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Influence of plant density and growth habit of common bean on leaf area development and N accumulation

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1	Influence of plant density and growth habit of common bean on leaf area
2	development and N accumulation
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22 ABSTRACT

- Crop yield requires leaf area to intercept solar radiation and to undertake photosynthesis, both of which depend on nitrogen (N) accumulation. Further, the amount of accumulated plant N at the beginning of seed fill serves as the reservoir for N required in synthesizing the proteins in developing seeds. For common bean <u>(Phaseolus vulgaris L.)</u>, resolution of the basic characteristics limiting production is challenging due tobecause of variation in plant growth-habit and in wide ranging plant spacing. Field experiments were undertaken at two low-latitude
- 29 locations with three plant growth-habit types and six plant densities to measure canopy leaf area
- and leaf N accumulation at the beginning of seed fill. Plant spacing of 20 plants m⁻² or more
- 31 was sufficient to result in equal leaf area and N accumulation for all six plant types at each
- 32 location. However, the low-altitude, higher-temperature location had lower accumulated leaf N
- and yield than the high-altitude, cooler-temperature location. These results indicate attention
- 34 needs to be given to physiological or agronomic approaches to overcome the negative impact of
- 35 high temperature on N accumulation by common bean.

36 Introduction

37 Leaf area development and nitrogen (N) accumulation in leaves prior to the initiation of 38 seed fill are each both essential processes impacting, respectively, the interception of solar radiation and the synthesis of photosynthesis components in the leaves to support high carbon 39 40 accumulation rates. The balance between the use of newly acquired N between formation of 41 new leaf area and accumulation of leaf N concentration to support photosynthesis influences the productivity of a crop (Sinclair and Horie, 1989). Further, once seed-fill is initiated, N in the 42 43 leaves and stems is transferred to the seeds contributing directly to yield formation. The total 44 amount of the N available in the leaves for transfer to the seeds can be quantitatively linked to seed yield as part of the 'self-destruction' process during seed growth (Sinclair and deWit, 45 1976). In common bean (Phaseolus vulgaris L.), Saberali et al. (2016) found a linear 46 47 relationship between seed yield and accumulated N.

Evaluating the use of N to support leaf area development and in-increaseing leaf N 48 49 concentration is challenging in common bean because of the confounding influence of differences in growth habit among genotypes and a range of plant densities under which the 50 crop is grown. In particular, four distinct growth habits have been identified in common bean: 51 52 type I, determinate bush; type II, indeterminate upright; type III indeterminate prostrate; and 53 type IV indeterminate climbing (Singh, 1981). Kueneman and Wallace (1979a) found leaf area index of common bean was to be lower in Type I genotypes than in Types II and III. They also 54 included plant density in their experiments, and found lower leaf area index at a plant density of 55 56 13.5 plants m⁻² than for canopies grown at densities of 18 and 38 plants m⁻². Their conclusion 57 was that "leaf area index was positively correlated with yield" at the highest plant density but the 58 correlation was more variable at lower plant densities.

In a subsequent study, Nienhuis and Singh (1985) found similar results. Type I 59 genotypes had fewer main stem nodes, i.e., fewer main stem leaves, and lower yield than Type 60 61 II and III. Maximum seed yield was not achieved in their study until plant density was 22 plants 62 m⁻² or more. In our previous report (Ricaurte et al., 2016) during the development of main stem node number and leaf area, leaf area index of two genotypes of Type I was slightly less than 63 that of genotypes of Types II and III. The node number increased linearly with cumulative 64 65 temperature units, and leaf area index increase was predicted to increase exponentially with 66 increase in node number.

