



## Source-sink Potential of Maize Inbreds Grown under Rainfed Condition in Bangladesh

D. Devnath<sup>1</sup>, M. Hasanuzzaman<sup>1</sup>, S. Jui<sup>1\*</sup>, M. A. K. Azad<sup>1</sup> and N. Dahal<sup>1</sup>

<sup>1</sup>Department of Genetics and Plant Breeding, Hajee Mohammad Danesh Science and Technology University, Dinajpur, Bangladesh.

### Authors' contributions

This work was carried out in collaboration among all authors. All authors read and approved the final manuscript.

### Article Information

#### Editor(s):

(1) Dr. K. S. Vinayaka, Kumadvathi First Grade College, India.

#### Reviewers:

(1) Sandeep Kumar Singh, Banaras Hindu University, India.

(2) Ajay Kumar, Agriculture Research Organization, Israel.

Complete Peer review History: <http://www.sdiarticle4.com/review-history/57556>

Original Research Article

Received 24 March 2020

Accepted 01 June 2020

Published 11 June 2020

### ABSTRACT

The experiment was conducted at the Plant Breeding Research Field, Department of Genetics and Plant Breeding, Hajee Mohammad Danesh Science and Technology University, Dinajpur, Bangladesh during the period of November, 2018 to March, 2019. The experiment was conducted at randomized complete block design with three replications comprised fifteen inbred lines. The individual plot size and spacing were 2 m x 3 m and 40 cm x 60 cm respectively. Plant height (cm), cob length (cm), cob weight (g), cob diameter (cm), kernels/cob, 100-kernel weight (g) and yield/plant (g) were recorded from randomly selected ten plants/plot. To assess source (leaves) and sink (kernels), leaves/plant were counted at four crop growth stages, 30 DAP, 45 DAP, 60 DAP and 75 DAP (days after planting). Moreover, root lengths were measured four times at same interval and leaf length was measured at 75 DAP only. Four inbred lines, IL 15, IL 32, IL 33 and IL 35 appeared high yield potential as compared to other lines and the highest yield (477.02 g/plant) was obtained from IL35. Three lines, IL32, IL33 and IL35 produced high root length and leaves/plant so, these lines apparently found outstanding for synthesizing more starch and providing mechanical support against lodging. Two characters, cob diameter and 100-kernel weight had high heritability. Plant height showed negative correlation with yield/plant, hence short stature plant is suitable for maize cultivation. The character, cob diameter alone exerted maximum direct effect (0.86273), followed by

\*Corresponding author: E-mail: sohanajui96@gmail.com;

cob length (0.82606) and kernels/cob (0.77719). The negative direct effect of plant height (-0.7021) was not counter balanced and compensated by other indirect effects. The inbred lines, IL32, IL33 and IL35 were promising for future breeding. The accumulation of starch in kernels was not uniform in rainfed condition which resulted average low kernel yield potential in maize inbred lines and may be improved through providing good agronomic requirements.

*Keywords: Leaves counting; root growth; yield potential; parent selection.*

## 1. INTRODUCTION

Maize (*Zea mays* L.) is a monophyletic originated large grain cereal grown around the world but its major production areas are located in temperate regions [1]. Its native land is Mexico, derived from the wild species Teosinte (*Zeamexicana*) possesses  $2n=24$  chromosomes. The Mexican Valley of Tehuacan, is considered the center of origin where it was found about 2750 years ago of BC. It is a source of both food and feed, and present conversion of biofuel from this starch rich crop creates a threat on food and feed security around the world. If maize is used as supplementary food in our daily diets, protein uptake would significantly increase [2]. As a whole, this crop has global impact, especially staple food for 1 billion people of sub-Saharan and different countries of Latin America [3]. It is also an important cereal in Asia but mainly used as feed for livestock; rapid economic growth along with urbanization compel the farmers of Indian sub-continent to incline maize cultivation predominantly for fodder and feed [4]. Bangladesh is the moderate producer of maize with a production of 28, 45,691 metric tons [5] and about 32,88,102 metric tons in almost 9,89,582 acres area [6]. This country is ideal for successful maize production because it can be grown everywhere around the year [7]. Being a short day crop, it sheds pollen well and matures quickly at a temperature ranges from 20°C to 30°C, prevailing in our country. In the year 2018-2019, The United States was the largest producer of maize with a production of 366.3million metric tons [8] and China and Brazil rounded off the top maize producing countries. It is one of the most important cereal crops in the world after wheat and rice [9] because of its multifarious use and high quantity grain yield per unit area [10], is contributing billions of dollars to the global economy [11]. The hybrid maize varieties currently cultivating throughout the country, are imported from India, China and Thailand. Development of outstanding parental lines rather than import of hybrid seeds may offer a sustainable maize production technology

through cultivating locally produced hybrid maize varieties [12].

