Kernel number as a positive target trait for prediction of hybrid performance under low-nitrogen stress as revealed by diallel analysis under contrasting nitrogen conditions

Xiuxiu Li¹⁾, Zhen Sun¹⁾, Xiaojie Xu¹⁾, Wen-Xue Li¹⁾, Cheng Zou¹⁾, Shanhong Wang¹⁾, Yunbi Xu^{1,2)} and Chuanxiao Xie^{*1)}

¹⁾ Institute of Crop Science, Chinese Academy of Agricultural Sciences, National Key Facility for Crop Gene Resources and Genetic Improvement, Beijing, China 100081

²⁾ International Maize and Wheat Improvement Center (CIMMYT), El Batan, Mexico

Environmental sustainability concerns make improving yield under lower N input a desirable breeding goal. To evaluate genetic variation and heterosis for low-N tolerance breeding, 28 F1 hybrids from a diallel scheme, along with their eight parental lines, were tested for agronomic traits including kernel number per ear (KNE) and grain yield per plant (GY), in replicated plots over two years under low-nitrogen (LN, without nitrogen application) and normal-nitrogen (NN, 220 kg N ha⁻¹) conditions. Taken together the heritability in this and our previous studies, the correlation with grain yield, and the sensitivity to the stress for target trait selection, KNE was a good secondary target trait for LN selection in maize breeding. KNE also showed much higher mid-parent heterosis than hundred-kernel weight under both nitrogen levels, particularly under LN, indicating that KNE contributed the majority of GY heterosis, particularly under LN. Therefore, KNE can be used as a positive target trait for hybrid performance prediction in LN tolerance breeding. Our results also suggest that breeding hybrids for LN tolerance largely relies on phenotypic evaluation of hybrids under LN condition and yield under LN might be improved more by selection for KNE than by direct selection for GY *per se*.

Key Words: maize, heterosis, genetic variation, low-nitrogen tolerance, nitrogen-use efficiency.

Introduction

Global food demands pose huge challenges for sustainable production of cereal crops including maize (*Zea mays* L.) (Tilman *et al.* 2002). Two major innovations to increase grain yield (GY) to meet these demands are the use of hybrid maize and application of nitrogen fertilizers (Hirel *et al.* 2007).

N fertilization has made great contributions to increased GY in maize (Hirel *et al.* 2001). However, some serious environmental problems have resulted from application of high-level N in light of the relatively low N-use efficiency of maize (Guo *et al.* 2010, Ju *et al.* 2009). Developing maize varieties with improved N-use efficiency (NUE) or low-N (LN) tolerance will be essential for reducing N fertilization costs and the negative impact of excessive N on the environment (Wu *et al.* 2011). Furthermore, heterosis has been successfully used to increase maize GY (Shull 1946). One would also expect that hybrids might better tolerate stress and undergo developmental transitions better than their inbred parents (Schnable and Springer 2013). There-

fore, fundamental genetic and agronomic knowledge will improve understanding of how to combine maize hybrid vigor with N stress tolerance and provide new strategies to address nitrogen-related issues.

When breeding hybrids for low-N stress tolerance, assessing genetic variation including heterosis, general combining ability (GCA), and specific combining ability (SCA) (Le Gouis et al. 2002) under both LN and normal N (NN) conditions is key to understanding and enhancing maize NUE traits. The heredity of and genetic variation in N nutrition has been of interest for over 7 decades (Abdel-Ghani et al. 2013, Coque and Gallais 2007, Harvey 1939, Presterl et al. 2003, Worku et al. 2012, Wu et al. 2011). However, reports on genetic variation in the response to N (Azeez et al. 2006, Bänziger and Cooper 2001, Bänziger et al. 2002, Bolaños and Edmeades 1996, Lafitte and Edmeades 1994a, 1994b, Worku et al. 2012) have focused on tropical germplasm because inadequate inorganic N supply limits maize production in tropical regions (Lafitte and Edmeades 1994a, 1994b). Recent N-associated heterosis analyses have also been centered mainly on tropical germplasm (Amiruzzaman et al. 2013, El-Badawy 2013). However, heterosis has been frequently demonstrated and widely used in temperate maize. The objectives of this study were to (i) evaluate the genetic variation of temperate hybrids and their parental

Communicated by D.S. Brar

Received March 13, 2014. Accepted October 13, 2014.

^{*}Corresponding author (e-mail: xiechuanxiao@caas.cn)

lines in response to nitrogen input levels, (ii) estimate associated quantitative genetic parameters, and (iii) analyze heterosis for low-N tolerance. These research findings would help us to design and optimize breeding programs for improving low-N tolerance and NUE in maize.

Materials and Methods

Maize germplasm

Eight parental inbred lines, 81865, Qi209, Zheng30, Qi318, Jinhuang96B, CA156, Qi205, and 89-1 (Table 1) were used to produce 28 crosses from an 8×8 partial diallel that did not include reciprocals. The elite hybrid, Zhengdan958, and its two parental lines, Chang7-2 and Zheng58, were also planted together as checks for hybrids and their parental inbred lines, respectively. All of these lines are temperate-adapted germplasm and were selected to represent the different heterotic groups that have been developed during the past decades in Chinese maize breeding programs (Xie *et al.* 2007). The control hybrid, Zhengdan958, was the one of most successful temperate hybrids and still the leading hybrid now in China with a total of growing area of over 33 Mha (http://baike.baidu.com/view/264391.htm Internet news in Chinese, accessed on June 9, 2013).

Field preparation and nutrient management

The experiment was conducted at two locations in China: at Shunyi, Beijing (40.2°N 116.5°E, 44 MASL elevation) from May to October in 2009 and 2010; and at Tonghua, Jilin (42.5°N 125.6°E, 320 MASL) from May to October in 2010 and 2011 (Table 2). Field soil samples from each of the two locations were collected from soil layers at 0 to 30 cm depth prior to fertilization and planting, and then mixed, air dried, and sieved to remove undecomposed organic materials. Soil nutrients were tested to determine organic matter (Walkley 1947), total N (Bremner 1996), total P (HClO₄-H₂SO₄ digestion), total K, Olsen-P (Olsen *et al.* 1954), NH₄ OAc-K (Van Reeuwijk 1992), and pH (pHS-3C pH meter, Nanjing) The water: soil ratio was 2.5 : 1). The data on nutrients along with geographic information for each location are both provided in Table 2. Pairwise t-test revealed no significant differences in soil nutrients or pH prior to establishment of LN and NN conditions at either location.