While several studies have documented N in leaves at the beginning of vegetative
growth, the results were only for leaf concentration, not <u>for</u> total accumulated N (Soratto et al.,
2017; Nascente et al, 2017; Pias et al., 2017; Barros et al., 2018). Further, these studies were

70 often focused on a single cultivar or a single plant density so the impact of these variables even 71 on leaf N concentration could not be fully evaluated. In the study by Saberali et al. (2016), for 72 example, the plant density was 40 plants m⁻², which is high for many cases of common bean 73 production. Therefore, the objective of the current study was to compare at the end of 74 vegetative development leaf area and leaf N amount of six common bean genotypes representing three growth habits grown at six plant densities. A key analysis of this study was 75 76 the relationship between accumulated leaf N prior to seed fill and final seed yield across 77 genotypes and plant densitiesy. 78 79 Materials and methods 80 81 Experimental Designdesign 82 Two field experiments as previously described by Ricaurte et al. (2016) were conducted in 83 southwestern Colombia. The first experiment was sown in a fertile mollisol (Typic Pellustert) soil 84 at the Centro Internacional de Agricultura Tropical (CIAT) in Palmira-Valle del Cauca (999 masl, 85 86 3.54° N, 76.30° W) on 13 October 2013, whereasile the second experiment was sown in an inceptisol (Typic Fulvudands) soil at an experimental station in Darién-Valle del Cauca (1570 87 masl, 3.92° N, 76.60° W) on 29 May 2014. Prior to sowing, seeds were inoculated with 88 Rhizobioium tropici CIAT 899 and treated with fungicide. Application of 60 kg ha⁻¹ phosphorus 89 was applied made to the soil at both locations at sowing. Non-limiting management was used 90 91 throughout the experiment, including irrigation. 92 Two genotypes were selected for the experiment from each of growth habit types I, II, 93 and III, so that six genotypes were included in the study. Table 1 gives contains information on 94 origin, growth habits habit- and seed size of the studied genotypes. 95 At both locations, a completely randomized block design was employed with treatments arranged as split plots, where sowing density (5, 10, 15, 20, 25, 35 plants m⁻²) and the six 96 97 genotypes were main and subplots, respectively. Individual plots were 4-m long, with 7 rows at 98 Darien and 8 rows at Palmira. The rows were spaced 0.6 m apart. There were three replicates at Darien and four replicates at Palmira. The plots were irrigated as needed. 99 100 101 Data collection

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Visual observations were made every 2 to 3 d on phenological development to assess whether 103 104 the plant density treatments impacted the duration to reach initiation of seed fill. The initiation of 105 seed fill was identified as Developmental Stage 8, i.e., p: Pod filling, which begins when 50% of 106 the plants in a plot are observed to be filling seeds in the first pod (Fernandez et al., 1986). To 107 assess a possible difference in development between the two experimental locations, 108 cumulative temperature units from date of sowing were calculated as the basis for comparing 109 occurrence of stage R8. Cumulative temperature units were obtained by summing daily 110 temperature units, which were determined using a Beta function described by Yan and Hunt (1999). The base, optimum, and critical temperatures in these calculations were set at 10, 25 111 112 and 36 °C, respectively, based on data from Wallace et al. (1991). 113 Destructive sampling of 0.3 m² area was done biweekly within all plots (Ricaurte et al.,

2016). For each plant, mainstem node number, branch number, leaf number, leaf area, and leaf weight were measured. In the results presented here, only data collected at the harvest prior to seed fill are presented.

117 A final harvest for seed yield was done at pod maturity. A length of 3 m was harvested 118 from two adjacent rows in the center of each plot for a total harvest area of 3.6 m^2 . Seeds were 119 removed from the pods and weighed. The seed water content was measured (MT-16 Grain 120 Moisture Tester, Agratronix, Streetsboro, OH, USA) and seed weight adjusted to 0.14 g H₂O g⁻¹ 121 mass.

122 For measurement of leaf N content, leaf samples excluding petioles were processed at the University of Florida's Forage Evaluation Lab. Ground samples were digested using a 123 124 modification of the aluminum block digestion procedure of Gallaher et al. (1975). N analysis 125 was done by semi-automated colorimetry using the procedure described by Hambleton (1977). Results were expressed as leaf N per unit leaf area (g N m⁻², i.e. specific leaf N) by multiplying 126 N mass concentration of N (g N g⁻¹ mass) by the ratio of leaf mass to leaf area for each sample. 127 Total plant leaf N per unit ground area (g N m⁻²) was calculated by multiplying leaf N per unit 128 leaf area by leaf area index. 129