Rapid change in climate, there is a need to produce high yielding hybrid maize varieties exploiting the locally developed inbred lines. The performance of inbred lines tested in rainfed condition is helpful in identification of superior inbred lines that may be exploited for the production of high yielding hybrid maize through appropriate incorporation in hybridization [13]. There is a strong relation between dry-matter production and grain yield in maize, minimum dry-matter production after harvest suggests low source at vegetative phase that eventually results low quantity grain yield due to limitation of sink capacity [14]. In addition, rapid maturation followed by bending down of leaves, indicate minimal photosynthetic product, sugar, is the principal source for the sink (kernels) [15], the physiological state of kernels ultimately suffers from possible utilization of sugar [16], which leads to develop infertile kernels in the cobs.

The efficiency of selection depends upon heritability and the genetic advance of individual character [17] and heritability estimates are useful for breeding of quantitative characters because it permits to determine the most effective selection strategy, breeding method used in a breeding program, is to predict gain from selection [18]. Information regarding interrelationship between quantitatively inherited characters and their direct and indirect effects on grain yield are of great importance for success in selection to be conducted in breeding programs [19]. In path coefficient analysis, grain yield is considered as dependent variable and the remaining characters are considered as independent variables [20]. Moreover, grain yield is a complex character which is influenced by several component characters and direct selection for grain yield is often not be effective, therefore phenotypic association [21] and cause and effects of the components are to be assessed through appropriate biometrical tools

[22]. Therefore, before going to test combining ability of the developed parental lines, heritability of yield enhancing characters and association of these characters were investigated in this study.

## 2. MATERIALS AND METHODS

### 2.1 Experimental Design and Field Operation

The field experiment was carried out at the Plant Breeding Research Farm, Department of Genetics and Plant Breeding, Hajee Mohammad Danesh Science and Technology University, Dinajpur, Bangladesh during the period from November, 2018 to March, 2019. Fifteen inbred lines were received from the departmental store and these lines were developed through successive selfing/inbreeding up to 7<sup>th</sup> generation. The experiment was conducted following randomized complete block design with three replications. The unit plot size and spacing were 3 m x 5 m and 20 cm x 60 cm respectively. The inbred lines were developed from exotic hybrid maize varieties through continuous selfing/inbreeding under the maize improvement program carried out by the Department of Genetics and Plant Breeding, Hajee Mohammad Danesh Science and Technology University, Dinajpur, Bangladesh. All agronomic requirements were fulfilled, except no supplementary irrigation as the strategy of the experiment was rainfed condition. Rouging was followed began from germination and continued prior to harvesting in the field. Just after attaining 5% flowering the individual plants were selfed or inbred to maintain homozygosity in the individual parental lines. The cobs were protected from bird attack through netting just after completion of artificial selfing/inbreeding. Data were recorded on plant height (cm), root length (cm) and leaves/plant at 30, 45, 60 and 75 DAP (days after planting) and leaf length (cm) at 75 DAP, cob length (cm), cob diameter (cm), cob weight (g), kernels/cob, 100-kernel weight (g) and yield/plant (g).

### 2.2 Statistical Analysis of the Recorded Data

Mean data were statistically analyzed for each character separately. The analysis of variance for each of the characters under study was performed by F test using GENSTAT 5.7.1 [23]. The significant differences for the studied characters among the inbreds were performed by

the Duncan's Multiple Range Test (DMRT) at 5% level of probability [24]. Genetic parameters were estimated following the method proposed by Johnson et al. [25] and correlation coefficient and path coefficient were estimated according to [26]. Genetic advance was calculated following formula given by Johnson et al. [25] and Allard [27] and as percent of mean by the formula of Comstock et al. [28].

## 3. RESULTS AND DISCUSSION

### 3.1 Crop husbandry and Evaluation on Inbred Lines

In the present study yield/plant along with eight yield related characters, plant height (cm) leaves/plant, leaf length (cm), cob length (cm), cob diameter (cm), cob weight (g), kernels/cob, 100-kernel weight (g) and root length (cm) were measured to assess the contributions of the characters on yield in maize inbred lines. The results obtained from the study along with discussion of the results are described below-

The means of yield and yield related characters are presented in Table 1. The means of a particular character against fifteen inbred lines are separated by DMRT. The mean plant height was 218.66 cm and lower plant height was recorded from four inbred lines, IL15, IL32, IL33 and IL35. During crop husbandry, earthing up was indispensable to protect the inbred lines from lodging and such intercultural operation was laborious and costly, therefore, short stature inbred lines are suitable in maize breeding program. Number of leaves/plant were obtained at four different intervals started from 30 DAP to 75 DAP, hence the interval was 15 days. It is observed from Table 2 that leaves/plant gradually increased with corresponding increase of DAP. The range of leaves/plant was 7.33 to 20.09 among the inbred lines. Since maize is a C4 plant, it utilized a 4-carbon sugar during the Calvin cycle and up took more CO<sub>2</sub> to synthesize the source (starch) through photosynthesis for the sink (kernels). The process of transferring assimilated source in the green tissues into the developing kernels was complicated the source versus sink question, as it was also affected by the environment, time, and the genetic makeup of the plants [29]. Among the fifteen inbred lines, IL32, IL33 and IL35 apparently produced remarkable leaves/plant while counted at 4 stages, suggest these lines are desirable to produce more source for the kernels (sink). The mean cob length was estimated to 17.47 cm and

