Replicated radiations in the South American Marsh Pitcher Plants (*Heliamphora*) lead to convergent carnivorous trap morphologies

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7 Abstract

- 8 **Premise:** The evolution of carnivorous pitcher traps across multiple angiosperm lineages
- 9 represents a classic example of morphological convergence. Nevertheless, no comparative study
 10 to-date has examined pitcher evolution from a quantitative morphometric perspective.
- 11 **Methods:** In the present study, we used comparative morphometric approaches to quantify the

12 shape space occupied by *Heliamphora* pitchers and to trace evolutionary trajectories through this

13 space to examine patterns of divergence and convergence within the genus. We also investigated

14 pitcher development, and in particular, how the packing of pitchers is affected by crowding, a

- 15 common condition in their natural environments.
- 16 Key Results: Our results showed that *Heliamphora* pitchers have diverged along three main 17 axes in morphospace: pitcher curvature, nectar spoon elaboration, and pitcher stoutness. Both 18 curvature and stoutness are correlated with pitcher size, suggesting structural constraints in 19 pitcher morphological evolution. Among the four traits (curvature, spoon elaboration, stoutness 20 and size), all but curvature lacked phylogenetic signal and showed marked convergence across 21 the phylogeny. We also observed tighter packing of pitchers in crowded conditions, and this
- 22 effect was most pronounced in curved, slender pitchers.

23 **Conclusions:** Overall, our study demonstrates that diversification and convergent evolution of

- 24 carnivory-related traits extends to finer evolutionary timescales, reinforcing the notion that
- 25 ecological specialization may not necessarily be an evolutionary dead end.

26 摘要 (Simplified Chinese)

27 食虫植物的瓶状捕虫器在不同被子植物谱系中的演化是形态趋同进化的经典案例。然而, 迄今为止还没有研究使用系统发育比较分析法与计量方法来研究瓶状捕虫器的演化。本研 28 究使用了上述手段来研究南美沼泽瓶子草(Heliamphora)的瓶状捕虫器是如何占据形态 29 空间,并在捕虫器的形态空间内追踪它们的演化轨迹,以研究沼泽瓶子草属捕虫器的趋异 30 与趋同的演化模式。同时,本研究还探讨了沼泽瓶子草属植株(捕虫瓶)的发育模式,特 31 别是在沼泽瓶子草在原生地的发育是如何受到种群密度过高(拥挤)的影响。本研究的结 32 果表明,沼泽瓶子草属捕虫瓶在形态空间里沿着三个主要形态轴分化:捕虫瓶的弯曲度、 33 蜜匙的精细程度和捕虫瓶的粗壮程度。捕虫瓶的弯曲度和粗壮度均与瓶子的大小相关,表 34 明捕虫瓶在形态进化中存在结构约束。在这四个形态特征中(捕虫瓶曲率、蜜匙精细程 35 度、捕虫瓶粗壮度与大小),除了捕虫瓶曲率,其他的形态特征都没有显示系统发育信 36

37 号。在南美沼泽瓶子草属的进化中,蜜匙精细程度、捕虫瓶粗壮度与大小都显示了趋同演
38 化。本研究还阐明了瓶子草植株在拥挤条件下,捕虫瓶会更加紧密地排列,而且这种效应
39 在有着弯曲、细长的捕虫瓶的物种中最为显著。总体来说,本研究显示与食肉有关的特征
40 的趋异和趋同演化发生在更微观的时间尺度上,也表明了生态专化或许并不是生物进化的
41 亚坦回

41 死胡同。

Keywords: Sarraceniaceae, carnivorous plants, pitcher plants, morphological evolution, phylogenetic comparative
 methods

