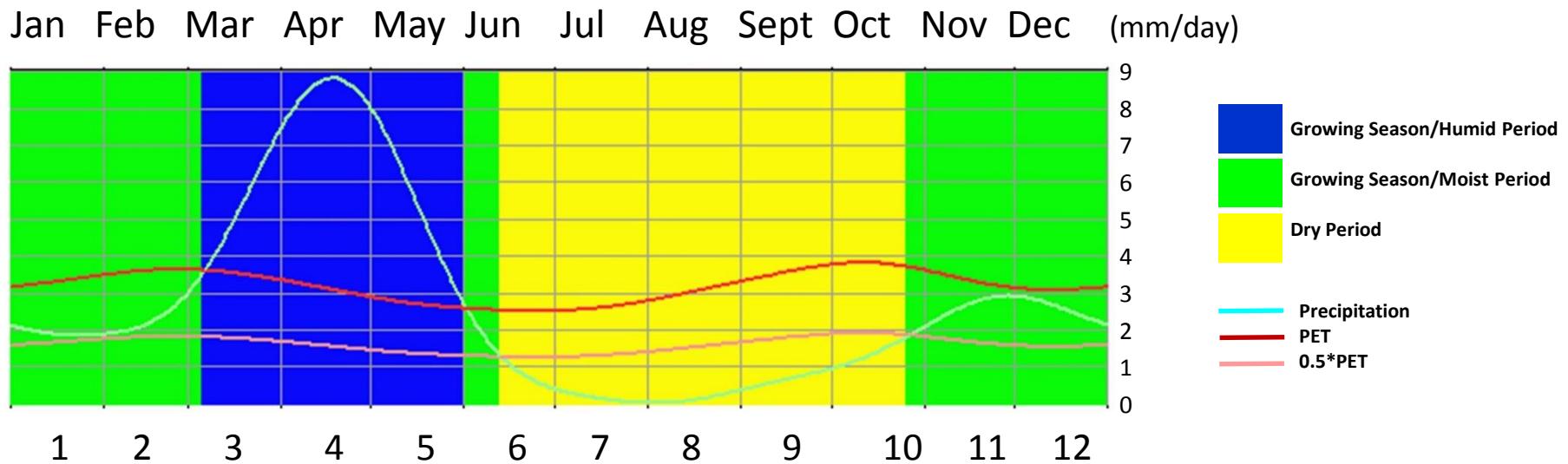


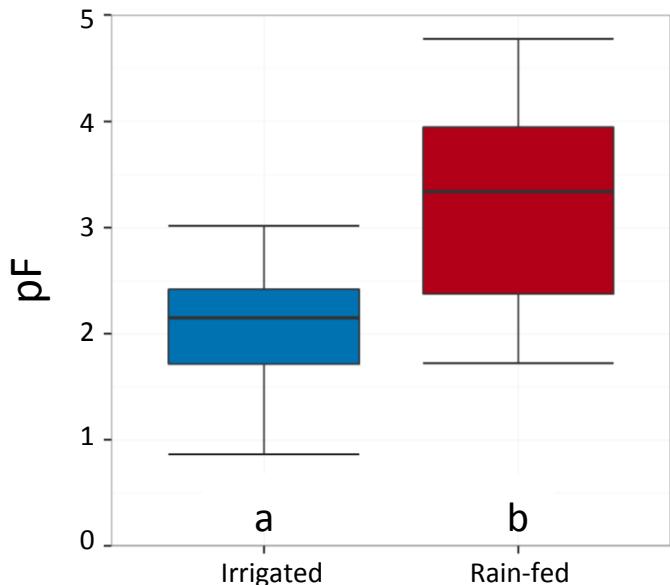
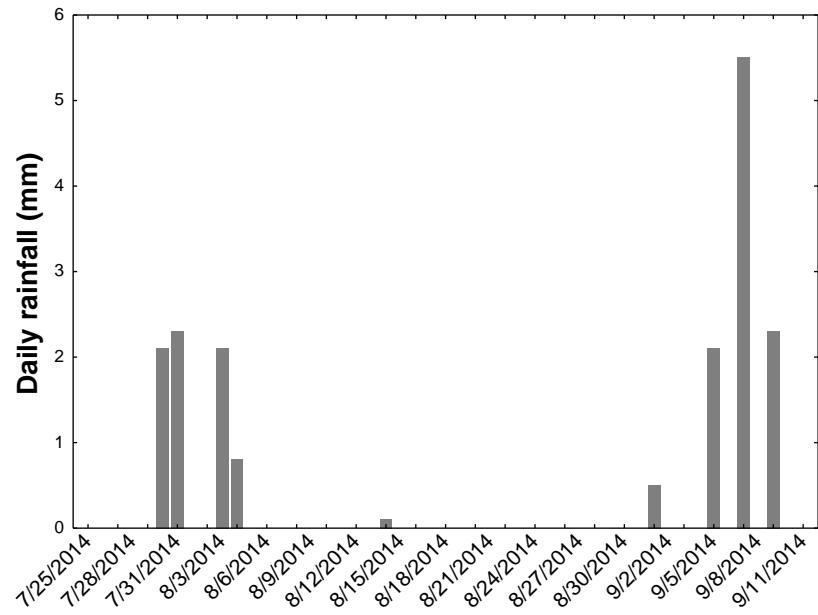
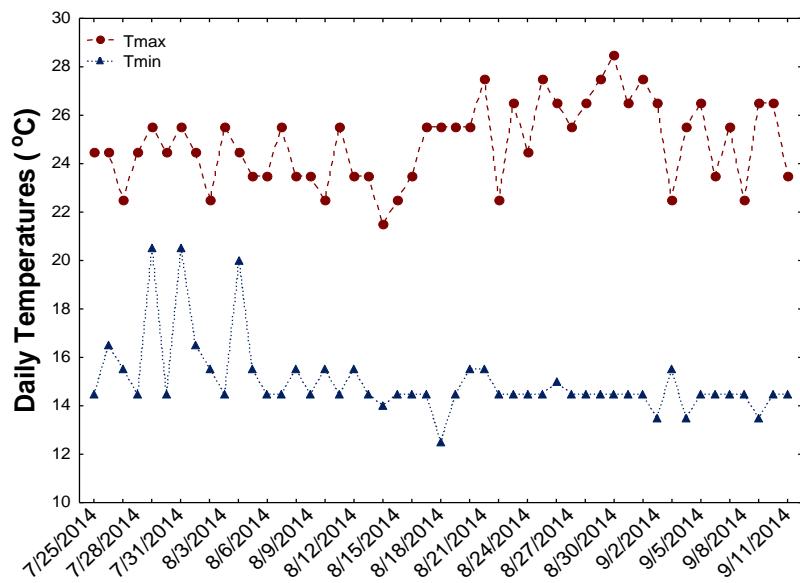
# Supplementary Information

Effect of seasonal drought on the agronomic performance of four banana genotypes (*Musa* spp.) in the East African highlands

**Brigitte Uwimana, Yasmín Zorrilla-Fontanesi, Jelle van Wesemael, Hassan Mduma, Allan Brown,  
Sebastien Carpentier and Rony Swennen**



