Biomimetic Surfaces and their Effect on

Bacterial Attachment, Adhesion and

Retention

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BIOMIMETIC SURFACES AND THEIR EFFECT ON BACTERIAL ATTACHMENT, ADHESION AND RETENTION

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Abstract	5
Introduction	8
Biomimetic surfaces	9
Properties of the leaf surface	
Topography and bacteria	
Surface Topography in nature	
Surface Topography quantification	
Physicochemistry of surfaces	13
Surface Chemistry	14
Escherichia coli	14
Listeria monocytogenes	
Microbial Co-cultures	16
Attachment, adhesion and retention	
Methods and materials	
Surface Topography	
Bacterial preparation	
Attachment (Wash) and Adhesion (Spray) assays	
Retention assays	20
Scanning Electron Microscopy	21
Statistics	21
Results	22
Surface Physicochemistry	22
Gibbs free energy (AGiwi)	22
Surface free energy (7,s)	22
Lifshitz-Van der Waals forces (_{?s} LW)	23
Lewis Acid-Base forces (7 ^{sAB})	23
Electron Acceptor (γ_s^+)	23
Electron donor (ys)	24
Surface Topographies	
Arithmetical mean deviation of the surface roughness (S _a)	
Root-mean-square deviation of surface topography (S_q)	
Mean peak to valley height of surface topography (S_{pv})	29
Colony-forming units/ mL (CFU/mL) analysis	

Contents

<i>E. coli</i> in monoculture	31
L. monocytogenes in monoculture	
Microbial Co-cultures	35
<i>E. coli</i> in co-culture	35
L. monocytogenes in co-culture	
SEM analysis	
<i>E. coli</i> in monoculture SEM analysis	
L. monocytogenes in monoculture SEM analysis	41
E. coli and L. monocytogenes co-culture SEM analysis	43
SEM images	45
<i>E. coli</i> in monoculture SEM images	45
L. monocytogenes in monoculture SEM images	49
E. coli and L. monocytogenes co-culture SEM images	54
Discussion	58
Physicochemistry effect on bacteria	59
Surface topography	61
Surface roughness and its effect on bacteria	61
Monoculture and co-culture bacterial assays	63
SEM enumeration	64
SEM cell clumping	64
Conclusions	65
Limitations of the study	66
Future work	67
References	69

Index of Figures

Figure 1 Surface energies of the prepared surfaces
Figure 2 Surface topographies of the prepared surfaces, Arithmetical mean deviation (Sa),
Root-mean-square deviation of surface topographies (Sq), Mean peak to valley height of
surface topographies (Spv)
Figure 3 E. coli attachment, adhesion and retention of biomimetic surfaces presented as
Log ₁₀ CFU/cm ²
Figure 4 L. monocytogenes attachment, adhesion and retention of biomimetic surfaces
presented as Log ₁₀ CFU/cm ²
Figure 5 E. coli co-culture attachment, adhesion and retention of biomimetic surfaces
presented as Log ¹⁰ CFU/cm ² 35
Figure 6 L. monocytogenes co-culture attachment, adhesion and retention of biomimetic
surfaces presented as Log ₁₀ CFU/cm ²
Figure 7 E. coli monoculture attachment, adhesion and retention scanning electron
enumeration analysis presented as Log ₁₀ CFU/cm ² 39
Figure 8 L. monocytogenes monoculture attachment, adhesion and retention scanning
electron enumeration analysis presented as Log ₁₀ CFU/cm ²
Figure 9 E. coli and L. monocytogenes monoculture attachment, adhesion and retention
scanning electron enumeration analysis presented as Log ₁₀ CFU/cm ²
Figure 10 SEM E. coli Tenderheart attachment, adhesion and retention46
Figure 11 SEM E. coli Leek attachment, adhesion and retention46
Figure 12 SEM E. coli Cauliflower attachment, adhesion and retention46
Figure 13 SEM <i>E. coli</i> Gladioli attachment, adhesion and retention47
Figure 14 SEM <i>E. coli</i> White cabbage attachment, adhesion and retention47
Figure 15 SEM <i>E. coli</i> Flat attachment, adhesion and retention47
Figure 16 SEM <i>L. monocytogenes</i> Tenderheart attachment, adhesion and retention51
Figure 17 SEM L. monocytogenes Leek attachment, adhesion and retention51
Figure 18 SEM L. monocytogenes Cauliflower attachment, adhesion and retention51
Figure 19 SEM L. monocytogenes Gladioli attachment, adhesion and retention
Figure 20 SEM <i>L. monocytogenes</i> White cabbage attachment, adhesion and retention52
Figure 21 SEM <i>L. monocytogenes</i> Flat attachment, adhesion and retention

Figure 22 SEM E. coli and L. monocytogenes Tenderheart attachment, adhesion and	
retention	5
Figure 23 SEM E. coli and L. monocytogenes Leek attachment, adhesion and	
retention	5
Figure 24 SEM E. coli and L. monocytogenes Cauliflower attachment, adhesion and	
retention	5
Figure 25 SEM E. coli and L. monocytogenes Gladioli attachment, adhesion and	
retention	5
Figure 26 SEM <i>E. coli</i> and <i>L. monocytogenes</i> White cabbage attachment, adhesion and retention	5
Figure 27 SEM <i>E. coli</i> and <i>L. monocytogenes</i> Flat attachment, adhesion and retention	5

Index of equations

Equation 1 Δ Giwi equation	18
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This thesis would have not been possible without the support from the micro team at MMU, my family, Dexter and Professor Kathryn Whitehead, who knows the difference between a carrot and a stick.

Abstract

Biofouling in the dairy industry accounts for billions of dollars in lost product each year. Surface properties, such as macro, micro and nano topography and hydrophobicity were analysed with bacteria in monoculture and co-culture to determine the surface characteristics that prevented biofilm formation. Replica biomimetic surfaces were made using dental wax from five different types of plant leaves (White Cabbage (Brassica oleracea capitate), Leek (Allium ampeloprasu), Tender Heart (Brassica oleracea), Cauliflower (Brassica oleracea var. botrytis), and Gladioli (Gladiolus); this included a flat surface wax control. Surface physicochemistry was determined using contact angle measurements and surface topography (S_a , S_q , and S_{pv}) using optical profilometry. Monoculture and co-culture bacterial attachment, adhesion and retention assays were carried out using Escherichia coli and Listeria monocytogenes and determined using colony-forming units/mL. Scanning Electron Microscopy provided quantitative cell counts (CFU/cm²). The results demonstrated that the Tenderheart leaf surface was the most hydrophobic with the highest surface free energy, highest γ_s^{AB} , most electron-donating and most electron-accepting surface. The Leek surface demonstrated the lowest surface free energy. The White cabbage surface was the most non-polar surface, with the least γ_s^{AB} properties, the least electron-accepting and least electron-donating surface. However, it had the highest S_a and S_q values. The Cauliflower leaf surface was the least hydrophobic and least nonpolar surface whilst the Gladioli surface was found to have the highest S_{pv} values. Finally, the flat surface showed the lowest S_a , S_q and S_{pv} values. Following the attachment, adhesion and retention assays, E. coli in monoculture did not show any trends between the surface properties and the number of cells retained. However, for L. monocytogenes in monoculture, following the attachment and retention assays the Flat surface showed the least number of cells (6 Log₁₀ CFU/cm² and 4.5 Log₁₀ CFU/cm² respectively). Following the adhesion and retention assays, the Gladioli surface (highest S_{pv} values) displayed the lowest numbers of L. monocytogenes cells (6 Log10 CFU/cm² and 3.7 Log10 CFU/cm² respectively). Use of the bacteria in co-cultures demonstrated that for both the attachment and retention assays, the Tenderheart surface (most hydrophobic) displayed the lowest number of cells (4.5 Log¹⁰ CFU/cm² and 3.4 Log¹⁰ CFU/cm² for *E. coli*, 5.1 Log₁₀ CFU/cm² and 3.9 Log₁₀CFU/cm² for *L.monocytogenes* respectively). SEM analysis did not correlate with the CFU/mL assays. However, with L. monocytogenes the flat surfaces (lowest roughness) retained the lowest numbers of cells (4.7 Log_{10} cells /cm²) and regarding the co-culture, the White cabbage surface (most hydrophilic) displayed the lowest number of cells when tested for bacterial attachment, adhesion and retention (4.1 Log₁₀ cells /cm², 4.5 Log₁₀ cells /cm² and 0 Log₁₀ cells /cm² respectively. These results demonstrate that when more topographically complex surfaces are analysed, the conclusions drawn between the effect of the surface properties on bacterial attachment, adhesion and retention from more uniform surfaces do not apply. Further, the processes of bacterial attachment, adhesion and retention are different and hence differentiation between these classifications needs to be clarified. It became apparent that the varying methods used produced a wide range of results and that the use of different bacteria in monoculture and co-culture affected the microbial assays. Hence, a new approach needs to be taken to understand the cell: surface interactions on complex surfaces.

Introduction

Biofouling is a major concern in many different industries. The dairy industry must prevent biofouling not only to stop product wastage but also to avert the distribution of contaminated product that can cause transmission of bacteria to consumers (Fu *et al.*, 2016). Biofouling in the dairy industry accounts for billions of dollars in lost product each year. Causes of this include protein denaturation, protein aggregation and bacterial fouling (Bansal and Chen, 2006).

Most of these industries rely on physical cleaning such as washing down dairy equipment regularly with cleaning products or clean in place procedures (Guozhen *et al.*, 2019). Modern methods of bacterial removal are either chemical or physical. Physical removal methods can consist of the use of steam or pressure, ultraviolet light exposure and manual scrubbing. Chemical removal methods can consist of the use of sodium hypochlorite, sodium hydroxide and novel methods such as bacteriocins, bacteriophages, essential oils, non-thermal plasma and quorum sensing inhibitors (Fister *et al.*, 2016; Galié *et al.*, 2018). In the UK outbreaks transmitted by food occur multiple times a year, often linked with raw drinking milk or improper handling of meat (Pennington, 2014). In 2016, an outbreak within mixed salad leaves resulted in the transmission of a strain of *Escherichia coli* to 161 people throughout England Scotland and Wales, with 62 of the patients needing hospital care. Unfortunately, this outbreak resulted in the death of two patients (Gobin et al., 2018). In 2017, a total of 135 cases of Listeriosis were reported in England and Wales of which

30.3% of patients died (Public Health England, 2018). Removing bacteria from industrial equipment is a costly process that results in equipment downtime, for example in the paper industry downtime and breakdowns caused by bacterial build-up can cost between \$2000 and \$10,000 in every instance (Bajpai, 2015). A possible method to prevent food spoilage is to prevent initial bacterial attachment and subsequent biofilm formation. The application

of biomimetic topographies may provide a solution to stop bacterial accumulation on surfaces in the food industry (Shahali *et al.*, 2019).

