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10-10-2011

# River drying lowers the diversity and alters the composition of an assemblage of desert riparian arthropods

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1	River drying lowers diversity and alters composition of a desert riparian
2	arthropod community
3	
4	Running title: Riparian arthropods along a drying river
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12	Keywords: River drying, drought, riparian, arthropod, water availability
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#### Summary 15 16 1. Many studies have shown negative effects of river drying on in-stream animals. However, the influence of river drying on riparian animals remains poorly studied. 17 18 2. We examined ground-dwelling riparian arthropod assemblages along a drying section of 19 the semi-arid San Pedro River in southeastern Arizona, USA. 20 3. We found strong differences in assemblage composition, taxon diversity and the 21 abundance of key taxa between dry and flowing sites, with higher diversity and 22 abundance of most taxa at flowing sites. 23 4. Changes in assemblage composition, taxon diversity, and abundance of representative 24 taxa were associated with a combined measure of water availability that included distance 25 to water and type of water. Other environmental variables showed a weaker association 26 with changes in these arthropod assemblages. 27 5. Thus, we found evidence that desert riparian arthropods are sensitive to river drying and 28 to reduction in water resources. Increases in drying along this river may reduce diversity 29 and the abundance of many groups of ground-dwelling arthropods leading to marked 30 shifts in community composition.

### 32 Introduction

Rivers around the world are drying with increasing frequency, including large rivers, such as the Yellow River in China and the Colorado in the USA (e.g. Zusman, 2000; Tockner & Stanford, 2002; Gleick, 2003; Pearce, 2006; Stone & Jia, 2006). Droughts related to climate change and increased human appropriation of water resources have led to declining ground and surface waters in many regions, contributing to river drying (e.g. Pool & Coes, 1999).

38 Much recent research has concentrated on the effects of river drying on in-stream aquatic 39 communities and ecosystems (e.g. Stanley et al., 1994; Feminella, 1996; Baldwin & Mitchell, 40 2000; Arthington et al., 2010; Larned et al., 2010; Ludlam & Magoulick, 2010). This work has documented strong effects of river drying on the dynamics of aquatic macroinvertebrates, and 41 42 their taxonomic richness, abundance and community composition (e.g. Clarke et al., 2010), as 43 well as on ecosystem processes such as litter decomposition (e.g. Leberfinger, Bohman & 44 Herrmann, 2010). Some research has suggested that, after rewetting, previously dry reaches are quickly re-colonized from nearby perennial reaches (Clarke et al., 2010) or potentially from 45 46 underground refugia or resistant life-stages (Stubbington et al., 2009), minimizing long-term 47 effects, whereas others have found lasting effects of drying events (Sponseller et al., 2010). One 48 recent study has implicated river permanence as a key factor controlling aquatic food chain 49 length, with much shorter food chains in rivers that dried at some point within the last decade 50 (Sabo *et al.*, 2010). Clearly, and as one might expect, river communities and ecosystems are 51 greatly influenced by river drying.

52 Terrestrial organisms living near the river may also be strongly linked to declining 53 surface and groundwater. For example, changes in riparian plant community composition and 54 diversity in the southwestern US are associated with altered flow regimes (Stromberg, Tiller &

55 Richter, 1996; Baird, Stromberg & Maddock, 2005; Lite, Bagstad & Stromberg, 2005; Lite &

56 Stromberg, 2005; Stromberg *et al.*, 2005; Stromberg *et al.*, 2007a). The composition of

57 dominant woody species shifts from cottonwood (Populus spp.) and willow (Salix spp.) to

58 tamarisk (*Tamarix* spp.), as spring floods are reduced by flow regulation, groundwater drops and

rivers dry (Rood & Mahoney, 1990; Stromberg et al., 2007b; Stromberg & Tellmann, 2009).

60 The diversity of herbaceous plants is also affected by these changes in flow (Stromberg *et al.*,

61 2005).

Recent research has sought to relate changes in riparian arthropod assemblages with 62 63 floods, river regulation and other changes in flow regime (Ellis, Crawford & Molles, 2001; 64 Cartron et al., 2003; Paetzold, Yoshimura & Tockner, 2008). For example, Lambeets et al. (2008) explored the associations between a suite of environmental factors and spider and carabid 65 66 beetle assemblages along the Meuse River in N. Europe, finding that the composition of these assemblages varied with flooding disturbance, vegetation density and siltation. Direct 67 68 examination of the effects of *river drying* on riparian arthropods is still needed. 69 In contrast to the rich literature showing effects of reduced flows and drying on aquatic 70 organisms and riparian plants, and some research relating the flood regime with riparian animal 71 communities, less is known about the effects of drying on riparian animals. Since the aquatic 72 and terrestrial components of riverine landscapes are strongly connected and riparian consumers 73 often depend on aquatic subsidies (Nakano & Murakami, 2001; Baxter, Fausch & Saunders, 74 2005), one would expect drying to influence riparian animal communities. However this effect 75 may be reduced compared to aquatic communities, considering the high productivity of terrestrial riparian zones (National\_Research\_Council, 2002) and the ability for some riparian 76

consumers to switch to terrestrial prey (Sabo & Power, 2002). Additionally, in dryland regions,

78	river drying may directly influence riparian communities through a decrease in water
79	availability-decreases have direct consequences for animal physiology (Hadley, 1994),
80	behaviour (Davis & Denardo, 2006) and trophic interactions (McCluney & Sabo, 2009). One
81	recent study also showed a decline in the abundance of riparian fishing spiders with river drying
82	in New Zealand (Greenwood & McIntosh, 2010). Drying was associated with a decline in
83	aquatic, but not terrestrial, prey. In the laboratory, these spiders were shown to be intolerant to
84	desiccation, which was alleviated by access to moist prey (similar to McCluney & Sabo, 2009).
85	Thus, the river provided this species with important food and water resources.
86	Here we ask how the distribution and diversity of riparian arthropods varies along a
87	drying section of the San Pedro River, a desert river in semi-arid southeastern Arizona, USA.
88	We expected that riparian arthropod assemblages would be influenced by river drying, but that
89	the magnitude of effects would vary among taxonomic groups. More specifically, riparian
90	predators that are known to rely on subsidies of aquatic invertebrates are likely to be most
91	strongly influenced by river drying, but the composition of the entire arthropod community could
92	also be altered through combinations of changes in water availability and trophic interactions.
93	Therefore, we (1) compared arthropod assemblage composition, diversity and abundance along
94	dry and flowing sections of a 6 km section of the river (Fig 1) and (2) examined associations
95	between biological responses and several environmental factors, including availability of river-
96	derived resources.

