Physiological responses to drought stress and recovery reflect differences in leaf function and microanatomy among grass lineages

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Abstract

Grasses are cosmopolitan, existing in many biome and climate types from xeric to tropical. Traits that control physiological responses to drought vary strongly among grass lineages, suggesting that tolerance strategies may differ with evolutionary history. Here, we withheld water from 12 species representing 6 tribes of grasses to compare how tolerant and intolerant species respond to drought in different grass lineages. We measured physiological, morphological, and microanatomical traits. Dominant lineages from tropical savannas, like Andropogoneae, tolerated drought due to above and belowground morphological traits, while temperate grasses utilized conservative leaf physiology (gas exchange) and microanatomy. Increased intrinsic water-use efficiency (iWUE) coincided with a larger number of stomata, resulting in greater water loss (with inherently greater carbon gain) and increased drought sensitivity. Inherent leaf and root economic strategies impacting drought response were observed in all species, resulting in either high SLA or SRL, but not both. Our results indicate that grasses subjected to severe drought were influenced by microanatomical traits (e.g., number of stomata and xylem area) which were shared within lineages. In addition, grasses recovered at least 50% of physiological functioning across all lineages and 92% within Andropogoneae species, illustrating how drought can influence functional responses across diverse grass lineages.









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16 ABSTRACT

Grasses are cosmopolitan, existing in many biome and climate types from xeric to tropical. Traits 17 that control physiological responses to drought vary strongly among grass lineages, suggesting 18 that tolerance strategies may differ with evolutionary history. Here, we withheld water from 12 19 species representing 6 tribes of grasses to compare how tolerant and intolerant species respond to 20 drought in different grass lineages. We measured physiological, morphological, and 21 microanatomical traits. Dominant lineages from tropical savannas, like Andropogoneae, tolerated 22 drought due to above and belowground morphological traits, while temperate grasses utilized 23 conservative leaf physiology (gas exchange) and microanatomy. Increased intrinsic water-use 24 efficiency (*iWUE*) coincided with a larger number of stomata, resulting in greater water loss 25 (with inherently greater carbon gain) and increased drought sensitivity. Inherent leaf and root 26 27 economic strategies impacting drought response were observed in all species, resulting in either high SLA or SRL, but not both. Our results indicate that grasses subjected to severe drought were 28 influenced by microanatomical traits (e.g., number of stomata and xylem area) which were 29 shared within lineages. In addition, grasses recovered at least 50% of physiological functioning 30 31 across all lineages and 92% within Andropogoneae species, illustrating how drought can influence functional responses across diverse grass lineages. 32 33

Key words: Poaceae, Drought Response, Phylogeny, Plant Functional Traits, Microanatomy,
 Ecophysiology, Leaf Economic Spectrum.

36 INTRODUCTION

Grasslands play a major role in regional carbon sequestration and water cycling because 37 grasses invest in extensive rooting systems and storage organs (Pendall et al. 2018; Veldman et 38 al. 2019). Carbon dynamics are highly influenced by water availability in grassland systems, 39 evident in drought years that result in decreased productivity (Fay, Carlisle, Knapp, Blair & 40 Collins 2003; Hoover & Rogers 2016; Carroll et al. 2021). Grasslands experiencing extreme 41 droughts can have reduced physiological functioning (Cook, Ault & Smerdon 2015; Hoover, 42 43 Duniway & Belnap 2015), increased invasibility from non-native species (Linder, Lehmann, Archibald, Osborne & Richardson 2018), disruption of fire intervals (Wilcox et al. 2020), and 44 loss of ecosystem functioning (i.e. productivity & species composition) (Eters, Tarks & 45 Ernandez 2014; Mainali et al. 2014; Knapp et al. 2020). While many grass species in grassland 46 47 ecosystems have evolved in the context of an inherently variable climate, future climate projections emphasize large shifts in water availability, resulting in extreme drought and deluge 48 49 events within the coming century (Stocker et al. 2013; Cook et al. 2015; Griffin-Nolan et al. 2019; Post & Knapp 2019; Knapp et al. 2020). While it is widely accepted that grasslands will 50 51 vary in drought response (ability to withstand shifts from equilibrium) and drought recovery (ability to regain equilibrium), modifications in precipitation seasonality and amount will have 52 53 sizable and diverse impacts on ecosystem function (Hoover, Knapp & Smith 2014; Volaire 2018; Maurer, Hallmark, Brown, Sala & Collins 2020). 54

55 Biophysical factors determining drought sensitivity in individual plant species include 56 precipitation and temperature variability (Knapp et al. 2015), while biotic factors such as plant productivity, species richness (Burri, Niklaus, Grassow, Buchmann & Kahmen 2018), and 57 potentially dominant species with associated functional traits, also play an important role (Avolio 58 et al. 2019). Furthermore, the history of different drought exposure in plant lineages is likely to 59 frame future drought responses within those lineages. For example, lineages of plant species 60 from arid and semi-arid regions have functional traits (narrow leaves, strict stomatal regulation, 61 absorptive rooting systems) that allow them to acquire and conserve water (Ocheltree et al. 62 2020), whereas lineages from tropical regions may have wider leaves and altered stomatal traits 63 that result in distinct water-use strategies (Liu et al. 2018; Buckley 2019). These evolutionary 64 tradeoffs have shaped functional differences across lineages and directly impact ecological 65 66 dynamics (Griffith et al. 2020). However, the extent of such evolutionary tradeoffs has not been

utilized to identify lineage-specific trait responses to extreme drought conditions. Even more
uncommon are investigations that combine physiology, microanatomy, morphology, and
structural data from grass species spanning several Poaceae tribes.

