2015

Mechanism of Inactivation by High Voltage Atmospheric Cold Plasma Differs between Escherichia coli and Staphylococcus aureus

Lu Han
Technological University Dublin, lu.han@dit.ie

Sonal Patil
Dublin Institute of Technology, sonalpatil81@gmail.com

Daniela Boehm
Dublin Institute of Technology, 4S2929@dit.ie

vladimir Milosavljević
Dublin Institute of Technology, vm@dit.ie

Patrick Cullen
Dublin Institute of Technology, pjcullen@dit.ie

See next page for additional authors

Follow this and additional works at: https://arrow.dit.ie/schfsehart

Part of the Biotechnology Commons, and the Other Microbiology Commons

Recommended Citation

This work is licensed under a Creative Commons Attribution-Noncommercial-Share Alike 3.0 License
Authors
Lu Han, Sonal Patil, Daniela Boehm, vladimir Milosavljević, Patrick Cullen, and Paula Bourke
Mechanism of Inactivation by High Voltage Atmospheric Cold Plasma Differs between *Escherichia coli* and *Staphylococcus aureus*

Running title: Inactivation Mechanism of Atmospheric Cold Plasma

Han, L.¹, Patil, S.¹, Boehm, D.¹, Milosavljević, V.¹, Cullen, P.J.¹,², & Bourke, P.¹#

1. School of Food Science and Environmental Health, Dublin Institute of Technology, Ireland

2. School of Chemical Engineering, UNSW, Sydney, Australia

Correspondence:

# Paula Bourke, Dublin Institute of Technology, Cathal Brugha Street, Dublin 1, Ireland.

Tel: +353 1 4027594

E-mail: paula.bourke@dit.ie
Abstract:

Atmospheric cold plasma (ACP) is a promising non-thermal technology effective against a wide range of pathogenic microorganisms. Reactive oxygen species (ROS) play a crucial inactivation role when air or other oxygen containing gases are used. With strong oxidative stress, cells can be damaged by lipid peroxidation, enzyme inactivation and DNA cleavage. Identifying ROS and understanding their role is important to advance ACP applications to a range of complex microbiological issues.

In this study, the inactivation efficacy of in-package, high voltage (80 kV\textsubscript{RMS}) ACP (HVACP) and the role of intracellular ROS were investigated. Two mechanisms of inactivation were observed where reactive species were found to either react primarily with the cell envelope or to damage intracellular components. \textit{E. coli} was inactivated mainly by cell leakage and low level DNA damage. Conversely, \textit{S. aureus} was mainly inactivated by intracellular damage with significantly higher levels of intracellular ROS observed and little envelope damage. However, for both bacteria studied, increasing treatment time had a positive effect on intracellular ROS levels generated.

Keywords:

High voltage atmospheric cold plasma, in-package, intracellular ROS, cell leakage, DNA damage, \textit{E. coli} and \textit{S. aureus}
INTRODUCTION

Atmospheric cold plasma (ACP) refers to non-equilibrium plasma generated at near ambient temperatures and pressure. They are composed of particles including free electrons, radicals, positive and negative ions, but are low in collision frequency of gas discharging compared to equilibrium plasma (1, 2). ACP technologies have been widely applied for many surface treatments and environmental processes. Recently they have been studied for food sterilisation and plasma medicine (2-5).

ACP provides inactivation effects against a wide range of microbes, mainly by the generation of cell-lethal reactive species (6-8). By discharging in air, groups of reactive species are generated, such as reactive oxygen species (ROS), reactive nitrogen species (RNS), ultraviolet (UV) radiation, energetic ions and charged particles (5). However, the inactivation efficacy can be varied by changing the working gases which results in different types or amounts of reactive species generated (9-11). ROS are often identified as the principal affecting species with relatively long half-life and strong anti-microbial effects, which are generated in oxygen containing gases (12).

ROS generated during plasma discharge in air or oxygen-containing mixtures are assemblies of ozone, hydrogen peroxide, singlet and atomic oxygen, while ozone is considered as the most microbicidal specie (13). With strong oxidative stress, cells are damaged by lipid peroxidation, enzyme inactivation and DNA cleavage. Generating plasma in air or a nitrogen containing gas mixture can also generate NO\textsubscript{x} species. However, a higher inactivation efficacy has been reported with the combined
application of NO and H$_2$O$_2$ on *E. coli* than a treatment with NO or H$_2$O$_2$ alone (14). Reactive nitrogen species are highly toxic and can lead to cell death by increasing DNA damage (15). One of the potential benefits of ACP as a sterilization or pasteurization technology is the reported low mutation level associated which may be attributed to the ‘cocktail’ of reactive species generated (16, 17). However, different patterns of cellular damage between Gram negative and positive bacteria were observed in former studies (18, 19). Moreover, the treatment parameter of mode of exposure has been previously described (13, 20), where the inactivation mechanism reported was similar in relation to direct or indirect exposure to the plasma. With regard to inactivation efficacy, indirect exposure to ACP had a reduced microbicidal effect where interaction with UV, electron beam, charged particles and other short-lived species was absent. However, the in-package treatment used in this study allows the contained recombination of reactive radicals, which could result in strong bactericidal effects, even with indirect exposure.