Biomimetic surfaces

Biomimetic surfaces are created to replicate all or some of the properties of surfaces found in nature, such as macro, micro and nano topography, unique surface structures and chemical interactions (Hwang et al., 2015). Biomimetic surfaces can be produced by many different methods. The use of photolithography is a common method due to its high level of detail (Boyan et al., 2017). The process of creating a biomimetic surface with photolithography starts with analysing a bioLogical surface with microscopy and extracting detailed X, Y and Z spatial data, which is then used to create a greyscale digital photomask. A UV light is applied to a surface coated with a photoresist layer which uses the mask as a template. The surface is developed with a solvent to remove material from the surface leaving the etched pattern behind (Kyle et al., 2016). These methods require specific equipment throughout making them cost-intensive and so not suited to the large-scale production. Fabrication via a mould is an alternative option which is far more cost-effective without compromising detail on the microscale (Wiedemeier et al., 2017). In more recent years 3D scanning and printing have become a far more economical alternative to biomimetic surface production. Multiple sets of quantitative surface characteristic data can be obtained with magnetic resonance imaging, topography analysis and ultrasound. This data gives information about the surface and the structure below that can be printed as individual layers with a resolution up to 50nm (Xiao et al., 2019; Vaidya and Solgaard, 2018).

Properties of leaf surfaces

The surface of some plant leaves have unique properties that allow them to repel debris that has settled on their surface. The most well known of this phenomenon is the Lotus effect. Lotus leaves and many other similar Nelumbonaceae have papillae on each epidermis cell on the adaxial side. These are narrow structural protrusions coated in epicuticular wax that minimize water to leaf contact area increasing surface hydrophobicity. The papillae's primary functions concerning hydrophobicity are to increase surface area and protect the epicuticular wax (Ensikat *et al.*, 2011). Water contact pressure also determines how the surface reacts to liquid; rain droplets travelling at an average speed of 3 m/s would on a more ridged surface carry enough energy to protrude into the sub-papillae air pockets thus negating the structural hydrophobicity (Koch *et al.*, 2008). However, due to the flexibility of the leaf itself the impact energy is absorbed and so the water droplet rolls off (Ensikat *et al.*, 2011). Surface with a water contact angle <90° is considered non-wettable, any surface with a water contact angle >90° is considered non-wettable, any surface with a water contact angle >150° is considered superhydrophobic (Law, 2014).

The epicuticular wax is the principal component of the surface's hydrophobicity, if it is removed the leaf contact angle reduces from 161° to 122° consequently lowering its status from superhydrophobic to non-wettable. The epicuticular wax is thought to be the most crucial component in plant leaf composition concerning bacterial attachment (Marcell and Beattie, 2002).

Topography and bacteria

The relationship between microbial attachment and surface topography is dependent on multiple variables (S_a , S_q and S_{pv}). Surface topography can be divided into three main categories representing the size of surface variation, nano (< 0.5 µm), micro (>0.5 µm – 10 µm) and macro (<10 µm) (Rajab *et al.*, 2017). Due to the presence of multiple roughness

structures on the leaf surface, any contaminating particles are carried away with the water droplets resulting in a self-cleaning surface (Yamamoto *et al.*, 2015).

Topographies of manufactured surfaces can vary greatly depending on the method of production, surfaces with an average peak to valley height (S_a) that is similar to specific bacterial cell size may retain more of that species bacteria than surfaces with a much smaller or larger S_a . Bacterial morphology also influences retention, rod-shaped cells may not become trapped in any surface features or pits due to their elongated shape compared to cocci bacteria (Whitehead and Verran, 2006).

Nano topography has been shown to influence bacterial adhesion, by altering a glass surface, resulting in a more wettable surface (44.8°- 41.6°), decreased S_a (2.1-1.3 mm) and other surface parameters. A three-factor reduction in adhered cells was observed per mm² (Mitik-Dineva *et al.*, 2008). Macroscopic topography variations have been shown to provide a preferential position for initial bacterial deposition from the surrounding medium (Lorenzetti *et al.*, 2015).

Surface Topography in nature

Papillae and waxes exist in nature to primarily prevent fouling of the plant and water loss (Barthlott *et al.*, 2017). "Bioinspired" surfaces have been developed that take inspiration from nature replicating topographical features such as nano spikes, nanowires, and nano grass, that use these features to create a bactericidal environment. The surfaces replicated cicada and dragonfly wings and gecko skin (Tripathy *et al.*, 2017). Each surface had a wide variation of wettability and bactericidal efficacy. Some of the surfaces created were observed rupturing the cell wall in what is known as the contact killing mechanism (Tripathy *et al.*, 2017).

Sharkskin is coated in denticles situated micrometre apart from one another that move with the underlying elastic skin beneath. It is thought that the combination of the surface texture and the denticle flexibility aids preventing bacterial attachment (Kesel and Liedert, 2007). A biomimetic surface called Sharklet has been developed mimicking shark skin microstructures and attributes with an aim to reduce bacterial attachment. Sharklet has been shown to reduce *E. coli* colony forming units (CFU) by 47% when compared to a smooth control surface concerning bacterial adhesion. A 77% reduction in colony size and an 80% reduction in migration were also reported when compared to the control surface (Reddy *et al.*, 2011). It is thought that sharklet can do this due to its ability to put mechanical stress on bacterial cells that attempt to adhere to it. In a process called mechanotransduction surface structures create stress on the cell membranes when contact is made causing the cell to expend more energy to stay in contact with the surface thus making it disadvantageous to do so resulting in cell detachment (Schumacher *et al.*, 2008). Sharklet is currently being applied in the medical industry for use in catheters and wound dressings (Magin *et al.*, 2016).

Surface Topography quantification

Surface topography or roughness can be quantified in two separate ways, the measuring of one or more cross-sectional 2D lines drawn across the surface which is represented with the letter R when presenting values or 3D scanning which has recently become a more available alternative in which a map of the measured surfaces is digitally created and analysed. The types measurements remain the same as when looking at a 2D line or a 3D scan however they are represented with the letter R and S retrospectively. R_a equal to the arithmetical mean height of a surfaces profile. However, if multiple R_a measurements are made of the same surface then R_a becomes indicative of the surface S_a (Lancashire, 2017). S_a is the most commonly used method of surface roughness quantification for quality control due to its ability to give a centre line average (CLA) and relative simplicity. It does not give information about surface wavelength and lacks the sensitivity to measure smaller surface

details. S_q presents a parameter known as the surface Root mean square which describes the standard deviation of the distribution of surface heights. S_q provides more accuracy when analysing large surface deviations when compared to S_a (Gadelmawla *et al.*, 2002). S_{pv} is described as the total peak to valley height of a surface, this gives a general indication of surface roughness. However, it does not consider the shapes of surface details that form the overall roughness (Etxeberria *et al.*, 2015). Surfaces designed to test bacterial affinity are typically uniform with structured peaks and troughs and so produce highly precise quantitative topographical data (Perera-Costa *et al.*, 2014). BioLogical surfaces do not adhere to such strict architecture and so present a larger relative deviation in topographic analysis when compared to man-made surfaces (Scardino *et al.*, 2009).

Physicochemistry of surfaces

The hydrophobicity of a surface is directly related to its wettability. Many factors affect surface wettability such as surface chemistry and micro and nano topography (Härth and Schubert, 2012);(Duta *et al.*, 2015). Surface wettability is determined by the mean left and right contact angle of a liquid placed on a surface. This data is analysed to determine the surface hydrophobicity which is presented in the form of the $\Delta Giwi$ (Van Oss, 1995). $\Delta Giwi$ is equal to the surface free energy (γ_s) of the sample (Barkai *et al.*, 2016). $\Delta Giwi$ represents the amount to which a polar attraction between surface and water is greater or smaller than the polar attraction water molecules have between themselves. When the net free interaction energy between a surface and water is less than zero the surface is classed as hydrophobic, when interaction energy is greater than zero the surface is classed as hydrophobic, when interactions and the Lewis acid-base surface energy (γ_s^{AB}) representing non-polar interactions must each obtained (Faten *et al.*, 2016). Polar acid-base interactions are comprised of electron donor and acceptor interactions (Rosairo *et al.*, 2001). Acid-base interactions are typically due to electron donor (γ_s) and electron acceptor (γ_s) interactions. If a surface does not possess electron donor or acceptor then it is described as apolar, conversely, if it possesses both then it is described as bipolar (Chibowski, 1992).

Surface Chemistry

Epicuticular wax plays a crucial role in pathogen prevention in plants. Forming a smooth or crystalline layer on the external surface of plant leaves this wax is composed of very-longchain aliphatics (Buschhaus *et al.*, 2007). The composition and 3D structure of the wax creates a hostile environment in the pollysphere for bacteria. Select species of bacteria (*Bacillus* sp., *Janibacter* sp., *Kocuria* sp., *Methylobacterium* sp., *Microbacterium* sp., and Staphylococcus sp.) possess the ability to produce biosurfactants that degrade the Epicuticular wax, this grants them access to nutrients and water on the leaf surface bellow (Zeisler-Diehl et al., 2018; Siriratruengsuk et al., 2017).

Escherichia coli

Escherichia coli is a Gram-negative, rod-shaped, facultative anaerobic bacterium. Some *E. coli* strains form symbiotic relationships in humans and animals by aiding in the production of vitamin K and preventing pathogenic bacteria colonising the gastrointestinal tract (Lim *et al.*, 2010). Other strains can cause outbreak and illness. Unpasteurised milk can act as a vector for the bacterium due to the faecal-oral transmission cycle of the bacteria and the location of udders in dairy animals (McClure and Hall, 2000). In 2016, the CDC reported that dairy accounted for 11% of all outbreaks in the US, although this figure represents only unpasteurized milk (CDC, 2016). In 1999, an outbreak was linked to *E. coli* present in pasteurized milk occurred in North Cumbria, UK and 114 people were affected, with 28 being admitted to hospital, resulting in the largest *E. coli* 0157 outbreak in England and Wales recorded at that time (Goh *et al.*, 2002).

