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Dispersal strength determines meta-community structure in a dendritic riverine network

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Citation	Cañedo-Argüelles, M., Boersma, K. S., Bogan, M. T., Olden, J. D., Phillipsen, I., Schriever, T. A., & Lytle, D. A. (2015), Dispersal strength determines meta-community structure in a dendritic riverine network. Journal of Biogeography, 42(4), 778–790. doi: 10.1111/jbi.12457
DOI	10.1111/jbi.12457
Publisher	John Wiley & Sons Ltd.
Version	Accepted Manuscript
Terms of Use	http://cdss.library.oregonstate.edu/sa-termsofuse



- 1 Original article. 2 Word count (abstract, text and references): 7,697. 3 Estimated number of journal pages required by figures and tables: 3. 4 5 Title: Dispersal strength determines meta-community structure in a dendritic riverine 6 network. 7 8 **Running header:** meta-community structure in a fragmented riverine network 9 Author names: Cañedo-Argüelles, M.^{ab}, Boersma, K.S.^{ac}., Bogan, M.T.^{ad}, Olden, J.D^e., 10 Phillipsen, I.^a, Schriever, T.A.^a, Lytle, D.A.^a. 11 12 13 Author addresses: 14 a Dept. of Integrative Biology, Oregon State Univ., 3029 Cordley Hall, Corvallis, OR 15 97331-2914, USA 16 b Grup de Recerca Freshwater Ecology and Management (FEM), Departament 17 d'Ecologia, Facultat de Biologia, Universitat de Barcelona (UB), Diagonal 645, 08028 18 Barcelona, Spain 19 c Current address: Department of Biology, University of San Diego, 5998 Alcala Park, 20 San Diego, CA 92110, USA 21 d Current address: Department of Environmental Science, Policy and Management, 22 University of California, Berkeley, CA 94720, USA 23 e School of Aquatic and Fishery Sciences, University of Washington, Seattle, WA 24 98105, USA 25 26 Correspondence: Miguel Cañedo-Argüelles. FEM Research Group, University of 27 Barcelona, Av. Diagonal 643, 08028, Barcelona. E-mail address: 28 mcanedo.fem@gmail.com 29 30 Abstract 31 Aim 32 Meta-community structure is a function of both local (site-specific) and regional 33 (landscape-level) ecological factors, and the relative importance of each may be 34 mediated by the dispersal ability of organisms. Here, we used aquatic invertebrate 35 communities to investigate the relationship between local and regional factors in 36 explaining distance decay relationships (DDRs) in fragmented dendritic stream 37 networks. 38 39 Location Dryland streams distributed within a 400 km² section of the San Pedro River basin, 40 41 southeastern Arizona, USA.
- 42
- 43 Methods

- 44 We combined fine-scale local information (flow and habitat characteristics) with
- 45 regional-scale information to explain DDR patterns in community composition of
- 46 aquatic invertebrate species with a wide range of dispersal abilities. We used a novel
- 47 application of a landscape resistance modeling approach (originally developed for
- 48 landscape genetic studies) that simultaneously assessed the importance of local and
- 49 regional ecological factors as well as dispersal ability of organisms.
- 50

51 **Results**

- 52 We found evidence that both local and regional factors influenced aquatic invertebrate 53 DDRs in dryland stream networks, and the importance of each factor depended on the 54 dispersal capacities of the organisms. Local and weak dispersers were more affected by 55 site-specific factors, intermediate dispersers by landscape-level factors, and strong
- 56 dispersers showed no discernable pattern. This resulted in a strongly hump-shaped
- 57 relationship between dispersal ability and landscape-level factors, where only moderate
- dispersers showed evidence of DDRs. Unlike most other studies of dendritic networks,
- 59 our results suggest that overland pathways, using perennial refugia as stepping-stones,
- 60 might be the main dispersal route in fragmented stream networks.
- 61

62 Main conclusions

- We suggest that using a combination of landscape and local distance measures can help
 to unravel meta-community patterns in dendritic systems. Our findings have important
 conservation implications, such as the need to manage river systems for organisms that
 span a wide variety of dispersal abilities and local ecological requirements. Our results
- 67 also highlight the need to preserve perennial refugia in fragmented networks, since they
- 68 may ensure the viability of aquatic meta-communities by facilitating dispersal.
- 69
- 70 Keywords: aquatic invertebrates, dendritic networks, distance decay relationship,
- 71 drought, dispersal, connectivity, landscape resistance, meta-community.

72 Introduction

- Studying patterns of biological diversity has been the foundation of numerous
 ecological pursuits over the past two centuries. Distance decay relationships (DDRs) –
 which describe the biogeographic phenomenon where taxonomic similarity between
 localities decreases or decays as the distance between them increases have received
- considerable interest among ecologists (Nekola & White, 1999). Indeed, this ecological
 pattern is encapsulated in Tobler's first law of geography, which states that 'everything
- pattern is encapsulated in 100fer s first law of geography, which states that everythin
 is related to everything else, but near things are more related than distant things'
- 80 (Tobler, 1970). DDRs have now been studied across a wide range of organisms and
- 81 environments (reviewed in Soininen *et al.* 2007), but are still relatively understudied in
- riverine ecosystems (Thompson & Townsend 2006, Leprieur *et al.* 2009, Brown &
- 83 Swan 2010, Bonada et al. 2012, Warfe et al. 2013). This is largely because streams and
- 84 rivers are organized as complex dendritic networks rather than simple linear systems
- 85 (Benda *et al.*, 2004; Campbell Grant *et al.*, 2007; Erős *et al.*, 2012), thus necessitating
- the incorporation of network connectivity to explore the interactions among
- 87 communities that are linked by dispersal (Fausch *et al.*, 2002).
- 88

The environmental phenomena that drive any particular DDR can be decomposed intolocal and regional factors. Local factors include site-specific attributes such as water