highest length was obtained from the line IL15 (21.83 cm), which was statistically similar with the results obtained in IL 33 and IL 35. Similar finding was reported by Rathore et al. [30]. The highest mean cob diameter was obtained from the line IL 33 (4.81 cm), which was statistically similar with the results obtained in IL 35. The cob diameter had effect to increase yield/plant (Table 1), therefore, it is an important character for improving maize yield potential. The highest kernels/cob (478.51) was recorded in IL 35 and it was significantly higher than other inbred lines, suggests superiority of the line for this character; breeders may consider the character during breeding programs. The character, 100-kernel weight though varied among the inbred lines in statistical sense, but variation was not remarkable, however, the highest 100-kernel weight (41.11 g) was obtained from the line IL35 and it was statistically similar to the line, IL 5 with 39.56 g; other lines were significantly lower than the two lines for the character. Source capacity was determined by the photosynthetic activity of the crop and by the availability of carbohydrate reserved and variation in final kernel weight reflected the interaction between source capacity and sink strength [31]. The resultant character, yield/plant was found the highest in IL35 with 477.02 g, which significantly higher than the yield production by other inbred lines. The second rank yield/plant was recorded from IL 32 with 451.10 g, which was significantly higher than the other thirteen inbred lines. The yield was increased in these inbred lines due to increase of kernels per cob as well as size of kernels [32,33] and the results of present study are favorably compared with the findings reported by several authors [34,35,36,37,38,39]. In conventional breeding, always interested to consider the final product like yield potential, therefore, the lines, IL35, IL 32 or even the third rank lines, like IL15 with 401.90 g/plant and IL 33 with 406.49 g/plant might exploit in future breeding programs to improve maize yield potential.

### 3.2 Contribution of Roots and Leaves during Crop Growth

High yield potential is directly related to high depth of root system [40], this information may apply to assess relationship of root system with shoot growth in maize inbred lines, necessary for breeding programs.

The mean values having same letter (s) were not different significantly at 5% level of probability. Root length along with number of leaves/plant

were recorded at four different stages of crop growth, like 30DAP, 45DAP, 60DAP and 75 DAP. The results presented in Table 2 revealed that there was a harmony between root length and leaves/plant at different stages and the values increased with corresponding increase of DAP. Since roots function as carrier of water and nutrients from soil to different parts of shoot, increase in root length suggests absorption of water and nutrients from different levels of soil. Furthermore, the elongated roots provided mechanical support to protect the crop from lodging. However, at 75DAP, noticeable root length vs leaves/plant were revealed by the lines, IL32 (root length vs leaves/plant= 67.30 cm vs 24.10), IL33 (root length vs leaves/plant= 61.16 cm vs 24.90) and IL35 (root length vs leaves/plant= 62.71 cm vs 25.12) (Table 2). The results suggest that these three lines provided strong mechanical support to keep the plants erect at pre-flowering stage of crop growth. Besides, long roots synthesized more cytokinins which later on translocated to different parts of shoot. There is a high scope to isolate desirable inbred lines against lodging considering long root length in maize [41].

### 3.3 Study on Genetic Parameters of Different Characters of Maize Inbred Lines

From the Table 3 it is revealed that difference between genotypic coefficient variation and phenotypic coefficient of variation for a particular character was very close, suggests inherent genetic potential of the inbred lines to express the characters but variation might be created by changing the growing condition. Heritability estimates were higher for the character like, 100-kernel weight (86.33%), followed by cob length (84.27%) but none of the characters showed appreciable heritability coupled with genetic advance, suggests low progress under selection in the evaluated inbred lines. Analysis of variance pertaining to plant height revealed highly significant differences among the maize inbred lines [42,43]. High heritability coupled with a high range of genotypic coefficient of variation (GCV) and phenotypic coefficient of variation (PCV) for plant height suggests that genetic improvement might be achieved through simple selection for the character [44,45,46]. Ilyas et al. [47] studied two sets of maize inbred lines and found that both heritability and genetic advance estimates were higher for grain yield and plant height, hence additive gene action was depicted. High heritability along with high genetic advance