70 Large interannual variation in precipitation is a feature of many grassland ecosystems and, in combination with CO_2 and temperature, has played a major role in the evolution and 71 biogeographic history of major grass lineages (Osborne 2008; Cleland et al. 2013; Cotton, 72 73 Cerling, Hoppe, Mosier & Still 2016). Importantly, the varying evolutionary histories of 74 grasslands have driven the evolution of different functional traits across the Poaceae phylogeny, likely accounting for differences in drought responses (Ocheltree et al. 2020; Knapp et al. 2020). 75 For example, leaf-level microanatomical trait variation and convergent evolution has resulted in 76 spatially separated photosynthetic tissues allowing for C₄ photosynthesis, which is heavily 77 78 expressed in Poaceae, and provides a physiological advantage that increases carbon assimilation 79 while reducing water loss via stomatal regulation (Taylor et al. 2010; Zhou, Helliker, Huber, 80 Dicks & Akcay 2018). While it is recognized that C_4 species are not inherently more drought tolerant than C₃ species (Ehleringer 2005; Nippert, Fay & Knapp 2007; Knapp et al. 2020), there 81 82 is evidence that increased WUE (water-use efficiency), inherent to C_4 species, can be advantageous when water is limiting (Lambert, Baer & Gibson 2011; Kimball, Gremer, Angert, 83 Huxman & Venable 2012; Leakey et al. 2019). For example, native species in the arid American 84 southwest, have the ability to initially tolerate the negative consequences of drought by 85 86 maintaining physiological functioning for prolonged periods of time (Thomey, Collins, Friggens, 87 Brown & Pockman 2014; Skelton, West & Dawson 2015). The ability of some species to maintain physiological functioning despite drying soils may be due to increased cuticle 88 thickness, decreased stomatal size and densities, less negative turgor loss point, and more 89 conservative growth strategies (specific leaf area, SLA; specific root length, SRL) (Reich 2014; 90 Habermann et al. 2019; Bertolino, Caine & Gray 2019; Ocheltree et al. 2020). Alternatively, the 91 production of cheaper leaves and roots (higher carbon to nitrogen ratio) and tight stomatal 92 regulation is associated with the ability to avoid desiccation and quickly recover once drought 93 breaks (Poorter, Niinemets, Poorter, Wright & Villar 2009; Lin et al. 2015; Garbowski et al. 94 2020). The ability to quickly resume pre-drought physiological function via rapid recovery may 95 or may not be associated with the ability to tolerate drought in the first place (Hoover et al. 2014; 96 97 Volaire 2018).

During the evolutionary development of Poaceae, separate lineages have evolved 98 different suites of traits, including fairly different water use strategies (Osborne 2008; Edwards, 99 Osborne, Stromberg & Smith 2010). For instance, the two most abundant monophyletic groups 100 of C₄ grasses - Andropogoneae (water spenders) and Chloridoideae (water savers) - vary in 101 water-use strategies because of distinct biogeographic histories (Taub 2000; Grass Phylogeny 102 103 Working II 2012; Griffith et al. 2020). Species in these lineages occupy warmer climates but 104 vary in global distribution as a function of precipitation availability: high in Andropogoneae and 105 low in Cynodonteae (Williams, Wilsey, Mcnaughton & Banyikwa 1998; Liu & Osborne 2015; Lehmann et al. 2019). There are many characteristics impacting water-use and drought response 106 associated with this ecological sorting, and they include morphological, physiological, and 107 anatomical traits. Morphological strategies and traits associated with water relations include the 108 109 production of fine roots to increase water absorption (McCormack et al. 2015; Roumet et al. 2016; Iversen et al. 2017), leaf rolling to decrease irradiance (Cardoso, Pineda, Jiménez, Vergara 110 111 & Rao 2015), and variations in growth form (caespitose and rhizomatous) (Blair, Nippert & Briggs 2014; Ott & Hartnett 2015). These traits are often related in terms of economics, 112 113 reflecting plant investment of carbon and nitrogen in both leaf and root structures (Pérez-Harguindeguy et al. 2013). More specifically, these morphological traits are framed by 114 underlying structures at the microanatomical level in leaf and root tissues (John et al. 2017). 115 Microanatomical leaf traits within and across families in Poaceae also have been observed to 116 117 influence physiological responses most often associated with hydraulics (xylem area/diameter; 118 resistance to cavitation) (Hacke, Sperry, Pockman, Davis & McCulloh 2001; Bachle & Nippert 2018, 2021). However, the aforementioned physiological, morphological, and anatomical traits 119 may not convey equal benefits in drought response or recovery across and within Poaceae 120 lineages. For these reasons, it is increasingly important to understand how diverse lineages of 121 grass species that vary in climate niches and evolutionary histories will respond to extreme 122 drought conditions 123