The F1 hybrids and their parental inbreds were evaluated separately under NN and LN conditions. The NN experiment was managed under well-fertilized conditions, so as to provide an ideal maize growing environment. For the NN experiment, 80.5 kg nitrogen (N) ha⁻¹ in the form of urea, 51 kg phosphorous ha⁻¹ in the form of calcium superphosphate, and 90 kg potassium ha⁻¹ as muriate of potash were applied as a basal dressing before planting, and a second application of 161 kg N ha⁻¹ and 90 kg potassium ha⁻¹ was made as a top dressing at V8 stage. For the LN experiment, no nitrogen was applied as either a basal dressing or top-dressing, but phosphorous and potassium were applied in the same manner as for the NN experiment.

All hybrids and their parental lines along with hybrid and parental line controls were planted in the field using a randomized block design with two replications. The hybrids

N	0.	Name	Pedigree	Note
Inbred lines	1	81865	Tropical germplasm, Mexico	
	2	Qi209	Qi205 × Ye107	
	3	Zheng30	$Zheng20 \times Ye478$	
2	4	Qi318	American hybrid 78599	
1	5	Jinhuang96B	Zhongzong3 C2 (synthetic)	
(5	CA156	pool 33 temperated adapted	
	7	Qi205	Wei ai 141 × Zhongxi017	
8	8	89-1	American hybrid 78599	
С	K1	Chang7-2	$(Huangzao4 \times Wei95) \times S901$	parental line check
С	K2	Zheng58	improved Ye478	parental line check
Hybrid C	K	Zhengdan958	Zheng58 × Chang7-2	hybrids check

Table 1. Parental inbred lines and their pedigree information, in addition to parental line and hybrid checks used in this study

Table 2. Mean with standard deviation (mean \pm SD) and pairwise t-tests for background level of soil nutrients between LN and NN at two locations

	Latituda longtituda					Soil nutrients		
Locations	and elevation	N treaments	рН	Organic matter (g/kg)	Total N (g/kg)	Olsen-P (mg/kg)	Total P (g/kg)	Total K (g/kg)
Tonghua, Jilin	42.5°N 125.6°E, 320 MASL	NN LN	$\begin{array}{c} 8.30 \pm 0.10 \\ 8.35 \pm 0.03 \end{array}$	24.69 ± 3.57 29.40 ± 1.12	$\begin{array}{c} 0.83 \pm 0.03 \\ 0.89 \pm 0.02 \end{array}$	$\begin{array}{c} 39.78 \pm 3.78 \\ 36.38 \pm 3.23 \end{array}$	$\begin{array}{c} 1.50 \pm 0.08 \\ 1.61 \pm 0.09 \end{array}$	31.37 ± 0.44 29.80 ± 14.96
Shunyi,	40.2°N 116.5°E,	Pairwise t-test NN	$\begin{array}{c} 0.28\\ 8.36\pm0.10\end{array}$	$0.11 \\ 20.48 \pm 2.04$	$\begin{array}{c} 0.08\\ 0.87\pm0.05\end{array}$	$0.01 \\ 17.23 \pm 12.01$	$\begin{array}{c} 0.01\\ 0.87\pm0.09\end{array}$	$0.43 \\ 41.20 \pm 3.41$
Beijing	44 MASL	LN Pairwise t-test	8.19 ± 0.21 0.21	17.72 ± 3.57 0.14	$\begin{array}{c} 0.88 \pm 0.03 \\ 0.36 \end{array}$	20.08 ± 0.15 0.36	$0.92 \pm 0.03 \\ 0.19$	$43.24 \pm 1.35 \\ 0.27$

LN, low nitrogen; NN, normal nitrogen; MASL, meters above sea level.

and the inbred lines were tested in different blocks to avoid plant height interactions between hybrids and inbred lines. Each entry was planted in two rows, each 4.0 m long, with 0.20 m space between plants and 0.5 m between rows. The borders of both the LN and NN field trials were protected by two rows of the CK hybrids (Table 1) for removing the border effect.

Measurement of agronomic traits

The number of leaves was scored by counting the number of nodes on the stalks of six plants. Senesced leaves were also counted by marking young leaves during early development. Plant height (PH) was recorded on ten plants per block after silking, as the distance between the ground surface and the first tassel branch of the same plant.

After harvest, shoot biomass (SB), grain yield per plant (GY), kernel number per ear (KNE) and hundred-kernel weight (HKW) were recorded trait by trait for 10 plants per block. SB for each entry was recorded at time of harvest on a whole plant dry weight basis as the average of three plants selected at random per block. Fallen leaves were not included in the shoot biomass measurement. And the same time, GY was recorded on a 100% dry matter basis. Harvest index (HI) was then calculated as: $HI = \frac{GY}{SB}$. KNE was obtained by multiplying the number of rows in the middle ear and the average kernel number per row together, measured on ten similar dry ears per block and averaged. HKW was recorded by weighing 100 kernels for three replications and averaging per block.

Statistical analysis

Differences among parental lines or hybrids were tested by analysis of variance (ANOVA) conducted for each measured trait using PROC GLM (SAS 1999) with the following model (1):

$$Y_{ijkl} = \mu + E_k + B(E)_{l(k)} + G_{ij} + G_{ij} \times E_k + \varepsilon_{ijkl}$$
(1)

where Y_{ijkl} is the observed value of a parental line or hybrid, μ is the population mean, E_k is the main effect of the environment, $B(E)_{l(k)}$ is the main effect of block within environment, G_{ij} is the genotypic effect (hybrids or parental inbred lines), $G_{ij} \times E_k$ is the genotype-by-environment interaction ($G \times E$), and ε_{ijkl} is the random residual effect. Environments and genotypes were considered fixed effects.

Broad-sense heritability (h_B^2) was estimated on a block basis (Harvey 1939) as Eq. (1):

$$h_B^2 = \frac{\sigma_G^2}{\sigma_G^2 + \sigma_{G \times E}^2 + \sigma_{\varepsilon}^2} \tag{1}$$

where σ_G^2 is genotypic variance, $\sigma_{G\times E}^2$ is the variance due to the $G \times E$ interaction, and σ_{ε}^2 is the error variance. The narrow-sense heritability (h_N^2) of hybrids was calculated based on variance in combining ability, which is derived by Eq. (2) from the variances of general combining ability (GCA) and special combining ability (SCA), according to Griffing's method 4 (Griffing 1956):

$$h_{N}^{2} = \frac{2\sigma_{GCA}^{2}}{2\sigma_{GCA}^{2} + \sigma_{SCA}^{2} + 2\sigma_{GCA \times E}^{2} + \sigma_{SCA \times E}^{2} + \sigma_{\varepsilon}^{2}} \quad (2),$$

where σ_{GCA}^2 , σ_{SCA}^2 , $\sigma_{GCA \times E}^2$, and $\sigma_{SCA \times E}^2$ were estimated from the subsequent diallel analysis.