97

# 98 *Methods*

99 Study System

100 The San Pedro River is one of the last free-flowing rivers in the western US, originating 101 in northern Mexico and flowing north across the US border for over 100 km until reaching the 102 Gila River, part of the Colorado River drainage. The San Pedro River experiences seasonal 103 periods of drying that have been exacerbated by recent droughts and increased withdrawals of 104 groundwater (Pool & Coes, 1999). Our research was conducted along a section that has dried 105 with increasing frequency in recent years, partly due to municipal pumping from the regional 106 aquifer and local climate change (Pool & Coes, 1999). Floodplain habitats along the San Pedro 107 are characterized by cottonwood (*Populus fremontii* S. Watson) and willow (*Salix gooddingii* 108 C.R. Ball) trees, whereas uplands are characterized by plants of the Chihuahuan desert scrub to 109 the south and Sonoran desert to the north. In places where surface water permanence has 110 declined and groundwater tables have fallen substantially, the floodplain is often dominated by 111 introduced Tamarisk (Tamarix chinensis Lour). The river is subject to dramatic changes in 112 stream flow and experiences severe floods in the late-summer rainy season. A wide floodplain 113 has developed along much of the river, which is often covered by riparian trees and a forest floor 114 blanketed with leaf litter.

115 The San Pedro River valley harbours a high diversity of birds (~100 breeding species and 116 another 250 migratory), mammals (~80 species), and reptiles and amphibians (~65 species), 117 including endangered species, such as the southwestern willow flycatcher (Empidonax traillii 118 extimus Phillips) (Stromberg & Tellmann, 2009). This river and other riparian areas in the 119 southwest appear to be important stop-over points for migrating birds (Skagen *et al.*, 1998). 120 Thus, this region has been identified as particularly important for conservation purposes by The 121 Nature Conservancy. A large section of the riparian corridor in the upper basin is designated as a 122 national conservation area and is managed by the US Bureau of Land Management. However,

the San Pedro is affected by land-use changes outside the conservation area, through effects of
regional groundwater pumping on baseflows within the river (Pool & Coes, 1999).
Understanding the effects of groundwater pumping on this river is a high priority (Stromberg &
Tellmann, 2009). Our study took place along the upper San Pedro, including both flowing and
drying sections of the river channel adjacent to Boquillas Ranch House (31°41'50.95" N,
110°10'57.15" W), near the town of Tombstone, AZ (Fig 1). This portion of the San Pedro, in

particular, is thought to be influenced by municipal groundwater pumping (Pool & Coes, 1999). *Methods*

131 We collected pitfall trap samples in the dry portion of the active channel along a 6 km 132 section of the river (Fig 1: see Supporting Information Text S1 for details of construction). 133 While most of the samples were collected in unmanipulated areas, 10 of the sampling locations 134 were near artificially constructed and maintained pools measuring approximately 1 m in diameter 135 (Fig 1; Text S1). The river was flowing at all sites on 14 May 2006, but had dried near many of 136 the sites by 25 June 2006. There were a total of 62 sites over the entire experimental period, but 137 only 35 sites on the final sampling date (25 June 2006; see Supporting Information Table S1). 138 While our initial sampling design was intended to include sites near the flowing river, natural 139 pools, artificial pools and dry sites, the river dried more quickly and completely than anticipated, 140 so we were forced to chose new natural pool and flowing sites over the course of the sampling 141 period, while abandoning some of our previous sites (due to sampling constraints). 142 Two pitfall traps were set at each site, with one within 0.5 m on either side of the water 143 body that was present at initial placement. Within a week of each sampling event (in between

144 each sampling event), we recorded the distance between each trap and the nearest water body

145 (flowing river, natural pool, artificial pool). We did not measure distances beyond 55 m,

146 considering these locations to be dry. Once a site had dried, it stayed dry, with no reinstatement 147 of flows during the study period. We also measured the distance from each trap to the nearest 148 channel bank, as our traps were generally located in the dry portion of the river channel with low 149 cover of litter and vegetation. This variable may be of importance, because arthropods may 150 inhabit banks or floodplain habitats with more litter and vegetation during the day, but move into 151 the channel at night. For instance, we have observed field crickets (Gryllus alogus Rehn) 152 moving from channel banks into the dry portion of the channel at sunset. We also measured the 153 percent cover of herbaceous vegetation within a 1 m diameter circle of the trap by visual 154 approximation in 5% categories. Further, we classified the substratum near the trap by visual 155 approximation of soil type and rock size classes. Later, these descriptors were categorized into a 156 structural rating between 0 and 5 (see Table 1). On the final sampling date, we also measured 157 the relative cover of leaf litter in four categories (none, low, medium or high). On this date, we 158 recorded the presence of cottonwood or willow trees near the traps.

