



## Comparison of selection indices in forage maize (*Zea mays* L.) hybrids in normal and salt stress conditions

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### Extended abstract

#### Introduction

To improve a complex character such as grain yield with low heritability, indirect selection through other characters and selection index based on different effective traits were used. Grain yield has quantitative heritance and can be affected by environment severely; therefore selection for genetic improvement only based on yield may have low efficiency. But selection based on proper index can be one of the most effective methods for indirect selection of yield and yield components simultaneously.

#### Materials and methods

In order to determine selection index for improvement of maize yield, 14 single cross maize hybrids (including 12 promising maize hybrids and KSC704 and KSC705 cultivars as control cultivars) were planted in two separate experiments (Saline stress and normal condition) based on randomized complete block design (RCBD) with four replication in Khorasan Razavi agricultural and natural resources research and education center (TOROQ Station and Abbas Abad Station), Mashhad Iran on 2017-2018. In this study silage yield, Dry Forage yield, number of total leaves, Ear Diameter, Ear Length, Ear Height, Kernel depth, anthesis silking interval (ASI) and Plant Height appearance was measured randomly from 10 sample. Then some of morphological and phenological traits, yield and yield components were recorded.

#### Results and discussion

The results of ANOVA showed significant differences between hybrids for many of measured traits in both conditions ( $P \leq 0.01$ ). Thus, selection will be effective due to existence of enough variation. The results of correlation, multiple regression and principle component analysis were used for identification of traits that are more effective on grain yield. Selection indices were calculated based on results of stepwise regression considering to phenotype, genotypic and economic values. Based on stepwise regression results in normal condition, Plant Height, Number of Ear, Dry Forage Yield, Days to anthesis, Number of Leaves totally could explain 77.84 percent of gain yield variation, then these traits were used to calculate selection index. In Saline stress condition, Number of Ear, Ear Length, Days to anthesis, Number of Leaves, Plant height could explain 76.90 percent of grain yield variation that these traits were used to calculate of selection index. Smith-Hazel and Pesk-Baker selection indexes for dry silage yield performance, leaf total number, number of cob, plants length and days to pollination in non-stressed condition and number of cob, days to pollination, leaf total number and plants length were calculated

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under stressed situation. Moreover, relative efficiency of selection and expected gain of selection index using the Smith – Hazel index was higher than the Pesk – Baker index. The highest relative efficiency of selection under non-stressed condition was measured in index number 5 (Smith – Hazel 5) while in saline stressed condition it was achieved in index number 4 (Smith - Hazel 4).

### Conclusion

In summary, by adjusting phenotype values in mentioned traits in index equivalent, the amount of each index was determined. Finally Based on grain yield and selection indices, 20 percent of the best genotypes were selected by each selection indices. The highest selection indices were obtained for the hybrids 1, 5, 2, 8 and 6 in normal condition and hybrids 13, 3, 4, 10 and 8 in saline condition.

**Keywords:** Agronomic traits, Breeding value, Relative utility, Selection

**Table 1. Analysis of soil saturated paste and water of research stations of Mashhad and Abbas-Abad stations, 2017**

Samples characters	CO <sub>3</sub> <sup>2+</sup>	HCO <sub>3</sub> <sup>-</sup>	CL <sup>-</sup>	SO <sub>4</sub> <sup>2+</sup>	Ca <sup>2+</sup>	Mg <sup>2+</sup>	Na <sup>+</sup>	Ca <sup>2+</sup>		pH	S.A.R	EC	
								Mg <sup>2+</sup>	K <sup>+</sup>				
m.eq / Lit										ds/m			
Soil	Mashad	0.0	2.5	23	-	12	14.4	9.1	16.4	2.88	7.9	6.2	1.54
	Abbas abad	16.12	3.75	170	-	32	37	155	69	1.77	8.03	26.3	21.9
Water	Mashad	0.0	3.7	3.2	2.9	2.8	3.8	3.2	6.6	0.04	7.2	1.7	1.05
	Abbas abad	0.0	5.8	35	12.21	3.6	9.6	39.6	13.2	0.21	7.39	15.4	4.95

**Table 2. Coefficient of traits in selection index in non-stress condition in single cross hybrids of maize (Torogh Mashhad station), 2017**

Traits	Smith- Hazel Index					Baker
	1	2	3	4	5	6
Wet Forage yield	1.426	0.013	0.76	1.01	-0.87	15.65
Dry Forage yield	-0.766	3.20	-0.36	-0.62	3.50	-7.59
Number of leaves	-15.17	-8.17	-7.55	-1.37	-6.58	-218.59
Number of ears	0.34	0.36	0.09	-0.19	0.61	3.35
Plant height	0.69	0.75	0.33	-0.03	0.77	8.35
Days to anthesis	0.92	1.90	0.42	-0.58	2.36	6.26

