EFFECT OF SOME INACTIVATION PHYSICAL AND CHEMICAL AGENTS ON VIABILITY, DNA POLYMORPHISM AND GENE TRANSFER ABILITY OF TWO *Bacillus thuringiensis* BACTERIOPHAGES

Youssef, M. A. H. and Amina A. Hassan

Genetics Dept., Fac. Of Agriculture, Zagazig University.

ABSTRACT

The inactivation behavior of two *Bacillus thuringiensis* bacteriophages (A8 and R3) was studied to evaluate their potential as viral indicators for disinfection by heat, UV-C, gamma rays, chlorine, isopropanol and ethanol. The heat effect was highly efficient, PFU/ ml was decreased from 12 log (non- treated) to 1 log at 55 °C for 90 min, while 60°C was completely inactivated at 20 min. It was suggested that this killing effect may due to nucleic acid denatured. UV-C and Gamma irradiation were also dramatically decreased of PFU/ml. The lowest effect of tested biocides was for chlorine which supplied as sodium hypochlorite, isopropanol come in the second order. The ethanol at 80% concentration was inactivated all phage A8 population in 90 min and phage R3 in 15 min. In general, phage R3 was sensitive to tested agents than phage A8.

The ability of two inactivated phages to transfer ampicillin resistance gene was studied, this ability was decreased with temperature increased. It was totally inactivated with 60 °C. Two phages were able to transfer the marker gene at all UV-C, Gamma and chlorine treatments. Isopropanol and ethanol were completely inactivated the process, it was suggested that this effect due to alteration of nucleic acid and structure of phages.

The effects of some physical agents (UV-C and gamma rays) only were studied to identify DNA polymorphism through RAPD-PCR analysis in comparison to the control of the two selected bacteriophages. The result of RAPD analysis indicated the appearance of DNA polymorphic bands in response to treatments with doses of all gamma irradiation. Some bands present in the control disappeared in treated bacteriophages or vice versa.

Keyword: Inactivation, Bacillus thuringiensis, bacteriophage, radiation, RAPD-PCR

INTRODUCTION

Bacteriophages have the potential to significantly affect microbial ecology in several distinct ways. Infection of host bacteria resulted in the death of susceptible cells and production of progeny virions. Under laboratory (optimal) conditions, the production of bacteriophage at the expense of host cells can be so efficient that dense, turbid cultures are rapidly cleared due to the lysis of bacterial cells. Thus, under some conditions, bacteriophages may act to destroy sensitive bacterial cell populations.

Bacteriophages are still a major problem in the field of industrial bacterial fermentation. Cheese being one of the most important products affected (Muller *et al.*, 2005). In addition to the potential for dynamic control of bacterial populations and resultant genetic changes exerted due to selection processes. Bacteriophages are likely to be important mediators of genetic

exchange processes in bacterial populations. The genetic exchange mediated by the direct transport of host genetic material encapsidated in virion particles (transduction) or possible stimulation of cell transformation as a results of cell lysis is both a potential accelerator and confounder of engineered bioprocesses. Evolution was stimulated by genetic exchanges which may aid the adaptation of remediative communities to stress and metabolism of xenobiotic compounds. However, the risk assessment of such process is complicated by the tendency of bacteriophages to disseminate genetic materials, both chromosomal and plasmid. Thermal treatment can be used for inactivation of phages, but some authors affirmed the high heat resistance of phages (Muller *et al.*, 2005).

Irradiation treatment has been proved to be a powerful tool in inactivating human pathogenic microorganisms in water, waste water and sludge, in food and medical products (Farooq *et al.*, 1993 and Lagunas-Solar, 1995). One of the main advantages of such treatment consists of the fact that since irradiation is a physical process, no chemicals have to be added.

Exposure to a chemicals (Biocides) is an effective means to reduce the microbial population, it is widely used method of disinfection and preservation in food and other industries (Peleg, 2002).

Random amplified polymorphic DNA (RAPD) fingerprinting is a modification of the polymerase chain reaction (PCR), which utilises a single arbitrarily-chosen primer to amplify a number of fragments from a given template DNA to generate a discrete "fingerprint" when resolved by gel electrophoresis. Alterations by as little as a single base in the primer sequence lead to marked alterations in the fingerprints generated with a given template under optimized conditions. By inference, single base alterations in the genomic template DNA may also lead to changes in the RAPD fingerprints. RAPD assay was first described by two independent groups (Welsh & McClelland, 1990 and Williams et al., 1990). RAPD assay is a technique widely used for detecting alterations in DNA sequences induced by mutagenic agents such as gamma rays and is a useful analysis tool for identifying polymorphisms in DNA sequences (Kaberi and Naresh, 1998 and El-Sherbeny et al., 2005). The advantage of RAPD over other techniques is the low technical input and small quantity of DNA needed for the analysis (Hernandez et al., 1999 and Manabe et al., 1999).

