INDUCTION OF MUTATION IN *NEUROSPORA CRASSA* WITH DIATHANE-M45 AND GENETICAL STUDIES OF SOME SELECTED MUTANTS

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Key words: Diathane-M45, Mutants, Neurospora crassa

Abstract

Diathane-M45 at 0.125 and 0.062% concentration induced nine different groups of mutants in *Neurospora crassa*. The mutants were cauliflower, fissure, mat, ropy, check, fluffy, conidial band, vigorous and buff which showed difference with Ema in their morphology. Complementation test of different morphological mutants showed positive results but complementation of similar morphological mutants were negative. Linkage study revealed clf (D₆-34) was linked with pyr-1A of linkage group IV.

Introduction

The most technically advantageous method of studying the organization and mode of action of gene or the genic inheritance in any organism is achieved by inducing artificial mutation in the organism at different loci.

Nowadays a good number of fungicides are used in various purposes such as mutagens, biological control etc. Diathane-M45 is a fungicide which contains 8000 g Mankojeb per kg (zinc, manganese ethylenebisdithiocarbamate). Bari and Mian⁽¹⁾ used diathane-M45 on peanut as biological control. Sultana⁽²⁾ and Rahman⁽³⁾ observed the mutagenic effect of pesticide (Fenom) on *N. crassa* and nine groups of mutants were obtained. Sandhu *et al.*⁽⁴⁾ and Bignami *et al.*⁽⁵⁾ studied the mutagenic activities of different fungicides, Haque and Shamsi^(6,7) worked with diathane-M45 against *Colletotrichum corchori* and *Macrophomina phaseolina*. In case of strong inhibition of the pathogen by the application of fungicides, deformation of the colony and mycelial aggregation were observed.⁽⁸⁾

Therefore, it was decided: (i) to induce mutation in *Neurospora crassa* by using diathane-M45, (ii) to study in detail a few selected and interesting mutants for their morphological changes by studying the linear growth, mycelial weight, time required for germination, fertility, mating type, heterocaryosis in comparison with the wild type Ema and (iii) to prepare extracts from the foreign marker in order to study in detail the segregation and linkage of selected mutants.

Materials and Methods

The experimental materials were *Neurospora crassa* Emerson 'a'/Ema (5297) and Emerson 'A'/EmA (5296). Ema was used for treating conidia with mutagen. Both Ema and EmA were used for crossing and checking the mating types of the mutants. Seven markers of linkage groups I-VII with 'A' mating types were used for the study of linkage of the mutants. These were leu-3(R156)A of linkage group I, trp-3(C83)A-Ext-2 of linkage group-II, ade-2(STL2)A of linkage group-III, pyr-1(H263)A of linkage group IV, ilv-1(16117)A of linkage group V, trp-2(75001)A of linkage group VI and arg-10(13317)A of linkage group VII. Diathane-M45 is a fungicide (Rohm & Hass company, France, Distributor Rhone-Polenc Bangladesh Ltd.). Active ingredient: 8000 g Mankozeb (zinc, manganese, ethylenebisdithiocarbamate) per kg was used as mutagen.

Vogel's minimal medium (VM)⁽⁹⁾ was used for the maintenance of cultures, linear growth and for the study of mycelial growth. Sorbose minimal media (SM) was used for single colony and spore isolation. Westergaard and Mitchell's crossing medium⁽¹⁰⁾ was used for crossing.

For preparation of different solutions and 5% survival of conidial colonies procedure of Mozmader *et al.*⁽¹¹⁾ was followed. Diathane-M45 solutions of 0.5, 0.25, 0.125, 0.0262, 0.03 per cent concentration were prepared which was denoted as D_3 , D_4 , D_5 , D_6 , D_7 , respectively.

Complementation test: The selected morphological mutants were subcultured three times to obtain fresh cultures. Small tubes were used for preparation of VM. Observations were taken after 3, 5, 7, 15 and 21 days for the growth of heterocaryons. For further confirmation, they were tested in liquid media. In every case there were duplication test with duplicate controls.

Results and Discussion

Diathane-M45 of 0.125% (D₅) and 0.062% (D₆) concentration produced 5.08 and 7.90% viable conidia respectively (Table 1). It showed significant mutagenic activity. Nine different groups of morphological mutants were obtained cauliflower (clf), fissue (fis), mat, ropy (ro), check, fluffy (fl), conidial band (con-band), vigorous (vg) and buff. The highest frequency (43.90%) was found for check mutant and the lowest frequency (2.43%) was found in cauliflower, vigorous and buff. The types of mutants obtained with diathane-M45 solution are slightly different from those found earlier with phenol⁽¹¹⁾ and formaldehyde vapour.⁽¹²⁾ Checked growth, vigorous growth, band and cauliflower types were obtained when formaldehyde vapour was used and in case of phenol, vigorous, mat checked, fluffy, buff and band types were obtained. These discrepancies could be due to the difference in mutagenic specificity of formaldehyde vapour, formaldehyde solution and phenol.

In case of diathane-M45 induced mutants the mycelial dry weight of mat (D₆-31), buff (D₆-33) and clf (D₆-34) was found less than Ema and lowest weight (0.0935g) was obtained in case of mat D₆-14 (Table 2). Of the diathane-M45 induced mutants after 72 hours the lowest linear growth found in mat D₆-14 (3.25 cm) and highest for mutant cauliflower D₆-43 (6.75 cm) whereas Ema showed growth of 19.45 cm (Table 3).

Concentration of diathane-M45	No. of germinated conidia	Mean	Percentage of viable conidia	
0% (Normal)	91	88.5	100	
	86			
$0.5(D_3)$	0	-	0	
	0			
$0.25(D_4)$	2	1.5	1.695	
	1			
$0.125(D_5)$	4	4.5	5.08	
	5			
$0.02626(D_6)$	7	7	7.90	
	7			
0.03(D7)	12	14.5	16.38	
	17			

Table 1. The percentage of viable conidia at different concentration of Diathane-M45.