44 Introduction

- 45 Pitcher plants are one of the most dramatic examples of convergent evolution in carnivorous
- 46 plants, with at least 6 independent origins in angiosperms (Fleischmann et al. 2018). The pitchers
- 47 are specialized leaves that form cavities filled with an often viscous liquid containing a suite of
- 48 digestive enzymes (Adlassnig, Peroutka, and Lendl 2011). Pitcher plants utilize these pitchers to
- 49 attract, drown, and consume small animal prey in order to obtain nutrients not available from the
- 50 abiotic environment (Juniper, Robins, and Joel 1989; Givnish et al. 2018).
- 51 Among the independently evolved pitcher plant lineages, *Nepenthes* (Caryophyllales),
- 52 Sarraceniaceae (Ericales), and *Cephalotus follicularis* (Oxalidales) are commonly considered as
- 53 the true pitcher plants due to their highly specialized carnivorous pitchers. Moreover, the true
- 54 pitcher plants are considerably older (all emerged during the Cretaceous) as compared to the
- 55 more recently evolved (< 3 mya) pitcher lineages in Poales (e.g. *Brocchinia reducta* and
- 56 Paepalanthus bromelioides), which have structurally simpler pitchers morphologically similar to
- 57 their tank forming relatives (Magallón et al. 2015; Cross et al. 2018). With the exception of the
- 58 monotypic Australian pitcher plant *C. follicularis*, *Nepenthes* (the tropical pitcher plants) and
- 59 Sarraceniaceae (the American pitcher plants) both represent true pitcher lineages with high
- species diversity, together comprising more than 98% of all extent pitcher plant taxa (McPherson
- 61 et al. 2011; McPherson and Schnell 2011; Fleischmann et al. 2018; Murphy et al. 2020).
- 62 In addition to dispersal and vicariance (Ellison et al. 2012; Naczi 2018; Murphy et al. 2020),
- 63 adaptive radiation played an important role in the diversification of *Nepenthes* and
- 64 Sarraceniaceae (Pavlovic 2012; Clarke et al. 2018; Thorogood, Bauer, and Hiscock 2018). The
- 65 emergence of pitcher traps has likely facilitated the diversification in Nepenthes and
- 66 Sarraceniaceae by allowing them to adapt to new ecological niches, resulting in dramatic
- 67 morphological variations in pitcher traps (Pavlovic 2012; Clarke et al. 2018; Thorogood, Bauer,
- and Hiscock 2018). Many Nepenthes species have diversified and further modified their pitcher
- 69 traps to specialize in novel symbiotic relationships (e.g. ant-mutualism in *N. bicalcarata*) or
- novel nutrient acquisition strategies (e.g. detritus-feeding in *N. ampullaria*, fecal-feeding in *N.*
- 71 *lowii*, and subterranean-feeding in *N. pudica*) (Thornham et al. 2012; Moran et al. 2012; Gilbert
- et al. 2022; Dančák et al. 2022). A similar radiation has occurred across Sarraceniaceae lineages,
- 73 giving rise to many species with novel trap types (e.g. lobster traps of *Darlingtonia californica*
- 74 and *Sarracenia psittacina*), trapping mechanisms (e.g. superhydrophilic hairy surface of
- 75 *Heliamphora nutans*), symbiotic relationship (e.g. *S. purpurea* and commensal *Wyeomyia*
- *smithii*), and nutrient specializations (Jaffe et al. 1992; Peterson et al. 2008; McPherson et al. 2011; Peterson et al. 2012; Nagri 2018). Pitcher membrils
- 77 2011; Bauer et al. 2013; Naczi 2018). Pitcher morphology in Sarraceniaceae and *Nepenthes* is
- also plastic to varying degree, affected by both biotic (e.g. crowding or competition from

neighboring plants) and abiotic factors (e.g. extreme microclimate) (Brewer 1999; McPherson et

al. 2011; McPherson and Schnell 2011; Clarke et al. 2018).

81 Despite the wide variety of morphologies associated with pitcher traps, few comparative studies

82 have focused on the morphological evolution of carnivorous traps, and none with quantitative

- 83 methods. Morphometric analyses allow us to define the spaces occupied by phenotypic
- combinations as well as those that are empty (Stayton 2019; Segall et al. 2020; Jardine et al.
- 85 2022) and, in the context of well supported phylogenies (Givnish et al. 1997), identify instances
- of morphological convergence (Papadopulos et al. 2013; Smith and Kriebel 2018). This
 quantitative shape information also sets the stage for rigorous testing of ecological and
- quantitative shape information also sets the stage for rigorous testing of ecological and
 evolutionary hypotheses (Smith and Kriebel 2018; Dellinger et al. 2019; Kriebel et al. 2022).
- 89 Here we focus on the evolution of pitcher morphologies in the South American marsh pitcher
- 90 plants *Heliamphora* (Sarraceniaceae). This genus is endemic to the Guiana Highlands, and is the
- 91 most species-rich genus in the family, with 23 extant species and several yet to be described

92 (McPherson et al. 2011). The major clades are thought to have emerged through both vicariance

- 93 and dispersal during the Miocene with more recent diversification in each clade driven by
- 94 vertical displacement during the Pleistocene glacial-interglacial cycles (Liu and Smith 2021).
- 95 The adult pitchers vary dramatically in pitcher shape, size, nectar spoon structure and many other
- 96 taxonomically important characters across species (Figure 1) (McPherson et al. 2011). It has
- 97 been suggested these diverse morphologies may relate to differences in microclimate (Nerz
- 2004), prey attraction strategies (Wistuba, Harbarth, and Carow 2001; Wistuba, Carow, and
- Harbarth 2002), and/or shifts in prey composition (Jaffe et al. 1992). Furthermore, structural
- stability of the pitcher could impose constraints on the morphological evolution of *Heliamphora*
- 101 pitchers, limiting available pitcher morphospace.

102 In the present study, we used comparative morphometric approaches to quantify the shape space

103 occupied by *Heliamphora* pitchers and to trace evolutionary trajectories through this space.

- 104 Building on previous phylogenetic work in *Heliamphora* (Liu and Smith 2021), we also tested
- 105 for phylogenetic signal in pitcher shape and investigated how pitcher shape evolution may be
- related to pitcher size. In particular, we hypothesize that aspects of shape evolution may be
- tightly correlated with height given the structural constraints imposed by supporting high volume
- 108 of pitcher fluid. In light of the dense packing in many natural populations (McPherson et al.
- 109 2011), we also considered how pitcher development responded to crowding and tested the
- 110 hypothesis that pitcher angle (the angle between two most recently developed pitchers) shrinks in
- 111 crowded conditions to allow for tighter packing. In interpreting these results, we discuss the
- potential ecological significance of *Heliamphora*'s diverse pitcher morphologies and ability to
- 113 modify pitcher development in response to varying conditions.
- 114

