

**Effect of Estrogen on Host-Pathogen
Interactions in *ex vivo* and *in vitro*
Models of the Inflammatory Phase
of Age-Related Impaired Healing**

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Mohamed El Mohtadi

Department of Life Sciences
Manchester Metropolitan University

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List of Abbreviations

Aβ	Amyloid β -peptide
Arg-1	Arginase
ATP	Adenosine tri phosphate
BCA	Bicinchoninic acid
BCG	<i>Mycobacterium bovis bacillus Calmette-Guerin</i>
BHI	Brain heart infusion
BSA	Bovine serum albumin
CAMs	Cellular adhesion molecules
CF	Cystic fibrosis
CFU	Colony-forming units
CMP	Common myeloid progenitor
CMR	Common myeloid progenitor
CMoP	Committed monocyte progenitor
CLRs	C-type lectin receptors
C5b-9	Membrane attack complex
CR	Complement receptors
CRID	Rac-interacting and binding domain
c-SRC	p60 Src tyrosine kinase
DHT	Dihydrotestosterone
DHEA	Dehydroepiandrosterone
DHEA-S	Dehydroepiandrosterone sulphate
DN	Dominant negative
DPBS	Dulbecco's phosphate-buffered saline
DPN	Diarylpropionitrile
ECM	Extra cellular matrix
E-coli	<i>Escherichia coli</i>
EDTA	Ethylenediaminetetraacetic acid
EGTA	Ethylene glycol-bis(β -aminoethyl ether)-N,N,N',N'-tetraacetic acid
EGF	Epidermal growth factor
ER	Membrane-bound estrogen receptor
ER-α	Estrogen receptor alpha
ER-β	Estrogen receptor beta
ERE	Estrogen response element
ESC	Endometrial stromal cells
FAK	Focal adhesive kinase
FBS	Foetal bovine serum
FGF-2	Fibroblast growth factor 2
FITC	Fluorescein isothiocyanate
FL1-A	Fluorescence parameter 1
FL2-A	Fluorescence parameter 2
FSC	forward scattered light

FTIR	Fourier Transform Infra-Red
GAP-s	GTPase activating proteins
GCS-F	Granulocyte colony-stimulating factor
GDI_s	Guanine nucleotide dissociation inhibitors
GEFs	Guanine nucleotide exchange factors
GFP	Green fluorescent protein
GM-CSF	Granulocyte-macrophage colony-stimulating factor
GDP	Guanosine diphosphate
HBCCM	Human blood cell culture medium
HEPES	(4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid)
HPBM	Human peripheral blood monocytes
HSCs	Hematopoietic stem cells
HRT	Hormone replacement therapy
ICs	Immune complexes
IgG	Immunoglobulin G
IGF-1	Insulin-like growth factor-1
INF-γ	Interferon gamma
IL-1β	Interleukin-1 beta
IL-4	Interleukin-4
IL-13	Interleukin-13
IL-8	Interleukin-8
IL-17	Interleukin-17
IL-6	Interleukin-6
iNOS	Inducible NO synthase
KGF	Keratinocyte growth factor
<i>K. pneumonia</i>	<i>Klebsiella pneumoniae</i>
LPS	Lipopolysaccharide
LTA	Lipoteichoic acid
LR	Langerhans cells
M	Molar
MBL	Mannose binding lectin
MCS-F	Macrophage colony-stimulating factor
MCP-1	Macrophage chemoattractant protein 1
MDR	Macrophage and dendritic cells precursor
MIF	Migration inhibitory factor
MIP	Macrophage inflammatory protein
MFI	Median fluorescence intensity
MHC	Major histocompatibility complex
MMPs	Matrix metalloproteinases
mL	Millilitre
MOI	Multiplicity of Infection
MRSA	Methicillin-resistant <i>Staphylococcus aureus</i>
MRS11	Methicillin-resistant <i>Staphylococcus aureus</i> strain 11
NA	Nutrient agar

NaCl	Sodium chloride
NC	Negative control
NET	Neutrophil extracellular trap
NF-κB	Nuclear transcription factor kappa beta
ng	Nanograms
NGF	Nerve-growth factor
NHS	National Health Service
NLRs	Nucleotide-binding oligomerisation domain-like receptors
NO	Nitric oxide
PAK 1	p21-activated kinases
PAMPs	Pathogen-associated molecular patterns
PAO1	<i>Pseudomonas aeruginosa</i> strain
PDGF	Platelet-derived growth factor
PKC	Protein kinase-C
PMA	Phorbol 12-myristate 13-acetate
PMSF	Phenylmethylsulfonyl fluoride
PPT	Propyl pyrazole triol
PRRs	Pattern recognition receptors
PTK2	Protein tyrosine kinase 2
qPCR	Real-time polymerase chain reaction
RIPA	Radio immunoprecipitation assay
RLRs	RIG-1 like receptors
ROS	Reactive oxygen species
RPMI	Roswell Park Memorial Institute
rpm	Revolutions per minute
RNAseq	RNA sequencing
RT	Room temperature
SDS	Sodium dodecyl sulfate
SEM	Scanning electron microscopy
SERDs	Selective estrogen receptor degraders
SERMs	Selective estrogen receptor modulators
SSC	Side scattered light
TBS-tween	Tris-buffered saline tween
TGF β	Transforming growth factors β
THAM	Tris (hydroxymethyl) aminomethane hydrochloride
THC	T helper cells
Th1	Type 1 helper T-cell
Th2	Type 2 helper T-cell
Tris-HCl	Tris-hydrochloride
TLR4	Toll-like receptors 4
TLRs	Toll-like receptors
TNF-α	Tumour necrosis factor alpha
TPBM	Theophylline, 8-[(benzylthio)methyl]
VEGF	Vascular endothelial growth factor

WASP	Wiskott Aldrich Syndrome Protein
μg	Microgram
μm	Micrometre
μL	Microliters
γδ	Gamma delta
°C	Degree Celsius

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Abstract

Chronic wounds in the elderly often become infected, leading to substantial morbidity and mortality. Impaired healing in the elderly is mediated by age-related changes in steroid hormones, particularly declining levels of estrogen with increasing age. Although the anti-inflammatory activity of estrogen has been defined, very little is known about the effects of estrogen deprivation (ageing processes) on bacterial clearance. The aim of this study was to determine the effect of ageing (estrogen deprivation) on the ability of *in vitro* human U937-derived macrophages and *ex vivo* human peripheral blood monocyte (HPBM)-derived macrophages to eliminate bacteria via phagocytosis.

Host-pathogen assays were used to measure macrophage-mediated phagocytosis of two major wound pathogens, methicillin-resistant *Staphylococcus aureus* (MRSA) and *Pseudomonas aeruginosa*, under *in vitro* and *ex vivo* conditions that model estrogen levels in the elderly, young adults and following exogenous estrogen supplementation. Epifluorescence, confocal and scanning electron microscopy were used to visualise host-pathogen interactions and protein mediators of phagocytosis were measured by immunoblotting. Estrogen at concentrations typical of youth or supraphysiological levels significantly ($P < 0.05$) increased the phagocytosis and effective killing of MRSA and *P. aeruginosa* in a dose-dependent manner compared to estrogen deprivation with significantly enhanced clearance of bacteria by M1 macrophages compared to M2 or M0 macrophages. Epifluorescence, confocal and scanning electron microscopy confirmed estrogen increases co-localisation of fluorescent GFP-*S. aureus* or mCherry-*P. aeruginosa* within macrophages and promotes bacterial internalisation. Activation of estrogen receptor (ER)-alpha (ER- α) mirrored the stimulatory effect of estrogen on phagocytosis whilst ER- α antagonism significantly ($n=6$; $P < 0.05$) blocked the phagocytic effect of estrogen. In contrast, activation of ER-beta (ER- β) had no significant ($n=6$; $P > 0.05$) effect on phagocytosis, confirming estrogen mediates bacterial clearance via specifically through ER- α . Immunoblotting analysis demonstrated that enhanced phagocytosis by estrogen is associated with altered levels of mediators involved in the actin

cytoskeleton of phagocytes including increased levels of FAK, Rac1, Cdc42 and RhoG, but reduced levels of RhoA.

Collectively the findings suggest estrogen may promote the resolution of wound bacterial infections during youth but this protection is lost as estrogen levels decline with increasing age, resulting in increased propensity and progression of wound infections in the elderly. Thus, novel wound dressings that provide local estrogen supplementation or selective activation of ER- α and/or specific targeting of downstream mediators of the actin cytoskeleton may provide effective treatment options for infected wounds in the elderly.

Declaration and Copyright Statements

Declaration

No portion of the work referred to in this thesis has been submitted in support of an application for another degree or qualification of this or any other university or institute of learning.

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This thesis is being submitted in partial fulfilment of the requirements for the degree of PhD.

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The author has successfully published the following manuscripts during the PhD research course at MMU:

Nicola R. Sproston, **Mohamed El Mohtadi**, Mark Slevin, William Gilmore and Jason J. Ashworth; The Effect of C-Reactive Protein Isoforms on Nitric Oxide Production by U937 Monocytes/Macrophages. *Frontiers in Immunology*. (2018) 9:1500.

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Samuel J. Davidson, Lisa I. Pilkington, Nina C. Dempsey-Hibbert, **Mohamed El Mohtadi**, Shiyang Tang, Thomas Wainwright, Kathryn A. Whitehead and David Barker. Molecular Synthesis and Biological Investigation of 5-Hydroxymethyl Dibenzyl Butyrolactones and Related Lignans. *Molecules* 2018, 23, 3057.

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Finally, to my sister Farah, It is really difficult to find what to write in this paragraph as I am struggling to find the strong words to express what I really want to say. You have been my sister, my mum, and my friend during the last few years. You believed in me from day one, and gave me everything I needed to succeed. Everything I have achieved in this PhD and in all my life is due to God and to you in the first place. You have always been a model to follow and a source of inspiration to me. I really hope you will be proud of your brother when you read this thesis. I love you.

Dedication

I dedicate chapter 6 in my thesis to Professor Kathryn Whitehead and Dr Nina Dempsey.

To my Almighty God, to all my family and close friends, to my lovely mum and dad, to Mina and Thabet, to everyone who believed in me, especially my sister Farah, this thesis is for you....

Chapter 1: Introduction

1.1 Acute Wound Healing

Acute wound healing is a complex and dynamic biological process divided into four sequential, overlapping phases; haemostasis, inflammation, tissue proliferation and remodelling of the tissue scar (Figure 1.1). Immediately after trauma, healing initiates with haemostasis (coagulation). This phase is characterised by the formation of a fibrin clot within minutes following injury to prevent further blood loss and provide a temporary, protective barrier over the wound. The inflammatory phase occurs within minutes after injury, with neutrophils being the first inflammatory cells recruited from circulation, followed by the recruitment of monocytes. The primary objective of inflammation is to localise and eradicate the causative irritant or stimuli (e.g. bacteria). Neutrophils peak in numbers at 24 to 36 hours post wounding whereas monocytes peak at around 5 to 7 days after injury. Monocytes undergo a series of changes to differentiate into tissue macrophages. Macrophages carry out phagocytosis and release cytokines that encourage the recruitment and activation of leukocytes at the injury site. Upon the clearance of apoptotic cells, macrophages undergo a phenotypic change from a pro-inflammatory to a reparative state that encourages the initiation of the proliferation phase (Mosser and Edwards, 2010).

Three to ten days after injury, the proliferation phase starts enabling granulation tissue formation, re-epithelialisation and angiogenesis. Proliferation is characterised by the creation of a new extracellular matrix (ECM) by fibroblasts, angiogenesis by endothelial cells and re-epithelialisation by keratinocytes. The final phase is remodelling of a mature tissue scar, which can take several months or, in some cases, up to a year post-injury. It is characterised by the remodelling of collagen and the vascular maturation of newly formed capillaries, allowing vascular density to return to normal within the wound (Guo and DiPietro, 2010).

For successful healing, wound repair requires progression through all four phases in the correct order and timeframe (Singer and Clark, 1999; Guo and DiPietro, 2010).

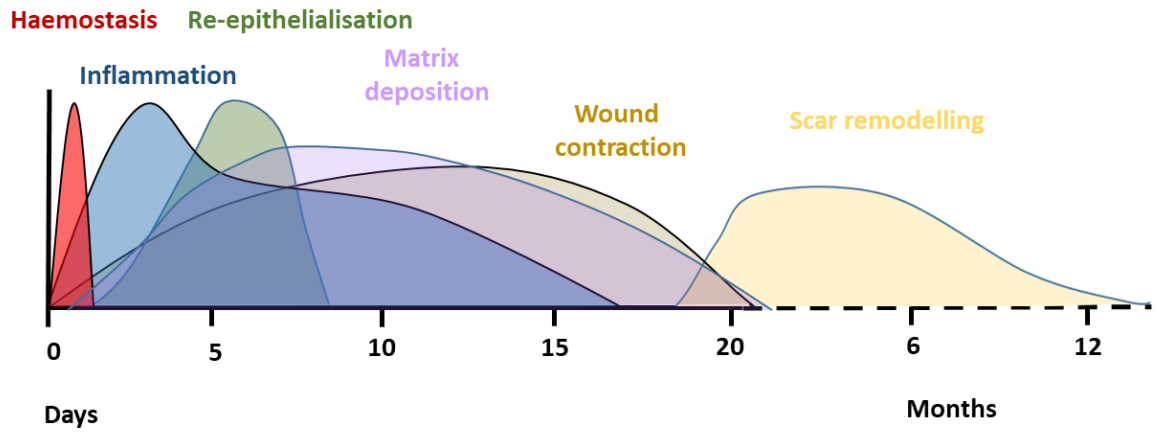


Figure 1.1 Diagrammatic illustration of typical timescale and phases of acute wound healing stages. 0 = day of wounding/injury. Figure drawn based on information in Gosain and DiPietro (2004) and Olczyk *et al.* (2014).

1.1.1 Haemostasis

Immediately after injury, degranulating platelets adhere to damaged blood vessels and start a haemostatic reaction, increasing the coagulation cascade to prevent extreme blood loss and to provide temporary protection for the wound against foreign bodies (Figure 1.2). This is accomplished by vasoconstriction and the creation of a platelet plug (Vaughan *et al.*, 2000; Weyrich and Zimmerman, 2004; Gilliver *et al.*, 2007). When the platelets come into contact with exposed collagen, they become adherent and activated, producing a platelet plug at the wound site. This involves a coagulation cascade in which fibrinogen is transformed to fibrin by thrombin, creating a clot of insoluble fibres bound to platelets (Fang *et al.*, 2005). The clot forms a provisional matrix rich in fibronectin that acts as a scaffold over which cells can migrate during the wound healing process. Platelets in the clot release a variety of pro-inflammatory cytokines and growth factors including platelet-derived growth factor (PDGF), transforming growth factors β (TGF- β), fibroblast growth factor 2 (FGF-2), vascular endothelial growth factor (VEGF) and epidermal growth factor (EGF) (Bauer *et al.*, 1985; Guo and DiPietro, 2010). These factors induce the recruitment of inflammatory cells from circulation and cells involved in the proliferative phase of wound repair including keratinocytes, endothelial cells and fibroblasts (Singer and Clark, 1999; Guo and DiPietro, 2010).

Platelet-fibrin clot forms to protect against excessive blood loss and/or foreign wound invaders

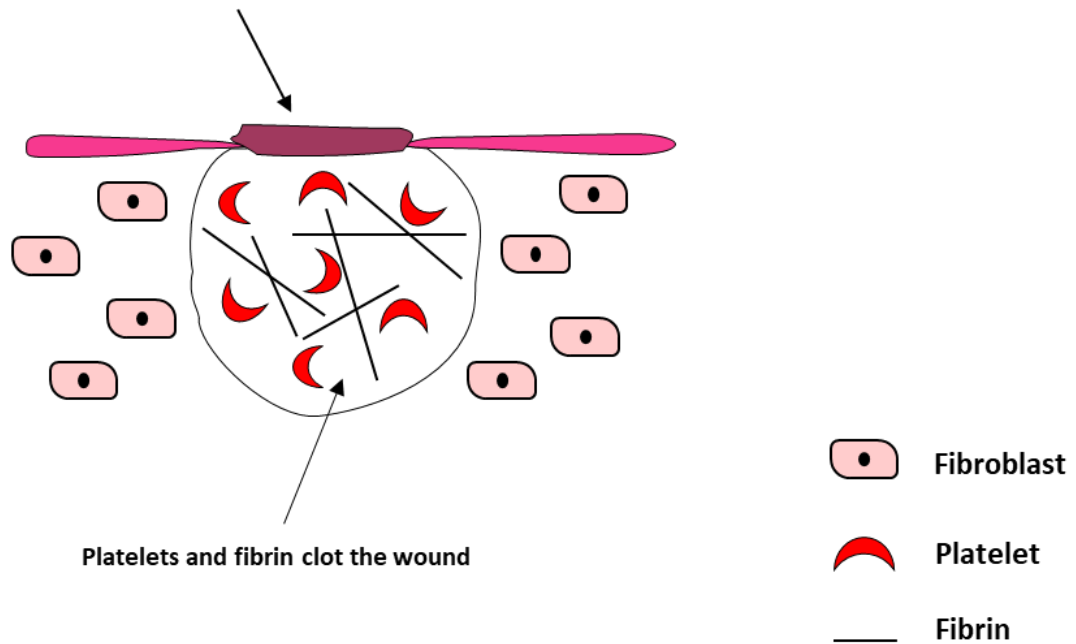


Figure 1.2. Schematic illustration of haemostasis in acute wound healing. Following an injury, platelets adhere to damaged blood vessels and start a haemostatic reaction, increasing the coagulation cascade to prevent extreme blood loss. A platelet plug forms that provides temporary protection against colonisation of the wound by microorganisms. The provisional fibrin matrix also provides a scaffold over which cells can migrate during the wound healing process. Figure drawn based on information in Singer and Clark (1999) and Guo and DiPietro (2010).

1.1.2 Inflammatory Phase

Following injury, the release of cytokines and chemokines by platelets and resident immune cells attracts inflammatory cells from circulation to the wound site (Figure 1.3). Large numbers of circulating neutrophils are recruited to the wound site by chemokine and cytokine expression of endothelial surface cellular adhesion molecules (CAMs), such as CD11 and CD18 (Ley *et al.*, 2007). Neutrophils are the first immune cells to arrive at the wound site within minutes of injury. They peak in numbers at 24 to 36 hours post-injury (Mosser and Edwards, 2010). Neutrophils start to remove foreign substances and invading microorganisms, such as bacteria, via the release of reactive oxygen species (ROS) and lysosomal enzymes, and degrade damaged matrix tissues by collagenases and proteinases (Singer and Clark, 1999). The majority of neutrophils are enclosed in the wound clot and are either eliminated with the eschar or by macrophages via phagocytosis (Newman *et al.*,

1982). In response to chemoattractants such as TGF- β , macrophage chemoattractant protein 1 (MCP-1), and macrophage inflammatory protein (MIP), monocytes from the bloodstream subsequently arrive at the wound area and differentiate into tissue macrophages, peaking in number around days 5 to day 7 post-injury (Lorenz and Longaker, 2003; Sen and Roy, 2008). Growth factors mainly TGF- β and residues of the remaining extra cellular matrix (ECM) including collagen, elastin, fibronectin, and thrombin induce circulating monocytes to adhere to the endothelium of blood vessels and migrate into the tissue. Macrophages replace neutrophils as the predominant inflammatory cells at the wound site and carry out the process of phagocytosis of invading microorganisms, removal of damaged tissues and dead neutrophils and the release of growth factors such as PDGF and TGF- β (Beanes *et al.*, 2003). Damaged extracellular matrix is degraded by the action of macrophage-derived proteolytic enzymes such as metalloproteases. Macrophages stimulate re-epithelialisation by releasing insulin-like growth factor-1 (IGF-1), keratinocyte growth factor (KGF) and EGF (Chen *et al.*, 2004) and angiogenesis via the secretion of VEGF (Shaw *et al.*, 1990). During wound clearance, macrophages experience a balanced switch from pro-inflammatory (classically activated) M1 macrophages to anti-inflammatory (alternately activated) M2 macrophages (Mosser and Edwards, 2008).

Adaptive immunity plays a major role in regulating inflammation and improving wound healing (Falanga, 2005). The adaptive immune response is late, but specific, compared to the innate immune response. It has the capacity to memorise invaders/pathogens and induce prompt reactions to later immunological tasks. The key cells linked with adaptive immune responses are T cells and B cells (Jameson *et al.*, 2002). They are initiated in reaction to antigens or cells presenting antigens. When activated, B cells produce specific antibodies that label pathogens for destruction, activate the complement system, ingest bacteria, and activate toxins. T cells are induced by the major histocompatibility complex (MHC) molecule that presents antigens. Antigens presented by MHC class I are detected by CD8+ cytotoxic T cells, whereas antigens presented by MHC class II are recognised by CD4+ helper T lymphocytes. CD8+ cells are more specialised in targeting cells than

CD4+ cells, which are more involved in cytokine production and paracrine signalling to adjacent cells. T cells are suggested to play a role in tissue repair (Jameson *et al.*, 2002). A specific group of T cells known as gamma delta ($\gamma\delta$) T cells, or dendritic epidermal T cells, are cells that are able to recognize antigens expressed by damaged and/or stressed keratinocytes (Jameson *et al.*, 2002). Research demonstrated that ($\gamma\delta$) T cells enhance re-epithelialisation via producing growth factors such as Keratinocyte growth factor-7 (KGF-7) and keratinocyte growth factor-10 (KGF-10). This has been reported in an animal study where $\gamma\delta$ T cell deficiency led to a delay in wound repair (Jameson *et al.*, 2002). CD4+ and CD8+ cells have also been shown to have an impact on the infiltration of neutrophils and macrophages to the wound site, which has an evident impact on the regulation of the inflammatory phase in wound healing. Furthermore, an induced deficit of CD4+ cells is linked with an increase in the levels of interferon gamma (IFN- γ), interleukin-1 beta (IL-1 β), interleukin-6 (IL-6), interleukin-17 (IL-17) and a diminution in interleukin-4 (IL-4). However, an induced deficit of CD8+ cells has the opposite effect on these inflammatory factors (Chen *et al.*, 2004). A balanced innate/adaptive immune reaction is essential for an appropriate inflammatory phase and wound healing.

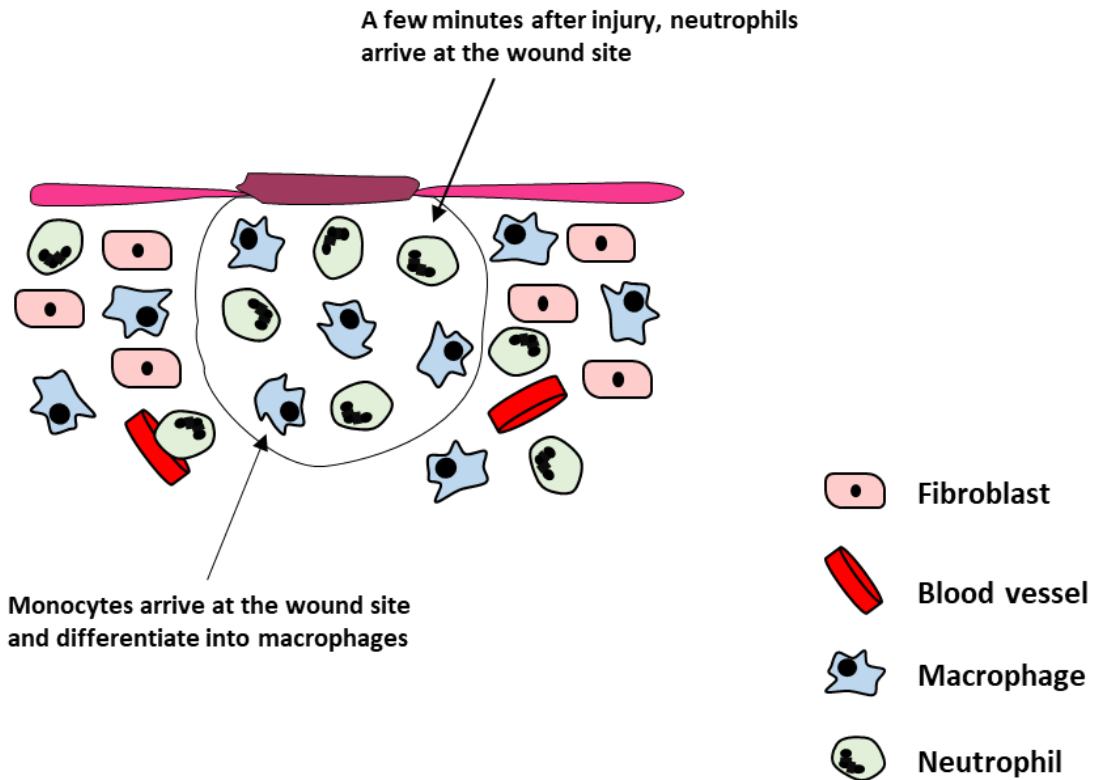


Figure 1.3. Representation of the inflammatory phase of acute wound healing. Cytokines released by platelets initially attract neutrophils and then monocytes to the wound site. Monocytes subsequently differentiate into tissue macrophages. Figure drawn based on information in Singer and Clark (1999) and Guo and DiPietro (2010).

1.1.3 Proliferative Phase

The third phase of the acute wound healing is called the proliferative phase. The major events during this phase are ECM formation, re-epithelialisation and angiogenesis (Kirsner and Eaglstein, 1993).