159 Arthropods were collected in pitfall traps lined with Tangle-trap<sup>™</sup> (The Tanglefoot 160 Company, Grand Rapids, MI, USA) on the bottom 4 cm of 473 mL (16 oz) party cups and left 161 open for 24 hours. Traps were processed by freezing, thawing, soaking in mineral oil and 162 filtering (0.5 µm) (see Text S1). Due to biases inherent in our collection and processing, we 163 excluded all arthropods less than 1 mm in length from our analysis as well as all mites and 164 Collembola. All remaining arthropods were typically identified to family. Ground beetles 165 (family: Carabidae) were identified to genus. We used Borrer et al. (1992), Ubick et al. (2005) 166 and Arnett & Thomas (2000) to identify our samples. We also categorized arthropods into 167 feeding guilds using Arnett & Thomas (2000), Hamback et al. (2009), Hering (1998) and general 168 knowledge (e.g. all spiders in our samples were assumed to be predators).

169 Statistical analysis

170 GENERAL DESCRIPTION—To assess the effect of river drying, we first quantified differences in 171 the riparian arthropod community between locations that were dry or adjacent to flowing river 172 (Fig 1). We then examined associations between environmental factors (Table 1) and the 173 arthropod community. This second analysis included additional sites near natural and artificial 174 pools (Fig 1; Table S1). In all cases, we conducted analyses at two taxonomic levels. First, we 175 examined all arthropod families collected in traps, including flying insects. Second, we 176 conducted analyses for genera in the ground beetle family Carabidae. We also examined total 177 abundance of all arthropods, arthropods known to be predators and those known to consume 178 some animal material (this included omnivores). Similar techniques were used for analyses of 179 both taxonomic groups. We note any differences below.

180 We typically collected two traps per sampling location, but for statistical analyses we 181 calculated mean values for each location. Levels of environmental characteristics measured for 182 each trap were also averaged. Thus we obtained single values for the abundance and 183 environmental characteristics of each location. Due to difficulties in maintaining our desired 184 river state categories in the face of rapid river drying, we sampled some sites repeatedly for all 185 four collections, while other sites were sampled only once, twice, or thrice. Statistical 186 approaches that dealt adequately with this incomplete repeated sampling were not available. 187 Thus, we chose only the final sampling date for all our statistical analyses (when there was 188 substantial variation in stream flow across sites).

We assessed and eliminated multi-collinearity using variance inflation factors (VIF)
following Meyers (1990) (see Text S1). Our environmental variables representing leaf litter and
the presence of riparian trees were collinear and leaf litter was removed from all analyses in

192 favour of presence of trees, which was likely to control the distribution of leaf litter, as well as 193 the availability of greenfall (freshly fallen moist green leaves). Distance to water and type of 194 water were also collinear. Rather than remove one of these variables from the analyses, we used 195 principal components analysis (PCA) in R v. 2.9.0 to combine these two factors into a single 196 factor representing river resources more generally (PC1). However, to improve interpretability, 197 we used the normalized inverse of the original values (max value - actual value) of distance to 198 water and created numerical categories of water type with dry sites equal to 1, artificial pools 199 equal to 2, natural pools equal to 3, and stream sites equal to 4. Thus, in general, most types of 200 river resources (i.e. drinking water and emergent insects) should increase with increasing values 201 of PC1.

ANALYSING MULTIVARIATE RESPONSES—We tested for differences in community composition between dry and flowing sites, as well as for associations between environmental factors and community composition using non-parametric permutational multivariate ANOVA (*adonis*) in the VEGAN package of R v. 2.9.0 (4999 permutations; see Text S1 for additional details). Data were natural log-transformed prior to all multivariate analyses.

207 To display differences in assemblage composition between flowing and dry sites, or 208 associations of assemblages with environmental variables, we used non-metric multidimensional 209 scaling (metaMDS) with the envfit vector plotting function in the VEGAN package of R v. 2.9.0. 210 For environmental association graphs, we calculated proximity to channel bank instead of 211 distance by taking the normalized inverse of the original values (max value – actual value). This 212 approach improved simplicity of interpretation of graphs, by making the direction of increase 213 correspond to increasing proximity to the channel bank (increasing proximity to leaf litter, 214 vegetation, etc). Thus taxonomic groups aligned along an environmental axis could be

interpreted as associated with this factor more clearly. All multivariate community compositiontechniques employed Bray-Curtis distances.

217 ANALYSING UNIVARIATE RESPONSES—Next, we tested for differences in total abundance, 218 diversity, the abundance of predators, the abundance of consumers known to eat at least some 219 animal material and the abundance of key taxa, between flowing and dry sites, using general 220 linear modelling (glm) in R v.2.9.0. We then performed similar analyses examining associations 221 between environmental factors (Table 1) and these same responses. We employed Shannon's 222 diversity and Pielou's evenness, in addition to richness, as our estimates of diversity. Pielou's 223 evenness is a good measure for our type of dataset and study question because it is relatively 224 insensitive to the effects of rare taxa (Beisel et al., 2003). Tests of mean total abundance and 225 diversity assumed a Gaussian distribution, whereas tests of taxa or functional group abundance 226 assumed a quasi-Poisson distribution, since these datasets often contained a large number of 227 zeros and this distribution is better for modelling such datasets (Potts & Elith, 2006). Total 228 abundance data were natural log transformed prior to analysis, whereas taxon abundance data 229 were rounded to the nearest integer prior to analysis (for the use of a quasi-Poisson distribution). 230 In all tests, we evaluate differences assuming a Type-I error rate of  $\alpha = 0.1$  given our low sample 231 sizes and our goal of identifying patterns rather than testing causal relationships; however, we 232 report exact P values so that patterns at the more conservative Type-I error rate ( $\alpha = 0.05$ ) can be 233 easily assessed.

234 Spatial autocorrelation

Due to the varying distances between our sampling locations, spatial correlation could be an important source of variation in our study. Currently, no good methods exist for detecting and correcting for spatial autocorrelation with multivariate community data (Gilbert & Bennett,

238 2010). However, for significant univariate responses (diversity, abundance), we report results of

239 Moran's I test for spatial autocorrelation (using the ape package of R v. 2.9.0), thus detecting

spatial effects that could be mechanistically investigated by future studies.