131 Data analysis

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The effects of density, sites and genotypes on all variables were explored using data collected from all six genotypes that were studied. Linear models were used to conduct an analysis according to a split-plot design with the whole-plot factor arranged in a randomized complete block design using the *nlme* package (Pinheiro and Bates, 2000) in the R statistical

software (R Core Team, 2015), following the guidelines in Zuur et al. (2013). In the analysis, 137 138 density was used as a factor (Density f, as a categorical variable with 6 levels) or as continuous covariate centered at the 25 plant m⁻² level (Density.c) to facilitate model parameter 139 140 interpretation and reduce collinearity between interacting model terms (Schielzeth, 2010). 141 Initially, a full model was fitted that included random effects of site, density, and genotype within 142 site and fixed effects for Site, Density.f and Genotype and their interactions. 143 The full model for each variable was then used to identify the appropriate variance structure within nlme, using Akaike's information criteria (AICc) corrected for sample size (AICc) 144 to select the most parsimonious form (data not shown), calculated using the AICcmodavg 145 146 package (Mazerolle, 2011). Model selection was done using ΔAICc values, which were calculated as the difference between the model with lowest AICc and every other model. 147 Assessment of these differences indicated that those models having $\Delta AICc < 2$ having strong 148 support, those with $\Delta AICc$ between 4-7 substantial support, and any model with $\Delta AICc > 10$ not 149 considered (Burnham and Anderson, 2002). The final variance structure selected had an 150 exponential or a power function of the fitted values to model the increasing variances with 151 152 increasing node numbers by site. In a second step, a full model without random terms using 153 generalized least squares was fit and compared using likelihood ratio tests to determine if the random terms were necessary in the model. In the current study, only variable specific leaf N 154 155 was analyzed using the random term formulation described previously. 156 With the appropriate variance structure, the final fixed-effect or regressor structure was

157 selected by fitting all possible models with 2^{nd} and 3^{rd} -order interactions and main effects, with 158 density either as a factor or continuous, and genetic effects by genotype, growth habit or 159 determinacy levels. Model selection was carried out using the procedure described previously 160 based on Δ AlCc values, and the top three models reported in each case. Finally, goodness-of-fit 161 measure root mean squared error (RMSE) were was also estimated for each of the final 162 models.

163

164 Results

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The environments at the two sites were different as expected. At the higher altitude of Darien, minimum and maximum temperatures were cooler and solar radiation during the growing

168 season was higher than at Palmira (Table 2). For Type I genotypes, the same cumulative

169 temperature at both locations was required to reach Stage 8. However, for Type II and Type III

170 genotypes Stage 8 was delayed by about an additional 100 °C cumulative temperature units at

Palmira as compared <u>with</u> Darien. A result of the difference between the two locations in
cumulative temperature units during the vegetative stages of <u>type-Type</u> II and <u>Type</u> III growth
habite plants was that the plants had about two more nodes at Stage 8 at Palmira than at
Darien.

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176

177 Leaf area

178 Comparison of leaf area development at the beginning of seed fill across growth habits and 179 plant density densities was a major objective of this study. As illustrated in Figure 1, with one 180 genotype for each of the three growth habits, the results for the two locations were quite similar. Leaf area per plant decreased with increasing plant density for each of the genotypes. Leaf 181 182 area per plant was highest at a plant density of 5 plants m⁻², with mean values greater than 0.3 183 m^2 plant¹. At the lower plant densities, leaf area per plant tended to be greater for types Types 184 II and III than for type Type I. At higher plant densities the leaf area per plant decreased linearly 185 with increasing plant density. All genotypes at the highest plant density had similar plant leaf 186 areas of less than 0.1 m² plant⁻¹. Model selection indicated the most parsimonious model for 187 plant leaf area included interaction terms for of Site × Density, f, Growth Habit × Density, f and Site x Growth Habit and their respective main effects. 188 189 Leaf area per unit ground area, i.e., leaf area index, increased from approximately 1.5 at 190 low plant densities to mean values of about 2.5 to 2.8 at high plant densities (as illustrated in

Figure 2). An increase in leaf area index of less than two-fold represents much less variation in leaf area index across plant densities than found for plant leaf area. At plant densities of 20 plants m⁻² and greater, leaf area was approximately about constant. Leaf area index was slightly greater for growth habit types Types II and III as compared withte type Type I in these Stage R8 data. Model selection indicated the best model for leaf area index included interactive effects of Site × Density.f + Site × Genotype, and all lower-order terms (Table 3).