1.1.3.1 ECM Formation

Granulation tissue formation occurs approximately between 3 and 10 days post-injury. The temporary matrix produced during haemostasis is replaced with a more defined granulation tissue, consisting of newly formed blood vessels and various cell types including fibroblasts, macrophages and granulocytes within loose connective tissue (El Ghalbzouri *et al.*, 2004; El Ghalbzouri and Ponec, 2004). Fibroblasts exert several functions during wound healing processes such as wound contraction and ECM deposition. Fibroblasts are activated by PDGF and FGF-2, and

are encouraged to proliferate by insulin-like growth factor 1 (IGF-1) and TGF- β 1 (Singer and Clark, 1999). TGF β 1, released by macrophages during inflammation and induced by mechanical pressure in the ECM, directs fibroblasts to differentiate into contractile myofibroblasts (Gabbiani, 1999; Serini and Gabbiani, 1999). Fibroblasts attach together and to the extracellular matrix, producing stress fibres of contractile actin bundles that close the wound (Hinz, 2007). Fibroblasts synthesise new ECM through the generation of collagen, fibronectin, elastin, glycosaminoglycans, proteoglycans and hyaluronic acid (Hinz, 2007).

1.1.3.2 Re-epithelialisation

Re-epithelialisation is the process of restoring an intact epidermis after cutaneous injury, and it involves an orderly series of events including the detachment, migration and proliferation of adjacent epidermal keratinocytes across the wound, the differentiation of the neo-epithelium into a stratified epidermis, and the restoration of an intact epidermal barrier function (Li *et al.*, 2007). Keratinocytes at the wound borders proliferate to replace the cells that migrate across the temporary wound matrix (Li *et al.*, 2007). The main factors inducing keratinocyte migration are EGF, keratinocyte growth factor (KGF), Insulin-like growth factor 1 (IGF-1) and nerve-growth factor (NGF) (Marikovsky *et al.*, 1993; Tokumaru *et al.*, 2000). Growth factors that induce re-epithelialisation are released by activated platelets during haemostasis and by inflammatory cells, mainly macrophages, during inflammation (Li *et al.*, 2007). Keratinocytes upregulate the production of proteolytic enzymes including matrix metalloproteinases (MMPs) that contribute to create a path between the scab and tissue below. Migration of cells then stops causing a rearrangement of the actin cytoskeleton (Jacinto *et al.*, 2001; Mayor and Carmona-Fontaine, 2010).

1.1.3.3 Angiogenesis

Angiogenesis is the revascularisation procedure that re-establishes a blood supply to regenerate tissues following wounding. Angiogenesis is induced by growth factors including members of the TGF- β family, VEGF, FGF-2 and PDGF, which are released during haemostasis, and by inflammatory cells in response to injury

(Adams and Alitalo, 2007). Recruited endothelial cells release proteolytic enzymes that facilitate their entrance to the wound site, where they multiply and assemble into tubular canals, and secrete MMPs, leading to lysis of adjacent tissue. Endothelial cells increase in number and reinforce their vessel walls by recruiting smooth muscle cells and pericytes, reinstating blood flow to the injured tissue (Clark *et al.*, 1982).

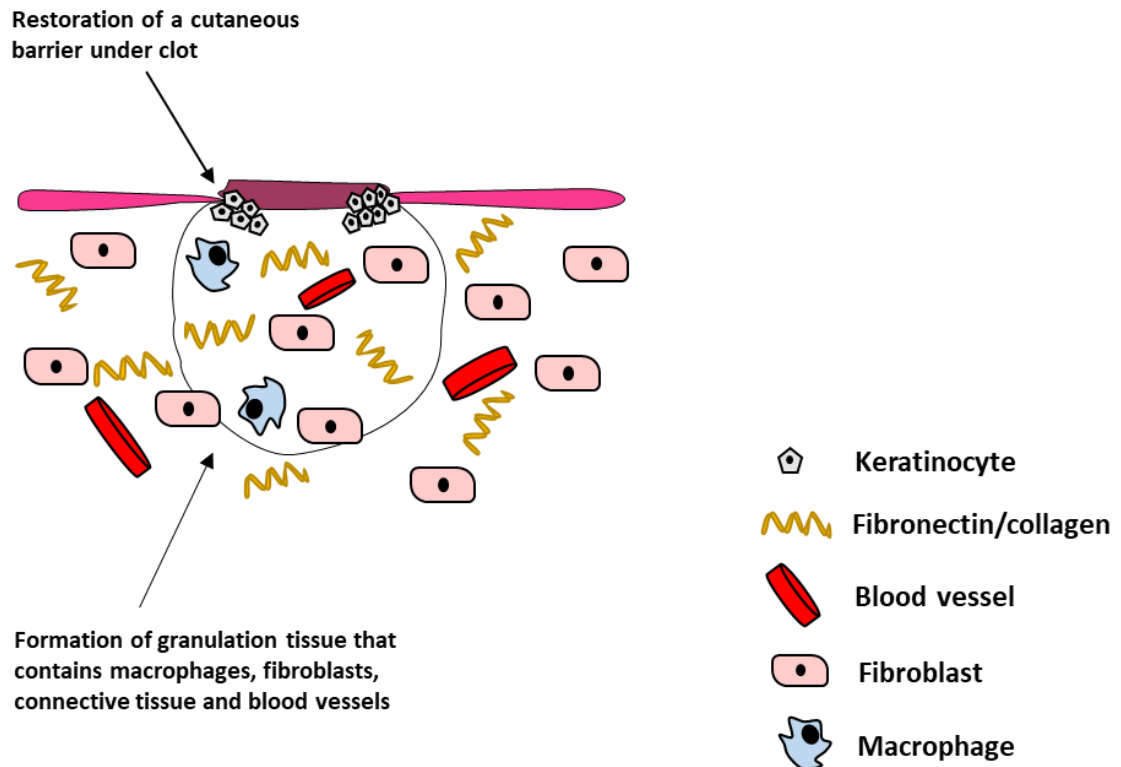


Figure 1.4. Schematic illustration of the proliferative phase in wound repair. As the inflammatory phase resolves, the overlapping proliferative phase includes the synthesis of granulation tissue that is characterised by blood vessel formation and contains recruited fibroblasts and macrophages. Keratinocytes at the wound edge migrate over the wound to restore the epidermis. Figure drawn based on information in Singer and Clark (1999) and Guo and DiPietro (2010).

1.1.4 Remodelling Phase

Remodelling is the last phase of wound repair that can endure for many months after wounding. It is characterised by a termination of the inflammatory response and the continuous remodelling of the ECM. The production/release of cytokines stops and inflammatory cells are removed from the wound area by apoptosis and/or phagocytosis (Serhan *et al.*, 2008). The new ECM is remodelled over several

months to generate a robust dermal structure in which collagen III is gradually replaced by collagen I leading to the formation of a mature tissue scar (Singer and Clark, 1999). Blood vessels also degenerate during the remodelling phase (Singer and Clark, 1999).

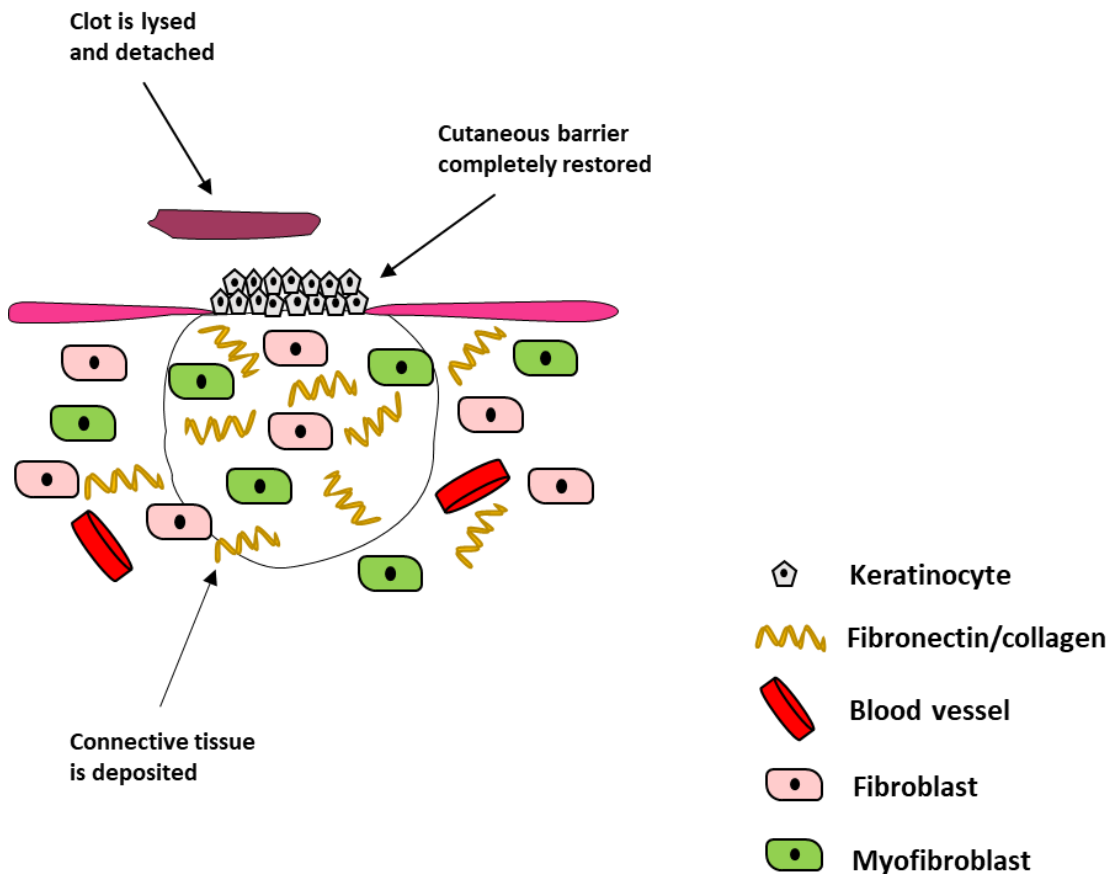


Figure 1.5. Schematic representation of the remodelling phase of wound healing. Blood vessels degenerate. The provisional extracellular matrix of fibronectin and fibrin is replaced with connective tissue, including collagen, and the formation of a mature scar. Figure drawn based on information in Singer and Clark (1999) and Guo and DiPietro (2010).

1.2 Abnormal Wound Healing

After an injury, wound healing is vital in re-establishing a new cutaneous barrier and preventing infection by invading foreign microorganisms. The speed and quality of wound repair are influenced by several factors, with age being a key factor that is known to markedly delay healing in the elderly (Ashcroft *et al.*, 2002).

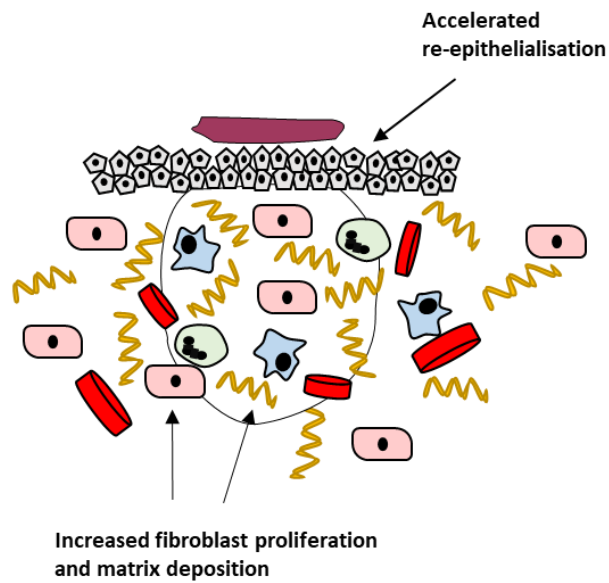
Pathological circumstances such as, diabetes, contribute directly to the development of chronic wounds, with foot ulcers being the most common cause of hospitalisation in diabetic patients. The treatments for chronic wounds are often

ineffective and place a substantial financial burden on the world's health organisations (Harding *et al.*, 2002; Boulton *et al.*, 2005).

1.2.1 Age-Related Impaired Healing

With increasing age, acute wound healing proceeds but becomes delayed. This detrimental change in acute wound healing in the elderly is called age-related impaired healing and is linked with an increase in skin fragility, reduced immune responses and cellular aging (Thomas, 2001). Delayed wound healing in the elderly is associated with delayed haemostasis (Ashcroft *et al.*, 1999b), prolonged and excessive inflammatory response, delayed re-epithelialisation, impaired angiogenesis, and reduced matrix deposition (Ashcroft *et al.*, 1997b; Ashcroft *et al.*, 2002). Although the inflammatory response becomes more pronounced with increasing age, the propensity for wound infections increases in the elderly (Ashcroft *et al.*, 2002; Cooper *et al.*, 2015), in part due to the delay in wound repair. Moreover, despite the pronounced inflammatory response, evidence suggests that increasing age may result in an impaired ability of inflammatory cells to eliminate bacteria from the wound site (Emori *et al.*, 1991; Thomas, 2001). It is suggested that an excessive presence of macrophages lead to an excessive matrix demolition due to the production of great amounts of nitric oxide (NO) and ROS which can delay wound healing and cause damage to the wound zone (Sen and Roy, 2008).

Young



Elderly

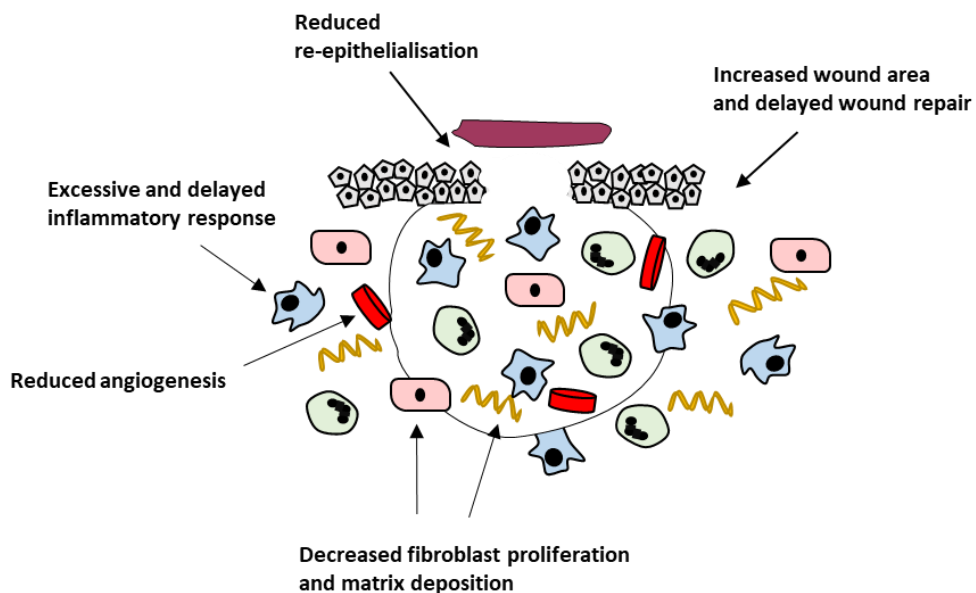


Figure 1.6. Schematic representation of the effect of age on wound healing. Age-related impaired healing is linked with delayed haemostasis, delayed and excessive inflammatory response and enhanced pro-inflammatory cytokine production, impaired platelet function and delayed re-epithelialisation, in addition to decreased fibroblast proliferation and matrix deposition. Figure drawn based on information in Gosain and DiPietro (2004).

1.2.2 Chronic Wounds

A chronic wound develops when a wound fails to proceed through an orderly set of wound healing phases within an expected timeframe. Chronic wounds can take

several months or even years to heal (if they heal at all) properly. Wounds are considered chronic if they do not heal within three months (Mustoe, 2005; Adeyi *et al.*, 2009). Chronic wounds are mainly linked to an underlying disease that is common in the elderly (over 65 years of age). Chronic wounds represent a major clinical issue that causes an enormous burden to healthcare services, demanding huge medical efforts and a substantial amount of healthcare funds (Harding *et al.*, 2002; Boulton *et al.*, 2005). Chronic wound treatment costs the UK National Health Service (NHS) about £5 billion per annum (Guest *et al.*, 2015). At present, effective therapies/treatments for chronic wounds are somewhat limited, making this an area of research that needs urgent attention. Chronic wounds are typically trapped within the inflammatory phase of wound repair and are characterised by an excessive, unabated inflammatory response that leads to tissue breakdown (Snyder, 2005; Taylor *et al.*, 2005; Fazli *et al.*, 2009). They are also characterised by a loss in the specific balance between the formation and degradation of the extracellular matrix (ECM), particularly collagen and elastin, with a gross shift to ECM destruction (Edwards *et al.*, 2004; Schönfelder *et al.*, 2005). Chronic wounds are characterised by increased pro-inflammatory cytokine production, decreased tissue growth factor secretion, reduced matrix production, delayed but excessive inflammation, decreased angiogenesis, delayed re-epithelialisation, bacterial infection and defective macrophage function (Hohn *et al.*, 1976; Harding *et al.*, 2002). There are several underlying pathologies associated with chronic wounds, with more than 90 % of chronic wounds being venous, diabetic or pressure ulcers (Boulton *et al.*, 2005). Moreover, it has been reported that diabetic foot ulcers and venous leg ulcers have an increased number of B cells and an inferior CD4+/CD8+ ratio in comparison with acute wounds, mainly due to a lower number of CD4+ cells (Moore *et al.*, 1997).



Figure 1.7. Patients with a chronic wound. A typical example of; (A) diabetic foot ulcer, (B) diabetic foot ulcer developing gangrene (Jeffcoate and Harding, 2003), (C) venous leg ulcer and (D) pressure foot sore (Grey *et al.*, 2006).

1.2.2.1 Venous Ulcers

Venous ulcers (Figure 1.7c) are most frequently observed in the elderly and represent approximately 70 % of chronic wounds of the lower leg (Baker *et al.*, 1991). Venous ulcers mainly develop in the gaiter area of the leg and signify a clinical manifestation of sustained venous hypertension, typically characterised by excessive vascularisation and unhealthy granulation tissue formation. At an advanced stage, venous ulcers can impair the function of the calf muscle (Cockett, 1955; Grey *et al.*, 2006).

Research has suggested the delay in healing and development of venous ulcers is connected with an inappropriately excessive inflammatory response (White and Ryjewski, 2005). It is thought that leukocytes recruited during the inflammatory phase are trapped in the microcirculation of the wounded leg (Moyses *et al.*, 1987) and release inflammatory mediators that lead to further recruitment and accumulation of leukocytes. Inflammatory cells release proteolytic enzymes that break down the dermal structure, leading to the development of an ulcer (Wysocki and Grinnell, 1990; Grinnell *et al.*, 1992).

1.2.2.2 Diabetic Ulcers

It has been reported that 15 % of diabetic patients in the UK are at risk of developing a chronic diabetic foot ulcer (Guo and DiPietro, 2010), and around 85 % of diabetic ulcer patients will require an amputation of their lower leg due to the development of gangrene (Figure 1.7b) (Apelqvist and Larsson, 2000; Boulton, 2004; Clinton and Carter, 2015). Diabetes is characterised by an alteration of blood sugar levels which can cause nerve damage leading to autonomic neuropathy and vasculopathy (microcirculation dysfunction) (Boulton, 2004). This can generate local hypertension and hypoxia, resulting in an ulcer due to tissue necrosis. A diabetic ulcer (Figure 1.7a) typically develops on feet and ankle regions that are repetitively exposed to mechanical pressures. Diabetic ulcers occur frequently in patients who develop diabetic neuropathy, because they are incapable of sensing and relieving cutaneous pressure and pain (Boulton *et al.*, 1983; Boulton, 2004). Like many chronic wounds, diabetic wounds exhibit impaired macrophage and neutrophil function, in turn increasing the risk of infection and prolonging healing time.

1.2.2.3 Pressure Ulcers

A pressure ulcer, also called a pressure sore (Figure 1.7d), is local damage of the skin and/or underlying tissues, resulting from pressure, or pressure combined with shear (Black *et al.*, 2007). Pressure ulcers are mostly hospital-acquired chronic wounds, occurring predominantly in the elderly and particularly in patients who remain stationary for a long period of time, such as paralysed or disabled patients (Allman *et al.*, 1995). Pressure ulcers are usually small ulcers caused by the

interruption of blood flow to the tissue by persistent pressure. The absence of oxygen at the pressure site results in necrosis of the tissue and the development of a pressure sore (Black *et al.*, 2007).

1.3 Wounds and Bacteria

1.3.1 Wound Infection

It is believed that low levels of bacteria in a wound have a beneficial impact on acute wound healing by stimulating the inflammatory response and improving granulation tissue formation (Robson, 1997). However, chronic wounds, that take months or years to heal, typically have microbes surviving and proliferating within the wound site. Research has indicated that all chronic wounds are subject to colonisation with a varied range of bacteria (Bowler and Davies, 1999). When the bacteria outweigh the host's immune system, the result is infection (Friedman and Su, 1984). It has been reported that the environment of chronic wounds is potentially favourable to bacterial infections, particularly when the wound tissue is ischaemic and dry (Rubinstein *et al.*, 1983; Whiston *et al.*, 1994).

1.3.2 Bacterial Species in Chronic Wounds

Numerous studies have demonstrated that the main bacterial species found in chronic wounds are *S. aureus*, *P. aeruginosa*, *S. epidermidis*, *Streptococcus spp*, *Enterococcus spp.* and coliform bacteria (Hansson *et al.*, 1995; Bowler, 1999; Gjødsbøl *et al.*, 2006; Kirketerp-Møller *et al.*, 2008; Rybtke *et al.*, 2015). Research has identified bacterial species present in infected surgical wounds of 676 patients. The most dominant pathogen was *S. aureus* (28.2 %), followed by *P. aeruginosa* (25.2 %), *E. coli* (7.8 %), *S. epidermidis* (7.1 %) and *Enterococcus faecalis* (5.6 %) (Giacometti *et al.*, 2000). The most frequent bacteria cultured from chronic wounds are illustrated in Table 1.1.

Table 1.1. Bacteria frequently associated with chronic wounds.

Information in table adapted from (Robson, 1997; Bowler and Davies, 1999).

Gram-positive		Gram-negative	
Aerobe	Anaerobe	Aerobe	Anaerobe
<i>Staphylococcus</i>	<i>Peptostreptococcus</i>	<i>Pseudomonas</i>	<i>Bacteriodes</i>
<i>Streptococcus</i>	<i>Clostridium</i>	<i>Escherichia</i>	<i>Prevotella</i>
<i>Enterococcus</i>		<i>Klebsiella</i>	<i>Fusobacterium</i>

It has been reported that chronic wounds are notably susceptible to *S. aureus* infections (Bowler and Davies, 1999; Beasley and Hirst, 2004). Along with anaerobic bacteria, *S. aureus* is known to be the most predominant bacteria found in chronic wounds, followed by *P. aeruginosa*, *Peptostreptococcus* spp. and *Bacteriodes* spp. (Louie *et al.*, 1976; Lookingbill *et al.*, 1978; Stephens *et al.*, 2003; Davies *et al.*, 2004).

Staphylococcus spp. and *Pseudomonas* spp. are the aerobic species most frequently detected in all wound types, with a great predominance in chronic wounds (Davies *et al.*, 2004).

1.3.2.1 *Staphylococcus aureus*

S. aureus is a facultative anaerobic Gram-positive cocci, typically 1 µm in diameter. It is commonly found on the skin, nose and in the respiratory tract (Stryjewski and Corey, 2014; Chambers, 2001). In normal healthy and immunocompetent people, *S. aureus* colonisation of the skin, intestinal tract, or nasopharynx is not pathogenic. However, it is considered the classic opportunist since it is frequently associated with wound infections and abscesses (Chambers, 2001; Missiakas and Schneewind, 2016).

Subpopulations of *S. aureus* comprise antibiotic resistant strains, such as methicillin-resistant *S. aureus* (MRSA). Penicillin was the antibiotic of choice to treat infections with *S. aureus*. After the acquisition of a resistance to penicillin, methicillin was introduced in 1959 as a treatment for penicillin-resistant *S. aureus* (Jevons, 1961; Jessen *et al.*, 1969). However, reports from the UK indicated the first case of MRSA just two years later (in 1961) and, during the period between the 1960s and the 1990s, hospital-acquired MRSA cases became increasingly common,

leading to a serious worldwide health problem (Enright *et al.*, 2002). MRSA is now recognised as a nosocomial pathogen (hospital acquired-MRSA) throughout the world (Diekema *et al.*, 2001). However, cases of MRSA in healthy, community-resident individuals have been recorded towards the end of the 1990s, with such infections referred to as community-acquired MRSA (Naimi *et al.*, 2001). MRSA has the ability to live and survive undetected in a host without provoking any symptoms (Stryjewski and Corey, 2014). Nevertheless, the pathogen can cause endocarditis, sepsis, and meningitis if it is introduced into the bloodstream, or if the host is incapable of developing a normal immune response (i.e. immunocompromised). Vancomycin was the only antibiotic used as a treatment for MRSA. However, resistance to this antibiotic has also been reported in recent years (Hiramatsu *et al.*, 1997; Cardona and Wilson, 2015).

MRSA is frequently linked with chronic wounds. Giacometti *et al.* (2000) reported that 54.4 % of *S. aureus* isolated from surgical wounds were discovered to be methicillin-resistant. The treatment of chronic wounds infected with MRSA represents a big challenge for the modern healthcare organisations (Beasley and Hirst, 2004). When chronic wounds are heavily infected with MRSA, or other hospital-acquired pathogens, treatment may necessitate aggressive medication with last line of defence antibacterial therapies (Beasley and Hirst, 2004).

