

Grouping of red bean genotypes based on the relationship between some quantitative and qualitative traits-using multivariate statistical methods

Archive of SID

KS31169

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(Amini *et al.*, 2000)

(*Phaseolus vulgaris* L.)

(Common bean)

(Koocheki and Banayan., 1994)

(Raffi and Nath, 2004)

(Lackey, 1983)

(Aggarwal and Singh, 1973)

(Anon., 2003)

(Bennet *et al.*, 1977)

(Schoonhoven and Voysest, 1993)

(Sarafi, 1978)

(Majnoon Hosseini, 1993)

(Yazdi Samadi and Abdmishani, 1996)

(Martin *et al.*, 1995)

(Amini *et al.*, 2002)

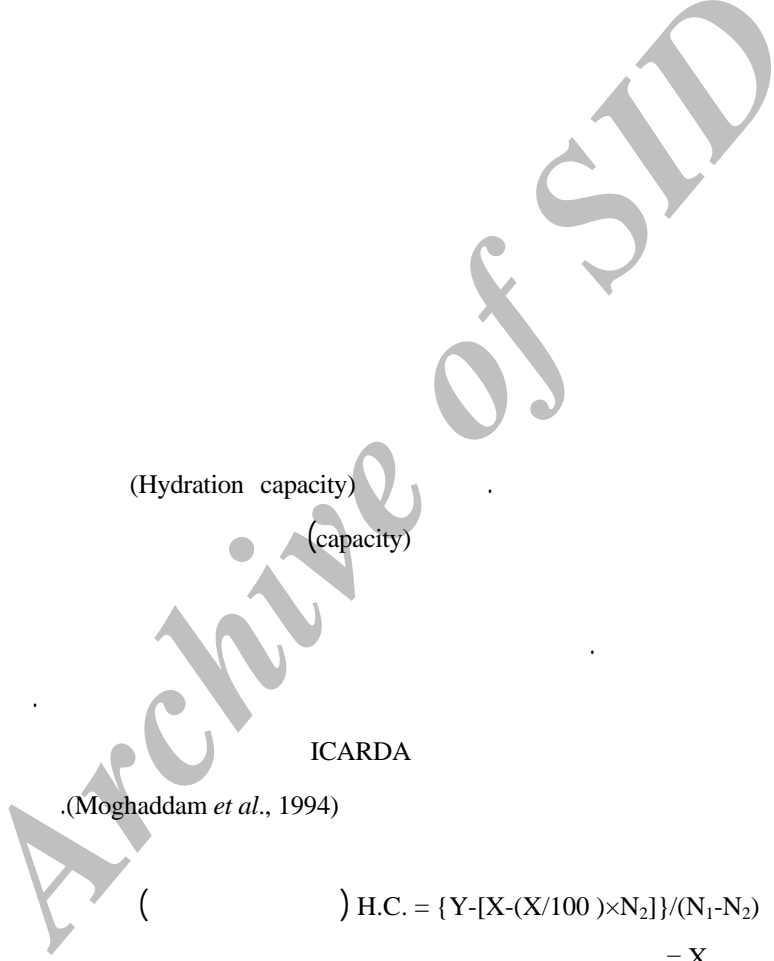
(Mirzaie Nadooshan, 1997)

(Santalla *et al.*, 1993)

CIAT ICARDA IPGRI
(Moghaddam *et al.*, 1994)

$$r = \frac{...}{...}^{**}$$

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Swelling

(Hydration capacity)
(capacity)

ICARDA

(Moghaddam *et al.*, 1994)

$$\begin{aligned} () \text{ H.C.} &= \{ Y - [X - (X/100) \times N_2] \} / (N_1 - N_2) \\ &= X & &= Y \\ & & &= N_2 & &= N_1 \end{aligned}$$

$$\begin{aligned} () \text{ S.C.} &= (Y_1 - Y_2) - \{ (X_1 - X_2) - [(X_1 - X_2) / N_1] \times N_2 \} / (N_1 - N_2) \\ &= Y_2 & &+ & &= Y_1 \\ = N_1 & & &= X_2 & &+ & &= X_1 \\ & & & & & & &= N_2 \end{aligned}$$

1- International Plant Genetic Resources Institute.
2- International Center for Agricultural Research in the Dry Areas.
3- Centro International de Agricultura Tropical.