**Table 1. Mean performance of yield and yield contributing characters of maize inbred lines**

Treatment	Plant height (cm)	Leave length (cm)	Cob length (cm)	Cob diameter(cm)	Cob weight(g)	Kernels/cob	100- kernel weight (g)	Yield/plant (g)
IL1	214.00 cd	65.77c	17.11 cd	4.43cd	145.88 f	288.21 f	38.24b	320.08 f
IL3	214.66 cd	71.33b	15.87 d-f	4.57 a-c	161.23e	324.77e	38.29 b	349.69 e
IL5	229.53 ab	73.67ab	14.75 f	4.60 a-c	186.78c	392.00c	39.56 a	155.40 j
IL6	224.83bc	66.00c	17.04 cd	4.27e	136.79 g	275.33f	37.11 de	304.31 g
IL7	214.66 cd	73.00ab	19.17c	4.59 a-c	161.20e	329.33e	38.19 bc	351.80 e
IL11	229.16 ab	72.00b	14.89 ef	4.57 a-c	105.72i	211.07gh	36.69 ef	254.81 h
IL15	211.33 e	71.33b	21.83a	4.60ab	187.61c	394.07 c	38.29 b	401.90 c
IL21	228.66 b	72.07b	16.17 de	4.45c	89.99 i	179.44 h	36.60 ef	245.09 h
IL22	214.06 cd	72.00b	17.50 cd	4.53 a-c	157.85e	319.67e	38.35 b	345.31 e
IL23	216.53 c	64.71 c	10.17 g	4.31 e	117.49h	230.33g	37.49 cd	192.13 j
IL26	233.33 a	72.47a	17.26 cd	4.30de	154.72e	331.67e	35.89 f	220.69 i
IL30	213.16 d	73.83ab	17.91 cd	4.51 bc	173.99 d	360.96 d	38.31 b	376.68 d
IL32	212.43 de	74.71 ab	20.18 b	4.33 bc	211.11 b	454.11 b	38.93b	451.10 b
IL33	211.80 e	74.52 ab	21.07a	4.81 a	187.72c	401.55 c	38.15bc	406.49 c
IL35	211.73 e	76.11 a	21.11 a	4.75 a	239.31 a	478.51 a	41.11a	477.02 a
LSD(0.05)	7.10	3.54	4.12	2.14	8.33	19.18	1.17	14.69
Mean	218.66	71.57	17.47	4.51	161.16	331.41	38.08	323.5

Here, IL=Inbred line

**Table 2. Root length vs Number of leaves/plant estimates at different days after planting of maize inbred lines**

Genotypes	30 DAP		45 DAP		60 DAP		75 DAP	
	RL	NLPP	RL	NLPP	RL	NLPP	RL	NLPP
IL1	18.60 e	7.76 de	32.00 c-f	11.33 de	42.86 de	16.33 bc	63.16 f	21.27 d
IL3	20.58 e	7.12 de	32.43 c-e	12.00 cd	43.83 c-e	16.00 c	65.16 d-f	19.61 e
IL5	16.06 f	5.91 f	30.66 e-g	10.00 g	41.83 e	13.00 f	61.88 f	17.28 g
IL6	19.74 e	7.33 c-e	33.20 bc	11.33 de	44.20 b-d	14.67 d	65.33 d-f	18.00 fg
IL7	21.66 c-e	7.67 b-d	34.80 ab	12.33 c	45.70 a-c	14.67 d	69.80 bc	17.90 fg
IL11	17.33 f	6.00 f	31.20 d-f	10.33 fg	42.43 de	13.33 ef	62.40 f	17.96 fg
IL15	23.16 ab	8.22 a-c	35.11 a	14.10 a	55.40 a	17.33 b	79.19 a-b	13.20 h
IL21	21.56 c-e	6.67 ef	31.90 c-f	10.88 ef	42.50 de	13.67 d-f	67.80 b-e	18.67 ef
IL22	21.10 de	7.37 c-e	33.83 bc	11.33 de	46.20 ab	14.33 de	64.80 d-f	18.33 f
IL23	15.89 f	7.07 de	29.20 gh	11.33 de	42.00 de	16.00 c	64.31 ef	21.90 cd
IL26	21.16 de	6.67 ef	33.83 bc	11.21 ef	45.96 a-c	16.00 c	67.26 c-e	21.00 d
IL30	22.00 b	7.16 b-d	33.03 b-d	11.67 c-e	44.06 b-e	17.00 bc	68.53 b-d	22.11 cd

Genotypes	30 DAP		45 DAP		60 DAP		75 DAP	
	RL	NLPP	RL	NLPP	RL	NLPP	RL	NLPP
IL32	22.54 b	7.44 b-d	28.53 h	12.11 c	34.80 f	17.00 bc	67.30 c-e	24.10 b
IL33	24.19 bc	9.11 a	28.46 h	13.61 b	34.95 f	19.67 a	61.16 ab	24.90 b
IL35	26.00 a	8.19 ab	31.26 fg	13.14 b	37.00 f	19.88 a	62.71 a	25.12 a
LSD(0.05)	2.37	0.88	1.28	0.91	2.11	1.31	3.44	1.81
Mean	20.77	7.32	31.96	11.78	42.38	15.92	66.05	20.09

Here, RL= Root length, NLPP= Number of leaves/plant, DAP= Days after planting

**Table 3. Estimates of coefficients of variation, heritability in broad sense, genetic advance and genetic advance in percentage of mean for various characters in maize inbred lines**

Characters	Mean	Coefficient of variation (%)		Heritability in broad sense (%)	Genetic advance	Genetic advance as percent of mean
		Genotypic	Phenotypic			
PH	218.66	10.35	11.47	67.77	14.56	21.09
LL	71.57	4.96	5.79	73.5	20.12	8.77
CL	17.47	22.97	23.66	84.27	8.82	45.95
CD	4.51	2.79	3.35	69.12	1.44	4.77
CW	161.16	23.95	24.09	68.77	2.59	49.02
KPC	331.41	25.34	26.59	78.02	20.96	51.68
HKW	38.08	2.72	3.93	86.33	8.29	5.21
YPP	323.5	37.74	38.33	68.99	2.08	57.34