1.3.2.2 *Pseudomonas aeruginosa*

P. aeruginosa is a Gram-negative bacillus, typically 3 µm × 0.5 µm in diameter. It is an opportunistic pathogen and a causative agent of a wide range of diseases in both immunocompromised and otherwise healthy patients (Passador *et al.*, 1993; Ryan and Ray, 2004). It is a typical-multidrug-resistant (MDR) organism known for its ubiquity, its fundamentally advanced antibiotic resistance mechanisms, and its link with serious diseases, particularly nosocomial infections such as sepsis (Cross *et al.*, 1983). It is also considered to be a serious danger to patients with impaired host defences, such as patients with burn wounds and cystic fibrosis (CF) (Hachem *et al.*, 2007).

P. aeruginosa has the ability to produce a biofilm in several environments (O'Toole and Kolter, 1998; Costerton *et al.*, 1999; Yoon *et al.*, 2002). Biofilms are surface-attached bacterial communities in which microbial aggregates are enclosed in a self-formed, extracellular polymeric matrix (Whitchurch *et al.*, 2002; Matsukawa and Greenberg, 2004; Fazli *et al.*, 2009). Biofilms have a specific complex architecture and structure with biochemical characteristics that lead to increased bacterial resistance against host immune mechanisms and antibiotics (O'Toole and Kolter, 1998; Costerton *et al.*, 1999; Mah *et al.*, 2003). Thus, it is usually difficult to treat such biofilms effectively with antibiotics since the biofilm structure appears to protect the bacteria from diverse environmental factors (Stover *et al.*, 2000). Biofilms of *P. aeruginosa* can be the cause of many chronic opportunistic infections in immunocompromised and elderly patients. It is the main cause of mortality and morbidity in CF patients, since it adapts, persists and produces biofilms in anaerobic CF environments, as opposed to the aerobic biofilms typically formed under laboratory conditions (Oliver *et al.*, 2000; Singh *et al.*, 2000).

P. aeruginosa are resistant to a wide range of antibiotics and has the ability to acquire additional resistance after failed treatments. Resistance of *P. aeruginosa* to many common first-line antibiotics, such as carbapenems, polymyxins, and more lately tigecycline, has been demonstrated (Strateva and Yordanov, 2009). However, such antibiotics are still in use in cases where resistance has not yet been determined. The antibiotic fluoroquinolone is one of the few antibiotics that are efficient against *P. aeruginosa*. In some hospitals, treatment with this drug is severely controlled to avoid the progress of resistant strains (Köhler *et al.*, 1997; Pesci *et al.*, 1999). Although *S. aureus* has been established as a main participant in chronic wounds, recent evidence has demonstrated that *P. aeruginosa* infects the deeper layers of the chronic wounds (Fazli *et al.*, 2011; Rybtke *et al.*, 2015).

1.3.3 Chronic Wounds and Biofilms

The growth of microorganisms in chronic wounds has been shown to take the form of irregularly shaped micro-colonies that can comprise hundreds of bacteria (Bjarnsholt *et al.*, 2008; Rybtke *et al.*, 2015).

It is becoming increasingly apparent that biofilms play an important role in the production and maintenance of the chronic wound environment (Davis *et al.*, 2008; James *et al.*, 2008a; Kirketerp-Møller *et al.*, 2008; Fazli *et al.*, 2009). Nevertheless, the role of biofilms in chronic wounds is still not entirely understood, but it is supposed that their presence could be one of many causes of non-healing wounds (Bjarnsholt *et al.*, 2005; Jensen *et al.*, 2007; Fazli *et al.*, 2009). Chronic impaired wounds are frequently colonised with biofilms of multiple bacterial species including *S. aureus* and *P. aeruginosa* (Clinton and Carter, 2015). Treatment of biofilm-infected wounds is extremely difficult due to the polymicrobial nature of these biofilms, and their high resistance to traditional antibiotic combinations (Clinton and Carter, 2015). Compared to normal wounds, chronic wounds are highly susceptible to colonisation with biofilms (James *et al.*, 2008b). This makes wound infection one of the main factors that delay wound healing in chronic wounds (James *et al.*, 2008b).

Antimicrobial genes of biofilms, composed of one or multiple bacterial species, can interact with each other leading to an increased virulence of the overall biofilm (James *et al.*, 2008b). Moreover, the metabolism of these multi-species biofilms can contribute to the development of antibacterial resistance, leading to auto-regeneration of the biofilm after treatment with antibiotics (Fauvert *et al.*, 2011). When established, biofilms become difficult to eradicate leading to a higher persistence of chronic wounds (James *et al.*, 2008b).

1.3.4 Bacterial Invasion of Host Tissues

Invading bacteria adhere to host cells, generally epithelial cells, and start colonising host tissues. Bacteria develop complex strategies to successfully invade and survive inside the host (Cossart and Sansonetti, 2004). The key mechanisms by which bacteria invade the host include adhesion to host cells, release of toxins causing tissue damage, and development of resistance against host antimicrobial defences, thus assuring effective growth and survival (Finlay and Falkow, 1989; Galan, 1994; Ofek and Doyle, 2012). Such resistance mechanisms are particularly important in establishing a persistent, chronic infection.

Bacteria bind to host cells using filamentous proteins named adhesins, such as pili and fimbriae. Lipopolysaccharide (LPS), M proteins, capsule, glycocalyx and lipoteichoic acids all act directly or indirectly as adhesins (Ofek and Doyle, 2012). It is known that the membrane surface of host cells and bacteria are negatively charged, leading them to repel each other (Gottenbos *et al.*, 2001). However, when bacteria use adhesins, their hydrophobicity changes, which is a key feature that allows them to adhere to host cells (Peschel, 2002). Bacteria release a range of molecules, such as enzymes and endotoxins, which cause host tissue damage. Endotoxins are produced by both Gram-positive and Gram-negative bacteria and act mainly to disturb host cell functions. Endotoxins, produced by Gram-negative bacteria including, *P. aeruginosa*, are normally expressed on their cell wall and are typically released in huge quantities after bacterial lysis (Shenep and Mogan, 1984). LPS is a component of the Gram-negative bacterial membrane and is the most common endotoxin (Ulevitch and Tobias, 1999). The over-reaction to LPS can cause severe inflammatory diseases such as sepsis (Peyssonnaud *et al.*, 2007)

1.3.4.1 Host-Pathogen Interactions

Research has largely described the immune system response to free-living (not in a biofilm arrangement) pathogenic infection. It is essential to understand the immune system's response to bacteria and apply that understanding to acute and chronic wound infections (Wolcott *et al.*, 2008; Mahla *et al.*, 2013).

The host has highly developed approaches to recognise and identify bacteria and other microorganisms. Innate immunity (Figure 1.8) is initiated by pathogen-associated molecular patterns (PAMPs), which are molecules expressed on the cell membrane of bacteria and microorganisms. PAMPs are recognised by pattern recognition receptors (PRRs) expressed on the membrane of immune cells (Hoffmann *et al.*, 1999; Janeway and Medzhitov, 2002; Broz and Monack, 2013; Fukata and Arditi, 2013). PAMPs, such as LPS, bind to PRRs including Toll-like receptors (TLRs), and initiate a signalling cascade normally, via activation of nuclear transcription factor kappa beta (NF- κ B) (Aliprantis *et al.*, 1999; Kopp and Medzhitov, 1999). This event initiates the immune system to release a variety of

pro-inflammatory cytokines, such as granulocyte colony-stimulating factor (G-CSF), granulocyte-macrophage colony-stimulating factor (GM-CSF), macrophage colony-stimulating factor (M-CSF), interferon-gamma (IFN- γ), tumour necrosis factor alpha (TNF- α), interleukin-1 (IL-1), interleukin-8 (IL-8) and interleukin-6 (IL-6) (Wolcott *et al.*, 2008). These cytokines, particularly IL-8, are responsible for regulating the recruitment and the activation of neutrophils, and upregulating antibacterial activity in macrophages and dendritic cells. As the innate immune response progresses, pathogenic invaders are labelled with complement, allowing neutrophils and macrophages to identify, bind, engulf and finally digest them (Shepherd, 1986). Neutrophils are then digested by tissue macrophages to avoid host tissue destruction that could result from excessive secretion of neutrophil-derived proteases.

The PRRs are distributed into four functional families: Toll-like receptors (TLRs), nucleotide-binding oligomerisation domain-like receptors (NLRs), C-type lectin receptors (CLRs), and RIG-1 like receptors (RLRs) (Kopp and Medzhitov, 1999). Several TLRs respond to constituents of Gram-positive and Gram-negative bacteria and stimulate the immune inflammatory reaction (Hoffmann *et al.*, 1999).

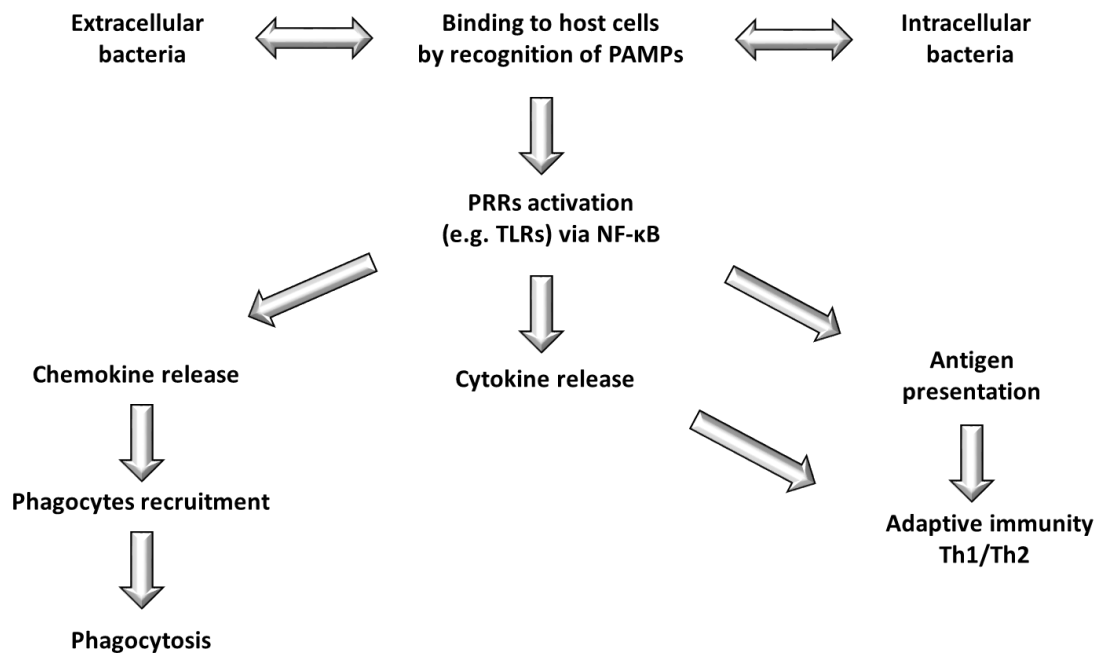


Figure 1.8. Innate immunity in host-pathogen interaction. When TLRs are activated, the innate immune response is initiated through the release of chemokines and cytokines. The acquired immune response is initiated by cytokines of Th1 and Th2, resulting in the production of antibodies. Figure drawn based on information in Basset *et al.* (2003).

The activated complement system (Figure 1.9) consists of three main pathways: classical pathway, alternative pathway and lectin pathway. Complement is a key player of the innate immune system, providing a quick, robust response against foreign invaders, but it also has a very significant impact on the acquired immune system involving T and B cells to remove invaders (Dunkelberger and Song, 2010). Thirty proteins are involved in the complement pathways (Sarma and Ward, 2011). One of their main roles is to lyse microorganisms and to encourage the production of inflammatory mediators that attract phagocytes and initiate the inflammatory response (Sarma and Ward, 2011). The main complement components involved in host cell recruitment at the infection site are C3a, C4a and C5a. Even though the complement system comprises several proteins, they all finally converge in the production of C3, which is the most abundant complement protein in the blood. C3 is converted into various active forms including C3a, C3b, C5a and the membrane attack complex (C5b-9) (Sarma and Ward, 2011).

The alternative pathway is an antibody independent pathway and represents a key element of innate immunity. It is initiated by molecules such as carbohydrates and

endotoxins found on foreign invaders (Qu *et al.*, 2009). The classical pathway depends on the production of antibodies, and is a key element of the adaptive specific immune system (Sarma and Ward, 2011). The lectin pathway is activated by the binding of either mannose binding lectin (MBL) or ficolin to carbohydrates present on the membrane of invaders, including bacteria, viruses and parasites (Sarma and Ward, 2011).

During bacterial infection, C3 is cleaved to opsonin (C3b), which binds to the pathogen to enable the immune phagocytic cells to remove the foreign invader (Walport, 2001; Sarma and Ward, 2011). Key cells of the innate immunity are epithelial, mast and dendritic cells in addition to phagocytic cells such as neutrophils, monocytes and macrophages.

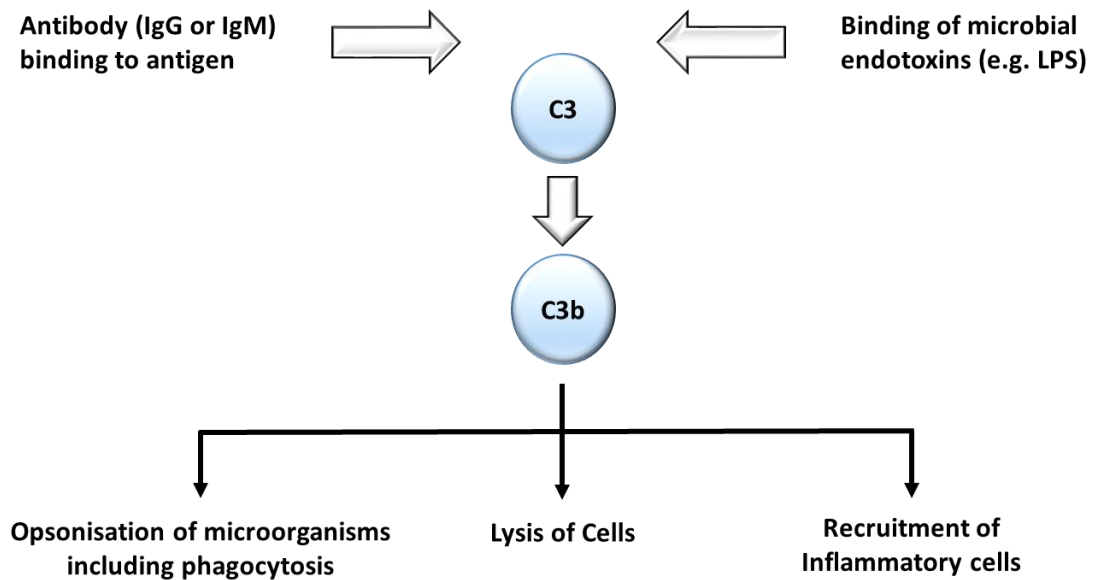


Figure 1.9. Representation of the complement system in host-pathogen interactions. Proteins of the complement system all converge at C3, which is then cleaved into C3b. C3b binds to the pathogen, resulting in its recognition and removal by immune cells, mainly via phagocytosis. Figure drawn based on information in Dumitru *et al.* (2000).

1.3.4.2 The Phagocytosis Process

Phagocytosis is a complex process whereby a phagocyte eliminates or engulfs large cargo $0.5 > \mu\text{m}$ by an actin-dependent mechanism (Mukherjee *et al.*, 1997; Aderem and Underhill, 1999). The term “professional phagocytes” was first introduced by Rabinovitch (1995). It includes a specialised subset of host cells such as monocytes, macrophages, neutrophils and dendritic cells (Ginhoux and Guilliams, 2016).

Phagocytosis serves as a system to clear away the cellular and non-cellular materials from the extracellular space. It also removes microorganisms, such as bacteria, via opsonisation (antibody or complement) binding or pathogen-specific receptors (Arandjelovic and Ravichandran, 2015).

Phagocytes are attracted to an infection site via chemotaxis in response to chemokines, cytokines such as IL-8, and bacterial elements such as LPS (Aliprantis *et al.*, 1999). The process of phagocytosis begins with recognition of foreign invaders by a phagocytic cell. Microorganisms, such as bacteria, associate with the surface of the phagocyte resulting in actin polymerisation (Freeman and Grinstein, 2016; Garcia-Gomez *et al.*, 2016), arrangement of the actin cytoskeleton followed by engulfment, phagosome and phagolysosome formation. Bacterial destruction is mediated by lysosomal enzymes and ROS (Pauwels *et al.*, 2017).

Many receptors contribute to phagocytosis with many particles being detected by several receptors such as antibody receptors (e.g. Fc receptor) and/or complement receptors (CR), in addition to the dual function of some receptors in internalisation and adhesion (Freeman and Grinstein, 2014; Freeman and Grinstein, 2016). It is important that immune cells can distinguish harmful invaders from self (Randow *et al.*, 2013), which is achieved through recognition of pathogen-associated particulates by phagocytic receptors (Santoni *et al.*, 2015; Silva-Gomes *et al.*, 2016). These particulates are recognised by pattern recognition receptors (PRRs) on the membrane of phagocytes (Broz and Monack, 2013; Fukata and Arditi, 2013). Pathogen-associated molecular patterns (PAMPs) that are recognised by PRRs include lipopolysaccharides (LPS) from Gram-negative bacteria, lipoteichoic acid (LTA) from Gram-positive bacteria, mannans from yeast and formulated bacterial peptides (Bæk *et al.*, 2016; Galinari *et al.*, 2017; Lam *et al.*, 2017; Pauwels *et al.*, 2017).

When the innate system is functioning appropriately, the eradication of bacteria and pathogens occurs quickly but when the system is compromised by host comorbidities, bacteria can become difficult to eradicate. After eliminating an invading pathogen, the immune system memorises the pathogen and develops

long-term defences against it, known as adaptive immunity (acquired immune response) (discussed in 1.1.2) (Wolcott *et al.*, 2008).

1.3.4.3 Monocytes/Macrophages

Monocytes are cells that circulate in the blood, spleen, and bone marrow, and represent around 10% of the entire population of human leukocytes (Geissmann *et al.*, 2010). They are the biggest white blood cells, and have distinctive morphological structures, such as irregular cell shape, a folded or kidney-shaped nucleus, and cytoplasmic vesicles. Monocytes remain circulating in the bloodstream for about 1 to 3 days before moving into tissues, where they differentiate into macrophages or dendritic cells. If they are not recruited into tissues by a stimulus, they die and are removed by other immune cells. Monocytes are created in the bone marrow from hematopoietic stem cells (HSCs) and mature over successive differentiation phases: the common myeloid progenitor (CMP) (Akashi *et al.*, 2000), the granulocyte-macrophage progenitor (GMP) (Akashi *et al.*, 2000), the macrophage and dendritic cell precursors (MDP) (Fogg *et al.*, 2006), and the committed monocyte progenitor (cMoP) (Hettinger *et al.*, 2013; Italiani and Boraschi, 2014). Monocytes are considered to be the main reservoir of myeloid precursors for the regeneration of tissue macrophages and dendritic cells. However, some subpopulations of macrophages and dendritic cells (for instance skin Langerhans cells (LC), lymphoid organ dendritic cells and brain microglia) are generated originally from the MDP instead of monocytes (Liu *et al.*, 2009; Boltjes and Van Wijk, 2014), and in some circumstances, they can be directly generated from the bone marrow (Ajami *et al.*, 2007).

Human peripheral blood monocytes are a heterogeneous population of cells (Passlick *et al.*, 1989; Italiani and Boraschi, 2014). Recent studies have indicated that human monocytes are divided into three functional subsets (Italiani and Boraschi, 2014). The characterisation of these subsets is under investigation since it remains largely unclear, as do the exact functions they play in inflammation *in vivo*, in comparison with activated macrophages (see below).

Macrophages are heterogeneous phagocytic cells that reside in lymphoid and non-lymphoid tissues of all adult humans and animals. When they are inactive (under normal healthy conditions), they represent up to 15 % of the entire number of immune cells. However, this percentage becomes higher in reaction to inflammatory stimuli. Depending on their tissue location, macrophages have various designations such as alveolar macrophages (lung), osteoclasts (bone), Küpffer cells (liver), histiocytes (connective tissue), and skin Langerhans cells (skin) (Gautier *et al.*, 2012). Macrophages exert the same functions in all tissues. They are crucial in the immune response to pathogens (by producing and directing the inflammatory response), in tissue development, in monitoring tissue changes, and in the maintenance of steady-state tissue homeostasis (by elimination of apoptotic cells, by remodelling and repairing tissues, and by the production of growth factors). Macrophages are equipped with a broad range of pathogen-recognition receptors that make them efficient at phagocytosis, and they induce the production of inflammatory cytokines (Lin *et al.*, 2008).

Macrophages are characterised by the ability to engulf and digest cellular debris, microbes, cancer cells, and any other substance that does not express on its surface the normal, healthy-cell specific proteins, via a process known as phagocytosis (Duque and Descoteaux, 2015). In addition to phagocytosis, macrophages play a crucial role in both innate (non-specific) and adaptive (specific) immunity as they have the capacity to present antigens to other immune cells, such as T cells, to release cytokine (pro-inflammatory/anti-inflammatory mediators) and to move in response to chemokines in a process called chemotaxis (Duque and Descoteaux, 2015). Macrophages are also crucial for the secretion of growth factors and cytokines. Growth factors and cytokines cause the attraction and division of cells involved in tissue repair (Werner and Grose, 2003), leading to angiogenesis, re-epithelialisation of wounds, and the formation of new extracellular matrix (Greenhalgh, 1998; Stashak *et al.*, 2004).

Evidence has shown that there are numerous, activated forms of macrophages (Mosser and Edwards, 2010). There are two subtypes of macrophages, designated as M1 and M2. M1 macrophages encourage inflammation through the secretion of

pro-inflammatory cytokines whereas M2 macrophages reduce inflammation and encourage tissue repair (Mills, 2012). Lipopolysaccharide (LPS) and interferon-gamma (IFN- γ) are typically present in infected wounds and lead to the generation of M1 macrophages. In contrast, M2 macrophages that stimulate tissue formation (e.g. during wound healing), are promoted by factors such as by interleukin-4 (IL-4) (Biswas and Mantovani, 2010; Sica and Mantovani, 2012).

Table 1-2. Main functions of tissue resident macrophages.

Apart from tissue-specific roles, tissue macrophages exert a series of common roles including removal of cellular debris, immune surveillance, wound repair, defence against microorganisms including bacteria, and the origination and resolution of inflammation. Figure adapted from Italiani and Boraschi (2014).

Tissue Macrophages	Tissue location	Functions
Heart macrophages	Heart	Surveillance.
Osteoclasts	Bone	Bone modelling and remodelling, support to haematopoiesis (Pollard, 2009; Blin-Wakkach <i>et al.</i> , 2014).
Microglia	Brain	Brain progress, immune surveillance, synaptic remodelling (Paolicelli <i>et al.</i> , 2011; London <i>et al.</i> , 2013).
Alveolar macrophages	Lung	Clearance of surfactant and inhaled pathogens (Maus <i>et al.</i> , 2002).
Intestinal macrophages	Gut	Tolerance to microbiota, defence against pathogens, intestinal haemostasis (Zigmond and Jung, 2013).
Bone marrow macrophages	Bone marrow	Reservoir of monocytes, waste removal (Davies <i>et al.</i> , 2013).
Langerhans cells	Skin	Immune surveillance.
Küpfper cells	Liver	Toxin elimination, lipid metabolism, erythrocyte, microbes' clearance, and cell debris clearance from blood (Klein <i>et al.</i> , 2007; Ganz, 2012).
Adipose tissue associated macrophages	Adipose tissue	Metabolism, adipogenesis, adaptive thermogenesis (Nguyen <i>et al.</i> , 2011).
Marginal zone, red pulp macrophages	Spleen	Erythrocyte clearance, iron recycling, uptake of microorganisms from blood (Den Haan and Kraal, 2012).
Inflammatory macrophages	All tissues	Protection against pathogens and harmful stimuli (Labonte <i>et al.</i> , 2014).
Healing macrophages	All tissues	Angiogenesis, branched morphology (Mantovani <i>et al.</i> , 2013).

U937 monocytes are an eternal human cell line commonly used to generate macrophage-like cells for *in vitro* studies. U937 cells exhibit characteristics of immature monocytes and, after conversion into macrophages, these cells display similar features and characteristics to human tissue macrophages (Minafra *et al.*, 2011). Phorbol 12-Myristate 13-Acetate (PMA), also known as TPA, is a phorbol ester used to transform monocytes into macrophage-like cells *in vitro*. PMA-treated U937 cells express high levels of CD11b, CD11c and protein kinase-C (PKC) (Sintiprungrat *et al.*, 2010). When treated with PMA, U937 monocytes upregulate their cell adhesion molecules, such as beta-2 integrins (CD11a, CD11c, CD18 and CD11b) allowing the cells to attach to cell culture flask surfaces. PMA triggers calcium and phospholipid-dependent isoforms of PKC and encourages cyclic AMP metabolism, initiating maturation into a macrophage-like morphology (Luscinskas *et al.*, 1994).