Enteric pathogenic *E. coli* infections can cause stomach cramps, bloody diarrhoea, fever and vomiting. *E. coli* 0157 has multiple virulence factors, one of which is the production of Shiga toxins which are bacteriophage encoded toxins that consist of an active subunit (A1) and five receptor binding subunits (B5). When bound to a host cell A1 is internalised into the cell cytoplasm where it inhibits cell protein synthesis, this can cause bloody diarrhoea or Haemolytic Uremic Syndrome (HUS) (Lim *et al.*, 2010). *E. coli* can also produce heatliable enterotoxin which stimulates membrane-bound adenylate cyclase in intestinal epithelial cells which leads to elevated levels of Cyclic adenosine monophosphate resulting in hypersecretion of electrolytes and water manifesting as diarrhoea in patients (Gyles, 1992).

Listeria monocytogenes

Listeria monocytogenes is a Gram-positive cocci bacterium in the firmicute family. It is a facultative anaerobe and is one of the most virulent foodborne pathogens (Chen *et al.*, 2014). Listeriosis is most commonly contracted from contaminated animal products especially soft cheeses and unpasteurised milk (Hanson *et al.*, 2019). Up to 30% of foodborne listeriosis cases in high-risk individuals such as the elderly and pregnant women can lead to meningoencephalitis, miscarriage or stillbirth and even death (Segado-Arenas *et al.*, 2018). *L. monocytogenes* infects via the oral route, once inside the hosts intestine it enters the epithelial cells and targets the liver where it will begin to multiply. In healthy patients, this is commonly the final step of the infection as a cell-mediated immune response initiates and removes the cells. However, in immunocompromised patients, the infection will continue to spread from the liver to the blood with the possibility of host death (Ramaswamy *et al.*, 2007).

L. monocytogenes contamination in the food industry is relatively common due to its virulence factors, it is a facultative halophile and can grow in temperatures as low as 0°C.

It can be extremely difficult to remove from equipment, utensils, and floors in food production factories (Huang *et al.*, 2016). Many aspects of the food industry operate at environmental temperatures between 4 and 12 °C when grown at this temperature *L. monocytogenes* displays a reduced affinity for biofilm formation (Bonaventura *et al.*, 2008). At these temperatures *L. monocytogenes* biofilms displaying sparse clusters of cells and a reduced amount of EPS. (Colagiorgi *et al.*, 2017).

Microbial Co-cultures

Mixed species biofilms are commonly found within the environment in which they often live in symbiosis and consequently enhancing the overall pathogenicity of clinically significant species (Madsen *et al.*, 2012). Symbiosis can increase the biofilms virulence factors subsequently enhancing its proclivity for attachment (Camargo *et al.*, 2017). Bacteria living in co-culture have been shown to have increased short-term mutation rates, increasing the possibility of antibiotic resistance (Frapwell *et al.*, 2018). Co-culture biofilms have also been shown to present increased resistance to chemical agents such as chlorhexidine when compared to monoculture biofilms, this was due to shared protection conferred by neutralizing enzymes or inhibitory molecules that are spread throughout the biofilm (Marsh *et al.*, 2011).

Attachment, adhesion and retention

Artificial surfaces commonly used in the food industry, such as polyethene, wood, glass rubber and stainless steel are subject to biofilm formation, this can lead to surface corrosion and alteration of product taste and smell (Galié *et al.*, 2018). There are four stages of biofilm formation, attachment, adhesion, retention and biofilm development. Each stage making

bacterial removal more challenging. These stages can be affected by a multitude of environmental factors, such as bacterial species, surface roughness and surface hydrophobicity.

Bacterial attachment is the initial stage of interaction between bacterial cell and surface. This process is controlled primarily by two main processes, Brownian motion and Van der Waals forces. Brownian motion is described as the particles move throughout a medium allowing initial contact to be made (Marshall, 1986). Van der Waals forces are weak intermolecular forces that unlike covalent bonding do not rely on chemistry, instead, they rely on electron configuration to attract atoms (Yannopapas and Vitanov, 2007). Bacterial attachment as a result of van der Waals forces is a three-step process, the first employs Lifshitz–van der Waals forces which are effective over several hundred nanometres, the second step utilises Lifshitz–van der Waals forces in conjunction with electrostatic interactions which takes place at a distance of 20nm. The third step occurs at a distance of 5nm where specific cell receptors begin the process of adhesion. Throughout the stages of attachment, bacteria still present Brownian motion and can be removed by weak fluid shear forces (Palmer *et al.*, 2007).

Bacterial adhesion creates a stronger bond between bacteria and surface. Bacterial adhesins are utilized to secure cells firmly to the structure. These adhesins can form part of bacterial appendages such as pili ad fimbriae. Multiple strains of *E. coli* poses P fimbriae which have the adhesin papG located at the fimbriae tip (Wullt, 2003). Multiple suggested theories are trying to fully explain the process of bacterial adhesion such as DVLO theory, Thermodynamic theory and Neuman's theory. An extended DVLO theory has been suggested that combines van der Waals forces, double layer interactions, acid-base interactions and hydrophobic interactions; however, this theory is not conclusive (Katsikogianni and Missirlis, 2004).

Bacterial retention becomes much more challenging due to bacterial cells utilizing surface topography and hydrophobicity, creating a more hospitable environment with the production of an extracellular polymeric matrix (EPS) (Bos *et al.*, 2000; Donlan, 2002). This step of surface colonisation requires more physical force for bacterial removal to be achieved. Bacterial retention is the final step before biofilm formation (Whitehead et al., 2015). Due to the nature of bacterial attachment the longer cells are present on a surface the more mechanisms that are in place to prevent cell detachment such as adhesins and EPS formation (Pizarro-Cerda and Cossart, 2006).

This research aimed to determine the effects of bioinspired surface properties on bacterial attachment, adhesion and retention. Surface properties, such as macro, micro and nano topography and hydrophobicity were analysed with bacteria in monoculture and co-culture to determine the surface characteristics that prevented biofilm formation.

Methods and materials

Biomimetic replicates coupons emulating five different types of plant leaves (White Cabbage (*Brassica oleracea capitate*), Leek (*Allium ampeloprasu*), Tender Heart (*Brassica oleracea*), Cauliflower (*Brassica oleracea var. botrytis*), and Gladioli (*Gladiolus*) and flat wax control were created by fabricating a silicone negative mould (Duosil silicone Shera; Germany) from the leaves by adhering them to a mould base with double-sided tape (3M). Dental wax (Kemdent Eco dental wax; UK) was poured into the negative mould creating a positive wax mould of each leaf. To create the individual coupons a 15 mm diameter steel hole punch (Trimming shop; UK) was used to create equally sized coupons.

Each biomimetic coupon was analysed with a goniometer (Krüss; Germany) using ultrapure water (BDH; UK), diiodomethane (Alfa Aesar; US) and formamide (VWR International;

US). A 5 µl drop of each liquid was applied to the surface. The left and right contact angle of the droplet was used to calculate the surface hydrophobicity with the Van Oss equation.

$$\Delta \mathbf{G} \mathbf{i} \mathbf{w} \mathbf{i} = -2\gamma_{iw} = -2 \begin{bmatrix} ((\gamma_i^{LW})^{1/2} - (\gamma_w^{LW})^{1/2})^2 + \\ 2 \begin{bmatrix} (\gamma_i^* \gamma_i^-)^{\frac{1}{2}} + (\gamma_w^* \gamma_w^-)^{\frac{1}{2}} - (\gamma_i^* \gamma_w^-)^{\frac{1}{2}} - (\gamma_w^* \gamma_i^-)^{\frac{1}{2}} \end{bmatrix} \end{bmatrix}$$

Equation 1. Δ Giwi equation (Van Oss, 1995)

Surface Topography

The coupons were measured with an optical light profilometer (Zygo; US) to find the S_a , S_q , and S_{pv} , a 3D scan of each surface was created and levelled to adjust for surface inclination. A surface cross-sectional line was also obtained from each surface. Images obtained with a maximum range of 160 µm x 160 µm in X and Y directions. For statistical analysis, three replicates for the same surface were measured five times each resulting in 15 measurements for each surface type.

Bacterial preparation

E. coli and/or *L. monocytogenes* were obtained from stock plates stored on tryptone soya agar (TSA) (Oxoid; US) at 4°C (Liebherr; Switzerland), placed into 10 mL of TSB and incubated for approximately 18 h at 37°C on an orbital shaker set at 150 rpm (New Brunswick Scientific; US). Cultures were then washed by centrifugation (Rotina 380, Hettich; Germany) at 1721 RCF three times, rinsing with sterile water in between. Cultures were diluted to an absorbance of 0.5 at 540nm on a spectrophotometer (+/- 10%) (Jenway; UK) equating to 3.38E+08 *E. coli* cells and 7.60E+08 *L. monocytogenes* cells per mL.

Attachment (Wash) and Adhesion (Spray) assays

The cell suspension was sprayed onto the biomimetic coupons using a compressed gas paint sprayer (Spraycraft; UK)for a duration of 5 s at a distance of 10 cm inside a class two laminar flow cabinet(Faster; Italy). Immediately after sterile distilled water was sprayed

onto the coupons for a duration of 5s at a distance of 10 cm. The spraying of both attachment and adhesion coupons was done in tandem, adhesion coupons were placed above the attachment as to prevent water dripping down and inadvertently washing them, both sets of coupons were adhered with double-sided tape in rows of four to a stainless steel tray which was angled at approximately 45° . The coupons from the spray assay were swabbed with 70% ethanol on their sides to remove residual bacteria. Each coupon was added to 2 mL of PBS (1 tablet per 100 mL)(Oxoid; US) and vortexed for one min. Coupons were extracted with sterile forceps and placed into a separate 2 mL of PBS and vortexed again. The contents of each of the two universals of PBS were then mixed, creating the solution to be used for serial dilutions. Each sample was diluted to 10^{6} in PBS by diluting at a ratio of 1:10. 3x10 µl drops were plated out from each dilution onto TSA, MacConkey agar (Oxoid; US) or oxford growth medium (Oxoid; US). The agar plates were incubated for 18 h at 37° C. A colony enumeration was then performed. The bacterial adhesion assay was methodologically very similar to the attachment assay except for the water rinse which did not occur.

Retention assays

Bacteria were prepared as in the bacterial preparation method. Each biomimetic coupon was submerged to 25 mL of solution for one hour at 37°C. Once incubated the cell suspension was poured off and 25 mL of sterile water was used to rinse the coupons. Each coupon was swabbed with 70% ethanol on the abaxial plane and its sides then added to 2 mL of PBS and vortexed for one minute, extracted with sterile forceps and placed into a separate two mL of PBS and vortexed again, each of the two universals of PBS were then mixed, creating the solution to be used for serial dilution. Samples were analysed as before.

In the case of monoculture assays, each experiment was run on TSA, however, for the coculture analysis *E. coli* was grown on MacConkey agar and *L. monocytogenes* was grown on Oxford growth medium. Water controls were performed following the exact protocol of the assays previously stated with the only exception being the exemption of any bacteria.