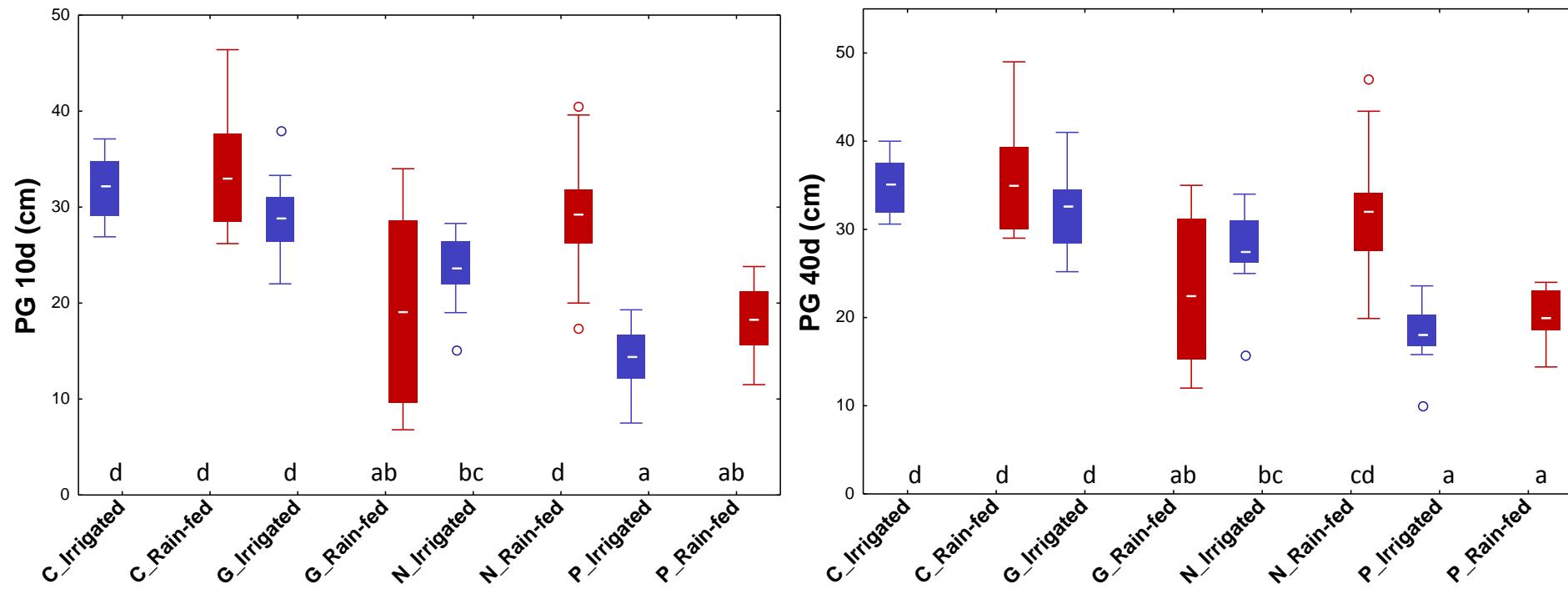
**Supplementary Figure S1.** Vegetation period for Arusha (Tanzania). Source: New\_LocClim ([http://www.fao.org/nr/climpag/pub/en3\\_051002\\_en.asp](http://www.fao.org/nr/climpag/pub/en3_051002_en.asp)). Data collected over the last 30 years, as accessed on May 31<sup>st</sup>, 2019. PET: potential evapotranspiration.

