162	0.000	0.000	-1.362	0.093	6	-6.916	30.773	6.603	0.012
	3	-5.550	NA	0.001	NA	NA	3	-12.315	31.830	7.661	0.007
	4	-3.975	-0.001	0.001	NA	NA	4	-11.014	32.133	7.963	0.006
	2	1.342	-0.001	NA	NA	NA	3	-12.783	32.766	8.597	0.004
	1	0.874	NA	NA	NA	NA	2	-14.852	34.276	10.106	0.002
	11	-5.738	NA	0.002	NA	-0.374	4	-12.116	34.337	10.168	0.002
	10	0.792	-0.001	NA	NA	0.431	4	-12.497	35.099	10.930	0.001
	12	-4.041	0.000	0.001	NA	-0.061	5	-11.009	35.351	11.181	0.001
	9	0.636	NA	NA	NA	0.176	3	-14.811	36.822	12.653	0.001
Dry	13	0.731	NA	NA	-1.447	0.872	4	-2.406	14.917	0.000	0.465
	5	2.050	NA	NA	-1.602	NA	3	-4.937	17.074	2.157	0.158
	14	0.779	0.000	NA	-1.386	0.894	5	-2.045	17.423	2.507	0.133
	15	0.402	NA	0.000	-1.421	0.846	5	-2.395	18.123	3.206	0.094
	7	-0.581	NA	0.001	-1.380	NA	4	-4.370	18.845	3.928	0.065
	6	2.114	0.000	NA	-1.556	NA	4	-4.752	19.609	4.693	0.044
	16	0.933	0.000	0.000	-1.397	0.907	6	-2.043	21.026	6.110	0.022
	8	-0.335	0.000	0.000	-1.365	NA	5	-4.282	21.897	6.981	0.014
	3	-5.911	NA	0.001	NA	NA	3	-9.433	26.066	11.149	0.002
	11	-5.171	NA	0.001	NA	0.757	4	-8.432	26.970	12.053	0.001
	9	-0.848	NA	NA	NA	1.230	3	-10.125	27.449	12.533	0.001
	10	-0.599	0.000	NA	NA	1.245	4	-9.153	28.412	13.495	0.001
	4	-5.384	0.000	0.001	NA	NA	4	-9.256	28.617	13.700	0.000
	12	-4.313	0.000	0.001	NA	0.843	5	-8.027	29.386	14.470	0.000
	1	0.848	NA	NA	NA	NA	2	-12.867	30.306	15.389	0.000
	2	1.107	0.000	NA	NA	NA	3	-12.162	31.524	16.608	0.000

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