Leaf weight per <u>unit</u> ground area at both locations was lowest at the 5 plants m⁻² density but tended to be stable at sowing densities of 10 plants m⁻² and above (Figure 3). However, leaf weight per ground area was different between the two locations. For growth habit types II and III, the leaf weights per ground area for the high densities at Palmira were about 60 to 70 g m⁻² and at Darien about 100 to 110 g m⁻². The leaf weight per ground area was slightly less for the type I genotypes as compared to types II and III. Model selection indicated the most

parsimonious model for plant leaf weight included genotype, Site and Density.f main effects, as
well Site × Density.f interaction term (Table 4).

206 Leaf N

207 Except for differences between locations, there was general stability in leaf N per unit leaf area 208 across growth habit types and plant density (Figure 4). However, leaf N per unit leaf area was somewhat elevated for the 5 plants m⁻² density at Darien as compared to-with the other plant 209 210 densities at that location. Mean leaf N per unit leaf area at Darien was commonly about 2.0 g N 211 m⁻², while whereas at Palmira it was roughly about 1.3 g N m⁻². The difference between the two 212 locations in leaf N per unit leaf area for each plant density was generally in the range of 0.4 to 213 1.0 g N m⁻². Model selection criteria indicated that a model containing main effects terms for 214 Site, Density.f and Growth Habit, as well as a-for_Site × Growth Habit interaction was the most 215 parsimonious.

Given the approximate stability in the leaf N per unit leaf area at each location at plant 216 density of 10 plants m⁻² and above, the pattern in total leaf N per unit ground area (Figure 5) is 217 roughly the same as the pattern in leaf area index. That is, the two lowest plant densities (5 and 218 10 plants m⁻²) tended to have somewhat lower leaf N per unit ground area than the high plant 219 220 densities. At the high plant densities at each location, N per unit ground area had fairly stable 221 values. Differences in leaf N per unit ground area existed between the two locations with an 222 average mean of 2.11 g N m⁻² lower at Palmira as compared to with Darien. This is in spite of the fact that the vegetative period for Type II and III genotypes was 100 TU longer at Palmira 223 224 than at Darien. Model selection criteria indicated that a model containing main effecte terms for 225 Site, Density f and Growth Habit, as well as a Site x Growth Habit interaction was the most 226 parsimonious (Table 5).

227 228 Discussion

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Carbon assimilation is required for seed development and the production of crop yield, and is
dependent of the interception of solar radiation by the leaves and the photosynthetic activity of
the leaves. However, in common bean these two variables associated with carbon
accumulation have not been found to be consistently associated with yield. Kueneman and
Wallace (1979a) only found a consistent correlation between yield and leaf area in common
bean at a 38 plants m⁻² density. In the study by Kueneman and Wallace (1979b), ILeaf
photosynthesis rate measured at one time during pod filling showed no correlation with yield for

the growth habits and plant densities tested by Kueneman and Wallace (1979b).

238 N accumulation by a crop is also essential in for producing yield because both leaf area 239 development and photosynthesis rate are dependent on N availability in the plant, and crucially, during yield formation N is quantitatively required in the formation of the components of the 240 241 growing seeds. No previous studies have, however, investigated N accumulation in common 242 bean to offer information about the N resource needed in seed yield formation. The lack of 243 insight in common bean about the N influence on yield may have resulted because of the challenge to account for the influence of plant growth habit and plant density, which are two 244 common variables in common bean production. This study was targeted to explore the variation 245 in leaf area development and N accumulation at the beginning of seed fill by genotypes 246 247 representing three growth types of common bean and plant densities of 5 to 35 plants m⁻². 248 The results of leaf N measurements indicated a sensitivity of leaf N accumulation to location. While leaf N per unit leaf area was fairly stable across all plant densities (Figure 4), 249 leaf N per unit leaf area at Palmira was less than that at Darien. The explanation of this 250 difference in leaf N concentration is likely to result from differences in N accumulation rates 251

252 between the two sites.