The objectives of this study are to investigate the inactivation of two selected bacteriophages, namely A8 and R3 (infecting *Bacillus thuringiensis*) by some physical and chemical agents and identify DNA polymorphism through RAPD-PCR analysis in comparison to the control of the two selected bacteriophages.

MATERIALS AND METHODS

This study was carried out in Microbial and Molecular Genetics Lab., Genetics Dept., Fac. of Agric. Zagazig Univ.

Bacteriophages, bacteria and culture conditions:

Two *Bacillus thuringiensis* bacteriophages (A8 and R3) and three *B. thuringiensis* isolates (Bt2, Bt3 and Bt8) were obtained from Microbial Genetics Lab., Genetic Dept. Faculty of Agric. Zagazig Univ. Bacteria were grown routinely in nutrient agar (NA) and nutrient broth (NB) media at 30 °C for overnight. Phage propagation and counts were also performed on NA medium at 30°C and overnight incubation (Hassan and Youssef, 2011). Phages were enumerated by using double- layer agar method and expressed as plaque – forming unit per milliliter (PFU/mI).

Heat treatments:

Temperature and time combinations used to identify the thermal effects on phages were 45, 50, 55 and 60°C for 20, 50, 80 and 120 min. Phage suspensions in NB medium were distributed to tubes with a volume of 5 ml. These tubes were placed into water bath previously increased to selected temperature. Tubes were transferred to an ice bath immediately to stop the phages inactivation. Subsequently, phages titers were determined. Control counts were identified with the relating phages that were not exposed to thermal treatment (Muller *et al.*, 2005).

UV irradiation:

The phage lysates (volume 10 ml) were irradiated in sterile Petri dishes (diameter 90 mm) under 260 nm UV- lamp at different times (0, 5, 15, 30, 60, 90 min). Phage PFU/ml was quantified immediately after irradiation. **Gamma irradiation:**

The samples were irradiated in Egyptian Atomic Energy Authority-Nuclear Research Center-Inshas Egypt. The dose rate at time of irradiation was 1.4 KGy/h. Ten ml of phage lysates were irradiated using 0, 200, 400, 600, 800 and 1000 KGy

Chemical treatments:

For chemical inactivation of phages, three biocides were used: chlorine (0.5 and 1.0 mg/ml), isopropanol and ethanol (40 and 80%) at different times (0, 15, 30, 60 and 90 min). Chlorine was supplied as sodium hypochlorite, residual chlorine was neutralized by the addition of sodium thiosulphate (Duran *et al.*, 2003). Both alcohols (isopropanol and ethanol) were mixed with phage suspensions in tubes with respect to final concentration of the biocide. At intervals, tubes were removed and the surviving phages were diluted and assayed. For each experiment, non-biocide- treated phages were used as controls (Buzrul *et al.*, 2007).

Bacteriophages ability to gene transfer:

With each previous experiments, treated and non- treated phages were subjected to studying their ability to transfer ampcillin resistant gene. One ml of phage suspension was mixed with 1.0 ml of overnight liquid culture recipient in test tube for 30 min at room temperature. The mixture was diluted, 0.1 ml was spreaded onto selective medium. After incubation time (2-5 days), number of transductants were recorded and transduction frequency was calculated (Hassan and Youssef, 2011).

DNA isolation:

Phages treated with UV and gamma rays only were subjected to DNA isolation and RAPD-PCR. Assays were performed as follows; two hundred

milliliters of treated phage lysate was incubated for 1 h at 37°C. Polyethylene glycol 8000 and NaCl were added to final concentrations of 10% (w/v) and 3% (w/v), respectively. After gentle mixing, the samples were incubated overnight at 4°C. Phages were pelleted by centrifugation at 10,000 rpm at 4°C for 10 min, and the supernatants were discarded. Phage pellets were resuspended in 100-µl of 50 mM Tris (pH 8.0). Phage suspensions were extracted two times with 100 µl of phenol and then twice with phenol-chloroform-isoamyl alcohol (25:24:1). Nucleic acids were precipitated with 50 µl of 3 M sodium acetate and 1 ml of 70% (vol/vol) ethanol and resuspended in 50 µl of Tris-EDTA buffer (Prior et al., 2007).