1.4 Estrogen

Estrogens are a group of sex steroid hormones which were first discovered by Edgar Allen and Edward Doisy in 1923 (MacCorquodale *et al.*, 1936). In humans, three main forms of estrogen are described; estradiol, estrone and estriol with 17- β -estradiol being the most potent and biologically active form. Estrogen is involved in several physiological roles such as the regulation of ovulation and fertility, development of female secondary sexual characteristics, synthesis of lipoproteins, regulation of insulin sensitivity, maintenance of bone mass, and conservation of cell growth and cognitive function (Cagnacci *et al.*, 1992). Estrogen is also important in males as it is involved in spermatogenesis (Gladen *et al.*, 1996; Miura *et al.*, 1999; O'donnell *et al.*, 2001), maintaining the density of bone (Stěpán *et al.*, 1989; Daniell, 1997) and cardiovascular health (Group, 1973; Eriksson *et al.*, 1989). Evidence has suggested that estrogen deficit is a key regulator of delayed wound healing in the elderly (Ashcroft *et al.*, 1997b; Ashcroft and Ashworth, 2003). Estrogen has protective anti-inflammatory properties in several tissues, such as the brain (Straub, 2007). Despite the anti-inflammatory effect of estrogen on the skin, mediated via macrophage migration inhibitory factor (MIF) (Hardman *et al.*, 2005), estrogen has been reported to stimulate wound repair independently from its anti-

inflammatory effects in both genders. Systemic and peripheral levels of estrogen decline with increasing age in both males and females (with this diminution highly noticeable in post-menopausal females) (Ashcroft *et al.*, 1997b). HRT-treated post-menopausal females heal acute wounds faster than their age-matched control counterparts, who take no estrogen supplementation (Ashcroft *et al.*, 1997b). Another study indicated that topical estrogen supplementation enhanced wound healing in elderly male and female patients, connected with a reduced inflammatory response (Ashcroft *et al.*, 1997b; Ashcroft *et al.*, 1999b). Accordingly, these findings collectively indicate that estrogen has notably beneficial effects on wound healing.

1.4.1 Estrogen Biosynthesis

Estrogens are produced from cholesterol (Figure 1.10). Cholesterol is converted by several enzymes to androgens, such as testosterone and androstenedione, which are then converted to estrogens through the action of the P450 enzyme aromatase, in the endoplasmic reticulum of estrogen-producing cells (Payne and Hales, 2004). In adipose tissues, androstenedione is converted to estrone whilst in ovarian granulosa cells testosterone is converted into estradiol. Aromatase is found in many peripheral tissues such as bone (Nawata *et al.*, 1995), adipose tissue (Simpson, 2000), brain (Azcoitia *et al.*, 2001) and vascular smooth muscle (Ling *et al.*, 2004). In females at the age of reproduction, systemic estrogen is produced mainly by the ovary. It is fundamentally biosynthesised in the granulosa cells of the ovarian follicles and the corpora lutea. In males, the gonad produces the largest quantity of systemic estrogen. However, a substantial amount of estrogen is also produced locally in peripheral tissues, acting in an autocrine and paracrine manner (Labrie *et al.*, 1998). A substantial amount of inactive steroid precursors including dehydroepiandrosterone (DHEA), its sulphate (DHEA-S), and androstenedione (4-dione) are produced by the adrenals and converted into active steroid hormones in peripheral tissues (Labrie *et al.*, 1998). Several peripheral human tissues, such as adipose tissue, bone and skin can produce active estrogens and androgens locally from conversion of adrenal-derived precursors (Nelson and Bulun, 2001). The reduction in the formation of DHEA-S by the adrenals during aging results in a

dramatic fall in the formation of androgens and estrogens in peripheral target tissues, a situation which could be associated with age-related diseases (Labrie *et al.*, 1998).

Estrogen is synthesised locally in both young and old females and becomes progressively more important after postmenopause, when systemic levels are lost. After the menopause, estrogen biosynthesis changes from a systemic source produced by the ovaries, to a local source, synthesised in peripheral tissues (Picard *et al.*, 2000). However, the rapid decline in the local production of estrogens with increasing age means peripheral estrogen is still insufficient to compensate for the loss in systemic estrogen levels.

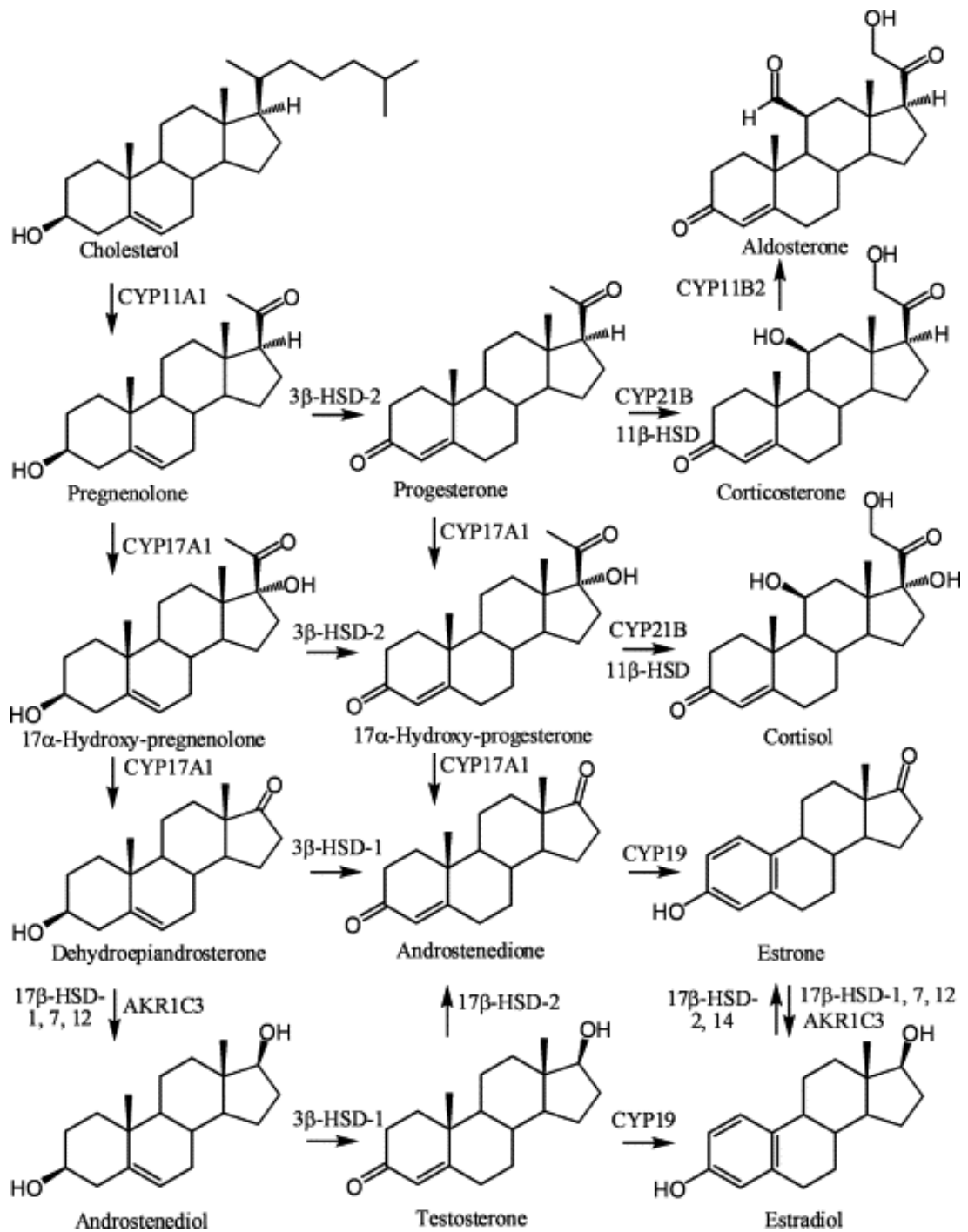


Figure 1.10. Estrogen biosynthesis from cholesterol and androgen precursors. CYP = Cytochrome P450, HSD = Hydroxysteroid dehydrogenase and AKR = Aldo-keto reductase enable hormone synthesis. Figure adapted from Blair (2010).

1.4.2 Estrogen Receptors

Estrogen has multiple functions, such as the regulation of haemostasis, inflammation, cell growth and differentiation. Over the past decades, the existence of two nuclear and membrane-bound estrogen receptor (ER) proteins have been

identified: ER-alpha (ER- α) and ER-beta (ER- β). ER- α was first discovered in 1958 (Jensen and Jacobson, 1960), whereas ER- β was first identified in rat prostate and ovary in 1996 (Mosselman *et al.*, 1996). The biological effects of estrogens are mediated by the binding of estrogen to ER homodimers or heterodimers (Matthews and Gustafsson, 2003), and subsequent activation or repression of gene transcription and signalling pathways, including ligand-dependent/independent pathways (Paige *et al.*, 1999), estrogen response element (ERE)-independent pathways and non-genomic pathways. The ligand-dependent pathway is the central pathway. Interestingly, estrogen signalling can directly target specific ER genes and upregulate the expression of mRNA (Gruber *et al.*, 2002; Ascenzi *et al.*, 2006). Recent research also suggests estrogen could have a direct influence on inflammatory cells, such as monocytes and macrophages, and other cells in the human skin such as keratinocytes, due to the nuclear and membrane-bound localisation of ER proteins. (Weusten *et al.*, 1986; Stimson, 1988; Cocchiara *et al.*, 1990). The response of these particular inflammatory cells depends on the local levels of estrogen and the maturity (stage of differentiation) of the cells (Ashcroft and Ashworth, 2003).

1.4.3 Effect of Estrogen on Skin

It is commonly accepted that the age-related reduction in estrogen levels is linked with skin degeneration. Estrogen deficiency results in detrimental effects on skin's appearance (dry fragile skin, sagging, wrinkling) with decreased thickness and collagen (Ashcroft *et al.*, 1999b; Shah and Maibach, 2001). Estrogen has been shown to have an opposite effect, improving collagen quality, improving skin thickness and augmenting vascularisation. The link between menopause and wound healing has been investigated where it was reported that systemic estrogen supplementation results in an intensification of collagen deposition in post-menopausal women (Savvas *et al.*, 1993; Ashcroft *et al.*, 1997b), whereas topical estrogen increases collagen I and III deposition in the skin of postmenopausal women, conserving skin thickness (Savvas *et al.*, 1993; Sauerbronn *et al.*, 2000). Another study on post-menopausal women reported that there was a reduction in collagen type I and mainly type III in the skin of post-menopausal women compared to pre-menopausal

women, and a decrease in type III/type I ratio within the dermis. This alteration is predominantly associated with estrogen deficiency (Affinito *et al.*, 1999; Horng *et al.*, 2017). It has also been reported that estrogen replacement therapy can improve skin elasticity by 5% per year (Brincat *et al.*, 1987). When locally applied on the skin of post-menopausal women, estradiol significantly increased the production of hydroxyproline and collagen fibres (Albright *et al.*, 1941; Affinito *et al.*, 1999; Sator *et al.*, 2001; Horng *et al.*, 2017).

Additionally, topical estrogen improves the fibres elasticity in the dermis (Albright *et al.*, 1941; Sator *et al.*, 2001). Estrogen also augments the deposition of glycosaminoglycans in the ECM, enhances skin turgor and water retention, stimulates keratinocytes and inhibits matrix metalloproteinase (MMP) synthesis (Brincat, 2000).

1.4.4 Estrogen and Wound Healing

Recent research has demonstrated the key role of sex-steroid hormones in inflammation and wound healing process (Guo and DiPietro, 2010; Gilliver *et al.*, 2007). This is particularly observed in age-related impaired wound healing research where there is an important impact of these hormones on the inflammatory response *in vivo*.

The impact of estrogen on wound healing was first studied in animals in 1947 (Sjövall, 1947; Horng *et al.*, 2017) and then in humans in 1953 (Sjöstedt, 1953; Horng *et al.*, 2017). Afterwards, there has been an accumulating body of evidence supporting that estrogen has a key impact on wound healing (Brincat *et al.*, 1987; Varila *et al.*, 1995) (Affinito *et al.*, 1999; Sauerbronn *et al.*, 2000; Mills *et al.*, 2005; Hardman and Ashcroft, 2008; Brufani *et al.*, 2009; W.-L. Lee *et al.*, 2013; Midgley *et al.*, 2016; Mukai *et al.*, 2016; Chenu *et al.*, 2017; Leblanc *et al.*, 2017; Horng *et al.*, 2017; Pepe *et al.*, 2017; Wilkinson and Hardman, 2017). It has been indicated that estrogen deficiency contributes to cutaneous aging and delayed and/or impaired wound healing (Mukai *et al.*, 2016; Wilkinson and Hardman, 2017). It has also been reported that estrogen replacement therapy could improve the skin elasticity by 5% in one year (Brincat *et al.*, 1987). When locally applied on the skin of

postmenopausal women, estradiol significantly increases the production of hydroxyproline and collagen fibres (Affinito *et al.*, 1999; Horng *et al.*, 2017). Similar effects on the skin and collagen fibres were observed with systemic oral estrogen replacement therapy (Sauerbronn *et al.*, 2000).

Variance in the human immune system between male and female subjects have been identified in some epidemiological and medical studies (McGowan *et al.*, 1975; Bone, 1992), with evidence indicating women have an improved immune system compared to men (Gulshan *et al.*, 1990; Wichmann *et al.*, 1996). This enhanced immune system in women is attributed to the hormonal regulation of the immune system. Other experiments have indicated that estrogen has an immune-improving impact during stress and by contributing resistance to several pathogenic infections (Yamamoto, 1999). Ashcroft *et al.* (1997b) have reported that both systemic and topical estrogen treatments enhance wound healing in elderly men and women. Estrogen has been shown to accelerate re-epithelialisation, encourage angiogenesis, promote matrix deposition and wound contraction, dampen the inflammatory response and the inhibit expression of pro-inflammatory cytokines and proteolytic mediators (Ashcroft and Ashworth, 2003).

1.4.4.1 Effect of Estrogen on the Inflammatory Phase of Wound healing

It is commonly known that age-related impaired healing is associated with an excessive and prolonged inflammatory response, linked with increased inflammatory cell migration and adhesion, and increased pro-inflammatory cytokines such as TNF α (Ashcroft and Ashworth, 2003). Recent research has indicated that chronic wounds are associated with elevated levels of elastase and MMPs, which are released by neutrophils/macrophages and linked with extreme tissue destruction (Gosain and DiPietro, 2004). Estrogen controls and dampens the early inflammatory response during acute wound healing by inhibiting neutrophil infiltration to the wound by reducing the expression of neutrophil adhesion molecules (Ashcroft *et al.*, 1999b). Estrogen causes a decrease in tissue-damaging proteases levels, such as elastase, and an increase in the content of collagen and fibronectin in the dermis (Ashcroft *et al.*, 1999b). Furthermore, estrogen increases

the oxidative metabolism of neutrophils, suggesting estrogen deprivation should lead to diminished phagocytic capability of neutrophils, an increased risk of infection and a postponement in healing (Ashcroft and Ashworth, 2003). Estrogen is also believed to have a direct influence on monocytes and macrophages, due to their possession of nuclear and membrane-bound ER (Weusten *et al.*, 1986; Suenaga *et al.*, 1996; Suenaga *et al.*, 1998). Macrophage migration inhibitory factor (MIF) is a pro-inflammatory cytokine released by monocytes, macrophages, neutrophils, endothelial cells and keratinocytes. This cytokine is known to be a major regulator of the effects of estrogen on wound healing (Gilliver *et al.*, 2010). Ashcroft *et al.* (2003) reported that mice with estrogen deficiency have higher MIF levels, resulting in an elevated inflammatory response and delayed wound healing, whereas MIF null-mice displayed enhanced wound healing, with lower inflammation and greater matrix formation. Estrogen causes a down-regulation of MIF expression leading to a decline in inflammation, enhanced matrix deposition, increased re-epithelialisation and an overall accelerated wound repair (Ashcroft and Ashworth, 2003; Ashcroft *et al.*, 2003).

1.4.4.2 Effect of Estrogen on the Proliferative Phase of Wound Healing

Estrogen improves the proliferative phase of wound healing by enhancing re-epithelialisation, wound contraction, granulation tissue construction and angiogenesis. Estrogen enhances the mitogenesis of keratinocytes and increases the speed of re-epithelialisation and wound contraction in post-menopausal women (Ashcroft *et al.*, 1997b). The rate of wound re-epithelialisation of post-menopausal women treated with HRT for more than 3 months was similar to levels of re-epithelialisation in pre-menopausal females, whereas a non-HRT post-menopausal group showed diminished re-epithelialisation. One feature of this improved re-epithelialisation following estrogen supplementation is due to increased proliferation of epidermal keratinocytes (Raja *et al.*, 2007). Estrogen promotes PDGF expression by monocytes and macrophages (Mendelsohn and Karas, 1999), leading to mitogenesis and chemotaxis of fibroblasts and a subsequent increase in wound contraction and ECM deposition (Seppä *et al.*, 1982). Estrogen also enhances fibrosis by increasing wound TGF- β 1 levels (Ashcroft *et al.*,

1997b; Ashcroft *et al.*, 1999b), resulting in enhanced formation of ECM, particularly collagen deposition (Ashcroft and Ashworth, 2003). Estrogen promotes angiogenesis, leading to increased granulation tissue (Iyer *et al.*, 2012) through a direct stimulation of endothelial cells (Rubanyi *et al.*, 2002). Estrogen increases endothelial cell attachment to laminin, collagen I and IV, laminin, and fibronectin. In addition, estrogen enhances the creation of capillary-like structures by endothelial cells, when positioned on a reconstructed basement membrane (Morales *et al.*, 1995).

1.4.4.3 Effect of Estrogen on the Remodelling Phase of Acute Wound Healing

Estrogen improves wound remodelling by controlling the degradation of collagen through wound proteases. Topical estrogen supplementation increases the deposition of collagen in during the remodelling phase of wound repair in elderly patients (Ashcroft *et al.*, 1997b; Ashcroft *et al.*, 1999b). Estrogen also acts by stimulating the expression of TGF- β 1 to improve collagen deposition in the dermis (Ashcroft *et al.*, 1997b; Ashcroft *et al.*, 1999b). Matrix collagen deposition at 7 and 84 days post-wounding was decreased in post-menopausal women lacking HRT treatment. In contrast, post-menopausal females who took HRT for more than 3 months had similar levels of matrix collagen deposition and wound remodelling as younger pre-menopausal females (Ashcroft *et al.*, 1997b; Ashcroft *et al.*, 1999b).

1.4.5 Estrogen and Chronic Wounds

Since systemic estrogens decline with age, it has been suggested that estrogen deprivation in the elderly could possibly be linked with pathological wound healing. Margolis *et al.* (2002) performed a case-cohort study to investigate the protective effects estrogen on chronic wounds. Patients aged over 65 years receiving HRT treatment were shown to be 30-40% less likely to develop a venous leg ulcer than age-matched patients lacking HRT supplementation (Margolis *et al.*, 2002). Chronic wounds are characterised by an excessive and chronic prolonged inflammation. High levels of inflammatory mediators, including TNF α , TGF β , IL-1 β , IL-6, IGF-1 and MMPs, present in chronic wound exudate (Ashcroft *et al.*, 1997b; Ashcroft *et al.*,

1999b) are downregulated via the action of estrogen (Straub, 2007; Wira *et al.*, 2015).

1.5 Aim and Objectives

Despite its anti-inflammatory activity in wound healing, the effect of estrogen on bacterial clearance from acute or chronic wounds remains largely unknown. Moreover, it is not known whether estrogen deprivation in the elderly exacerbates chronic wound infections. However, there is a vital need to develop novel therapeutic strategies to treat bacterial wound infections in the elderly and one exciting approach is to enhance innate host responses so the immune system is better able to eradicate potential pathogens. Thus, the aim of this study was to determine the effects of estrogen on the phagocytic function of macrophages using *in vitro* and *ex vivo* host-pathogen models of the inflammatory phase of age-related impaired healing.

Specific objectives of this study were to:

- Generate an *in vitro* model of HPBM and U937 monocyte differentiation into M0-like, M1-like and M2-like macrophages.
- Determine the effect of estrogen on the phagocytosis of Gram-positive (MRSA) and Gram-negative (*P. aeruginosa*) wound pathogens by U937- and HPBM-derived macrophages.
- Compare the effect of age-related changes in estrogen levels on the phagocytic activity of M0-like, M1-like and M2-like macrophages against Gram-positive and Gram-negative bacteria.
- Explore the role of ER- α and ER- β in mediating the effects of estrogen on the phagocytosis of bacteria by pro-inflammatory M1-like macrophages.

- Investigate the effect of estrogen on downstream mediators of phagocytosis including regulating protein levels of FAK, Rac1, RhoA, Cdc42 and RhoG in pro-inflammatory M1-like macrophages.

Chapter 2: Materials and Methods

2.1 Materials

Alexa Fluor 647-conjugated anti-human MMR/CD206 (BioLegend, UK)

Bicinchoninic acid assay reagent (Sigma-Aldrich, UK)

Bovine serum albumin (Fisher Scientific, UK)

Brain heart infusion agar (Oxoid, UK)

Brain heart infusion broth (Oxoid, UK)

Chloramphenicol (Sigma-Aldrich, UK)

Diarylpropionitrile (Sigma-Aldrich, UK)

Dimethylsulphoxide (Sigma-Aldrich, UK)

Dulbecco's phosphate-buffered saline (Sigma-Aldrich, UK)

Estrogen (Sigma-Aldrich, UK)

Ethanol (Fisher Scientific, UK)

Ethylenediaminetetraacetic acid (Fisher Scientific, UK)

Ethylene glycol-bis(β -aminoethyl ether)-N,N,N',N'-tetraacetic acid (Fisher Scientific, UK)

EZ-ECL chemiluminescence detection kit (Biological Industries IBH Ltd, USA)

Fluorescein isothiocyanate (FITC)-conjugated anti-human CD11c antibody (Clone Bu15; BioLegend, UK)

Fluorescein isothiocyanate (FITC)-conjugated anti-human CD197 (CCR7) antibody (BioLegend, UK)

Foetal bovine serum (Lonza, UK)

Fulvestrant (Sigma-Aldrich, UK)

Gentamicin (Sigma-Aldrich, UK)

Glutaraldehyde (Sigma-Aldrich, UK)

Green fluorescent protein (GFP)-*S. aureus* strain SH1000 (Kind gift from Paul Williams, Professor of Molecular Microbiology, Centre for Biomolecular Sciences, Faculty of Medicine & Health Sciences, University of Nottingham, UK)

Human blood cell culture medium (Cell Applications Inc, San Diego, USA)

Human peripheral blood CD14⁺ monocytes (Cell Applications Inc, San Diego, USA)

Hydrogen chloride (Thermo Fisher Scientific, UK)

Interferon gamma (Sigma-Aldrich, UK)

Interleukin-4 (Sigma-Aldrich, UK)

Interleukin-13 (Sigma-Aldrich, UK)

Lipopolysaccharide from the membrane of *Escherichia coli* (Sigma-Aldrich, UK)

MCherry- *P. aeruginosa* strain PAO1 (Kind gift from Paul Williams, Professor of Molecular Microbiology, Centre for Biomolecular Sciences, Faculty of Medicine & Health Sciences, University of Nottingham, UK)

Methanol (Thermo Fisher Scientific, UK)

Methicillin Resistant *Staphylococcus aureus* strain 11 (Hospital isolates, Manchester, UK)

Mouse primary antibody to Cdc42 (Abcam, UK)

Mouse primary antibody to Cyclophilin (Abcam, UK)

Mouse primary antibody to Fak (Abcam, UK)

Mouse primary antibody to Rac1 (Abcam, UK)

Mouse primary antibody to RhoA (Abcam, UK)

Mouse primary antibody to β -actin (Abcam, UK)

Mouse primary antibody to β -tubulin (Abcam, UK)

Nitrocellulose membrane (GE Healthcare Life Sciences, UK)

Nutrient agar (Oxoid, UK)

Nutrient broth (Oxoid, UK)

Paraformaldehyde (Sigma-Aldrich, UK)

Penicillin – streptomycin (Lonza, UK)

Peroxidase-HRP anti-mouse IgG (produced in A9044 rabbit) (Sigma-Aldrich, UK)

Peroxidase-HRP anti-rabbit IgG (produced in goat) (Sigma-Aldrich, UK)

pH meter (Hanna Instruments, UK)

Phalloidin–tetramethylrhodamine B isothiocyanate conjugate from *Amanita phalloides* (Fluka, Germany)

Phorbol 12-myristate 13-acetate (Applichem, Germany)

Propylpyrazoletriol (Sigma-Aldrich, UK)

Protease inhibitor cocktails (Sigma-Aldrich, UK)

Pseudomonas aeruginosa strain PAO1 (Hospital isolates, Manchester, UK)

Pyrazole triol (Sigma-Aldrich, UK)

Rabbit primary antibody to RhoG (Abcam, UK)

Roswell Park Memorial Institute Medium (RPMI) (Lonza, UK)

Saline tablets (Sigma-Aldrich, UK)

Skimmed milk powder (Premier Foods Group, UK)

Sodium chloride (Thermo Fisher Scientific, UK)

Sodium deoxycholate, phenylmethylsulfonyl fluoride (Thermo Fisher Scientific, UK)

Sodium dodecyl sulfate (Sigma-Aldrich, UK)

Sodium hydroxide (Fisher Scientific, UK)

Tamoxifen (Sigma-Aldrich, UK)

Tetracycline (Sigma-Aldrich, UK)

Theophylline, 8-[(benzylthio)methyl]-(7Cl,8Cl) (Sigma-Aldrich, UK)

Trypan blue (Lonza, UK)

Trypsin EDTA (Lonza, Belgium)

Tris base (Fisher Scientific, UK)

Tris hydrochloride (Fisher Scientific, UK)

Tris (hydroxymethyl) aminomethane (THAM) hydrochloride (Fisher Scientific, UK)

Triton X-100 (Lonza, Belgium)

Tween-20 (Fisher Scientific, UK)

U937 Human monocytic cell line (Health Protection Agency Culture Collections, UK)

2.2 Methods

2.2.1 Preparation of RPMI-1640 Complete Medium

Foetal bovine serum (FBS) was heat-inactivated in a water bath maintained at 56 °C for 30 minutes. Roswell Park Memorial Institute (RPMI)-1640 (L-Glutamine, 25 mM HEPES) supplemented with 10% FBS and 100 I.U./mL penicillin-streptomycin was prepared under aseptic conditions and referred to as RPMI-1640 complete medium throughout the study.