Thus, the inactivation mechanism of ACP is a possible result of the reactive species actions, which correlate to process and system parameters. Reactive species reactions with Gram negative and positive bacteria are potentially different. To prove this hypothesis, this study compared the inactivation mechanism of HVACP against *E. coli* and *S. aureus* to expand understanding of the possible different patterns of damage against Gram negative and Gram positive bacteria, especially the action of reactive oxygen species. The interactive effects of intracellular ROS generation and DNA damage with treatment time were examined in conjunction with spectral diagnostics.
of the in package process to elucidate the mechanism.

MATERIALS AND METHODS

Bacterial Strains and Growth Conditions

The bacterial strains used in this study were Escherichia coli NCTC 12900 (non-toxigenic O157:H7) and Staphylococcus aureus ATCC 25923. Strains were chosen to represent both Gram negative and Gram positive bacteria and to facilitate comparison with other studies. They are pathogens of relevance to the food industry in addition to their multi-drug resistance and high rate of mutations (21, 22). E. coli NCTC 12900 was obtained from the National Collection of Type Cultures of the Health Protection Agency (HPA, UK), and S. aureus was obtained from the microbiology stock culture of the School of Food Science and Environmental Health, Dublin Institute of Technology. Strains were maintained as frozen stocks at -70 °C in the form of protective beads, which were plated onto tryptic soy agar (TSA, Scharlau Chemie, Barcelona, Spain) and incubated overnight at 37 °C to obtain single colonies before storage at 4 °C.

Preparation of Bacterial Cell Suspensions

Cells were grown overnight (18 h) by inoculating isolated colonies of respective bacteria in tryptic soy broth without glucose (TSB-G, Scharlau Chemie, Barcelona, Spain), at 37 °C. Cells were harvested by centrifugation at 8,720 g for 10 min. The cell pellet was washed twice with sterile phosphate buffered saline (PBS, Oxoid LTD, UK). The pellet was re-suspended in PBS and the bacterial density was determined by measuring absorbance at 550 nm using McFarland standard (BioMérieux,
Marcy-l'Étoile, France). Finally, cell suspensions with a concentration of $10^8$ CFU ml$^{-1}$ were prepared in PBS.

**HVACP system configuration**

The dielectric-barrier discharge (DBD) HVACP system used in this study consists of a high voltage transformer (with input voltage 230 V at 50 Hz), and a voltage variac (output voltage controlled within 0–120 kV) (Figure 1). HVACP discharge was generated between two 15-cm diameter aluminium electrodes separated by two perspex dielectric layers (10 mm and 1mm thickness). The system was operated at high voltage level of 80 kV$_{\text{RMS}}$ at atmospheric pressure. Voltage and input current characteristics of the system were monitored using an InfiniVision 2000 X-Series Oscilloscope (Agilent Technologies Inc., USA). A polypropylene container, which acted as both a sample holder and an additional dielectric barrier, was placed between the two perspex dielectric layers. The distance between the two electrodes was kept constant (2.2 cm) for all experiments.

**HVACP treatment**

For direct plasma treatment, 10 ml of bacterial cell suspensions in PBS were aseptically transferred to a sterile plastic petri dish, which was placed in the centre of the polypropylene container, between the electrodes. For indirect plasma treatment, a separate container was used, where the sample petri dish was placed on the upper left corner of the container, outside the plasma discharging area. Each container was sealed in a high barrier polypropylene bag (B2630; Cryovac Sealed Air Ltd, Dunkan, SC, USA) using atmospheric air as a working gas for HVACP generation. Bacterial
samples were then treated with HVACP at 80 kV\textsubscript{RMS} for 1, 3 and 5 min. After HVACP treatment, samples were subsequently stored at room temperature for either 0, 1 or 24 h (23). Ozone concentrations were measured using GASTEC gas tube detectors (Product # 18M, Gastec Corporation, Kanagawa, Japan) immediately after treatment and also after 1 or 24 h storage. Containers were kept sealed to ensure the retention of contact with generated reactive species during post-treatment storage. Microbiological analysis were immediately applied after respective post-treatment storage. All experiments were carried out in duplicate and replicated twice.

**Microbiological Analysis**

To quantify the effects of plasma treatment, 1 ml of treated samples were serially diluted in maximum recovery diluent (MRD, Scharlau Chemie, Barcelona, Spain) and 0.1 ml aliquots of appropriate dilutions were surface plated on TSA. 1 ml and 0.1 ml of the treated sample was spread onto TSA plates as described by EN ISO 11290-2 method (ISO 11290-2, 1998). The limit of detection was 1 Log CFU ml\textsuperscript{−1}. Plates were incubated at 37 °C for 24 h and colony forming units were counted. Any plates with no growth were incubated for up to 72 h and checked for the presence of colonies every 24 h. Results are reported in Log CFU ml\textsuperscript{−1} units.