Primers:

A set of 20 primers was analyzed based on the accurate amplified bands profiles of DNA fingerprinting using selected five different primers as shown in (Table 1).

Table 1.	Sequence and operon codes of the random primers used to)
	detection of variations in treated bacteriophages.	

Primer codes	Sequence (5 to 3)
OPA-06	GGT CCC TGA C
OPC-08	TGG ACC GGT G
OPC-20	ACT TCG CCA C
OPD-03	GTC GCC GTC A
OPD-05	TGA GCG GAC A

Amplification reaction mixture:

The amplification conditions were conducted based on Williams *et al.*, (1990) with some modifications. The reaction was prepared using 25µl per tube, containing 2µl DNA of each treatment (20 µg), 1 unit of *Taq* DNA polymerase enzyme, 2µl 10X buffer, 2 µl MgCl₂ (25 mM), 2µl dNTP_s (2.5 mM of each), 2 µl primer (10 pmol) and 14.8µl H₂O.

DNA amplification cycles:

The temperature cycling program used with a Perkin-Elmer Gene Amp PCR system (model 2400) was as follows: one cycle at 94°C for 5 min followed by 30 cycles consisting of one step of denaturation (94°C) for 1 min, one step of annealing (35°C) for 1 min, followed by one step of synthesis (72°C) for 2 min and a final extension step consisting of 72°C for 7 min and finally 4°C infinitive(Hassan and Youssef, 2011)..

Band analysis:

Reaction products were analyzed by electrophoresis on 1.4% agarose gels, stained with ethidium bromide and photographed under UV light. The synthetic DNA, ladder 100 bp (Pharmacia) was employed as molecular markers for bands molecular weight. Each amplified band profile was defined by the presence or absence of bands at particular positions on the gel. Profiles were considered different when at least one polymorphic band was identified. Fragments were scored based on standard marker using GelAnalyzer 3 (Egygene) software.

RESULTS AND DISCUSSION

Physical agents inactivation:

Fig. 1(a and b) shows the thermal inactivation of phages A8 and R3. Temperature time combination was used in these experiments. Two phages were highly influenced, increase of treatment time for each individual temperature leading to inviability of two phages. The PFU/ml was decreased from 12 log (non- treated) to 1 log at 55 °C for two phages. Totally inactivated was found by the heat treatment at 60 °C for all treatment times. Killing of two phages may be due to denaturation of nucleic acid (Bryant *et al.*, 2007). The results presented here are consistent with the results reported in literature (Quiberoni *et al.*, 2003; Bonnie, 2005, Muller *et al.*, 2005 and Buzrul *et al.*, 2007). In general, it is well known that heat effect of phages is species dependent (Muller *et al.*, 2005). The mode of action of heat is denatured capsid proteins, essential enzymes and nucleic acids if temperatures are high to enough (Bonnie, 2005). Bryant *et al.* (2007) though that DNA itself was denatured because, heat treatment showed notable killing of phage T4.

Fig. 1(c and d) shows the effect of UV and gamma rays on phages A8 and R3 viability. UV radiation is divided into UV-C (below 280 nm), UV-B (280-320 nm) and UV-A (320-390 nm). UV used in this study was from the type UV-C. Bacteriophages were irradiated at different times (dose, 0 - 90 min), which dramatically decreased PFU/ml. Phage A8 was reduced from 10 log to 2 log, while R3 from 10 log to 1 log. It can be concluded high correlation was appeared between the loss of infectivity and the increase of UV dose. These results are agreed with Sommer et al. (2001) and Said et al. (2009). Ultraviolet light disinfection was being increasingly in treatment of both waste- water and potable drinking water since such treatment does not produce disinfectant by- products (Said et al., 2009). The effectiveness of UV light in biological inactivation are due to the direct DNA damage by inducing the formation of photoproducts pyrimidine dimmer. The accumulation of DNA photoproducts caused cell death if the lesion blocks DNA synthesis and RNA transcription or can be mutagenic if the lesion is by passed by DNA polymerase. This photodimerization process has also been observed with uracil in RNA viruses. However, UV disinfection is noted to have some problems, one of which is reactivation. In response to UV damage, bacteria have developed different repair pathways, including photoenzymatic repair (PER), nucleotide excision repair and recombinational repair. The process of photoreactivation was originally described as the restoration of infectivity to UV- irradiated bacteriophages upon exposure to visible light in the presence of host bacteria (Markus et al., 1997).

For most DNA- viruses, the absence of associated photolyase genes precludes their ability to utilize visible light for repairing UV light-induced genomic damage (Wommack and Colwell 2000). Therefore, any DNA repair dependent on photoreactivation is conditional on successful infection of a host which can provide the necessary enzyme.