Scanning Electron Microscopy

Throughout the multiple assay's samples were created in parallel and stored in 4% a glutaraldehyde (Agar Scientific, UK) solution then rinsed with sterile distilled water and dried for SEM analysis (Zeiss supra 40VT, Germany) to be used in conjunction with the CFU analysis. Three separate images were taken of each coupon; a cell enumeration was performed and used to ascertain the cells per cm². Images were analysed at a magnification of 5000x.

Statistics

Independent samples tests were used to compare all sets of data in individual categories, data was considered to be significant if P=<0.05. One way ANOVA and Kruskal-Wallis tests were used to compare all data collected. Statistical Product and Service Solutions (SPSS) was used to perform all statistical tests for this experiment.

Results

Surface Physicochemistry

The physicochemistry was carried out on the surfaces to determine the different surface free energy parameters of the surface topographies.

Gibbs free energy (⊿Giwi)

Analysis of the $\Delta Giwi$ results demonstrated that the results were not significantly different from one another. The Tenderheart surface (-3.6 $\Delta Giwi$) displayed as the least hydrophobic surface (Figure 1a) which was closely followed by the Cauliflower leaf surface (-9.9 $\Delta Giwi$).

The Leek surface (-87 $\Delta Giwi$) and the White cabbage surface (-87.4 $\Delta Giwi$) presented as the most hydrophobic surfaces (Figure 1a). The Gladioli (-58.1 $\Delta Giwi$) and Flat (-65.1 $\Delta Giwi$) surfaces appeared central in the range of data. Thus, the Tenderheart leaf surface was the most hydrophobic, whilst the Cauliflower leaf surface was the least hydrophobic.

Surface free energy (γ_s)

A significant difference was found when comparing the surface free energy of the White cabbage (37.7 γ_s) surface against the Leek (21.2 γ_s) (P= 0.028) and Flat (34.0 γ_s) (P=0.024) surfaces. The Tenderheart (67.6 γ_s) surface demonstrated the highest surface free energy value. The Gladioli (25.5 γ_s) surface displayed the second-lowest γ_s value after the Leek surface. The Cauliflower *leaf* surface (40.8 γ_s) surface displayed the second-highest γ_s value after the Tenderheart surface (Figure 1b). In summary, the Tenderheart leaf surface demonstrated the highest surface free energy whilst the Leek surface demonstrated the lowest surface free energy.

Lifshitz-Van der Waals forces (y_sLW)

The White cabbage surface $(37.4 \gamma_s^{LW})$ when compared to the Gladioli surface $(23.2 \gamma_s^{LW})$ (P= 0.049) and the Cauliflower surface $(15.9 \gamma_s^{LW})$ (P= 0.033) showed a significant difference. The Flat surface $(32 \gamma_s^{LW})$ and the Tenderheart leaf surface $(31.5 \gamma_s^{LW})$ displayed with the second and third highest surface γ_s^{LW} . The Leek surface $(21.0 \gamma_s^{LW})$ displayed centrally in the range of data (Figure 1c). Hence, the White cabbage surface demonstrated the most non-polar surface, whilst the Cauliflower leaf surface demonstrated the least nonpolar surface.

Lewis Acid-Base forces (γ_s^{AB})

No significant difference was found when surfaces were compared to one another. The Tender heart surface $(36.2 \gamma_s{}^{AB})$ and Cauliflower leaf surfaces $(25.9 \gamma_s{}^{AB})$ showed the highest Lewis acid-base value. The Leek $(0.3 \gamma_s{}^{AB})$, White cabbage $(0.3 \gamma_s{}^{AB})$, Gladioli, $(2.3 \gamma_s{}^{AB})$ and Flat $(2.0 \gamma_s{}^{AB})$ surfaces all displayed low values (Figure 1d). In summary, the Tenderheart plant surface demonstrated the highest $\gamma_s{}^{AB}$, whilst the White cabbage surface demonstrated the least $\gamma_s{}^{AB}$.

Electron Acceptor (γ_s^+)

No significant difference was found when comparing surface electron acceptor values (γ_s^+). The Tenderheart (15 γ_s^+) surface displayed the highest electron acceptor value, with the Cauliflower (14.1 γ_s^+) surface showing a slightly lower value. The Leek (0.2 γ_s^+) surface showed the lowest value, closely followed by the White cabbage (0.6 γ_s^+) surface. The Gladioli (3.0 γ_s^+) and Flat (2.0 γ_s^+) surfaces displayed relatively low values also (Figure 1e). Thus, the Tenderheart leaf surface was the most electron-accepting surface whilst the White cabbage surface was the least.

Electron donor (γ_s)

No significant difference was found when comparing surface electron donor values (γ_s^{-}). The Tenderheart (21.8 γ_s^{-}) surface displayed the highest electron donor values, which was followed by the Cauliflower (11.0 γ_s^{-}) surface. The Leek (0.1 γ_s^{-}), Gladioli (0.4 γ_s^{-}) and Flat (0.5 γ_s^{-}) surfaces displayed relatively lower values. The White cabbage (0.1 γ_s^{-}) surface displayed the lowest value (Figure 1f). In summary, the Tenderheart leaf surface was the most electron-donating, whilst the White cabbage surface was the least.







¹b) surface free energy



1c) Lifshitz-van der Waals



1d) Lewis acid-base



Figure 1 Surface energies of the prepared surfaces

Surface Topographies

The topography of the surfaces was analysed to determine roughness values.

Arithmetical mean deviation of the surface roughness (S_a)

When comparing surface S_a several comparisons were statistically significant. When compared to the Tender heart (2.1 µm) surface the Cauliflower (3.4 µm) (P=0.000), Gladioli (2.6µm) (P=0.020), Flat (0.6 µm) (P=0.000) surfaces all showed a significant difference. When compared to the Leek surface the Leek (1.6 µm) surface the Cauliflower (P=0.000), Gladioli (P=0.033), White cabbage (3.8 µm) (P=0.021), and Flat (P=0.000) surfaces all showed a significant difference. When compared to the Cauliflower surface the Gladioli (P=0.021), White cabbage (P=0.000) and Flat (P=0.000) surfaces all showed a significant difference. When compared to the Gladioli surface the White cabbage (P=0.001), and flat (P=0.000) surfaces showed a significant difference. When compared to the White cabbage surface the flat (P=0.000) showed a significant difference. When compared to the White cabbage surface the flat (P=0.000) showed a significant difference. When compared to the White cabbage surface the flat (P=0.000) showed a significant difference. When compared to the White cabbage surface the flat (P=0.000) showed a significant difference. When compared to the White cabbage surface the flat (P=0.000) showed a significant difference. When compared to the White cabbage surface the flat (P=0.000) showed a significant difference. When compared to the White cabbage surface the flat (P=0.000) showed a significant difference. White cabbage, Cauliflower and Gladioli presented with high Sa values. The Flat surface displayed the lowest Sa. In summary, the White cabbage surface was found to have the highest Sa value and the Flat surface showed the lowest (Figure 2a).

Root-mean-square deviation of surface topography (S_q)

When comparing surface S_q a number of comparisons were significant. When compared to the Tenderheart (2.2µm) surface the Cauliflower (4.3 µm) (P=0.000) surface showed a significant difference. When compared to the Leek (2 µm) surface the Cauliflower (P=0.000), Gladioli (3.9 µm) (P=0.000), White cabbage (5.2 µm) (P=0.044), and Flat (0.8 µm) (P=0.000) surfaces all showed a significant difference. When compared to the Cauliflower surface the Gladioli (P=0.008), White cabbage (P=0.000) and Flat (P=0.000) surfaces all showed a significant difference. When compared to the Gladioli surface the White cabbage (P=0.001) and Flat (P=0.00) surfaces showed a significant difference. When compared to the White cabbage surface the flat (P=0.00) surface showed a significant difference. In summary, the White cabbage surface presented with the highest Sq values while the Flat surface displayed the lowest. (Figure 2b).

Mean peak to valley height of surface topography (S_{pv})

When comparing surface S_{pv} the Tender heart (54.3 µm) surface compared with the Cauliflower (57.8 µm) (P=0.002) and Flat (6.7 µm) (P=0.000) surfaces there were significant differences demonstrated. The Leek (38.1 µm) surface compared with the Cauliflower (P=0.000) and Flat (P=0.000) surfaces also showed significant differences. The Cauliflower surface compared with the White cabbage (52.9 µm) (P=0.014), Gladioli (60.3 µm) (P=0.019) and Flat (P=0.000) surfaces showed significant differences, as well as the White cabbage surface compared with the Flat (P=0.000) surface. In summary, the Gladioli surface was found to have the highest mean peak to valley height of surface topography and the Flat surface had the lowest (Figure 2c).





Figure 2 Surface topographies of the prepared surfaces a) Arithmetical mean deviation (Sa) b) Root-mean-square deviation of surface topographies (Sq) c) Mean peak to valley height of surface topographies (Spv)

Colony-forming units/ mL (CFU/mL) analysis

Surfaces were analysed with attachment, adhesion and retention assays to determine the effect varying topographies had on pre-biofilm surface colonisation.

E. coli in monoculture

Attachment assays

No significant difference was found when comparing the numbers following the bacterial attachment assays. The Leek ($6.3 \text{ Log}_{10} \text{ CFU/cm}^2$) surface displayed the highest number of *E. coli* cells attached. The Gladioli ($6.2 \text{ Log}_{10} \text{ CFU/cm}^2$) and Flat ($6.3 \text{ Log}_{10} \text{ CFU/cm}^2$) surfaces displayed centrally in the range of data. The Cauliflower ($6.1 \text{ Log}_{10} \text{ CFU/cm}^2$) surface displayed the lowest number of *E. coli* cells attached, followed by the Tenderheart ($4.12 \text{ Log}_{10} \text{ CFU/cm}^2$) and White cabbage ($6.12 \text{ Log}_{10} \text{ CFU/cm}^2$) surfaces. In summary, the Cauliflower surface was found to prevent attachment of *E. coli* cells the most and the Leek surface displayed the highest number of cells attached (Figure 3).