**Detection of reactive oxygen species after plasma treatment**

DCFH (2',7'-dichlorodihydrofluorescein) is a cellular assay probe widely used for fluorescence detection of intracellular ROS. It revealed the concentration of ROS in HVACP treated samples.

After HVACP treatment and subsequent storage, cells were incubated with DCFH-DA
(2',7'-dichlorodihydrofluorescein diacetate, Sigma Aldrich Ltd, Dublin, Ireland) at a final concentration of 5 μM in PBS for 15 min at 37 °C. Two hundred μL aliquots of each sample were transferred into 96 well fluorescence microplate wells (Fisher Scientific, UK) and measured by Synergy™ HT Multi-Mode Microplate Reader (BioTek Instruments Inc.) at excitation and emission wave lengths of 485 and 525 nm.

**Optical emission spectroscopy**

Optical emission spectroscopy (OES) of the discharge within empty packages was acquired with an Edmund Optics UV Enhanced Smart CCD Spectrometer with an optical fibre input. UV Enhanced Smart CCD Spectrometers have been optimized for maximum performance in the ultraviolet and near UV region, and for multichannel operation with ultra-low trigger delay. The spectral resolution of the system was 0.6 nm.

The fibre optic from the spectrometer was placed facing towards the package to allow the light to cross the centre of the side wall of the polypropylene container. The fibre had a numerical aperture of 0.22 mm and was optimized for use in the ultraviolet, visible and near infrared portion of the spectrum with a wavelength range of 200 – 920 nm. A 5 mm diameter lens collected light from a column across the diameter of the package and focused it onto a 200 μm multi-mode fibre. The other end of the 2 m long fibre was connected to the spectrometer.

**Cell membrane integrity**

Membrane integrity was examined by determination of the release of intracellular materials absorbing at 260 and 280 nm (A_{260} and A_{280}) (24). Untreated (bacterial cells
in PBS) and HVACP-treated samples were centrifuged at 13,200 g for 10 min. Untreated controls were used to determine the release of any intracellular material before HVACP treatment. Two hundred µL supernatant of each sample was transferred into UV-transparent microtitre plate (Corning Life Science, US) wells and measured by Synergy™ HT Multi-Mode Microplate Reader at 260 nm and 280 nm.

**DNA damage**

To further examine intracellular damage, double-strand DNA (dsDNA) concentrations were investigated after 24 h storage, which provided adequate reaction time between ROS and cell components. SYBR Green I, [2-\([\text{N-(3-dimethylaminopropyl)}-\text{N-propylamino}]-4-[2,3\text{-dihydro}-3\text{-methyl-(benzo-1,3-thiazol-2-yl)}]-\text{methylidene]-1-phenyl-quinolinium}\), is a highly sensitive detector of dsDNA and can be used to quantify nucleic acids. SYBR Green I has been widely used in fluorescence analysis, real-time PCR and biochip applications. (25) In this study, it was used as an indicator of DNA damage with a digested cell solution. Lysozyme and lysostaphin hydrolyse the bacterial cell wall by breaking 1-4 bonds between N-acetyl-\(\beta\)-D-glucosamine (NAG), N-acetyl-\(\beta\)-D-muramic acid (NAM) and polyglycine cross-links present in the peptidoglycan (26).

Following HVACP treatment *E. coli* samples were incubated with 100 µg mL\(^{-1}\) lysozyme at 37 °C for 4 h to break the cell envelope and release the intracellular DNA. Because of the different cellular structures in Gram positive bacteria, *S. aureus* samples were incubated with 100 µg mL\(^{-1}\) lysozyme and 10 µg mL\(^{-1}\) lysostaphin at 37 °C for 4 h. Cell digestion effects were verified by colony counts on TSA plates. Cells
without HVACP treatment were digested and used as positive control group, while untreated cells without digestion were used as negative controls. The bacterial envelope was considered as completely digested when the survival rate was below the detection level.

After cell digestion, solutions were incubated with SYBR Green I (1:10,000, Sigma Aldrich Ltd, Dublin, Ireland) at working concentration (1:1) for 15 min at 37 °C. 200 µL aliquots of each sample were transferred into 96 well fluorescence microplate wells (Fisher Scientific, UK) and measured by Synergy™ HT Multi-Mode Microplate Reader at excitation and emission wave lengths of 485 and 525 nm.

**Scanning Electron Microscopy**

Bacterial samples in PBS exposed to plasma for 1 min treatment with a post-treatment storage time of 1 or 24 h were selected for SEM analysis. This was based on a noticeable difference in plasma inactivation efficacy with respect to post-treatment storage time. Bacterial cells were prepared as described by Thanomsub *et al.* 2002 with minor modifications (27, 28). Samples were then examined visually by using a FEI Quanta 3D FEG Dual Beam SEM (FEI Ltd, Hillsboro, USA) at 5 kV.