When genotypic effects for traits in hybrids are significant according to ANOVA, diallel analysis of combining ability is then conducted according to Griffing's method 4 (Griffing 1956) using a special SAS program for diallel analysis with missing crosses developed by Wu and Matheson (2000) that considers environment and genotype as fixed effects. The response Y_{ijkl} of the cross between parents *i* and *j* was modeled as model (2):

$$Y_{ijkl} = \mu + E_k + B(E)_{l(k)} + GCA_i + GCA_j + SCA_{ij} + GCA_i \times E_k + GCA_j \times E_k + SCA_{ij} \times E_k + \varepsilon_{ijkl}$$
(2),

where μ , E_k , $B(E)_{l(k)}$ and ε_{ijkl} are the same as in model (1); GCA_i and GCA_j are the GCA of parents *i* and *j*; SCA_{ij} is the SCA of hybrid *ij*; $GCA_i \times E_k$, $GCA_j \times E_k$, and $SCA_{ij} \times E_k$ are interactions between GCA and SCA and the environment.

The relative importance of GCA and SCA was estimated as the ratio: $\frac{2K^2_{GCA}}{2K^2_{GCA} + K^2_{SCA}}$, modified from Baker (1978), where K^2_{GCA} is the quadratic form (analogous to a variance component, but referring to a fixed effect) derived from the mean square of the GCA effect and K^2_{SCA} is the quadratic form of the SCA effects. As this ratio approaches unity, GCA becomes more important, and the predictability of the performance of a specific hybrid could be based on GCA alone.

The variance of the SCA effect of parent $i (S_{si}^2)$ indicates the differences in SCA among a series of hybrids involving one line, estimated as Eq. (3) (Griffing 1956):

$$S_{si}^{2} = \frac{1}{P-2} \sum_{i \neq j} SCA_{ij}^{2} - \frac{P-3}{P-2} \sigma_{\varepsilon}^{2}$$
(3)

The low-N tolerance index (*LNTI*) was estimated by Eq. (4) to evaluate the ability of parental lines and hybrids to tolerate low nitrogen in terms of the traits measured.

$$LNTI = \frac{LN}{NN} \times 100\% \tag{4}$$

A higher *LNTI* indicates a stronger ability to adapt to low nitrogen.

Heterosis (%) was calculated over the high-parent and mid-parent values. High-parent heterosis (*HPH*) was calculated using Eq. (6) (Hill *et al.* 1987). Similarly, mid-parent heterosis (*MPH*) was estimated using Eq. (7).

$$HPH(\%) = \frac{F_1 - HP}{HP} \times 100 \tag{6}$$

$$MPH(\%) = \frac{F_1 - MP}{MP} \times 100 \tag{7}$$

 F_1 is the performance of each hybrid; HP indicates the performance of the best parent and MP represents the mean performance of two parental inbred lines. For each hybrid, the difference between the hybrid and the mean of its two parents as well as the best parent was tested by a t-test to determine whether *MPH* or *HPH* was significant (Le Gouis *et al.* 2002).

The relationship between the *LNTI* of parental inbred lines and that of hybrids was measured using PROC REG in SAS (SAS 1999). Similarly, correlations between SCA, heterosis, and least square mean values of hybrids, as well as correlations between performance of hybrids and the mean performance of two parents for all measured traits were calculated using PROC CORR in SAS (SAS 1999).

The correlation between inbred line and hybrid performance was estimated by linear regression analysis of the GCA values on inbred performance.

Results

The effect of low nitrogen on agronomic performance of the hybrids and their parental inbreds

Significant differences were found between LN and NN trials for most of the measured traits except of leaf number (LEAF) and harvest index (HI) for both parental lines and their half-diallel cross hybrids (Table 3). Among these traits, GY and SB showed the largest differences between LN and NN treatments. For example, average GY of the parental inbred lines and hybrids under LN was only 74.1% and 64.0% of those under NN, with an average reduction of 25.9% and 36.0%, respectively. Additionally, the differences between the inbred lines and hybrids were also very significant due to heterotic effect. The hybrids showed significantly higher phenotypic performance than their parental lines under both NN and LN conditions, indicating significant heterosis. For

Table 3. Differences between parental lines and diallel cross F1s under low- (LN) or normal-nitrogen (NN) conditions

		LN							NN						
		LEAF	PH	HKW	KNE	GY	SB	HI	LEAF	PH	HKW	KNE	GY	SB	HI
		(number)	(cm)	(g)	(number)	(g)	(g)	(%)	(number)	(cm)	(g)	(number)	(g)	(g)	(%)
Parental lines	Avg.	18.2	193.9	26.6	155.6	41.1	108.0	32.7	18.9	210.8	30.5	181.4	55.5	156.8	34.9
	Max.	20.8	239.5	36.5	251.6	72.2	134.4	51.4	22.3	277.7	36.8	271.8	83.8	192.0	52.6
	Min.	16.0	170.9	18.6	87.5	22.5	87.2	13.8	17.0	191.8	23.5	81.2	23.8	106.3	14.0
	CV (%)	7.7	10.9	21.0	33.2	48.5	10.2	35.0	10.4	12.1	13.9	23.3	38.3	19.4	31.2
	Avg. LN/NN (%)	96.4	92.0	87.2	85.8	74.1	68.8	93.9							
	Avg.	19.3	280.5	29.9	366.3	109.3	217.5	46.7	20.1	298.8	35.2	483.7	170.7	335.8	51.6
	Max.	21.3	339.4	38.3	445.8	149.9	323.0	54.1	23.8	356.4	42.5	578.6	208.7	428.6	61.1
F1s	Min.	17.3	223.3	24.2	182.8	53.3	154.0	30.3	18.0	242.5	29.3	351.6	114.7	278.5	38.9
	CV (%)	6.0	9.5	13.5	15.3	17.7	18.7	8.4	6.1	9.6	11.3	13.6	14.5	11.2	10.2
	Avg. LN/ NN (%)) 95.8	93.9	85.0	75.7	64.0	64.8	90.5							
Parental	Lines/F1s (%)	94.4	69.1	89.1	42.5	37.6	49.6	70.1	93.8	70.6	86.7	37.5	32.5	46.7	67.5

LN, low nitrogen; NN, normal nitrogen; LEAF, the final number of leaves; PH, plant height; HKW, hundred-kernel weight; KNE, kernel number per ear; GY, grain yield; SB, shoot biomass; HI, harvest index; Avg., average; Max., maximum; Min., minimum.

