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Cytotoxicity of the Fungicides Azoxystrobin and Difenconazole in Root Tips of *Allium cepa* L.

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Abstract: Non-target effects of two broad spectrum, foliar spray, systemic fungicides, Amistar (azoxystrobin 25% EC) and Score (difenconazole 25% EC) in terms of cytotoxicity were investigated at concentrations ranging from 0.44 - 2200 μg (a.i.) mL^{-1} . The test material used was the root meristems of *Allium cepa*. At the recommended dose for field application (2.2 μg (a.i.) mL^{-1}), Difenconazole depressed mitotic index by 4.305% while Azoxystrobin showed a decrease of 1.282% over untreated control at 12 h exposure period. The extent of chromosomal abnormalities has direct relationship with the concentration of the active ingredients and treatment time. The fungicide treated root meristems tended to recover from the cytotoxic effects when they were transferred to distilled water. The rate of recovery as indicated by increasing mitotic index and decreasing incidence of cytological abnormalities was highly pronounced in Azoxystrobin treated roots when compared with those treated with Difenconazole.

Key words: Amistar, score, chromosomal abnormalities, mitotic index, cytotoxicity

Introduction

Increased use of pesticides for insect, weed and disease control in the past decade has proved the fact that certain agricultural chemicals may cause changes, which include inhibition of cell division, induction of chromosomal abnormalities and chromosomal damage. Chromosomal aberrations induced by agrochemicals in crop plants is widely used as an indicator of genetic damage. Grant (1978) selected root meristems as experimental systems as they are very sensitive to environmental changes and they represent normal plant-cell populations. It is a short-term assay and can be used with minimum space requirements. A number of workers have demonstrated the cytotoxic effects of different agrochemicals on plant species (Mousa, 1982; Ahmad and Yasmin, 1992; Mosuro *et al.*, 1999; Chandra *et al.*, 2002).

As new fungicides may induce many mitotic anomalies, they should undergo a rigorous testing for cytotoxic/mutagenic activity before their release due to the seriousness of the consequences.

The cytological effects of different agrochemicals on plant species have been studied by many workers (Grover and Tyagi, 1980; Njagi and Gopalan, 1981; Mousa, 1982; Amer and Ali, 1983; Soriano, 1984; Amer and Farah, 1985; Amer and Ali, 1986; Kumar and Sinha, 1989; Adam *et al.*, 1990). Fungicides are known to induce mutation and are proved to be potential mutagens (Sahu *et al.*, 1981). Many of the fungicides and their metabolic derivatives have been reported to be both carcinogenic and mutagenic (Kumar and Banerjee, 2001; Chandra *et al.*, 2002). Higher plants have been used as testorganisms for studying the effects of genotoxic substances in the environment.

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Devi *et al.* (1991) analyzed the long-term effects of fungicides on both mitotic and meiotic systems in *Allium cepa* and concluded that they can induce chromosomal aberrations. With increasing concentrations of the pesticides lindane, pirimiphos methyl, glyphosate and 2,1- metachlor: atrazine in *A. cepa* root cells (Mosuro *et al.*, 1999), endocel and monocil in *Vicia faba* root tip cells (Singh, 2001) and trifluralin in *Vicia faba* root cells (Chandra *et al.*, 2002), a decline in mitotic index was observed. These are in accordance with the result obtained with the root tip cells of *Allium cepa* treated with aluminium sulphate (Sreedevi and Bindu, 2004). Davids (1973) reported a reduction in mitotic index accompanied by inhibition of DNA synthesis by diethyl sulphate. Contradictory to this, malathion tended to increase the mitotic index of root tips of *V. faba* (Zakia *et al.*, 1990).

Mitotoxicity and clastogenicity effects were induced in onion by a variety of insecticides and pesticides like quinalphos, monocrotophos, thriam, parathion and malathion (Bhanja *et al.*, 1988; Devi *et al.*, 1991; Kiranmani *et al.*, 1994).

Different cytological aberrations viz., chromosome fragments at mitotic metaphase and chromatin bridges, fragments, laggards and bridges with fragments and/or laggards at mitotic anaphase were observed in the pesticide treated *V. faba* seeds in frequencies significantly higher than those in the control (Singh, 2001).

Many type of mitotic aberrations induced by pesticides such as binucleate cells, c-metaphase, polyploidy cell, tripolar anaphase, chromatin bridges and lagging chromosomes were observed by Chandra *et al.* (2002).