241 **Results** 

#### 242 Total abundance and abundance of functional groups

243 We found significantly higher abundance of all known predators at flowing sites than at 244 dry sites (Table 2: Table S3) and predators were positively associated with river resources (i.e. 245 water and/or aquatic invertebrates) and negatively associated with distance to the nearest channel 246 bank (higher near channel bank; Table 3; Table S4). Significant spatial autocorrelation was 247 apparent for these relationships (Tables S5 and S6). There were no significant differences in the 248 total abundance of all arthropods, or the abundance of consumers known to consume at least 249 some animal material, nor did we find any association of predators with other environmental 250 variables (Table S3, Table S4).

251 Arthropod families

Shannon's diversity and familial richness were higher at flowing than dry sites (Table 2,
Table S3) and there was a significant positive association between familial richness and river
resources, but no other associations (Table 3, Table S4). These relationships did not show spatial
autocorrelation (Tables S5 and S6).

Assemblage composition also differed significantly between dry and flowing sites (F = 2.23, df = 1,16, P = 0.021; Fig 2; Table 2; Table S7) and there was a significant association between assemblage composition and river resources (Fig 3; Table 3; Table S7). These community differences seem to have been driven by higher abundances of field crickets (Gryllidae), wolf spiders (Lycosidae), rove beetles (Staphylinidae) and shore bugs (Saldidae) at flowing sites (Table 2, Table S8) and positive associations of shore bugs (Saldidae), pygmy mole
crickets (Tridactylidae), and pygmy grasshoppers (Tetrigidae) with river resources (Table 3,
Table S9).

264 In addition to associations with river drying and river resources, several taxa showed 265 significant associations with other environmental factors. Wolf spiders (Lycosidae) and shore 266 bugs (Saldidae) were more abundant nearer to the channel bank, but pygmy grasshoppers 267 (Tetrigidae) were more abundant farther from the channel bank (Table 3, Table S9). Both 268 pygmy mole crickets (Tridactylidae) and pygmy grasshoppers (Tetrigidae) were more abundant 269 with less herbaceous ground cover (Table 3, Table S9), but pygmy grasshoppers were also more 270 abundant where cottonwood and willow trees were in close proximity. Pygmy grasshoppers 271 were the only family to respond to substrate complexity, with greater abundance at intermediate 272 complexity (e.g. coble bars with small to medium sized rocks and sand or gravel, see Table 1; 273 Table 3; Table S9). There were no significant associations between environmental factors and 274 the abundance of Gryllidae, Carabidae, Formicidae, Elateridae, Noctuidae, Linyphiidae, 275 Anthicidae, Staphylinidae and Rhaphidophoridae (Table S9). Only the relationship between 276 wolf spiders (Lycosidae) and distance to the channel bank showed significant spatial 277 autocorrelation (Table S6).

278 Carabid genera

Similarly to the results for the diversity of arthropod families, Shannon's diversity and
the richness of carabid beetle genera were higher at flowing sites than at dry sites (Table 2, Table
S11) and there was a positive association between river resources and generic richness.
Differing from the family-level results, Shannon's diversity and Pielou's evenness of carabid

283 genera were additionally positively related to river resources. Shannon's diversity and generic

richness were also higher near the channel bank (Table 3, Table S12). All of these relationships 284 285 showed significant spatial influence except for the difference in Shannon's diversity between dry 286 and flowing sites and the association between Pielou's evenness and water resources. 287 Like for the patterns of familial assemblage composition (and carabid diversity), we 288 found a significant difference in carabid assemblage composition between dry and flowing sites 289 (F = 33.78, df = 1, 16, P = 0.000; Fig 4; Table 3; Table S10) and significant associations between 290 carabid assemblages and river resources (Fig 5, Table 3, Table S10). River related differences in 291 carabid assemblages were likely driven by higher abundances of the carabid beetle genera 292 Brachinus, Agonum, Lachnophorus, Chlaenius and Bembidion at flowing than dry sites (Table 2, 293 Table S13), along with positive associations of these genera with river resources (Table 3, Table 294 S14). Lower abundance of the genus Syntomus at flowing sites (Table 2, Table S13) and 295 negative associations of this genus with river resources (Table 3, Table S14) also contributed to 296 differences in carabid assemblages. Differences in Lachnophorus, Bembidion and Syntomus 297 between dry and flowing sites showed significant spatial autocorrelation, as did associations of 298 Brachinus, Agonum, Lachnophorus, Chlaenius, Bembidion and Syntomus with river resources. 299 Differing from the results for familial assemblage composition, carabid genera 300 assemblage composition was found to be additionally associated with distance from the channel 301 bank and substratum complexity (Fig 5, Table 3, Table S10). The significant influence of the 302 channel bank on composition was likely driven by higher abundances of *Brachinus*, *Agonum*, 303 Lachnophorus, Chlaenius, Bembidion and Schizogenius near the channel bank (Table 3, Table 304 S14). Additionally, the higher abundance of *Lachnophorous* at low to intermediate substratum

306 relationship between this factor and carabid assemblage composition. Some carabid genera were

complexity (e.g. sand or gravel and small rocks, see Table 1; Table S14) likely contributed to the

305

307 related to environmental variables in ways that did not manifest in assemblage differences.

308 *Schizogenius, Chlaenius* and *Agonum* were found to be negatively associated with percent

309 herbaceous ground cover (Table 3, Table S14) and *Schizogenius* was higher without cottonwood

310 or willow trees present (Table 3, Table S14). All of these relationships showed significant

311 spatial influence (Table S6).