- 91 chemistry and habitat structure that serve as filters, excluding some taxa and favoring
- 92 others (Poff, 1997; Townsend *et al.*, 1997). Regional factors include landscape-level
- 93 features that facilitate or impede the movement of organisms across landscapes. These
- 94 features may include the dendritic structure of stream networks (Fausch *et al.*, 2002;
- 95 Benda *et al.*, 2004), the spatial arrangement of suitable habitat patches across the
- landscape (Campbell Grant *et al.*, 2007; Erős *et al.*, 2012; Phillipsen & Lytle, 2013),
- 97 and the simple Euclidean distance between sites. Therefore, a full understanding of the
- 98 ecological processes underlying DDR patterns in stream networks must account at least
- 99 for three main factors: environmental filters, dispersal of organisms and network100 topology.
- 101

102 Environmental harshness can influence the role of local and regional forces shaping

biodiversity patterns (Brown *et al.*, 2011; Heino, 2011). For example, unstable

104 environments (e.g. aquatic habitats that experience severe, recurrent droughts) show a

105 high degree of niche filtering, allowing only those species adapted to the local

106 conditions to persist (Poff, 1997; Chase, 2007). Therefore, harsh environmental

- 107 conditions may cause meta-communities to be structured by local factors (Urban, 2004)
- 108 and DDRs may not meet the expectation of decreasing community similarity with
- 109 increasing distance.
- 110

111 Dispersal of aquatic-obligate riverine organisms is highly constrained by flow

112 connectivity (Fausch et al., 2002; Hughes, 2007; Schick & Lindley, 2007). In contrast,

- aquatic organisms that can disperse overland, such as flying forms of adult aquatic
- 114 insects, can move both along drainages and across drainage divides (Bilton *et al.*, 2001;
- 115 Petersen *et al.*, 2004). Therefore, variability in dispersal mode and ability will determine

- the extent to which local and regional factors structure assemblages of organisms
- 117 (Bohonak & Jenkins, 2003; Cottenie & De Meester, 2004) and may be reflected in the
- shape of the DDR. For example, studies on aquatic invertebrate meta-communities in
- stream networks have reported that increasing dispersal strength results in a weakening
- 120 of DDRs due to relaxation of dispersal limitation (Thompson & Townsend, 2006;
- 121 Brown *et al.*, 2011). Very high dispersal rates could cause the homogenization of
- 122 communities, since organisms can disperse to all available habitats and only the
- 123 strongest competitors will survive (Kneitel & Miller, 2003; Leibold et al., 2004).
- 124

The network topology, or spatial structure of the river network, also has important implications for dispersal and resulting meta-community structure (Muneepeerakul *et al.*, 2008; Auerbach & Poff, 2011). The use of within-network and overland dispersal pathways by aquatic organisms largely depends on the connectivity between the habitat branches, with the loss of connectivity constraining within-network dispersal (Fagan, 2002). Therefore, within-network DDR should be weaker when populations are disjunct within the river network, as is the case with headwater specialist organisms.

132

133 Although critical for understanding the potential mechanisms shaping DDRs (Brown et 134 al., 2011), landscape resistance to the dispersal of organisms has been largely neglected 135 in meta-community analyses (Moritz et al., 2013). Landscape resistance quantifies "distances" between communities that may yield more biologically informative DDRs 136 137 than straight-line Euclidean distance, such as those associated with barriers to dispersal 138 (e.g. high mountains or cliffs). To the present date, only Euclidean and network distance 139 (i.e. the distance between sites along the riverine dendritic network) have been applied 140 to stream networks, which fails to consider more realistic landscape variables in DDR 141 analyses (McRae et al., 2008).

142

143 In this study, we present a novel application of a landscape resistance modeling approach, originally developed for landscape genetic studies, to understand local and 144 145 regional drivers of community structure. Dryland streams were used as a model system 146 to test how environmental stability, dispersal capacity and network topology interact to structure aquatic meta-communities in dendritic networks. These streams experience 147 148 frequent droughts and floods, which lead to strong niche filtering of stream organisms 149 (Lytle, 2002; Lytle & Poff, 2004), with perennial habitats serving as refugia for species that need water during their entire life cycle to survive (Bogan & Lytle, 2011; Phillipsen 150 151 & Lytle, 2013). Therefore, the high temporal and spatial variation in environmental 152 conditions may disrupt expected longitudinal patterns of species' replacement along the 153 network (Bogan et al., 2013). In these systems DDRs are expected to be weak or non-154 existent, with adjacent sites showing very different aquatic assemblages due to large 155 among-site variation in environmental conditions (e.g. one site may flow year-round, 156 while an adjacent site may flow only during rainy seasons). We focused on aquatic 157 invertebrates because they possess a wide range of dispersal capacities (Bilton et al., 158 2001; Bohonak & Jenkins, 2003) and are present over a wide range of environmental

159 conditions (Rosenberg & Resh, 1993; Merritt *et al.*, 2008), as exemplified by the great
160 diversity of biological traits that they exhibit (Statzner *et al.*, 2004; Poff *et al.*, 2006).

161

162 The aim of our study was to use DDRs to investigate the relationship between local and 163 regional factors in explaining aquatic meta-community structure in fragmented dendritic 164 networks. Here we define fragmentation as the loss of surface water connectivity along 165 the river network. During low precipitation periods (typically in late Spring and 166 Summer) some dryland streams experience very low flow, with sections becoming 167 disconnected pools separated from one another by dry stream reaches. We used fine-168 scale local environmental variables and landscape resistance metrics to quantify the 169 influence of local and regional drivers on DDRs for groups of species with different 170 dispersal capacities. The following hypotheses were tested:

171

172 H1: DDRs should be weak for very strong and very weak dispersers (at the ends of the 173 dispersal gradient) because meta-communities of weak dispersers show little spatial 174 structure and meta-communities of strong dispersers are homogenized by competition. 175 We predicted that DDR would be strongest in species with moderate dispersal strength. 176 H2: Due to high network fragmentation in dryland streams (i.e. longitudinal flow 177 disruption during long dry seasons), no significant DDR should be found when using 178 network distance. We predicted that network distance would have low explanatory 179 power due to high spatial and temporal stream fragmentation, while our approach that 180 considers landscape resistance to dispersal would provide greater explanatory power. 181 H3: Due to strong niche filtering (i.e. high environmental heterogeneity), DDRs 182 associated with flow and environmental characteristics should be consistently stronger 183 than DDRs associated with landscape resistance variables, regardless of the dispersal 184 strength of the organisms. We predicted that DDRs associated with flow and 185 environmental characteristics would be significant, regardless of species' dispersal 186 abilities. 187

188 Methods

189 *Study site and field sampling*

190 We sampled aquatic invertebrates at 28 sites across seven dryland streams distributed within a 400 km² section of the Upper San Pedro River basin, southeastern Arizona, 191 192 USA (Figure 1; Schriever et al., in press). Streams in the area generally have perennial 193 flow in montane headwaters, intermittent flow in upper alluvial fan reaches, ephemeral 194 flow lower on alluvial fans, and then alternating perennial and intermittent reaches in 195 valley rivers (Bogan et al., 2013). We distributed our sample sites among perennial, 196 intermittent and ephemeral reaches (classification follows Levick et al., 2008), but used 197 a continuous flow metric to quantify permanence (see below). Sites were sampled three 198 times a year (March/April, August/September, and November/December) between 2009 199 and 2011. The number of sites and samples collected differed among streams because 200 not all sites had flow or all microhabitats during each sampling event. The study period 201 spanned numerous dry seasons, several periods of ephemeral flows from summer 202 monsoon rains, and one period of intermittent flows resulting from a wet winter. The

203 majority of the sampling occurred during the fall (Nov/Dec) and winter (Mar/Apr)
204 seasons for a total of 144 site x sampling event combinations.

204 205

206 Both riffle and pool microhabitats were sampled at each site, when present. For riffle samples we disturbed 0.33 m^2 of stream substrate to a depth of 5 cm while capturing 207 invertebrates immediately downstream with a D-net (500-µm mesh). Pool samples 208 209 consisted of sweeping the entire pool area including water column, surface, and pool benthos with a D-net at an effort of 10 seconds for every 1 m^2 of pool habitat (following 210 211 Bogan & Lytle, 2007). Abundances from replicate microhabitat samples collected from 212 the same site during the same sampling event (e.g. three riffles in November) were 213 summed for each taxon and divided by the number of replicates to acquire relative 214 abundances. Samples were preserved in 95% ethanol and invertebrates were identified 215 in the laboratory to the finest taxonomic level practical, usually to genus or species for 216 insects (including Chironomidae) and family or order for non-insects.

217

218 During each visit, we measured water temperature, pH (Whatman pH Indicators,

219 Whatman International, Maidstone, England) and conductivity (Milwaukee waterproof 220 EC meter C65; Milwaukee Instruments, Rocky Mount, NC, U.S.A.), visually estimated 221 canopy cover and benthic substrate on a percent cover scale (0 - 100%; substrate 222 categories: silt, sand, gravel, cobble and bedrock). We measured the timing and duration 223 of streamflow through the deployment of 15 electrical resistance sensors (Jaeger & 224 Olden, 2012), each representing the hydrologic conditions at the nearest location of 225 invertebrate sampling. The sensors logged the presence or absence of water in the 226 stream channel at 15 minute intervals from April 2010 to December 2011. From the 227 sensor data, we calculated four hydrologic metrics for each site: % flow permanence by 228 year, mean % flow permanence by season (spring = March-June; monsoon = July-September; fall = October – November; winter = December-April), mean duration 229 230 (number of days) of zero flow periods (ZFP) each year, and total number of ZFP each 231 year. For the two flow permanence metrics and duration of ZFP, we summed 15-minute 232 time periods of both wet and dry conditions for the sampling period and for individual 233 zero flow periods, converting the time unit to either days or years as appropriate for the final stream flow metric. We used an average of 2010 and 2011 flow data to estimate 234 235 flow conditions for the November 2009 invertebrate sampling period (16 samples) that 236 occurred prior to the deployment of sensors. Flow permanence is the percentage of time 237 a given reach is wetted or flowing, while the duration of ZFP indicates how long (in 238 days) a given reach is dry during each drying event. For example, a site with permanent 239 stream flow would have a flow permanence of 100% and would receive a value of 0 for 240 ZFP duration. While these metrics were all calculated from the flow sensor records, 241 they were designed to characterize distinct components of the hydrologic regime that 242 may influence aquatic invertebrate occurrence and abundance.

243

244 Distance measures

We used four regional distance metrics (Table 1, Fig. 2), two of which described to the physical distance between sites (geographic and network distance) and two of which 247 described the resistance of the landscape to dispersal (topography and perennial 248 distance). *Geographic distance* is simply the straight-line Euclidian distance between 249 two sites as determined from map coordinates. Network distance was generated via a 250 least-cost path analysis in ArcGIS 9.3 (Environmental Systems Research Institute, 251 Redlands, CA). For this variable, only one pathway connects each pair of sites, and this 252 pathway is restricted to the stream network. *Topography distance* assumes that dispersal 253 occurs along concave corridors such as streambeds, dry gullies, or low saddle points 254 along mountain ridges. Flying and crawling insect adults are likely to follow these 255 relatively cool and moist pathways to disperse from one wetted site to another (Bogan 256 & Boersma, 2012; Phillipsen & Lytle, 2013). Perennial distance assumes that isolated 257 perennial freshwater habitats act as stepping-stones for dispersal among communities in 258 fragmented dendritic networks. For example, in arid landscapes perennial habitats are 259 known to be critical for the survival of certain aquatic species when rivers cease to flow 260 during droughts (Chester & Robson, 2011).