Adhesion assays

No significant difference was found when comparing the results of the bacterial adhesion experiments of *E. coli* in monoculture. The Leek (6.7 $\text{Log}_{10} \text{ CFU/cm}^2$) surface displayed the highest *E. coli* cells adhered, flowed by the Gladioli (6.6 $\text{Log}_{10} \text{ CFU/cm}^2$) surface. The White cabbage (6.5 $\text{Log}_{10} \text{ CFU/cm}^2$) and Cauliflower (6.57 $\text{Log}_{10} \text{ CFU/cm}^2$) surfaces were found to be central in the range of data. The Flat (6.26 $\text{Log}_{10} \text{ CFU/cm}^2$) surface displayed the lowest number of cells adhered flowed by the Tenderheart (2.3 $\text{Log}_{10} \text{ CFU/cm}^2$) surface. In summary, the Flat surface was found to prevent adhesion of *E. coli* cells the most while the Leek surfaces displayed the highest number of cells adhered flowed by the Tenderheart (Figure 3).

Retention assays

Several significant differences were found when comparing the bacterial retention assays. The Cauliflower (4.4 Log_{10} CFU/cm²) surface compared with the Tenderheart (4.2 Log_{10} CFU/cm²) (P=0.046), Leek (4.6 Log_{10} CFU/cm²) (P=0.046) White cabbage (3.6 Log_{10} CFU/cm²) (p=0.052) and Flat (5.4 Log_{10} CFU/cm²) (P=0.021) surfaces was all found to be significant. No significant difference was found when the Gladioli (4.7 Log_{10} CFU/cm²) surface was compared to all other surfaces. The Flat surface presented with the highest number of *E. coli* cells retained, while the White Cabbage presented with the lowest. In summary, the White cabbage surface was found to prevent retention of *E. coli* cells the most while the Flat surfaces retained the highest number of cells (Figure 3).





L. monocytogenes in monoculture

Attachment assays

When compared to the White cabbage (6.0 $\text{Log}_{10} \text{ CFU/cm}^2$) surface, the Gladioli (6.1 $\text{Log}_{10} \text{ CFU/cm}^2$) (P=0.046) and Tender heart (6.2 $\text{Log}_{10} \text{ CFU/cm}^2$) (P=0.0009) surfaces showed significant differences concerning the numbers of bacteria attached. The Cauliflower (6 $\text{Log}_{10} \text{ CFU/cm}^2$) surface-displayed central in the range of data. In summary, the Leek surface displayed the most *L. monocytogenes* cells attached while the Flat surface showed the least (Figure 4).

Adhesion assays

When compared to the White cabbage (6.8 Log_{10} CFU/cm²) surface, the Tender heart (6.3 Log_{10} CFU/cm²) (P=0.004), Leek (6.8 Log_{10} CFU/cm²) (P=0.04), and Flat (6.7 Log_{10} CFU/cm²) (P=0.042) surfaces showed significant differences in the results as did the Gladioli (6 Log_{10} CFU/cm²) surface when compared to the Tender heart surface (P=0.012), in relation to bacteria adhered. In summary, the White cabbage surface displayed the highest number of *L. monocytogenes* cells adhered, while the Gladioli surface displayed the least (Figure 4).

Retention assays

No significant difference could be found when comparing cells retained. The Leek (5 Log_{10} CFU/cm²) surface retained the highest number of *L. monocytogenes* cells, followed by the Tenderheart (4.5 Log_{10} CFU/cm²) and Flat (4.5 Log_{10} CFU/cm²) surfaces. The Gladioli (3.7 Log_{10} CFU/cm²) surface displayed the lowest number of retained *L. monocytogenes* cells, followed by the White cabbage (4.1 Log_{10} CFU/cm²) and Cauliflower (4.3 Log_{10} CFU/cm²) surfaces. In summary, the Leek surface displayed the highest number of cells retained while the Gladioli surface displayed the lowest (Figure 4).



Figure 4 *L. monocytogenes* attachment, adhesion and retention of biomimetic surfaces presented as Log_{10} CFU/cm².

Microbial Co-cultures

Isolation of CFU/cm² of each species of bacteria was obtained from a 1:1 culture using two varieties of selective growth media.

E. coli in co-culture

E. coli was isolated from co-culture using MacConkey agar.

Attachment assays

The White cabbage (5.5 Log^{10} CFU/cm²) surface was found to be significant when compared to the Tender heart (4.5 Log^{10} CFU/cm²) (P=0.046), Leek (5.1 Log^{10} CFU/cm²) (P=0.046) and Gladioli (6.2 Log^{10} CFU/cm²) (P=0.046) surfaces in relation to *E. coli* cells attached in co-culture. The Gladioli surface displayed the most *E. coli* cells attached followed by the White cabbage and Cauliflower (5.4 Log^{10} CFU/cm²) surfaces. The Tenderheart surface displayed the least amount of *E. coli* cells attached, followed by the Leek and Flat (5.2 Log^{10} CFU/cm²) surfaces. In summary, the Gladioli surface displayed the lowest (Figure 5).

Adhesion assays

When compared to the Leek (6.2 Log^{10} CFU/cm²) surface, the Tender heart (5.1 Log^{10} CFU/cm²) (P=0.049) surface showed a significant difference between the number of cells adhered. The Leek surface displayed the highest number of *E. coli* cells adhered in co-culture, followed by the Gladioli (6.1 Log_{10} CFU/cm²) and Cauliflower (6 Log_{10} CFU/cm²) surfaces. The Tenderheart surface displayed the lowest number of *E. coli* cells adhered in co-culture followed by the White cabbage (5.6 Log^{10} CFU/cm²) and Flat (5.9 Log^{10} CFU/cm²). In summary, the Leek surface displayed the highest number of cells adhered, while the Tenderheart surface showed the least (Figure 5).
Retention assays

When compared to the Tender heart (3.4 Log^{10} CFU/cm²) surface, the White cabbage (4.7 Log^{10} CFU/cm²) (P=0.049) and Flat (4.8 Log^{10} CFU/cm²) (P=0.030) surfaces showed a significant difference in the number of cells in relation *to E. coli* retention within the coculture. Significant differences were also observed when comparing the Cauliflower (4.61 Log^{10} CFU/cm²) surface to the Gladioli (6.18 Log^{10} CFU/cm²) (P=0.033) surface. The Gladioli surface displayed the highest number of *E. coli* cells retained, followed by the Flat and White cabbage surfaces. The Tenderheart surface displayed the lowest number of *E. coli* cells retained, follower surfaces. In summary, the Gladioli surface displayed the highest number of cells retained while the Tenderheart surface displayed the lowest (Figure 5).



Figure 5 *E. coli* co-culture attachment, adhesion and retention of biomimetic surfaces presented as Log^{10} CFU/cm².

L. monocytogenes in co-culture

L. monocytogenes was isolated from co-culture using Oxford growth medium.

Attachment assays

In relation to bacterial attachment, White cabbage (6.2 Log_{10} CFU/cm²) showed a significant difference when compared to Tenderheart (5.1 Log_{10} CFU/cm²) (P=0.043) and Gladioli (6.3 Log_{10} CFU/cm²) (P=0.046). The Gladioli surface displayed the highest number of *L. monocytogenes* cells attached, followed by the White cabbage surface. The Tenderheart surface presented with the lowest number of *L. monocytogenes* cells attached followed by the leek (5.5 Log_{10} CFU/cm²), Cauliflower (5.7 Log_{10} CFU/cm²) and Flat (5.8 Log_{10} CFU/cm²) surfaces. In summary, the Gladioli surface displayed the highest number of cells attached while the Tenderheart surface displayed the least (Figure 6).

Adhesion assays

No significant difference was found when comparing *L. monocytogenes* cell adhesion in co-culture. The Flat (6.6 Log_{10} CFU/cm²) surface displayed the highest number of *L. monocytogenes* cells adhered, followed by the Leek (6.6 Log_{10} CFU/cm²), Gladioli (6.6 Log_{10} CFU/cm²) and Cauliflower (6.5 Log_{10} CFU/cm²) surfaces. The Tenderheart (6.1 Log_{10} CFU/cm²) surface displayed the lowest number of cells adhered, followed by the White cabbage (6.3 Log_{10} CFU/cm²) surface (Figure 6). In summary, the Flat surface displayed the highest number of cells adhered, while the Tenderheart surface displayed the least.

Retention assays

No significant difference could be found when comparing all surfaces concerning bacterial retention with *L. monocytogenes* in co-culture. The Gladioli (5.1 Log_{10} CFU/cm²) surface displayed the highest number of *L. monocytogenes* cells retained, followed by the Flat (4.8

 Log_{10} CFU/cm²) and Cauliflower (4.6 Log_{10} CFU/cm²) surfaces. The Tenderheart 3.9 Log_{10} CFU/cm²) surface showed the lowest number of *L. monocytogenes* cells retained, followed by the Leek (4.1 Log_{10} CFU/cm²) and White cabbage (4.3 Log_{10} CFU/cm²) surfaces. In summary, the Gladioli surface displayed the highest number of cells retained, while the Tenderheart surface displayed the least (Figure 6).



Figure 6 *L. monocytogenes* co-culture attachment, adhesion and retention of biomimetic surfaces presented as Log_{10} CFU/cm².

SEM analysis

Surfaces were analysed with attachment, adhesion and retention assays to determine the effect varying topographies has on pre-biofilm surface colonisation; SEM was used to determine cells /cm².

E. coli in monoculture SEM analysis

Attachment assays

None of the results compared in relation to the numbers of bacteria observed following SEM of the bacterial attachment showed a significant difference. The Leek (5.9 Log₁₀ cells /cm²) surface presented with the highest number of *E. coli* cells attached, followed by the Tenderheart (5.7 Log₁₀ cells /cm²), Gladioli (5.7 Log₁₀ cells /cm²) and Flat (5.6 Log₁₀ cells /cm²) surface. The White cabbage (4.7 Log₁₀ cells /cm²) surface presented with the lowest number of *E. coli* cells attached, followed by the Cauliflower (5.5 Log₁₀ cells /cm²) surface. In summary, the Leek surface displayed the highest number of cells attached, while the White cabbage surface displayed the least (Figure 7).

Adhesion assays

None of the results compared in relation to bacterial adhesion showed any significant difference. The Flat (6.3 Log_{10} cells /cm²) surface displayed the highest number of *E. coli* cells adhered, followed by the Gladioli (6.1 Log_{10} cells /cm²), White cabbage (6.0 Log_{10} cells /cm²) and Tenderheart surfaces (5.9 Log_{10} cells /cm²). The Cauliflower (5.4 Log_{10} cells /cm²) surface displayed the least amount of *E .coli* cells adhered closely followed by the Leek (5.7 Log_{10} cells /cm²) surface. In summary, the Flat surface displayed the Highest number of cells (Figure 7).