Here, we conducted a robust assessment of physiological and anatomical traits from multiple grass lineages in response to and following recovery from drought. The species under investigation were selected based on divergent drought responses within lineages. We performed a dry-down experiment to impose severe drought on 12 species of grasses across 6 tribes within the Poaceae lineage. We withheld water in order to assess various physiological, morphological,

and microanatomical trait responses to drought, as well as (above and belowground) productivity 129 data, to capture both drought response and recovery. We hypothesized that: (1) species within 130 tribes will exhibit a similar response to drought sensitivity (duration in drought), based on similar 131 evolutionary histories and drought traits specific to withstanding long periods of low water 132 availability; (2) species within tribes will also exhibit similar responses in drought recovery, 133 based on shared evolutionary histories and functional traits that serve to quickly utilize resources 134 when available; and (3) leaf-level microanatomical traits would best describe species (within and 135 136 across tribes) response to, and recovery from drought due to the constraints of structures that influence water transport and availability. 137

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139 MATERIALS AND METHODS

140 Twelve grass species from six tribes were grown from seeds obtained from the USDA Germplasm Resources Information Network or locally sourced from the Konza Prairie 141 142 Biological Station. Species include: Paspalum juergensii, Paspalum notatum, Festuca ovina, Panicum virgatum, Setaria viridis, Urochloa ruziziensis, Andropogon gerardii, Sorghastrum 143 144 nutans, Danthonia spicata, Rytidosperma semiannulare, Bouteloua dactyloides, and Bouteloua gracilis (accession information in Supplemental table 1). Species were selected to represent 145 146 different major lineages of the family Poaceae (Cynodonteae, Andropogoneae, Paniceae, Danthonieae, Poeae, and Paspaleae), and included both C₃ (BEP and PACMAD clades) and a 147 148 range of C₄ species. In addition, we intentionally chose species (within the same tribe) that were 149 previously reported to have varying responses (tolerant and sensitive) to low soil moisture. Seeds were germinated in 868.5 cm³ size pots with a mix of potting soil and general-purpose sand with 150 a ratio of 2:1 soil to sand and placed in a Kansas State University greenhouse under ambient 151 conditions and raised to maturity throughout 2016 – 2018. Each pot was inoculated with a 152 153 handful of Konza Prairie soil. After reaching maturity, the samples were subjected to 100% water reduction (referred to 'dry-down'), simulating an extreme drought, as previously described 154 (Qiu, Bachle, Nippert & Ungerer 2020b; Qiu et al. 2020a). During the dry-down, samples were 155 156 monitored daily and placed into categorical conditions based on their physiological state: "Initial", "Stressed", and "Recovery". Physiological leaf traits were monitored daily and 157 included: leaf-level net photosynthetic rates (A_n ; μ mol m⁻² s⁻¹), stomatal conductance (g_s ; mol m⁻² 158 s⁻¹), transpiration (E: mmol m⁻² s⁻¹), and instantaneous water use efficiency (*iWUE*; A_n/E) 159

calculated as the ratio between A_n and E). Data was collected with a LI-6400 system (LiCOR 160 Biosciences Inc., Lincoln, NE, USA) equipped with an LED light source (light intensity 161 maintained at 2000 μ mol m⁻²s⁻¹) CO₂ concentration at 400 μ mol mol⁻¹, and relative humidity at 162 ambient levels (35-50%). Physiological states were determined by relative rates of A_n . The 163 condition: "Initial" was measured on Day 1 (first day of drought after being watered the previous 164 day) in order to avoid biased measurements from saturated soils. When samples reached near 165 stomatal closure and extremely low photosynthetic rates ($A_n < 25\%$) of Day 1 A_n ("Initial"), they 166 were categorized into the new condition "Stressed". At this point, water was re-applied to soil 167 saturation after the pertinent data were collected. Plants were allowed two days to recover before 168 post-drought physiological data was collected ("Recovery"). 169

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171 <u>Economic trait measurements</u>

After physiological data were collected in the "Recovery" period, above and 172 173 belowground tissues were harvested to determine productivity of all species and samples that were subjected to dry-down conditions. Leaf-level economic and microanatomical data were 174 175 collected from samples that included all non-droughted individuals but excluded P. juergensii, P. notatum, or F. ovina due to the lack of samples. The leaf tissue data included: Leaf area (LA; 176 cm²), specific leaf area (SLA, leaf area divided by dry mass; cm² g⁻¹), and leaf dry-matter content 177 (*LDMC*, fresh leaf mass divided by dry mass; $g g^{-1}$). SLA and *LDMC* were analyzed with the 178 179 standardized rehydration method (Garnier, Shipley, Roumet & Laurent 2001; Pérez-180 Harguindeguy et al. 2013), while LA data were obtained by processing images in ImageJ (Rasband 1997). Roots were washed and cleaned of debris for digital root imaging; analysis of 181 root images was completed with a root imaging software (WinRhizo; Regent Instruments, Inc., 182 Nepean, Ontario, Canada). Root imaging provided the following traits: total root length (cm), 183 root diameter (mm), and specific root length (SRL, root length divided by dry mass; cm g⁻¹). 184 After scans were completed, above and belowground biomass samples were dried for 48 hours at 185 65°C and weighed for productivity comparisons. 186