(Arghamee and Bozorgnia, 1991) Ward (ANOVA)
Moghaddam *et al.* $g = \left[\frac{1}{2}n \right]^{\frac{1}{2}}$
(al., 1994) (Varimax)
PATH 2 SAS Version 8 , SPSS (Principle Coordinate Analysis)

Table1. Code of red bean genotypes.

Row	Code Line	Row	Code Line	Row	Code Line
1	KS31101	6	KS31106	11	KS31111
2	KS31102	7	KS31107	12	KS31138
3	KS31103	8	KS31108	13	KS31139
4	KS31104	9	KS31109	14	KS31169
5	KS31105	10	KS31110	15	KS31170

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(Ibrahimi *et al.*, 2001)

(Raffi and Nath, 2004)

1- Ward's Minimum – Variance Method.

Table 2. Analysis of variance for different traits in 15 red bean genotypes.

S.O.V.	df	Mean Square								
		Days to emergence	Days to maturity	Days to first pod maturity	Filling duration	Days to 50% podding	Days to 50% flowering	Nod no. per main Shoot	Sub shoot No.	Plant height
Block	2	2.16 ^{ns}	107.5 ^{**}	5.70 ^{ns}	208.7 ^{**}	7.11 ^{ns}	36.87 ^{**}	2.16 ^{ns}	1.68 ^{ns}	111.4 ^{**}
Genotype	14	6.74 ^{**}	25.06 ^{**}	45.12 ^{ns}	35.91 ^{**}	19.28 ^{ns}	26.47 ^{**}	6.74 ^{**}	0.756 ^{ns}	129.9 ^{**}
Error	28	1.00	5.442	27.33	6.44	21.41	2.209	1.00	0.454	30.19

(Table 2. continued)

Protein %	Mean Square										
	100 Seed weight	Seed yield	Pod no. per Plant	Pod weight	Seed no. per Plant	Internode diameter	Internode length	Seed diameter	Seed weight	Seed length	Seed no. per Pod
1096.6 ^{**}	12.74 ^{ns}	9.14 ^{ns}	2.14 ^{ns}	13.45 ^{ns}	50.04 ^{**}	2.01 ^{ns}	0.066 ^{ns}	0.025 ^{ns}	0.405 ^{**}	0.017 ^{ns}	0.352 ^{ns}
189.57 ^{**}	47.18 [°]	4.88 ^{ns}	4.82 ^{ns}	8.55 ^{ns}	129.7 ^{**}	0.707 ^{ns}	0.239 ^{ns}	0.387 ^{ns}	0.597 ^{**}	1.99 ^{**}	0.437 ^{ns}
41.05	20.13	6.09	5.411	8.18	0.351	0.729	0.119	0.277	0.127	0.307	0.347

(Table 2. continued)

Plant type	Mean Square										
	Time before swelling	Time after swelling	Swelling capacity	Hydration capacity	Pod length	Pod tail length	Seed Scent	Seed appearance	Seed taste	Seed texture and structure	
0.023 ^{ns}	21.18 ^{ns}	167.6 ^{ns}	0.74 [°]	0.003 ^{ns}	1.25 [°]	0.028 ^{ns}	0.114 ^{ns}	1.11 ^{**}	6.45 ^{ns}	0.425 ^{ns}	
2.77 ^{ns}	281.4 ^{ns}	627.7 ^{ns}	0.26 [°]	0.062 ^{**}	1.03 ^{**}	0.07 [°]	1.18 ^{**}	0.771 ^{**}	18.06 ^{ns}	1.28 ^{**}	
0.002	312.8	493.1	0.140	0.011	0.342	0.03	0.212	0.257	23.70	0.466	

* و ** :
:ns

* and **: Significant at the 5% and 1% levels of probability, respectively.
ns:Non-significant.

Table 3 . Stepwise regression for the traits used into the final model for 15 red bean genotypes.