Gamma ray was used at different doses: 0, 200, 400, 600, 800 and 1000 Gry. Phage A8 was decreased from 7 log to 2 log, while phage R3 from

9 log to 1 log. This indicated that nucleic acid is the main target of ionizing radiation, but in viruses the coat also plays an important role. The microorganisms may be affected either by direct (single and double strand breaks) or by indirect reaction (radicals originating from water, mainly OH free radicals) (Sommer *et al.*, 2001).

Fig. 1. Effect of some physical agents, thermal (a,b) and radiation (c,d) inactivation on viability of phages A8 and R3.

Effect of UV radiation on DNA polymorphism using RAPD-PCR

RAPD technique was shown to be able in detect mutations (Atienzar *et al.*, 2002, El-Sherbeny *et al.*, 2005, Atienzar and Jha, 2006 and Dhakshanamorthy *et al.*, 2011) and genetic variation at intraspecific level (Palágyi *et al.*, 2004) even though the most common application of RAPD, which has been to determine taxonomic variation and genetic mapping (Dhakshanamorthy *et al.*, 2011). It is suggested that with proper optimization, RAPD could be a sensitive and reliable technique to detect a wide range of DNA damage (Atienzar and Jha, 2006).

In this study, 10 ml of phage lysats were irradiated using UV-C for different times including 0, 5, 15, 30, 60 and 90 min. RAPD assay was employed to detect genomic alterations and it could clearly show that some DNA changes had occurred in treated bacteriophages.

J. Agric. Chem. and Biotechn., Mansoura Univ. Vol. 3 (8), August, 2012

Evaluation of possibility mutation changes in bacteriophage DNA after the influence of UV-C by RAPD-PCR showed that five out of 20 primers could reveal some polymorphism in the amplified DNA pattern. The total number of amplified fragments from all primers for untreated and each treated bacteriophage with UV-C ray are summarized in Table 2. Each of five different random primers yielded from 3 to 5 DNA fragments whose molecular size ranged from approximately 200 to 900 bp. The total number of amplified RAPD fragments after using all five primers was 22 bands with an average of 4.4 fragments / primer. The number of polymorphic bands through each primer ranged from 2 to 5 bands per primer with an average of 3.0 polymorphic bands per primer. The total number of polymorphic amplicons produced by the 5 primers was 15 thus, representing a level of polymorphism of 68.18% at all. Primer OPD-03 produced the lowest number of fragments (3 bands) among the primers while Primers OPA-06. OPC-20 and OPD-05 produced the highest number of fragments (5 bands). Primer OPC-08 produced the lowest percentage of polymorphism (50%), while primer OPC-20 produced the highest percentage of polymorphism (100%). It was shown that single nucleotide substitutions in 10- mer primers indeed can be reflected on the amplicons' profiles. The main changes observed in the RAPD profiles have been resulted both in an appearance or disappearance of different bands with variation of their intensity as well. These effects might be connected with structural rearrangements in DNA caused by different types of DNA damages.

The fragment size of the PCR products generated by primer OPD-05 (Figs. 2 and 3) ranged from approximately 200 to 900 bp. Bands with molecular size approximately 350 and 900 bp appeared under the control and treatment for 5 min of UV-C ray in bacteriophage R3 and disappeared under other treatments, while in bacteriophage A8 the two bands disappeared. The appearance of new bands could be attributed to mutation rather than DNA damage (Atienzar and Jha, 2006).

In this study, additional or missing DNA bands were detected due to UV-C radiation and possibly consequent pyrimidin dimmer formation. UV-radiation could produced several major types of DNA lesions such as cyclobutane-type, pyrimidine dimers and the 6-4 photoproducts (Sinha and Hader 2002). Other important types of DNA damages such as protein cross-links, DNA strand breaks, deletion or insertion of base pairs could also be induced by UVirradiation.

UV radiation can change many aspects of plant processes at the physiological and DNA levels (Danon and Gallois, 1998). UV-C radiation has often been used to study various physiologically relevant responses to DNA damages, and in particular it has been shown to induce apoptosis in animal cells. UV-C radiation has also been used to induce apoptosis-like changes in *Arabidopsis thaliana* (Danon and Gallois, 1998). Jones and Kortenkamp (2000) compared RAPD fingerprints of non-mutant and mutant samples in bacteria and cultured human cells exposed to UV radiation. A dose response in the number of polymorphic bands was seen with UV irradiation of E. coli.