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188 <u>Microanatomy trait measurements</u>

The newest mature leaf was used for microanatomical analysis prior to the initiation of
drought from the following species: *Setaria viridis*, *Urochloa ruziziensis*, *Danthonia spicata*,

191 Rytidosperma semiannulare, Bouteloua dactyloides, and Bouteloua gracilis. Sorghastrum 192 nutans, Andropogon gerardii, and Panicum virgatum were collected from parent populations in the field at peak physiological performance. Festuca ovina, Paspalum notatum, and Paspalum 193 *juergensii* samples were not included in these analyses due to sample loss. Microanatomical 194 samples, roughly 30 mm in length, were collected (4 - 8 samples per species; n = 33) by clipping 195 leaf tissue and placing them into a fixative FAA (10% formalin / 5% glacial acetic acid / 50% 196 ethanol (use 95% EtOH) / 35% DI water) under a vacuum. Tissues were then cut (cross 197 198 sectioned) to 4µm in thickness with a Leica RM2135 microtome (Leica Biosystems, Newcastle, UK), and mounted in paraffin at Kansas State's College of Veterinary Medicine Histopathology 199 lab. Tissue was stained with Safranin-O and Fast Green (Ruzin 2000), cover slipped, and imaged 200 on a Zeiss 880 confocal microscope (Carl Zeiss, Walldorf, Germany) at 10X and 20X when 201 202 necessary with a multitrack configuration, digital dual-bypass filters and a GaAsP detector (Fig. 1). Microanatomical data were collected using IMAGEJ software (Rasband 1997) by analyzing 203 204 two tissue regions from either side of the midrib between two major vascular bundles which were then averaged together from each leaf sample (Bachle & Nippert 2018, 2021). Here, the 205 206 total subsampled area is referred to as the cross - sectional area or the area between two major vascular bundles (CSA). Microanatomical traits collected from subsamples include: xylem area 207 $(X_{area}; \mu m^{-2})$, xylem diameter $(X_{diameter}; \mu m)$, t/b (xylem wall thickness/ $X_{diameter}$; xylem resistance 208 to cavitation; $\mu m \mu m^{-1}$); while stomatal count (S_{count}) was collected from the whole-leaf cross 209 210 section. In this study, we did not collect stomatal densities - as that would entail epithelial peels 211 or impressions, therefore we do not equate density measurements and interpretations with the S_{count} . Instead, we utilized S_{count} to inform how many stomata are serving major and minor 212 vascular bundles within the whole-leaf cross section. 213

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215 <u>Data analysis</u>

The selected traits were averaged by species and separated into the three physiological 'stages' (Initial, Stressed, and Recovery) based on physiological responses. We included tribe as a factor to investigate differences among lineages. We considered using a phylogenetic generalized linear mixed model (PGLMM), however, our goal was to not control for phylogeny, but rather to determine if lineages with different traits have evolved different drought tolerance strategies. All data were checked to meet assumptions of normality before analyses began.

Comparisons among tribes and dry-down 'stages' were analyzed using mixed-effect model 222 223 ANOVA with physiological data used as the response variables and tribe and condition as predictor variables. Tests were performed with the lmer function within the lme4 package (Bates, 224 225 Maechler, Bolker & Walker 2015). To assess bivariate relationships between plant functional traits, we performed simple regression analyses (using the 'lm' function). Non-parametric data 226 were analyzed via Kruskal-Wallis rank sum test paired with a post hoc pairwise Wilcox test. We 227 also used Akaike's information criterion, adjusted for small sample size AICc model selection to 228 229 determine the most impactful trait parameters determining drought response using the "MuMIn" package (Grueber, Nakagawa, Laws & Jamieson 2011; Bartoń 2018). All data were analyzed in 230 the statistical program R V3.5.3 (R Core Team 2020). In order to summarize the relationships 231 and range of physiological, functional, and microanatomical diversity represented in our dry-232 down sample, we conducted a Principal Component Analysis (PCA) using the "prcomp" 233 function within the "stats" library on the mean trait data across species, which cumulatively 234 235 explained 72% of the variation in traits (Fig. 2). Not all traits were measured for every species, and so we focused on key traits coming from each of the data types we measured. The purpose of 236 237 the PCA was to visually explore the multivariate relationships among species in multivariate space, and only includes species that had all functional trait data (excluding *P. notatum*, *P.* 238 239 *juergensii*, and *F. ovina*).