#### 312 **Discussion**

313 Recent global changes are drastically altering the distribution of water resources. One 314 result has been dewatering of rivers around the world (Gleick, 2003; Pearce, 2006). Little 315 information is available about how these drying events influence riparian animal communities. 316 Here we show that river drying is associated with alterations of riparian arthropod assemblage 317 composition, a decline in taxon diversity and reductions in the abundance of several taxa. 318 Further, direct access to river-related resources (e.g. water, aquatic food) may help explain these 319 patterns, since this factor is more strongly correlated with differences in these arthropod 320 communities than other habitat characteristics. These associations are apparent for riparian 321 arthropod families and appear strong when focussing on genera within the ground beetle family 322 (Carabidae). Thus, this study suggests that many desert riparian arthropods are sensitive to river 323 drying events and changes in water availability and thus may require the existence of at least 324 some perennial flows within a river network for persistence.

While we cannot separate whether drinking water or emergent insects were more important in driving the response of arthropods to river resources, insect emergence from rivers in this region has been found to be relatively low at the end of June, with peak emergence generally occurring earlier in the year (Hagen, 2010). This suggests water may be directly important as a resource for this arthropod community in June when dry conditions prevail, in

addition to the important effects of emergence of aquatic insects documented in other systems
(e.g. Paetzold, Bernet & Tockner, 2006) and likely still important here. For example, pygmy
grasshoppers in the family Tetrigidae are known to eat mostly moist, recently stranded algae
(Bastow *et al.*, 2002) and bombardier beetles in the carabid genus *Brachinus* are known to have
ectoparasitic larvae of aquatic diving beetles in the family Dytiscidae (Juliano, 1985). Both of
these taxa were positively associated with river resources in our study. These examples highlight
the important roles rivers play as both water and food resources for riparian arthropods.

#### 337 Desert riparian zones harbour cosmopolitan species

338 Drying may be particularly common along desert rivers (Kingsford, 2006). Many species 339 living in desert uplands (Noy-Meir, 1974; Polis & Seely, 1990; Davis & DeNardo, 2007) or 340 within rivers (Meffe & Minckley, 1987; Beche, McElravy & Resh, 2006; Kingsford, 2006) have 341 traits that help them cope with the challenges of highly variable desert environments. Therefore, 342 one might be tempted to predict that desert riparian zones contain species that are well adapted 343 and relatively insensitive to drying events. However, many of the riparian arthropods in our 344 study are distributed along rivers throughout North America. For instance, the most common 345 large spider collected in our study, the beach wolf spider, Arctosa litorallis Hentz, is widely 346 distributed across North America (Ubick et al., 2005; Punzo, 2006). Most of the carabid beetle 347 genera we found are also widely distributed throughout North America, although individual 348 species within these genera may or may not be widely distributed. For instance, bombardier 349 beetles (Carabidae: Brachinus) and the genus Syntomus, which contains only one species in 350 North America, Syntomus americanus Dejean, are widely distributed across the continent (Arnett 351 & Thomas, 2000). Thus, desert riparian zones often harbour a cosmopolitan fauna dominated by 352 taxonomic groups also found in more mesic environments. This result is consistent with

353 previous evidence that riparian zones contain different species than uplands (Sabo et al., 2005b). 354 In fact, floodplain forests along desert rivers may act as oases, showing more mesic 355 environmental characteristics than do adjacent uplands (Skagen et al., 1998; Sabo et al., 2008). 356 The mixture of widely and locally distributed taxa among those collected, suggests the 357 intriguing hypothesis that the relative endemism of each taxonomic group may be a good 358 predictor of the degree to which it is influenced by river drying or to which it is associated with 359 measures of water availability, with more endemic species being less influenced by drying. 360 However, our data do not support this hypothesis. For example, the beach wolf spider 361 (Lycosidae: A. littoralis Hentz), which made up most of the individuals collected in the family 362 Lycosidae, is widespread (Punzo, 2006), but the family Lycosidae did not show a significant 363 association with river resources (though abundance was higher overall at flowing sites). 364 Similarly, the carabid beetle, S. americanus Dejean, is found throughout North America (Arnett 365 & Thomas, 2000), but its abundance was higher at dry sites and farther from water bodies. On 366 the other hand, the carabid beetle *Lachnophorus elegantulus* Dejean (the only representative of 367 this genus in this area) is most commonly found in the southwestern US (Arnett & Thomas, 368 2000), but its abundance was lower at dry locations and farther from water bodies. Therefore, 369 the responses of each taxonomic group to river drying and the association with environmental 370 characteristics are not strictly predictable by regional distribution. This suggests a) some species 371 most abundant in desert regions are still sensitive to river drying (e.g. *L. elegantulus* Dejean), 372 particularly along historically perennial rivers like the San Pedro and b) that species interactions 373 (e.g. McCluney & Sabo, 2009) or life history constraints (e.g. Juliano, 1985) may be modulating 374 the observed response to river drying.

375 Floodplain versus dry channel habitats

376 Generally, we found that predator abundance, some metrics of diversity, and abundance 377 of some families and carabid genera were higher nearer the channel bank (Table 3). This 378 suggests an important positive aspect of the transition zone between floodplain and channel, such 379 as the availability of increased structure or vegetation, the overlap of species between these two 380 habitat types, or decreased danger of flow-related disturbance. However, we observed some taxa 381 that were more abundant farther from the channel bank, with less herbaceous cover, or without 382 cottonwood trees nearby. While these preferences for more open environments may be due to 383 energy balance requirements or predator avoidance (e.g. wolf spiders were higher near channel 384 banks), the preference may also reflect our choice of study location. Our research focussed on 385 the river channel itself and did not sample the wide floodplains that occur along this river. These 386 floodplains probably harbour a different suite of arthropods that may be better suited to habitats 387 with leaf litter, herbaceous vegetation and shade. For instance, the large wolf spider A. littoralis 388 Hentz, is abundant in the river channel along the San Pedro, but is rarely found in floodplain 389 habitats, which instead are dominated by the wolf spider Hogna antelucana Montgomery, which 390 in turn is rarely found in the river channel (K. McCluney and J. Sabo, personal observations). 391 Spatial autocorrelation

Significant spatial autocorrelation was observed for some of our responses, but not others, with spatial effects more commonly observed for the abundance of particular taxa and less commonly for patterns of diversity and predator abundance (Table S5 and S6). In some cases, local migration may cause spatial autocorrelation and may explain patterns instead of the relevant site characteristics. However, we note that, in comparisons between dry and flowing sites, all four flowing sites were clumped together and separated from dry sites (Fig 1). Thus, significant spatial autocorrelation would be likely to occur, even if differences in responses were

399 completely associated with river drying. However, our analysis was concerned with identifying 400 patterns of variation of riparian arthropods along this drying river rather than testing for causal 401 relationships. A related study that manipulated water resources along this river segment as it 402 dried, and avoided these spatial issues, found similar patterns for diversity, composition, and 403 biomass of some of these taxa (McCluney, 2010).