**A****B****C**

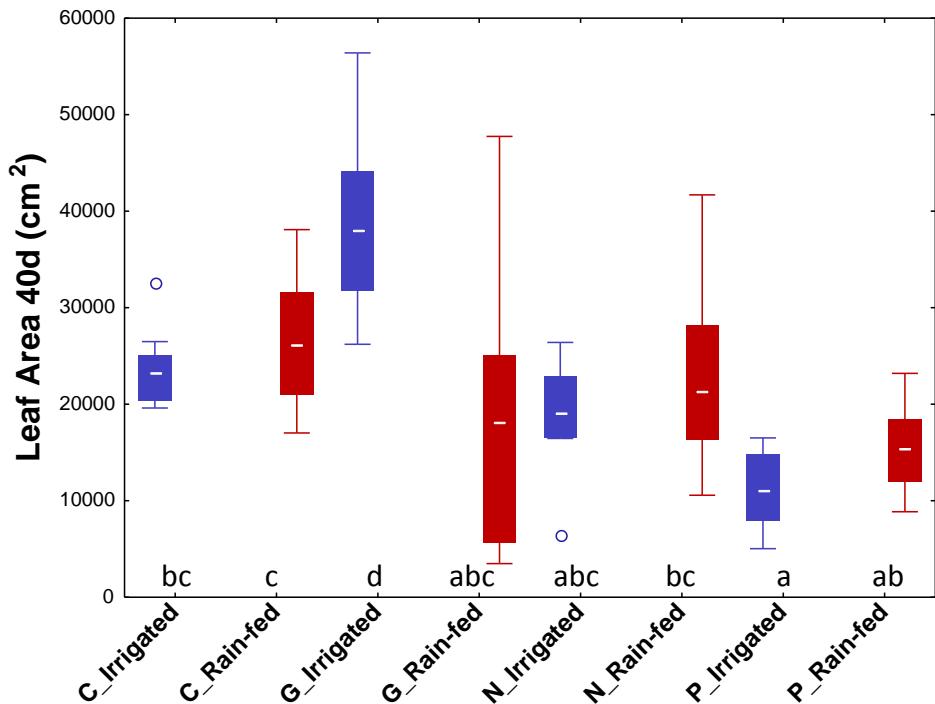
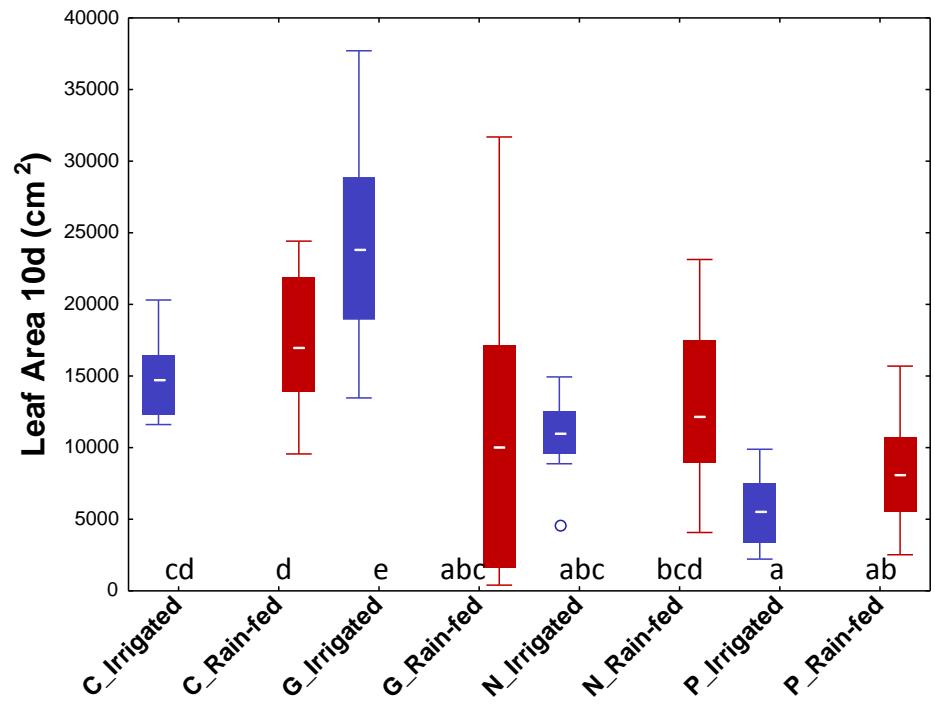
**Supplementary Figure S2.** **A)** Boxplot for soil water retention (pF values) of irrigated vs. rain-fed plots measured between 30 and 45 days after stopping irrigation. Different letters indicate significant differences based on t-test (a-b,  $\alpha=0.05$ ). N=8/8 (irrigated/rain-fed). **B-C)** Weather data between 25<sup>th</sup> July (stop of irrigation) and 11<sup>th</sup> September (48 days after stopping irrigation), 2014: B) daily cumulative rainfall, C) daily temperatures (maximum, and minimum).