253 Figure 6 shows tThe correlation between final seed yield and the amount of leaf N per 254 unit ground area accumulated prior to seed fill is shown in Figure 6. The results for Palmira 255 emphasize the importance of leaf N accumulation per unit ground area being positively associated with increasing yield. That is, low leaf N accumulation per unit ground area was 256 257 associated with low yield. The correlation between increasing yield and accumulated N per unit ground area is consistent with the results reported by Saberali et al. (2016) for an experiment in 258 259 which N fertilizer treatment accounted for differences in accumulated N. In the experiment at 260 Palmira, results from low plant density were the source of low leaf N per unit ground area(<3 g 261 N m⁻²) and low yield. On the other hand, at Darien leaf N accumulation per unit ground area was greater than about 3 g N m⁻² at all plant densities and yield was stable with increasing 262 263 accumulation of leaf N per unit ground area. These results indicate that a threshold of N accumulation per unit ground area is required to maximize yield. 264

There are two possible bases for the difference in N accumulation between sites. One basis may be the lower soil organic matter content at Palmira than at Darien so the smaller reservoir for soil uptake may have provided less N to developing leaves. A-<u>The</u> second basis for the lower leaf N accumulation at Palmira may result from the temperature difference between the two locations and possible sensitivity of soil N accumulation and/or symbiotic N₂ fixation in common bean to high temperature. Palmira is a lower altitude location with an

271 average maximum temperature of 31.1 °C during the experiment (Table 2) as compared to-with 272 average maximum temperature of 25.5 °C at Darien. 273 Published experiments reports have showned decreases in N2 fixation rates in 274 Phaselous when temperature exceeded about 30 °C. Pankhurst and Sprent (1976) found maximum N₂ fixation rate $\frac{1}{2}$ French bean occurred at 20 °C and virtually no N₂ fixation was 275 found when temperature was increased to 35 °C. Hernandez-Armenta et al. (1989) found N2 276 277 fixation rate at 32 °C was substantially less than the rate at 26 °C. Similarly, Piha and Munns 278 (1987) found a very large decrease in N₂ fixation rate by common bean at 34 °C daytime 279 temperatures as compared to-with 28 °C. Therefore, high temperature a Palmira may have 280 resulted in less N₂ fixation, low leaf N per unit leaf area (Figure 4), low leaf N accumulated per unit ground area (Figure 5), and lower yield (Figure 6). 281 282 Overall, growth habit type had little impact on the leaf area index and plant N content at 283 the initiation of seed fill for cases where when plant density was 20 plants m⁻² or more. Therefore, growth-habit type based on these field studies did not appear to be a major factor for 284 285 direct physiological improvement for in carbon and N accumulation in common bean as long as 286 plant density was at least 20 plants m⁻². However, a major finding from these field results 287 experiments was the deficiency in N accumulation at Palmira as compared to-with Darien was associated with lower seed yields. These results indicate attention may need to be given to the 288 289 negative impact of high temperature on N accumulation by common bean, at least for the six 290 genotypes included in this study. Lower leaf N per unit area is likely associated with lower leaf photosynthesis rate and radiation use efficiency (Sinclair and Horie, 1989). Crucially, the lower 291 292 total leaf N means that less N is available for transfer to the seeds during seed-fill to achieve 293 higher vields (Sinclair and deWit, 1976). 294 295 296 References Barros RLN, Oliveira LBD, Magalhaes WBD, Pimentel C. 2018. Growth and yield of common 297 298 beanas affected by seed inoculation with rhizobium and nitrogen fertilization. Exp. Agric. 299 54, 16-30. Burnham KP, Anderson DR. 2002. Model Selection and Multimodel Inference: A Practical 300 301 Information-Theoretic Approach. Springer-Verlag, NY. Fernandez F, Gepta P, Lopez M. 1986. Stages of development of the common bean plant. 302 Centro Internacional de Agricultura Tropical (CIAT), Cali, Columbia. 32p. 303

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352 **Table 1.** Information on common bean genotypes sown in both experiments.