404 Conclusion

405 Overall, we found evidence that short-term river drying can substantially influence 406 riparian arthropod communities and that direct access to the resources associated with river water 407 may be an important part of the effect. In general, we found a decrease in diversity and in 408 abundance of many taxa associated with river drying, with an increase in only one carabid beetle 409 genus. Riparian arthropods make up an important component of the diet of higher consumers, 410 such as birds, skunks and foxes (Sabo, Soykan & Keller, 2005a; Soykan, 2007). Thus, they are 411 important in the conservation of the entire suite of organisms living along the river. Achieving a 412 balance between human and non-human water needs in this region and other similar regions will 413 require an understanding of the connections between groundwater pumping and in-stream and 414 riparian ecology. Our study helps provide some of the first information linking groundwater 415 withdrawals to the invertebrate resource base of riparian animal communities.

417	
418	Acknowledgments
419	We wish to thank S. Anderson of Gray Hawk Nature Center for accommodation and
420	support and the US Bureau of Land Management for permission to work on lands under their
421	management. We thank Y. Marusenko, L. Thompson, C. Soykan and N. McCluney for
422	assistance in the field and laboratory and M. Tseng for assistance with identification. We thank
423	J. Oksanen for statistical advice on spatial autocorrelation and two anonymous reviewers and A.
424	Hildrew for helpful comments and editing. This study was funded by EPA STAR grant FP-
425	91649201 to K. McCluney and the Brian Daniel Corrigan Foundation.

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- 588
- 589
- 590

#### 591 Tables

### 592

TABLE 1. Description of each environmental factor measured.

Factor	Description		
Substratum Complexity	Score out of 5: Sand/Clay/Silt = 1, Gravel = 2, Rocks<10cm = 3, Rocks 10		
	40  cm = 4, Rocks> $40  cm = 5$ (If the rocks were in clay, it lowered the score		
	by 1)		
Distance to Water	Distance between the trap and the nearest water source (up to 55 m)		
Type of Water	Classification into categories of flowing stream, natural pool, artificial poo		
	dry site.		
Water Resources	Principal component 1 from principal components analysis (PCA) of		
	distance to water and type of water. However, original factors were		
	converted so that water resources increase with increasing values of PC1		
	(see Methods).		
Distance to Channel Bank	Distance between the trap and the nearest channel bank. Proximity to		
	channel bank was calculated for interpretability in graphs (see Methods).		
Percent Herbaceous Cover	Visual estimates of herbaceous ground cover within 1 m diameter of each		
	trap, divided into 5% categories		
Litter	Not examined due to collinearity with other factors.		
Presence of Cottonwood or	Whether or not cottonwood or willow trees were found near the trap (clos		
Willow Trees	enough for leaves to commonly fall near the trap)		

5	0	7	
J	7	/	

TABLE 2. Significant results for differences between flowing and dry sites on 25 June 2006. Grey highlighting indicates significance only at  $\alpha = 0.1$ , while all other reported metrics are significant at  $\alpha = 0.05$ .

	Type of site with higher			
Metric	level of metric	F	df	р
Abundance of known predators	Flowing	24.352	1, 16	0.000
Families				
Community composition	NA	2.234	1,16	0.021
Shannon's diversity	Flowing	5.350	1, 16	0.034
Richness	Flowing	5.983	1, 16	0.026
Gryllidae abundance	Flowing	5.062	1,16	0.039
Lycosidae abundance	Flowing	10.023	1, 16	0.005
Staphylinidae abundance	Flowing	4.780	1, 16	0.044
Saldidae abundance	Flowing	23.822	1, 16	0.000
Carabid Genera				
Community composition	NA	33.777	1, 16	0.000
Shannon's diversity	Flowing	12.0829	1, 16	0.003
Richness	Flowing	27.346	1, 16	0.000
Brachinus abundance	Flowing	8.042	1, 16	0.012
Agonum abundance	Flowing	48.13	1, 16	0.000
Lachnophorus abundance	Flowing	48.13	1, 16	0.000
Bembidion abundance	Flowing	8.033	1,16	0.012
Syntomus abundance	Dry	17.418	1,16	0.001
Chlaenius abundance	Flowing	4.197	1,16	0.057

598

600

TABLE 3. Significant associations between environmental factors and response variables on 25 June 2006. Grey highlighting indicates significance only at  $\alpha = 0.1$ , while all other reported metrics are significant at  $\alpha = 0.05$ . For F and df see Tables S4, S7, S9, S10, S12 and S14.