115 MATERIALS AND METHODS

116 **Taxon sampling and phylogenetic framework**

- 117 We sampled 23 described and 2 additional yet to be described taxa of *Heliamphora* for this study
- 118 (Appendix S1, Table S1; see the Supplementary Data with this article). All specimens were

- 119 grown under controlled environmental conditions [photoperiod 15hr per day (full sun);
- temperature range 12–25 °C; relative humidity > 80%], resembling the climate of their native
- 121 habitats (Jaffe et al. 1992; Adlassnig, Pranjic, et al. 2010). Species determinations of the living
- 122 collections were confirmed by comparing their morphological traits to the original descriptions
- 123 (Bentham 1840; Nerz and Wistuba 2000; Carow, Wistuba, and Harbarth 2005; Fleischmann,
- 124 Wistuba, and Nerz 2009; McPherson et al. 2011).
- 125 We used the recently published phylogeny of 24 of the 25 *Heliamphora* taxa (Liu and Smith
- 126 2021) to reconstruct the evolution of adult pitcher shape, maximum adult pitcher size and pitcher
- 127 angle. This phylogeny lacks *H. macdonaldae*, which is thought to be sister to *H. tatei* based on
- similarities in pitcher morphology and distributional range (Maguire 1978; McPherson et al.
- 129 2011; Liu and Smith 2021). We manually included this species in the tree (Figure 1) by placing it
- 130 sister to *H. tatei* and setting their split to 0.75 mya, which is the average node depth of other
- 131 sister pairs in the W lineage, i.e., (*H. ceracea, H. hispida*) and (*H. neblinae, H. parva*).

132 Morphometric analysis of pitcher shapes

- 133 Data for maximum pitcher size (i.e. the maximum length of adult pitchers observed growing
- 134 under optimal conditions, *in situ*) were obtained from the monograph of *Heliamphora*
- 135 (McPherson et al. 2011). Adult pitcher shape was characterized using images of fully opened and
- 136 matured pitchers taken from living individuals. For each taxon, a single adult pitcher from the
- individual sampled for the phylogeny was photographed in a lateral view, providing a two-
- dimensional representation of the adult pitcher shape along the proximal-distal axis. The lateral
- 139 view captured shape variation among major structural components involved in carnivorous
- functions, such as the shapes of nectar spoon and pitcher body associated with prey attraction
- 141 and prey capture/retention, respectively (Figure 2). These images were converted into silhouettes
- in Affinity Designer 1.10.5, which were then transformed into outlines using the R package
 Momocs (Bonhomme et al. 2014). Also using Momocs, the outlines were converted into lists of
- 144 two-dimensional coordinates describing the polygons and then centered.
- A set of six landmarks were place on each outline to delineate pitcher functional zones (Figure 2)
- according to McPherson et. al. (2011). In *Heliamphora* pitchers, Zone 1 (landmarks 2-4)
- 147 corresponds to the nectar spoon, which is responsible for attracting prey. Zone 2 (landmarks 1, 2,
- 148 4, 5) represents the upper section of the pitcher above the narrow band and plays a role in prey
- 149 capture and retention. Zone 3 (landmarks 1, 5) consists of a narrow restriction of the pitcher
- 150 interior, located close to the waterline of the pitcher fluid. The position of the drainage hole
- 151 (indicated by landmark 1; the drainage hole is further adapted into drainage slit in E1 clade)
- determines the maximal level of pitcher fluid in the pitcher (landmarks 1, 5), which is crucial in maintaining carnivorous and other ecological functions in *Heliamphora* (see discussion). Zone 4
- maintaining carnivorous and other ecological functions in *Heliamphora* (see discussion). Zone 4
 (landmarks 1, 5, 6) comprises the bottom part of the pitchers and is involved in prey digestion as
- 155 well as nutrient absorption. Despite morphological diversity, these functional zones are
- 156 homologous and easily identifiable across *Heliamphora* species, making them reliable reference
- 157 points for the landmarks and preventing any distortion of the outlines.
- 158 Using the landmarks, shape variation of adult pitchers was quantified using elliptical Fourier
- analysis (EFA) that decomposed the outlines into 21 harmonics, each with four coefficients.
- 160 After aligning the outlines using Full Generalized Procustes alignment in Momocs, a principal

- 161 component analysis (PCA) was used to summarize the 84 coefficients resulting from the EFA.
- 162 Taxa were plotted along the PC axes to illustrate their distribution in morphospace. Blomberg's
- 163 K (Blomberg, Jr, and Ives 2003) for the shape PCs (PC1, PC2, and PC3) and adult pitcher size
- were estimated to infer phylogenetic signals in pitcher shapes and size by using 1000
- 165 randomizations of the data across the phylogeny to test if the value was significantly different
- 166 from K=0 (no phylogenetic signal).
- 167 Finally, we used ancestral state estimation to visualize the expected ancestral pitcher shape in
- 168 *Heliamphora*. We estimated the ancestral values for each of the three shape PCs and for
- 169 maximum pitcher size using the 'fastAnc 'function in R package phytools (Revell 2012). We
- 170 visualized the reconstructed shape given the estimated ancestral PC values using Momocs, and
- 171 scaled the size of the pitcher based on the estimated ancestral size.