Amistar (azoxystrobin 25% EC) and Score (difenoconazole 25% EC) are the two broad spectrum, foliar, systemic fungicides, yet to be released to the farmers and planters in India by Syngenta India Limited, Mumbai. The objective of the present study was to investigate their non-target effects in terms of cytotoxicity at concentrations ranging from 0.44 - 2200 μg (a.i.) mL^{-1} . For both the fungicides, manufacturer's recommended dose for foliar spray is 2.2 μg (a.i.) mL^{-1} .

Materials and Methods

Test Material

Bulbs of *Allium cepa* L. var. Co.15 were used as the test material. They were obtained from the Horticulture Division, Tamil Nadu Agricultural University, Coimbatore, India. All sets of measurement were repeated by conducting a separate set of measurements on a separately executed experiment. Means of two sets of experiments are statistically analyzed.

Fungicide Treatment

Healthy and uniform bulbs of *Allium cepa* were selected from the same collection. The outer scales were removed from the bulbs and apices of the root primordia exposed. Bulbs were then allowed to sprout in wet sand for 72 h at $25\pm 1^\circ\text{C}$ in dark. When 10-15 roots emerged upto 0.3 to 0.5 cm long, the roots were excised and transferred to the fungicide solutions of concentrations ranging from 0.44 to 2200 μg (a.i.) mL^{-1} of Azoxystrobin and Difenoconazole and incubated at 2 different time intervals i.e., 12 and 24 h. A set retained in distilled water served as control. The experiment was conducted at room temperature ($25\pm 1^\circ\text{C}$) and three bulbs were used for each treatment.

Recovery Treatments

The fungicide treated root tips were recovered by transferring to distilled water and incubating for different time intervals viz., 0 (without any recovery period), 12, 24, 48 and 72 h.

Cytological Observations

Treated root tips were transferred to the fixative (3:1 alcohol: acetic acid) for a minimum period of 24 h. Root tips were hydrolyzed in 1N HCl at 60°C for 5 min and squashes were made in 2% acetocarmine. Mitotic index was computed (Mosuro *et al.*, 1999) by determining the mitotic cell frequency at the root tip cells as:

$$\text{Mitotic index} = \frac{\text{Number of dividing cells}}{\text{Total number of cells observed}} \times 100$$

Percentage of cells showing chromosomal abnormalities such as chromosomal non-orientation, star metaphase, stickiness, clumps, rings, univalents, breaks, bridges, laggards, chromosome fragments, multipolarity, micronuclei, binucleate cells, giant cells, trinucleate cells, nuclear vacuole and chromatin elongation were recorded at the appropriate mitotic stages.

Results

Mitostatic Effect

Table 1 gives the mitotic indices in *Allium cepa* root mersiterns treated with Azoxystrobin and Difenonazole. Generally, Difenonazole was more cytotoxic over Azoxystrobin. The mitotic index decreased significantly with increasing concentrations of the fungicides and the duration of the exposure. At the recommended dose for field application (2.2 µg (a.i.) mL⁻¹), Difenonazole depressed mitotic index by 4.305% while Azoxystrobin showed a decrease of 1.282% over untreated control at 12 h exposure period. At the highest concentration of 2200 µg (a.i.) mL⁻¹, the 24 h treatment with Difenonazole totally blocked mitosis. But in the case of Azoxystrobin, a very low percentage (2.098%) of mitosis was observed. The mitotic index recovered slowly when the treated root tips were incubated in distilled water over varying periods viz., 12, 24, 48 and 72 h.

Chromosomal Abnormalities

Data on chromosomal aberrations induced by the fungicides Azoxystrobin and Difenonazole in the root tip cells of *Allium cepa* are presented in Table 2. The extent of chromosomal abnormalities is directly related to the concentration of the active ingredients and treatment time. Both the fungicides induced maximum percentage of abnormalities during ana-telophase stage (Table 2). Of the two fungicides, Difenonazole induced the highest number of abnormalities in the dividing cells.

Non-orientation of chromatids, star metaphase, clumping, ring formation, univalents and breaks (gaps) were the abnormalities noted during metaphase stage (Table 2). Breaks, with a per cent frequency of 4.60 and clumping with 4.31 were the highest abnormalities observed in the cells treated with Azoxystrobin and Difenonazole respectively. Univalents were the infrequently observed abnormality at and above 44 µg (a.i.) mL⁻¹ in Azoxystrobin and 22 µg (a.i.) mL⁻¹ in Difenonazole.

During ana-telophase stage (Table 2), bridges, laggards, fragments, multipolar cells and micronuclei cells were commonly observed. Cells treated with the highest concentrations of Difenonazole and Azoxystrobin showed respectively 6.53 and 3.69% of micronuclei closely followed by multipolar cells.