240

241 **RESULTS**

242 Drought responses differed among species to the experimentally induced dry-down. Physiological viability, defined as maintaining at least 25% of the initial photosynthetic rate, 243 ranged from 4-33 days (Fig. 3A). Drought duration (days in drought before re-watering) was 244 similar among species within tribes, but varied significantly across tribes (Fig. 3A, P < 0.001). 245 246 Species within the tribe Cynodonteae (B. dactyloides & B. gracilis) were physiologically viable for the longest time during the dry-down, reaching the "stressed" stage after 30 days in drought 247 (Fig. 3A, Supporting Table S2), whereas the most drought-sensitive species were within 248 Paspaleae and Poeae. These tribes were similar (P > 0.05), reaching the 'stressed' stage more 249 than 20 days before the Cynodonteae tribe (Fig. 3A, Supporting Table S2). The recovery of grass 250 species and tribes following re-watering displayed a more variable response (Fig. 3B). There 251 were no statistical differences among tribes in their recovery dynamics (P > 0.05), though 252

significant differences were observed among species and within tribes (P < 0.05). S. nutans and 253 254 B. gracilis were the only species that that exceeded pre-drought photosynthesis levels after recovery (114% and 121% of initial A_n , respectively - Fig. 3B). Festuca ovina and B. dactyloides 255 were the only species that did not regain at least 50% of 'Initial' physiological measurements. 256 Several species within Paspaleae, Paniceae, Danthoneae, and Andropogoneae lineages did not 257 fully recover (100%) to 'Initial' physiological levels within the experimental timeframe. 258 However, all recovered to at least 50% of Day 1 measurements (Fig. 3B). Andropogoneae 259 260 species displayed the highest drought recovery, regaining on average over 92% of physiological functioning while the only species measured in Poeae (F. ovina) was the least resilient, with only 261

262 42% recovery of pre-drought A_n following re-watering.

Leaf economic trait data were collected at the conclusion of the dry-down when recovery 263 264 data was collected for each individual sample. Production of aboveground biomass was observed to vary significantly at both tribe and species level (Supporting Table S4, S5; P < 0.001). SLA 265 was statistically similar within Andropogoneae, Cynodonteae, and Danthonieae (P > 0.05) while 266 SLA within Paniceae species displayed significant variation (P < 0.05), ranging from 29 cm²g⁻¹ 267 (P. virgatum) – 143 cm²g⁻¹ (U. ruziziensis). Similarly, LDMC was statistically similar across 268 tribes, except for two species within Paniceae (P < 0.05) (S. viridis and U. ruziziensis) 269 270 (Supporting Table S2). The production of fine root length (diameter < 0.5 mm) differed among tribe (P < 0.0001) and species (P < 0.0001). Significant differences in SRL were observed across 271 272 tribes and species as well (P < 0.001, Supporting Table S2). All species within their respective tribes were found to have statistically similar *SRL* except for species in Paniceae (P < 0.05), 273 Paspaleae (P < 0.05), and Danthonieae (P < 0.01) due to the 294.99 cm g^{-1} difference in SRL. 274 Most microanatomical traits displayed significant differences among species across tribes 275 276 (P < 0.05) but typically had reduced variability between species within the same tribe X_{area} was 277 statistically different between tribes (P < 0.001) with the exception of Danthonieae and Cynodonteae (P = 0.485). Andropogoneae (589.308 μ m²) had the largest X_{area} and was 5x larger 278 than the smallest X_{area} found in Cynodonteae (108.957 μ m²) (P < 0.01; Supporting Table S3). 279 While there were significant species differences found across all tribes (P < 0.001; Supporting 280 Table S3) there were no observable species differences within tribe (P > 0.05). $X_{diameter}$ reflected 281 a similar pattern to that of X_{area} : significant differences between tribes (P < 0.001) and 282 statistically similar values within tribes (P > 0.15; Supporting Table S3) (consistent with 283

phylogenetic niche conservatism). Xylem resistance to cavitation (t/b) differed significantly

across tribes (P < 0.001), species (P < 0.001), but not among species within a tribe (P > 0.05;

Supporting Table S3). Stomata within the subsampled area (S_{count}) showed high variation; the

significant differences among tribes (P < 0.01) are likely attributed to Paniceae, which had higher

288 *S_{count}* (Supporting Table S3).

There were few statistically significant relationships explaining drought responses and 289 recovery among tribes (Supporting Fig. S1), except for X_{area} and S_{count} . Surprisingly, given the 290 large volume of literature designating *iWUE* as a pivotal functional trait reflecting drought 291 tolerance, there was no statistically significant correlation of *iWUE* with drought resistance or 292 resilience (P > 0.05). However, stomatal number was significantly negatively correlated with 293 drought duration (P < 0.01) (Fig. 4), X_{area} (P < 0.05), and *iWUE* (P < 0.05) (Supporting Fig. S1). 294 295 Results also indicate a differentiation between sample productivity and economic growth 296 strategies (Fig 5). A significant relationship was observed when comparing above and belowground biomass (Fig 5A), yielding a tight positive relationship (P < 0.001; $r^2 = 0.825$). Yet, 297 when above and belowground economic strategies (SLA and SRL, respectively) were calculated, 298 a breakdown in the previous relationship was observed (P > 0.05; $r^2 = 0.008$) (Fig. 5B). While 299 SLA displayed no bivariate relationships with other traits, LDMC correlated with t/b (P < 0.01) 300 301 and S_{count} (P < 0.05) (Supporting Fig. S1). The AICc model selection process indicated how the selected functional traits influence both drought resistance and resilience. The model explaining 302 303 the greatest variation in drought sensitivity included Tribe, S_{count}, *iWUE*, *LDMC*, and *SLA*. 304 However, the best explanation for variation in drought resilience included a single anatomical trait: S_{count}. 305