Retention assays

Concerning *E. coli* retention, the comparison of the Tenderheart (4.7 Log_{10} cells /cm²) and Leek (0 Log_{10} cells /cm²) (P=0.000) surfaces was found to be significant. The Cauliflower (4.8 Log_{10} cells /cm²) surface displayed the highest number of *E. coli* cells retained, followed by the White cabbage (4.8 Log_{10} cells /cm²) and Tenderheart surfaces. The Leek surface displayed the lowest number of *E. coli* cells, followed by the Gladioli (4.5 Log_{10} cells /cm²) and Flat (4.5 Log_{10} cells /cm²) surfaces. In summary, the Cauliflower displayed the highest number of cells retained, while the Leek displayed the lowest (Figure 7).



Figure 7 *E. coli* monoculture attachment, adhesion and retention scanning electron enumeration analysis presented as Log₁₀ CFU/cm².

L. monocytogenes in monoculture SEM analysis

Attachment assays

A significant difference was found when comparing the Leek (6.1 Log_{10} cells /cm²) and Gladioli (6.8 Log_{10} cells /cm²) surface to the Flat (5.6 Log_{10} cells /cm²) (P=0.029), (P=0.025) surface. The Gladioli surface displayed the highest number of *L. monocytogenes* cells after an attachment assay, followed by the Cauliflower (6.5 Log_{10} cells /cm²), Tenderheart (6.4 Log_{10} cells /cm²) and Leek surfaces. The Flat surface displayed the lowest number of *L. monocytogenes* cells attached, followed by the White cabbage (5.9 Log_{10} cells /cm²) and Leek surface displayed the highest number of *cells* surfaces. In summary, the Gladioli surface displayed the highest number of cells attached, while the Flat surface displayed the lowest (Figure 8).

Adhesion assays

When comparing bacterial adhesion none of the surfaces showed a significant difference in the number of bacteria adhered to the different surfaces. The Gladioli (6.8 Log₁₀ cells /cm²) surface displayed the highest number of *L. monocytogenes* cells present after an adhesion assay, followed by the Flat (6.7 Log₁₀ cells /cm²), White cabbage (6.7 Log₁₀ cells /cm²), Tenderheart (6.6 Log₁₀ cells /cm²) and Cauliflower (6.5 Log₁₀ cells /cm²) surfaces. The Leek (6.3 Log₁₀ cells /cm²) surface presented with the lowest number of *L. monocytogenes* cells present. In summary, the Gladioli surface displayed the highest number of cells adhered, while the Leek surfaces displayed the lowest (Figure 8).

Retention assays

A significant difference was found when comparing the numbers of bacteria retained on the Tenderheart (5.9 Log_{10} cells /cm²) surface to the White cabbage (5.2 Log_{10} cells /cm²) (P=0.023) surface. The Cauliflower (6.1 Log_{10} cells /cm²) surface displayed the highest number of *L. monocytogenes* cells present after a retention assay, followed by the Tenderheart and Gladioli (5.8 Log_{10} cells /cm²) surfaces. The Flat (4.7 Log_{10} cells /cm²)

surface displayed the lowest number of *L. monocytogenes* cells present after a retention assay, followed by the White cabbage and Leek ($5.6 \text{ Log}_{10} \text{ cells / cm}^2$) surfaces. In summary, the Cauliflower surface displayed the highest number of cells retained, while the Flat surface displayed the lowest (Figure 8).



Figure 8 *L. monocytogenes* monoculture attachment, adhesion and retention scanning electron enumeration analysis presented as Log_{10} CFU/cm².

E. coli and L. monocytogenes co-culture SEM analysis

Attachment assays

A significant difference was found when comparing the White cabbage (4.1 Log_{10} cells /cm²) surface to the Gladioli (6 Log_{10} cells /cm²) (P=0.046) and Flat (4.9 Log_{10} cells /cm²) (P=0.046) surfaces. The Cauliflower surface displayed the highest number of bacterial cells attached, followed by the Gladioli, Tenderheart (5.6 Log_{10} cells /cm²) and Flat surfaces. The White cabbage surface displayed the lowest number of bacterial cells attached followed by the Leek (4.6 Log_{10} cells /cm²) surface. In summary, the Cauliflower surface displayed the lowest highest number of cells attached, while the White cabbage surface displayed the lowest (Figure 9).

Adhesion assays

No significant difference could be found when comparing bacterial adhesion of the cocultures. The Tenderheart (6.6 Log_{10} cells /cm²) surface displayed the highest number of bacterial cells adhered followed by the Leek (6.3 Log_{10} cells /cm²), Flat (6.2 Log_{10} cells /cm²) surface. The White cabbage (4.5 Log_{10} cells /cm²) surface displayed the lowest number of bacterial cells, followed by the Cauliflower (5.6 Log_{10} cells /cm²) and Gladioli (5.6 Log_{10} cells /cm²) surfaces. In summary, the Tenderheart surface displayed the highest number of cells adhered, while the White cabbage surface displayed the lowest (figure 9). *Retention assays*

No significant difference was found when comparing surfaces after the bacterial retention assays. The Flat (4.9 Log_{10} cells /cm²) surface displayed the highest number of bacterial cells retained. Followed by the Gladioli (4.5 Log_{10} cells /cm²) surface. The White cabbage (0 Log_{10} cells /cm²) surface displayed the lowest number of bacterial cells retained, followed by the Tenderheart (4.1 Log_{10} cells /cm²), Leek (4.2 Log_{10} cells /cm²) and Cauliflower (4.2

 Log_{10} cells /cm²) surfaces (Figure 9). In summary, the Flat surface displayed the highest number of cells retained, while the White cabbage displayed the lowest (Figure 9).



Figure 9 *E. coli* and *L. monocytogenes* monoculture attachment, adhesion and retention scanning electron enumeration analysis presented as Log_{10} CFU/cm².

SEM images

E. coli in monoculture SEM images

Attachment assays

After an attachment assay on the Tenderheart surface, *E. coli* cells appeared in small shallow surface recesses (~2 μ m). No cell clumping was observed (Figure 10a). On the Leek surface following an attachment assay, *E. coli* cells appeared clumped in relatively large groups in large surface recesses. Small groups of cells were also observed on smaller negative surface features (Figure10d). On the Cauliflower surface ensuing an attachment assay, cells were found to be mostly spaced arbitrarily on the surface; however, a small group of cells was observed in a shallow negative surface feature (Figure 10g). Following an attachment assay on the Gladioli surface cells were observed in large shallow ridges along the surface. A small group of cells was observed on top of a large protruding topographical structure (Figure 10j). On the White cabbage surface after an attachment assay, no cells were observed in the image (Figure m). On the Flat surface ensuing an attachment assay, no cells were observed in the image (Figure 10p).

Adhesion assays

On the Tenderheart surface after an adhesion assay, cells appeared to be grouped in small clumps predominantly on lower sections of the topography, however, a small clump of cells was also observed in a more elevated and exposed area of the topography (Figure 10b). Following an adhesion assay on the Leek surface, a small clump of cells was observed on the surface with no apparent connection to topography (Figure 10e). On the Cauliflower surface, after an adhesion assay cells appear to be placed on the sides of ridge structures on the surface, small clumping was observed (Figure 10h).

Ensuing an adhesion assay on the Gladioli surface, cells were observed in small clumps on the side of large protruding surface features, with small clumps of cells in large deep surface structures (Figure 10k). Following an adhesion assay on the White cabbage surface, cells appeared to show a clear affinity for deep surface features, with large clumps of cells present in them throughout. Small clumps of cells were also observed on the more exposed areas of the surface (Figure 10n). After an adhesion assay on the Flat surface, cells were observed in sheltered areas of the surface, with these areas presenting with the largest clumps present on the surface, however, an even coating of cells can be seen throughout the image (Figure 10q).

Retention assays

Following a retention assay on the Tenderheart surface, only one independent cell was observed on the surface, no large surface features were observed (Figure 10c). After a retention assay on the Leek, cauliflower and gladioli surfaces no cells were observed in the image (Figure 10f,i,l). Ensuing a retention assay on the White cabbage surface, two independent cells were observed in a large shallow negative surface feature (Figure 10o).). Following a retention assay on the Gladioli surface, no cells were observed in the image (Figure 10r).

In summary, an increased frequency of cells clumping was observed when cells were within proximity to larger negative and positive surface features like those present on the Gladioli and White cabbage surfaces. Smoother surfaces like the Flat surface showed cells predominately independent from one another. Surfaces analysed after attachment and adhesion assays generally displayed more cell clumping than surfaces after a retention assay (Figure 10).



a)

b)

c)

Figure 10 SEM E. coli Tenderheart attachment (a), adhesion (b) and retention (c)



Figure 11 SEM E. coli Leek attachment (a), adhesion (b) and retention (c)



Figure 12 SEM E. coli Cauliflower attachment (a), adhesion (b) and retention (c)



Figure 13 SEM E. coli Gladioli attachment (a), adhesion (b) and retention (c)







Figure 15 SEM *E. coli* Flat attachment (a), adhesion (b) and retention (c)

L. monocytogenes in monoculture SEM images

Attachment assays

After an attachment assay on the Tenderheart surface cells were observed clumped in multiple sheltered areas on the surface, with no cells present on more exposed areas (Figure 11a). Following an attachment assay on the Leek surface cells were observed to be arbitrarily spaces on the surface with surface topography having little to no visible influence (Figure 11d). Ensuing an attachment assay on the Cauliflower surface cells were found to be arranged in numerous large clumps on the side of surface ridges, with little cells in the lower grooves beneath (Figure 11g). After an attachment assay on the Gladioli surface cells were observed in large clumps showing an affinity for large positive topographical features, with fewer cells present on lower and smooth areas in the image (Figure 11j). Following an attachment assay on the White cabbage surface cell clumps were observed with a slight affinity shown for areas with a more varied microtopography (Figure 11m). Ensuing an attachment assay on the Flat surface very few cells were observed in the image, no large surface features were observed (Figure 11p).

Adhesion assays

After an adhesion assay on the Tenderheart surface, a large clump of cells was observed in a large negative topographical feature. A smaller clump of cells was seen on a less prominent feature. Little to no cells were observed in more exposed areas in the image (Figure 11b). Following an adhesion assay on the Leek surface cell clumping was observed throughout the image with cells showing little to no affinity for surface structures (Figure 11e). Ensuing an adhesion assay on the Cauliflower surface minimal cell clumping was observed, with cells showing an affinity to adhere to lower regions of the large surface feature present in the image (Figure 11h). After an adhesion assay on the Gladioli surface, small amounts of cell clumping was observed in small microtopography features and areas within proximity to larger surface features (Figure 11k). Following an adhesion assay on the White cabbage surface, cell clumping was observed, and cells appear arbitrarily spaced on the relatively smooth surface (Figure 11n). Ensuing an adhesion assay on the Flat surface large amounts of independent cells were observed with minimal clumping, no large surface features were observed (Figure 11q).