**Statistical Analysis**

Statistical analysis was performed using SPSS 22.0 (SPSS Inc., Chicago, U.S.A.). Data represent the means of experiments performed in duplicate and replicated at least twice. Means were compared using analysis of variance (ANOVA) using Fisher’s Least Significant Difference-LSD at the 0.05 level.
RESULTS

Effect of treatment time and post-storage time on plasma inactivation efficacy

The inactivation efficacy of HVACP against *E. coli* NCTC 12900 and *S. aureus* ATCC 25923 is shown in Tables 1 and 2. Inactivation was related to both treatment time and post-treatment storage time.

After 1 min exposure of HVACP, *E. coli* samples were decreased by around 2 log cycles in conjunction with 24 h post treatment storage. When treatment time was increased to 3 min, bacterial populations were undetectable for both 1 and 24 h storage times. Without post-treatment storage, approximately 3.6 and 2.3 log cycle reductions were detected with direct and indirect exposure after 3 min treatment, but further extending treatment time to 5 min resulted in 6 log cycle and at least 8 log cycle reductions for direct and indirect exposure respectively (Table 1, p≤0.05).

A similar trend of HVACP inactivation was recorded for *S. aureus*. With 24 h storage, all treatment times used led to undetectable levels of bacterial population, irrespective of the mode of exposure. Increasing treatment time, from 1 min to either 3 or 5 min, yielded undetectable levels, with direct and indirect exposure, respectively, after 1 h storage. With no post treatment storage time, populations declined by approximately 1.8 and 6.1 log cycles by increasing treatment time from 1 min to 5 min with direct exposure (Table 2, p≤0.05). Similar effects were achieved with indirect exposure.

Effect on cell membrane integrity

The absorbance of 260 and 280 nm which is commonly used for quantification of DNA and protein concentration, can also indicate the release of intracellular DNA and
protein and loss of cell integrity (24). Different trends between *E. coli* and *S. aureus* were observed from their absorbance measured at 260 nm following plasma treatment (Figure 2 and 3).

For *E. coli*, all absorption curves showed similar trends (Figure 2). With 24 h post-treatment storage, a sharp increase in absorbance followed by a steady stage indicated that the cell integrity was compromised within 1 min of HVACP treatment. In the case of 0 and 1 h post treatment storage samples, a sharp increase at 1 min of treatment was followed by a gradual increase in the absorbance as a function of treatment time (*p* ≤ 0.05). In contrast, no leakage was recorded for *S. aureus*, even after 5 min treatment (Figure 3, *p* > 0.05). However, a small increase in absorbance was observed for the 24 h post treatment storage sample group for both control and treated samples. Similar trends were observed at 280 nm (data not shown).

**Reactive oxygen and nitrogen species**

The emission spectrum is presented in Figure 4 (a). Analysis of the discharge was carried out in air at 80 kV_{RMS} over the range of 200 - 920 nm. Distinct peaks obtained in the near UV and visible regions corresponded to strong emissions from N$_2$ and N$_2^+$ excited species. The ozone concentration inside package after HVACP treatment was investigated using colorimetric tubes, which revealed its correlation with treatment and post-treatment storage time (Table 3). The in-package ozone densities were similar for each bacterial sample. Treatment time and post-treatment storage time had positive and negative effects respectively on the ozone concentration detected. Detected ozone concentration were not significantly different from containers of *E.*
coli or S. aureus samples with same treatment parameters. No ozone was detected in either treatment condition after the 24 h post-treatment storage time. In air DBD-ACPs, the well-known generation–depletion cycle of ozone is interlinked to that of nitrogen oxides through several gas-phase reactions that generate N₂O, NO and O atoms starting from O₂ and N₂ (29). In Figure 4 (b), one of the major emission intensity of second positive N₂ system from empty box and sample packages, where other major peaks had similar results (data not shown).

The concentrations of ozone and nitrogen oxides (O₃, NO₂, NO₃, N₂O₄) for this set-up were quantified using absorption spectroscopy (OAS) and are reported elsewhere (29). The measurements of ozone using the gas detectors compare with those reported using OAS.

The oxidant-sensing fluorescent probe, DCFH-DA, is a nonpolar dye, which is converted into the nonfluorescent polar derivative DCFH by cellular esterases and switched to highly fluorescent DCF when oxidized by intracellular ROS and other peroxides (30). It has been widely used for intracellular detection with fluorescence analysis. The fluorescence signal correlated with the intracellular ROS density. Figure 5 shows the intracellular ROS density results of E. coli and S. aureus in PBS, where a similar trend of ROS generation in response to HVACP is demonstrated for both bacteria. With regard to the effect of mode of exposure, with indirect treatment the ROS density increased gradually as a function of treatment time from 1 min to 5 min, by comparison with direct treatment where ROS density was lower with prolonged treatment.
DNA damage

Figure 6 presents the dsDNA quantity of *E. coli* and *S. aureus* before and after HVACP treatment. The control group from both bacteria obtained similar signal strength, which proved a similar initial DNA amount from samples. However, different signal levels were observed from the two treated strains. *E. coli* samples showed a reduction of fluorescence signal which correlated with treatment time. However, there was only a trace of fluorescence signal from *S. aureus* samples after treatment (*p* ≤ 0.05).