Table 4. Variance analysis of parental lines and their diallel cross F1 hybrids under two nitrogen levels in two locations across two years

T	C				LN				NN						
Types	Source	LEAF	PH	HKW	KNE	GY	SB	HI	LEAF	PH	HKW	KNE	GY	SB	HI
	Genotype	5.20*	2863.77**	65.12**	21682.29**	2133.59**	6284.09**	0.012**	6.63**	2807.15**	62.89**	24999.81**	2613.65**	6057.49**	0.015**
	Env.	0.81	7941.65**	994.01**	2580.03	21633.11**	65712.76**	0.290**	0.81	391.73	39.6	509.63	3747.83*	5495.26	0.093**
	Rep.	1.45*	4331.74**	45.03	7539.64*	229.31	4220.85	0.001	1.13	287.58	1.77	1266.36	655.08	637.60	0.002
F1 hybrids	Genotype*Env.	0.06	170.88	26.29*	3660.65*	784.85	3175.65**	0.006*	0.25	219.93	23.17	2616.01	927.38	1717.58	0.003
	Error	0.43	150.42	14.84	1638.87	739.36	1385.59	0.00	0.57	125.27	14.47	1785.43	646.05	2283.60	0.002
	h_B^2	0.822	0.833	0.365	0.678	0.335	0.261	0.265	0.791	0.828	0.326	0.759	0.379	0.332	0.579
	h_N^2	0.683	0.639	0.315	0.436	0.101	0.047	0.011	0.476	0.745	0.245	0.547	0.133	0.077	0.413
	Genotype	8.93**	1642.61**	129.12**	17921.77**	1219.49**	767.46*	0.064**	14.42**	3810.04**	86.35**	16462.04**	1727.41**	3625.80**	0.049**
	Env.	0.50	3130.40**	124.39**	79693.63**	794.36*	117.05	0.024*	0.28	390.60*	4.54	16066.28*	1287.42*	5067.48*	0.008
Parental	Rep.	0.63	14.60	17.81*	23.64	117.83	333.07	0.009	0.16	305.33	6.72	42.37	437.07	687.12	0.014
lines	Genotype*Env.	0.07	137.16*	30.50**	9991.51*	86.08	595.27	0.005	0.07	107.95	45.99**	1946.11	456.10	730.90	0.008
	Error	0.20	46.32	4.37	1515.44	121.26	274.60	0.005	0.58	83.94	9.76	2490.93	174.47	613.85	0.005
	h_B^2	0.917	0.813	0.641	0.416	0.694	0.221	0.753	0.855	0.907	0.407	0.584	0.552	0.528	0.640

LN, low nitrogen; NN, normal nitrogen; LEAF, the final number of leaves; PH, plant height; HKW, hundred-kernel weight; KNE, kernel number per ear; GY, grain yield; SB, shoot biomass; HI, harvest index; Geno., genotype; Env., environment; Rep., replication; h_B^2 : broad-sense heritability; h_N^2 : narrow-sense heritability. *,** indicates significance at 5% and 1% levels of probability, respectively.

instance, the inbred lines had only 37.6% of the average GY of hybrids, indicating that the heterosis effect was very strong for grain yield under LN stress.

Genetic variation in response to LN stress revealed by analysis of variance

Significant differences were observed among both the eight parental lines and their hybrids for all measured traits under both nitrogen levels (Table 4). Furthermore, both environmental variance and the $G \times E$ interaction for most of the measured traits were more frequent and significant for both hybrids and inbred lines under LN than NN. This suggests that LN treatment in this study at each location across years increased phenotypic variance because of the increase in effects due to environment and $G \times E$ interaction. This is also an indication of differential responses to LN stress among tested hybrids and their parental inbred lines.

Broad-sense heritability for KNE was relatively high for both parental lines and hybrids under both LN and NN conditions. Narrow-sense heritability of KNE for hybrids under both LN and high NN conditions was higher than that of GY and its component trait HKW. These data indicate that KNE could respond better to selection than GY for a breeding program directed at maintaining or improving grain yield under LN stress.

and SCA variation across these lines were significant for all measured traits under both LN and NN conditions (Table 5). GCA × Env interactions and SCA × Env interactions were not significant for most traits. These results indicate that genetic effects were the main contributors to combining ability variance. Under both LN and NN conditions, PH, KNE, GY, SB, and HI showed high HPH and MPH, indicating that they would be good traits to target for exploiting heterosis (Table 6). GY can be dissected into its components KNE and HKW, and average HPH and MPH for KNE were much higher than those for HKW under both conditions, and were even larger under LN than under NN. Therefore, the KNE component contributes the majority of heterosis for GY under both LN and NN conditions, but particularly under LN. GY exhibited the largest range of heterosis among measured traits, ranging from 38.7% to 501.1% and from 71.9% to 436.2% for HPH under LN and NN, respectively, and from 70.7% to 517.2% and from 101.4% to 502.6% for MPH under LN and NN, respectively. The LEAF trait exhibited the lowest range of heterosis; from -8.0% to 9.9% and -9.0% to 23.2% for HPH under LN and NN, respectively, and from -0.7% to 15.6% and from -1.4% to 24.1% for MPH under LN and NN, respectively. The ranges of values for heterosis for all traits except LEAF increased concomitantly with the lower nitrogen levels under the LN treatment, indicating that the hybrids showed much larger differential responses to LN stress compared to their parental lines.

Diallel analysis

GCA and SCA were also analyzed in this study. GCA

The GCA effect, which determines the average

 Table 5.
 Mean squares for general combining ability (GCA) and specific combining ability (SCA) under low-nitrogen and normal-nitrogen conditions

		LEAF	PH	HKW	KNE	GY	SB	HI
	GCA	14.7**	7287.4**	159.5**	46553.5**	3876.3**	7809.4**	0.014**
	SCA	1.3**	882.7**	26.3	11441.2**	1416.0*	5656.0**	0.012**
LN	$GCA \times Env.$	0.009	231.2	25.7	3120.1	821.1	3330.1*	0.009*
	$SCA \times Env.$	0.08	150.2	26.5	3883.2**	769.9	3112.1*	0.005
	Ratio	0.66	0.53	0.60	0.34	0.35	0.15	0.13
	GCA	14.8**	8588.1**	158.5**	61286.9*	1881.1*	7795.69**	0.037**
	SCA	3.3**	480.6**	23.5	10058.1**	2915.3**	5341.8*	0.006**
NN	$GCA \times Env.$	0.2	340.5*	56.5**	3622.5	1610.0*	1377.8	0.005*
	$SCA \times Env.$	0.3	169.6	9.4	2201.6	646.3	1857.5	0.002
	Ratio	0.38	0.74	0.65	0.46	0.06	0.17	0.48

LEAF: the final number of leaves; PH, plant height; HKW, hundred-kernel weight; KNE, kernel number per ear; GY, grain yield; SB, shoot biomass; HI, harvest index; Env., environment; * ,** Significant at 5% and 1% levels of probability, respectively; Ratio: $2K^2_{GCA}/(2K^2_{GCA} + K^2_{SCA})$.