2.2.2 Estrogen Treatment Preparation

Estrogen (17 β -estradiol) was prepared to final concentrations of 1×10^{-7} M, 1×10^{-8} M, 1×10^{-9} M and zero M in antibiotic-free RPMI-1640 complete medium or human blood cell culture medium (HBCCM). Concentrations of estrogen were chosen according the previously published models of ageing to represent physiological levels of estrogen (1×10^{-8} M) typical of young human adults, supraphysiological levels of estrogen (1×10^{-7} M) to model estrogen supplementation, estrogen deprivation (1×10^{-9} M) to represent declining levels found in elderly humans and an absolute absence of estrogen (zero M) (Ashcroft and Ashworth, 2003; Hardman *et al.*, 2005; Sproston *et al.*, 2018).

2.2.3 Cell Culture

U937 monocytes were cultured under aseptic conditions at 37 °C and 5% CO₂ using RPMI-1640 complete medium. CD14⁺ human peripheral blood monocytes (HPBM) were cultured in HBCCM under aseptic conditions at 37 °C and 5% CO₂. The U937 and HPBM cell suspensions were centrifuged at 500 *g* for 7 minutes before the medium was aspirated. U937 monocytes and HPBM were maintained at 0.5×10^6 cells/mL and 1×10^6 cells/mL respectively by resuspension in fresh medium every other day. Sterile filtered 0.4% trypan blue dye was used to check cell viability according to published methodologies (Mir *et al.*, 1991) using a 1:1 ratio of cell suspension to trypan blue and counting the number of non-viable cells that took up

the blue dye using a TC10 automated cell counter (Bio-Rad, USA). The viability of the cells was above 80% for all experiments.

2.2.4 Differentiation of Monocytes into M0 Macrophages

HPBM and U937 monocytes at a concentration of 1×10^6 viable cells/mL were differentiated into macrophage-like cells by incubating with 50 ng/mL phorbol 12-myristate 13-acetate (PMA) in RPMI complete medium for 24 hours at 37 °C and 5% CO₂. Cells were washed twice with complete medium and then incubated for a further 48 hours in PMA-free complete medium to obtain resting macrophages known as M0 macrophages (Daigneault *et al.*, 2010; Rios de la Rosa *et al.*, 2017).

2.2.5 Generation of M1 and M2 Macrophages

U937 monocytes and HPBM were cultured as described in 2.2.3. Cells were differentiated into resting M0 macrophages as described in 2.2.4. The medium was aspirated and the cells washed twice with complete medium. M1 and M2 macrophages were generated using the method adapted from Rios de la Rosa *et al.* (2017). For M1 differentiation/polarisation, M0 macrophages were incubated with 100 ng/mL lipopolysaccharide (LPS) and 20 ng/mL interferon gamma (IFN- γ) in PMA-free complete medium for 24 hours at 37 °C and 5% CO₂. The adherent cells were washed twice with LPS/IFN- γ -free complete medium and then incubated with complete cell culture medium for 4 days at 37 °C and 5% CO₂, with medium changes every 48 hours to obtain classically activated resting M1 macrophages. For M2 differentiation/polarization, M0 macrophages were incubated for 24 hours with 20 ng/mL interleukin-4 (IL-4) and 20 ng/mL interleukin-13 (IL-13) in complete medium. The adherent cells were washed twice with IL-4/IL-13-free complete medium prior to incubation in complete medium for 4 days, with medium changes every 48 hours to obtain alternatively activated resting M2 macrophages.

2.2.6 Flow Cytometry

2.2.6.1 Confirming the Differentiation of Monocytes into M0 Macrophages

Differentiation of U937 and HPBM monocytes into M0 macrophages was confirmed by flow cytometry via analysis of CD11c⁺ surface marker expression. Monocytes and M0 macrophages were seeded in 1 mL of RPMI/HBCCM complete medium at a density of 1×10^6 cells/mL in 12-well plates according to methods described in 2.2.3 and 2.2.4 respectively. Adherent macrophages were detached from six replicate wells by removing the supernatant and adding 250 μ L trypsin EDTA for 3 minutes at 37 °C and 5% CO₂. The trypsin EDTA was neutralised by adding 250 μ L RPMI/HBCCM complete cell culture medium before being washed (centrifugation of well constituents at 500 *g* for 7 minutes and resuspension of the cell pellet) in 1 mL complete medium. The monocytes (6 replicate wells) and detached macrophages (6 replicate wells) were then centrifuged at 500 *g* for 7 minutes and supernatants discarded prior to fixing pelleted cells with 200 μ L 4% paraformaldehyde in Dulbecco's phosphate-buffered saline (DPBS) for 10 minutes at room temperature (RT). Cells were washed twice in 200 μ L DPBS. Half the fixed macrophage samples (*n* = 3) and half the fixed monocyte samples (*n*=3) were stained for 30 minutes at RT with FITC-conjugated anti-human CD11c antibody diluted 1:40 with 10% FBS in DPBS. The remaining fixed monocyte (*n* = 3) and fixed macrophage (*n* = 3) samples were prepared as unstained negative controls by incubating at RT for 30 minutes with 10% FBS in DPBS lacking antibody. Unbound antibody was removed from samples by two sequential wash steps in DPBS, before resuspension in 500 μ L DPBS. CD11c surface marker expression was assessed on 10,000 events (live, individual cells) with a BD Accuri C6F1 cytometer (BD Biosciences, USA). Data were analysed with BD Accuri C6 Software after gating events in the forward scattered channel (FSC)/side scattered channel (SSC) and fluorescence parameter 1 (FL1-A) windows. The average percentage CD11c⁺ cells (%) and median fluorescence intensity (MFI) were determined relative to unstained negative control U937/HPBM monocytes.

2.2.6.2 Polarisation of M0 Macrophages into M1/M2 Macrophages

Polarisation of U937 and HPBM M0 macrophages into M1/M2 macrophages was confirmed by flow cytometry via analysis of CD197⁺ and CD206⁺ surface marker expression respectively. M0 macrophages were seeded in 1 mL of RPMI/HBCCM complete medium at a density of 1×10^6 cells/mL in 12 well plates according to methods described in 2.2.4. M0 macrophages were polarised into M1 (n = 6 replicate wells) and M2 (n = 6 replicate wells) macrophages as described in 2.2.5. Adherent M0/M1/M2 macrophages (n = 6 replicate wells) were detached using trypsin, fixed with 4% paraformaldehyde and washed twice with DPBS as described in 2.2.6.1. Supernatants were discarded and half the M0/M1/M2 samples (n = 3 replicate wells) were stained for 30 minutes at RT with FITC-conjugated anti-human CD197 (CCR7) antibody diluted 1:50 with 10% FBS in DPBS, washed twice with 500 μ L DPBS before being incubated with Alexa Fluor 647-conjugated anti-human MMR/CD206 antibody (1:50 with 10% FBS in DPBS) for 30 minutes at RT. The remaining M0/M1/M2 samples (n = 3 replicate wells) were incubated at RT for 1 hour with 10% FBS in DPBS lacking any antibodies (unstained negative controls). Two sequential wash steps were performed in DPBS prior to resuspension in 500 μ L DPBS as described in 2.2.6.1. Samples were analysed as described in 2.2.6.1 after gating events in the FSC, SSC, FL1-A and fluorescence parameter 2 (FL2-A) windows. The average percentage CD197⁺/CD206⁺ cells (%) and median fluorescence intensity (MFI) were determined relative to the unstained M0 macrophages.

2.2.7 Bacterial Culture Preparation

Methicillin-resistant *Staphylococcus aureus* (MRSA) strain 11, *Pseudomonas aeruginosa* strain PAO1, green fluorescent protein (GFP) labelled-*S. aureus* strain SH1000WT and mCherry-*P. aeruginosa* strain PAO1 were used in this project. When required, cultures of MRS11 and PAO1 were inoculated onto nutrient agar (NA) plates and incubated at 37 °C for 24 hours. Cultures of GFP-*S. aureus* were inoculated onto brain heart infusion (BHI) agar plates supplemented with 10 μ g/mL chloramphenicol and incubated at 37 °C for 24 hours. Cultures of mCherry- *P. aeruginosa* strain PAO1 were inoculated onto NA plates supplemented with 1.25

$\mu\text{g}/\text{mL}$ tetracycline and incubated at $37\text{ }^{\circ}\text{C}$ for 24 hours. The bacterial inoculated agar plates were stored at $4\text{ }^{\circ}\text{C}$ and replaced every four weeks.

When required for experimental assays, MRSA 11 and *P. aeruginosa* PAO1 were cultured in nutrient broth (NB) overnight at $37\text{ }^{\circ}\text{C}$ on an orbital shaker (New Brunswick Scientific, USA). Ten-fold serial dilutions of each broth were prepared in saline and $100\text{ }\mu\text{L}$ aliquots were inoculated onto duplicate NA plates with subsequent incubation at $37\text{ }^{\circ}\text{C}$ overnight, whilst the broths were refrigerated overnight at $4\text{ }^{\circ}\text{C}$ to inhibit further bacterial growth. The colony-forming units (CFU) counted on the NA plates were used to calculate the density of bacteria (CFU/mL) in each NB, taking into account the dilution factor. The neat broths were centrifuged at 3500 rpm for 10 minutes and the bacterial cell pellets washed twice with saline prior to re-suspension at 2×10^6 , 2×10^5 , and 2×10^4 CFU/mL in saline.

GFP-*S. aureus* strain SH1000WT was cultured in BHI broth supplemented with $10\text{ }\mu\text{g}/\text{mL}$ chloramphenicol and incubated at $37\text{ }^{\circ}\text{C}$ overnight on an orbital shaker. Ten-fold serial dilutions of the broth were prepared in saline and $100\text{ }\mu\text{L}$ aliquots of each dilution were inoculated onto duplicate BHI agar plates supplemented with $10\text{ }\mu\text{g}/\text{mL}$ chloramphenicol with subsequent incubation at $37\text{ }^{\circ}\text{C}$ overnight. The neat broth was refrigerated overnight at $4\text{ }^{\circ}\text{C}$ to inhibit further bacterial growth. The CFU counted on the agar plates were used to calculate the density of bacteria (CFU/mL) in the neat broth, taking into account the dilution factor. The bacterial broth was centrifuged at 3500 rpm for 10 minutes and the cell pellet washed twice with saline prior to re-suspending the bacteria at 2×10^5 CFU/mL in saline.

Red mCherry-*P. aeruginosa* strain PAO1 was cultured in NB supplemented with $1.25\text{ }\mu\text{g}/\text{mL}$ tetracycline and incubated at $37\text{ }^{\circ}\text{C}$ overnight on an orbital shaker. Ten-fold serial dilutions of the broth were prepared in saline and $100\text{ }\mu\text{L}$ aliquots of each dilution were inoculated onto duplicate NA plates supplemented with $1.25\text{ }\mu\text{g}/\text{mL}$ tetracycline with consequent incubation at $37\text{ }^{\circ}\text{C}$ overnight. The neat broth was refrigerated overnight at $4\text{ }^{\circ}\text{C}$ to inhibit further bacterial growth. The CFU counted on the agar plates were used to calculate the density of bacteria (CFU/mL) in the neat broth, taking into account the dilution factor. The bacterial broth was

centrifuged at 3500 rpm for 10 minutes and the cell pellet washed twice with saline prior to re-suspending the bacteria at 2×10^5 CFU/mL in saline.

2.2.8 *In vitro* and *Ex vivo* Host-Pathogen Interaction Assays

2.2.8.1 Interaction of M0 Macrophages with MRSA and PAO1

M0 resting macrophages generated from HPBM and U937 monocytes were prepared in 24-well plates as described in 2.2.4. The medium was aspirated and replicate wells ($n = 6$) were treated with different concentrations (1×10^{-7} M, 1×10^{-8} M, 1×10^{-9} M and zero M) of estrogen in corresponding antibiotic-free medium (HBCCM or RPMI-1640 for macrophages derived from HPBM and U937 respectively) for 24 hours at 37 °C and 5% CO₂. In addition, replicate ($n = 6$) blank control wells consisting of antibiotic-free HBCCM or RPMI-1640 medium, but lacking macrophages, were prepared and incubated for 24 hours at 37 °C and 5% CO₂. Replicate negative control wells ($n = 6$) were prepared by treating macrophage-containing wells with 1×10^{-7} M bovine serum albumin (BSA) in corresponding antibiotic-free medium and incubating for 24 hours at 37 °C and 5% CO₂. The supernatant was removed from all wells prior to treating with 1 µg/mL LPS and 100 ng/mL IFN-γ in corresponding antibiotic-free medium for 2 hours at 37 °C and 5% CO₂.

The supernatant was removed from all wells before inoculating the macrophages with 1×10^4 CFU of bacteria in a total volume of 100 µL antibiotic-free medium containing 1 µg/mL LPS and 100 ng/mL IFN-γ (1:1). Plates were incubated for 3 hours at 37 °C and 5% CO₂ to enable host-pathogen interactions (phagocytosis) to occur.

Following the 3 hours host-pathogen interaction period, the supernatant of each well was collected. M0 macrophages were released from each well by adding 450 µL trypsin EDTA and incubating for 3 minutes at 37 °C and 5% CO₂. The trypsin EDTA was neutralised by adding 450 µL antibiotic-free medium to each well. The constituents of each well (900 µL) were collected and added to the previously collected supernatant (100 µL) and thoroughly mixed. Aliquots (100 µL) of each

sample were then inoculated onto NA plates in duplicate and incubated at 37 °C overnight. The number of CFU formed on the agar plates were counted to calculate the bacterial recovery in each sample following the period of host-pathogen interaction.

2.2.8.2 Interaction of M1/M2 Macrophages with MRSA and PAO1

Classically activated resting M1 macrophages and alternatively activated resting M2 macrophages were generated from HPBM and U937 macrophages as described in 2.2.5. The medium was aspirated and 3-hour host-pathogen interaction assays were conducted as described in 2.2.8.1.

2.2.9 Interaction of M0 Macrophages with GFP- *S. aureus*

M0 U937 macrophages were prepared in 50 mL cell culture flasks as described in 2.2.4. The medium was discarded and the cells washed with complete RPMI medium. Flasks were treated with different concentrations (1×10^{-7} M, 1×10^{-8} M, 1×10^{-9} M and zero M) of estrogen in antibiotic-free RPMI medium for 24 hours at 37 °C and 5% CO₂. A negative control flask was prepared by treating with 1×10^{-7} M BSA and incubating at 37 °C and 5% CO₂ for 24 hours. The supernatants in all flasks were discarded before treating all flasks with 1 µg/mL LPS and 100 ng/mL IFN-γ (all in antibiotic-free RPMI medium) for 2 hours at 37 °C and 5% CO₂.

The supernatants were discarded and each flask was then inoculated with 1×10^5 CFU of GFP-*S. aureus* in a total volume of 10 mL antibiotic-free RPMI medium containing 1 µg/mL LPS and 100 ng/mL IFN-γ. All flasks were incubated for a 3-hour period of host-pathogen interaction at 37 °C and 5% CO₂ prior to collecting the supernatant of each flask. Adherent M0 macrophages were released from each flask by adding 5 mL trypsin EDTA and incubating for 3 minutes at 37 °C and 5% CO₂ with gentle detachment using a sterile cell scraper. A volume of 5 mL antibiotic-free RPMI medium was added to each flask to neutralise the EDTA and the contents of each flask (10 mL) was combined with the previously collected supernatant (10 mL) and thoroughly mixed. Each sample (20 mL) was centrifuged at 500 g for 7 minutes and the cell pellet re-suspended in 5 mL antibiotic-free RPMI medium.

A countess II FL cell counter (Life technologies, USA) was used to detect the GFP signals and capture images ($n = 10$) of bacterial internalisation within macrophages from each flask. The phagocyte density, number of internalised bacteria, total number of macrophages and total bacterial counts were determined from images in order to calculate the total number of phagocytes/mL and the ratio of bacterial internalisation/recovery.

2.2.10 Gentamicin Protection Assay

U937 monocytes (1×10^6 cell/well) were seeded in a 24-well plate and differentiated into resting M0 macrophages as previously described in 2.2.6.1. Cells were treated with estrogen (1×10^{-7} M, 1×10^{-8} M, 1×10^{-9} M and zero M) or 1×10^{-7} M BSA and stimulated with LPS ($1 \mu\text{g/mL}$) and IFN- γ (100 ng/mL) before being incubated with 1×10^4 CFU of MRSA 11 and *P. aeruginosa* PAO1 for 3 hours as described in 2.2.8.1. Protection gentamicin assay was performed according to methods detailed by Hockenberry *et al.* (2016). After a three-hour incubation period, the supernatant ($100 \mu\text{L}$) was collected and the wells were treated with 0.1% Triton X-100 ($100 \mu\text{L}$) in DPBS for 5 minutes at 37°C and 5% CO_2 to release the membrane-adherent bacteria. The supernatant was collected and the cells were washed 10 times with DPBS before being incubated with $50 \mu\text{g/mL}$ gentamicin in antibiotic-free RPMI medium for 1 hour to eradicate extracellular and membrane-bound gentamicin-sensitive bacteria. The supernatant was discarded and the cells were washed 10 times with DPBS. The supernatant was discarded and cells were detached after incubation with $400 \mu\text{L}$ Trypsin EDTA at 37°C and 5% CO_2 for 3 minutes. A volume of $400 \mu\text{L}$ antibiotic-free RPMI medium was added to each well to neutralise the trypsin EDTA. The constituents of each well ($800 \mu\text{L}$) were then collected and added to the previously collected supernatants ($200 \mu\text{L}$) and thoroughly mixed. CFU/mL were counted after plating $100 \mu\text{L}$ aliquots on duplicate NA plates and incubating at 37°C overnight.

2.2.11 Visualising Host-Pathogen Interactions

2.2.11.1 Fluorescence Microscopy

Silicon wafers (Montco Technologies, USA) were cut at a diameter of 1 cm² and sterilised with 70% ethanol for 30 minutes before being rinsed with dH₂O and washed with DPBS for 10 minutes. U937 monocytes (0.5 x 10⁶ cell/well) were seeded and differentiated into adherent M0 macrophages on sterile silicon wafers inside a 12- well plate. Cells were treated with/without estrogen (1 x 10⁻⁷ M) or 1 x 10⁻⁷ M BSA, stimulated with LPS (1 µg/mL) and IFN-γ (100 ng/mL), and incubated with GFP *S. aureus* or mCherry *P. aeruginosa* (both at 1 x 10⁵ CFU) using methods described in 2.2.8.1. The supernatant from each well was removed and the cells were fixed with 4% paraformaldehyde in DPBS for 10 minutes at RT. The supernatants were discarded and the cells washed three times with DPBS before being incubated with Triton X-100 in DPBS (0.1%) for 5 minutes at RT. The cells were washed three times with DPBS and treated with a blocking buffer (3% BSA in DPBS) for 1h at RT. The supernatants were aspirated and the cells were immediately treated with phalloidin–tetramethylrhodamine B isothiocyanate conjugate from *Amanita phalloides* (1:200 in blocking buffer) and incubated in the dark at RT overnight. The supernatant was removed and the cells were washed three times with DPBS. The silicon wafers with attached cells were removed using sterile tweezers and attached to a microscope slide using double-sided tape. Samples were observed at 100X on a Nikon E600 epifluorescence microscope (Nikon, Japan) and at 63X under a Leica TSC SPE1000 confocal microscope (Leica Microsystems, UK). Z-stack analysis was performed on samples to localise the bacterial internalisation by macrophages.

2.2.11.2 Scanning Electron Microscopy

Sterile silicon wafers of 1 cm² diameter were washed with DPBS for 10 minutes before being placed in 12-well plates. Adherent U937 M0 macrophages were generated on the silicon wafers. Cells were treated with/without estrogen (1 x 10⁻⁷ M) or 1 x 10⁻⁷ M BSA, stimulated with LPS (1 µg/mL) and IFN-γ (100 ng/mL), and incubated with 1 x 10⁵ CFU of MRSA 11 or PAO1 for 1 hour at 37 °C and 5% CO₂

using methods described in 2.2.8.1. Following the 3-hour period of host-pathogen interaction, the silicon wafers from each well were collected and fixed with 2.5% glutaraldehyde in DPBS. Samples were incubated at 4 °C overnight. The silicon wafers were washed twice in DPBS, then soaked in 20% methanol/ethanol for 30 minutes, 40% methanol/ethanol for 30 minutes, 60% methanol/ethanol for 30 minutes, 80% methanol/ethanol for 30 minutes, and finally in 100% methanol/ethanol for 30 minutes twice before being dried overnight in a vacuum-assisted desiccator (Sigma-Aldrich, UK). A Supra 40VP scanning electron microscope (Zeiss, Germany) was used to capture scanning electron microscopy (SEM) images using SmartSEM software (Carl Zeiss Ltd, Germany) to compare the internalisation of MRSA strain 11 or *P. aeruginosa* strain PAO1 by M0 macrophages with/without presence of estrogen.

2.2.12 Estrogen Receptor Stimulation/Blockade

HPBM and U937 monocytes were differentiated into M1 macrophages as described in 2.2.5. Replicate wells ($n = 6$) were treated with 1×10^{-7} M BSA, 1×10^{-7} M estrogen, zero M estrogen, 1×10^{-6} M tamoxifen, 1×10^{-6} M propyl pyrazole triol (PPT), and 1×10^{-6} M diarylpropionitrile (DPN) (all prepared in complete RPMI medium) and incubated at 37 °C and 5% CO₂ for 24 hours. Treatments were aspirated and the cells washed with DPBS. A volume of 500 µL of complete RPMI medium was added to each well and incubated for 24 hours at 37 °C and 5% CO₂. Further replicate wells ($n = 6$) containing M1 macrophages were treated with 1×10^{-6} M fulvestrant and 1×10^{-6} M theophylline, 8-[(benzylthio)methyl] (TPBM) in complete RPMI medium for 24 hours. Treatments were aspirated and the cells washed with DPBS. Cells were treated with 1×10^{-7} M estrogen and incubated at 37 °C and 5% CO₂ for 24 hours.

Host-pathogen interaction assays were then performed for 3 hours in all wells as described in 2.2.8 and bacterial (MRSA11 and PAO1) recovery determined.

2.2.13 Investigating Mediators of Host-Pathogen Interactions by Immunoblotting

2.2.13.1 Preparation of Buffers

TBS-Tween buffer, adjusted to pH 7.4

- 10 mM Tris-base
- 140 mM Sodium chloride (NaCl)
- 0.1% Tween-20
- Distilled water (dH₂O)

Blocking buffer, adjusted to pH 7.4

- 1% BSA in TBS-Tween

RIPA buffer

- 140 mM Sodium chloride (NaCl)
- 25 mM Tris-hydrochloride (Tris-HCl)
- 1 mM Ethylenediaminetetraacetic acid (EDTA)
- 0.5 mM Ethylene glycol-bis(β -aminoethyl ether)-N,N,N',N'-tetraacetic acid (EGTA)
- 0.5% Sodium deoxycholate, phenylmethylsulfonyl fluoride (PMSF)
- 1% Triton X-100
- 0.1% Sodium dodecyl sulfate (SDS)
- 0.1% Protease inhibitors cocktail
- Ultrapure water

2.2.13.2 Protein Extraction and Quantification

U937 monocytes were differentiated into M1 macrophages in three T25 flasks (1 x 10⁷ cells/flask) using methods described in 2.2.5. The medium in all flasks was discarded and adherent M1 macrophages were treated with 10 mL of 1 x 10⁻⁷ M BSA (negative control), zero M estrogen or 1 x 10⁻⁷ M estrogen in RPMI complete medium in a single flask and incubated at 37 °C and 5% CO₂ for 24 hours. The treatments were aspirated and cells washed with DPBS before incubation with 3 mL

trypsin EDTA for 3 minutes at 37 °C and 5% CO₂ to initiate cell detachment. A volume of 3 mL complete RPMI medium was added to the flasks to neutralise the trypsin EDTA. Cells were transferred into 50 mL centrifuge tubes and centrifuged at 500 *g* for 7 minutes. Supernatants were discarded and 200 µL of ice-cold radio immunoprecipitation assay (RIPA) buffer containing protease inhibitor cocktail (1:1000) was added the cell pellets in order to lyse the cells. Samples were transferred to -80 °C for 30 minutes. Cells were defrosted at 37 °C and sonicated 10 times for 10 seconds, mixing well between each period of sonication. Samples were then centrifuged for 5 minutes at 12000 *g* and the lysates were collected on ice. The total protein content of each sample was quantified using the bicinchoninic acid (BCA) protein quantification assay according to the manufacturer's instructions. Briefly, a 1 mg/mL stock solution of BSA in dH₂O was used to provide a set of protein standards (0.8, 0.6, 0.4 and 0.2 mg/mL). A volume of 25 µL of each BSA standard was added to a 96-well plate with dH₂O (zero mg/mL BSA) used as a blank. A working BCA solution, consisting of reagents A and B (1:50) was prepared as described in the BCA kit. A volume of 200 µL of this solution was added to all the BSA standards and incubated at 37 °C for 30 minutes before measuring the absorbance at 562 nm. A standard curve of absorbance against protein concentration was generated from the protein standards and used to determine the concentration of total protein in cell lysates via interpolation. Protein samples were normalised to 20 mg/mL prior to storage at -80 °C for later use.