Name of the culture	Weight of the empty filter paper (g)	Weight of the filter paper with dried mycelia (g)	Weight of dried mycelia (g)	Mean
Mat (D ₆ -14)	$0.921 \\ 0.921$	1.019 0.1811	$0.098 \\ 0.089$	0.0935
Fissure (D ₆ -31)	$0.895 \\ 0.922$	$1.009 \\ 1.02$	$0.114 \\ 0.098$	0.106
Buff (D ₆ -33)	$0.934 \\ 0.999$	$1.093 \\ 1.197$	$0.159 \\ 0.198$	0.1785
Cauliflower (D ₆ -34)	$0.892 \\ 0.890$	$1.002 \\ 1.011$	$0.110 \\ 0.121$	0.1155
Ema	$0.921 \\ 0.922$	1.287 1.332	$0.366 \\ 0.410$	0.388

Mating types and fertility of mutants were checked by crossing with Mat, check, ropy, fissure, buff, cauliflower and vigorous both Ema and EmA separately. Perithecia in large number were formed only with EmA in all cases. Spores were shed in all the crosses but the mutants vary in their fertility (Table 4). Single spore of fertile crosses was isolated and classified into mutant and wild types. The mutant and wild progenies were segregated in the ratio of 1 : 1. This finding suggests that the obtained mutants might be the results of changes in the nuclear genes (Table 5).

Name of the	Growth in cm after the following hours											
culture	24	Mean	36	Mean	48	Mean	60	Mean	72	Mean	84	Mean
Mat (D ₆ -14)	0.1	0.15	0.3	0.4	0.95	0.9	1.90	1.95	3.1	3.25	4.2	4.3
	0.2		0.5		1.00		2.00		3.4		4.4	
Fissure (D ₆ -31)	0.5	0.4	1.2	1.05	3.4	3.25	5.1	5.15	6.00	6.00	6.9	6.95
	0.3		0.9		3.1		5.2		6.00		7.0	
Buff (D_6 -33)	0.3	0.35	0.8	0.95	2.90	2.95	4.9	4.95	5.7	5.75	6.4	6.6
	0.4		1.1		3.00		5.0		5.8		6.8	
Cauliflower	1.2	1.3	1.8	1.9	3.9	3.95	5.9	5.85	6.8	6.75	7.8	7.85
(D_6-34)	1.4		2.0		4.0		5.8		6.7		7.9	
Ema	2.1	2.2	4.9	4.95	11.0	10.0	15.8	15.75	19.5	19.45	24.1	24.1
	2.3		5.0		10.9		15.7		19.4		24.1	

Table 3. Linear growth of Ema and four selected mutant.

Table 4. Mating type and fertili	ty of some mutant.
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Cross	Days to initiation of perithecia	Mating type	Shedding days	Fertility
$D_{6}-14 \times Ema$	-		-	Excellent
$D_{6}-14 \times EmA$	7	a	15	
(mat)				
$D_{6}-21 \times Ema$	-		-	Excellent
$D_{6}-21 \times EmA$	15	a	21	
(check)				
$D_{6}-28 \times Ema$	-		-	Poor
$D_{6}-28 \times EmA$	15	а	27	
(ropy)				
$D_{6}-31 \times Ema$	-		-	Poor
$D_{6}-31 \times EmA$	10	а	27	
(fissure)				
$D_{6}-32 \times Ema$	-		-	Poor
$D_{6}-32 \times EmA$	10	a	27	
(fissure)				
D_{6} -33 × Ema	-		-	Excellent
$D_{6}-33 \times EmA$	10	a	21	
(buff)				
$D_{6}-34 \times Ema$	-		-	Excellent
$D_{6}-34 \times EmA$	7	a	15	
(cauliflower)				
$D_{6}-43 \times Ema$	-		-	Poor
$D_{6}-43 \times EmA$	7	а	27	
(vigorous)				
$D_{6}-50 \times Ema$	-		-	Poor
$D_{6}-50 \times EmA$	7	a	27	
(ropy)				

Complementation test revealed that complementation did not take place in case of mutants of a particular group. Each group therefore, probably was produced as a result of mutation at a single locus. But complementation took place among the different morphological mutants indicating that the mutants of different groups were produced as a result of mutation at separate loci.

Cross	No. of spores isolated	No. of wild types isolated	No. of mutant isolated	Segregation ratio
	Isolatoa	oj pos isolatoa	15014004	wild : mutant
$D_6\text{-}21\times EmA$	198	96	102	1:1.052
(check)				
$D_{6}-28 \times EmA$ (ropy)	208	102	106	1:1.039
$D_{6}-43 \times EmA$ (vigorous)	230	110	120	1:1.090

Table 5. Segregation ratio of four mutants.

Table 6. Linkage of cauliflower (D₆-34a) mutant.

Marker used	Linkage group	Crossing name	Total isolates	Progenies	% of the progenies	Inference on linkage
руг-1 (Н 263)А	IV	D_{6} -34 × EmA	146	Wild = 4 pyr-1 = 62 clf = 62 clf+pyr = 18	Wild = 2.75 pyr-1 = 42.46 clf = 42.46 clf+pyr = 12.32	Linked with pyr-1 of linkage group IV

The selected mutants induced with diathane-M45 were crossed with markers of seven linkage groups. Then linkage of the fertile crossed were studied and the distances of the mutants were calculated and obtained that mutants clf (D_{6} -34) linked with pyr-1, linkage group-IV (Table 6).

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(Manuscript received on 11 October, 2004; revised on 24 April, 2005)