Scanning Electron Microscopy

From the SEM results (Figure 7), more visible damage was evident on *E. coli* surfaces than *S. aureus*, indicating cell breakage effects for *E. coli* inactivation, while HVACP treatment caused irregular shape and cell shrinkage in *S. aureus*.

Proposed Inactivation Mechanism

Figure 8 illustrates the proposed mechanism of action of ACP with Gram negative and Gram positive bacteria based on the results described here for *E. coli* and *S. aureus*. After HVACP treatment, generated reactive oxygen species, associated with process and system parameters, attack both cell envelope and intracellular components. For Gram negative cells the cell envelope is the major target of ROS. Reactions of ROS with cell components cause disruption of the cell envelope and result in leakage, with some possible damage of intracellular components (e.g. DNA). For Gram positive cells the intracellular components are the major target of ROS. Reactions of ROS will cause severe damage of intracellular components (e.g. DNA), but not cell leakage.
Lower intracellular ROS in Gram negative bacteria can be result of both ROS depletion by cell envelope components and the cell leakage.

DISCUSSION

From the results of inactivation efficacy, there is clearly a strong effect of increasing treatment time, even without post treatment storage time. However, a surviving population could be below the detection limit with recovery possible during storage under some treatment and storage conditions. No further enrichment procedures were employed in this study. Incorporating a post-treatment storage time increased the inactivation efficacy significantly, especially with 24 h post-treatment storage time, which could be attributed to the amount of reactive species generated and their extended reaction time with bacteria (Tables 1 and 2). Similar results have been observed in our former studies (18). A post treatment storage time with retained antimicrobial efficacy has two-fold potential advantage, whereby the initial exposure could be minimal with enhanced efficacy during storage which is compatible with treatment of sensitive samples. Additionally a post treatment storage stage is compatible with many industrial processes. However, with applications to the food, beverage and pharmaceutical industries in mind, the strong oxidative effect with long HVACP exposure time could adversely affect some ingredients by inducing surface oxidation, which has been observed from ozone food sterilization technologies (31). A challenge for developing HVACP applications in the food industry is to optimize the dose or gas mixtures applied to ensure control of microbiological risks whilst
maintaining food quality characteristics.

A hypothetic mechanism of action of HVACP against *E. coli* and *S. aureus* were concluded as shown in Figure 8. Different reaction mechanism with ROS and cell components are discussed below from reactive species and cell damage results.

The leakage studies recorded pointed to different modes of action. High leakage levels were observed with all treatment and post-treatment storage steps for *E. coli* (p≤0.05), but not in *S. aureus* (p>0.05) (Figure 2 and 3). The cell wall of Gram positive bacteria consists of peptidoglycan with tight structure and strength, while Gram negative bacteria are covered by a thin layer of peptidoglycan and an outer membrane of lipopolysaccharide. During plasma treatment, generated ROS can react with both lipopolysaccharide and peptidoglycan thus breaking the molecule structure by damaging C-O, C-N and C-C bonds. (32-34) However, an obvious leakage was only observed from *E. coli*. With the higher lipid content, lipid peroxidation may have taken place on lipopolysaccharides and resulted in the breakage of the cell envelope. (19) This could suggest that reactive species reacted with the cell wall in different patterns. Reactions with other cell wall components, such as peptidoglycan, could be also involved. Furthermore, Figure 7 visually illustrates the difference between *E. coli* and *S. aureus* after HVACP treatment and further supports our hypothesis on the pattern of damage. The effect of shrinkage but not breakage has also been reported on another Gram positive bacteria, *L. monocytogenes* (35).

As a main inactivation species, the ozone level inside the package showed strong correlation with treatment time and post-treatment storage time, but not with the type
of bacteria in the sample (Table 3). However, the fluorescent signal recorded for *S. aureus* was three times that of *E. coli*, thus indicating a much higher intracellular ROS density in *S. aureus* than for *E. coli* (Figure 5, *p*≤0.05). A similar time correlated ROS generation was reported by other researchers using a plasma jet treatment. Intracellular ROS increased over 5 min of treatment by air plasma from a jet (36), with a similar trend reported on generation of RNS (37). Plasma treatment time determines the input energy during discharging. As the key reactive species for oxygen containing working gases, the generation of ROS consumes most of the energy in air plasma. It has been suggested that in-package ROS can penetrate cell membranes by active transport across the lipid bilayer or transient opening of pores in the membrane (3). This could explain the correlation between treatment time and ozone/ intracellular ROS. The mode of exposure also adds complexity, where an obvious difference in reactive species was observed from OES and DCFH DA assay according to mode of exposure (Figure 4 and 5). Lower reactive species levels were detected from samples exposed to direct plasma than the indirectly exposed samples. This could be due to the quenching effect of liquid between electrodes on the ionizing of gases. However, similar inactivation levels and cell components damage were recorded. During direct treatment, undetectable ROS, mostly very short lived and transient species, might react immediately with cell components and be transformed. It appears cells were damaged by the relatively long lived species associated with indirect treatment, such as higher ozone levels.