<b>Response Metric</b>	<b>Environmental Factor</b>	<b>Relationship direction</b>	р
Abundance of known predators	Water Resources	Positive	0.074
Abundance of known predators	Distance to Channel Bank	Negative	0.011
Families			
Community composition	Water Resources	NA	0.020
Richness	Water Resources	Positive	0.090
Lycosidae abundance	Distance to Channel Bank	Negative	0.025
Saldidae abundance	Water Resources	Positive	0.003
Saldidae abundance	Distance to Channel Bank	Negative	0.005
Tridactylidae abundance	Water Resources	Positive	0.034
Tridactylidae abundance	Percent Herbaceous Cover	Negative	0.036
Tetrigidae abundance	Water Resources	Positive	0.000
Tetrigidae abundance	Distance to Channel Bank	Positive	0.003
Tetrigidae abundance	Percent Herbaceous Cover	Negative	0.001
Tetrigidae abundance	Presence of Trees	Positive	0.046
Tetrigidae abundance	Substrate Complexity	Intermediate	0.020
Carabid Genera			
Community composition	Water Resources	NA	0.000
Community composition	Distance to Channel Bank	NA	0.072
Community composition	Substrate Complexity	NA	0.002
Shannon's diversity	Water Resources	Positive	0.000
Shannon's diversity	Distance to Channel Bank	Negative	0.007
Richness	Water Resources	Positive	0.000
Richness	Distance to Channel Bank	Negative	0.000
Pielou's Evenness	Water Resources	Positive	0.014
Brachinus abundance	Water Resources	Positive	0.004
Brachinus abundance	Distance to Channel Bank	Negative	0.077
Agonum abundance	Water Resources	Positive	0.000
Agonum abundance	Distance to Channel Bank	Negative	0.000
Agonum abundance	Percent Herbaceous Cover	Negative	0.000
Lachnophorus abundance	Water Resources	Positive	0.000

Lachnophorus abundance	Distance to Channel Bank	Negative	0.000
Lachnophorus abundance	Substratum Complexity	Intermediate	0.095
Bembidion abundance	Water Resources	Positive	0.006
Bembidion abundance	Distance to Channel Bank	Negative	0.054
Syntomus abundance	Water Resources	Negative	0.075
Chlaenius abundance	Water Resources	Positive	0.001
Chlaenius abundance	Distance to Channel Bank	Negative	0.000
Chlaenius abundance	Percent Herbaceous Cover	Negative	0.027
Schizogenius abundance	Distance to Channel Bank	Negative	0.063
Schizogenius abundance	Percent Herbaceous Cover	Negative	0.001
Schizogenius abundance	Presence of Trees	Negative	0.002

## 603 Figure Legends

- FIG 1. Maps of the region of Southeastern AZ, USA where the study took place and the
- 605 distribution of sampling sites along the river on the final sampling date. The direction of flow is
- from south to north. Sampling regions are noted by broader lines following the river course,
- 607 with narrow solid lines indicating dry sites, broad dashed lines indicating a mix of dry sites,
- 608 natural pools, and artificial pools, and broad solid lines indicating flowing stream sites. The
- 609 Sierra Vista metropolitan area, a region of groundwater withdrawal, is just upstream of the
- 610 sampling locations.

- 612 FIG 2. Non-metric multi-dimensional scaling plot of the difference in familial assemblage
- 613 composition between flowing and dry sites (D = Dry, S = Flowing Stream). Grey text and
- 614 symbols refer to each family in the analysis. Only families with greater than one individual
- 615 across all sites are labelled.
- 616

617	FIG 3. Non-metric multi-dimensional scaling plot showing the association between each
618	environmental factor and familial assemblage composition. The rays show the direction and
619	importance of variation of each factor along the first two axes (WaterR = Water Resources,
620	ProxBank = Proximity to the Channel Bank, Pcov = Percent Herbaceous Ground Cover,
621	SubstrScore = Structural Complexity of the Substratum; the variable indicating the presence of
622	trees near the trap is not shown since the labels overlapped with the origin and were
623	uninformative). Grey text and symbols refer to each family in the analysis. See Table S7 for $R^2$
624	values for each factor. Only families with greater than one individual across all sites are labelled.
625	

- 626 FIG 4. Non-metric multi-dimensional scaling plot of the difference in carabid genera assemblage
- 627 composition between flowing and dry sites (D = Dry, S = Flowing Stream). Grey text refers to
- 628 each genus in the analysis. Only genera with greater than one individual across all sites are
- 629 labelled.
- 630

631 FIG 5. Non-metric multi-dimensional scaling plots showing the association between each

632 environmental factor and carabid genera assemblage composition. The rays show the direction

and importance of variation of each factor along the first two axes (WaterR = Water Resources,

634 ProxBank = Proximity to the Channel Bank, Pcov = Percent Herbaceous Ground Cover,

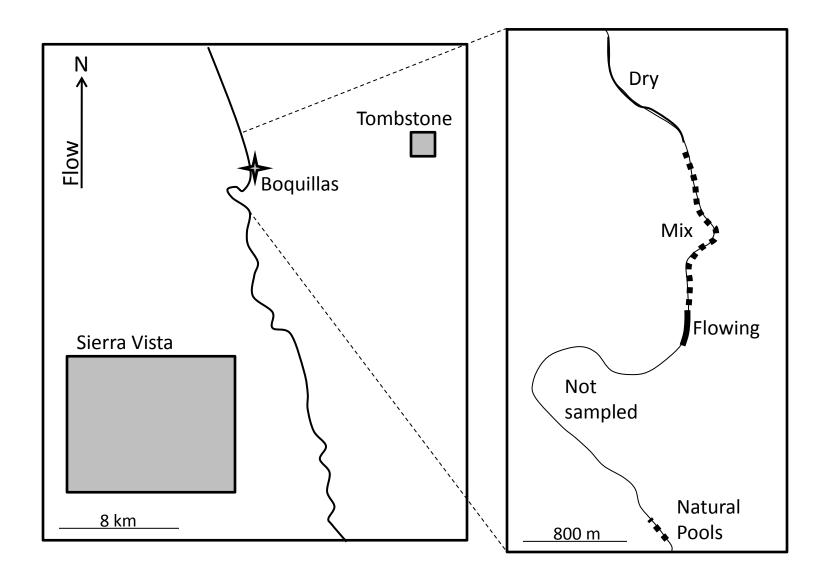
635 SubstrScore = Structural Complexity of the Substratum, CWoWNear = Presence of Cottonwood