The abnormalities noted during interphase stage (Table 2) were giant cells, nuclear vacuolation, chromatin elongation, binucleate cells and trinucleate cells, of which chromatin elongation was more frequent with a frequency of 4.414% in Difenonazole and 2.357% in Azoxystrobin. The frequency

Table 1: Impact of the fungicides Amistar and Score on the mitotic index in *Allium cepa*

Conc. (μg (a.i.) mL^{-1})	Treatment period (h)	Recovery period (h)	Cells observed	Mitosis cells	Mitotic index (%)		
Amistar	0	12	0	586	79	13.481	
			12	585	80	13.675	
			24	560	78	13.928	
			48	562	79	14.056	
	0	24	0	584	80	13.698	
			12	589	81	13.752	
			24	564	80	14.184	
			48	563	80	14.209	
	0.44	12	0	562	71	12.633	
			12	555	71	12.792	
			24	519	69	13.294	
			48	541	74	13.678	
		0.44	24	0	589	77	13.073
				12	606	81	13.366
				24	615	83	13.495
				48	618	87	14.077
1.10	12	0	525	65	12.380		
		12	560	70	12.500		
		24	564	72	12.765		
		48	524	68	12.977		
	1.10	24	0	515	70	13.592	
			12	540	68	12.592	
			24	498	64	12.851	
			48	477	62	12.997	
1.46	12	0	582	77	13.230		
		12	487	67	13.757		
		24	624	76	12.179		
		48	512	63	12.304		
	1.46	24	0	544	69	12.683	
			12	545	70	12.844	
			24	540	72	13.330	
			48	540	72	13.330	
2.20	12	0	715	89	12.447		
		12	612	78	12.745		
		24	674	89	13.204		
		48	700	94	13.428		
	2.20	24	0	699	96	13.733	
			12	450	54	12.000	
			24	472	58	12.288	
			48	515	64	12.427	
4.40	12	0	523	67	12.810		
		12	475	61	12.842		
		24	475	61	12.842		
		48	475	61	12.842		
	4.40	24	0	623	76	12.199	
			12	672	83	12.351	
			24	614	79	12.866	
			48	507	66	13.017	
4.40	12	0	572	75	13.111		
		12	598	65	10.869		
		24	524	52	10.992		
		48	420	47	11.190		
4.40	24	0	428	49	11.448		
		12	428	49	11.448		
		24	428	49	11.448		
		48	515	61	11.844		

Table 1: Continued

Conc. ($\mu\text{g a.i.mL}^{-1}$)	Treatment period (h)	Recovery period (h)	Cells observed	Mitosis cells	Mitotic index (%)	
22.00	24	0	521	54	10.364	
		12	552	60	10.869	
		24	564	62	10.992	
		48	584	65	11.130	
		72	592	69	11.655	
	12	0	587	60	10.221	
		12	563	59	10.479	
		24	571	62	10.858	
		48	619	68	10.985	
		72	657	73	11.111	
		24	0	672	62	9.226
			12	684	64	9.356
24	545		53	9.724		
48	704		68	9.659		
72	418		42	10.047		
44.00	12	0	675	62	9.185	
		12	664	62	9.337	
		24	572	54	9.440	
		48	542	53	9.778	
		72	570	57	10.000	
	24	0	705	64	9.078	
		12	621	57	9.178	
		24	648	62	9.567	
		48	654	64	9.785	
		72	607	60	9.884	
		12	0	601	46	7.653
			12	577	45	7.798
24	612		48	7.843		
48	615		50	8.130		
72	567		48	8.465		
220.00	24	0	521	39	7.485	
		12	587	48	7.666	
		24	474	37	7.805	
		48	619	49	7.915	
		72	724	59	8.149	
	12	0	625	32	5.120	
		12	631	33	5.229	
		24	669	37	5.530	
		48	654	38	5.810	
		72	606	36	5.940	
		24	0	654	33	5.045
			12	621	32	5.152
24	640		35	5.468		
48	687		38	5.531		
72	509		30	5.893		
2200.00	12	0	653	20	3.062	
		12	659	22	3.338	
		24	623	22	3.531	
		48	661	24	3.630	
		72	625	24	3.840	
	24	0	572	12	2.097	
		12	660	16	2.424	
		24	646	17	2.631	
		48	587	16	2.725	
		72	707	21	2.970	