**Table 3. Coefficient of traits in selection index in salt stress condition in single cross hybrids of maize (Abbas- abad station), 2017**

Traits	Smith- Hazel					Baker
	1	2	3	4	5	6
Wet Forage yield	27.81	10.30	44.96	-0.55	20.42	-56.35
Ear length	14.55	4.90	22.88	-0.35	10.52	-26.37
Number of leaves	14.59	4.81	22.49	0.21	10.11	-24.42
Number of ear	-53.36	-18.22	-84.85	1.25	-38.92	99.34
Plant height	2.19	0.64	3.34	-0.04	1.56	-3.25
Days to anthesis	-71.15	-24.03	-112.85	1.61	-51.87	130.53

**Table 4. Economical values of traits for calculating selection index in normal conditions in maize hybrids (Torough Mashhad station), 2017**

Traits	Economical values of traits for calculating selection index					
	Smith= Hazel					Baker
	1	2	3	4	5	6
Wet Forage yield	1	1	0.6	1	0	8.89
Dry Forage yield	1	6.88	0.45	-0.27	6.88	3.89
Number of leaves	1	2.71	0.12	-0.25	2.71	0.25
Number of ear	1	0.57	0.24	0.44	0.57	4.44
Plant height	1	0.16	0.49	-0.02	0.16	15.82
Days to anthesis	1	-1.04	0.48	0.18	-1.04	1.84

Economical values above are: 1- yield of unit weight (no, 1), 2- coefficient of entered traits in standard stepwise regression, 3-heritability of traits entered in regression model, 4- correlation coefficient between entered traits in regression model with yield, 5- like the state 2, but zero instead 1 for yield, 6-Baker index (root of genotypic variance of traits)

**Table 5. Economical values of traits for calculating selection index in saline conditions in maize hybrids (Abbas- abad station), 2017**

Traits	Economical values of traits for calculating selection index					
	Smith= Hazel					Baker
	1	2	3	4	5	6
Wet Forage yield	1	1	0.66	1	0	2.24
Dry Forage yield	1	0.33	0.13	0.023	-0.33	0.7
Number of leaves	1	-0.62	0.46	0.074	-0.62	0.52
Number of ear	1	0.44	0.68	0.743	0.442	3.25
Plant height	1	0.04	0.19	0.543	0.04	5.18
Days to anthesis	1	-0.172	0.5	-0.2	-0.172	1.86

Economical values above are: 1- yield of unit weight (no, 1), 2- coefficient of entered traits in standard stepwise regression, 3-heritability of traits entered in regression model, 4- correlation coefficient between entered traits in regression model with yield, 5- like the state 2, but zero instead 1 for yield, 6-Baker index (root of genotypic variance of traits)

**Table 6. Combined analysis of variance for forage yield and related traits of maize single cross hybrids in saline and normal conditions, 2017**

S.O.V	df	Wet forage yield	Dry forage yield	Forage quality index	Kernel number/row	Rows number	Ear length	number of Leaves
Environment(E)	1	83543.8**	11271.73**	0.052**	11418.017**	51.689**	1706**	32.86**
rep/envirment	6	113.86	53.15	0.0044	18.64	0.963	4.35	0.733
Genotype(G)	13	180.78**	31.42**	0.0027 <sup>ns</sup>	61.58**	2.283 <sup>ns</sup>	6.15**	0.959**
G × E	13	126.04**	32.136**	0.0038*	37.82**	2.388 <sup>ns</sup>	2.18 <sup>ns</sup>	0.440 <sup>ns</sup>
Error	78	27.42	9.184	0.0016	12.48	1.809	2.443	0.382
Cv (%)		11.80	20.05	13.37	10.40	8.68	8.60	4.45

**Table 6. Continued**

S.O.V	df	Stem diameter	Ear height	Plant height	Tassel length	Days to anthesis	Days to silking
Environment(E)	1	67.504**	101066.86**	56532.71**	638.74**	440.036**	252.00**
rep/envirment	6	13.03	253.57	290.93	11.420	1.750	1.637
Genotype(G)	13	8.76**	63.58*	543.45**	15.133 <sup>ns</sup>	27.783**	25.981**
G × E	13	5.41*	92.50**	565.94**	23.24 <sup>ns</sup>	0.0357 <sup>ns</sup>	0.000 <sup>ns</sup>
Error	78	2.698	33.44	189.08	15.069	3.609	4.650
Cv (%)		6.83	11.03	6.51	9.37	3.24	3.47

\* significant differences \*\* and <sup>ns</sup>, respectively at 0.05 and 0.01 probability levels

**Table 7. Means comparison test for forage yield and related traits of maize single cross hybrids in saline and normal conditions-2017**