**Supplementary Table S1.** Results of the analysis of variance (ANOVA) on early plant growth parameters and mean leaf temperature measured at 10, 40 and 47-48 days after stopping irrigation. <sup>1</sup>Overall residual; \* $p\leq 0.05$ , \*\*\* $p\leq 0.001$ , ns: not significant ( $p>0.05$ ); d: days after stopping irrigation.

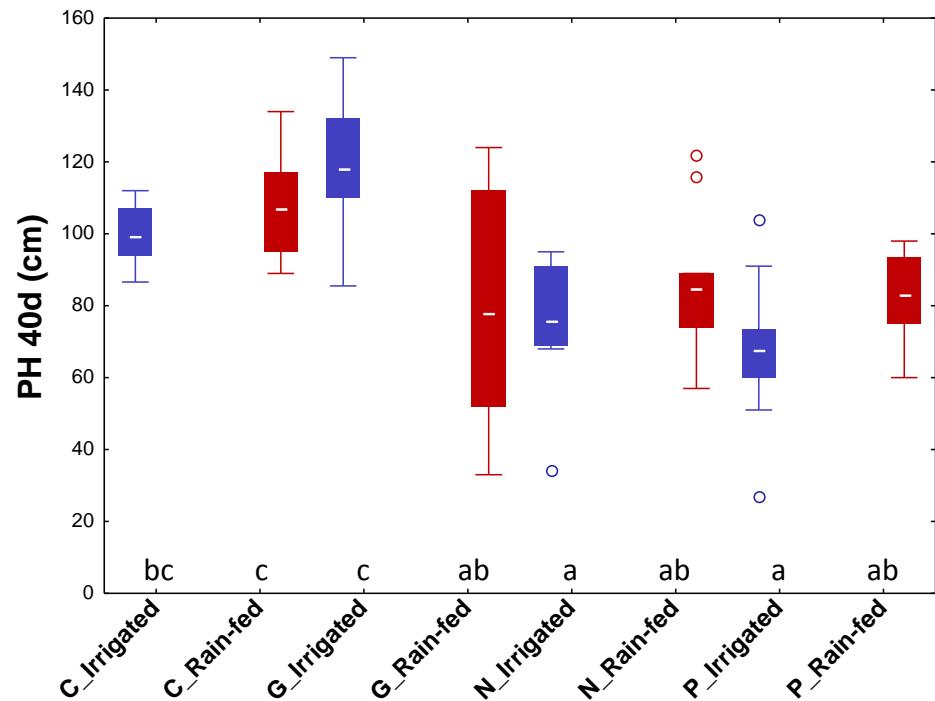
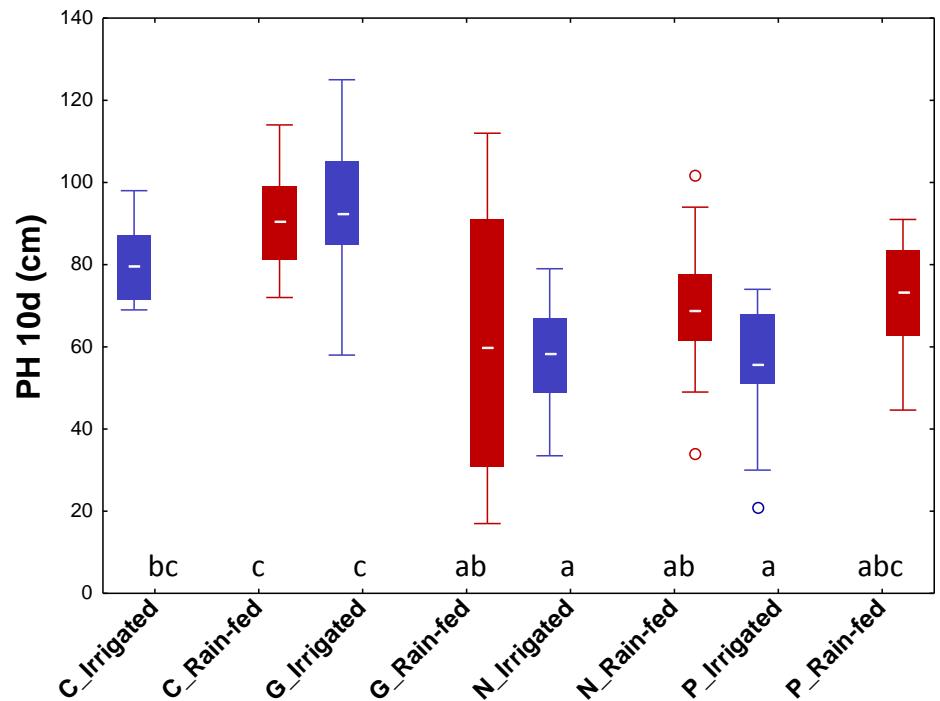
Trait	Time of measurement	Source of variation	Mean squares
Pseudostem height growth (cm)	(10d – 40d)	Treatment	303.76 <sup>ns</sup>
		Genotype	269.44 <sup>ns</sup>
		Genotype × Treatment	35.64 <sup>ns</sup>
		Residual <sup>1</sup>	58.02
Pseudostem girth growth (cm)	(10d – 40d)	Treatment	10.65***
		Genotype	7.65 <sup>ns</sup>
		Genotype x Treatment	1.05 <sup>ns</sup>
		Residual <sup>1</sup>	4.08
Leaf growth (m <sup>2</sup> )	(10d – 40d)	Treatment	7.77 <sup>ns</sup>
		Genotype	74.78*
		Genotype × Treatment	66.90*
		Residual <sup>1</sup>	9.860
Leaf temperature (°C)	47d, 48d	Treatment	470.85 <sup>ns</sup>
		Genotype	18.57 <sup>ns</sup>
		Genotype × Treatment	2.44 <sup>ns</sup>
		Residual <sup>1</sup>	7.24