Table 2.	Number of	amp	licons	and the	number	of p	olymorph	nic ba	nds
	produced	by	each	RAPD	primer	for	control	and	all
	bacterioph	age	treated	l with UV	-C ray.				

	bacteriopriage treated with OV-C lay.														
		bact	A8 erio	} pha	ige			R3 bacteriophage		R3 bacteriophage			of s	ic	%
Primers	C 0trol	5 min	15 min	30 min	60 min	90 min	COtrol	5 min	15 min	30 min	60 min	90 min	Total No. c amplicon	Polymorph amplicon	Polymorphis
OPA-06	4	3	5	4	5	5	4	3	5	3	5	3	5	3	60
OPC-08	4	4	4	4	2	4	4	3	4	З	3	4	4	2	50
OPC-20	З	3	2	3	2	4	3	4	5	З	3	4	5	5	100
OPD-03	3	3	2	3	3	3	3	3	3	2	3	3	3	2	66.66
OPD-05	4	4	3	3	3	3	3	3	3	3	3	3	5	3	60
Total													22	15	68.18
Average													4.4	3.0	



Fig. 2. RAPD profile using primer OPD-05 with genomic DNA isolated from bacteriophage A8 exposed to UV-C radiation for different time periods. Lane M, molecular marker; lane C, control (without UV-C exposure); lane 3, 5 min exposure; lane 4, 15 min; lane 5, 30 min; lane 6, 60 min and lane 7, 90 min.



Fig. 3. RAPD profile using primer OPD-05 with genomic DNA isolated from bacteriophage R3 exposed to UV-C radiation for different time periods. Lane M, molecular marker; lane C, control (without UV-C exposure); lane 3, 5 min exposure; lane 4, 15 min; lane 5, 30 min; lane 6, 60 min and lane 7, 90 m

Effect of gamma ray on DNA polymorphism using RAPD-PCR

Gamma irradiation produced oxygen radicals generated by the hydrolysis of water (Puchala and Schessler, 1993) and could induce chromosomal rearrangements (Nagy et al., 1997). In this study, 10 ml of phage lysats were irradiated using different doses of gamma rays including 0, 200, 400, 600, 800 and 1000 KGy. RAPD assay was employed to detect genomic alterations and it could clearly show that some DNA changes had occurred in treated bacteriophages. Five primers chosen from the initial screening process were OPA-06, OPC-08, OPC-20, OPD-03 and OPD-05 (Table 1). Total number of amplified fragments from all primers for untreated and each treated bacteriophages with gamma ray are summarized in Table 3. Each of five different random primers yielded from 3 to 5 DNA fragments whose molecular size ranged from approximately 285 to 1000 bp. The total number of amplified RAPD fragments after using all five primers was 22 bands with an average of 4.4 fragments / primer. The number of polymorphic bands through each primer ranged from 1 to 4 bands per primer with an average of 2.4 polymorphic bands per primer. The total number of polymorphic amplicons produced by the 5 primers was 12, which representing a polymorphism level of 54.54% at all. Primer OPD-03 produced the lowest number of fragments (3 bands) and produced the lowest percentage of polymorphism (33.33%) among the primers, while primer OPC-20 produced the highest percentage of polymorphism (80%).

In this study, RAPD assay showed changes in number and presence or absence of some amplified bands between control and treated

bacteriophages. The result of RAPD analysis indicted the appearance of DNA polymorphic bands in response to treatment with doses of all gamma irradiation. Some bands present in the control disappeared in treated bacteriophages or vice versa. Differences were apparent as the appearance of new amplified bands or the disappearance of some bands. The disappearance of bands in the treated bacteriophages may be related to such events as DNA damage (single or double strand breaks, modified bases site, oxidized bases, and bulky adducts), DNA-protein cross links, point mutations and complex chromosomal rearrangements induced by gamma irradiation (Ginchner et al., 2008). However the appearance of new bands could be attributed to mutation rather than DNA damage (Hadrys et al., 1992 and Atienzar and Jha, 2006). The fragment size of the PCR products generated by primer OPA-06 (Figs. 4 and 5) ranged from approximately 285 to 950 bp. Molecular size of bands was approximately 700 and 950 bp as appeared under all the effects of gamma irradiation in bacteriophage A8 and disappeared under control, while in bacteriophage R3 two bands were appeared under irradiation by 800 and 1000 Gy. The appearance of new bands could be attributed to mutation rather than DNA damage (Hadrys et al., 1992 and Atienzar and Jha, 2006).