Table 1: Continued

Conc. ($\mu\text{g (a.i.) mL}^{-1}$)	Treatment period (h)	Recovery period (h)	Cells observed	Mitosis cells	Mitotic index (%)		
0	12	0	586	79	13.481		
		12	585	80	13.675		
		24	560	78	13.928		
		48	562	79	14.056		
	24	24	72	542	77	14.206	
			0	584	80	13.698	
			12	589	81	13.752	
			24	564	80	14.184	
	0.44	12	48	563	80	14.209	
			72	584	84	14.385	
			0	786	105	13.358	
			12	450	61	13.555	
24		24	24	532	73	13.721	
			48	612	86	14.052	
			72	693	98	14.141	
			0	418	53	12.679	
		12	12	12	534	69	12.921
				24	598	78	13.043
				48	612	81	13.235
				72	657	90	13.698
1.10	12	0	446	51	11.434		
		12	473	55	11.627		
		24	481	57	11.850		
		48	499	60	12.024		
	24	24	72	515	63	12.233	
			0	452	49	10.840	
			12	412	45	10.922	
			24	517	57	11.025	
		12	12	48	546	62	11.355
				72	580	67	11.551
				0	411	43	10.462
				12	428	46	10.747
1.46	24	24	576	63	10.937		
		48	612	68	11.111		
		72	687	78	11.353		
		0	422	43	10.189		
	12	12	12	479	49	10.229	
			24	509	55	10.805	
			48	569	62	10.896	
			72	430	47	10.930	
		24	24	0	745	76	10.201
				12	633	65	10.268
				24	598	62	10.402
				48	696	74	10.632
2.20	12	72	669	72	10.762		
		0	425	39	9.176		
		12	531	50	9.416		
		24	554	54	9.747		
	24	24	48	460	46	10.000	
			72	612	62	10.130	
			0	690	67	9.710	
			12	721	71	9.847	
		12	12	24	712	71	9.971

Table 1: Continued

Conc. (μg (a.i.) mL^{-1})	Treatment period (h)	Recovery period (h)	Cells observed	Mitosis cells	Mitotic index(%)	
22.00	24	48	725	73	10.068	
		72	708	73	10.310	
		0	473	42	8.879	
		12	488	44	9.016	
		24	509	48	9.430	
		48	536	52	9.701	
	12	12	72	568	56	9.859
			0	446	32	7.174
			12	455	33	7.252
			24	482	37	7.676
			48	529	41	7.750
			72	571	48	8.406
44.00	24	0	622	44	7.073	
		12	679	49	7.216	
		24	699	52	7.439	
		48	409	31	7.579	
		72	720	55	7.638	
		0	498	21	4.216	
	12	12	12	528	23	4.356
			24	549	25	4.553
			48	606	30	4.950
			72	631	35	5.546
			0	477	20	4.192
			12	561	24	4.278
220.00	24	24	584	25	4.280	
		48	490	22	4.489	
		72	624	29	4.647	
		0	677	16	2.363	
		12	537	14	2.607	
		24	598	18	3.010	
	12	12	48	652	22	3.374
			72	468	17	3.632
			0	440	9	2.045
			12	511	11	2.152
			24	678	16	2.359
			48	488	12	2.459
440.00	12	72	657	18	2.739	
		0	499	4	0.801	
		12	463	4	0.863	
		24	502	5	0.996	
		48	487	5	1.026	
		72	515	6	1.165	
	24	12	0	635	2	0.314
			12	660	3	0.454
			24	625	3	0.480
			48	583	4	0.686
			72	527	4	0.759
			0	417	2	0.479
2200.00	12	12	579	1	0.172	
		24	472	2	0.423	
		48	541	2	0.369	
		72	445	2	0.449	
		0	620	-	-	
		12	619	-	-	
	24	12	24	628	-	-
			48	654	-	-
			72	1080	1	0.092

Note: '-' indicates the absence of mitosis

Analysis of variance for mitotic index

SV	DF	SS	MS	F
Replicate	2	0.00432	0.00216	<1
Treatment	199	10730.29414	53.92108	2230.64**
Concentration ©	9	9004.23850	1000.47094	41388.06**
Recovery (R)	4	58.45474	14.61368	604.55**
Treatment time (T)	1	17.81996	17.81996	737.19**
Fungicides (F)	1	1199.94304	1199.94304	49639.94**
CXR	36	4.57359	0.12704	5.26**
CXT	9	6.82187	0.75799	31.36**
CXF	9	406.82844	45.20316	1869.99**
RXT	4	0.25144	0.06286	2.60*
RXF	4	1.27629	0.31907	13.20**
TXF	1	5.57115	5.57115	230.47**
CXRXT	36	3.58605	0.09961	4.12**
CXRXF	36	3.96384	0.11011	4.55**
CXTXF	9	14.30183	1.58909	65.74**
RXTXF	4	0.26226	0.06556	2.71*
CXRXTXF	36	2.40115	0.6670	2.76**
Error	398	9.62083	0.02417	
Total	599	10739.91929		

cv = 1.88%, ** = Significant at 1% level, * = Significant at 5% level

Table 2: The extent of chromosomal aberration in the root meristem of *Allium cepa* treated with the fungicides Amistar and Score