Here, PH= Plant height, LL= Leave length, CL = Cob length, CD = Cob diameter, CW = Cob weight, KPC = Kernel per cob, HKW = Hundred kernel weight and YPP = Yield per plant

**Table 4. Genotypic ( $r_g$ ) and phenotypic ( $r_p$ ) correlation coefficient among pairs of characters in maize inbred lines**

Characters	LL		CL		CD		CW		KPC		HKW		YPP	
	$r_g$	$r_p$	$r_g$	$r_p$	$r_g$	$r_p$	$r_g$	$r_p$	$r_g$	$r_p$	$r_g$	$r_p$	$r_g$	$r_p$
PH	0.63	0.91	-0.72*	-0.81*	-0.54	-0.77	-0.69*	-0.57*	-0.7*	-0.21*	-0.43*	-0.62*	-0.79	-0.48
LL			0.52**	0.89**	0.59	0.37	0.62*	0.28*	0.63**	0.81**	0.41*	0.72*	0.56**	0.70**
CL					0.49*	0.71*	0.66**	0.69**	0.68**	0.35**	0.4	0.11	0.69**	0.53**
CD							0.51**	0.45**	0.49**	0.72**	0.56	0.24	0.57**	0.55**
CW									0.99**	0.65**	0.76**	0.35**	0.81**	0.89**
KPC											-0.69	-0.42	0.8**	0.88**
HKW													0.6**	0.79**

\*\*Indicate significant at 5% level of probability, Here, PH= Plant height, LL= Leave length, CL = Cob length, CD = Cob diameter, CW = Cob weight, KPC = Kernel per cob, HKW = Hundred kernel weight and YPP = Yield per plant

was reported by Kinfе and Tsehayе [48] for plant height and grain yield. Bekele and Rao [49] recorded high heritability with high genetic advance for 100-kernel weight in maize. Whatever, heritability against yield/plant and genetic advance even as percentage of mean were lower in magnitudes, invariably indicates yield was influenced by yield influencing characters along with growing condition, breeding method may apply for discriminant function to integrate appropriate character in selection to get gain in next generation.

### 3.4 Characters Association and Partitioned of Association in Relation to Kernel Yield

Correlation coefficients were measured both at genotypic and phenotypic levels (Table 4). The results where genotypic correlation coefficient was higher than corresponding phenotypic correlation coefficient suggest inherent potential of the inbred lines to make such association between the pair of characters concerned, while phenotypic correlation coefficient was higher than corresponding genotypic correlation coefficient, suggests the association may be changed upon changing the growing conditions. All the yield related characters showed positive

and significant associations both at genotypic and phenotypic levels, except plant height, which exhibited negative and non-significant association with yield/plant. Plant height had significant but negative correlation with the yield at both genotypic and phenotypic levels [50]. A significant correlation was revealed between plant height and grains per cob [51].

The results suggest that the selected characters, except plant height had prominent effect on increasing yield/plant maize inbred lines. Since genotypic variation and environmental variation are covered by phenotypic variation, only phenotypic correlation coefficients of different characters with yield/plant were portioned into direct and indirect effects (Table 5) and the cause and effect of the characters are presented by the Fig. 1. Among the selected characters, cob diameter exerted maximum direct effect (0.86273), followed by cob length (0.82606) and kernels/cob (0.77719). Prakash et al. [52] reported positive direct of cob diameter on yield/plant in maize. Munawar et al. [53] reported that cob length had high positive direct effect on grain yield. Furthermore, cob length (cm) and cob diameter (mm) had direct and positive effect on grain yield/plant [54]. The negative indirect effect of plant height was not counter balanced

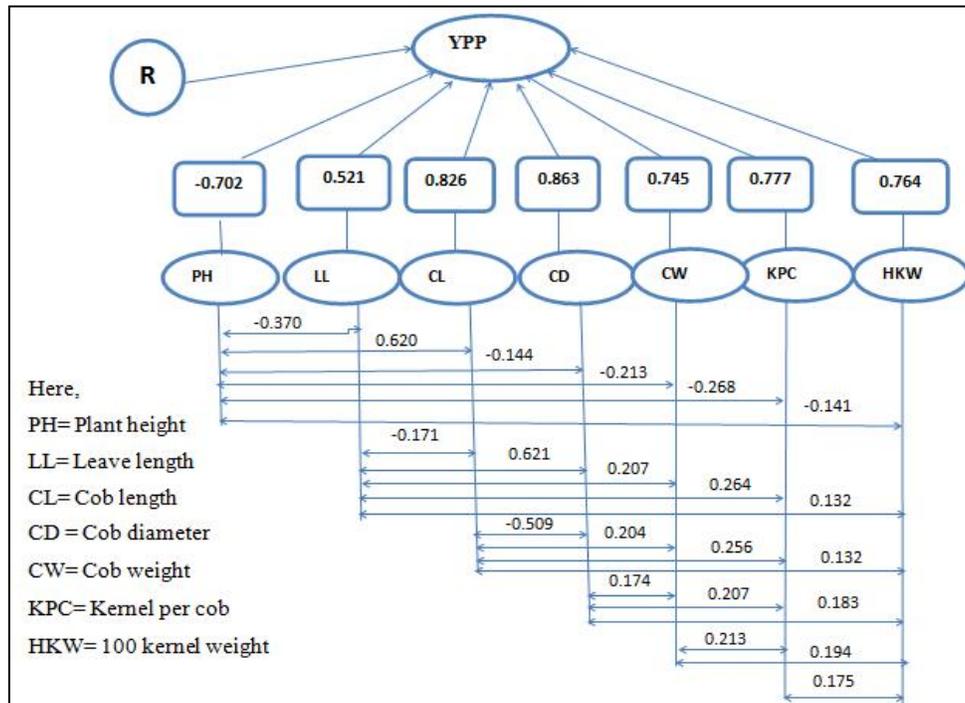


Fig. 1. Path diagram showing yield contributing characters on yield per plant in maize