2.2.13.3 Immunoblotting

A volume of 5 µL of each of the protein samples (*n* = 6) was blotted onto nitrocellulose membranes. BSA at a concentration of 1 mg/mL (*n* = 6) was used as a negative control. The membranes were allowed to dry before being treated with the blocking buffer and then incubated at RT on a rocking shaker (Cole-Parmer, UK) for one hour with rocking at 50 rpm. The buffer was discarded and the membranes were treated with primary antibodies specific for the proteins Rac1, Fak, RhoG, RhoA, Cdc42 and β-tubulin (Table 2.1) overnight at 4 °C with rocking at 30 rpm. The primary antibodies were recovered (stored at 4 °C for further use) and the membranes washed five times for 5 minutes each with TBS-Tween buffer. The

membranes were treated with the horseradish peroxidase (HRP) anti-mouse IgG (produced in the A9044 rabbit) or HRP anti-rabbit IgG (produced in goat) secondary antibodies as appropriate, made in blocking buffer (1:1000), and incubated at RT for 1 hour with rocking at 50 rpm. When the primary antibody was raised in mouse, a rabbit anti-mouse secondary antibody was used. When the primary antibody was raised in rabbit, a goat ant-rabbit secondary was used (Table 2.1). The secondary antibodies were removed and membranes were washed five times for 5 minutes with TBS-Tween buffer. A working EZ-ECL chemiluminescence detection solution, consisting of reagents A and B (1:1) was prepared in the dark according to the manufacturer's instructions provided in the EZ-ECL kit. The membranes were covered with EZ-ECL working solution and incubated in the dark for 2 minutes at RT. Excess solution was aspirated and the membranes were visualised with a Chemidoc Touch Transilluminator (Bio-Rad, USA) using the Image Lab™ Touch Software at 1-minute exposure times. Dot blots were measured and analysed using ImageJ Software (Version 1.48).

Table 2-1. Primary and secondary antibodies used for immunoblotting.

Primary antibody	Working concentration	Host primary species	Secondary antibody	Host secondary species
β-tubulin	1:5000	Mouse	HRP anti-mouse IgG	Rabbit
Rac1	2 µg/mL	Mouse	HRP anti-mouse IgG	Rabbit
Fak	0.5 µg/mL	Mouse	HRP anti-mouse IgG	Rabbit
RhoG	1:500	Rabbit	HRP anti-rabbit IgG	Goat
RhoA	2 µg/mL	Mouse	HRP anti-mouse IgG	Rabbit
Cdc42	1:500	Mouse	HRP anti-mouse IgG	Rabbit

**Chapter 3: Effect of 17 β -estradiol
on the Phagocytosis of MRSA and
P. aeruginosa by Human M0
Macrophages**

3.1 Introduction

3.1.1 Estrogen Promotes Wound Healing

Active estrogens are secreted by the gonads into the bloodstream, with smaller amounts of active hormones produced by the adrenals. However, a substantial amount of inactive steroid precursors, including DHEA, its sulphate DHEA-S and androstenedione (4-dione) are produced by the adrenals and converted locally into active estrogens (such as 17 β -estradiol) in peripheral tissues such as adipose tissue, bone and skin (Labrie *et al.*, 1998; Nelson and Bulun, 2001). The reduction in the formation of DHEA-S by the adrenals during ageing results in a dramatic fall in the formation of active estrogens in peripheral target tissues, a situation which is strongly associated with age-related processes such as age-related impaired healing (Labrie *et al.*, 1995).

Estrogen declines significantly in postmenopausal women, due to a rapid decline in ovarian secretion of estrogens and a loss of both adrenal-derived estrogen and its precursors with increasing age. In men, estrogen declines rapidly with increasing age, largely due to a loss of adrenal-derived estrogen and its precursors. It is believed that the decline in estrogen in both sexes with increasing age results in detrimental effects on skin appearance (dry skin, sagging, wrinkling) and wound healing (Hardman *et al.*, 2007). However, the detrimental effect of hormone-driven ageing (estrogen deprivation) on the development and progression of wound infections remains largely unknown.

Ageing is known to be a key risk factor for impaired wound healing, with the declining production of estrogen in the elderly being the main key regulator of age-related delayed wound healing (Hardman and Ashcroft, 2008). The reduction in estrogen that occurs with increasing age has a pronounced effect on cutaneous healing in both genders. Ashcroft *et al.* (1997a) and Ashcroft *et al.* (1999a) reported that estrogen deficiency causes a significant delay in wound healing, with topical and systemic estrogen replacement reversing this delay. Moreover, estrogen supplementation promoted age-related wound healing in both elderly males and females (Ashcroft *et al.*, 1999a; Guo and DiPietro, 2010).

Estrogen has been shown to accelerate re-epithelialisation, promote angiogenesis, enhance matrix deposition and wound contraction, dampen the inflammatory response (Ashcroft and Ashworth., 2003). In particular, the key role of estrogen in regulating inflammation has been extensively reviewed (Guo and DiPietro, 2010; Gilliver *et al.*, 2007), with estrogen supplementation decreasing the inflammatory response in elderly subjects of both genders (Ashcroft *et al.*, 1997b; Ashcroft *et al.*, 1999a). Delayed wound healing in the elderly is associated with a delayed but prolonged and excessive inflammatory response, with dysregulated expression of pro-inflammatory cytokines and proteolytic mediators. However, despite the pronounced inflammatory response, evidence suggests increasing age may result in an impaired ability of inflammatory cells to eliminate bacteria from the wounds (Emori *et al.*, 1991; Thomas, 2001). Indeed, the propensity for wound infections increases in the elderly, in part due to the delay in wound repair (Cooper *et al.*, 2015).

Impaired wounds often become colonised with a variety of microbial species, such as *S. aureus*, *P. aeruginosa*, *Proteus* species and anaerobic bacteria (Gjødsbøl *et al.*, 2006; Kirketerp-Møller *et al.*, 2008; Rybtke *et al.*, 2015). When innate immunity functions appropriately, the eradication of wound bacteria is generally successful without intervention. However, if the host immune system becomes compromised by comorbidities, particularly in the elderly, clearance of wound bacteria can fail in the absence of effective treatments. In particular, if wounds become heavily infected with MRSA, *P. aeruginosa* or other hospital-acquired pathogens, treatment might necessitate aggressive medication with last line of defence antibacterial therapies (Beasley and Hirst, 2004). Indeed, the treatment of wounds infected with MRSA or *P. aeruginosa* represent a big challenge for the modern healthcare organisations (Beasley and Hirst, 2004).

This study used *in vitro* and *ex vivo* assays to investigate the effects and potential mechanisms by which estrogen can affect the clearance of MRSA and/or *P. aeruginosa*. Previous work suggests the sex steroid hormones, particularly estrogen and testosterone, play a role in regulating host responses to infection. Findings indicate testosterone acts as an immunosuppressor, whereas estrogen activates the

immune system (Cutolo *et al.*, 2004). These effects are linked with sexual dimorphism found in bacterial infections, where males (human and animal) are more vulnerable to bacterial infections than females (Klein and Roberts, 2010). In addition, changes in estrogen levels during menstrual cycles and pregnancy can influence bacterial infections (García-Gómez *et al.*, 2012). Administration of sex steroid hormones, such as estrogen, in models of ovariectomized mice can control the progression of bacterial infections, acting as a supplement to antibiotic therapies. In line with this, some bacteria have developed potential mechanisms to degrade sex steroid hormones or reduce their effectiveness (García-Gómez *et al.*, 2012).

3.1.2 Macrophages and Wound Healing

Macrophages are amongst the main inflammatory cell types present in age-related impaired wounds where inflammation is typically pronounced and prolonged. The phenotype of macrophages changes during the phases of wound healing (Mosser and Edwards, 2010). Following injury, pro-inflammatory macrophages, known as “M1” macrophages, arrive at the wound site in order to remove foreign invaders such as bacteria, debris and dead cells. During acute wound healing, as the tissue starts to heal, M1 macrophages undergo a process of conversion into macrophages that promote anti-inflammatory effects, referred to as “M2” macrophages, to resolve inflammation (Koh and DiPietro, 2011). Macrophages also have key roles in vascularisation, arranging themselves nearby restored blood vessels in order to support their fusion and stability (Ogle *et al.*, 2016). Macrophages release matrix metalloproteinases (MMPs) to breakdown the provisional extracellular matrix prior to undergoing apoptosis in order to bring about the deposition of mature matrix in the dermis as the wound progresses through the proliferative phase of wound repair (Vannella and Wynn, 2017). However, in non-healing wounds, macrophages persist in a pro-inflammatory state and the transition to an anti-inflammatory phenotype is blocked (Hesketh *et al.*, 2017).

Macrophages in the skin are either derived from a resident tissue-macrophage population established before birth or from circulating bone marrow-derived

monocytes that are recruited to the site of injury and differentiate into macrophages (Malissen *et al.*, 2014; Vannella and Wynn, 2017). When the wound site is infiltrated by microorganisms such as bacteria, monocyte-derived macrophages are systemically recruited within 24 hours post-injury in order to carry out phagocytosis (Minutti *et al.*, 2017).

Tissue-resident macrophages serve as early indicators of an injury or foreign invading pathogens (Malissen *et al.*, 2014) by recognizing PAMPs such as LPS via binding to TLRs. Responses to PAMPs include initial recruitment of neutrophils to the wound area within minutes of the injury to eliminate foreign invaders (Malissen *et al.*, 2014). Monocytes are subsequently recruited to the wound area to intensify the inflammatory response and become the predominant inflammatory cell type at around 3-7 days post-injury (Davies *et al.*, 2013). Dermal tissue-resident macrophages can be identified by numerous surface markers, such as CD64, CD11c, CD14, CD16, CD68, CD71 and CCR5 (Malissen *et al.*, 2014). Close to the resolution of injury, dermal tissue macrophages self-renew, and eliminate apoptotic cells as the tissue returns to homeostasis (Davies *et al.*, 2013).

Dendritic cells in the skin can also be derived from monocytes (e.g. Langerhans cells). Langerhans cells and macrophages share many similarities such as expressing similar surface markers (e.g. F4/80, CD14, and IL-10), which makes it difficult to distinguish between these two types of cell (Malissen *et al.*, 2014). Many researchers consider Langerhans cells as a type of tissue-resident macrophage, as they also have a very similar gene expression profile (Davies *et al.*, 2013; Doebel *et al.*, 2017; Minutti *et al.*, 2017). Interestingly, it has been reported that there is a link between healing diabetic foot ulcers and augmented numbers of Langerhans cells (Stojadinovic *et al.*, 2013). However, the exact function of Langerhans cells in wound repair (particularly impaired wound healing) has not been clarified yet (Stojadinovic *et al.*, 2013).

3.1.3 Effect of Estrogen on Macrophages

Estrogen is believed to be a key player in regulating immune responses against bacterial infections (Cutolo *et al.*, 2004). Known for its anti-inflammatory

properties, estrogen is reported to resolve excessive inflammation by directly effecting inflammatory responses of monocytes and macrophages (Ashcroft *et al.*, 2003; Guo and DiPietro, 2010). Previous research reported that ovariectomized mice expressed high numbers of macrophages in injured tissues compared to normal healthy mice during the wound healing process (Ashcroft *et al.*, 2003).

Macrophage migration inhibitory factor (MIF) is a pro-inflammatory cytokine, produced by many cell types (Calandra and Roger, 2003). MIF induces the production of other cytokines and growth factors by inflammatory cells such as TNF- α , IL-1 and IL-6 (Ashcroft *et al.*, 2003; Hardman *et al.*, 2005). MIF is a key pro-inflammatory mediator inhibited by estrogen during wound healing (Ashcroft *et al.*, 2003; Hardman *et al.*, 2005). MIF is upregulated in human and murine impaired wounds *in vivo* (Ashcroft *et al.*, 2003; Hardman *et al.*, 2005; Emmerson *et al.*, 2009), whereas MIF is decreased in wounds following exposure to estrogen (Ashcroft *et al.*, 2003). Recent research indicated that estrogen inhibits MIF production by tissue macrophages (Ashcroft *et al.*, 2003; Hardman *et al.*, 2005).

TNF- α is a key pro-inflammatory cytokine released by activated macrophages and is associated with chronic wounds in the elderly (Emmerson *et al.*, 2009). After a pathogenic infection, bacterial clearance is reliant on effective regulation of TNF- α , and abnormal expression of the cytokine may cause substantial morbidity and mortality (Billiau and Vandekerckhove, 1991). Uncontrolled TNF- α production is also associated with the progression of several autoimmune diseases, such as rheumatoid arthritis, which are more dominant in females than males (Zandman-Goddard *et al.*, 2007). TNF- α wound levels are reduced in healthy premenopausal females and young males in comparison with postmenopausal females and elderly males (Vural *et al.*, 2006). Research suggests that estradiol suppresses the production of TNF- α in murine macrophages (Deshpande *et al.*, 1997; Srivastava *et al.*, 1999).

Long-time exposure of *ex vivo* macrophages to estrogen augments the expression of iNOS and the production of numerous cytokines, particularly IL-1 β , IL-12 and IL-6 following TLR4 activation by LPS. Other studies have reported anti-inflammatory effects of estrogens on macrophages *in vitro* (Deshpande *et al.*, 1997; Ghisletti *et*

al., 2005). For example, a 2-hour estrogen treatment of macrophages (RAW 264.7 cell line) *in vitro* caused a substantial reduction in the expression of the inflammatory mediators IL-1 α , IL-6, and TNF- α by splenic macrophages following TLR4 activation (Deshpande *et al.*, 1997; Ghisletti *et al.*, 2005).

In summary, estrogen accelerates wound healing and is associated with a reduced inflammatory response. Estrogen inhibits the recruitment of monocytes to the wound and reduces the production of macrophage-derived pro-inflammatory mediators. However, the precise role estrogen plays in macrophage-mediated bacterial clearance has not been elucidated to date. Thus, this study used *in vitro* and *ex vivo* models of host-pathogen interaction to investigate estrogen-mediated phagocytosis of bacteria by human U937- and HPBM-derived macrophages. Macrophage-mediated phagocytosis of two major wound pathogens, MRSA and *P. aeruginosa*, were assessed under *in vitro* and *ex vivo* conditions that model estrogen levels in the elderly (estrogen deprivation), physiological concentrations in young adults and following exogenous estrogen supplementation. U937 and HPBM-derived macrophages are routinely used as flexible models of macrophages to study numerous aspects of the inflammatory response; including chemotaxis, phagocytosis, apoptosis and cytokine production (Hall, 2017).

3.2 Aim and Objectives

3.2.1 Aim

To investigate the effects of age-related changes in estrogen levels on the phagocytic function of M0 macrophages using *in vitro* and *ex vivo* models of host-pathogen interactions.

3.2.2 Objectives

- To compare the phagocytic function of M0 U937- and HPBM-derived macrophages under *in vitro* and *ex vivo* conditions of physiological (typical of youth) and supraphysiological levels of estrogen, estrogen deprivation and the absolute absence of estrogen.

- To determine the effect of estrogen on the phagocytosis of Gram positive (MRSA) and Gram negative (*P. aeruginosa*) wound pathogens by U937- and HPBM-derived M0 macrophages.
- To visualise estrogen-mediated internalisation of Gram positive (MRSA) and Gram negative (*P. aeruginosa*) bacteria by U937- and HPBM-derived M0 macrophages.

3.3 Methods

3.3.1 Generation of U937- and HPBM-Derived M0 Macrophages

The differentiation of U937 and HPBM monocytes into macrophages was confirmed by flow cytometry as described in Section 2.2.6.1. U937 and HPBM monocytes were cultured as described in Sections 2.2.3. Cells were treated with/without 50 ng/mL PMA as described in Section 2.2.4 to induce differentiation of monocytes into macrophages. Cells were then stained with FITC-conjugated anti-CD11c antibody as described in 2.2.6.1. The expression of CD11c (relative MFI and % fluorescence) on untreated control monocytes was then compared with PMA-treated M0 macrophages by flow cytometry as described in Section 2.2.6.1.

3.3.2 Effect of Estrogen on Bacterial Growth

Before conducting host-pathogen assays to test the effect of estrogen on the macrophage-mediated phagocytosis of MRSA and PAO1, it was important to investigate its direct effect on the growth of both pathogens.

MRSA 11 and PAO1 were cultured in NB overnight as described in Section 2.2.7. Neat broths were centrifuged at 3500 rpm for 10 minutes and the bacterial cell pellets washed twice with saline before re-suspension at 1×10^4 CFU/mL (see Section 2.2.7) in 1 mL saline alone (zero M estrogen) or 1mL saline containing either 1×10^{-8} M estrogen or 1×10^{-7} M BSA (negative control). Bacterial suspensions were then incubated at 37 °C on an orbital shaker for 3 hours. Aliquots (50 μ L) of each sample were then inoculated onto NA plates in duplicate and incubated at 37 °C overnight. The number of CFU formed on NA plates were counted in order to calculate the growth of bacteria in each sample.

3.3.3 Effect of Estrogen on the Clearance of MRSA/PAO1 by M0 Macrophages

3.3.3.1 Phenol Red-Free Medium Controls

In order to determine whether the phenol red component of cell culture medium had an effect on estrogen-mediated phagocytosis of MRSA/PAO1 by M0

macrophages, *in vitro* host-pathogen assays were conducted using RPMI medium containing/lacking phenol red. U937 monocytes were cultured in phenol red-free RPMI medium or RPMI complete medium using methods described in Section 2.2.3. M0 macrophages generated in Section 3.3.1 were then treated with physiological estrogen (1×10^{-8} M), prepared in either phenol-free RPMI medium or RPMI complete medium, for 24 hours and then incubated with MRSA or PAO1 for 3 h. Recovered bacterial colonies were then determined as described in Section 2.2.8.1.

3.3.3.2 Multiplicity of Infection (MOI)

The effect of estrogen on phagocytosis over a range of starting bacterial inoculations was assessed for both MRSA or PAO1. U937 M0 macrophages generated in Section 3.3.1 were treated with 1×10^{-8} M estrogen for 24 hours prior to incubation for 3 hours with different concentrations (1×10^3 CFU, 1×10^4 CFU, 1×10^5 CFU and 1×10^6 CFU) of MRSA or PAO1 to give initial bacteria: macrophage ratios of 0.001, 0.01, 0.1 and 1. Recovered bacterial colonies were then determined as described in Section 2.2.8.1.

3.3.3.3 *In vitro* and *Ex Vivo* Host-Pathogen Interaction Assays

U937 and HPBM M0 macrophages generated in Section 3.3.1 were treated in wells with different concentrations (1×10^{-7} M, 1×10^{-8} M, 1×10^{-9} M and zero M) of estrogen for 24 hours as described in Sections 2.2.2 and 2.2.8.1. Cells were incubated with MRSA or PAO1 (1×10^4 CFU) for 3 hours and bacterial recovery was determined as described in Section 2.2.8.1.

3.3.4 Interaction of Estrogen-Treated U937 M0 Macrophages with GFP- *S.aureus*

U937 M0 macrophages generated in Section 3.3.1 were incubated in flasks with different concentrations (1×10^{-7} M, 1×10^{-8} M, 1×10^{-9} M and zero M) of estrogen for 24 hours as described in Sections 2.2.2 and 2.2.8.1. Cells were then incubated with GFP- *S. aureus* (1×10^5 CFU/mL) for 3 hours prior to detection of GFP signal as described in Section 2.2.9.

3.3.5 Gentamicin Protection Assay

In order to confirm that internalisation of MRSA and PAO1 by U937 and HPBM M0 macrophages was leading to bacterial killing rather than simply bacterial sequestration, a gentamicin protection assay was conducted to determine bacterial survival within macrophages using methodologies described in Sections 2.2.10 and 3.3.1.

The gentamicin protection assay is commonly used in host-pathogen research due to the incapability of gentamicin to penetrate the membrane of eukaryotic cells (Elsinghorst, 1994). The gentamicin protection assay was performed to investigate the effect of estrogen on the killing of intracellular MRSA and PAO1 as gentamicin was previously described to eradicate extracellular bacteria, but not intracellular bacteria due to its poor ability to permeate the macrophage membrane (Hamad *et al.*, 2010; Hockenberry *et al.*, 2016). Moreover, incubating cells with gentamicin kills bacteria that fail to penetrate cells, and any bacterial recovery obtained following lysis of the macrophages must be due to bacterial survival within the macrophages. The gentamicin protection assay is a very sensitive technique that can allow the detection of even single colonies of bacteria in cells (Elsinghorst, 1994).

3.3.6 Visualising Host-Pathogen Interactions

Internalisation of GFP *S. aureus* or mCherry- *P. aeruginosa* by U937 and HPBM M0 macrophages, with/without estrogen treatment, was confirmed by fluorescence microscopy as described in Section 2.2.11.1. Host-pathogen interactions between U937 or HPBM M0 macrophages and MRSA/PAO1, with/without estrogen treatment, were visualised by SEM as described in Section 2.2.11.2.

3.4 Results

3.4.1 Differentiation of Monocytes into M0 Macrophages

According to previous research, monocytes (including U937 monocytes) can differentiate into macrophage-like cells via stimulation with PMA through PKC activation (Gidlund *et al.*, 1981; Koeffler, 1983; Kiley and Parker, 1995).

Figure 3.1 illustrates alterations in the morphology of U937 cells after treatment with PMA at 50 ng/mL for 24 hours. Round floating monocytes were transformed into adherent cells characterised by increased adhesion, formation of clumps and ruffled cell membranes.

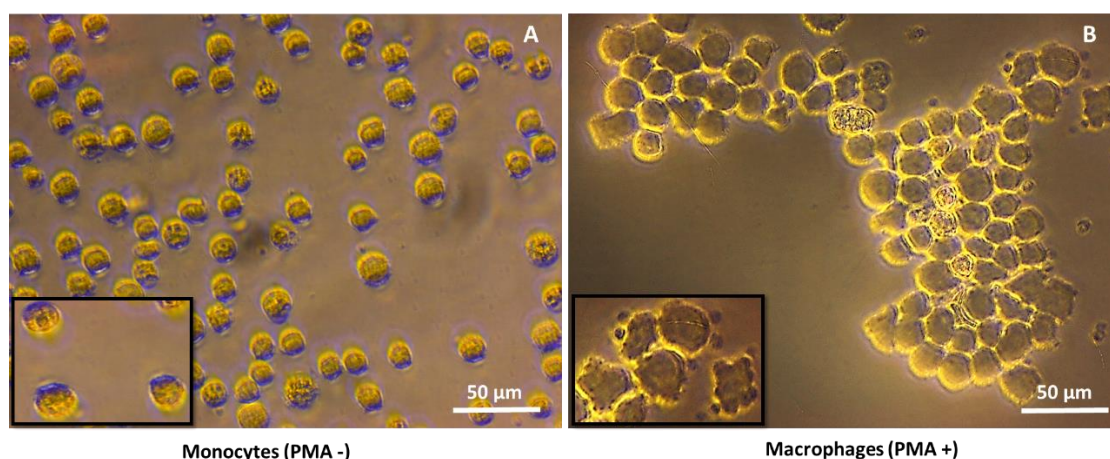


Figure 3.1. Morphological changes of U937 monocytic cells after treatment with 50 ng/mL PMA for 24 hours. Round floating monocytes (A) were differentiated into adherent cells (B) that formed clumps and had ruffled cell membranes. Images were taken using an Axiovert 40C inverted phase contrast microscope. Inserts show enlargements of highlighted cells.

The conversion of U937 and HPBM monocytes into a distinct population of macrophages was confirmed via detection of the FITC-conjugated anti-human CD11c surface marker (Rios de la Rosa *et al.*, 2017) by flow cytometry (Section 2.2.6.1). U937 PMA-differentiated cells were almost exclusively (99.6%) CD11c⁺ (Figure 3.2) whereas untreated control U937 monocytes predominantly lacked the CD11c surface marker (0.2% CD11c⁺). The CD11c MFI was significantly ($P < 0.001$) higher in PMA-treated U937 cells compared to untreated control U937 monocytes, confirming PMA transformed U937 monocytes into adherent U937 M0 macrophages.

There was high expression of CD11c by PMA-differentiated HPBM cells (76% CD11c⁺), whereas the CD11c macrophage surface marker was almost absent (0.3% CD11c⁺) in untreated control HPBM (Figure 3.3). The MFI of CD11c was significantly ($P<0.05$) greater in PMA-treated HPBM cells compared to untreated control HPBM, confirming the transformation of HPBM into adherent M0 macrophages had occurred following PMA treatment.