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307 DISCUSSION

Historically, water use efficiency (*WUE*) has been used as a seminal physiological trait to explain why some species persist and others succumb to drought (Fay *et al.* 2002; Leakey *et al.* 2019; Lavergne *et al.* 2019; Yang *et al.* 2021). However, the data presented here illustrate that using this physiological trait by itself may misconstrue the interpretation of drought responses in grass species (Fig. 3A; Supporting Fig. S1) (Morgan *et al.* 2011). Physiological, morphological (*SLA*, *SRL*, *LDMC*), and anatomical data (xylem area, stomata number, t/b) in combination help provide a more nuanced perspective on grass responses to drought, and should be integrated in

order to more appropriately identify mechanisms of water stress across diverse grass lineages. 315 Shared phylogenetic and biogeographic histories have resulted in unique adaptations and trait 316 development in Poaceae that are reflected in the patterns of global distribution and subsequent 317 responses to drought (Christin, Freckleton & Osborne 2010; Liu & Osborne 2015; 318 Watcharamongkol, Christin & Osborne 2018). Here, we show considerable variation in 319 320 functional traits among diverse species within Poaceae in response to drought and recovery. In 321 addition, these data illustrate that the traits and adaptations that confer an ability to withstand 322 extreme drought conditions are not the same as the traits and adaptations that confer an ability to recover from drought, as they are driven by the coordination of different factors (Fig. 2; Fig. 3). 323

The grass species measured here displayed variable sensitivity to the dry-down, visible in 324 both drought response and recovery (Fig. 3). The physiological responses observed during the 325 326 dry-down were statistically related to variability in anatomical features, mainly those influencing water relations at the leaf level (Fig. 4). We also found a separation between productivity and 327 328 economic strategies in both above and belowground tissues of the selected Poaceae species (Fig. 5), reflecting a trade-off in leaf and root growth economic strategies. Differences in economic 329 330 traits indicated water-use strategies varied in drought responses (Funk et al. 2017; Reich & Flores-Moreno 2017; Cui, Weng, Yan & Xia 2020). For example, plants with lower SLA and 331 higher SRL traits are likely to have lower metabolic costs and grow in resource-poor 332 environments with an increased ability to acquire resources (Cornelissen et al. 2003; Pérez-333 334 Harguindeguy et al. 2013). These trait strategies should allow for sustained physiological 335 tolerance and quicker drought conditions. While evolutionary relatedness guides physiological, morphological, and anatomical traits in determining drought responses, drought recovery was 336 mainly driven by S_{count} alone. The number of stomata within the subsampled cross-sectional area 337 (CSA) should not be interpreted as a stomatal density measurement because the subsampling 338 method we used was not limited to a defined leaf area. Instead, S_{count} indicates the number of 339 stomata that are supplying CO_2 to major and minor vascular bundles, which may be a more 340 direct mechanistic comparison to that of stomatal densities. Stomatal density, rather, considers a 341 leaf's surface as a uniform and homogeneous surface and has little to no consideration for 342 microanatomical traits. Such microanatomical trait measurements (i.e., mesophyll area, bundle 343 sheath area, and the diffusion distance through mesophyll) are important to consider due to their 344

influence on carbon assimilation and subsequent water loss (Esau 1953; Sack & Frole 2006;
Ocheltree, Nippert & Prasad 2011; Lundgren *et al.* 2019).

Low soil moisture negatively impacts growth, increases xylem tension, and decreases 347 carbon assimilation (Lemoine, Griffin-Nolan, Lock & Knapp 2018; Jardine, Thomas & Osborne 348 2021). The ability to mitigate and recover from drought is based on anatomical and physiological 349 traits (Taylor, Ripley, Woodward & Osborne 2011; Olson, Anfodillo, Gleason & McCulloh 350 351 2020). While the impacts of severe drought on the physiology of grassland species have been 352 observed in previous research, few studies combined physiological, whole-leaf, and microanatomical trait data (Fry et al. 2013; Liu & Osborne 2015). As was previously mentioned, 353 354 we intentionally chose species (within the same tribe) that were previously reported to have varying responses to low soil moisture. However, our results indicate that closely related grasses 355 356 can respond similarly to decreasing soil moisture (Fig. 3A; Fig. 4) but display variable responses 357 when water becomes available (Fig. 3B). This variability supports previous claims that drought 358 responses within a functional type (i.e. C_4 grasses) are not uniform and vary due to a myriad of reasons (i.e. evolutionary histories, functional trait, etc.) (Liu & Osborne 2015; Griffith et al. 359 360 2020). The diversity in physiological responses among species has been observed to protect individuals and populations while protecting ecosystem functioning from detrimental effects of 361 drought (Mori, Furukawa & Sasaki 2013; Kreyling et al. 2017; Roberts, Twidwell, Angeler & 362 Allen 2019). 363