261

262 We generated the four regional distance measures from landscape data layers obtained 263 from the Arizona State Land Department (www.land.state.az.us). Data layers used in 264 our analyses included a digital elevation model (DEM; 10 m resolution), the stream 265 network of the region (from the National Hydrology Dataset), and a map of perennial 266 stream habitats. The latter was constructed using data for the San Pedro River watershed 267 from the Nature Conservancy (www.azconservation.org) combined with observations 268 from field studies in the region (e.g. Bogan & Lytle, 2007; Bogan et al., 2013). We used 269 ArcGIS 9.3 to generate new data layers and to calculate the distance (km) between all 270 pairs of sites. The distances related to landscape resistance (topography and perennial 271 distances) were generated from the GIS data layers in the form of pixelated maps (i.e. 272 rasters). Each raster map was used as input for the program CIRCUITSCAPE (McRae, 273 2006). CIRCUITSCAPE calculates the resistance of the landscape to dispersal between 274 each pair of sites (analogous to electrical resistance in a circuit diagram), allowing for 275 multiple pathways between sites. This pairwise resistance is a summation of the 276 resistances of individual pixels in the input map. Pixels with high input values are 277 hypothesized to offer high resistance to movement, and vice versa. Thus, pairwise 278 resistances from CIRCUITSCAPE model the structural connectivity of communities, 279 based on the landscape/habitat feature represented by the input map. We used the 280 original values of the map pixels to assign resistance values to the raster maps. Using 281 the original pixel values is more conservative than assigning relative costs of landscape 282 features based on expert opinion (a practice that some have questioned; Spear et al. 283 2010). Before running the CIRCUITSCAPE analysis, we transformed the original 284 values of the maps so that they were all on the same scale (1 for lowest resistance, 10 285 000 for highest resistance; results were qualitatively similar for different values of 286 highest resistance). We performed a separate CIRCUITSCAPE analysis for both 287 topography and perennial distances, generating their two independent data sets of all 288 pairwise resistance distances as output. See Phillipsen & Lytle (2013) for an example of 289 CIRCUITSCAPE output in a population evolution context and how this relates to the 290 underlying distance metrics.

- 291
- In addition to the four regional distance measures, two local ecological distance
- 293 measures were calculated: habitat distance and flow regime distance. *Habitat distance*
- was calculated as the dissimilarity between the multivariate centroids of each pair of
- sites based on their environmental characteristics, including canopy cover, conductivity,
- 296 pH, and % of bedrock, cobble, gravel, sand and fines. Flow regime distance was
- 297 calculated as the dissimilarity between the multivariate centroids of each pair of sites
- from a composite of flow metrics: % flow permanence in year of sample, % flow
- permanence by season, duration of zero flow periods each year (mean) and total number of zero flow periods each year. All variables were normalized (mean = 0; SD = 1)
- 301 before analysis.
- 302
- 303 Statistical analyses

304 Prior to analyses, we placed each of the 225 aquatic invertebrate taxa into one of four 305 categories: weak, local, moderate and strong dispersers (Appendix S1). Weak dispersers 306 (17 taxa) are aquatic obligates that spend nearly all of their life cycle within the stream 307 (e.g. Abedus herberti). Local dispersers (142 taxa) have flying adult stages but can only 308 travel short distances due to their short life cycles and/or weak flying musculature (e.g. 309 Hydrobaenus sp.). Moderate dispersers (64 taxa) have flying adult stages that can travel 310 long distances but cannot cover the entire geographic range of our study (e.g. Enochrus 311 aridus). Strong dispersers (10 taxa) are powerful fliers that can travel between any of 312 the sites in our studied geographic range (e.g. Lethocerus medius). These categories 313 were derived from a trait database specific to the study region built from over 80 314 publications from primary literature, existing databases and expert judgment (Schriever 315 et al., in press). Abundance data were log (x+1) transformed and then used to calculate 316 the Chao dissimilarity index among all pairs of sites (using the function vegdist in the 317 vegan package: Minchin et al., 2013). The Chao index was the most appropriate 318 dissimilarity index to use because each dispersal ability group had a different number of 319 taxa; it is intended to account for the effect of unseen shared species and thus reduce 320 sample-size bias (Chao et al., 2005). Habitat distance, flow regime distance and the four 321 landscape distances (Table 1) were used as independent explanatory variables of Chao's index for each group of species (weak, local, moderate and strong dispersers). Spearman 322 323 correlation tests were performed between all pairs of the explanatory variables. For 324 those variables showing strong correlation (i.e. Spearman's rho > 0.5 and p-value <325 0.01), we used partial Mantel tests (mantel function in vegan package) to compare 326 community data to the explanatory variable of interest while controlling for the 327 correlated variable (Legendre & Legendre, 2012). The differences in the relationship 328 between community dissimilarity and each of the distance metrics (geographic, 329 network, topography, perennial, flow regime, and habitat distances) across dispersal 330 classes was tested through an analysis of covariance (ANCOVA; Legendre & Legendre, 331 2012) with Chao dissimilarity as the dependent variable, each driver as a covariate, and 332 the dispersal class as the grouping factor. 333