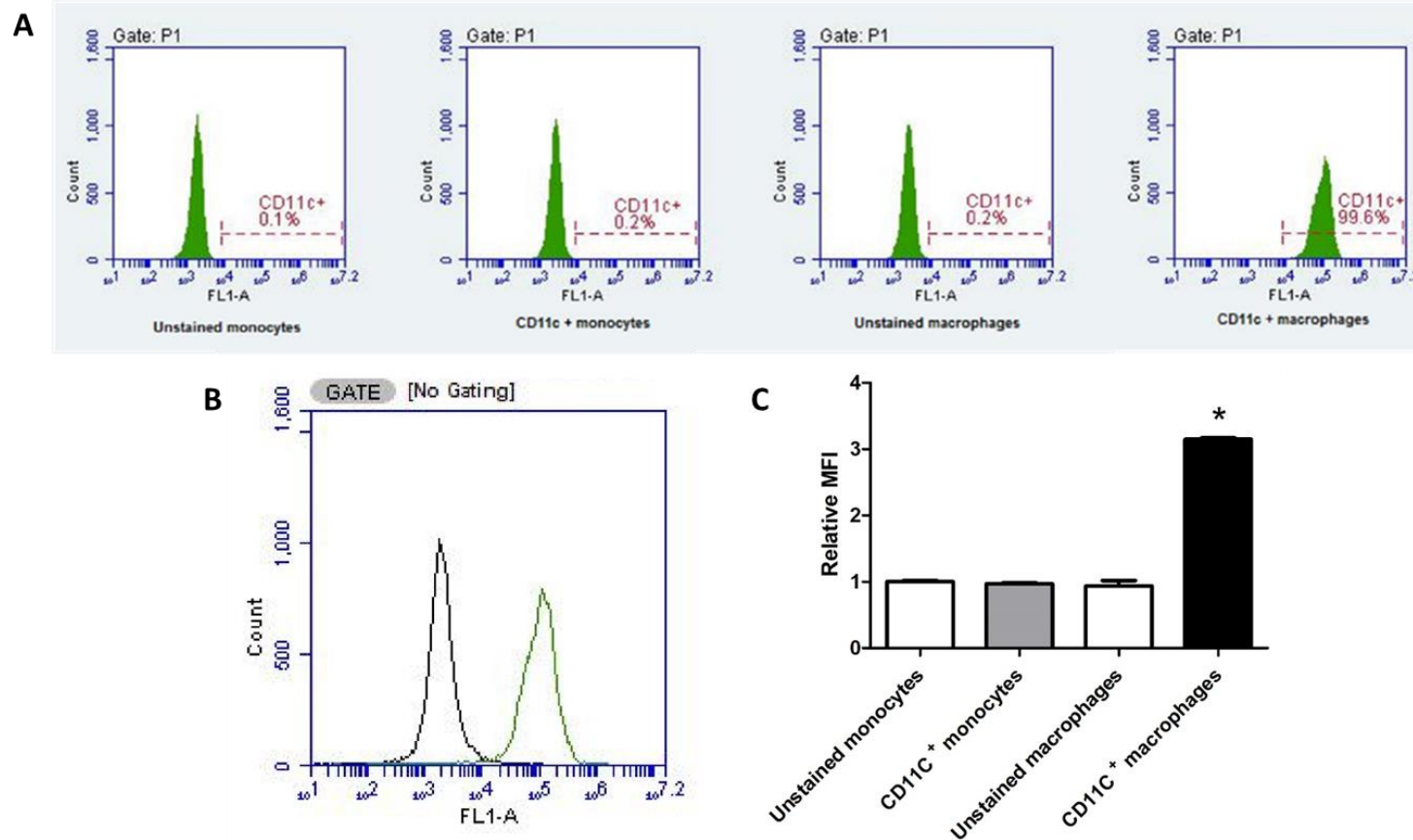


Figure 3.2. Flow cytometry confirming U937 monocytes differentiation into Mo macrophages. The conversion of U937 monocytic cells into macrophages was established via detection of the CD11c surface marker by flow cytometry. Phorbol 12-myristate 13-acetate (PMA)-differentiated cells almost exclusively expressed the CD11c macrophage marker, whereas monocytes predominantly lacked the CD11c surface marker (A). Two distinct populations of U937 cells were detected (B) with significantly ($P < 0.01$) higher median fluorescence intensity (MFI) from U937 PMA-treated cells compared to undifferentiated U937 monocytes (C). Data are presented as average of $n = 3$ experiments. * Indicates significant difference (One-way ANOVA) in MFI ($P < 0.01$). Error bars represent the standard error of the mean (SEM).

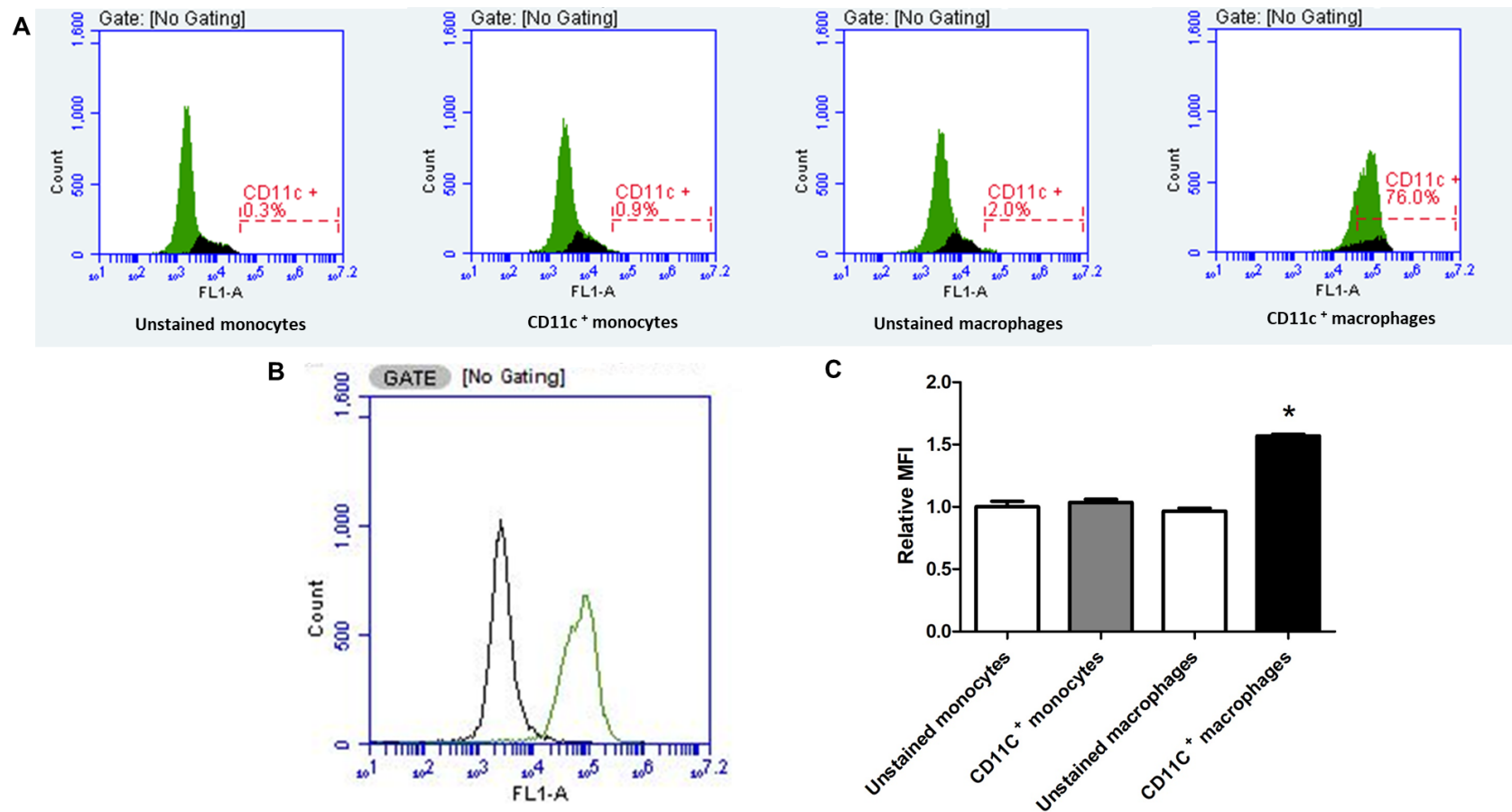


Figure 3.3 Flow cytometry confirming HPBM differentiation into M0 macrophages. The differentiation of HPBM into macrophages was assessed via detection of the CD11c surface marker by flow cytometry. Phorbol 12-myristate 13-acetate (PMA)-differentiated cells highly expressed CD11c whereas monocytes predominantly lacked the CD11c surface marker (A). Two distinct cell populations were detected (B) with significantly ($P < 0.05$) higher median fluorescence intensity (MFI) observed on HPBM PMA-treated cells compared to untreated monocytes (C). Data are presented as average of $n = 3$ experiments. * Indicates significant difference (One-way ANOVA) in MFI ($P < 0.05$). Error bars represent the standard error of the mean (StEM).

3.4.2 Effect of Estrogen on the Growth of MRSA and PAO1

The effect of estrogen (1×10^{-7} M) on the growth of MRSA and PAO1 colonies after 3 hours incubation was investigated compared to a BSA (1×10^{-7} M) negative control and a untreated saline control. Estrogen had no significant effect on the MRSA or PAO1 growth ($P>0.05$) (Figure 3.4A and Figure 3.4B) compared to the controls. These results indicated that estrogen had no direct effect on the growth of MRSA or PAO1 compared to the controls.

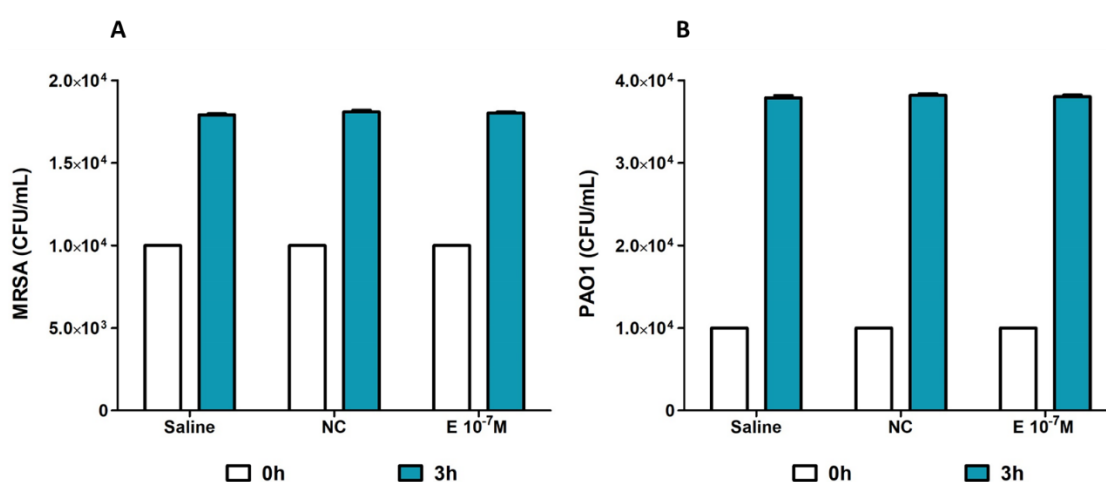


Figure 3.4. Effect of estrogen on MRSA and PAO1 growth. Estrogen (1×10^{-7} M) had no significant ($P>0.05$) effect on the growth of MRSA (A) or PAO1 (B) compared to the 1×10^{-7} M bovine serum albumin (BSA) negative control (NC) or the untreated saline control. Data are presented as average of $n = 24$ experiments.

3.4.3 The Effect of Phenol Red on Bacterial Clearance

The effect of phenol red on the clearance of MRSA and PAO1 by U937 M0 macrophages with/without estrogen treatment was investigated. The findings confirmed that phenol red in the RPMI cell culture medium had no significant effect on the MRSA (A) and PAO1 (B) recovery compared to RPMI medium lacking phenol red ($P>0.05$), regardless of the presence of estrogen (Figure 3.5). Physiological (1×10^{-8} M) estrogen levels significantly decreased MRSA and PAO1 recovery compared to the negative control ($P<0.05$), both in the presence or absence of phenol red.

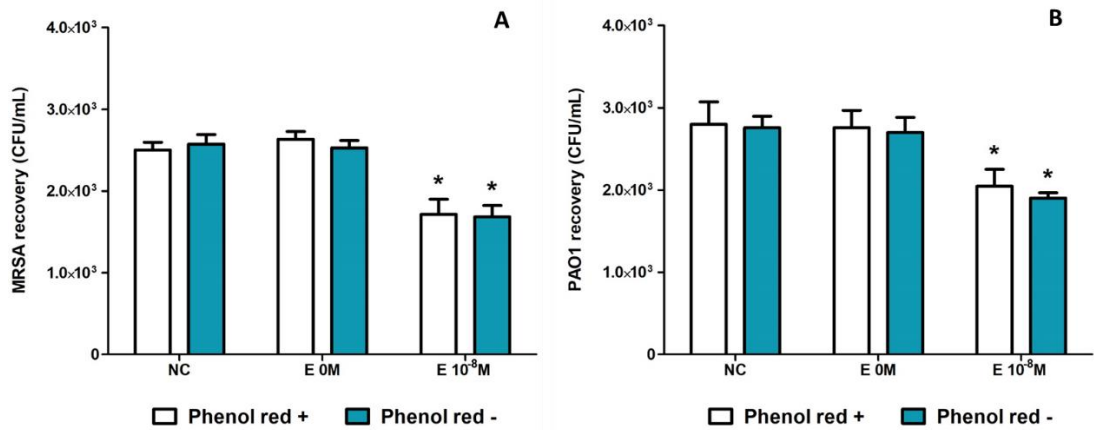


Figure 3.5. Effect of phenol red on bacterial clearance by M0 macrophages. Phenol red in cell culture medium had no significant effect on the recovery of MRSA (A) or PAO1 (B) compared to the absence of phenol red ($P > 0.05$), regardless of the presence/absence of estrogen (1×10^{-8} M). Physiological (1×10^{-8} M) estrogen levels significantly ($*$: $P < 0.05$) decreased the MRSA (A) and PAO1 (B) recovery compared to the negative control (NC), with or without phenol red. Data are an average of $n = 10$ experimental repeats with error bars representing the standard error of the mean (StEM).

3.4.4 Multiplicity of Infection (MOI)

The effect of physiological estrogen (1×10^{-8} M) on the uptake of MRSA by M0 macrophages, *in vitro*, was investigated. Cells were incubated with a range of initial MRSA inoculations (1×10^3 CFU, 1×10^4 CFU, 1×10^5 CFU and 1×10^6 CFU) (Figure 3.6). The absence of estrogen had no significant effect ($P > 0.05$) on the MRSA recovery compared to negative control. Physiological estrogen significantly decreased ($P < 0.05$) the MRSA recovery, in comparison with the BSA control with all MOIs. The bacterial recovery of MRSA significantly increased ($P < 0.05$) following estrogen (1×10^{-8} M) supplementation as the MOI [bacteria : macrophage ratio] increased from 0.001 to 1. However, the MRSA recovery followed a similar pattern throughout, regardless of the MOI ratio, with estrogen consistently increasing ($P < 0.05$) MRSA clearance at all MOIs.

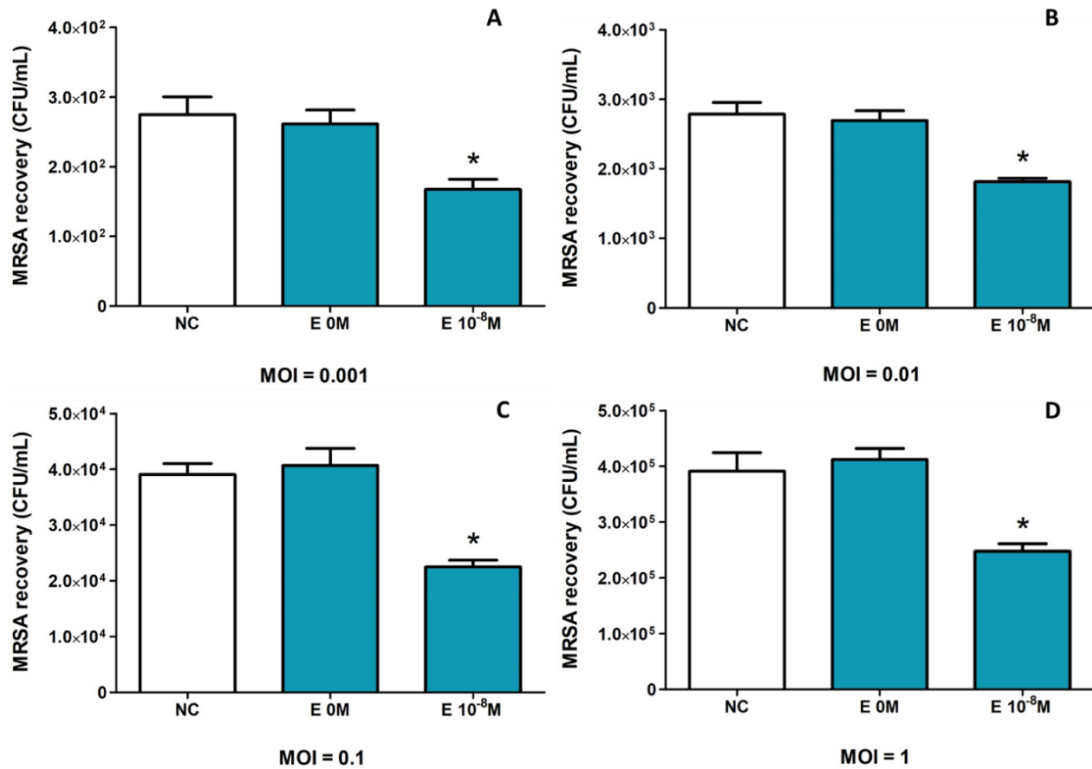


Figure 3.6. Effect of physiological estrogen on the recovery of MRSA by M0 macrophages *in vitro*. Cells were incubated with MRSA at MOIs of 0.001 (A), 0.01 (B), 0.1 (C) and 1 (D). Following treatment of macrophages with physiological estrogen (1×10^{-8} M), MRSA recovery significantly decreased ($P < 0.05$), at all MOIs, in comparison with the absence (zero M) of estrogen and the negative control (NC). Data represent mean recovery (CFU/mL) \pm standard error of the mean (StEM), $n = 10$. *: $P < 0.05$, indicate significant differences in the MRSA/PAO1 recovery compared to the NC.

Figure 3.7 illustrates the effect of physiological estrogen (1×10^{-8} M) on the internalisation of PAO1 by macrophages. After incubating the macrophages with a varied range of initial PAO1 colonies (1×10^3 CFU, 1×10^4 CFU, 1×10^5 CFU and 1×10^6 CFU), the absence of estrogen had no significant effect ($P > 0.05$) on the recovery of in comparison with the BSA control. Interestingly, physiological estrogen significantly decreased ($P < 0.05$) the PAO1 recovery, in comparison with the NC control at all MOIs. The PAO1 recovery significantly increased ($P < 0.05$) following estrogen (1×10^{-8} M) supplementation as the MOI [bacteria : macrophage ratio] increased from 0.001 to 1. However, the PAO1 recovery followed a comparable pattern throughout, regardless of the MOI ratio, with estrogen consistently increasing ($P < 0.05$) PAO1 clearance at all MOIs.

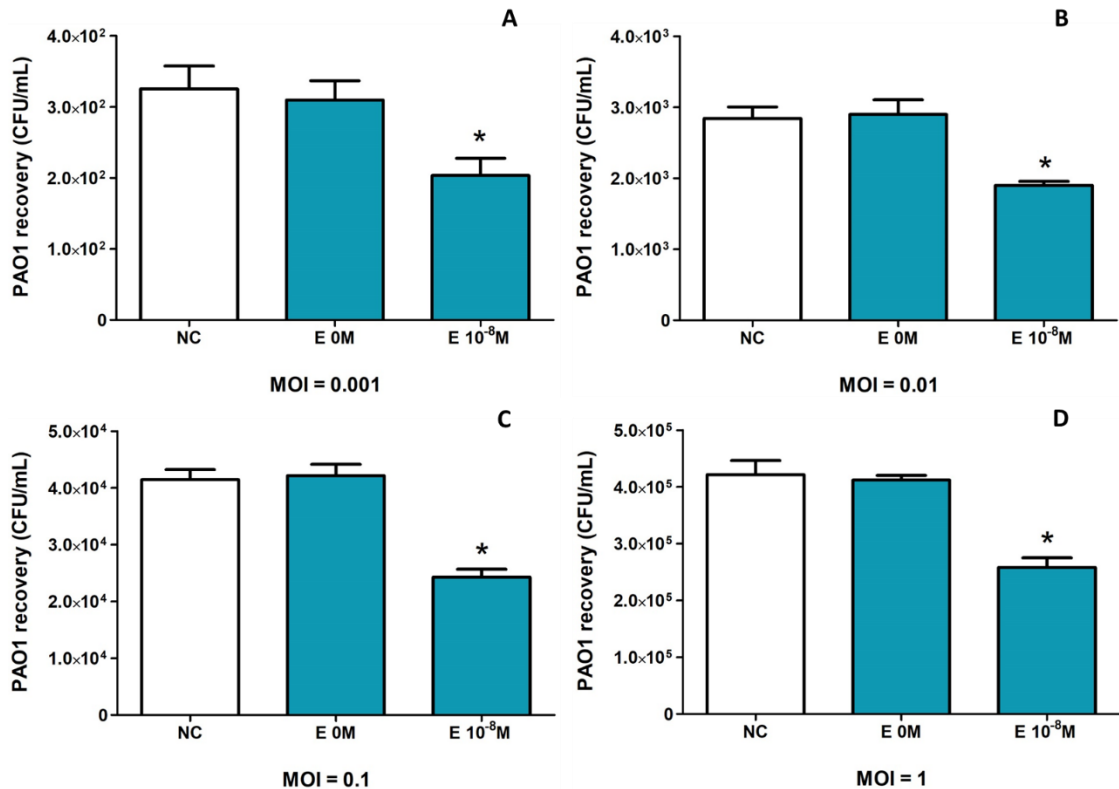


Figure 3.7. Physiological estrogen decreases the recovery of PAO1 by M0 macrophages *in vitro*. U937-derived M0 macrophages were incubated with PAO1 at MOIs of 0.001 (A), 0.01 (B), 0.1 (C) and 1 (D). Physiological estrogen (1×10^{-8} M) significantly decreased ($P < 0.05$) the PAO1 recovery at all MOIs, in comparison with the absence (zero M) of estrogen and the negative control (NC). Data represent mean recovery (CFU/mL) \pm standard error of the mean (StEM), $n = 10$. *: $P < 0.05$, indicate significant differences in the MRSA/PAO1 recovery compared to the NC.

3.4.5 Effect of Estrogen on MRSA and PAO1 Clearance

3.4.5.1 Effect of Estrogen on MRSA and PAO1 Clearance by U937 Macrophages

The effect of estrogen on the clearance of MRSA and PAO1 by U937 M0 macrophages was investigated. The absolute absence of estrogen (zero M) showed no significant ($P > 0.05$) change in the number of recovered MRSA or PAO1 colonies (Figure 3.8A and Figure 3.8B) compared to the BSA negative control (NC).

Estrogen significantly ($P < 0.01$) reduced the recovery of MRSA colonies compared to the negative control in a dose-dependent manner. The MRSA recovery (Figure 3.8A) was significantly ($P < 0.01$) reduced from 1886 CFU/mL in the NC to 1759 CFU/mL, 1295 CFU/mL and 936 CFU/mL with estrogen at a concentrations of 1×10^{-9} M, 1×10^{-8} M and 1×10^{-7} M respectively. The MRSA recovery decreased significantly

($P < 0.01$) between increasing estrogen concentrations, with physiological estrogen concentrations (1×10^{-8} M) significantly reducing MRSA recovery compared to estrogen deprivation, and supraphysiological estrogen concentrations (1×10^{-7} M) significantly reducing MRSA recovery compared to physiological estrogen levels.

Physiological estrogen (1×10^{-8} M) and supraphysiological estrogen (1×10^{-7} M) significantly ($P < 0.05$) reduced the PAO1 recovery (Figure 3.8.B) from 2410 CFU/mL in the NC to 2105 CFU and 1960 CFU respectively. The PAO1 recovery also decreased significantly ($P < 0.05$) between physiological and supraphysiological estrogen levels.