172 Effects of crowding on pitcher angle

173 We used living collections of *Heliamphora* species to characterize the angle of pitcher

- 174 emergence and examine the effect of crowding on this angle. Depending on the availability of
- 175 material, we sampled 3 to 63 individuals per taxon for all 25 taxa. These individuals were
- 176 sampled from our collections and those of G. Pipis; in possessing adult pitchers, these plants
- were three or more years of age. For each sampled individual, images were taken directly above
- each growing point, i.e., where new pitchers are constantly produced (Appendix S2, Figure S1).
- For each growing point, a line was drawn from the growing point (rosette center) to the tip of the nectar spoon of the youngest (the most recently fully developed) pitcher. Similarity, a second
- 181 line was drawn for the second youngest pitcher. Pitcher angle (PA) was measured as the angle
- between the two straight lines using the software ImageJ (Schneider, Rasband, and Eliceiri
- 183 2012). We replicated these measurements for both juvenile and adult pitchers, since pitchers are
- 184 dimorphic in *Heliamphora* (McPherson et al. 2011) and may respond differently to crowding. To
- 185 test whether PA is affected by the presence of neighboring individuals (i.e., if crowding reduces
- 186 PA), phylogenetic paired t-tests were performed for both adult and juvenile pitchers using the
- ¹⁸⁷ 'phyl.pairedttest 'function in R package phytools (Revell 2012). Similarly, the effect of
- 188 developmental stage on PA was also tested with phylogenetic paired t-tests. In addition to 189 phylogenetic t-tests, regular t-tests were also conducted for each comparison. An additional chi
- 189 phylogenetic t-tests, regular t-tests were also conducted for each comparison. An additional chisquared test was performed to test whether the drastic bending observed in juvenile pitchers was
- associated with the presence of neighbors. Similar to the PCs, we computed Bloomberg's K for
- PA in adult and juvenile individuals growing with and without neighbors, and then tested if the
- 193 value was significantly different from zero.

194 Model testing for correlations among morphological traits

- 195 We used phylogenetic generalized linear models (PGLM) (Paradis and Claude 2002) to test the
- 196 relationship between shape variation and maximum pitcher size in adult pitchers. For some
- 197 combinations of shape PCs and size, we predicted a non-linear relationship. Thus, we compared
- 198 linear and quadratic models for each pair and present results for the best-fitting model (that with
- 199 the lowest AIC score). We fit these models using the R package 'phylolm 'with 1000 bootstraps
- 200 (Ho and Ane 2014). We assumed a correlation structure based on the Brownian motion (BM)
- 201 process, which models stochastic evolution in those traits. Similarly, the relationships between
- shape variation and pitcher angle were also tested using PGLM in adult pitchers growing singly

- and ones growing with neighbors present. Assuming BM process, the PCs were regressed against
- 204 pitcher angle with PGLM. Taxa with incomplete pitcher angle data due to sampling limitations
- 205 were dropped from the analysis.

206 **RESULTS**

207 Morphospace of Heliamphora Pitchers

208 The PCA of harmonic coefficients from the elliptic Fourier analysis captured 79% of shape 209 variation (SV) of adult pitchers in the first three PCs (Figure 3). PC1, accounting for 33.4% of 210 SV, describes pitcher body curvature or concavity. While the negative and positive PC1 values 211 described concave and convex pitcher shapes, respectively, PC1 values closer to zero represented 212 pitchers with straighter bodies. PC2, comprising 29.5% of SV, describes the elaboration (shape, relative size, and position) of nectar spoon with negative and positive values representing erect 213 214 vs. and elaborated structures, respectively. PC3, explaining 16.1% of SV, is related to pitcher 215 stoutness, or the general pitcher body length-to-width ratio. On the PC3 axis, negative values 216 described more slender pitcher body shapes whereas positive values corresponded to bulkier

- body shapes.
- 218 All of the measured pitcher traits, except for curvature, lacked a significant phylogenetic signal.
- 219 Specifically, we found Blomberg's K was not significantly different from zero for nectar spoon
- elaboration (PC2, K = 0.06, p = 0.886), pitcher stoutness (PC3, K = 0.09, p = 0.391), or
- maximum pitcher size (K = 0.12, p = 0.198), consistent with the frequent patterns of
- 222 convergence across the phylogeny (Figure 4). By contrast, significant phylogenetic signal was
- detected in pitcher curvature (PC1, K = 0.18, p = 0.018) (Figure 4). Looking at the phylogeny,
- 224 pitcher curvature appears to be conserved in the W+E1 and E2a clades but not in other lineages.
- In the W+E1 clade, all species were associated with convex pitcher bodies to varying degrees
- 226 (PC1 > 0.02), with the exception of *H. chimantensis*, which was associated with a straighter
- pitcher body (PC1 = -0.00321). In E2a, all adult pitchers were associated with very concave pitcher body shapes (PC1 < -0.04).
- 229

230 Relationships between Pitcher Shape and Size

- 231 Model comparisons indicated a strong quadratic relationship between pitcher curvature (PC1)
- and maximum pitcher size (Appendix S1, Table S2). As pitchers becomes straighter (i.e., less
- concave or convex), the maximum pitcher size generally increases (Figure 5), consistent with the
- 234 expectation that more erect pitchers can accommodate more pitcher fluid (see discussion). A
- strong linear relationship was found between pitcher stoutness (PC3) and maximum pitcher size,
- with larger pitchers being stouter (Figure 5). No relationship was found between nectar spoon
- elaboration (PC2) and maximum pitcher size (Figure 5).