334 We fitted linear models to each distance metric, and performed F-tests to assess model performance. Models were tested for linearity using the diagnostics plots for 335 336 generalized linear models (Appendix S2). We used the glm.diag.plots function in R, 337 which makes a plot of jackknife deviance residuals against linear predictor, normal 338 scores plots of standardized deviance residuals, plot of approximate Cook statistics 339 against leverage/(1-leverage), and case plot of Cook statistic. After validating the 340 models we used an information-theoretic approach to compare the contribution of 341 different explanatory variables that best described differences in invertebrate 342 community composition. We derived the log-likelihood for each model and calculated 343 Akaike's information criterion (AIC; Akaike, 1973; Burnham & Anderson, 2002) to 344 rank the models from lowest to highest AIC. We only compared single variable models 345 and the combination of local environmental distance metrics (flow regime and habitat) 346 and landscape distance metrics (geographic, topography, perennial and network 347 distances), since our aim was to compare the importance of local versus regional filters 348 for aquatic invertebrate meta-communities across a gradient of dispersal strength. 349 Information for the rest of the models is shown in Appendix S3. Once the models were 350 ranked, additional information-theoretic metrics were calculated. The difference 351 between the AIC of a particular model and the AIC of the estimated best-fitting model 352 (i.e. the model with the lowest AIC) is Δ AIC. We also calculated Akaike weights, the 353 probability that the model is actually the best-fitting of the candidate models. The sum 354 of Akaike weights across the models is 1.0. When the weight of the model with the 355 lowest AIC is not close to 1.0, there is evidence for model selection uncertainty. We 356 accounted for the non-independence of our data (represented by pairwise distances 357 among sites) by using an R-squared approach for fixed effects in a linear mixed model 358 to adjust for the inflation of sample size (Edwards et al., 2008). Since both the AIC and 359 mixed model approaches yielded similar results, we only report the adjusted R-squares 360 for fixed effects in the Supplementary Material. The selection of AIC over adjusted R-361 squares was based on the limited use of adjusted R-square values in model building, due 362 to the lack of diagnostic and selection tools for linear mixed models (Edwards et al., 363 2008). All analyses were conducted in R version 3.0.2 (R Core Team, 2013) and 364 significance was assigned at P <0.05. Bonferroni correction was used to adjust p-values 365 for multiple comparisons.

366

367 Results

368 The six distance metrics displayed weak pairwise correlations. Only geographic, topography and perennial distance were significantly correlated (Fig. 3). Habitat and 369 370 flow regime distances significantly explained community dissimilarity, regardless of the species' dispersal abilities (Table 2). Geographic, topography and perennial distances 371 372 significantly explained community dissimilarity for all dispersal groups except for the 373 weak dispersers, and had a higher explanatory power for strong dispersers (Table 2). 374 Network distance was not a significant predictor for any dispersal group. Associations 375 between community dissimilarity and the explanatory variables varied considerably 376 among the four dispersal groups (Fig. 4). The relationship between community

dissimilarity and the six distance metrics differed across the dispersal categories for allbut one metric (network distance; Table 3).

379

380 Partial Mantel tests revealed that: (1) geographic distance was significantly correlated 381 with the composition of strong dispersing taxa after controlling for topography, (2) 382 topography was significantly correlated with the composition of moderate dispersers 383 while accounting for geographic distance and (3) topography and perennial distance 384 correlated with the composition of moderate dispersers while controlling for each other 385 (Table 4). Of all the models, local drivers (habitat distance plus flow regime distance) demonstrated the strongest relationships (i.e. lowest AIC for single variable models) 386 387 with the composition of weak, local and strong dispersers, while regional drivers 388 (geographic, topography, perennial and network distances) best-explained moderate 389 dispersers (Table 5).

390

391 Discussion

392 We used distance decay relationships to examine the importance of local and regional 393 drivers of aquatic invertebrate meta-community structure in dryland streams. DDRs 394 have been used to examine the interaction of processes operating at local and regional 395 scales for a wide range of organisms and ecosystems (Cottenie, 2005; Soininen et al., 396 2007), including stream networks (Thompson & Townsend, 2006; Leprieur et al., 2009; 397 Brown & Swan, 2010; Warfe et al., 2013). However, past studies have largely not 398 explored the relative roles of local- and regional-scale landscape drivers within 399 connected networks. The only published study that used DDRs to address the effect of 400 hydrological connectivity on stream meta-communities showed, rather counterintuitively, that the loss of connectivity enhanced DDR in a variety of organisms 401 402 (Warfe et al., 2013). However, environmental conditions in that study were not 403 independent of geographic distance; therefore dispersal limitation and niche partitioning

- both played a role in shaping assemblage structure. Our results suggest that
- fragmentation impacts DDRs by altering the viable dispersal pathways, with organisms
 dispersing overland instead of using the stream network. As we hypothesized, meta-
- dispersing overland instead of using the stream network. As we hypothesized, meta-community structure was determined by three main factors: niche filtering due to local
- 408 among-habitat differences, dispersal ability of the species and landscape resistance
- 409 (geographic distance, topography and availability of perennial refugia).
- 410
- 411 Niche filtering due to among-habitat differences
- 412 We hypothesized that high environmental heterogeneity would generate significant
- 413 environmental DDRs for all dispersal categories. Indeed, we found that the local filters
- 414 of habitat and flow regime had a stronger effect on community dissimilarity than
- 415 regional filters across all dispersal classes, except for moderate dispersers. At
- 416 intermediate levels of dispersal, organisms might be able to survive harsh
- 417 environmental conditions such as floods and droughts by escaping and finding refugia
- 418 (Velasco & Millan, 1998; Lytle, 1999), but at the same time they might not be able to
- 419 reach all the available habitats, preventing meta-community homogenization via mass
- 420 effects (Kneitel & Miller, 2003; Leibold et al., 2004). The significant influence of local

421 filters on meta-community patterns was not surprising given the high temporal and 422 spatial variation in environmental conditions in the study region. Highly variable 423 seasonal and interannual precipitation patterns and various geomorphic settings (e.g. 424 bedrock canyons, alluvial fans) interact to create a patchy stream landscape. In these 425 dryland streams, perennial reaches are adjacent to intermittent reaches, and intermittent 426 reaches with seasonal flow in a wet year can be completely dry during the same period 427 in a dry year (Jaeger & Olden, 2012, Bogan et al., 2013). The amount of water and how 428 it is distributed within the year (i.e. frequency and timing of droughts and floods) have 429 important consequences for water quality and habitat variables (e.g. canopy cover, river 430 substratum). Accordingly, these local habitat filters should be extremely strong in 431 dryland streams. Our results validate previous studies that linked these patchy 432 environmental conditions to disruptions in the longitudinal patterns of stream 433 invertebrate communities in the region (Bogan et al., 2013). They are also in agreement 434 with a recent study from northwestern Australia, which found that flow and channel 435 width best explained invertebrate meta-community patterns across a range of perennial 436 and intermittent streams (Warfe et al., 2013). Moreover, the importance of niche 437 filtering in structuring meta-communities has been demonstrated for a variety of 438 ecosystems (Soininen et al., 2007), including ponds (Urban, 2004; Chase, 2007) and 439 streams (Thompson & Townsend, 2006; Brown & Swan, 2010). Since the loss of 440 connectivity among stream reaches results from high flow heterogeneity (leading to 441 high environmental heterogeneity), niche filtering can be expected to be strong in 442 fragmented stream networks.