Table 6.	Heterosis	for measure	d traits	under	two	nitrogen	levels
----------	-----------	-------------	----------	-------	-----	----------	--------

-															
					LN				NN						
		LEAF	PH	HKW	KNE	GY	SB	HI	LEAF	PH	HKW	KNE	GY	SB	HI
HPH	Aver.	1.5	33.9	-0.9	114.3	159.8	89.4	39.3	1.5	28.2	4.2	99.5	192.7	97.1	43.7
	Max.	9.9	69.4	40.6	239.7	501.1	176.4	170.7	23.2	47.4	38.2	133.3	436.2	163.5	111.7
	Min.	-8.0	10.9	-27.7	39.7	38.7	33.3	-9.4	-9.0	15.3	-12.5	56.6	71.9	46.2	-13.0
	Aver.	6.5	42.7	11.3	147.2	206.3	100.9	68.3	7.1	37.8	14.0	118.8	257.4	120.1	69.3
MPH	Max.	15.6	72.2	51.9	267.3	517.2	193.6	195.6	24.1	53.0	46.8	150.8	502.6	193.0	135.0
	Min.	-0.7	21.0	-12.7	59.5	70.7	49.9	12.6	-1.4	25.5	-4.7	83.1	101.4	53.8	19.9

LEAF, the final number of leaves; PH, plant height; HKW, hundred-kernel weight; KNE, kernel number per ear; GY, grain yield; SB, shoot biomass; HI, harvest index; Aver., average; Max., maximum; Min., minimum; LN, low nitrogen; NN, normal nitrogen; HPH, high-parent heterosis; MPH, mid-parent heterosis.

Table 7. General combining ability effects (GCA) and variance of specific combining ability $(SCA)(S_{si}^2)$ of parent i for all measured traits under two N levels at two sites

GCA				LN							NN			
$S^2{}_{Si}$	LEAF	PH	HKW	KNE	GY	SB	HI	LEAF	PH	HKW	KNE	GY	SB	HI
1	1.43**	38.38**	3.80**	7.61	21.03**	32.22**	-0.0017	1.65**	37.90**	2.24**	-8.49	6.79	37.47**	-0.0328**
	0.02	180.47	6.99	1433.15	12.62	837.89	0.0015	0.45	21.23	6.81	2796.70	345.96	1162.62	0.0011
2	-0.24	-0.38	3.66**	-82.56**	-15.58**	-27.46**	-0.0353**	-0.22	11.35**	3.60**	-73.46**	-1.03	4.05	0.0071
	0.17	101.26	1.42	5173.06	263.78	542.77	0.0024	0.80	64.39	2.16	2418.59	388.23	530.87	0.0001
3	-0.58**	-10.20**	1.32	17.97*	10.70*	21.07**	0.0232*	-0.54**	-16.24**	2.97**	-8.18	6.15	1.67	0.0255**
	0.10	216.37	0.06	597.10	50.77	1469.72	0.0005	0.95	105.51	2.50	584.33	558.59	839.53	0.0003
4	0.02	4.59	-0.65	1.91	4.22	-2.88	0.0209	0.11	-3.53	0.01	-12.15	-1.53	0.96	-0.0128
	0.40	142.23	0.21	27.70	139.90	801.82	0.0006	0.47	123.31	0.35	88.31	458.91	1374.88	0.0001
5	-0.48**	-10.02**	-2.62**	84.54**	10.22*	2.12	0.0435*	-0.64**	-12.49**	-2.63**	83.33**	6.95	-12.34	0.0514**
	0.06	23.66	1.13	5662.20	162.89	687.07	0.0002	0.00	18.91	3.10	2272.00	203.76	131.00	0.0003
6	-0.21	-16.34**	-0.75	2.96	-17.09**	-19.18*	-0.0141	-0.67**	-19.47**	-1.31	24.09**	-8.33	-30.76**	0.0028
	0.04	237.92	4.72	596.05	92.06	443.95	0.0028	0.01	125.19	6.29	1759.83	533.21	622.28	0.0010
7	-0.90**	-19.77**	-2.69	42.72**	-3.26	2.80	0.0003	-0.43**	-17.88**	-2.15**	66.64**	10.03	-1.12	0.0365**
	0.44	68.42	2.66	833.07	212.55	932.65	0.0013	1.28	64.42	1.41	1631.57	164.87	233.81	0.0001
8	0.97**	13.75**	-2.07*	-39.21**	-10.23	-8.68	-0.0368**	0.74**	20.37**	-2.73**	-71.78**	-19.03**	0.07	-0.0777**
	0.28	304.19	4.13	1902.60	22.09	1160.75	0.0015	0.43	208.93	1.86	2043.33	217.27	479.14	0.0011

LEAF, the final number of leaves; PH, plant height; HKW, hundred-kernel weight; KNE, kernel number per ear; GY, grain yield; SB, shoot biomass; HI, harvest index; LN, low nitrogen; NN, normal nitrogen; GCA, general combining ability; SCA, specific combining ability.

performance of a parent in a series of crosses, was calculated for each parental line for all traits along with their corresponding SCA variances (Table 7). Combined data analysis provides average performance values and reveals the outstanding hybrids with high SCA variance for a given line. Jinhuang96B has the highest GCA along with high SCA variance for KNE, which makes it the best line to use for LN tolerance as defined by this component of yield. Conversely, the parental line 2 Qi209 had the lowest GCA but high SCA, which means it could be useful for selection of elite hybrids based on high SCA estimates alone although with less confidence.



Fig. 1. Correlations between midparent and hybrid performance for each trait under LN and NN. LEAF: the final number of leaves; PH, plant height; HKW, hundred-kernel weight; KNE, kernel number per ear; GY, grain yield; SB, shoot biomass; HI, harvest index.

Correlation between inbred line per se and hybrid performance

the correlation between hybrids and mid-parents was positive and significant for all traits under NN (Fig. 1). However, for the GY related traits, only KNE and HKW were significantly positively correlated under LN. Under LN, KNE and PH both fit a linear regression with high correlation coefficients (R^2) of 0.6688 and 0.7838 (Fig. 2) between *per se* and hybrid performance, respectively. However, under NN, R^2 for KNE decreased to 0.3832. In summary, KNE may be a positive trait with predictive value for the relationship between performances of an inbred line *per se* and in hybrids for a maize low-N tolerance breeding program. However, inbred lines and hybrids should be evaluated under LN conditions to identify inbreds with good breeding value and outstanding hybrids for low-N tolerance.