**Supplementary Figure S3.** Boxplots for pseudostem girth per genotype and per treatment at 10 and 40 days after stopping irrigation. Different letters indicate significant differences based on comparison over genotypes and treatments (LSD;  $\alpha=0.05$ ). Boxes represent the interquartile range (25%-75%) and whiskers the non-outlier range. Horizontal lines inside boxes indicate the mean values, Outliers are indicated as dots. C: ‘Cachaco’, G: ‘Guyod’, N: ‘Nakitengwa’, P: ‘Pahang’. N = 10/10 (irrigated/rain-fed). PG: pseudostem girth. d: days after stopping irrigation.



**Supplementary Figure S4.** Boxplots for leaf area per genotype and per treatment at 10 and 40 days after stopping irrigation. Different letters indicate significant differences based on comparison over genotypes and treatments (LSD;  $\alpha=0.05$ ). Boxes represent the interquartile range (25%-75%) and whiskers the non-outlier range. Horizontal lines inside boxes indicate the mean values, Outliers are indicated as dots. C: ‘Cachaco’, G: ‘Guyod’, N: ‘Nakitengwa’, P: ‘Pahang’. N =10/10 (irrigated/rain-fed). d: days after stopping irrigation.



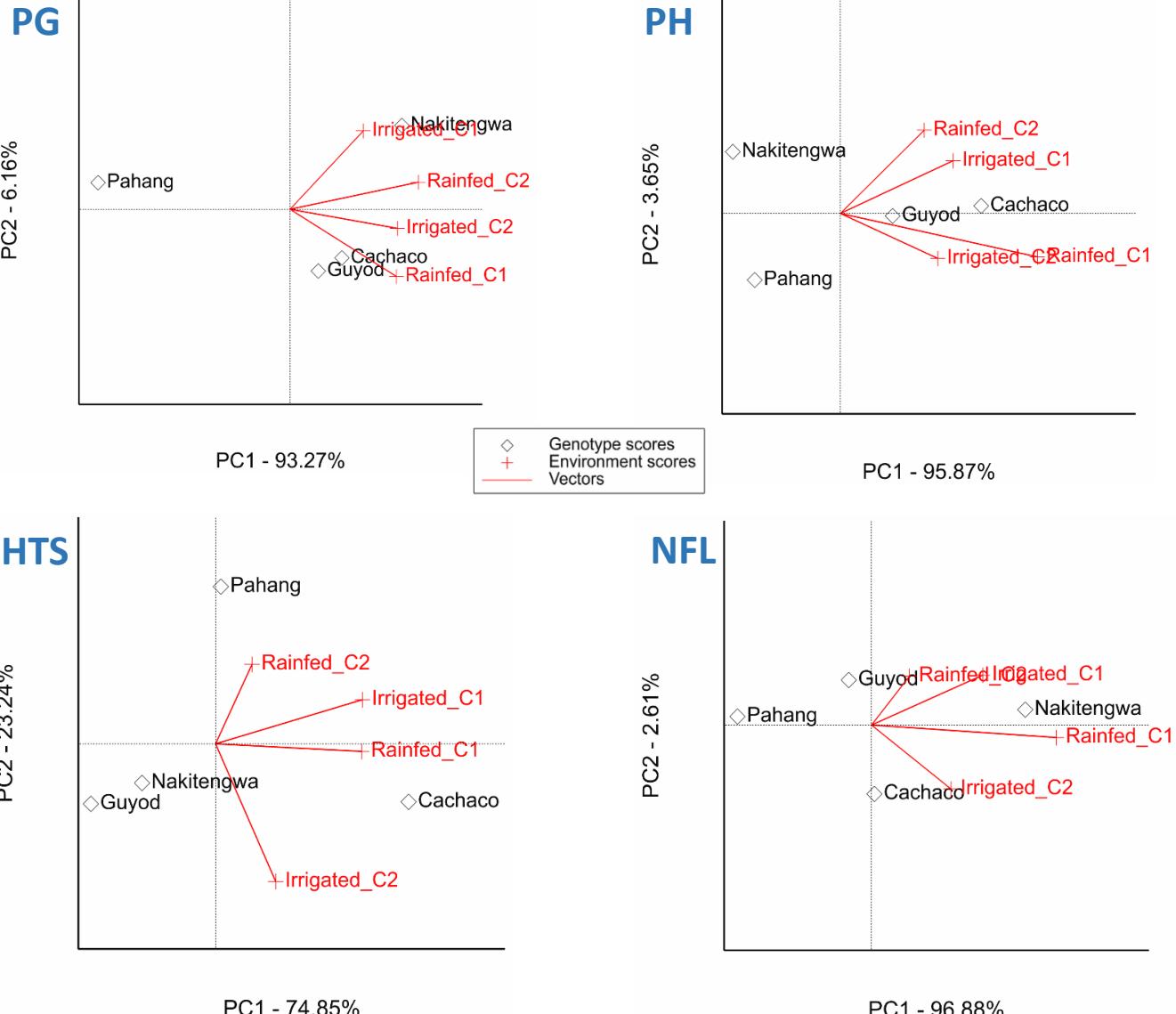
**Supplementary Figure S5.** Boxplots for pseudostem height per genotype and per treatment at 10 and 40 days after stopping irrigation. Different letters indicate significant differences based on comparison over genotypes and treatments (LSD;  $\alpha=0.05$ ). Boxes represent the interquartile range (25%-75%) and whiskers the non-outlier range. Horizontal lines inside boxes indicate the mean values, Outliers are indicated as dots. C: ‘Cachaco’, G: ‘Guyod’, N: ‘Nakitengwa’, P: ‘Pahang’. N =10/10 (irrigated/rain-fed). PH: pseudostem height. d: days after stopping irrigation.