Irradiation was well known as the best physical mutagen which dissociates water molecules and produces hydroxyl radicals and causes oxidative damage (Ginchner *et al.*, 2008). The free radicals interact with biomolecules including DNA and scavenge electrons from them. This imposes damage to the structure and the activity of DNA. When Taq polymerase encounters a damaged DNA, there could be a number of outcomes such as blockage, bypass and possible dissociation of the enzyme-DNA which will cause the loss of bands. However, the appearance of new bands could be attributed to different DNA structural changes (Breaks, transpositions, deletion etc) (Danylchenko and Sorochinsky, 2005) rather than DNA damage (Atienzar and Jha, 2006).

Although genetic basis of the polymorphisms generated by RAPD is not well-defined, the RAPD assay proved useful a means of discovering genomic alterations induced by gamma rays. Gamma rays irradiation leads to increasing the level of DNA break formation. These different types of DNA damages could be detected by changes in RAPD profiles (Senthamizh et al., 2008). El-Sherbeny et al. (2005) used gamma rays and fast neutrons to induce mutation in Hansenula anomala and Rhodotorula rtubra and employed RAPD assay to track genomic changes and they reported that RAPD could detect changes in genetic make up of both yeasts. Wendt et al. (2001) used the RAPD markers to study the effect of gamma irradiation on potato. Ganapathi et al.(2008) studied the effect of gamma irradiation on banana using RAPD-DNA analysis. They observed changes in the DNA bands, where the main changes in the RAPD profiles of the present investigation were the appearance or disappearance of different bands with variation in their intensity. These effects might be due to the structural rearrangements in DNA caused by different types of DNA damages.

	bacteriophage treated with gamma ray.															
		A8 bacteriophage					R3 bacteriophage						of S	ic	% u	
Primers	COtrol	200 KGy	400 KGy	600 KGy	800 KGy	1000 KGy	COtrol	200 KGy	400 KGy	600 KGy	800 KGy	1000 KGy	Total No. o amplicons	Polymorph amplicons	Polymorphisr	
OPA-06	3	5	5	5	5	5	3	3	3	3	5	5	5	2	40	
OPC-08	4	4	4	3	4	4	4	3	4	4	3	4	4	2	50	
OPC-20	3	3	2	3	5	4	4	4	3	3	3	4	5	4	80	
OPD-03	3	3	2	3	2	3	3	3	3	3	3	2	3	1	33.33	
OPD-05	5	3	3	4	3	4	4	3	3	3	4	4	5	3	60	
Total													22	12	54.54	
Average													4.4	2.4		

Table 3. Number of amplicons and the number of polymorphic bands produced by each RAPD primer for control and all bacteriophage treated with gamma ray.

Fig. 4. RAPD profile using primer OPA-06 with genomic DNA isolated from bacteriophage A8 exposed to different doses of gamma rays. Lane M, molecular marker; lane C, control(without gamma rays exposure); lane 3, 200 KGy ; lane 4, 400 KGy; lane 5, 600 KGy; lane 6, 800 KGy and lane 7, 900 KGy.

Fig. 5. RAPD profile using primer OPA-06 with genomic DNA isolated from bacteriophage R3 exposed to different doses of gamma rays. Lane M, molecular marker; lane C, control(without gamma rays exposure); lane 3, 200 KGy ; lane 4, 400 KGy; lane 5, 600 KGy; lane 6, 800 KGy and lane 7, 900 KGy.

Chemical inactivation:

The first chemical agent was chlorine, it supplied as sodium hypochlorite. Residual chlorine was neutralized by the addition of sodium thiosulphate. Two concentrations of chlorine were used (0.5 and 1.0 mg/ml). In this study Fig. 6 (a and b) shows intermediate effects of chlorine on A8 and R3 phages. PFU/ml was decreased 2 log for phage A8 and 4 log for phage R3.

Previous studies were showed differences in the log_{10} reduction of the various viruses and bacteriophages. According to Maillard *et al.* (1998) phage F116, active against *P. aeruginosa* PAO1, was highly sensitive to low concentrations (5 and 75 ppm) of sodium hypochlrite, achieving at least a 4 log_{10} . Studies on *L.helveticus* (Quiberoni *et al.*, 1999) and *S. thermophilus* phages (Binetti and Reinheimer, 2000) showed a complete inactivation when 100 ppm residual – free chlorine was supplied.

Inactivation of phages B40- 8 and SS13 were not significantly different, but phages SR 51, MY 2 and Ø X 174 were a greater inactivation (Duran *et al.*, 2003). Quiberoni *et al.* (2003) reported that sodium hypochlorite at 200 - 400 ppm inactivated the tested phages completely except phage Ib3 which was only destroyed after treatment with 1200 ppm, so, the effectiveness of sodium hypochorite was phage dependents.