443

444 Dispersal ability

445 Aquatic invertebrates are known to have very different dispersal capacities, ranging 446 from a few meters to thousands of kilometers (Kovats et al., 1996; Bilton et al., 2001; 447 McCauley, 2006). Given that meta-community structure is highly dependent on 448 geographic scale (Brown et al., 2011; Maloney & Munguia, 2011; Nekola & McGill, 449 2014), differences in dispersal can be expected to affect DDRs (Nekola & White 1999). 450 Increasing dispersal ability is expected to enhance community similarity among sites 451 and reduce beta diversity among habitat patches (Shurin et al., 2009). Recent studies on invertebrate meta-communities in stream networks have provided evidence for this 452 453 pattern, with DDR being weakened by increasing dispersal strength (Thompson & 454 Townsend, 2006; Brown & Swan, 2010; Bonada et al. 2012). However, we found a 455 more complex unimodal pattern, with DDR peaking at intermediate dispersal strength 456 for different measures of landscape resistance. This pattern might result from 457 assemblages of weak dispersers showing no spatial structure due to dispersal limitation 458 while assemblages of strong dispersers are more homogeneous across the landscape due 459 to the absence of dispersal restrictions (Kneitel & Miller, 2003; Leibold et al., 2004). 460 Figure 5 illustrates this pattern by showing the explanatory power of local (habitat and 461 flow distances) and landscape (geographic distance, topography and availability of 462 perennial refugia) filters along the dispersal strength gradient.

464 Our results could be influenced by the lower flow connectivity and environmental 465 stability in our dryland study system when compared to more mesic stream systems. In 466 low connectivity systems, weak dispersers are highly isolated, leading to species 467 distributions ruled by ecological drift and niche filtering (Hu et al., 2006). Therefore, in 468 these fragmented systems, low connectivity coupled with differing environmental 469 conditions can lead to adjacent sites having very different assemblages of weak 470 dispersal species. Previous investigations on the flightless aquatic obligate Abedus 471 herberti within our study area reported strong population genetic structure, with 472 populations within the same stream drainage (less than 5 km apart) showing significant 473 genetic differentiation (Finn et al., 2007; Phillipsen & Lytle, 2013). This same pattern 474 has been observed for the blackfly *Prosimulium neomacropyga* in isolated alpine 475 headwater streams (Finn & Poff, 2011). On the other side of the spectrum, extremely 476 strong dispersers can break down geographic barriers, occurring in all suitable habitats 477 (Townsend et al., 2003; McCauley, 2006; Thompson & Townsend, 2006; Brown & 478 Swan, 2010; Bonada et al. 2012). This would explain the decrease in the explanatory 479 power of landscape variables over meta-community structure at the upper end of the 480 dispersal strength gradient in the present study (Fig. 5). We suggest that regional drivers 481 should be important predictors of meta-community structure up to a certain dispersal 482 distance threshold, beyond which dispersal is strong enough to break the limitation 483 imposed by geographical barriers.

484

485 Distance among sites

486 Network distance did not significantly affect community dissimilarity for any of the 487 four dispersal groups, as we hypothesized. This contradicts the general rule of aquatic 488 invertebrates using the stream network as the main 'highway' for dispersal (Petersen et 489 al., 2004). While evidence supporting the ideas of the stream channel as the primary 490 dispersal route and restricted overland dispersal between catchments continue to 491 accumulate in the literature (Hughes, 2007; Brown & Swan, 2010; Rouquette et al., 492 2013), previous investigations within our study area suggest that dryland streams might 493 be exceptions to this rule. Many aquatic invertebrate species in our study region 494 disperse laterally from stream corridors in search of other wetted habitats (Bogan & 495 Boersma, 2012). Additionally, Bogan et al., (2013) reported an interruption of the river 496 continuum, where invertebrate communities in distant headwater and lowland perennial 497 streams were more similar to one another than to those in intervening intermittent 498 reaches. Furthermore, Phillipsen & Lytle (2013) found no significant relation between 499 network distance and population genetic structure of Abedus herberti. Instead, they 500 found that topography best explained genetic structure and suggested that overland 501 dispersal resulted from flood-escape behavior (Lytle, 1999; Lytle et al., 2008), where 502 individuals crawl from streams during floods and accidentally end up in adjacent 503 drainages.

504

We found significant (but generally weak) DDRs for perennial habitat distance in all
cases and for topography distance in all cases except weak dispersers. This supports the
hypothesis of overland dispersal (flight and crawling) being the main dispersal pathway

- 508 for aquatic invertebrates in highly fragmented stream networks, such as those inhabiting
- 509 dryland regions. Similarly, Campbell Grant *et al.* (2010) found evidence of high
- 510 overland dispersal rates in newly metamorphosed juveniles of stream salamanders and
- 511 suggested that the salamanders followed that dispersal strategy to increase population
- 512 persistence across isolated headwater streams.
- 513

514 Conclusion

515 Our DDR analyses suggests that in highly heterogeneous stream networks, where 516 environmental conditions vary greatly across space and time, local factors (i.e. niche 517 filtering) may swamp regional influences (i.e. landscape filters) on aquatic invertebrate 518 meta-community structure. However, this interaction between local and regional factors 519 is dependent on a species' dispersal capacity, which determines their ability to colonize 520 suitable habitats. Using a combination of landscape and local distance measures, we 521 found evidence that local and weak dispersers were affected by local-scale factors, 522 intermediate dispersers were affected by landscape-level factors, and strong dispersers 523 showed no discernable pattern. This resulted in a hump-shaped relationship between 524 dispersal ability and landscape-level factors, where only moderate dispersers showed 525 significant DDRs. Stream corridors may not be a primary dispersal pathway in these 526 networks, where frequent drought and flood disturbances generate habitat patches with 527 low connectivity. Overland pathways, using perennial refugia as stepping-stones, might 528 be the main dispersal route for aquatic invertebrates in these dryland stream networks 529 (Phillipsen & Lytle, 2013).