Conc. ($\mu\text{g (a.i.) mL}^{-1}$)	Treatment period (h)	Recovery period (h)	Aberrant cells (%)			Total aberrant cells (%)		
			Meta phase	Ana-telphase	Inter phase			
Amistar 0	12	0	-	-	-	-		
		12	-	-	-	-		
		24	-	-	-	-		
		48	-	0.160	-	0.160		
		72	0.348	0.200	-	0.548		
		24	0	-	-	-		
	24	12	-	-	-	-		
		24	-	-	-	-		
		48	-	0.167	-	0.167		
		72	0.613	0.334	-	0.947		
		0.44	12	0	1.200	2.914	0.427	4.541
				12	0.905	2.464	0.512	3.620
24	0.459			1.505	0.242	2.206		
24	48		-	1.470	-	1.470		
	72		-	0.189	-	1.189		
	0		1.438	3.287	0.742	5.467		
1.10	12	12	1.428	2.826	0.634	4.888		
		24	1.405	2.256	0.207	3.868		
		48	0.884	1.720	0.174	2.782		
		72	0.666	1.020	-	1.686		
		24	0	1.300	2.958	0.603	4.861	
			12	1.298	2.941	0.546	4.708	
	24		0.666	2.343	0.289	3.298		
	24	48	-	1.689	0.242	1.931		
		72	-	1.492	-	1.492		
		0	1.535	3.821	0.750	6.106		
		12	1.515	3.298	0.649	5.462		
		24	1.431	2.600	0.602	4.633		

Table 2: Continued

Conc. ($\mu\text{g (a.i.) mL}^{-1}$)	Treatment period (h)	Recovery period (h)	Aberrant cells (%)			Total aberrant cells (%)
			Meta phase	Ana-telphase	Inter phase	
1.46	12	48	0.915	1.834	0.176	2.925
		72	0.705	1.047	0.169	2.281
		0	1.858	3.339	0.945	6.142
		12	1.680	3.159	0.910	5.749
		24	1.226	2.862	0.401	4.489
	24	48	0.571	2.352	0.250	3.173
		72	-	1.913	0.229	2.142
		0	2.985	4.964	1.428	9.377
		12	2.540	3.361	1.228	7.129
		24	2.225	2.739	0.817	5.781
2.20	12	48	1.873	1.900	0.784	4.557
		72	1.089	1.190	0.537	2.816
		0	2.189	3.517	1.462	7.168
		12	1.890	2.818	1.376	6.084
		24	1.437	2.773	1.035	5.245
	24	48	0.884	2.138	0.550	3.572
		72	0.598	1.875	0.444	2.917
		0	3.130	5.165	1.764	10.029
		12	3.048	3.621	1.557	8.226
		24	2.874	2.788	1.127	6.789
4.40	12	48	1.923	1.986	0.823	4.732
		72	1.219	1.871	0.544	3.634
		0	3.252	4.319	2.355	9.986
		12	3.184	3.910	1.769	8.863
		24	2.000	3.339	1.075	6.414
	24	48	1.323	2.604	0.602	4.529
		72	1.037	1.880	0.550	3.467
		0	4.306	5.194	2.721	12.221
		12	4.242	4.914	2.276	11.432
		24	3.481	4.194	1.197	8.872
22.00	12	48	2.304	2.285	1.146	5.735
		72	1.701	2.133	0.639	4.473
		0	5.122	5.391	3.050	13.563
		12	4.408	4.687	2.895	11.990
		24	3.625	3.716	2.127	9.468
	24	48	2.514	2.896	1.016	6.426
		72	1.807	2.310	0.616	4.733
		0	5.775	5.483	3.201	14.459
		12	5.588	4.507	2.910	13.005
		24	4.572	4.225	2.600	11.397
44.00	12	48	3.906	2.554	1.174	7.634
		72	2.486	2.233	1.707	6.426
		0	5.339	6.317	3.653	15.309
		12	4.705	6.210	3.245	14.160
		24	3.827	5.335	3.095	12.251
	24	48	3.063	2.842	1.873	7.778
		72	1.914	2.439	1.384	5.737
		0	7.573	6.853	3.861	18.287
		12	7.434	6.359	3.441	17.234
		24	6.557	5.190	3.434	15.181
220.00	12	48	5.734	3.151	2.030	10.915
		72	5.094	2.503	1.824	9.421
		0	8.333	9.644	6.382	24.359
		12	7.833	9.138	5.040	22.011