Retention assays

After a retention assay on the Tenderheart surface, multiple independent cells were observed in the image, with some showing an affinity for sheltered grooves to which the cells are aligned to fit within (Figure 11c). Following a retention assay on the Leek surface, multiple independent cells were observed with some aligned in surface grooves and some retained on the side of surface structures (Figure 11f). Ensuing a retention assay on the Cauliflower surface, large clumps of cells were observed in the more sheltered areas in the image, some independent cells were observed also (Figure 11i). After a retention assay on the Gladioli surface, large amounts of cell clumping was observed on the lateral faces of multiple surface features, with some small clumps of cells present on the flatter areas of the image and on the peaks of the large surface structures (Figure 11l). Following a retention assay on the white cabbage surface, small amounts of arbitrarily spaced independent cells were observed on the surface with no cell clumps present (Figure 11o). Ensuing a retention assay on the Flat surface, only one cell was observed in the image, no large surface features were observed (Figure 11r).

In summary, an increased frequency of cells clumping was observed when cells were within proximity to larger negative and positive surface features like those present on the Gladioli and White cabbage surfaces. Smoother surfaces like the Flat surface showed cells predominately independent from one another. Surfaces analysed after attachment and adhesion assays generally displayed more cell clumping than surfaces after a retention assay (Figure 11).



Figure 16 SEM *L.monocytogenes* Tenderheart attachment (a), adhesion (b) and retention (c)



Figure 17 SEM *L.monocytogenes* Leek (a), adhesion (b) and retention (c)



Figure 18 SEM *L.monocytogenes* Cauliflower attachment (a), adhesion (b) and retention (c)



Figure 19 SEM *L.monocytogenes* Gladioli attachment (a), adhesion (b) and retention (c)



Figure 20 SEM L.monocytogenes White cabbage attachment (a), adhesion (b) and retention (c)



Figure 21 SEM L.monocytogenes Flat (a), adhesion (b) and retention (c)

E. coli and L. monocytogenes co-culture SEM images

Attachment assay

After an attachment assay on the Tenderheart surface, cells were observed in small clumps arbitrarily spaced throughout the image; no large clumps of cells were observed (Figure 12a). Following an attachment assay on the Leek surface, no cells were observed in the image (Figure 12d). Ensuing an attachment assay on the Cauliflower surface cells were observed to be independent and showed an affinity of more exposed rougher areas of the topography (Figure 12g). After an attachment assay on the Gladioli surface clumps of cells were observed at the base of large surface features, with independent cells present in the flatter space in-between (Figure 12j). Following attachment assays on the White cabbage and Flat surfaces, no cells were observed in the image (Figure 12m,p).

Adhesion assays

After an adhesion assay on the Tenderheart surface, a high frequency of cells in small clumps was observed, with an affinity for more sheltered areas (Figure 12b). Following an adhesion assay on the Leek surface cells were observed to be independent of one another and arbitrarily spaced throughout the image (Figure 12e). Ensuing an adhesion assay on the Cauliflower surface, no cells were observed in the image (Figure 12h). After an adhesion assay on the Gladioli surface, only one cell was observed in the image, however, it was observed to be within the proximity of a large surface structure (figure 12k). Following an adhesion assay on the White cabbage surface, only one cell was observed in the image, however, it was observed to be in the deepest surface groove in the image (Figure12n). Ensuing an adhesion assay on the Flat surface, a large clump of cells was observed in the centre of the image, with large amounts of independent cells throughout. The Large clump appeared to be situated in a shallow negative surface feature (Figure 12q).

Retention assays

After a retention assay on the Tenderheart surface, no cells were observed in the image (Figure 12c). After a retention assay on the Leek surface no cell clumping was observed on the relatively flat surface (Figure 12f). Following retention assays on the Cauliflower, Gladioli, White cabbage and Flat surfaces no cells were observed in the image (Figure 12i,l,o,r).

In summary, an increased frequency of cells clumping was observed when cells were within proximity to larger negative and positive surface features such as those present on the Gladioli and white cabbage surfaces. Surfaces without large features such as the flat surface showed cells predominately independent from one another while having seemingly no impact on cell numbers. Surfaces analysed after attachment and adhesion assays generally displayed more cell clumping than surfaces after a retention assay (Figure 12).



Figure 22 SEM *E. coli* and *L.monocytogenes* Tenderheart (a), adhesion (b) and retention (c)



Figure 23 SEM E. coli and L.monocytogenes Leek (a), adhesion (b) and retention (c)



Figure 24 SEM E. coli and L.monocytogenes Cauliflower (a), adhesion (b) and retention (c)



Figure 25 SEM *E. coli* and *L.monocytogenes* Gladioli (a), adhesion (b) and retention (c)



Figure 26 SEM *E. coli* and *L.monocytogenes* White cabbage (a), adhesion (b) and retention (c)



Figure 27 SEM E. coli and L.monocytogenes Flat (a), adhesion (b) and retention (c)

Discussion

Due to the ever-approaching obsolescence of antibiotics and increased consumer attitude towards the environment and use of harsh chemical cleaning products within it, new alternative methods for reducing bacterial fouling in the food industry need to be established. Extensive research has been carried out of surface roughness and physicochemistry and if an ideal combination of factors can be established in relation to bacterial fouling the benefits will far exceed the food industry (Cheng *et al.*, 2019; Carniello *et al.*, 2018; Wu *et al.*, 2018; Song *et al.*, 2015). This work looked at different surface characteristics in the form of biomimetic plant leaf surfaces and how their structure and attributes affect the way bacteria begin the process of biofilm development. Bacterial attachment, adhesion and retention were analysed separately to ascertain how each distinct process is influenced by varying surface characteristics.

Physicochemistry effect on bacteria

Surface physicochemistry governs surface liquid interactions, dictating how liquids and bacterial cells behave on surfaces. Surface chemistry and topography are the primary influencers of surface physicochemistry (Giovambattista *et al.*, 2009). A trend between increased hydrophobicity and reduced levels of bacterial attachment, adhesion and retention of *E. coli* in co-culture after CFU/mL enumeration was established in agreement with (Fadeeva *et al.*, 2011) who showed a similar trend. Hydrophobic surfaces have been shown to reduce the contact area of the interface between cell and surface, thus decreasing the adhesion force present (Yuan *et al.*, 2017). Gibbs surface free energy ($\Delta Giwi$) equates to surface hydrophobicity which is calculated using the Van Oss-Chaudhury-Good equation of the interaction between the surface when immersed in water (Absolom *et al.*, 1983).

Free surface energy (γ_s) when compared to *E. coli* attachment and adhesion in monoculture and *L. monocytogenes* attachment and retention in monoculture exhibited a negative correlation. This is seen in related works (Zhang *et al.*, 2015) where a lower difference in surface and bacterial cell γ_s resulted in increased levels of bacterial adhesion. The connections between free surface energy and bacterial attachment, adhesion and retention are thought to be a result of interactions between the free surface energy of the bacterial cell, the free surface energy of the surface and the surface tension of the suspending medium (Bollen, 1995). The greater the difference in the surface energy between the surface and bacterial cell, the greater the adhesion force present (Pringle and Fletcher, 1983). A relatively low surface free energy has been shown to be the optimum level for the prevention of bacterial growth on surfaces (Pereni *et al.*, 2006). Surface energy is equal to the sum of γ_s^{AB} and γ_s^{LW} forces, whilst γ_s^{AB} is equal to the sum of the γ_s^- and γ_s^+ parameters (Chibowski, 2011). A trend was seen in increased *L. monocytogenes* adhesion co-culture assays after CFU/mL enumeration that correlated to an increased Lifshitz van der Waals (γ_s^{LW}). It has been suggested that when non-polar $\gamma_s{}^{\rm LW}$ forces are increased, bacterial attachment and adhesion will also increase (Carniello et al., 2018). The relationship between bacterial surface interaction and γ_s^{LW} is dependent on a multitude of factors, such as the chemical and physical properties of the bacteria, surface, and water solutions (Absolom et al., 1983). Polar surface energy is notably important concerning bacterial adhesion. A low γ_s^{AB} is thought to correlate to less bacterial adhesion (Yuan *et al.*, 2017). γ_s^{AB} forces are thought to increase with a decrease of separation to a surface, hydrostatic forces are thought to bridge this gap resulting in γ_s^{AB} being one of the primary forces of bacterial adhesion (Mao *et al.*, 2011). This was not seen when comparing bacterial attachment, adhesion and retention possible due to bacterial cell γ_s^{AB} repelling the surface (Ista and López, 2013). It has been shown that bacterial cells possess an affinity for surfaces with a high Lewis acid value, which is reliant on the ratio of γ_s^+ and γ_s^- resulting in a positive number (Fontaine *et al.*, 1996; Lee, 1996). This was seen in only the negatively charged Lewis base surface which presented the lowest values of attachment, adhesion and retention in E. coli and L. monocytogenes co-culture after CFU/mL enumeration.

Due to the surfaces displaying both γ_s^- and γ_s^+ values all surfaces were observed to be bipolar (Chibowski, 1992). Under typical, conditions bacterial cell walls are electron donors and can be repelled by surfaces that present a high γ_s^- . If a surface has a γ_s^+ of approximate to its γ_s^- value then only the net difference should be taken as the surface charge (Hamadi *et al.*, 2005). The Tender Heart surface presented as the only surface with a negative net charge, which displayed the lowest levels of attachment, adhesion and retention in *E. coli* and *L. monocytogenes* co-culture after CFU/mL enumeration.

Surface topography

Surface topography has been suggested to enhance already present physicochemical interactions on the surface (Marmur, 2003). Surfaces with increased roughness can present a contact angle that deviates substantially from that of a smoother surface (M. W. England *et al.*, 2016). This deviation can be a result of the surface being homogeneous to wetting, allowing all the surface features to make direct contact with water described with the Wenzel equation (Marmur, 2003). Alternatively, deviation in surface contact angle can be a result of surface heterogeneity, relying on the Cassie–Baxter model which accounts for diminished contact between surface and liquid due to the presence of air bubbles (Marmur, 2003).