After plasma discharging, the ozone concentrations in the gas phase were determined
to be independent of the type of bacteria, while intracellular ROS levels were strongly correlated with both process parameter and target bacteria characteristic. This could contribute to the different reaction and diffusion patterns of ROS to the cells. Based on the absorbance results at 260 nm in Figure 2 and 3, HVACP generated ROS could react with the cell wall rather than entering the cell in *E. coli* samples, whilst ROS accumulated inside the *S. aureus* cells.

*E. coli* samples showed a reduction of fluorescence signal of DNA correlating with treatment time in Figure 6. This trend elucidated that DNA damage has a plasma dose dependent pattern. There was only a trace of fluorescence signal from *S. aureus* samples post treatment, indicating greater DNA damage than with *E. coli*. It has been reported that plasma induced oxidative stress damage in *S. aureus* is due to intracellular oxidative reactions (38).

Overall, treatment time and post-treatment storage time had strong effects on inactivation efficacy against *E. coli* and *S. aureus* in this study, with a lower impact observed for mode of plasma exposure. The amount of reactive species generated, including ozone, has been correlated with inactivation efficacy (12, 36, 39-41).

Among the reactive species generated during HVACP treatment, ROS contributed as major antimicrobial factors. Their concentrations were governed by plasma dose and applied gas compositions (18). The generation of ozone as an indicator of ROS showed a time-dependent pattern, while intracellular ROS had a similar trend. During penetration, ROS could react with the lipid content in the cell membrane and cause certain damage. Compared with Gram positive bacteria, the membrane of Gram
negative bacteria was more vulnerable. Visible damage as a result of plasma exposure was previously observed for *E. coli* (13).

A much higher intracellular ROS density detected in *S. aureus* showed the probable penetration of reactive species within the cell. At the same time, higher concentrations of reactive species overall could lead to more intracellular damage to cell components such as DNA, which was clearly noted in this study. Since the total amount of ROS generated using any system or process setting is around the same level and is independent of the target bacteria characteristics, it is apparent that less cell envelope damage may be associated with more intracellular damage.

In this study, the HVACP inactivation efficacy of *E. coli* and *S. aureus* bacteria was correlated with process and system parameters (i.e. treatment time or post-treatment storage time). These determined the amount and reaction time of reactive species, which were the essential factors of antimicrobial reactions. Two different possible mechanisms of inactivation were observed in the selected Gram negative and Gram positive bacteria. Reactive species were either reacting with cell envelope or damaging intracellular components. *E. coli* was inactivated by cell envelope damage induced leakage, while *S. aureus* was mainly eliminated by intracellular damage.

Additionally, the different cell damage mechanisms might due to different type of reactive species with regard to the mode of exposure. These findings are critical for the successful development of plasma applications where the system and process parameters can be nuanced in relation to the target risk characteristics presented.

**Acknowledgements**
The research leading to these results has received funding from the European Community’s Seventh Framework Program (FP7/2207-2013) under grant agreement number 285820.

Conflict of interest

No conflict of interest.


14. Boxhammer V, Morfill GE, Jokipi J, Shimizu T, Klämpfl T, Li YF,


28. **Thanomsub B, Anupunpisit V, Chanphetch S, Watcharachaipong T,**


Table 1. Surviving cell numbers of *E. coli* NCTC 12900 with respect to treatment and post-treatment storage time