Table 2: Continued

Conc. ($\mu\text{g (a.i.) mL}^{-1}$)	Treatment period (h)	Recovery period (h)	Aberrant cells (%)			Total aberrant cells (%)
			Meta phase	Ana-telphase	Inter phase	
440.00	24	24	6.925	6.940	4.238	18.103
		48	6.440	6.349	3.118	15.907
		72	5.295	5.116	2.075	12.486
		0	9.888	10.130	6.483	26.501
		12	8.522	9.942	5.846	24.310
		24	8.216	7.167	4.370	19.753
	12	48	7.115	6.613	3.345	17.073
		72	6.981	4.373	2.895	14.249
		0	10.493	10.490	6.967	27.950
		12	10.339	9.948	5.646	25.933
		24	9.422	8.042	5.566	23.030
		48	8.823	7.283	3.762	19.868
2200.00	24	72	7.363	5.158	2.443	14.964
		0	10.693	10.668	8.597	29.958
		12	10.567	10.178	7.865	28.610
		24	10.153	8.170	6.352	24.675
		48	9.830	7.344	5.443	22.617
		72	8.998	5.281	4.878	19.157
	12	0	14.482	11.550	7.784	33.816
		12	13.879	10.355	7.392	31.626
		24	13.015	10.104	6.054	29.173
		48	12.500	8.244	4.743	25.487
		72	11.928	6.867	4.572	23.367
		0	15.533	11.571	9.090	36.194
24	12	14.358	11.216	8.857	34.431	
	24	13.822	10.320	8.139	32.281	
	48	13.494	8.468	6.250	28.212	
	72	12.588	8.965	4.954	26.507	

Conc. ($\mu\text{g (a.i.) mL}^{-1}$)	Treatment period (h)	Recovery period (h)	Aberrant cells (%)			Total aberrant cells (%)
			Meta phase	Ana-telphase	Inter phase	
Score						
0	12	0	-	-	-	-
		12	-	-	-	-
		24	-	-	-	-
		48	-	0.160	-	0.160
		72	0.348	0.200	-	0.548
		0	-	-	-	-
	24	12	-	-	-	-
		24	-	-	-	-
		48	-	0.167	-	0.167
		72	0.613	0.334	-	0.947
		0	1.864	4.838	3.344	10.046
		12	1.851	4.172	3.066	9.089
0.44	12	24	0.956	3.812	2.742	7.51
		48	0.875	3.264	2.108	6.247
		72	0.713	2.419	1.182	4.314
		0	3.153	4.956	3.597	11.706
		12	2.608	4.261	3.459	6.869
		24	2.285	4.245	3.225	9.755
	24	48	1.840	3.824	3.073	8.737
		72	1.553	3.802	2.272	7.627

Table 2: Continued

Conc. ($\mu\text{g (a.i.) mL}^{-1}$)	Treatment period (h)	Recovery period (h)	Aberrant cells (%)			Total aberrant cells (%)		
			Meta phase	Ana-telphase	Inter phase			
1.10	12	0	2.173	8.024	4.826	15.023		
		12	1.848	7.922	4.606	14.376		
		24	1.639	7.363	4.032	13.034		
		48	1.250	5.958	3.051	10.259		
		72	0.956	4.379	2.292	7.627		
	24	0	4.301	8.253	5.597	18.151		
		12	3.913	8.126	4.936	16.975		
		24	3.245	7.633	4.487	15.365		
		48	2.409	6.336	4.248	12.993		
		72	2.281	5.829	3.030	11.14		
		1.46	12	0	6.109	9.533	5.513	21.155
				12	5.282	8.829	4.981	19.092
24	5.219			7.589	4.385	17.193		
48	4.460			7.038	3.114	14.612		
24	72		4.578	5.909	2.469	12.956		
	0		6.200	12.056	5.623	23.879		
2.20	12	12	6.197	11.475	5.150	22.822		
		24	5.989	11.147	5.122	22.258		
		48	5.067	8.724	4.656	18.447		
		72	4.599	6.637	3.322	14.558		
	24	0	7.066	10.588	6.613	24.267		
		12	6.919	9.815	5.160	21.894		
		24	5.928	9.112	4.408	19.448		
		48	5.008	8.009	4.025	17.042		
		72	4.789	6.986	3.151	14.926		
		0	7.710	14.215	6.739	28.664		
		12	7.042	13.265	6.257	26.564		
		24	6.054	11.168	5.077	22.299		
4.40	12	48	5.353	10.526	4.829	20.708		
		72	4.618	9.452	3.385	17.455		
		0	7.942	12.476	9.621	30.039		
		12	6.486	11.660	8.931	27.077		
	24	24	6.021	11.389	8.207	25.617		
		48	5.502	10.434	7.929	23.865		
		72	5.273	10.162	6.250	21.685		
		0	7.753	14.455	10.243	32.451		
		12	7.500	14.018	9.121	30.639		
		24	6.932	13.231	8.456	28.619		
		48	5.882	12.997	8.067	26.946		
		72	4.918	11.111	6.373	22.402		
22.00	12	0	9.262	14.312	10.139	33.713		
		12	8.566	13.867	9.906	32.339		
		24	7.508	13.452	8.464	29.424		
		48	7.246	13.082	8.179	28.507		
	24	72	7.065	12.152	7.610	26.827		
		0	9.965	14.513	10.491	34.969		
		12	8.808	14.062	10.299	33.177		
		24	8.075	13.852	9.725	31.652		
		48	7.760	13.184	8.874	29.818		
		72	6.944	12.758	8.260	27.962		
		44.00	12	0	10.499	16.298	11.015	37.812
				12	10.309	15.672	10.535	36.516
24	9.863			15.313	9.962	35.138		
48	9.764			14.428	9.210	33.402		