238 Relationship of pitcher angle to growth conditions and pitcher shape

- 239 Our results show that *Heliamphora* growing in crowded conditions tend to put out new pitchers
- at a tighter angle than those in isolated conditions. A significant difference in pitcher angle (PA)
- 241 was detected between adult pitchers growing in isolated vs. crowded conditions with both
- 242 phylogenetic and non-phylogenetic t-tests (phylogenetic mean difference [PMD] = 13.64°, p =

- 243 0.002; p < 0.0001 for regular t-test, Figure 6). In juveniles, a significant difference was detected
- 244 in regular t-test (p < 0.0001) but not in the phylogenetic t-test (PMD = 6.36°, p = 0.199).
- 245 Moreover, we found juvenile pitchers tend to bend unnaturally when their growth was obstructed
- 246 by neighboring pitchers (chi-squared test, p < 0.001), possibly due to the more slender and 247
- parallel-to-ground body shape. No significant difference in PA was found between juvenile and
- 248 adult pitchers within each condition (crowded or not) using either t-test (Figure 6). Raw 249
- measurements for all taxa, stages and conditions are shown in Appendix S2, Figure S2. No 250 phylogenetic signal was detected in PA across developmental stages and crowding conditions (K
- 251 = 0.12, p = 0.481 [juvenile, not crowded]; K = 0.11, p = 0.353 [juvenile, crowded]; K = 0.10, p = 0.481 [juvenile, not crowded]; K = 0.10, p = 0.481 [juvenile, not crowded]; K = 0.10, p = 0.481 [juvenile, not crowded]; K = 0.10, p = 0.481 [juvenile, not crowded]; K = 0.10, p = 0.481 [juvenile, not crowded]; K = 0.11, p = 0.353 [juvenile, not crowded]; K = 0.10, p = 0.481 [juvenile, not crowded]; K = 0.11, p = 0.353 [juvenile, not crowded]; K = 0.10, p = 0.481 [juv
- 252 0.494 [adult, not crowded], K = 0.09, p = 0.597 [adult, crowded]).
- 253 To investigate the association between pitcher shape and PA, similar PGLM analyses were
- 254 performed for each crowding condition using data for adult pitchers. The analyses indicated a
- 255 strong quadratic relationship between pitcher curvature and pitcher angle only in crowded
- 256 conditions (Appendix S1, Table S2). When crowded, species with straighter pitcher bodies
- 257 generally put out new pitchers at a wider angle than those with curvier bodies (Appendix S2,
- 258 Figure S3). Similarly, a strong linear relationship between pitcher stoutness and pitcher angle
- 259 was only found in crowded condition. No relationship was found between nectar spoon
- 260 elaboration (PC2) and pitcher angle in either condition, indicating that, in crowded conditions,
- the degree of pitcher angle reduction depends on the morphology of the pitcher body rather than 261 262 the morphology of nectar spoon.
- 263

264 DISCUSSION

265 Divergent and convergent evolution of pitcher morphologies

- 266 Our morphospace analysis revealed three main shape axes across which *Heliamphora* pitchers
- 267 have diversified, namely in curvature (PC1), nectar spoon elaboration (PC2), and pitcher
- 268 stoutness (PC3). In accordance with the generally low phylogenetic signal in these traits, we
- 269 observed that most clades contain a diversity of shapes, e.g. curved and straight, with and
- 270 without elaborate spoons (Figure 1). Heliamphora pitchers also vary widely in size (almost five-
- 271 fold difference across the genus), even between closely related species (e.g., the sympatric H.
- 272 chimantensis at 35cm and H. pulchella at 10cm). Across these axes, Heliamphora species are
- 273 clustered in distinct regions of morphospace, e.g., curved with elaborate spoons or straight and
- 274 slender (Figure 3; Appendix S2, Figure S4; Appendices S3 and S4).
- 275 Accompanying these patterns of divergence in shape and size among close relatives, we also
- 276 documented frequent convergence across the phylogeny. For example, small pitchers (< 20 cm)
- 277 evolved convergently in three lineages (Figure 4). Interestingly, these small pitcher species can
- 278 be found across all major areas of distribution but none of them occur sympatrically (e.g. H.
- 279 hispida, Neblina Massif; H. puchella, Chimantá Massif; H. minor, Auyán Tepui; H. nutans and
- 280 H. arenicola, different parts of Eastern Tepui Chain). Such widespread convergence is often
- 281 associated with adaptation to a common ecological niche (Donoghue et al. 2022), and previous
- 282 authors suggest that small pitchers may be favored in habitats with suboptimal growth
- 283 conditions, e.g., shallow or infertile substrates, heavy shade, and periodic drought or heat stress