Discussion

Development of hybrid varieties and application of N fertilizers are two of the most important proven approaches to meeting increased future demands for maize. Breeding for improved maize low-N tolerance can be expected to also effect positive genetic gains under high N because approximately two-thirds of genetic gains for grain yield at high N can be explained by improvements in grain yield at low N (Haegele *et al.* 2013). Moreover, GY of modern maize hybrids depends largely upon sowing density, as GY has been shown to be a function of plant population density and high planting-density tolerance, rather than intrinsic yield potential (Duvick 2005, Tollenaar and Lee 2002). At higher plant populations, low-N stress should be more severe due to competition for resources including N. Therefore, LN stress tolerance results from an important intersection of grain



Fig. 2. Observed and predicted KNE and PH across 28 F1 diallel crosses at NN (A, B) and LN (C, D) conditions calculated for each cross from the GCA effect of two parents and over-all means. KNE: kernel number per ear; PH: plant height.

yield, NUE, and heterosis in maize. However, the success of maize breeding for high NUE or low-N tolerance largely relies on acquiring relevant knowledge and understanding the genetic control of these traits at the population level. For example, the responses of hybrids and their parental lines to nitrogen input levels in terms of genetic variation and the magnitude of heterotic effects are important parameters for improving NUE and low N tolerance. Understanding the relevant kinds of genetic variation might direct breeders to the appropriate traits to target during line development or test crossing and reveal to what extent hybrid performance can be predicted based on parental lines and their testcrosses.

Genetic variability reveals KNE as a breeding target for LN tolerance

ANOVA revealed significant differences among these eight parental lines and their hybrids for all traits under two nitrogen levels (Table 4). However, the significances of the variances among traits differed between N levels. The heritability of each of these traits can explain their genetic contribution to the differences among variances. Under both N levels, genetic variability in the hybrids is higher than that in the parental lines for all traits except LEAF and HI (Table 4). Under LN conditions, the magnitude of the genetic variability decreased in the parental lines for all traits except HKW, but increased in the hybrids for all traits except LEAF. Compared with parental lines, hybrids showed lower heritability for all traits except KNE under NN, indicating that the performance of parental lines is more stable genetically than that of their hybrids under NN. LEAF showed the highest heritability, while KNE showed highest heritability among all measured yield traits in both parental-line and hybrid-based analyses except for parental line under the LN condition (Table 4). However, our previous report using a larger panel of inbred lines (n = 189), which included all parental lines in this study, tested under both LN and HN conditions revealed that KNE had higher heritability than any other yield component traits under LN (Wu et al. 2011). The genetic parameters characterized among those border genetic basis should be more credibly to justify that KNE was a good selecting trait which respond to LN selection at maize breeding. In addition to heritability, the sensitivity of the secondary trait other than GY per se was very important for improving the precision and demonstrating the degree of stress to crop to low nitrogen stress (Bänziger 2002). In this study, KNE was found to have a larger coefficient variation (CV) than HKW under LN and NN conditions in both parental lines and hybrids (Table 3), indicating that KNE shows a larger range of variation in both parental lines and hybrids under two nitrogen conditions. KNE, as well as GY, was more sensitive to nitrogen than HKW in both parental lines and hybrids (Table 3). For example, average KNE of the parental inbred lines and hybrids under LN was only 85.8% and 75.7% of those under NN, while HKW was 87.2% and 85.0%, respectively. In addition, KNE showed a more significant correlation with grain yield than HKW in both parental lines and hybrids under LN stress. For example, the correlation coefficients (R^2) between GY and KNE are 0.68 and 0.60 for parental lines and hybrids respectively under the LN condition, while HKW are only 0.48 and 0.35 for parental lines and hybrids, respectively (data not shown). Therefore, concerning the heritability in this and

previous studies, the correlation with grain yield, and the sensitivity to the stress, for target trait selection, KNE was better than HKW as a secondary target trait for LN selection in maize breeding. A primary goal of NUE breeding and production is to maintain high crop productivity under lower N application and to determine whether it is possible to select good genotypes that are adapted to LN, NN, or both nitrogen conditions (Hirel et al. 2007). As a result, we could achieve ideal GY under LN stress. Thus, in terms of the agronomic performance of inbred lines under LN, KNE would be the best target trait for selection. Evaluation of hybrid performance also revealed that KNE had higher heritability than other GY component traits under both LN and HN conditions, predicting good response to selection for KNE as a target trait for improvement of N stress tolerance in future hybrid breeding programs.

Genotype by environment interaction

Furthermore, parental lines showed more significant $G \times E$ interaction than did hybrids. The $G \times E$ was significant for HKW, KNE, GY, and HI under both N levels. For hybrids, the environment effect was significant only for GY and HI, while $G \times E$ was not significant for most of the measured traits under NN. These results indicate that inbred lines were more susceptible to environmental factors, particularly under LN, and that the performance of the hybrids should be relatively more stable. Therefore, hybrid evaluation under LN should be more reliable than the evaluation of parental lines in breeding programs for LN tolerance.

Heterosis for LN tolerance and its breeding implications

Maize hybrid vigor for grain yield is determined by the heterosis associated with plant biomass, kernel number, and harvest index (Munaro *et al.* 2011). Heterosis revealed in this study for HPH and MPH for PH and KNE tended to be relatively greater under LN conditions but lower under NN conditions (Table 6), consistent with a previous report (Zaidi *et al.* 2003). In this study, heterosis levels for HKW, GY, SB, and HI were greater under NN conditions. In particular, heterosis results for GY agreed with the conclusions from a previous study on a tropical maize population (Gama *et al.* 2002). Further, low heterosis was found for LEAF under both LN and NN conditions in this study, as also found for leaves above the ear in a previous report (Mehta and Sarkar

1992). Our findings suggest that leaf number is a developmental trait with large genotypic differences but low heterosis effects. Under NN, environmental effects in parental lines were significant for PH, KNE, GY, and SB; however, under LN, environmental effects were significant for PH, HKW, KNE, GY, and HI. MPH for KNE was much higher than that for HKW at both N levels, but particularly under LN. Therefore, KNE contributed the majority of GY heterosis under both LN and NN conditions, but especially under LN.

Prediction of hybrid performance from parental lines

Using inbred line information to predict hybrid performance could reduce the need for hybrid evaluation. An early study reported that the contribution of mid-parent performance for GY was comparatively greater than mid-parent heterosis under both low and normal nitrogen conditions (Zaidi et al. 2003), which encouraged us to explore the correlation between hybrid and mid-parent performance for all measured traits. A similar result for PH was obtained in an earlier study, with a lower correlation observed under severe low nitrogen stress than under optimal conditions (Zaidi et al. 2003) and stronger correlations observed for HKW under low-N stress (Bolaños and Edmeades 1996). However, correlations between performances of parent and hybrid have often been reported to increase under extreme environmental conditions (Good et al. 2004). The extent of hybrid performance that we could predict will determine whether this approach will be useful in breeding programs. Our results suggest that both KNE and PH can fit the linear regression with a high correlation coefficient under LN conditions (Fig. 2). Actually, most of the traits studied here, except HI, show positive correlations under N stress (data not shown). Linear regression revealed a correlation between parental lines and hybrids for low-N tolerance (Table 8). Among all the traits measured here, the regression coefficients between hybrids and parents estimated for the low-N tolerance index (LNTI) were only significant for HKW ($R^2 = 0.2334$) and SB ($R^2 = 0.2770$). This indicates that the nitrogen sensitivity of HKW and SB in hybrids is more related to that of their parents than are the nitrogen sensitivities of other traits we studied. However, our experiment revealed that for most of the other traits we studied, low-N tolerance in hybrids had no certain correlation with that of their parental lines. Based on the definitions of GCA and SCA (Griffing 1956), the