530

531 Our DDR approach has the potential to generate timely management insights, such as 532 the importance of preserving perennial habitat patches in fragmented river networks. 533 Meta-communities of weak dispersal species are highly disconnected in dryland stream 534 networks. Since most of these species depend on perennial water sources for their 535 survival (Bogan & Lytle, 2011; Hermoso et al., 2013), intensifying droughts may have 536 effects at both meta-population and meta-community levels. Our results suggest that 537 perennial habitat patches may facilitate dispersal of aquatic invertebrates and thereby 538 may ensure the long-term viability of populations. Conserving perennial habitats is of vital importance in dryland streams, and it will become increasingly important in basins 539 540 experiencing flow reduction due to warmer temperatures and increased anthropogenic 541 water use (Marshall et al., 2010). Given the different responses of invertebrate meta-542 communities to our various geographic and environmental distance measures, we 543 encourage future investigations to incorporate multiple regionally relevant measures of 544 landscape resistance into their studies. Further research is needed to better understand 545 how environmental stability affects the balance between local and regional factors 546 structuring meta-community patterns in dendritic networks, including work at different 547 spatial scales and degrees of fragmentation.

548

549 Acknowledgments

550 Funding was provided by the U.S. Department of Defense - Strategic Environmental

551 Research and Development Program (RC-1724 and RC-2203). We thank Richard Van

- 552 Driesche for help with invertebrate identification, Dr. Narcís Prat for help with
- 553 Chironomidae taxonomy, Sheridan Stone for logistical support at Fort Huachuca, AZ,
- 554 Brooke Gebow and the Nature Conservancy for access to streams and lodging at the
- 555 Ramsey Canyon Preserve, AZ, and to Frank McChesney for access to the Babocomari
- 556 River, AZ. Daniel Trovillion, Karima Ibrahim, and Katherine Delf assisted with sample
- 557 processing. Miguel Cañedo-Argüelles was supported by a Spanish Government Scholar
- 558 Cátedra Principe de Asturias Grant.
- 559

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- 759 760

761 SUPPORTING INFORMATION

Additional Supporting Information may be found in the online version of this article: 763

Appendix S1: aquatic invertebrate taxa list (sampled in 28 sites across seven dryland streams distributed within a 400 km² section of the Upper San Pedro River basin, southeastern Arizona, USA) including their assignation into one of four dispersal categories: weak, local, moderate and strong dispersers.

- Appendix S2: diagnostics plots for generalized linear models fitted to each distance metric and aquatic
 invertebrates' dispersal category.
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Appendix S3: sum of residuals, likelihood ratios and AIC values for all combinations of explanatory
variables (habitat, flow regime, geographic, topography, perennial and network) for each dispersal group
(weak, local, moderate and strong).

776 BIOSKETCHES

- Miguel Cañedo-Argüelles is an aquatic ecologist, interested in the response of aquatic
 ecosystems to disturbance and how aquatic organisms are connected through the
 landscape.
- 780781 Author contributions (in alphabetical order)
- 782 Designed research (MC, DL, IP), performed research (DL, MB, JO, KB), analyzed data
 783 (MC, IP), wrote the paper (all authors).
- 784
- 785 Editor: Ladle, Richard.
- 786

787 TABLES

Table 1: Details of the four regional distance metrics used in this study.

Distance metric	Explanation	Hypothesized relationship to species flow
Geographic distance	Straight-line distance between sites in two-dimensional space.	Dispersal increases when the geographic distance between a pair of sites decreases.
Topography	Pairwise resistances between sites based on low resistance of map pixels with concave topography and high resistance of pixels with convex topography.	Dispersal is highest in areas with strongly concave topography. Dispersal is lowest across areas with strongly convex topography.

Perennial	Pairwise resistances between sites based on low resistance of map pixels in patches of perennial freshwater habitats and high resistance of pixels in the matrix between these patches.	Dispersal increases in the presence of perennial freshwater habitats.
Network distance (stream network)	Pairwise least-cost paths between sites that strictly follow the stream/river network. Only one path exists between any pair of sites.	Dispersal occurs only within the stream/river network.

Table 2: Adjusted R-square, F statistic and p-value for each combination of aquatic invertebrate dispersal
category (weak, local, moderate, and strong) and explanatory environmental (habitat distance and flow
regime distance) and spatial (geographic, topography, perennial and network distance) variables.
Bonferroni correction was used to adjust p-values for multiple comparisons. Significant relationships
(p<0.05) are shown in bold.

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Explanatory variable		Wea	k		Loca	1		Modera	ate		Stron	ıg
	Adj R ²	F	p-value	Adj R ²	F	p-value	Adj R ²	F	p-value	Adj R ²	F	p-value
Habitat	0.13	51.1	<0.01	0.36	195.1	<0.01	0.08	33.3	<0.01	0.15	61.1	<0.01
Flow regime	0.09	37.2	<0.01	0.25	119.8	<0.01	0.08	29.6	<0.01	0.11	46.4	<0.01
Geographic distance	0	2.4	0.120	0.03	10.5	0.001	0.07	29.1	<0.01	0.09	37.7	<0.01
Topography	0	0	0.964	0.06	25.3	<0.01	0.24	109.2	<0.01	0.02	9.6	0.002
Perennial	0.02	8	0.005	0.1	40.9	<0.01	0.27	129.3	<0.01	0.01	5.6	0.019
Network distance	0	1.7	0.195	0	0.6	0.457	0	0.1	0.788	0	0.5	0.481

Table 3: Results of the ANCOVA analyses with Chao dissimilarity value as the dependent variable, each
 distance metric as a covariate, and the aquatic invertebrate dispersal category (weak, local, moderate, and

strong) as the grouping factor. Significant relationships (p<0.05) are shown in bold.