Environment	Wet forage yield	Dry forage yield	Forage quality index	Kernel number/row	Rows number	Ear length
Normal	71.695 <sup>a</sup>	25.148 <sup>a</sup>	0.326 <sup>a</sup>	44.086 <sup>a</sup>	16.171 <sup>a</sup>	22.088 <sup>a</sup>
Salt stress	17.071 <sup>b</sup>	5.084 <sup>b</sup>	0.282 <sup>b</sup>	23.892 <sup>b</sup>	14.813 <sup>b</sup>	14.280 <sup>b</sup>

**Table 7. Continued**

Environment	Number of leaves	Stem diameter	Ear height	Plant height	Tassel length	Days to anthesis	Days to silking
Normal	13.364 <sup>b</sup>	23.29 <sup>b</sup>	82.464 <sup>a</sup>	233.646 <sup>a</sup>	43.82 <sup>a</sup>	60.54 <sup>a</sup>	63.625 <sup>a</sup>
Salt stress	14.448 <sup>a</sup>	24.842 <sup>a</sup>	22.485 <sup>b</sup>	188.713 <sup>b</sup>	39.042 <sup>b</sup>	56.57 <sup>b</sup>	60.625 <sup>b</sup>

The means with at least one common alphabet letter had no significant differences

**Table 8. Stepwise regression with yield as dependent variable and other traits as independent variables in maize genotypes under non-stress conditions (Torogh Mashhad station), 2017**

Df	Step 1		Step 2		Step 3		Step 4		Step 5	
	Regression	Error	Regression	Error	Regression	Error	Regression	Error	Regression	Error
	1	54	2	53	3	52	4	51	5	50
Enterd trait	(X1)		(X2)		(X3)		(X4)		(X5)	
	Plant height		number of ear		Dry Forage yield		Days to anthesis		number of leaves	
MS	2307.59	76.9	2036.25	45.1	1559.01	34.3	1222.47	30.8	1006.05	28.4
F	29.99**		45.16**		45.41**		39.65**		35.13**	
(R <sup>2</sup> )	35.71		63.02		72.38		75.67		77.84	
$Y = 70.30 + 6.88(X1) - 2.712(X2) + 0.571(X3) + 0.155(X4) - 1.041(X5)$										

ns. \*, \*\*: non-significant, significant at 0.05 and 0.01 probability level, respectively

**Table 9. Stepwise regression with yield as dependent variable and other traits as independent variables in maize genotypes under salt stress conditions (Abbas- abad station), 2017**

Df	Step 1		Step 2		Step 3		Step 4		Step 5	
	Regression	Error	Regression	Error	Regression	Error	Regression	Error	Regression	Error
	1	54	2	53	3	52	4	51	5	50
Enterd trait	(X1)		(X2)		(X3)		(X4)		(X5)	
	number of ear		Ear length		Days to anthesis		number of leaves		Plant height	
MS	160.03	3.48	88.11	3.24	61.95	3.12	48.58	3.02	40.65	2.90
F	45.93**		27.16--		19.85**		16.10**		14.02**	
R <sup>2</sup>	55.26		64.70		70.12		72.66		76.90	
$Y = 14.16 + 0.330(X1) - 0.620(X2) + 0.442(X3) + 0.04(X4) - 0.172(X5)$										

ns. \*, \*\*: non-significant, significant at 0.05 and 0.01 probability level, respectively

**Table 10. Expected genetic gain for each traits by improved index, correlation between index with additive value, expected gain and relative efficiency of selection index based on 10 percent of selection intensity (k=1.76) in normal condition (Torogh Mashhad station), 2017**

index	Genetic gain of traits						(R <sub>HI</sub> )	(ΔH)	(RE)
	Dry Forage yield	Wet Forage yield	Number of leaves	Number of Ear	Plant height	Days to anthesis	Correlation between index with additive value	Expected gain	Relative efficiency of selection index
1	13.07	4.24	-0.24	2.84	25.99	1.39	0.9	47.30	0.49
2	14.45	4.45	-0.25	2.55	22.84	-1.38	0.74	50.92	1.20
3	13.02	4.35	-0.23	2.68	26.04	1.40	0.9	23.82	0.45
4	12.08	6.95	-0.01	2.31	28.57	4.07	0.71	11.36	0.77
5	16.45	4.08	-0.34	3.01	23.33	-2.93	3.93	35.66	2.05
Baker	0.04	0.01	0.001	0.01	0.1	0.003	0.92	1.76	0.49