- (Wistuba, Carow, and Harbarth 2002; Givnish et al. 2018). Small pitchers impose lower 284
- 285 construction costs (Karagatzides and Ellison 2009; McPherson et al. 2011) and are more drought
- 286 and heat tolerant due to the reduced transpiration from the pitcher and evaporation from the
- 287 pitcher fluid (Adlassnig, Pranjic, et al. 2010).
- 288 Instances of small pitcher species occurring sympatrically with species of larger pitchers (e.g., H.
- 289 hispida and H. ceracea; H. pulchella and H. chimantensis; H. nutans and H. glabra) can be
- 290 found across all major distributional areas (McPherson et al. 2011), suggesting a potential case of
- 291 adaptive radiation into different ecological niches (Givnish et al. 1997). Sympatric pairs of small
- 292 and tall pitcher plant species may be associated with different capture strategies and,
- 293 consequently, prey composition (e.g. crawling insects like ants for short pitchers and flying
- 294 insects for tall pitchers), as has been shown in Sarracenia (Gibson 1983; Givnish 1989) and
- 295 Brocchinia (Givnish et al. 1997). Overall, the diversity of habitats and prey types provided by the
- 296 topologically complex Pantepui region combined with the geological isolation between tepuis 297 may account for the repeated diversification of pitcher size in each clade and the resulting
- 298
- convergence across the genus.
- 299 Similarly marked convergence is also observed in aspects of pitcher shape, although the
- 300 underlying drivers are less clear. The degree of elaboration in the nectar spoon may relate the
- 301 available prev and the reward needed to attract them (Jaffé et al. 1995; Plachno, Swiatek, and
- 302 Wistuba 2007). Differences in shape have been related to the surrounding vegetation, for
- 303 example, with *Heliamphora* growing through dense vegetation tending to have straight pitchers
- 304 to protrude above the rest to capture prey (Brewer-Carias 1972; Jaffe et al. 1992; McPherson et
- 305 al. 2011). The stoutness of the pitcher may be more directly related to the overall size of the
- 306 pitcher and may represent a developmental constraint (see below).

307 Gaps in pitcher morphospace due to evolutionary constraints

- 308 Unoccupied regions in the space of possible phenotypes are often attributed to constraints on
- 309 evolution, be they structural, ecological or developmental [e.g. Vernescu and Ryser 2009;
- 310 Stayton 2019; Chitwood and Mullins 2022]. We observed several gaps in pitcher space
- 311 represented the absence of certain trait combinations, such as large and curvy pitchers or small
- 312 and slender pitchers. The strong correlations between size and shape, in particular curvature and
- 313 stoutness (Figure 5), are consistent with structural constraints, i.e., that stout and/or curved 314 pitchers are limited in height while straight, slender pitchers can remain erect at greater heights.
- We postulate that, compared to species with straight body shapes, taxa with curvy shapes are less 315
- 316 structurally stable due to the center of their pitcher mass being further away from the growing
- 317 point on the rosette which provides structural support. Nevertheless, the potential structural
- 318 instability resulting from the curvy shape or large size can be mitigated through the provision of
- 319 structural support from nearby pitchers, vegetation, or organic debris [e.g. H. ionasi (McPherson
- 320 et al. 2011)].
- 321 This general trade-off between pitcher size and shape may allow some conservation of pitcher
- 322 volume, as pitchers can achieve similar volume by expanding laterally or vertically. Pitcher
- 323 volume is expected to be under strong selection as maintaining a high level of fluid is crucial to
- carnivory in Heliamphora (Jaffe et al. 1992; Nerz 2004; Bauer et al. 2013). At a high level, the 324
- 325 pitcher fluid is able to effectively rise along the pubescent inner pitcher wall up to the rim by

326 capillary action, forming superhydrophilic (slippery) trapping surface that facilitates prey capture

327 (Jaffe et al. 1992; Bauer et al. 2013). The regulation and maintenance of the pitcher fluid level

328 might be important in other ecological functions, such as thermal regulation, water reservoir

during hot and dry periods, and microhabitat for commensal microbiomes to facilitate digestion

330 (Jaffe et al. 1992; Nerz 2004; Adlassnig, Pranji, et al. 2010). We expect that there are also lower

limits on pitcher fluid level, bounded by the minimum volume needed to allow for carnivorous

332 functions.

333 Plasticity in pitcher development

334

Like all leaves, the pitcher development is plastic and can be affected by various environmental

factors (Brewer 1999; Ellison and Gotelli 2002; Fukushima et al. 2021). This is the first study to

examine plasticity in *Heliamphora* pitcher rosette development, specifically focusing on a

338 common natural condition — crowding due to the presence of neighboring plants. In nature,

Heliamphora populations are often crowded and restricted to habitats with suitable growth

340 conditions, such as small depressions on tepui summits and open clearings in the montane forests

- 341 (McPherson et al. 2011). Our study found that rosettes in such crowded conditions put out new
- 342 pitchers at tighter angle while those in growing without neighbors grow out at wider angle. This

effect is apparent in both juvenile and adult pitchers (Figure 6). In adult pitchers, the effect is

dependent on pitcher curvature and pitcher stoutness (Appendix S1, Table S2; Appendix S2,

Figure S3), suggesting the degree of angle reductions is further constrained by pitcher body

346 shape.