Table 2: Continued

Conc. ($\mu\text{g (a.i.) mL}^{-1}$)	Treatment period (h)	Recovery period (h)	Aberrant cells (%)			Total aberrant cells (%)
			Meta phase	Ana-telophase	Inter phase	
220.00	24	72	9.409	14.223	9.090	32.722
		0	10.245	16.830	11.185	38.26
		12	9.893	15.939	10.839	36.671
		24	9.554	15.510	10.035	35.099
		48	8.888	15.212	9.586	33.686
		72	8.116	15.192	9.433	32.741
	12	0	12.565	17.973	11.463	42.001
		12	12.244	17.069	10.839	40.152
		24	11.985	16.719	10.211	38.915
		48	11.700	15.930	9.466	37.096
		72	11.131	15.654	9.219	36.004
		0	13.365	18.312	11.564	43.241
440.00	24	12	12.581	18.048	11.433	42.062
		24	12.186	17.927	10.498	40.611
		48	11.725	17.420	9.810	38.955
		72	11.224	17.413	9.568	38.205
		0	15.855	22.262	11.599	49.716
		12	15.068	22.118	11.054	48.24
	12	24	14.828	21.601	10.851	47.28
		48	14.056	21.497	10.305	45.858
		72	13.438	20.255	9.841	43.534
		0	16.369	23.260	11.892	51.521
		12	16.047	23.135	11.774	50.956
		24	15.834	23.017	11.623	50.474
2200.00	12	48	15.077	23.000	11.245	49.322
		72	14.310	22.997	11.229	48.536
		0	17.697	23.679	11.892	53.268
		12	17.406	23.021	11.658	52.085
		24	17.247	22.638	11.433	51.318
		48	16.928	21.654	11.269	49.851
	24	72	16.267	21.080	11.183	48.53
		0	17.719	25.911	12.437	56.067
		12	17.030	25.818	12.355	55.203
		24	16.300	25.787	12.353	54.44
		48	15.950	25.697	12.248	53.895
		72	14.740	25.301	12.203	52.244

Note: '-' indicates absence of any abnormality

Analysis of variance for chromosomal abnormalities

SV	DF	SS	MS	F
Replicate	2	0.1445	0.0722	3.35*
Treatment	119	133279.3394	669.7454	31014.49**
Concentration ©	9	75880.6171	8431.1797	390430.05**
Recovery (R)	4	4064.3157	1016.0789	47052.46**
Treatment time (T)	1	744.0852	744.0852	34457.01**
Fungicides (F)	1	46295.3341	46295.3341	2143838.74**
CXR	36	154.5242	4.2923	198.77**
CXT	9	40.2674	4.4742	207.19**
CXF	9	5442.5012	604.7224	28003.41**
RXT	4	2.3755	0.5939	27.50**
RXF	4	41.8246	10.4562	484.20**
TXF	1	16.9391	16.9391	784.41**
CXRXT	36	47.3399	1.3150	60.89**

Analysis of variance for chromosomal abnormalities

SV	DF	SS	MS	F
CXRXF	36	436.7338	12.1315	561.78**
CXTXF	9	88.5057	9.8340	455.39**
RXTXF	4	1.3784	0.3446	15.96**
CXRXTXF	36	22.5972	0.6277	29.07**
Error	398	8.5946	0.0216	
Total	599	133288.0785		

cv = 0.7%, ** = Significant at 1% level, * = Significant at 5% level

of giant cells was the lowest and were observed at and above the concentration of 44 μg (a.i.) mL^{-1} in Azoxystrobin and 4.4 μg (a.i.) mL^{-1} in Difenconazole.