	Standardized coefficients		SS	R ²	R ² Partial	F
	B	Std Error				
Intercept	-16.236	1.592	2.116			103.87 ^{**}
Weight Pod	0.271	0.040	0.928	0.637	0.637	45.49 ^{**}
100 Seed Weight	0.241	0.019	3.006	0.713	0.076	147.45 ^{**}
Pod/ Plant No.	0.369	0.067	0.622	0.867	0.154	30.51 ^{**}
Width Seed	1.238	0.149	1.403	0.935	0.068	68.84 ^{**}
Length Internode	-0.554	0.179	0.195	0.964	0.029	9.59 [*]
Seed/Pod No.	0.171	0.019	1.644	0.987	0.029	80.64 ^{**}
Node/Main shoot	-0.158	0.057	0.157	0.994	0.007	7.74 [*]

* and **: Significant at the 5% and 1% levels of probability, respectively. . . / / :*** و *

Table 4. Phenotypical correlation among seed yield with used traits in regression model.

Row	Traits	8	7	6	5	4	3	2
1	Seed Yield	0.252 ^{NS}	0.789 ^{**}	0.351 ^{NS}	-0.540 [*]	-0.067 ^{NS}	0.193 ^{NS}	0.0359 ^{NS}
2	Node No/ Shoot	-0.412 ^{NS}	-0.107 ^{NS}	0.761 ^{**}	0.311 ^{NS}	-0.292 ^{NS}	0.392 ^{NS}	
3	Seed No./Pod	-0.278 ^{NS}	-0.026 ^{NS}	0.256 ^{NS}	-0.081 ^{NS}	-0.215 ^{NS}		
4	Seed Width	0.510 [*]	-0.287 ^{NS}	-0.557 ^{**}	-0.012 ^{NS}			
5	Internode Length	0.006 ^{NS}	0.469 ^{NS}	-0.018 ^{NS}				
6	Pod No./ Plant	0.346 ^{NS}	0.133 ^{NS}					
7	Pod Weight	-0.026 ^{NS}						

* and **: Significant at the 5% and 1% levels of probability, respectively. / / :*** و *
 - (Number 8 is 100 Seed Weight). ()

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(/) (Santalla *et al.*, 1993)

(Bennet *et al.*, 1977)

Table 5. Path analysis of quantitative traits in 15 red bean genotypes.

Quantitative traits	Pod Weight	100 Seed weight	Pod no. per Plant	Seed width	Internode length	Seed no. per Pod	Nod no. per Shoot	Total Correlation
Pod Weight	1.065	0.012	0.108	0.306	0.611	0	0.013	0.797
100 Seed Weight	0.042	0.497	0.281	0.560	0.008	0.007	0.050	0.254
Pod no. per Plant	0.213	0.172	0.813	0.613	0.023	0.008	0.091	0.351
Seed width	0.447	0.254	0.454	1.099	0.015	0.006	0.036	0.068
Internode Length	0.756	0.003	0.015	0.014	1.302	0.002	0.037	0.541
Seed no.per Pod	0.042	0.138	0.208	0.237	0.105	0.029	0.047	0.193
Node no. per Shoot	0.172	0.204	0.619	0.322	0.405	0.011	0.120	0.035
Residual effect= -1.069								/ :

Table 6. Eigen values and factors variances in quantitative traits.

Factors	Eigen value	Variance ratio	Cumulative variance
Factor 1	5.586	0.207	0.207
Factor 2	4.629	0.171	0.378
Factor 3	4.351	0.161	0.539
Factor 4	3.106	0.115	0.654
Factor 5	2.098	0.078	0.732
Factor 6	1.610	0.059	0.791
Factor 7	1.435	0.053	0.845
Factor 8	1.329	0.049	0.894
Others	0.210	0.106	1.000

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Table 7. Factor analysis by varimax rotation for quantitative traits in 15 red bean genotypes.