347 To visualize how the plastic differences in pitcher angle would translate to spacing of pitchers

348 and rosettes, we created two sets of diagrams based on two angles of emergence (140° and 125°,

Appendix S2, Figure S5). The wider angle, roughly the mean estimated in uncrowded conditions

350 (Figure 6), leads to evenly spaced pitchers in the mature rosette (Figure 7 A & B). Mature

rosettes typically comprise five active pitchers (e.g, Butschi, Huber, and Ammann 1989; Wistuba et al. 2005; Wrazidlo 2019; Golos 2020), and we added a sixth pitcher to demonstrate how a new

pitcher would overlap with an older inactive pitcher (Figure 7 B). Furthermore, we envisioned

how a clonal cluster of rosettes emerging from the rhizomes of a single plant could fill a

confined space, such as a depression in the rock substrate (Figure 7 C). When we simulated the

356 same developmental process with the narrower pitcher angle observed under crowded conditions

357 (Figure 7), we found that individual pitchers on each rosette exhibited greater overlap (Figure 7

358 D & E), but more new rosettes could be packed into the confined space (Figure 7 F), resulting in

increased pitcher density. While these visual models are only hypothetical, they illustrate how

pitcher angle will have direct consequences for the packing of pitchers into confined spaces and

361 likely affect fitness through access to light, rainwater, and prey resources.

362 It is possible that the increased density of pitchers could lead to a decreased prey capture rate per 363 nitcher due to competition from surrounding nitchers, as exidenced in Samaconia (Ciberr 1092)

pitcher due to competition from surrounding pitchers, as evidenced in *Sarracenia* (Gibson 1983;
 Givnish 1989). *Heliamphora* often form dense clonal populations, as they have a tendency to

365 produce new growing points (rosettes) along the rhizome of existing rosettes, whether in situ

366 (Wistuba, Carow, and Harbarth 2002; Fleischmann and Grande Allende 2011) or in cultivation

367 (Ziemer 1979; McPherson et al. 2011). Even if the individual pitchers receive less prey during

368 crowding, the entire clonal population may benefit from the tight packing of rosettes with higher

369 total prey capture and higher growth by maximizing area for photosynthesis.

370 While we chose to focus on the effects of crowding, other biotic and abiotic factors may 371 influence pitcher development and represent adaptive plasticity. For example, Heliamphora 372 grown under drought stress produce smaller pitchers, a response which may serve to minimize 373 transpiration (McPherson et al. 2011). Pitchers growing in heavy shade typically lose 374 carnivorous activity and primarily function as photosynthetic structures, as indicated by the 375 elongated pitcher bodies, and the absence of pitcher coloration, nectar spoon structure, or 376 attractants (McPherson et al. 2011). Similar plastic responses are found in *Darlingtonia*, 377 Sarracenia and all other pitcher plants, suggesting pitcher plasticity commonly associated with 378 trade-offs between carnivory and photosynthesis (Givnish et al. 1984; Ellison and Gotelli 2002; 379 Ellison and Farnsworth 2005; Pavlovič and Saganová 2015; Givnish et al. 2018; Fukushima et al. 380 2021). Further ecological studies should be conducted to investigate and better understand the 381 dynamic interplay of pitcher development, carnivory, and photosynthesis. Furthermore, it would

- be interesting to investigate whether crowding has a similar effect on pitcher angle in the closely related *S. purpurea* and *S. psittacina*, which also have a similar rosette growth form to
- 384 *Heliamphora*.
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387 CONCLUSION

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389 Both divergent and convergent evolution of Heliamphora pitcher morphologies was likely a 390 result of adaptive radiation across the dynamic Pantepui landscape over the last 20 million years 391 (Liu and Smith 2021). We observed repeated diversification of pitcher shape and size within 392 geographically structured clades as well as frequent convergence in form across the entire genus. 393 This convergence upon similar combinations of shapes and sizes suggest that these suites of 394 traits may be adaptive, as has been proposed for convergently evolved pitcher trait combinations 395 in other carnivorous plant lineages (Clarke et al. 2018; Thorogood, Bauer, and Hiscock 2018). 396 Many traits beyond shape and size contribute to pitcher function (e.g., volatiles, pigmentation, 397 external nectaries, pubescence), and we predict that these traits may show similar patterns of 398 convergence and correlated evolution. While many studies have demonstrated phenotypic and 399 even genomic convergence associated with carnivory across angiosperms (e.g. Givnish et al. 400 1997; Fukushima et al. 2017; Bittleston et al. 2018; Clarke et al. 2018), our study further 401 contributes to this understanding by demonstrating that the diversification and convergent 402 evolution of carnivory-related traits can occur on finer evolutionary timescales. The radiation of 403 pitcher forms in Sarraceniaceae, including Heliamphora, reinforces the notion that extreme 404 ecological specialization, such as the carnivorous lifestyle, may not necessarily be an evolutionary dead end (Tripp and Manos 2008; Wim A. Ozinga 2012; Day, Hua, and Bromham 405 406 2016; Thorogood, Bauer, and Hiscock 2018).

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- 408

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423 Author Contributions

424

- 425 Sukuan Liu: Conceptualization, Data curation, Formal analysis, Funding acquisition,
- 426 Investigation, Methodology, Project administration, Resources, Software, Validation,
- 427 Visualization, Writing original draft, Writing review & editing. Stacey Smith:
- 428 Conceptualization, Formal analysis, Investigation, Methodology, Project administration,
- 429 Resources, Supervision, Writing review & editing.
- 430

431 Data Availability

- 432 Raw pitcher angle measurements, pitcher shape outlines, modified *Heliamphora* phylogeny, and
- 433 R Markdown file containing scripts used in this study can be accessed via "Figshare"
- 434 (<u>https://doi.org/10.6084/m9.figshare.23646414.v3</u>).
- 435