Table 8. Parent-offspring correlations for the low-N tolerance index (LNTI) between the parental lines and their diallel cross F1 hybrids

Traits -	F1	, MP	F1	, HP	F1,	LP	F1, (HP LP)			
Traits	Reg.	R-square	Reg.	R-square	Reg.	R-square	Reg. (HP)	Reg. (LP)	R-square	
LEAF	0.135	0.0031	0.231	0.0117	-0.019	0.0001	0.282	-0.119	0.0149	
PH	0.054	0.0075	0.213	0.0185	0.023	0.0044	0.207	0.003	0.0186	
HKW	0.073	0.0039	-0.185	0.0387	0.329	0.1039	-0.370	0.492*	0.2334	
KNE	-0.030	0.0035	-0.059	0.0243	0.043	0.0075	-0.097	0.105	0.0584	
GY	-0.089	0.0181	-0.187	0.1106	0.057	0.0104	-0.264*	0.172	0.1878	
SB	0.403	0.1215	0.127	0.0204	0.588**	0.2600	-0.135	0.675**	0.2770	
HI	0.097	0.0432	0.071	0.0387	0.079	0.0278	0.054	0.040	0.0435	

*,** Significant at 5% and 1% levels of probability, respectively; Reg., Regression coefficient; MP, mid-parent; HP, high parent; LP, low parent.

Traits			LN			NN						
Traits	r(F1,MPH)	r(F1,HPH)	r(SCA,MPH)	r(SCA,HPH)	r(F1,SCA)	r(F1,MPH)	r(F1,HPH)	r(SCA,MPH)	r(SCA,HPH)	r(F1,SCA)		
LEAF	0.47*	0.16	0.79**	0.61**	0.43*	0.29	0.05	0.82**	0.74**	0.56**		
PH	0.66**	0.46*	0.53**	0.61**	0.46*	0.37	-0.40*	0.95**	0.42*	0.35		
HKW	0.55**	0.50*	0.47*	0.44*	0.53**	0.62**	0.64**	0.51**	0.52**	0.51**		
KNE	0.36	0.17	0.40*	0.25	0.59**	0.57**	0.38	0.77**	0.64**	0.50**		
GY	0.55**	0.42*	0.44*	0.40*	0.63**	0.28	0.31	0.49*	0.45*	0.89**		
SB	0.91**	0.84**	0.79**	0.75**	0.80**	0.52**	0.53**	0.66**	0.47**	0.79**		
HI	0.09	0.08	0.46*	0.44*	0.81**	-0.09	0.11	0.41*	0.35	0.55**		

Table 9. Correlations among F1 and MPH, HPH, and SCA for traits measured under two nitrogen levels

LEAF: the final number of leaves; PH, plant height; HKW, hundred-kernel weight; KNE, kernel number per ear; GY, grain yield; SB, shoot biomass; HI, harvest index; LN, low nitrogen; NN, normal nitrogen.

correlations between MPH or HPH and F1 performance and between SCA and F1 performance for many traits might also be used for parent-hybrid performance prediction. A large number of positive correlations were found between these combinations of parameters in this study (Table 9), so it does seem that both GCA and SCA play important roles in F1 performance. Finally, these results provide further evidence that prediction of hybrid performance based on that of parental lines, although a major challenge in low-N tolerance breeding, may be amenable to the type of analysis presented here.

Conclusions

From a breeder's point of view, selection of the appropriate maize hybrids to be cultivated under LN input will be facilitated by identification of high GCA and SCA effects. Improving low-N tolerance in maize can be expected to also result in genetic gain for yield under high N. Genetic variation and heterosis analysis showed that genetic improvement of hybrids for LN tolerance is an effective approach to enhancing maize production under lower N input settings. Selection for KNE, rather than for GY itself, may be more effective for genetic improvement of yield under LN, or LN tolerance. Therefore, evaluation of hybrids under LN is essential for breeding hybrid maize for LN tolerance.

Acknowledgements

This study was supported by Chinese National "863" Program (Grant #2012AA10A306 and #2012AA101104) and National Science Foundation of China (Grant #31361140364).

Literature Cited

- Abdel-Ghani, A.H., B. Kumar, J. Reyes-Matamoros, P.J. Gonzalez-Portilla, C. Jansen, J.P. San Martin, M. Lee and T. Lübberstedt (2013) Genotypic variation and relationships between seedling and adult plant traits in maize (*Zea mays* L.) inbred lines grown under contrasting nitrogen levels. Euphytica 189: 123–133.
- Amiruzzaman, M., M. Islam, L. Hasan, M. Kadir and M. Rohman (2013) Heterosis and combining ability in a diallel among elite inbred lines of maize (*Zea mays* L.). Emir. J. Food Agric. 25: 132– 137.

- Azeez, J.O., M.T. Adetunji and S.T.O. Lagoke (2006) Response of lownitrogen tolerant maize genotypes to nitrogen application in a tropical alfisol in northern Nigeria. Soil Till. Res. 91: 181–185.
- Bänziger, M. and M. Cooper (2001) Breeding for low input conditions and consequences for participatory plant breeding: Examples from tropical maize and wheat. Euphytica 122: 503–519.
- Bänziger, M., G.O. Edmeades and H.R. Lafitte (2002) Physiological mechanisms contributing to the increased N stress tolerance of tropical maize selected for droought tolerance. Field Crops Res. 75: 223–233.
- Baker, R. (1978) Issues in diallel analysis. Crop Sci. 18: 533-536.
- Bolaños, J. and G.O. Edmeades (1996) The importance of the anthesissilking interval in breeding for drought tolerance in tropical maize. Field Crops Res. 48: 65–80.
- Bremner, J.M. (1996) Nitrogen-total. *In*: Sparks, D.L., A.L. Page, P.A. Helmke and R.H. Loeppert (eds.) Methods of Soil Analysis Part 3. Chemical Methods, Soil Science Society of America, American Society of Agronomy, Madison, WI, pp. 1085–1121.
- Coque, M. and A. Gallais (2007) Genetic variation for nitrogen remobilization and postsilking nitrogen uptake in maize recombinant inbred lines: heritabilities and correlations among traits. Crop Sci. 47: 1787–1796.
- Duvick, D.N. (2005) The contribution of breeding to yield advances in maize (*Zea mays* L.). *In*: Donald, L.S. (ed.) Advances in Agronomy, Academic Press, pp. 83–145.
- El-Badawy, M.E.M. (2013) Heterosis and combining ability in maize using diallel crosses among seven new inbred lines. Asian J. Crop Sci. 5: 1–13.
- Gama, E.G., I.E. Marriel, P. Gurmares, S.N. Parentoni, M.X. Santos, C.A.P. Pacheco, W.F. Meireles, P.H.E. Ribeiro and A.C. Oliveira (2002) Combining ability for nitrogen use in a selected set of inbred lines from a tropical maize population. Revista Brasileira de Milho e Sorgo 1: 68–77.
- Good, A.G., A.K. Shrawat and D.G. Muench (2004) Can less yield more? Is reducing nutrient input into the environment compatible with maintaining crop production? Trends Plant Sci. 9: 597–605.
- Griffing, B. (1956) Concept of general and specific combining ability in relation to diallel crossing systems. Aust. J. Biol. Sci. 9: 463– 493.
- Guo, J.H., X.J. Liu, Y. Zhang, J.L. Shen, W.X. Han, W.F. Zhang, P. Christie, K.W. Goulding, P.M. Vitousek and F.S. Zhang (2010) Significant acidification in major Chinese croplands. Science 327: 1008–1010.
- Haegele, J.W., K.A. Cook, D.M. Nichols and F.E. Below (2013) Changes in nitrogen use traits associated with genetic improvement for grain yield of maize hybrids released in different decades. Crop Sci. 53: 1256–1268.