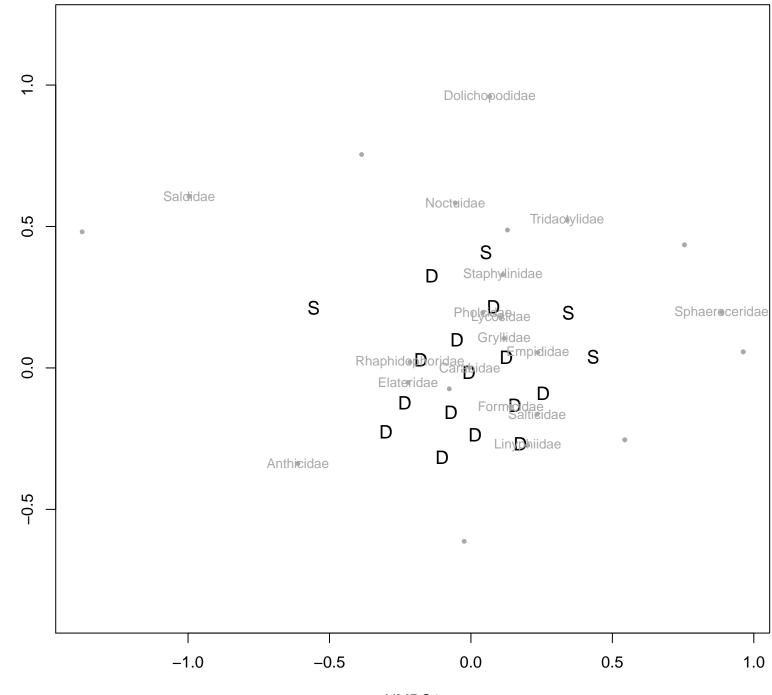
636 or Willow Trees Near the Trap, with the suffix -Y indicating yes, -N indicating no, and no suffix

637 representing a single site with missing information). Grey text refers to each genus in the

analysis. See Table S10 for  $R^2$  values for each factor. Only families with greater than one

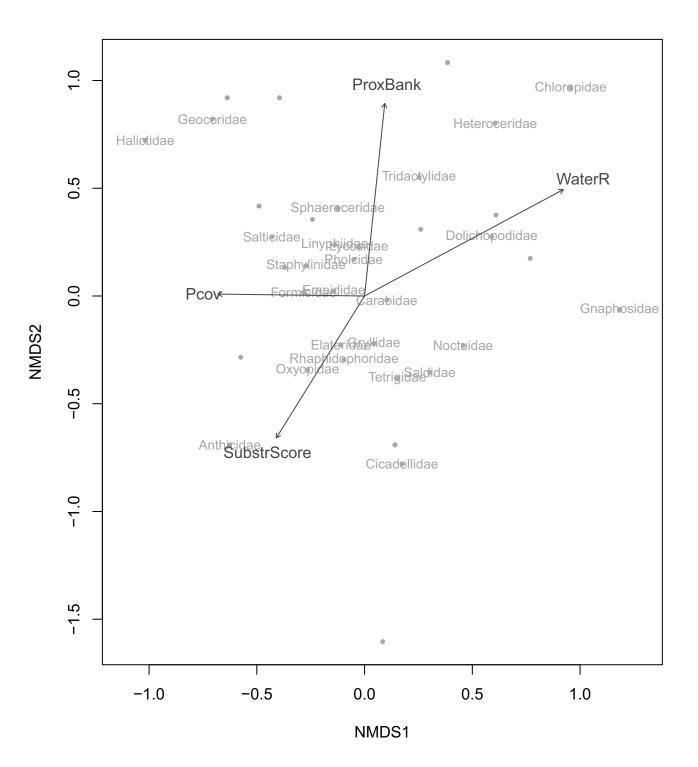
639 individual across all sites are labelled.

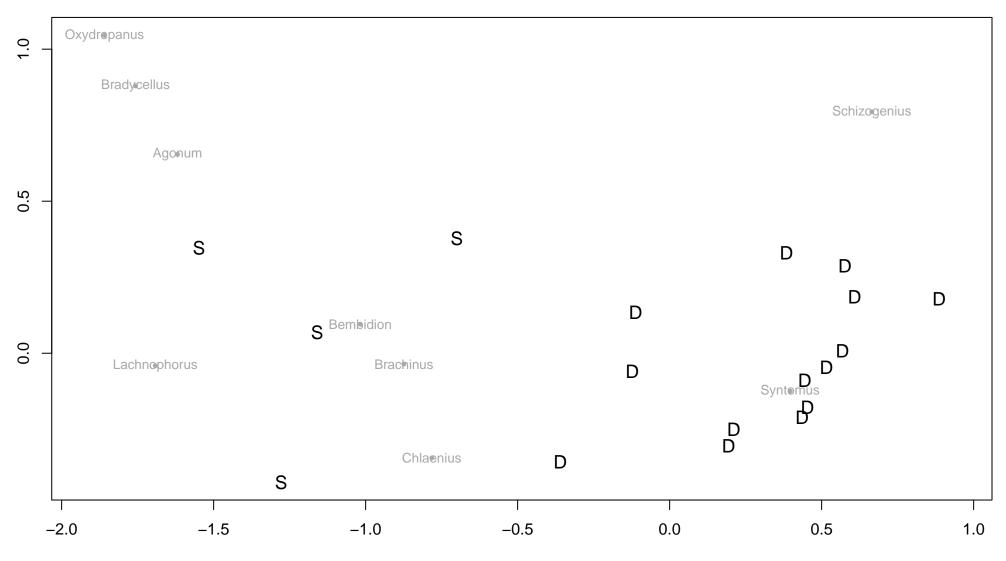




NMDS2







NMDS1

NMDS2