436 Supporting Information

- 437 Additional Supporting Information may be found online in the supporting information section at
- 438 the end of the article (Appendix S1, Table S1: Sampling information; Appendix S2, Table S2:
- 439 Model fit results; Appendix S2, Figure S1: Pitcher Angle Measurements; Appendix S2, Figure
- 440 S2: Pitcher Angle by Taxa; Appendix S2, Figure S3: Associations between Adult Pitcher Shapes
- 441 Variations and Pitcher Angle in Different Crowding Conditions; Appendix S2, Figure S4: Adult
- 442 Pitcher Morphosapce of PC1 and PC3; Appendix S2, Figure S5: Developmental Timeline of
- 443 *Heliamphora* Adult Pitcher Rosettes Associated with Wider and Sharper Angles; Appendix S3:
- 444 Morphospace of PC1, PC2, and PC3; Appendix S4: Morphospace of PC1, PC3, and Maximum445 Pitcher Size).
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664 Figure Legends

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667 668 Figure 1. Left: Adult Pitcher Morphologies of the reconstructed ancestral 669 Heliamphora (ANC) and extant species grouped by clades. Right: Phylogeny of 670 Heliamphora adapted and modified from Liu and Smith, 2021. Taxa names in each 671 clade are ordered aesthetically from left to right. W: H. hispida, H. ceracea, H. tatei, H. 672 neblinae, H. macdonaldae, and H. parva. E1: H. pulchella, H. minor var. pilosa, H. 673 ciliata, H. huberi, and H. chimantensis. E2a: H. heterodoxa, H. collina, H. sp. 674 "Angasima", H. sp. "Akopán", H. sarracenioides, and H. purpurascens. E2b: H. 675 exappendiculata, H. glabra, H. uncinata, and H. folliculata. E3: H. nutans, H. arenicola, 676 H. ionasi, and H. elongata. The ancestral Heliamphora pitcher morphology was 677 generated based on estimated ancestral values for each of the three shape PCs and 678 the estimated ancestral size and maximum pitcher size. Note that the ancestral pitcher 679 is based on the expected values at the root, which carry a degree of uncertainty. 680 681 Figure 2. Landmarking examples of three Heliamphora species: H. ciliata (left), H. 682 pulchella (middle) and H. sarracenioides (right). The adult pitcher functional zones 683 (McPherson et al. 2011) were delineated by six landmarks indicated above. 684 Morphologically, *Heliamphora* pitchers are similar to those of *Sarracenia* and 685 Darlingtonia, except that they lack a covering at the pitcher openings to prevent 686 rainwater from overfilling the pitchers. Instead, Heliamphora evolved the drainage hole 687 (further modified into drainage slit in E1 clade, indicated by landmark 1) to allow 688 rainwater to accumulate in the pitcher (thus maintaining a high level of pitcher fluid. 689 with maximum fluid level shown with the dotted blue line and roughly align with 690 landmark 1 and 5) while letting the excess to drain out and prevent the pitchers from 691 becoming overfilled and unstable (Bauer et al. 2013). 692 693 Figure 3. A. Adult pitcher morphospace of pitcher curvature (PC1) and nectar 694 **spoon elaboration (PC2)** with outlines of each taxa and the reconstructed ancestral 695 Heliamphora adult pitcher (darker gray). Taxa are colored by clades: W (orange), E1 696 (red), E2a (blue), E2b (green), E3 (purple). B. Share variation along PC axes. For 697 each of the first three PCs, the mean is shown along with shapes corresponding to ±2 698 standard deviation (SD). + 2 SD was chosen to better visualize the morphological 699 variation along shape PC axes. 700 701 Figure 4. Traitgrams of pitcher curvature (PC1), nectar spoon elaboration (PC2), 702 pitcher stoutness (PC3), and maximum pitcher size. Branches are colored by 703 clades. Examples of pitcher and nectar spoon morphologies are shown in the

- 704 traitgrams.
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708 Figure 5. Associations between adult pitcher shape variations and maximum

709 **pitcher size in pitcher morphospace.** Taxa are represented by solid circles and

- colored by clades. Regression lines were drawn based on the best fit phylogenetic
- generalized linear models. Example pitcher shapes (corresponding to the taxa
- encircled) are shown in each morphospace.
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714 Figure 6. Effect of crowdedness on pitcher angle across developmental stages.

715 Both phylogenetic and simple unpaired t-tests were conducted for each comparison with

test statistics shown in black and gray, respectively. Note the angles shown above each

condition were averages pooled from all samples, with potential bias toward species

- that were over sampled.
- 719

720 Figure 7. Morphologies of *Heliamphora* rosettes associated with wider (A, B) and

721 sharper pitcher angles (D, E). For each pitcher angle, rosettes are illustrated with

three (A, D) to six pitchers (B, E) to demonstrate the effect of shading from overlapping

pitchers. In each rosette, pitchers are numbered increasingly from the oldest to the

youngest pitcher (most active pitcher). Hypothetical scenario illustrating the effect of

725 wider (C) and sharper pitcher angles (F) on optimal packing in crowded

726 **conditions.** In this hypothetical scenario, the wider and sharper pitcher angles,

respectively, allowed the populations to fit in up to 10 and 12 rosettes when space is

128 limited. The sharper pitcher angle allowed *Heliamphora* population to occupy available

space more efficiently, with an increase of 20% more active pitchers.