Quantitative Traits	Factor1	Factor2	Factor3	Factor4	Factor5	Factor6	Factor7	Factor8
Plant height	0.106	-0.033	0.804	0.146	-0.207	-0.059	0.175	-0.142
Sub shoot No.	0.215	0.003	-0.119	-0.563	0.156	0.005	0.509	0.541
Nod no. per main shoot	0.425	-0.633	0.304	0.026	0.261	-0.413	0.148	0.042
Seed no. per Pod	0.146	-0.161	-0.638	0.564	0.118	0.176	-0.149	-0.087
Days to emergence	0.057	-0.318	-0.244	0.421	0.371	0.311	-0.321	0.190
Days to 50% flowering	0.910	-0.217	-0.035	0.171	0.058	0.013	-0.217	-0.051
Days to 50% poding	0.640	-0.518	-0.289	0.139	-0.210	0.075	0.090	-0.201
Days to first pod maturity	0.242	-0.328	-0.391	-0.028	-0.117	0.724	0.173	-0.171
Days to maturity	0.251	-0.022	-0.118	0.028	0.063	0.924	0.090	0.056
Seed filling duration	-0.571	0.168	-0.067	-0.134	0.003	0.761	0.110	0.091
Pod length	-0.120	0.865	0.129	0.023	0.344	-0.031	-0.139	0.025
Pod tail length	0.012	0.344	0.069	-0.428	0.303	0.614	0.285	0.028
Seed length	0.296	0.847	-0.007	0.076	0.006	-0.031	0.174	-0.093
Seed width	-0.522	0.020	-0.176	0.395	-0.294	-0.321	-0.215	-0.529
Seed diameter	0.421	-0.383	-0.211	-0.015	0.156	0.303	0.093	-0.514
Internode diameter	-0.046	0.381	0.763	-0.153	0.041	-0.195	0.121	0.076
Internode length	-0.011	0.263	0.839	-0.151	-0.357	0.076	-0.130	-0.056
Seed no. per Plant	0.777	-0.155	-0.081	-0.145	0.535	0.056	0.051	0.144
Pod no. per Plant	0.290	-0.495	0.148	-0.167	0.562	-0.196	0.129	0.366
Pods weight	0.076	0.308	-0.256	-0.154	0.770	0.061	-0.019	-0.031
Seed yield	-0.104	0.066	-0.236	0.039	0.936	0.055	-0.106	-0.159
100 Seed weight	-0.859	0.161	0.142	0.121	0.131	0.128	-0.129	-0.220
Hydration capacity index	0.097	0.158	-0.083	0.895	-0.061	-0.173	-0.168	0.084
Swelling capacity index	0.069	0.003	-0.017	0.972	-0.064	-0.025	0.075	-0.023
Time before swelling	0.086	0.029	0.059	-0.126	-0.182	0.058	0.914	0.068
Time after swelling	-0.408	-0.186	0.422	-0.014	0.333	0.134	0.686	-0.104
Protein %	-0.029	-0.097	-0.128	0.097	-0.102	0.057	-0.007	0.907
Total factors	4.139	3.391	3.219	3.152	3.006	3.005	2.150	2.081

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Table 8. Principle coordinate analysis on quantitative and qualitative traits.

Traits	Dim1	Dim2	Traits	Dim1	Dim2
Plant height	-0.037	0.067	Seed no. per Plant	0.065	-0.100
Nod no. per main shoot	0.027	-0.034	Seed no. per Pod	0.084	-0.017
Swelling capacity index	0.099	0.351	Pod no. per Plant	0.038	-0.024
Hydration capacity index	0.197	0.039	Pods weight	0.051	-0.026
Days to emergence	0.059	-0.002	Seed yield	0.063	0.016
Days to 50% flowering %	0.065	-0.040	100 Seed weight	0.022	0.124
Days to 50% ponding %	0.041	-0.027	Sub-shoot no.	0.0006	0.047
Days to first pod maturity	0.027	0.002	Plant type	0.054	-0.126
Days to maturity	0.041	-0.004	Protein percentage	0.049	-0.028
Seed filling duration	0.018	0.028	Seed appearance	0.033	0.059
Pod length	0.043	0.017	Seed Scent	0.084	-0.076
Pod tail length	0.033	-0.005	Seed taste	0.056	0.001
Seed texture and structure	0.017	0.094	Seed length	0.021	0.022
Internode diameter	0.00005	0.013	Seed width	0.044	0.038
Internode length	0.0003	0.043	Seed diameter	0.030	0.022
Time before swelling	-0.214	-0.107	Time after swelling	-0.099	0.078