Surface roughness and its effect on bacteria

Surface roughness is described as the deviation from the average line of the surface (BiolinScientific, 2019). A correlation between a high Arithmetical mean deviation (S_a) and low levels of bacterial retention in monoculture after CFU/mL enumeration was established. This is in disagreement with literature, which states that a reduced surface S_a will display a reduced level of bacterial colonisation (Dantas *et al.*, 2016). Retention is the first step in the process of biofilm formation that EPS production occurs (Jayathilake *et al.*, 2017). However, since surface S_a has been shown to not affect EPS production, this may suggest that external factors could influence the relationship between S_a and bacterial retention throughout all or some of the process, or that EPS production plays less of a role in bacterial retention than previously thought (Pizarro-Cerda and Cossart, 2006; Najafinobar, 2011). A trend was seen that agrees with other works (Dantas *et al.*, 2016). In relation to S_a and *L. monocytogenes* adhesion in monoculture after CFU/mL enumeration, the surface that

presented the highest S_a showed the highest levels of bacterial adhesion, although the surface with the lowest levels of *L. monocytogenes* after an adhesion assay also displayed a high S_a possibly indicating the influence of external factors such as S_q , S_{pv} or surface physicochemical properties (Marmur, 2003).

Increased root-mean-square deviation of surface topography (S_q) has been shown to decrease bacterial adhesion and inhibit biofilm formation in other works (Singh *et al.*, 2011). This was seen when bacterial retention of *E. coli* in monoculture after CFU/mL enumeration and bacterial retention in co-culture after SEM enumeration were compared to S_q . Little is known of the direct mechanisms that result in this correlation, it is possible that surface S_q is linked to surface physicochemistry values thus creating this effect (Marmur, 2003).

A negative correlation was seen when comparing a high surface S_q and S_a to $\gamma_s^{AB} \gamma_s^{LW}$, γ_s^+ and γ_s^- , this is possibly due to relatively high variations in surface topography distorting physicochemistry results by the trapping of air in micro and nano features on the surface (Marmur, 2003).

A trend was established between a high mean peak to valley height of surface topography (S_{pv}) and a high number of bacterial cells present after attachment and retention assays using *E. coli* and *L. monocytogenes* in monoculture. A surface with a high S_{pv} has a higher degree of variation from the S_a mean surface roughness line. This variation could be due to large protruding surface features or large surface depressions, in the case of the Gladioli surface, it was found to likely be both after analysis with SEM. Recent literature has shown that given that concave topography structures are of comparable size to bacterial cells, an increased number of bacterial cells could likely be due protection from shear forces and turbulent flow (Helbig *et al.*, 2016). The assays observed in this trend all included a rinse step that was applied resulting high enough levels of shear force to affect bacterial cells on

the surface. The approximate rinse shear force for the attachment assay was 70 kPa (Sharpe, 2011). The approximate rinse shear force for the retention assay was calculated to be 48 kPa ($p_d = 1/2 \rho v^2$). Both are values that far exceed the minimum levels of shear force needed to affect bacterial attachment and retention by a factor of tens of thousands (Nejadnik *et al.*, 2008).

Monoculture and co-culture bacterial assays

Results taken from monoculture enumerations were highly varied from each other in relation to the surface that displayed the highest and lowest number of bacterial cells present, whereas samples tested with co-culture presented relatively equivalent results when comparing *E. coli* and *L. monocytogenes* cells attached, adhered and retained, indicating that the relationship between the two species may be an influencing factor. This is likely a result of less experimental variation due to samples being taken from the same samples prepared in the same instance. It is thought that *L. monocytogenes* and *E. coli* do not have a significant effect on each other's growth rate while growing in co-culture (Mellefont *et al.*, 2008). Results from the co-culture assays after a CFU/mL enumeration appear to be approximately ten times lower than their monoculture counterparts. This showed a cell reduction of five times what was to be expected due to cell counts being halved when grown on selective agar. Indicating a competitive relationship between the two species of bacteria disagreeing with some works (Mellefont *et al.*, 2008). It is possible that while interacting with the surfaces stress was added to the bacteria resulting in the loss of an interspecific equilibrium (Khare and Tavazoie, 2015). This is also conferred by the larger gap between

the number of cells recovered and counted via SEM and CFU enumeration after a retention assay, likely due to the amount of time both species were able to interact with one another.

SEM enumeration

Cell enumeration results acquired by the analysis of SEM images differ greatly from results obtained via CFU/mL enumeration. Some SEM results directly contradict relevant literature such as bacterial adhesion analysis with co-culture (Yuan *et al.*, 2017). This could be due to potential reasons such as, all samples were treated with glutaraldehyde which is used as a fixative before SEM analysis, which has been shown to provide no increased protection from shear force (Kooten *et al.*, 1992). This would account for the reduced cell numbers seen throughout the SEM samples due to the method used to wash glutaraldehyde off the surface. Topographies with large surface structure have been shown to protect bacteria from shear force (Nejadnik *et al.*, 2008) explaining why a reduction of bacterial cells is not seen as prominently in surfaces with a raised S_{pv} .

SEM cell clumping

Surfaces with less topographical features and lower overall roughness displayed reduced amounts of cell clumping when compared to surfaces with large prominent surface structures. This was only demonstrated for surfaces that had been subject to a rinse step (attachment and retention assays). Surfaces that have not undergone a rinse step showed greater amounts of cell clumping. This is likely explained by the presence of large surface features which shield cells by aiding in the prevention of cell removal via shear force after rinsing (Nejadnik *et al.*, 2008). The effect of shear force on bacterial retention was not expected as it is thought that EPS provides a layer of protection from shear force to cells (Park *et al.*, 2011). Protection can be conferred from rinsing even in relatively small shallow surface features. This is likely due to the bacteria becoming flush with the surface topography thus reducing the shear force applied to the cells(Lazzini et al., 2019).

SEM negative control

The SEM images that had not been treated bacteria were not visualised since the bacteria were evident on the surface structures, and no advantage would have been gained from analysing these as the value would have been zero since they had been moulded using aseptic processing.

Conclusions

The results demonstrated that the Tenderheart leaf surface was the most hydrophobic with the highest surface free energy, highest γ_s^{AB} , most electron-donating and most electronaccepting surface. The Leek surface demonstrated the lowest surface free energy. The White cabbage surface was the most non-polar surface, with the least γ_s^{AB} properties, the least electron-accepting and least electron-donating surface. However, it had the highest S_a and S_q values. The Cauliflower leaf surface was the least hydrophobic and least nonpolar surface whilst the Gladioli surface was found to have the highest S_{pv} values. Finally, the flat surface showed the lowest S_a , S_q and S_{pv} values. Following the attachment, adhesion and retention assays, *E. coli* in monoculture did not show any trends between the surface properties and the number of cells retained. However, for *L. monocytogenes* in monoculture, following the attachment and retention assays, the Flat surface showed the least number of cells. Following the adhesion and retention assays, the Gladioli surface (highest S_{pv} values) displayed the lowest numbers of *L. monocytogenes* cells. Use of the bacteria in co-cultures demonstrated that for both the attachment and retention assays, the Tenderheart surface (most hydrophobic) displayed the lowest number of cells. SEM analysis did not correlate with the CFU/mL assays. However, for all the assays with *L. monocytogenes*, the flat surfaces (lowest roughness) retained the lowest numbers of cells, and for the co-culture, the White cabbage surface (most hydrophilic) displayed the lowest number of cells. These results demonstrate that when more topographically complex surfaces are analysed, the conclusions drawn between the effect of the surface properties on bacterial attachment, adhesion and retention from more uniform surfaces do not apply. Further, the processes of bacterial attachment, adhesion and retention are different and hence differentiation between these classifications needs to be clarified. It was clear that the different methods gave different results and that the use of different bacteria in monoculture and co-culture also affected the microbial assays. Hence, a new approach needs to be taken to understand the cell: surface interactions on complex surfaces.

Limitations of the study

Results taken from bacterial cell SEM enumeration were dissimilar to CFU/mL enumeration data. This may have been due to time constraints and only three SEM images were taken of each sample. The use of a handheld goniometer although convenient may have produced contact angle data that was slightly inaccurate. The Wenzel method although simpler does not account for air pockets present underneath the droplet. The application of a desktop goniometer would likely produce more reliable consistent results that would not have to be mapped by hand thus which was often the case thus increasing the chance of human error. The application of the more complex Cassie–Baxter model would also likely give more reliable results.

Future work

The work described in this study looks at the relationship between surface roughness and hydrophobicity compared to bacterial attachment, adhesion and retention. No information on cell grouping, density or cell spacing in co-culture was isolated from the results obtained. The application of multifractal analysis could prove invaluable in the understanding of the greater relationship between bacteria surface interactions (Wickens et al., 2014). Fractal analysis is a method of surface topography analysis that presumes surface topography is self-similar on all scales and that scale is described by a value termed the fractal dimension. Fractal analysis has the potential to bring together multiple values from varying surface scales into one quantifiable value. This would allow a holistic point of view to be attained and analysed in relation to surface topographical analysis. A multifractal is described as a fractal that cannot operate in one dimension and a continuous spectrum of dimensions are required for the object to exist (Brown and Scholz, 1985; Wickens et al., 2014). The combination of fractal analysis and 3D imaging software has the potential to describe cell clumping, dispersion and density data that can explain the relationship between individual topographical features and bacterial surface interactions at every stage of biofilm formation (Dazzo and Yanni, 2017).

To gain a more comprehensive understanding of all steps of biofilm development the last step in this process must be analysed. Once bacterial are retained on a surface cell proliferation occurs resulting in biofilm formation (Whitehead and Verran, 2015). A crystal violet assay would allow the quantification of biofilms on biomimetic surfaces in monoculture and co-culture (Peeters *et al.*, 2008). SEM analysis used in conjunction with this assay could give more comprehensive detailed results such as spacing and distribution data (Badha *et al.*, 2019). The application of confocal analysis would allow biofilm structure and architecture to be understood providing images that differentiate between cells in co-

culture displaying their intra-biofilm distribution and organisation (Reichhardt and Parsek, 2019).

Conditioning film analysis assays would prove insightful towards a more complete understanding of biofilm surface interactions. In nature biofilms do not form without there first being a conditioning film present, conditioning film composition directly affects how a biofilm will develop on a surface (Whitehead and Verran, 2015). Analysis with Fourier Transform InfraRed (FTIR) microscopy could give a detailed composition of conditioning films which could be used in conjunction with biofilm analysis (Humbert and Quiles, 2011). Cell surface hydrophobicity analysis could give a more complete understanding of surface physicochemistry by displaying the opposing side of interactions described by data obtained with a goniometer. Microbial adhesion to hydrocarbon (MATH) assay would be able to describe this data (Rosenberg, 2006).

Although a general knowledge of the chemical composition of the dental wax is known, a detailed understanding may prove to be highly insightful in relation to any influence chemicals present have on bacterial growth. Analysis with Fourier-transform infrared spectroscopy (FTIR) would provide this insight.

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