Genotype	Gene pool	Growth habit	Seed size (g 100 seeds ⁻¹)
CAL 96	Andean	I	Large (56.5)
ICA Quimbaya	Andean	I	Large (47.3)
DOR 364	Mesoamerican	Ш	Small (20.4)
SER 118	Mesoamerican	II	Medium (29.2)
G 21212	Mesoamerican	Ш	Medium(29.8)
Carioca	Mesoamerican	Ш	Small (23.4)

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356 Table 2. Average minimum and maximum temperatures, daily temperature units (DTU), and

solar radiation during the experimental period at the Darien and Palmira, Colombia, study sites.

		358	Standard err	or of the me	ean is given	i in paranth	esisparenthesis.
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	Temperatur	e	DTU	Solar Radiation		
Site	Min (°C)	Max (°C)	(°C)	(MJ m ⁻² day ⁻¹)		
Darein	16.6 (0.92)	25.5 (1.2)	12.5 (0.6)	18.6 (3.1)		
Palmira	19.1 (0.93)	31.1 (1.9)	14.6 (0.4)	14.2 (3.2)		

359

360

361 Table 3. Model selection table for leaf area index at the onset of seed fill (R8). The columns

362 present number of estimated parameters for each model (K), Akaike's information criteria

- 363 corrected for sample size (AICc), difference in AICc of model with lowest AICc and other
- 364 modelts (ΔAICc), Akaike's weights (wAICc), log likelihood (LL), and root mean square error
- 365 <u>(RMSE).</u>

-

Form	κ	AICc	ΔAICc	wAICc	LL	RMSE
Site*Genotype + factor(Density)*Site	24	470.51	0	0.95	-208.61	0.62
Site*Genotype + factor(Density)	19	477.84	7.33	0.02	-218.28	0.63
Site*factor(Density) + Genotype	19	478.26	7.75	0.02	-218.49	0.65

366

369	Table 4. Model selection table for plant leaf weight per unit leaf area at the onset of seed fill
370	(R8). The columns present number of estimated parameters for each model (K), Akaike's
371	information criteria corrected for sample size (AICc), difference in AICc of model with lowest
372	AICc and other modelts (ΔAICc), Akaike's weights (wAICc), log likelihood (LL), and root mean
373	square error (RMSE).

Form	Κ	AICc	ΔAICc	wAICc	LL	RMSE
Site*factor(Density) + Genotype	18	2264	0	0.76	-1112.5	24.9
Site*factor(Density) + Growth.Habit	15	2267.7	3.78	0.12	-1117.9	25.1
Site*factor(Density) + Site*Genotype	23	2268.7	4.71	0.07	-1108.9	24.6

 Table 5. Model selection table for plant leaf N weight per unit ground area at the onset of seed

fill (R8). The columns present number of estimated parameters for each model (K), Akaike's

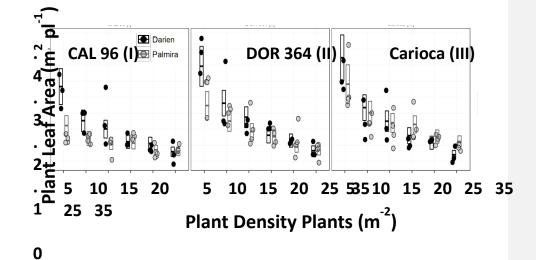
information criteria corrected for sample size (AICc), difference in AICc of model with lowest

AICc and other modelts (ΔAICc), Akaike's weights (wAICc), log likelihood (LL), and root mean square error (RMSE).