J. Agric. Chem. and Biotechn., Mansoura Univ. Vol. 3 (8), August, 2012

The results in Fig. 6 (c and d), (e and f) shows isopropanol and ethanol inactivation respectively, concentration of 80% was more efficient than 40% for both phages. It is fined a dose response between the concentration and time of treatment. Phage R3 is sensitive than A8, it is totally inactivation at 90 min. Isopropanol is less effective than ethanol. Ethanol at concentration of 80% inactivated all phage A8 population in 90 min and phage R3 in 15 min. These results are consistent with the results of other authors, Quiberoni *et al.* (2003) found that 10% concentration of isopropanol had no effect on phage viability and only low reductions were observed after 45 min with 50% or 100%, ethanol at a concentration of 100% produced the fastest inactivation of viral particles. Buzrul *et al.* (2007) reported that ethanol at a concentration 75% inactivated all the phage populations at 1 min, however, isopropanol at the same concentration completely inactivated the population at 15 min.

Fig. 6. Effect of some chemical agents on viability of phages A8 and R3,(a,b):chlorine inactivation,(c,d):isopropanol inactivation, (e,f):ethanol inactivation.

These results indicated that the effect of physical or chemical agents on bacteriophages are phage dependent. Phage A8 is less affected than phage R3. Ethanol at concentration 80% and 60°C are more efficient treatments, it more lethal than other tested treatments for *Bacillus thuringiensis* bacteriophages A8 and R3.

Inactivation of phages ability to gene transfer:

Phages treated with previous physical or chemical agents were subjected to study of their ability to transfer ampicillin resistant gene. Results in Tables 4, 5, 6 and 7 shows number of transduction frequency at some treatments. For thermal inactivation 50, 55, 60°C at 20 and 120 min only were chosen. The ability of two phages in gene transfer was reduced when temperature was increased and completely inactivated at 60°C. Ability of UV and gamma inactivated phages to transduce was found at all treatments, this may be due to reactivation repair mechanism of phage nucleic acid in the presences of bacterial host (Markus *et al.*, 1997). But, transduction frequency was decreased from 10^{-5} to 10^{-8} .

Effect of chlorine on gene transfer was low efficient, transduction frequency was decreased from 5.6×10^{-5} to 1.3×10^{-5} . In contrast, isopropanol and ethanol completely inactivated the ability of two phages on gene transfer (data not shown). It may suggest that, this effect was due to alteration of nucleic acid and structure of phages. Maillard *et al.* (1996 a and b) found that peractic acid altered the nucleic acid and the viral capsid of the phage F116. Isopropanol at a concentration of 100% showed a similar effect on the same phage (Maillard *et al.*, 1995 and Maillard *et al.*, 1996 a and b).

Table 4.	Effect o	of thermal	inactivation	on A8	and	R3	phages	ability	to
	gen	e transfer	-						

Transduction frequency	20	min	120 min			
Temperature (°C)	A8	R3	A8	R3		
50	6.1x 10 ⁻⁶	5.4 x10 ⁻⁶	3.6 x10 ⁻⁷	2.2x10 ⁻⁷		
55	3.5 x10 ⁻⁶	2.4 x10 ⁻⁶	3.2 x10 ⁻⁸	1.2 x 10 ⁻⁸		
60	0	0	0	0		
Transduction fraguency of non-tract	ad phagaa	A0 _ 7 0 v 10) ⁻⁵ D2 _ 22 y	10 ⁻⁵		

Transduction frequency of non-treated phages: A8 = 7.8 x 10⁻⁵ , R3 = 2.3 x10⁻⁵

Table 5. Effect of UV inactivation on the ability of A8 and R3 phages to gene transfer.

Dece (min)	Transduction frequency								
Dose (mm)	A8	R3							
0	5.6 x 10 ⁻⁵	1.3 x10⁻⁵							
5	5.1 x 10 ⁻⁵	1.2 x10 ⁻⁵							
15	4.2 x 10 ⁻⁵	1.0 x 10- ⁵							
30	9.8 x 10 ⁻⁷	4.0 x 10-'							
60	1.3 x 10 ⁻⁷	1.1 x 10 ⁻⁷							
90	4.2 x 10 ⁻⁸	1.0 x 10 ⁻⁸							

	Transductio	on frequency							
Dose (Gry)	A8	R3							
0	3.7 x 10 ⁻⁵	3.8 x10 ⁻⁵							
200	3.8 x 10 ⁻⁵	1.9 x10 ⁻⁵							
400	1.2 x 10 ⁻⁶	8.7 x 10 ⁻⁶							
600	9.8 x 10 ⁻⁷	6.1 x 10-'							
800	1.3 x 10 ⁻⁷	7.0 x 10 ⁻⁷							
1000	2.0 x 10 ⁻⁸	2.5x 10 ⁻⁸							

Table 6. Effect of gamma inactivation on the ability of A8 and R3 phages to gene transfer.