<table>
<thead>
<tr>
<th>Post-treatment storage time (h)</th>
<th>Mode of Plasma Exposure</th>
<th>Plasma treatment time (min)</th>
<th>Cell density (Log$_{10}$ CFU/ml)</th>
<th>SD*</th>
<th>Cell density (Log$_{10}$ CFU/ml)</th>
<th>SD*</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Direct</td>
<td>0</td>
<td>8.0$^a$</td>
<td>0.0</td>
<td>8.0$^a$</td>
<td>0.0</td>
</tr>
<tr>
<td></td>
<td></td>
<td>1</td>
<td>7.6$^a$</td>
<td>0.1</td>
<td>7.3$^b$</td>
<td>0.1</td>
</tr>
<tr>
<td></td>
<td></td>
<td>3</td>
<td>4.3$^b$</td>
<td>0.1</td>
<td>5.7$^c$</td>
<td>0.1</td>
</tr>
<tr>
<td></td>
<td></td>
<td>5</td>
<td>2.1$^c$</td>
<td>0.7</td>
<td>ND$^{*d}$</td>
<td>0.0</td>
</tr>
<tr>
<td></td>
<td>Indirect</td>
<td>0</td>
<td>8.0$^a$</td>
<td>0.0</td>
<td>8.0$^a$</td>
<td>0.0</td>
</tr>
<tr>
<td></td>
<td></td>
<td>1</td>
<td>7.2$^d$</td>
<td>0.1</td>
<td>7.1$^b$</td>
<td>0.2</td>
</tr>
<tr>
<td></td>
<td></td>
<td>3</td>
<td>ND$^{e}$</td>
<td>0.0</td>
<td>ND$^{d}$</td>
<td>0.0</td>
</tr>
<tr>
<td></td>
<td></td>
<td>5</td>
<td>ND$^{e}$</td>
<td>0.0</td>
<td>ND$^{d}$</td>
<td>0.0</td>
</tr>
<tr>
<td></td>
<td></td>
<td>0</td>
<td>8.0$^a$</td>
<td>0.0</td>
<td>8.0$^a$</td>
<td>0.0</td>
</tr>
<tr>
<td></td>
<td></td>
<td>1</td>
<td>5.9$^{df}$</td>
<td>0.1</td>
<td>6.1$^{be}$</td>
<td>0.8</td>
</tr>
<tr>
<td></td>
<td></td>
<td>3</td>
<td>ND$^{e}$</td>
<td>0.0</td>
<td>ND$^{d}$</td>
<td>0.0</td>
</tr>
<tr>
<td></td>
<td></td>
<td>5</td>
<td>ND$^{e}$</td>
<td>0.0</td>
<td>ND$^{d}$</td>
<td>0.0</td>
</tr>
</tbody>
</table>

Different letters indicate a significant difference at the 0.05 level between different treatment times and post-treatment storage times.

Critical controls were provided as 0 min treated samples with 0, 1 and 24 h post-treatment storage.

SD*: Standard deviation

ND*: Under detection limit
Table 2. Surviving cell numbers of *S. aureus* ATCC 25923 with respect to treatment and post-treatment storage time

<table>
<thead>
<tr>
<th>Post-treatment storage time (h)</th>
<th>Plasma treatment time (min)</th>
<th>Mode of Plasma Exposure</th>
<th>Cell density (Log$_{10}$ CFU/ml)</th>
<th>SD*</th>
<th>Cell density (Log$_{10}$ CFU/ml)</th>
<th>SD*</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td>Direct</td>
<td>Indirect</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>0</td>
<td>7.9$^a$</td>
<td>0.2</td>
<td>7.9$^a$</td>
<td>0.2</td>
<td></td>
</tr>
<tr>
<td></td>
<td>1</td>
<td>6.1$^b$</td>
<td>0.3</td>
<td>5.8$^b$</td>
<td>0.3</td>
<td></td>
</tr>
<tr>
<td></td>
<td>3</td>
<td>5.4$^c$</td>
<td>0.6</td>
<td>5.3$^c$</td>
<td>0.1</td>
<td></td>
</tr>
<tr>
<td></td>
<td>5</td>
<td>1.8$^d$</td>
<td>0.2</td>
<td>1.7$^d$</td>
<td>0.1</td>
<td></td>
</tr>
<tr>
<td></td>
<td>0</td>
<td>7.8$^a$</td>
<td>0.2</td>
<td>7.8$^a$</td>
<td>0.2</td>
<td></td>
</tr>
<tr>
<td></td>
<td>1</td>
<td>4.3$^{bf}$</td>
<td>0.0</td>
<td>2.0$^{bf}$</td>
<td>0.0</td>
<td></td>
</tr>
<tr>
<td></td>
<td>3</td>
<td>ND$^e$</td>
<td>0.0</td>
<td>ND$^e$</td>
<td>0.0</td>
<td></td>
</tr>
<tr>
<td></td>
<td>5</td>
<td>ND$^e$</td>
<td>0.0</td>
<td>ND$^e$</td>
<td>0.0</td>
<td></td>
</tr>
<tr>
<td></td>
<td>24</td>
<td>7.8$^a$</td>
<td>0.2</td>
<td>7.8$^a$</td>
<td>0.2</td>
<td></td>
</tr>
<tr>
<td></td>
<td>1</td>
<td>ND$^e$</td>
<td>0.0</td>
<td>ND$^e$</td>
<td>0.0</td>
<td></td>
</tr>
<tr>
<td></td>
<td>3</td>
<td>ND$^e$</td>
<td>0.0</td>
<td>ND$^e$</td>
<td>0.0</td>
<td></td>
</tr>
<tr>
<td></td>
<td>5</td>
<td>ND$^e$</td>
<td>0.0</td>
<td>ND$^e$</td>
<td>0.0</td>
<td></td>
</tr>
</tbody>
</table>

Different letters indicate a significant difference at the 0.05 level between different treatment times and post-treatment storage times.