**Supplementary Table S2.** Pearson's correlations among the traits evaluated in cycle 2 under irrigated and rain-fed treatments. Trait abbreviations are as given in Table 2 of the manuscript. \* $p\leq 0.05$ , \*\* $p\leq 0.01$ , \*\*\*  $p\leq 0.001$ , n.s.: not significant ( $p>0.05$ ).

Category	Trait	Treatment	PG	PH	HTS	NFL	FTH	BW	FFI	NH
Vegetative growth	PH	Irrigated	0.10 <sup>ns</sup>	-						
		Rain-fed	0.39*	-						
	HTS	Irrigated	0.20 <sup>ns</sup>	0.56***	-					
		Rain-fed	-0.38*	-0.10 <sup>ns</sup>	-					
	NFL	Irrigated	0.28 <sup>ns</sup>	-0.18 <sup>ns</sup>	-0.23 <sup>ns</sup>	-				
		Rain-fed	0.11 <sup>ns</sup>	0.16 <sup>ns</sup>	-0.10 <sup>ns</sup>	-				
Maturity	FTH	Irrigated	-0.22 <sup>ns</sup>	0.41**	-0.11 <sup>ns</sup>	-0.04 <sup>ns</sup>	-			
		Rain-fed	-0.34 <sup>ns</sup>	-0.14 <sup>ns</sup>	0.42*	0.03 <sup>ns</sup>	-			
	BW	Irrigated	0.74***	0.05 <sup>ns</sup>	0.14 <sup>ns</sup>	0.35*	-0.08 <sup>ns</sup>	-		
		Rain-fed	0.89***	0.41*	-0.32 <sup>ns</sup>	0.17 <sup>ns</sup>	-0.33 <sup>ns</sup>	-		
Fruit yield	FFI	Irrigated	0.75***	-0.14 <sup>ns</sup>	0.17 <sup>ns</sup>	0.32 <sup>ns</sup>	-0.45**	0.88***	-	
		Rain-fed	0.87***	0.34 <sup>ns</sup>	-0.38*	0.13 <sup>n</sup>	-0.51**	0.97***	-	
	NH	Irrigated	0.00 <sup>ns</sup>	-0.33*	-0.35*	-0.21 <sup>ns</sup>	-0.32 <sup>ns</sup>	-0.02 <sup>ns</sup>	0.07 <sup>ns</sup>	-
		Rain-fed	0.42*	0.02 <sup>ns</sup>	-0.44*	0.29 <sup>ns</sup>	-0.31 <sup>ns</sup>	0.49**	0.51**	-
NF	Irrigated	Irrigated	0.17 <sup>ns</sup>	-0.68***	-0.41**	0.01 <sup>ns</sup>	-0.48**	0.18 <sup>ns</sup>	0.31 <sup>ns</sup>	0.64***
		Rain-fed	0.30 <sup>ns</sup>	-0.45**	-0.36*	0.38*	-0.07 <sup>ns</sup>	0.27 <sup>ns</sup>	0.27 <sup>ns</sup>	0.58***

**Supplementary Table S3.** Results of ANOVA on expected yield for the four *Musa* genotypes under irrigation and rain-fed treatments in cycle 1. <sup>1</sup>Overall residual.

<b>Source of variation</b>	<b>Mean squares</b>	<b>p-value</b>
Genotype	1323.77	<0.001
Treatment	130.61	0.34
Genotype x Treatment	89.06	0.04
Residual <sup>1</sup>	20.46	



**Supplementary Figure S6.** Genotype and genotype by environment interaction (GGE) biplots for vegetative growth agronomic traits based on the best linear unbiased estimators (BLUEs) from ANOVA in the four cycle – treatment combination environments and the four *Musa* genotypes. PG: plant girth; PH: plant height; HTS: height of the tallest sucker; NFL: number of functional leaves; C1: first crop cycle, C2: second crop cycle; PC1: first principal component, PC2: second principal component.