**Table 11. Expected genetic gain for each traits by improved index, correlation between index with additive value, expected gain and relative efficiency of selection index based on 10 percent of selection intensity (k=1.76) in saline condition (Abbas- abad station), 2017**

index	Genetic gain of traits						(R <sub>HI</sub> )	(ΔH)	(RE)
	Dry Forage yield	Wet Forage yield	Number of leaves	Number of Ear	Plant height	Days to anthesis	Correlation between index with additive value	Expected gain	Relative efficiency of selection index
1	13.16	15.31	13.44	-15.82	13.28	-15.81	6.15	-33.98	0.94
2	4.76	5.13	4.78	-6.28	4.76	-5.25	2.26	-10.93	1.14
3	0.92	0.92	0.90	-0.41	0.89	-0.91	12.37	-53.99	0.97
4	82.12	-86.55	-82.08	112.03	-82.01	88.45	0.19	0.57	1.26
5	73.12	74.69	72.58	-101.18	72.75	-75.95	3.93	-24.80	0.95
Baker index	-43.82	-45.95	-43.74	58.63	-43.73	46.89	1.86	-58.30	1.19

**Table 12. Yield, selection indices and rank of each genotype (numbers in parenthesis) in normal condition in maize (Torogh Mashhad station), 2017**

Genotype	Yield (ton/ha)	Index					
		Smith Hazel					Baker
		1	2	3	4	5	6
1	83.51(2)	17.53	301.43	-2.90	-95.28	383.50	-566.56
2	79.15(4)	38.71(1)	316.68(2)	7.35(1)	-94.54	397.96(3)	-297.59(1)
3	80.49(3)	27.73	323.48(1)	2.85	-94.44	402.71(1)	-395.53(4)
4	66.15	29.78(5)	284.95	3.89(4)	-88.90(3)	361.88	-413.93
5	85.62(1)	32.31(3)	316.21(3)	4.04(3)	-95.27	398.24(2)	-375.52(3)
6	72.25	33.99(2)	307.19(5)	5.78(2)	-91.91	385.61(5)	339.10(2)
7	74.80(5)	14.91	290.74	-3.73	-91.97	369.93	-580.19
8	74.74	19.30	310.14(4)	-2.37	-97.43	394.05(4)	-553.33
9	86.47	11.34	271.03	-5.99	-91.66	351.73	-655.21
10	64.34	15.28	266.07	-4.03	-91.79(5)	346.45	-624.02
11	70.91	28.02	302.22	2.05	-94.63	384.18	-436.88
12	64.87	30.19(4)	283.94	3.52(5)	-90.60(4)	363.07	-412.84(5)
13	59.49	12.18	264.56	-5	-89.62(2)	343.13	-642.98
14	58.94	-1.41	258.60	-11.17	88.35(1)	335.46	-794.75

\*The selected genotypes for each index are 20 percent of the best ones (5 genotypes). The no inside parenthesis are the rank of each genotypes

**Table 13. Yield, selection indices and rank of each genotype (numbers in parenthesis) in saline condition in maize (Abbas- abad station), 2017**

Genotype	Yield (ton/ha)	Index					
		Smith Hazel				Baker	
		1	2	3	4	5	6
1	18.23(4)	-4275.38(4)	-1469.38(4)	-6823.71(4)	105.91	-3134.36(4)	8058.14
2	14.98	-4111.85(1)	-1412.78(1)	-6562.59(1)	102.02	-3014.72(1)	7747.84
3	21.67(1)	-4532	-1557.54	-7231.52	111.56(2)	-3320.98	8537.87(2)
4	19.41(2)	-4474.91	-1537.75	-7140.41	110.12(3)	-3279.45	8429.43(3)
5	19.32(3)	-4339.06(5)	1492.45(5)	-6926.54(5)	107.88	-3181.3(5)	8186.80
6	16.44	4403.21	-1512.38	-7025.41	108.92	-3227.04	8290.28
7	17.34	-4241.22(3)	-1458.38(3)	-6769.51(3)	104.87	-3109.08(3)	7997.94
8	16.88	-4446.27	-1526.78	-7093.68	109.59(5)	-3258.33	8368.03(5)
9	13.75	-4380.59	-1505.38	-6991.03	108.62	-3211.27	8254.81
10	15.08	-4451.38	-1527.86	-7101.06	109.66(4)	-3261.99	8372.84(4)
11	15.49	-4445.54	-1526.43	-7092.22	109.56	-3257.73	8365.71
12	16.19	-4439.73	-1525.49	-7084.33	109.21	-3253.84	8362.30
13	17.88(5)	-4771.29	-1637.55	-7640.44	117.39(1)	-3495.66	8972.14(1)
14	16.34	-4112.98(2)	-1414.47(2)	-6566.06(2)	102.55	-3016.17(2)	7760.21

\*The selected genotypes for each index are 20 percent of the best ones (12 genotypes). The no inside parenthesis are the rank of each genotypes