**Table 5. Phenotypic path coefficients of different characters with correlation coefficients ( $r_p$ ) of different characters of maize inbred lines**

Characters	PH	LL	CL	CD	CW	KPC	HKW	YPP
PH	<b>-0.7021</b>	0.36972	0.61955	-0.1435	-0.2133	-0.2678	-0.14132	-0.48
LL	-0.8725	<b>0.52073</b>	-0.1709	0.62115	0.20718	0.26403	0.13249	0.70**
CL	-0.6766	0.29682	<b>0.82606</b>	-0.509	0.20414	0.25649	0.13249	0.53**
CD	-0.7644	0.37493	-0.4874	<b>0.86273</b>	0.17367	0.20745	0.18254	0.55**
CW	-0.9215	0.2541	0.21346	0.19174	<b>0.74468</b>	0.21342	0.19436	0.89**
KPC	-0.8925	0.16451	0.26172	0.19449	0.20163	<b>0.77719</b>	0.17567	0.88**
HKW	-0.849	0.03433	0.12173	0.17488	0.25289	0.29044	<b>0.76441</b>	0.79**

Residual effect=0.039, the bold figures in the diagonal indicate direct effects and other figures are indirect effects. Here, PH= Plant height, LL= Leave length, CL = Cob length, CD = Cob diameter, CW = Cob weight, KPC = Kernel per cob, HKW = Hundred kernel weight and YPP = Yield per plant

and compensated by other indirect effects exerted by other characters that eventually developed negative and non-significant correlation with kernel yield/plant. Plant height showed negative direct effect on grain yield/plant [55]. The residual effect was only 0.039, suggests that 96.1% variation had included during path analysis of the characters. The source-sink relationship during kernel development period is the key information in selection of superior maize genotypes [56] and this information is relevant to identify optimal methods for crop modelling and to select breeding strategies [57].

#### 4. CONCLUSION

In Bangladesh the area under hybrid maize cultivation is increasing due to its high yield and reasonable market price as compared to other cereals. The agricultural entrepreneurs are showing interest to invest capital in this sector but most of the hybrid seeds are importing from China, India and Thailand expending huge amount of foreign currency every year. Our locally developed hybrids are still low yield potential compared to exotic hybrids. Due to absence of outstanding parental lines different agencies of Bangladesh are yet not become successful to produce a hybrid exceeding the demand of the farmers for foreign hybrid seeds. Considering these issues, superior parental lines have been developed to have better combinations for the development of outstanding hybrid maize varieties. A consistent contribution of kernel yield components and source-sink relationship became more limiting in rainfed condition, suggests optimized irrigation during kernel filling stage of the inbred lines of maize. Finally, it is concluded, the inbred lines appeared as promising may exploit either by developing hybrid varieties and synthetic varieties in future

through applying appropriate breeding programs for enhancing yield potential of hybrid maize cultivation in Bangladesh.

#### COMPETING INTERESTS

Authors have declared that no competing interests exist.

#### REFERENCES

- Patil SM, Kumar K, Jakhar DS, Rai A, Borle UM, Singh P. Studies on variability, heritability, genetic advance and correlation in maize (*Zea mays* L.). Int. J. Agric. 2016;9(6):1103-1108.
- Uddin H, Rashid MH, Akhter S. Relative profitability of maize production under different farm size groups in Kishoregonj District of Bangladesh. Progressive Agric. 2010;21(1-2):247-52.
- Gupta HS, Agrawal PK, Mahajan V, Bisht GS, Kumar A. Quality protein maize for nutritional security: Rapid development of short duration hybrids through molecular marker assisted breeding. Curr. Sci. 2009;96:230-237.
- Prasanna BM, Vasal SK, Kassahun B, Singh NN. Quality protein maize. Curr. Sci. Assoc. 2001;81:1308-1319.
- Bangladesh Bureau of Statistics (BBS). Yearbook of Agricultural Statistics-2018. 28<sup>th</sup> Series. Statistics and Informatics Division (SID). Ministry of Planning, Government of the People's Republic of Bangladesh; 2019.
- Bangladesh Bureau of Statistics (BBS). Yearbook of Agricultural Statistics-2017. 28<sup>th</sup> Series. Statistics and Informatics Division (SID). Ministry of Planning,

