Inheritance and Transfer of Root Rot (Pythium) Resistance to Bean Varieties

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Background

Bean root rot caused by several Pythium species is a relatively recent problem in East and Central Africa that is increasing in importance. Yield losses of up to 70% in popular commercial bean cultivars have been reported in Rwanda. In order to curb the effects of this disease, identification of resistance and transfer of resistance to bean genotypes preferred by resource poor small holder farmers is an important consideration.



Root rot symptoms in the field

Objectives

To identify sources of resistance within major market class, determine the nature of inheritance and through breeding transfer the resistance into well-adapted market class bean cultivars.

Materials and Methods

Populations have been developed between resistant and susceptible well adapted cultivars (Fig.1). F₂ populations of SCAM 80-CM/15, RWR 719 and MLB 4989A were used in this study.Evaluation of bean plants was done in a root rot hot spot nursery in Vihiga Western Kenya in August 2001. F₂ progenies alongside their parents were evaluated for root rot severity in a CRBD based on a 1-9 CIAT scale (Abawi and Pastor-Corrales., 1990). Fifty plants from each plot were sampled from parental plots while 150 plants were sampled from F₂ generations. Chi- square values (X²) were calculated based on actual number of plants counted in each of the resistant and susceptible categories.



Results

Table 1. Number of plants that showed resistant and susceptible reaction to bean root rot in the field

Susceptible	Resistant 🔶	RWR 719		SCAM 80-CM/15		MLB 49-89 A	
	Generation	Resistant	Susceptible	Resistant	Susceptible	Resistant	Susceptible
	Pl	50	_	50	_	50	2
GLP2	P2	8	42	8	42	6	44
GLP 585	P2	2	43	8	42	8	42
CAL96	P2	4	46	2	48	3	47
URUGEZI	P2	2	48	0	50	4	46
GLP2	F2	108	38	120	25	103	29
GLP 585	F2	124	28	104	22	98	46
CAL96	F2	105	33	95	48	107	32
URUGEZI	F2	109	42	90	31	115	30

P1 = resistant parent and 2 = susceptible paren

Susceptible	Resistant	RWR 719		SCAM80-CM/15		MLB 49-89 A	
		X²	Р	X²	Р	X ²	Р
GLP2	3:1	0.22**	0.70-0.50	4.94 ^{NS}	0.05 - 0.10	3.05**	0.10-0.05
GLP 585	3:1	3.74**	0.10 - 0.05	7.55 ^{NS}	0.01 - 0.001	3.71**	0.10 - 0.05
CAL96	3:1	1.24**	0.20 - 0.10	5.55 ^{NS}	0.05 - 0.01	1.28**	0.20 - 0.10
URUGEZI	3:1	0.57**	0.05 - 0.03	6.00 ^{NS}	0.05 - 0.10	1.74**	0.20 - 0.10

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Conclusion

Bean root rot score of F_2 plants from crosses involving resistant parents RWR 719 (KK22) and MLB 49-89 (KK15) segregated in the ratio approximately 3:1. The segregation of F_2 progenies derived from resistant parent SCAM- 49-89 A (KK8) did not fit any expected ratio in all the crosses. This indicated that the resistant gene in RWR 719 and MLB 49-89A is most likely a single dominant gene. The mode of resistance in SCAM-cm/15 could not be predicted. Further evaluation in the screen house and laboratory (using protein and DNA polymorphism) involving the parents, F_1 , F_2 and backcrosses is underway and will provide useful information for mechanisms of resistance in the studied genotypes. More field screening of the F_{2-3} families of resistant progenies derived from RWR 719 and MLB 49-89A to check goodness of fit for the ratio 1:2:1 (resistant : segregating: susceptible) would be confirmatory for the mechanism of resistance in the two cultivars that could be utilized in a breeding program

Reference:

Abawi, G.S. and Pastor-Corrales, M.A.; (1990). Root rots of bean in Latin America and Africa: diagnosis, research, methodologies and management strategies. CIAT, Cali, Colombia, 114 pp. Abawi, G.S. and Pastor-Corrales, M.A.; (1990). Abawi, G.S. and Pastor-Corrales, M.A.; (1990).

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 XX = values significant at 1 degrees of freedom, 95% confidence level and X² < 3.84

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