364 Given the fundamental role that past evolutionary histories have played in shaping 365 current species distributions (Fox et al. 2018; Folk, Siniscalchi & Soltis 2020), species that exhibit variable responses to ecosystem disturbances would benefit more than species that 366 maintain static responses (Isbell & Wilsey 2011; Isbell et al. 2015). For instance, species that 367 were more drought resistant (Cynodonteae) are broadly represented in the mixed and shortgrass 368 prairies of North America, regions that are known to have less rainfall and more frequent drought 369 (Carroll et al. 2021). In contrast, the drought-sensitive species (Paspalum) are from tropical 370 locations where moisture is typically not the most limiting resource. In addition, Cynodonteae 371 372 were also observed to have fewer stomata and decreased gas exchange rates compared to Paniceae and Paspalum species, leading to less water loss (Fig. 4; Supporting Table S2, S3). 373 Therefore, it stands to reason that phylogenetically dissimilar species evolving under different 374 375 environmental constraints would exhibit disparate drought response to the imposed dry-down,

while more closely related species would respond more uniformly (Fig. 3A; Fig. 4). Our data 376 377 also displays a clear indication of an evolved plasticity in physiological responses to variable climate conditions, as native grasses typically occupy regions with similar climate variability 378 379 (Bachle, Griffith & Nippert 2018). Grasses sampled in this experiment were severely desiccated and recovered >50% of pre-drought physiological functioning and in several cases, physiological 380 rates that were 20-30% higher than the initial state (Fig. 3B), highlighting a potentially unique 381 characteristic of grasses across lineages. The ability to quickly acquire water and other nutrients 382 383 following drought disturbances likely facilitates grasses competing with other neighboring functional types with deeper access to water (Holdo, Nippert & Mack 2018; Kulmatiski, Beard, 384 Holdrege & February 2020). 385

Water availability directly impacts plant physiological responses, which are constrained 386 387 by internal anatomical machinery (Fig. 1) (Christin et al. 2013; Bachle & Nippert 2021). For example, the spatial separation of C₄ photosynthesis allows for a reduced stomatal conductance 388 389 and decreased water loss, leading to higher water-use efficiencies (Farguhar & Sharkey 1982; Ehleringer & Monson 1993; Berry & Patel 2008; Way, Katul, Manzoni & Vico 2014). This 390 391 characteristic is often assumed to underlie success under conditions of increased aridity. Our findings, however, do not support this claim (Supporting Fig. S1). *iWUE* was not observed to 392 393 directly aid in drought sensitivity or recovery of grasses in this study but it was positively related to the number of stomata (Supporting Fig. S1), indicating here, that the presence of more stomata 394 395 is associated with higher *iWUE*. This counterintuitive result does not indicate a more drought 396 tolerant strategy, rather, it communicates higher gas exchange rates resulting in greater loss of water leading to desiccation (Xu & Zhou 2008; Bertolino et al. 2019; Knapp et al. 2020). 397 However, previous research has indicated that stomatal patterning, morphology, and densities 398 can greatly influence/alter physiological responses to water stress (Nunes, Zhang & Raissig 399 400 2020; Harrison, Arce Cubas, Gray & Hepworth 2020). Figure 4 clearly indicates species with more stomata have an increased sensitivity (decreased resistance) to drought, which may require 401 reevaluations of previously held claims regarding the functional significance of WUE (Condon 402 2004). 403

404 Xylem characteristics have also been previously shown to impact an individual's water–
405 use (Wahl & Ryser 2000; He *et al.* 2020; Olson *et al.* 2020). Xylem area is a commonly
406 measured trait because it corresponds with the amount of water that can be transported at any

given time. Here, our results indicate two water transport strategies. Larger xylem (X_{area}) 407 408 decreases drought resistance while displaying a positive relationship with recovery, when excluding A. gerardii and S. nutans (Supporting Fig. S1). This strategy enables individuals with 409 larger X_{area} to transport greater amounts of water, when available (seen in recovery). But, 410 drought conditions can lead to increased tension on the water column inside xylem vessels, 411 ultimately increasing the potential for embolism formation during drought conditions (McCulloh, 412 413 Domec, Johnson, Smith & Meinzer 2019). Previous research has highlighted how increased 414 thickness of xylem wall tissue with smaller diameter lumen (t/b) can protect from embolism events in water limiting conditions (Blackman et al. 2018; Guérin et al. 2020); however, our data 415 do not corroborate such findings (Supporting Fig. S1). Anatomical traits were observed to 416 contain large variation, which we can contribute to two main factors: (1) Our sample size was 417 418 relatively small, due to the time - consuming nature of anatomical studies; and (2) microanatomical traits are complex in nature and have large variability among individuals and 419 420 within grass leaves (Fig. 1) (Ocheltree et al. 2011; Bachle & Nippert 2021).