- Government of the People's Republic of Bangladesh; 2018.
7. Bangladesh Bureau of Statistics (BBS). Yearbook of Agricultural Statistics-2016. 28th Series. Statistics and Informatics Division (SID). Ministry of Planning, Government of the People's Republic of Bangladesh; 2017.
  8. Shahbandeh M. Corn production worldwide 2018/2019, by country; 2020.
  9. Golbashy M, Ebrahimi M, Khorasani SK, Choukan R. Evaluation of drought tolerance of some corn (*Zea mays* L.) hybrids in Iran. African J. Agril. Res. 2010; 5(19):2714-2719.
  10. Akbar M, Shakoor MS, Hussain A, Sarwar M. Evaluation of maize 3 way crosses through genetic variability, broad sense heritability, characters association and path analysis. J. Agril. Res. 2008;46(1): 39-45.
  11. FAO (Food and Agriculture Organization). FAO corporate document repository: Economic and Social Development Department; 2012.  
Available: <http://www.fao.org/agriculture/food/grain/in/developing/countries.html> (Accessed 14 April 2013)
  12. Azad MAK, Biswas BK, Alam N, Alam SS. Genetic Diversity in Maize (*Zea mays* L.) Inbred Lines. The Agriculturists. 2012; 10(1):64-70.
  13. Taiwo OP, Nwonuala AI, Isaiah BF. Variability in yield and yield components of selected pro-vitamin a maize (*Zea mays* L.) Varieties in a Humid Environment of Port Harcourt, Nigeria. Asian J. Agril. Hort. Res. 2020;19:1-0.
  14. Zhang H, Turner NC, Poole ML. Source-sink balance and manipulating sink-source relations of wheat indicate that the yield potential of wheat is sink-limited in high-rainfall zones. Crop and Pasture Sci. 2010; 61(10):852-861.
  15. Egile DB. Variation in leaf starch and sink limitations during seed filling in soybean. Crop Sci. 2000;39:1361-1368.
  16. Burton WJ. Effects of defoliation on seed protein concentration in normal and high protein lines of soybean. Crop Sci. 2004; 72:131-139.
  17. Bilgin O, Korkut KZ, Başer I, Dağlioğlu O, Öztürk I, Kahraman T, Balkan A. Variation and heritability for some semolina characteristics and grain yield and their relations in durum wheat (*Triticum durum* Desf.). World J. Agril. Sci. 2010;6(3):301-308.
  18. Singh BD. Plant breeding: principles and methods. Kalyani publishers; 2015.
  19. Rutkoski JE. Estimation of Realized Rates of Genetic Gain and Indicators for Breeding Program Assessment. Adv. Agron. 2019;157:217-249.
  20. Singh AK, Mishra SP, Parihar R. Studies on genetic variability parameters on grain yield and its yield attributing characters in maize (*Zea mays* L.). Int. J. Curr. Microbiol. Appl. Sci. 2018;7(09):2148-2150.
  21. Hallauer AR, Miranda JB. Quantitative Genetics in Maize Breeding. (2<sup>nd</sup>edn.), Iowa State University Press, Iowa, Ames, USA; 2016.
  22. Dewey DR, Lu KA. Correlation and path-coefficient analysis of components of crested wheatgrass seed 1. Agron. J. 2017;51:515-518.
  23. Cochran WG, Cox GM. Experimental Designs. 2<sup>nd</sup>edn. Wiley, New York; 1960.
  24. Gomez KA, Gomez AA. Statistical Procedure for Agricultural Research (2<sup>nd</sup> edn.). International Rice Research Institute. A Willey International Science Publication. 1984;28-192.
  25. Johnson HW, Robinson HF, Comstock RE. Estimates of genetic and environmental variability in soybeans. Agron. J. 1955; 47(7):314-318.
  26. Singh RK, Chaudhary BD. Biometrical methods in quantitative genetic analysis. Biometrical Methods in Quantitative Genetic Analysis. 1979;10:304.
  27. Allard R. Principles of plant breeding. John Wiley and Sons Inc., New York; 1975.
  28. Comstock R, Robinson H, Gowen, J. Estimation of average dominance of genes, Heterosis. 1952;494-516.
  29. Seebauer JR, Singletary GW, Krumpelman PM, Ruffo ML, Below FE. Relationship of source and sink in determining kernel composition of maize. J. Expt. Bot. 2010; 61(2):511-9.
  30. Rathore DM, Singh K, Singh BP. Effect of nitrogen and plant population on the yield attributes of maize. Indian J. Agric. Res. 1976;10(2):79-82.
  31. Shekoofa A, Emam Y, Pessaraki M. Source-sink manipulation effects on maize