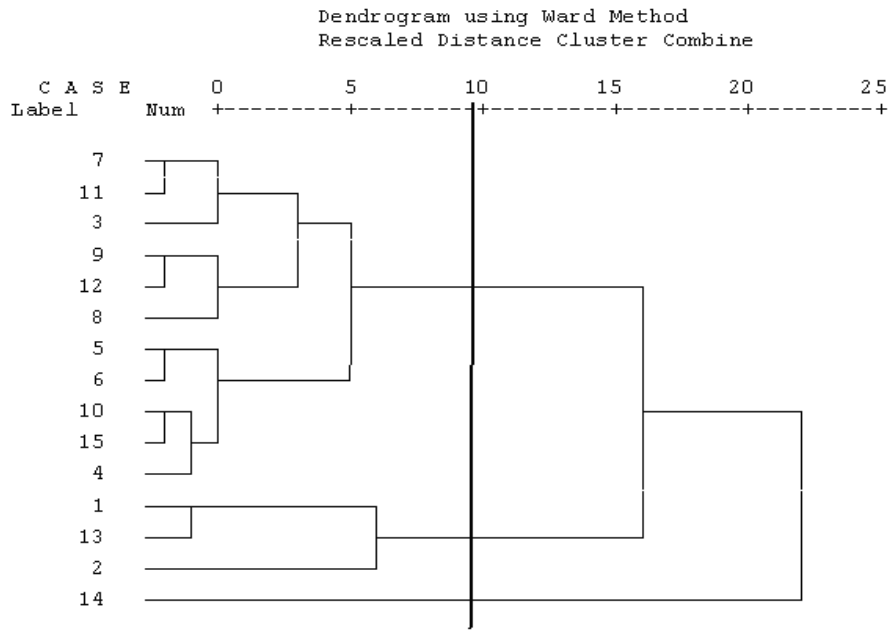
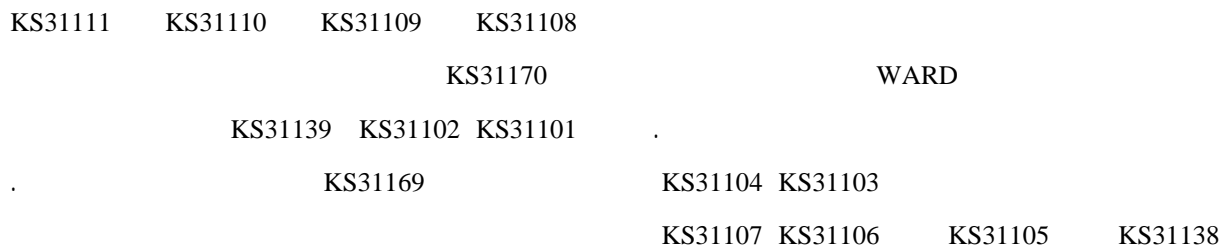


Figure 1. Dendrogram for 15 genotypes by Ward method.



(Schoonhoven and Voysest, 1993)

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Grouping of red bean genotypes based on the relationship between some quantitative and qualitative traits-using multivariate statistical methods

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ABSTRACT

Mohammadi, A., M. R. Bihamta, M. Soluoki and H. R. Dorri. Grouping of red bean genotypes based on the relationship between some quantitative and qualitative traits-using multivariate statistical methods. **Iranian Journal of Crop Sciences**. 10(2): 178-190.

To study the relationship between some quantitative and qualitative traits in red bean, 15 red bean genotypes were studied in experimental field of Faculty of agriculture, the University of Tehran in 2004 cropping season using a randomized complete block design with three replications. Necessary scores and measurements were made and multivariate statistical analyses were performed for different quantitative and qualitative traits. Analysis of variance of data revealed high genetic variation for concerned traits among red bean genotypes. Seven quantitative traits were used in stepwise regression model which included quantitative attributes of seed and plant morphological traits. Among these attributes pod weight in plant and length of internode with highest correlation coefficients had direct effects of 1.605 and -1.302, respectively. In factor analysis, 89% of total variation was explained by eight factors which were divided in two sets: The primary factors included; seed yield related; quantitative morphological and physiological traits and the secondary factors comprised; cooking quality related traits and plant type. Cluster analysis grouped the 15 red bean genotypes in three distinctive groups. KS31169 genotype had the least similarities with the other genotypes; therefore, it would be expected that crosses made between this genotype and genotypes of the first group will develop desirable variation in segregating populations for breeders.

Keywords: Red bean, Quantitative traits, Qualitative traits, Seed yield, Multivariate method, Cluster analysis.

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