While microanatomical traits and leaf-level physiological rates provide key mechanistic 421 422 insights into drought sensitivity and resilience, whole-plant traits are more easily observable and require less detailed scientific instrumentation and training (John et al. 2017; Reich & Flores-423 424 Moreno 2017). Whole-plant traits illustrate broader growth strategies by the individual, such as resistance or avoidance of detrimental growth conditions. Our results indicate a linear 425 426 relationship between above and belowground productivity (Fig. 5A), indicating a constant 427 proportional investment by the selected grasses. However, when comparing two widely utilized traits within the leaf and root economic spectrum (SLA and SRL), the previous relationship 428 breaks down to reveal tradeoffs in grass growth strategies (Fig. 5B). Individuals that invest in a 429 root system designed for quick absorption of water and nutrients (high SRL) can only produce an 430 inexpensive leaf (low SLA), while more 'expensive' leaves (high SLA) appear to be associated 431 with a less economically efficient rooting strategy (Fig. 5A). This finding highlights the inability 432 of grasses to produce tissues at the upper limits of the leaf and root economic spectrum. 433 However, it's important to recognize and consider the potential for convoluted interpretations of 434 economic traits, as they are the amalgamation of several underlying traits (Pérez-Harguindeguy 435 et al. 2013; Bachle & Nippert 2021). In addition, there could be grasses, which were not 436 437 investigated here, that invest in high SLA and SRL. One of these complex traits is leaf dry matter 438 content (*LDMC*), which was correlated with the thickening of xylem vessel walls or xylem
439 reinforcement: *t/b*. This thickening of water transport tissues (or xylem reinforcement) increases
440 the strength at which the water column can be under tension, allowing for a more negative water
441 potential while decreasing the likelihood of cavitation, a physiological indicator of drought stress
442 (McCulloh *et al.* 2019).

There is currently a dearth of available functional trait data in grasses, an 443 underrepresented functional type in trait databases given the importance of grass species for food 444 445 and forage and their geographical coverage. Results from this study indicate the need for increased collection of grass functional traits across a diverse assemblage of species within a 446 functional type. Plant functional types are often used in ecosystem models to more easily group 447 plants by common features; however, as our results indicate, this may lead to poor 448 449 parameterization and model output as such functional types do not account for phylogenetic relatedness. Our AICc selection indicates that microanatomical traits (specifically stomatal 450 451 count), physiological, economic traits, and phylogeny were essential to understanding species ability to withstand drought, while stomatal count was the best explanation for recovery 452 453 responses. The trait data from these lineage-specific responses to drought have potential consequences for how different grasslands are represented and forecast in Earth System Models 454 455 (Still, Cotton & Griffith 2018; Griffith et al. 2020).

The evolutionary histories of lineages within Poaceae have led to the development of 456 457 unique morphology (leaves and roots), anatomy (stomatal shape and photosynthetic tissue 458 arrangement), and physiology; these traits have allowed this functional type to dominate much of the surface cover on every continent (excluding Antarctica). Phylogenetic divergences and 459 subsequent trait adaptations have led to contrasting responses and recovery from drought (Fig. 460 3). Anatomical traits were key in explaining physiological drought response and recovery, 461 specifically traits concerned with water usage. Surprisingly, species that exhibited increased 462 *iWUE* were more prone to quicker desiccation, which is most likely due to the same individuals 463 maintaining larger numbers of stomata and higher overall rates of gas exchange (Supporting Fig. 464 S1). Interestingly this same trait (S_{count}) was responsible for recovery from drought, which did 465 allow for faster drought recovery. This study underscores the importance of collecting a myriad 466 of in-depth trait data from several Poaceae lineages to better understand the mechanisms that 467 468 describe drought responses and recovery.

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- 752 Figure Legends
- **Figure 1:** Leaf cross-sections of each major grass tribe stained with Safranin Red and Fast
- 754 Green. Top left, Andropogoneae; top right, Cynodonteae; bottom left, Danthonieae; bottom right,
- 755 Paniceae. Image taken with a Zeiss 880 confocal microscope.
- **Figure 2:** Principal components analysis (PCA) of mean trait values (in red text) of species in
- the dry down phase of the experiment. This PCA provides a summary of species in multivariate
- trait space using the first two PC axes, which together account for 72% of the trait variation. PC1
- vas most associated with variation in water use and rooting strategies whereas PC2 was
- 760 primarily associated with photosynthetic rate. Information concerning PCA axes importance and
- subsequent loadings are located in Supporting Table 6. Andropogoneae (light red), Cynodonteae
- (grey), Danthonieae (green), and Paniceae (blue); each point is a species mean.
- **Figure 3:** A) Number of days each species and tribe lasted before stomatal closure and
- rewatering occurred. B) The physiological recovery (A_n) compared to Day 0 or Initial physiology
- 765 (measured here as a percent). Dashed line signifies a complete 100% recovery of physiological
- function (i.e., A_n at or above its initial value). Andropogoneae (light red), Cynodonteae (grey),
- 767 Danthonieae (green), Paniceae (blue), Paspaleae (yellow), Poeae (red); each point is a species
- 768 mean and \pm SE.
- **Figure 4.** Relationship between stomatal count per entire leaf cross-section and days in drought
- before "Recovery". Andropogoneae (light), Cynodonteae (grey), Danthonieae (green), and
- Paniceae (blue); each point is a species mean and \pm SE.
- **Figure 5:** A) The relationship between leaf biomass and root biomass. B) Specific leaf area
- against specific root length. Andropogoneae (light red), Cynodonteae (grey), Danthonieae
- (green), Paniceae (blue), Paspaleae (yellow), Poeae (red); each point is a species mean and \pm SE.
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776 **Conflict of interest statement**

- We declare that the submitted work was not carried out in the presence of any personal,
- professional or financial relationships that could potentially be construed as a conflict of interest.