- kernel quality. J. Pl. Nutri. 2013;36(9): 1401-11.
32. Borrás L, Slafer GA, Otegui ME. Seed dry weight response to source–sink manipulations in wheat, maize and soybean: A quantitative reappraisal. Field Crops Res. 2004;86:131–146.
  33. Rao IM, Beebe SE, Polania J, Grajales M, Cajiao C, Ricaurte J. Evidence for genotypic differences among elite lines of common bean in the ability to remobilize photosynthates to increase yield under drought. J. Agric. Sci. 2017;155:857–875.
  34. Simeonov N, Tsankova G. Effect of fertilizers and plant density on yield of maize hybrids with two ears. Rasteniiev'dni Nauki. 1990;27(8):14-8.
  35. Narayanaswamy MR, Veerabadran V, Joyanthi C, Chinnuswamy C. Plant density and nutrient management for rainfed maize in red soil. Madras Agril. J. 1994;81(5):248-251.
  36. Hassen H. Effect of defoliation on yield components of maize and under sown forage. Agril. Trop. 2003;18(1/2):5-7.
  37. Zewdu T. Effect of defoliation and intercropping with forage legumes on maize yield and forage production. Trop. Sci. 2003;43(4):204-207.
  38. Li-Xiangjun, AnpingInanaga S, Enejj AE, Ali AM. Mechanisms promoting recovery from defoliation in determinate and indeterminate soybean cultivar. J. Food Agric. Env. 2005;3(3/4):178-183.
  39. Chaudhary AN, Latif MI, Haroon, Ur-Rasheed M, Jilani G. Profitability increase in maize production through fertilizer management and defoliation under rain fed cropping. Int. J. Bio. Biotech. 2005;2(4): 1007-1012.
  40. Koffler NF. A profundidade do sistema radicular e o suprimento de água às plantas no Cerrado. Piracicaba: POTAFOS. (Informações Agronômicas, 33). 1986; 12.
  41. Bray AL, Topp CN. The quantitative genetic control of root architecture in maize. Pl. Cell Physiol. 2018;59(10):1919-1930.
  42. Noor M, Rahman H, Durrishahwar MI, Shah SMA, Ullah I. Evaluation of maize half-sib families for maturity and grain yield attributes. Sarhad J. Agric. 2010;26:545-549.
  43. Stromberg DC, Campton WG. Ten cycles of full-sib recurrent selection in maize. Crop Sci. 1989;29:1170-1172.
  44. Mahmood Z, Malik SR, Akhtar R, Rafique T. Heritability and genetic advance estimates from maize genotypes in ShishiLusht a valley of Krakurm. Int. J. Agric. Biol. 2014;6:790-791.
  45. Mustafa HS, Ahsan M, Aslam M, Ejaz-ul-Hasan QA, Bibi T, Mehmood T. Genetic variability and traits association in maize (*Zea mays* L.) Accessions under drought stress. J. Agric. Res. 2013;51:231-238.
  46. Vashistha A, Dixit NN, Dipika, Sharma SK, Marker S. Studies on heritability and genetic advance estimates in Maize genotypes. Biosci. Discov. 2013;4:165-168.
  47. Ilyas M, SA Khan, Awan SI, Rehman S. Assessment of heritability and genetic advance in maize (*Zea mays* L.) under natural and water stress conditions. Sarhad J. Agric. 2019;35(1):144-154.
  48. Kinfe Hll Tsehaye Y. Studies of heritability, genetic parameters, correlation and path coefficient in elite maize hybrids. Academic Res. J. Agril. Sci. Res. 2015;3(10):296-303.
  49. Bekele A, Rao TN. Estimates of heritability, genetic advance and correlation study for yield and its attributes in maize (*Zea mays* L.). J. Pl. Sci. 2014; 21:4-6.
  50. Khatun F, Begum S, Motin A, Yasmin S, Islam MR. Correlation coefficient and path analysis of some maize (*Zea mays* L.) hybrids. Bangladesh J. Bot.1999;28:9–15.
  51. Alaei, Y. Correlation analysis of corn genotypes morphological traits. Int. Res. J. Appl. Basic Sci. 2012;3:2355-2357.
  52. Prakash R, Ravikesavan R, Vinodhana NK, Senthil A. Genetic variability, character association and path analysis for yield and yield component traits in maize (*Zea mays* L.). Electron. J. Pl. Bred. 2019;10(2):518-524.
  53. Munawar M, Shahbaz M, Hammad G, Yasir M. Correlation and path analysis of grain yield components in exotic maize (*Zea mays* L.) hybrids. Int. J. Sci.: Basic Appl. Res. 2013;12(1):22-27.
  54. Huda MN, Hossain MS, Sonom M. Genetic variability, character association and path analysis of yield and its component traits in maize (*Zea mays* L.). Bangladesh J. Pl. Bred. Genet. 2016;29(1):21-30.

55. Selvaraj CI, Nagarajan P. Interrelationship and path-coefficient studies for qualitative traits, grain yield and other yield attributes among maize (*Zea mays* L.). *Int. J. Pl. Bred. Genet.* 2011;5(3):209-223.
56. Bonelli LE, Monzon JP, Cerrudo A, Rizzalli RH, Andrade FH. Maize grain yield components and source-sink relationship as affected by the delay in sowing date. *Field Crops Res.* 2016;198:215-25.
57. Hall AJ. Wither crop physiology? In 'Crop physiology: applications for genetic improvement and agronomy'. (Eds V Sadras, D Calderini). 2009;545-570.

© 2020 Devnath et al.; This is an Open Access article distributed under the terms of the Creative Commons Attribution License (<http://creativecommons.org/licenses/by/4.0>), which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.

*Peer-review history:*

*The peer review history for this paper can be accessed here:  
<http://www.sdiarticle4.com/review-history/57556>*