Recovery of the fungicide treated cells in distilled water significantly decreased the number of cytological abnormalities. Further, a negative relationship was observed with the frequency of abnormal cells and the recovery period. The recovery from cytological abnormalities is more pronounced in Azoxystrobin treated cells when compared with those of Difenconazole treatment.

Discussion

It is obvious that many of the agrochemicals have cytotoxic and mutagenic properties and are environmentally hazardous (Burnett *et al.*, 1980). The present investigation examines mitodepressive and cytotoxic activities of the two foliar sprays, Azoxystrobin and Difenconazole in the root tip meristems of *Allium cepa*. This plant was selected as the test material in the present study because of its low chromosome number and larger chromosomal size. *Allium* species are favourable cytological materials as they also have the advantage of being available round the year and can be easily handled and cultivated (Kihlman, 1971). The cytological effects of Azoxystrobin and Difenconazole on the root cells were examined on the basis of changes in mitotic index and other induced abnormalities. A strong, dose dependant impact is obvious in terms of decline in mitotic index with increasing concentration and duration of exposure (Table 1).

Mitotic inhibition and reduction in mitotic index by fungicides Vitavax-200 and Dithane S-60 were reported by Al-Najjar and Soliman (1980) on wheat. Similar results were obtained in *Vicia faba* with Triflurain (Chandra *et al.*, 2002). Such a reduction in mitotic activity could be due to the inhibition of DNA synthesis which is considered as one of the major prerequisites for a cell to divide (Zakia *et al.*, 1990).

Cytological abnormalities induced by the two fungicides, Azoxystrobin and Difenconazole were similar to aberrations induced by other pesticides and chemical mutagens. The common abnormalities encountered during metaphase were non-orientation, star metaphase, clumping, ring formation, univalents and breaks (gaps). The ana-telophase abnormalities observed were bridges, laggards, fragments, multipolar cells and micronuclei cells. Giant cells, nuclear vacuolation, chromatin elongation, binucleate cells and trinucleate cells were observed during interphase. The frequencies of the different types of abnormalities were significantly influenced by the fungicides, their concentration and the exposure period (Table 2). The anomalies observed in the present study were also recorded by several workers in the pesticide treated root tip meristems (Adam *et al.*, 1990; Chand *et al.*, 1991; Ahmad and Yasmin, 1992; Kumar and Kumar, 2000; Singh, 2001; Chandra *et al.*, 2002).

Inhibition of spindle formation might have led to disturbed meta- and anaphase. The error of spindle organization may even lead to split or multipolar spindle. Many workers have reported the role of certain pesticides in spindle formation (Amer and Mikhael, 1986;

El-Khodary *et al.*, 1987, 1989). Stickiness and clumping of metaphase and bridge of anaphase have been attributed to the formation of dicentric chromosomes as a result of breakage and reunion (Sinha, 1989). Ring chromosomes in low frequency have been observed following treatments with Azoxystrobin and Difenoconazole. Kaur and Grover (1985) also reported low frequency ring chromosomes in root tip cells of barley treated with Anthio, Ekalux, Phendal and Rogar.

One of the frequent abnormality noticed in the present study was the appearance of ana-telophase bridges involving one or more chromosomes. They may be due to the general stickiness of chromosomes at metaphase (Al-Najjar and Soliman, 1980). Not all the bridges, especially those that appear at low concentration of the pesticides, are due to chromosome stickiness but may be due to breakage and reunion of chromosomes (El-Khodary *et al.*, 1990).

The occurrence of micronuclei has been regarded as reliable parameter for the clastogenicity/mutagenicity of an agent (Auerbach, 1976). Micronuclei may originate from lagging chromosome or from acentric fragments, which were observed in the mitotic stages (El-Khodary *et al.*, 1989; Ahmad and Yasmin, 1992). The occurrence of binucleate cells in the interphase indicates that the fungicides inhibited the cell plate formation. Binucleate cells were also noticed in organophosphorous pesticides treated root meristems of *Allium* and *Hordeum* (Grover and Malhi, 1988).

When a set of fungicide treated root tip meristems were transferred to distilled water and incubated for varying periods, the cells tended to recover from the fungitoxic effect (Table 1-2). The rate of recovery as indicated by increasing mitotic index and decreasing incidence of cytological abnormalities was highly pronounced in Azoxystrobin treated roots when compared with those treated with Difenoconazole. Similar recovery in *A. cepa* root meristems treated with Asulum, MSMA, Chlorpyrifos and Endosulfan (Rao and Rao, 1980) is already reported.

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