Critical controls were provided as 0 min treated samples with 0, 1 and 24 h post-treatment storage.

**SD**: Standard deviation

**ND**: Under detection limit
Table 3. In-package ozone concentration after different HVACP treatment and post-treatment storage time with both *E. coli* and *S. aureus* samples

<table>
<thead>
<tr>
<th>Post-treatment storage time (h)</th>
<th>Plasma treatment time (min)</th>
<th>Ozone concentration (ppm)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td>Direct</td>
</tr>
<tr>
<td>0</td>
<td>1</td>
<td>1600</td>
</tr>
<tr>
<td></td>
<td>3</td>
<td>2400</td>
</tr>
<tr>
<td></td>
<td>5</td>
<td>4200</td>
</tr>
<tr>
<td>1</td>
<td>1</td>
<td>100</td>
</tr>
<tr>
<td></td>
<td>3</td>
<td>180</td>
</tr>
<tr>
<td></td>
<td>5</td>
<td>330</td>
</tr>
<tr>
<td>24</td>
<td>1</td>
<td>ND</td>
</tr>
<tr>
<td></td>
<td>3</td>
<td>ND</td>
</tr>
<tr>
<td></td>
<td>5</td>
<td>ND</td>
</tr>
</tbody>
</table>

ND*: Non-detectable
Figure 1. A schematic diagram of the DIT120+ HVACP device.
Figure 2. Absorbance of HVACP treated *E. coli* NCTC 12900 suspension in PBS at 260 nm with different post-treatment storage times.

Data points at 0 min treatment time refer to untreated control stored with 0, 1, 24 h in PBS.

1, 3, 5 min treatment at 80 kV\textsubscript{RMS} with 0, 1, 24 h post-treatment storage

(■ 0 h post-treatment storage time; ♦ 1 h post-treatment storage time; ▲ 24 h post-treatment storage time)

(Solid line: direct exposure; Dotted line: indirect exposure)
Figure 3. *S. aureus* ATCC 25923 absorbance at 260 nm after HVACP treatment in PBS. Data points at 0 min treatment time refer to untreated control stored with 0, 1, 24 h in PBS. 1, 3, 5 min treatment at 80 kV$_{RMS}$ with 0, 1, 24 h post-treatment storage (■ 0 h post-treatment storage time; ♦ 1 h post-treatment storage time; ▲ 24 h post-treatment storage time) (Solid line: direct exposure; Dotted line: indirect exposure)
Figure 4. Emission spectrum of dielectric barrier discharge atmospheric cold plasma operating in air under atmospheric pressure

(a) Emission spectrum of empty box

(b) Emission intensity at 336.65 nm (■ Empty box; ▲ Direct exposure; ♦ Indirect exposure.)
Figure 5. *E. coli* NCTC 12900 and *S. aureus* ATCC 25923 Intracellular ROS density assay by DCFH DA

1, 3, 5 min treatment at 80 kV<sub>RMS</sub> with 0 h post-treatment storage

( ■ *E. coli* NCTC 12900; □ *S. aureus* ATCC 25923)

* indicate a significant difference at the 0.05 level between *E. coli* and *S. aureus*
Figure 6. *E. coli* NCTC 12900 and *S. aureus* ATCC 25923 DNA quantification assay by SYBR Green 1

1, 3, 5 min treatment at 80 kV$_{RMS}$ with 24 h post-treatment storage

(*E. coli* NCTC 12900; *S. aureus* ATCC 25923)

* indicate a significant difference at the 0.05 level between *E. coli* and *S. aureus*
Figure 7. SEM images of control and treated cells with 80 kV$_{RMS}$ 1 min indirect plasma exposed following 24 h post-treatment storage

(a) Untreated *S. aureus* ATCC 25923

(b) Treated *S. aureus* ATCC 25923

(c) Untreated *E. coli* NCTC 12900

(d) Treated *E. coli* NCTC 12900
Figure 8. Proposed mechanism of action of HVACP with Gram negative and positive bacteria

a, b, c the proposed inactivation mechanism of Gram negative bacteria: a, structure of Gram negative bacteria before treatment, cell envelope consists of thin layer of peptidoglycan and lipopolysaccharide; b, ACP generated ROS attacking both cell envelope and intracellular components, where cell envelope is the major target; c, inactivation mainly caused by cell leakage, with some DNA damage possible.

c, d, e the proposed inactivation mechanism of Gram positive bacteria: c, structure of Gram positive bacteria before treatment, cell envelope consist a thick rigid layer of peptidoglycan; d, ACP generated ROS attacking both cell envelope and intracellular components, where intracellular materials are the major targets; e, inactivation mainly caused by intracellular damage (eg. DNA breakage), but not leakage.