Form	Κ	AICc	∆AICc	wAICc	LL	RMSE
Density.f + Site × Growth.Habit	12	784.73	0	0.54	-379.71	1.35
Density.f + Site × Genotype	18	787.23	2.5	0.15	-374.15	1.34
Density.f + Site × Determinacy	10	787.92	3.19	0.11	-383.5	1.37

Figure 1. Leaf area per plant at each plant density for one genotype of each growth habit type at both experimental locations. The datum for each replicate is presented. The line in the bars represents the estimate for the value from the best fit of the model and the bars are the

389 standard error of this value.



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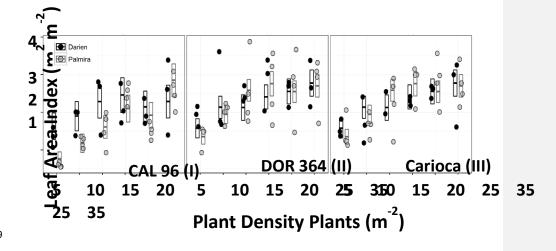
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393

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Figure 2. Leaf area index at each plant density for one genotype of each growth habit type at both experimental locations. The datum for each replicate is presented. The line in the bars represents the estimate for the value from the best fit of the model and the bars are the

398 standard error of this value.



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401

402 Figure 3. Total leaf weight per ground area at each plant density for one genotype of each

403 growth habit type at both experimental locations. The datum for each replicate is presented.

404 The line in the bars represents the estimate for the value from the best fit of the model and the

405 bars are the standard error of for this value.

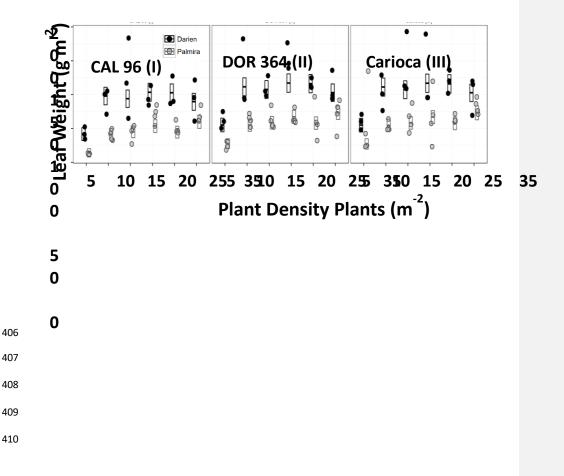
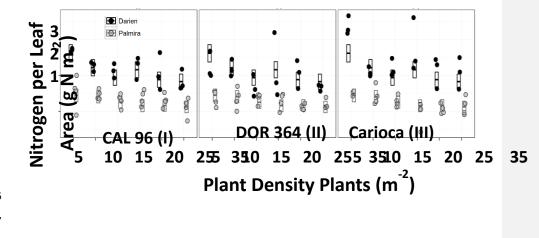


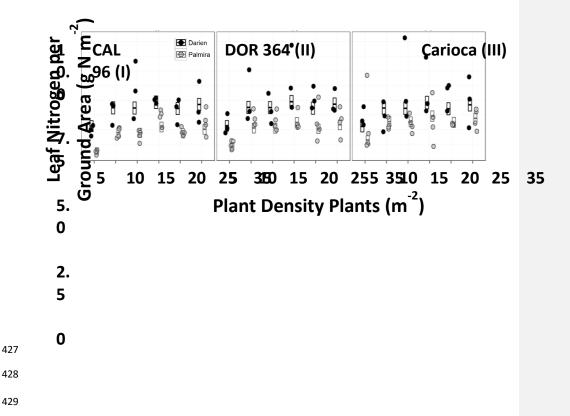
Figure 4. Leaf N per leaf area at each plant density for one genotype of each growth habit type at both experimental locations. The datum for each replicate is presented. The line in the bars represents the estimate for the value from the best fit of the model and the bars are the

414 standard error of for this value.



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Figure 5. Total leaf N per ground area at each plant density for one genotype of each growth habit type at both experimental locations. The datum for each replicate is presented. The line in the bars represents the estimate for the value from the best fit of the model and the bars are the standard error of for this value.



430 Figure 6. Graph of final bean yield vs. accumulated leaf N per ground area for each of the two

431 locations and six genotypes.