Table 7. Effect of chlorine inactivation on the ability of phage A8 and R3 to gene transfer.

Concentration (mg/ml)	Transduction frequency							
concentration (ing/iii)	A8	R3						
Non- treated phage	5.6 x 10 ⁻⁵	5.3 x 10⁵⁵						
0.5	3.8 x 10⁻⁵	3.5 x 10⁵⁵						
1.0	3.1 x 10⁻⁵	1.3 x 10⁻⁵						

In this study a very high diversity of phage inactivation was distinguished for physical and chemical tested agents against *Bacillus thuringiensis* bacteriophages A8 and R3 which were used as a model in this investigation. The kind of information obtained was important to outline the most effective thermal and some chemical treatment for inactivation of bacteriophages in laboratories, industries and environments.

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تأثير بعض العوامل الفيزيانية والكيميائية المثبطة على حيوية و تعدد صور الحامض النووى والقدرة على النقل الجينى لأثنين من بكتيريوفاجات الباسلس ثيورنجنسس محمد أبو بكر حسن يوسف وأمينة أحمد حسن قسم الورائة – كلية الزراعة – جامعة الزقازيق

تم دراسة السلوك التثبيطي لأثنين من بكتيريوفاجات بكتيريا الباسلس ثيورنجنسس هما A8, R3 وذلك لتقييمها كدليل فيروسي للتعقيم بالحرارة ، الأشعة البنفسجية ، أشعة جاما ، الكلورين ،الأيزوبروبانول و الايثانول.

كان تأثير الحرارة أكثر فاعلية، حيث نقص العدد من 12 لوغاريتم الى 1 لوغاريتم على درجة حرارة 55°م لمدة 90 دقيقة ،بينما أحدثت درجة الحرارة 60°م تثبيط كامل لكلا الفاجين عند 20 دقيقة، ويمكن إقتراح أن هذا الثاثير المميث يعزى الى تغيير فى طبيعة الحامض النووى . كما أدت العوامل الفيزيائية (الأشعة البنفسجية و أشعة جاما) الى نقص مثير فى أعداد الفاجات

أما أقل العوامل المختبرة في التأثير فكان الكلورين ويأتى الأيزوبروبانول في المرتبة الثانية ، أما الإيثانول بتركيز 80 % فقد أدى الى تثبيط كل عشيرة الفاج A8 بعد 90 دقيقة وعشيرة الفاج R3 بعد 15 دقيقة فقط. وبصفة عامة لوحظ أن الفاج R3 كان أكثر حساسية للعوامل المختبرة .

وقد تم دراسة تأثير بعض العوامل الفيزيائية (أشعة جاما و أشعة الـ UV-C) فقط بجر عات مختلفة لتوضيح الاختلافات على مستوى الـ DNA باستخدام تكنيك الـ RAPD-PCR بمقارنة البكتريوفاجات المعاملة مع الكنترول، وقد أوضحت النتائج وجود اختلافات فى الحزم نتيجة المعاملة و ذلك بظهور حزم فى البكتريوفاج المعامل دون الكنترول أو العكس. تم دراسة قدرة الفاجات المعاملة على نقل جين المقاومة للأمبسيللين ، وقد لوحظ أن هذه القدرة تقل

تم دراسة قدرة الفاجات المعاملة على نقل جين المقاومة للأمبسيللين ، وقد لوحظ أن هذه القدرة تقل بزيادة درجة الحرارة وتم تثبيط هذه العملية تماما فى الفاجات التى عوملت بدرجة حرارة 60° م ، ولكن استطاعت الفاجات المعاملة بالأشعة البنفسجية و أشعة جاما و الكلورين نقل الجين. أما الأيزوبروبانول و الايثانول فقد ثبطوا العملية تماما. يقترح أن هذا التأثير يعزى الى احداث تغيير فى تركيب الفاجات و أيضا فى تركيب الحامض النووى.

قام بتحكيم البحث

أ.د / خليفه عبد المقصود زايد
كلية الزراعة – جامعة المنصورة
أ.د / السيد ابراهيم محجوب
كلية الزراعة – جامعة الزقازيق