- Harvey, P.H. (1939) Hereditary variation in plant nutrition. Genetics 24: 437–461.
- Hill, R.R. Jr., E.C. Bashaw, C.R. Funk, A.E. Foster, J.A. Lee, D.A. Sleper, N.L. Taylor, A.R. Hallauer, C.M. Brown, R.A. Forsberg *et al.* (1987) Principles of cultivar development. Volume 2. Crop species. Macmillan publishing company, p. 556.
- Hirel, B., P. Bertin, I. Quilleré, W. Bourdoncle, C. Attagnant, C. Dellay, A. Gouy, S. Cadiou, C. Retailliau, M. Falque *et al.* (2001) Towards a better understanding of the genetic and physiological basis for nitrogen use efficiency in maize. Plant Physiol. 125: 1258–1270.
- Hirel, B., J. Le Gouis, B. Ney and A. Gallais (2007) The challenge of improving nitrogen use efficiency in crop plants: towards a more central role for genetic variability and quantitative genetics within integrated approaches. J. Exp. Bot. 58: 2369–2387.
- Ju, X.T., G.X. Xing, X.P. Chen, S.L. Zhang, L.J. Zhang, X.J. Liu, Z.L. Cui, B. Yin, P. Christie, Z.L. Zhu *et al.* (2009) Reducing environmental risk by improving N management in intensive Chinese agricultural systems. Proc. Natl. Acad. Sci. USA 106: 3041–3046.
- Lafitte, H.R. and G.O. Edmeades (1994a) Improvement for tolerance to low soil nitrogen in tropical maize I. Selection criteria. Field Crops Res. 39: 1–14.
- Lafitte, H.R. and G.O. Edmeades (1994b) Improvement for tolerance to low soil nitrogen in tropical maize II. Grain yield, biomass production, and N accumulation. Field Crops Res. 39: 15–25.
- Le Gouis, J., D. Beghin, E. Heumez and P. Pluchard (2002) Diallel analysis of winter wheat at two nitrogen levels. Crop Sci. 42: 1129–1134.
- Mehta, H. and K.R. Sarkar (1992) Heterosis for leaf photosynthesis, grain yield and yield components in maize. Euphytica 61: 161–168.
- Munaro, E.M., K.E. D'Andrea, M.E. Otegui, A.G. Cirilo and G.H. Eyhérabide (2011) Heterotic response for grain yield and ecophysiological related traits to nitrogen availability in maize. Crop Sci. 51: 1172–1187.
- Olsen, S.R., C.V. Cole, F.S. Watanabe and L.A. Dean (1954) Estimation of available phosphorus in soils by extraction with sodium bicarbonate. USDA Circ. 939. U.S. Govt. Print. Office, Washington.
- Presterl, T., G. Seitz, M. Landbeck, E.M. Thiemt, W. Schmidt and H. Geiger (2003) Improving nitrogen-use efficiency in European

maize: estimation of quantitative genetic parameters. Crop Sci. 43: 1259–1265.

- SAS (1999) SAS User's Guide: Version 8, SAS Institute. Inc, Cary, NC.
- Schnable, P.S. and N.M. Springer (2013) Progress toward understanding heterosis in crop plants. Annu. Rev. Plant Biol. 64: 71–88.
- Shull, G.H. (1946) Hybrid seed corn. Science 103: 547–550.
- Tilman, D., K.G. Cassman, P.A. Matson, R. Naylor and S. Polasky (2002) Agricultural sustainability and intensive production practices. Nature 418: 671–677.
- Tollenaar, M. and E.A. Lee (2002) Yield potential, yield stability and stress tolerance in maize. Field Crops Res. 75: 161–169.
- Van Reeuwijk, L.P. (1992) Procedures for soil analysis. 3rd edn. ISRIC, Wageningen, p. 9.
- Walkley, A. (1947) A critical examination of a rapid method for determining organic carbon in soils: effect of variations in digestion conditions and of inorganic soil constituents. Soil Sci. 63: 251– 264.
- Worku, M., M. Bänziger, G. Schulte auf'm Erley, D. Friesen, A.O. Diallo and W.J. Horst (2012) Nitrogen efficiency as related to dry matter partitioning and root system size in tropical mid-altitude maize hybrids under different levels of nitrogen stress. Field Crops Res. 130: 57–67.
- Wu, H. and A. Matheson (2000) Analysis of half-diallel mating design with missing crosses: theory and SAS program for testing and estimating GCA and SCA fixed effects. Silvae Genet. 49: 130–137.
- Wu, Y., W. Liu, X. Li, M. Li, D. Zhang, Z. Hao, J. Weng, Y. Xu, L. Bai, S. Zhang *et al.* (2011) Low-nitrogen stress tolerance and nitrogen agronomic efficiency among maize inbreds: comparison of multiple indices and evaluation of genetic variation. Euphytica 180: 281–290.
- Xie, C., S. Zhang, M. Li, X. Li, Z. Hao, L. Bai, D. Zhang and Y. Liang (2007) Inferring genome ancestry and estimating molecular relatedness among 187 Chinese maize inbred lines. J. Genet. Genomics 34: 738–748.
- Zaidi, P.H., G. Srinivasan and C. Sanchez (2003) Relationship between line *per se* and cross perfomance under low nitrogen fertility in tropical maize (*Zea mays* L.). Maydica 48: 221–231.