	F statistic	p-value
Habitat	5.11	<0.01
Flow regime	2.69	0.045
Geographic		
distance	12.41	<0.01
Topography	13.44	<0.01
Perennial	9.82	<0.01
Network distance	0.23	0.873

Table 4: Partial Mantel test results among those distance metrics that showed strong correlation (Fig. 3).

 $BO3 \qquad GEO = geographic distance; TOP = topography; PRN = perennial distance; * = 0.01$

0.01; ns = not significant.

Comparison	Control matrix	r p
Weak dispersers vs GEO	ТОР	-0.12 ns
Local dispersers vs GEO	TOP	-0.02 ns
Moderate dispersers vs GEO	TOP	-0.11 ns
Strong dispersers vs GEO	TOP	0.28 **

Weak dispersers vs TOP	GEO	0.08	ns
Local dispersers vs TOP	GEO	0.2	ns
Moderate dispersers vs TOP	GEO	0.43	**
Strong dispersers vs TOP	GEO	-0.09	ns
Weak dispersers vs PRN	ТОР	0.19	ns
Local dispersers vs PRN	TOP	0.22	ns
Moderate dispersers vs PRN	TOP	0.33	**
Strong dispersers vs PRN	ТОР	0.04	ns
Weak dispersers vs TOP	PRN	-0.11	ns
Local dispersers vs TOP	PRN	0.09	ns
Moderate dispersers vs TOP	PRN	0.26	*
Strong dispersers vs TOP	PRN	0.11	ns

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Table 5: ΔAIC (Akaike's information criterion) and weight for single variable models and the

807 combinations (in italics) of habitat distance and flow regime distance (local distance metrics) and

808 geographic, topography, perennial and network distance (regional distance metrics). The lower the AIC,

809 the higher the explanatory power of the model for each of the four dispersal categories (weak, local,

810 moderate and strong). Lowest AIC of single variables and combinations of variables for each dispersal

811 category are marked in bold.

	Weak		Local	Local Moderate		Strong		
	ΔAIC	Weight	ΔAIC	Weight	ΔAIC	Weight	ΔAIC	Weight
Habitat	15	< 0.001	60	< 0.001	98	< 0.001	20	< 0.001
Flow regime	27	< 0.001	112	< 0.001	102	< 0.001	32	< 0.001
Geographic distance	60	< 0.001	205	< 0.001	102	< 0.001	40	< 0.001
Topography	63	< 0.001	191	< 0.001	35	< 0.001	67	< 0.001
Perennial	55	< 0.001	177	< 0.001	20	< 0.001	71	< 0.001
Network distance	61	< 0.001	215	< 0.001	130	< 0.001	76	< 0.001
Local drivers	0	1	0	1	86	< 0.001	0	1
Regional drivers	53	< 0.001	179	< 0.001	0	1	37	< 0.001

814 FIGURE CAPTIONS

- 815 Figure 1 Map of sampling localities for stream invertebrates in southeastern Arizona (USA). Inset map
- 816 shows the location of the study area. Continuous blue line = perennial streams. Short-dashed green line =
- 817 intermittent streams. Long-dashed red line = ephemeral streams. Black circles represent sampling
- 818 localities. The map is based on a DEM (digital elevation model) at 10 m resolution.



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Figure 2: Hypothetical scenarios of species dispersal among sites in dryland streams based on each of the four regional distance metrics. In each scenario, the locations of three hypothetical communities are shown as white circles in a generic mountain landscape. Streams are depicted by dotted lines and thick black lines with arrowheads represent bi-directional species flow between pairs of communities. The paths of species dispersal in each scenario are determined by the hypothesized resistance to dispersal associated with the given landscape variable. In the Topography scenario, for example, the underlying hypothesis is that dispersal is easiest in areas with strongly concave topography. Thus, gene flow is 828 expected to be highest through areas with concave topography (shown as grey polygons in the figure). See the Table 1 for more detail on each of the regional distance metrics.



Figure 3: Pairwise correlations among all local and landscape distance metrics calculated between 28
sites across seven dryland streams distributed within a 400 km² section of the Upper San Pedro River
basin, southeastern Arizona, USA. HAB = habitat distance; GEO = geographic distance; TOP =
topography; FLO = flow regime distance; NTW = network distance; PRN = perennial distance. Blue
color indicates a positive correlation, while red color indicates a negative correlation. The intensity of the
color indicates the strength of the correlation. Spearman's rho are shown inside each box. * = 0.01 < p-
value < 0.05; ** = 0.001 < p-value < 0.01; *** = p-value < 0.001.



Figure 4: Distance decay relationships for each dispersal group of stream invertebrates sampled at 28
sites across seven dryland streams distributed within a 400 km² section of the Upper San Pedro River
basin, southeastern Arizona, USA. WEAK = weak dispersers; LOCAL = local dispersers; MODERATE
moderate dispersers; STRONG = strong dispersers.



846 Figure 5: Explanatory power of three different groups of distance metrics over stream invertebrates' 847 community dissimilarity among 28 sites across seven dryland streams distributed within a 400 km² 848 section of the Upper San Pedro River basin, southeastern Arizona, USA. Groups of distances: a) habitat 849 & flow distances; b) geographic, topography & perennial distances; and c) network distance. Group a 850 represents local environmental filters and group b represents regional landscape filters. The network 851 distance is shown separately as having a very low power to predict community dissimilarity. The 852 explanatory power is plotted against different categories of dispersal strength of stream invertebrates: 853 weak, local, moderate and strong (see methods section for a description of each category).



- Habitat and flow distances
- Geographic, topographic and perennial distances
- Network distance