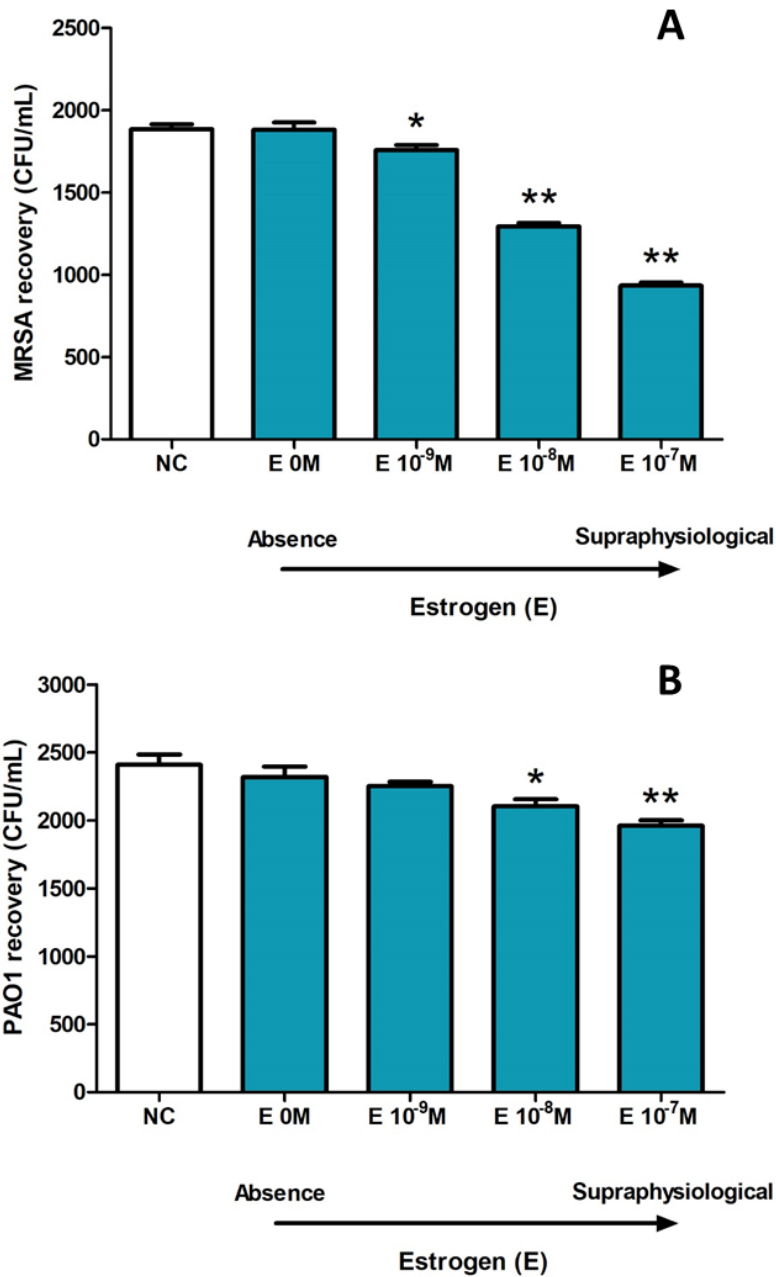


Figure 3.8. Estrogen enhances the uptake of MRSA11 (A) and PAO1 (B) by U937 M0 macrophages. Following treatment of U937 M0 macrophages with estrogen (1×10^{-7} M, 1×10^{-8} M) MRSA (A) and PAO1 (B) recovery significantly decreased in a concentration-dependant manner in comparison to the absence (zero M) of estrogen and the negative control (NC). Data represent mean recovery (CFU/mL) \pm standard error of the mean (StEM), $n = 48$. *: $P < 0.05$, **: $P < 0.01$ indicate significant differences in the MRSA/PAO1 recovery compared to the NC.

3.4.5.2 Effect of Estrogen on MRSA and PAO1 Clearance by *Ex Vivo* M0 Macrophages

Host-pathogen assays were repeated with human peripheral blood monocytes (HPBM) in order to determine whether estrogen has similar effects on the MRSA

and PAO1 recovery with macrophages derived from primary human monocytes. In the absence of estrogen (zero M estrogen) there was no significant difference ($P>0.05$) in the clearance of MRSA or PAO1 compared to the negative control (Figure 3.9.A and Figure 3.9.B). In contrast, estrogen treatment significantly reduced ($P<0.05$) MRSA recovery compared to the negative control in a dose-dependent manner. Physiological (1×10^{-8} M) and particularly supraphysiological (1×10^{-7} M) levels of estrogen were highly effective at increasing MRSA clearance, significantly decreasing ($P<0.01$) MRSA recovery (1775 CFU/mL, 1246 CFU/mL respectively) compared to the negative control (2692 CFU/mL). Estrogen had similar effects on the clearance of PAO1 when treated with physiological (1×10^{-8} M) or supraphysiological (1×10^{-7} M) levels, significantly ($P<0.05$) reducing PAO1 recovery (2252 CFU/mL and 1817 CFU/mL respectively) compared to the negative control (2631 CFU/mL).

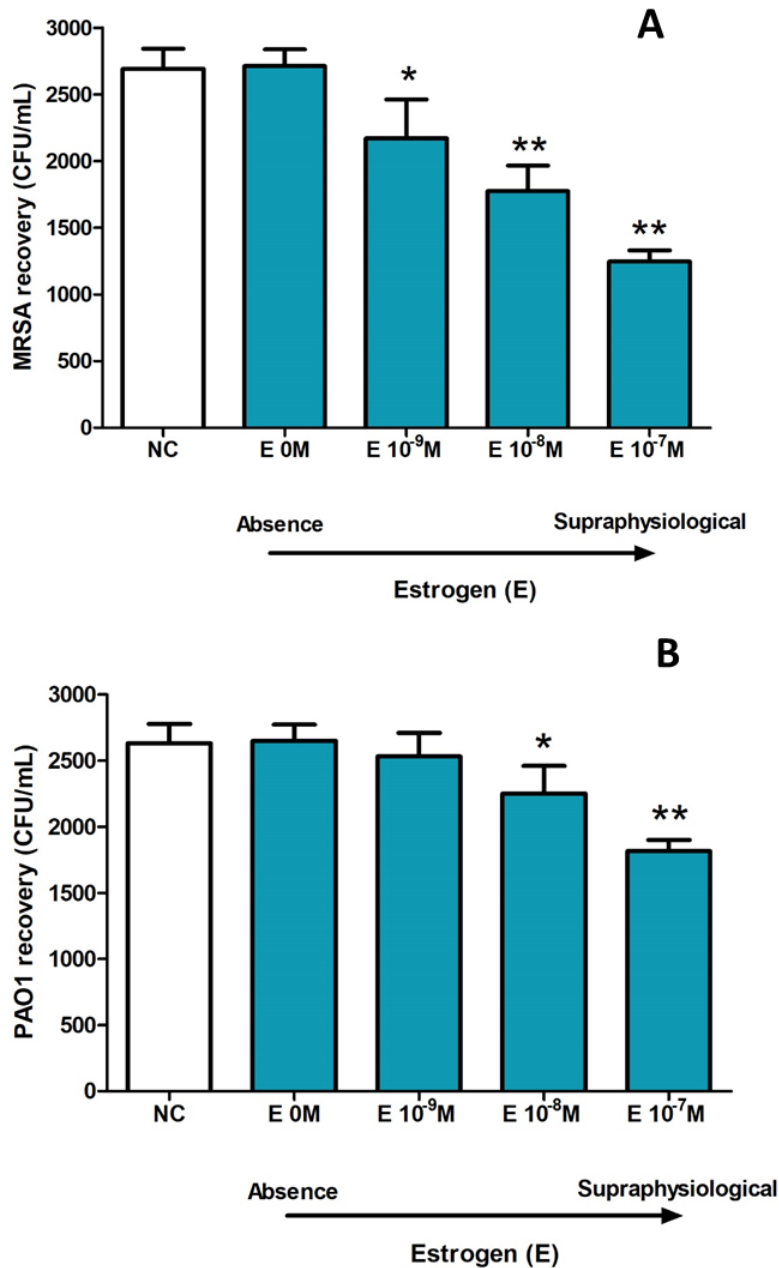


Figure 3.9. Estrogen enhances the uptake of MRSA11 (A) and PAO1 (B) by HPBM M0 macrophages. Following treatment of HPBM M0 macrophages with estrogen (1×10^{-7} M, 1×10^{-8} M) the recovery of MRSA/PAO1 was significantly reduced in comparison to the negative control (NC). Error bars represent the mean \pm standard error of the mean (SEM), $n = 6$. *: $P < 0.05$, **: $P < 0.01$ indicate significant differences in the MRSA/PAO1 recovery compared to the NC.

3.4.6 GFP *S. aureus* Host-Pathogen Assay

3.4.6.1 Effect of Estrogen on The Recovery of GFP- *S. aureus*

The effect of estrogen on the recovery of fluorescent GFP-labelled *S. aureus* colonies was investigated. U937-derived M0 Macrophages were incubated with

GFP-*S. aureus* for 3 hours. The bacterial colonies were tracked via detection of GFP signals inside and outside the macrophages using a countess II fluorescent cell counter. Images (n = 10) of bacterial internalisation within macrophages were captured. The phagocyte density, number of internalised bacteria, total number of macrophages and total bacterial counts were determined from images in order to calculate the bacterial recovery, total number of phagocytes/mL and the ratio of bacterial internalisation.

Consistent with previous results, physiological and supraphysiological levels of estrogen significantly reduced the *S. aureus* recovery ($P < 0.05$) in a concentration-dependant manner in comparison with the absence of estrogen and BSA controls (Figure 3.10).

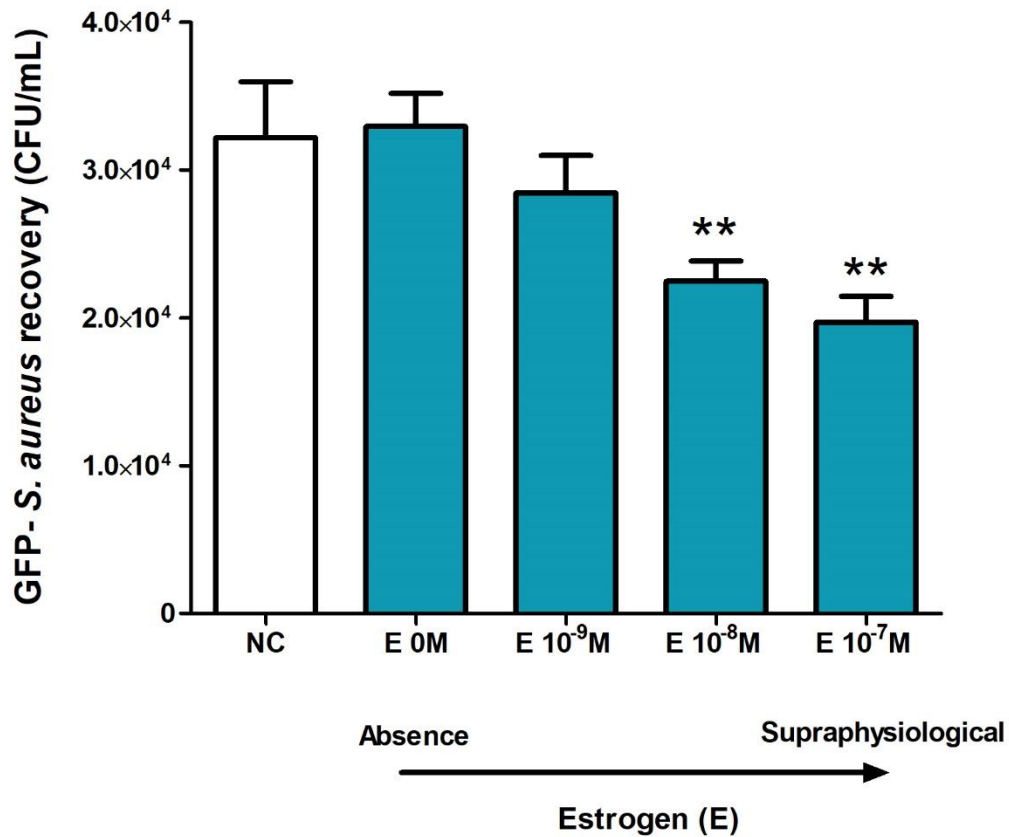


Figure 3.10. Estrogen increases the uptake of GFP-*S. aureus* by U937 M0 macrophages. Following treatment of U937 M0 macrophages with physiological and supraphysiological estrogen (1×10^{-9} M, 1×10^{-7} M), the *S. aureus* recovery was significantly reduced in a concentration-dependant manner in comparison to the absence (zero M) of estrogen and the BSA control (NC). Data represent mean recovery (CFU/mL \pm standard error of the mean (StEM), $n = 10$). **: $P < 0.01$ indicate significant differences in the GFP-*S. aureus* recovery compared to the NC.

3.4.6.2 Effect of Estrogen on Phagocyte Density

The effect of estrogen on the number of active phagocytes after the 3-hour period of host-pathogen interaction was investigated (Figure 3.11). Stimulation of cells with 1×10^{-8} M or 1×10^{-9} M estrogen did not have any significant effect ($P=0.16$ and $P=0.99$ respectively) on phagocyte density (3.0×10^5 phagocytes/mL and 2.8×10^5 phagocytes/mL respectively) compared to the negative control (2.3×10^5 phagocytes/mL) (Figure 3.11). However, supraphysiological (1×10^{-7} M) levels of estrogen significantly increased ($P < 0.01$) the number of active phagocytes (3.7×10^5 phagocytes/ mL) in comparison to the negative control (NC).

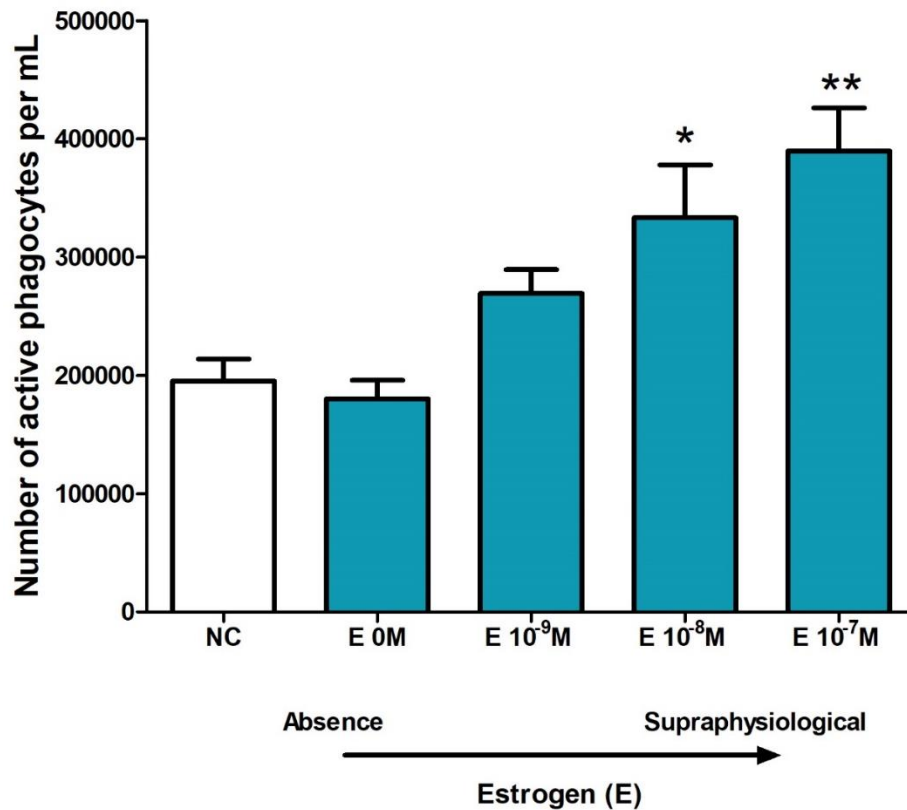


Figure 3.11. Estrogen Promotes an Increase in the Number of Active Phagocytes. Supraphysiological (1×10^{-7} M) levels of estrogen significantly (*: $P < 0.05$, **: $P < 0.01$) increased the detection of GFP signals inside macrophages compared to the negative control (NC). Data indicate mean number of active phagocytes per mL ($n = 8$) \pm standard error of the mean (StEM).

3.4.6.3 Effect of Estrogen on the Proportion of Total GFP *S. aureus* Internalised by U937-Derived Macrophages

Figure 3.12 illustrates the proportion of total GFP- *S. aureus* internalised by U937-derived macrophages following treatment with 17β -estradiol. When the cells were treated with zero M and 1×10^{-9} M estrogen, there was no significant ($P = 0.89$) effect on the proportion of GFP- *S. aureus* internalised (0.58) in comparison to the negative control (NC). However, when U937-derived macrophages were treated with physiological (1×10^{-8} M) or supraphysiological (1×10^{-7} M) levels of estrogen, there was a significant increase ($P < 0.05$) in the proportion of internalised bacteria (0.66 and 0.78 respectively) compared to the NC.

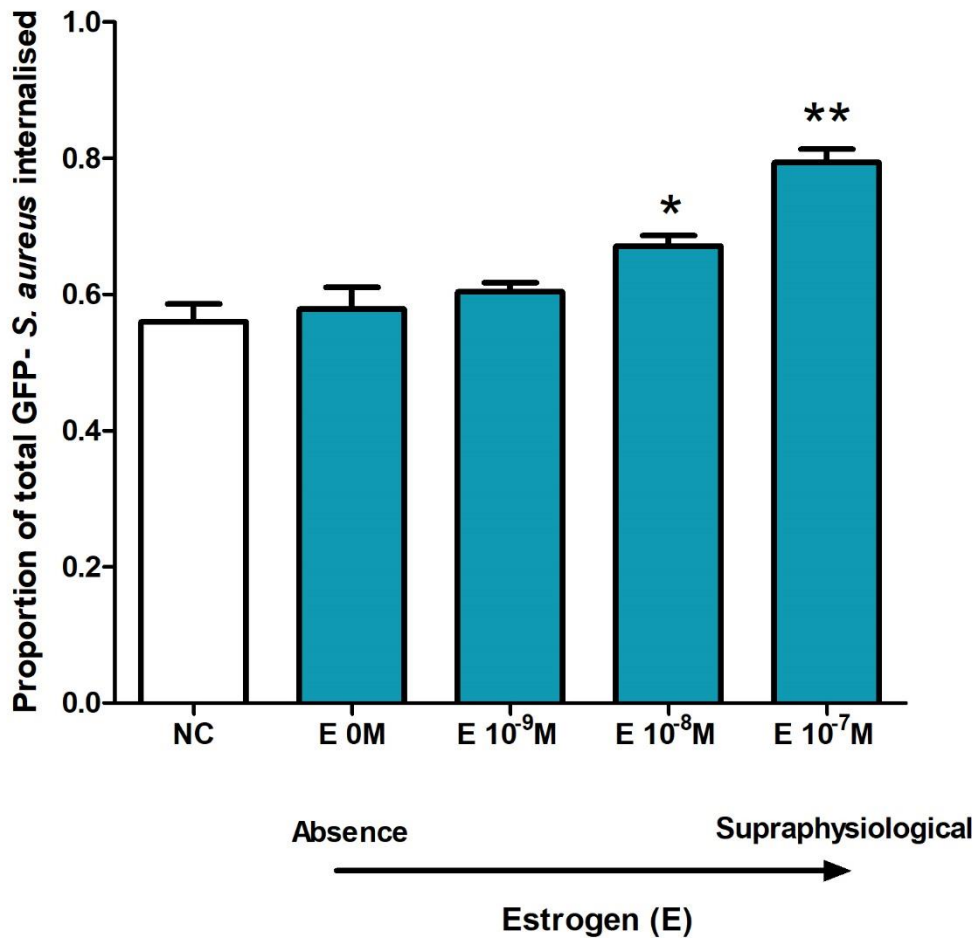


Figure 3.12. Effect of estrogen on the ratio of GFP- *S. aureus* internalisation. Data are presented as average of $n = 8$ experiments. * $P < 0.05$, **: $P < 0.01$ Indicate significant differences (One-way ANOVA) in bacterial recovery ($P < 0.05$). Error bars represent the StEM.

3.4.7 Effect of Estrogen on Bacterial (MRSA and *P. aeruginosa*) Killing by U937-Derived Macrophages.

In order to determine the effect of estrogen on bacterial killing by U937-derived macrophages, a gentamicin protection assay was performed. The gentamicin protection assay is commonly used to confirm the killing of internalised bacteria by phagocytes, or recover viable bacteria from inside phagocytes when pathogens are able to evade host cell responses and grow within phagocytes (Elsinghorst, 1994). The assay utilised gentamicin to eliminate non-internalised (extracellular) bacteria after a 3-hour period of host-pathogen interaction, followed by the lysis of washed macrophages to recover internalised bacteria. Gentamicin is unable to pass across the host cell membrane so only bacteria that have been successfully internalised

(phagocytosed) by immune cells but remain viable within phagocytes at the end of the host-pathogen interaction are recovered by the assay (Elsinghorst, 1994., Waldbeser *et al.*, 1994; Hess *et al.*, 2004; Oelschlaeger, 2010).

Similar to previous assays, physiological (1×10^{-8} M) and supraphysiological (1×10^{-7} M) levels of estrogen significantly ($P < 0.05$) decreased the recovery of viable internalised MRSA and PAO1 by U937 M0 macrophages in a dose-dependent manner (Figure 3.13) compared to the BSA (NC) and untreated controls. The data confirmed estrogen was not simply promoting internalisation of bacteria, but was also increasing the killing of phagocytosed MRSA and PAO1 by U937-derived macrophages. The absence of estrogen or estrogen deprivation (1×10^{-9} M) did not significantly ($P > 0.05$) effect the killing of internalised MRSA and PAO1 by U937-derived macrophages compared to the NC.

Physiological (1×10^{-8} M) and supraphysiological estrogen (1×10^{-7} M) significantly ($P < 0.05$) reduced the MRSA recovery (Figure 3.12.A) from 976 CFU/mL in the NC to 787 CFU and 692 CFU respectively. The PAO1 recovery also decreased significantly ($P < 0.05$) from 1019 CFU/mL in the BSA control to 822 CFU/mL and 692 CFU/mL following treatment with physiological (typical of youth) and supraphysiological levels of estrogen respectively.

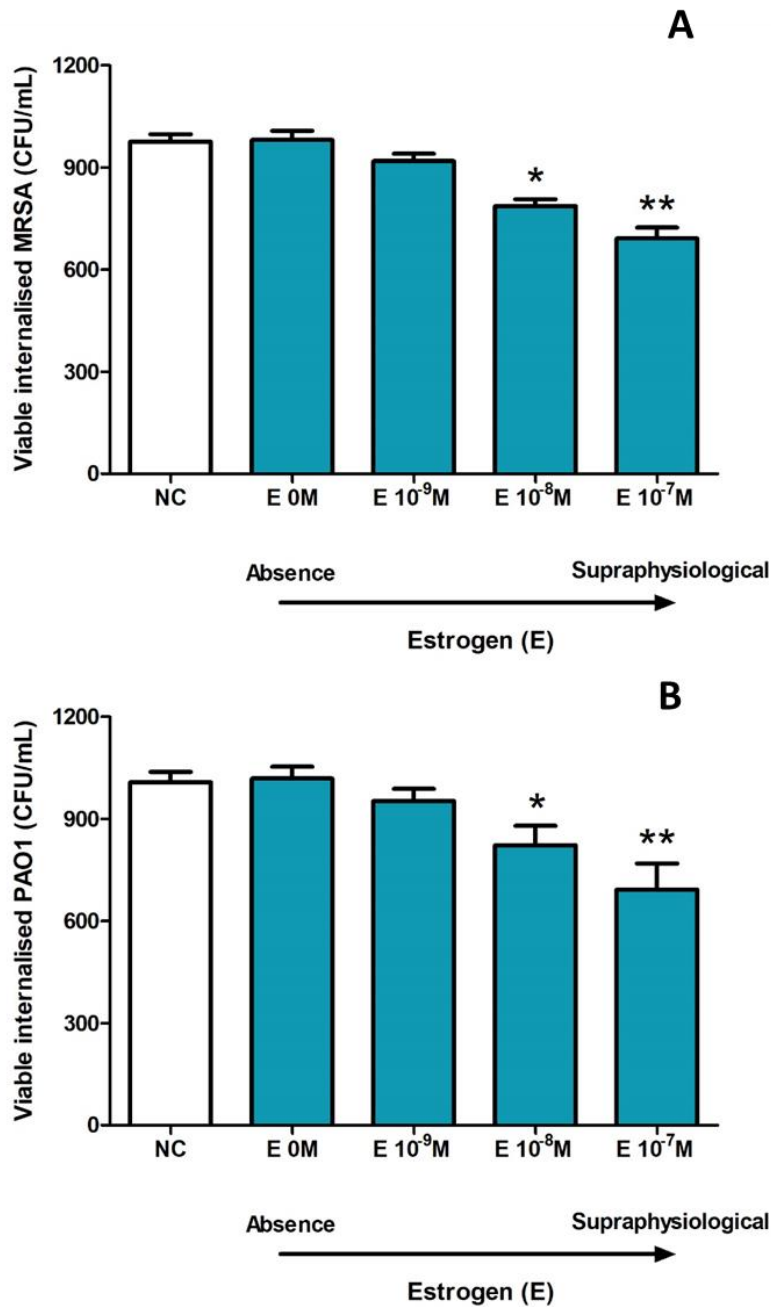


Figure 3.13. Estrogen promotes the killing of MRSA (A) and PAO1 (B) by U937-Derived Macrophages. Physiological (1×10^{-8} M) and supraphysiological (1×10^{-7} M) levels of estrogen significantly (*: $P < 0.05$, **: $P < 0.01$) promoted the killing of internalised MRSA (A) and PAO1 (B) by M0 macrophages compared to the negative control (NC). Data indicate mean ($n = 6$) viable internalised bacteria (CFU/mL) \pm standard error of the mean (StEM).

3.4.8 Visualisation of Bacterial Internalisation

3.4.8.1 Epifluorescence Microscopy

In order to visualise and compare the phagocytosis of pathogens, with and without estrogen treatment, U937-derived macrophages were incubated with GFP *S. aureus*

or mCherry *P. aeruginosa* prior to staining for phalloidin–tetramethylrhodamine B isothiocyanate conjugate from *Amanita phalloides* (Figure 3.14.A), or labelling with CD18 mouse monoclonal antibody and stained for goat anti-mouse IgG (H+L) superclonal secondary antibody, alexa fluor® 488 conjugate (Figure 3.14.B). Images were captured using a 100X objective on a Nikon E600 epifluorescence microscope (Section 2.2.11.1).

The absence of estrogen (A2 and B2) had no effect on the bacterial internalisation compared to the BSA negative controls (A1 and B1). However, it is clear that estrogen treated macrophages internalised more of the *S. aureus* and *P. aeruginosa* colonies (A3 and B3) than untreated cells.

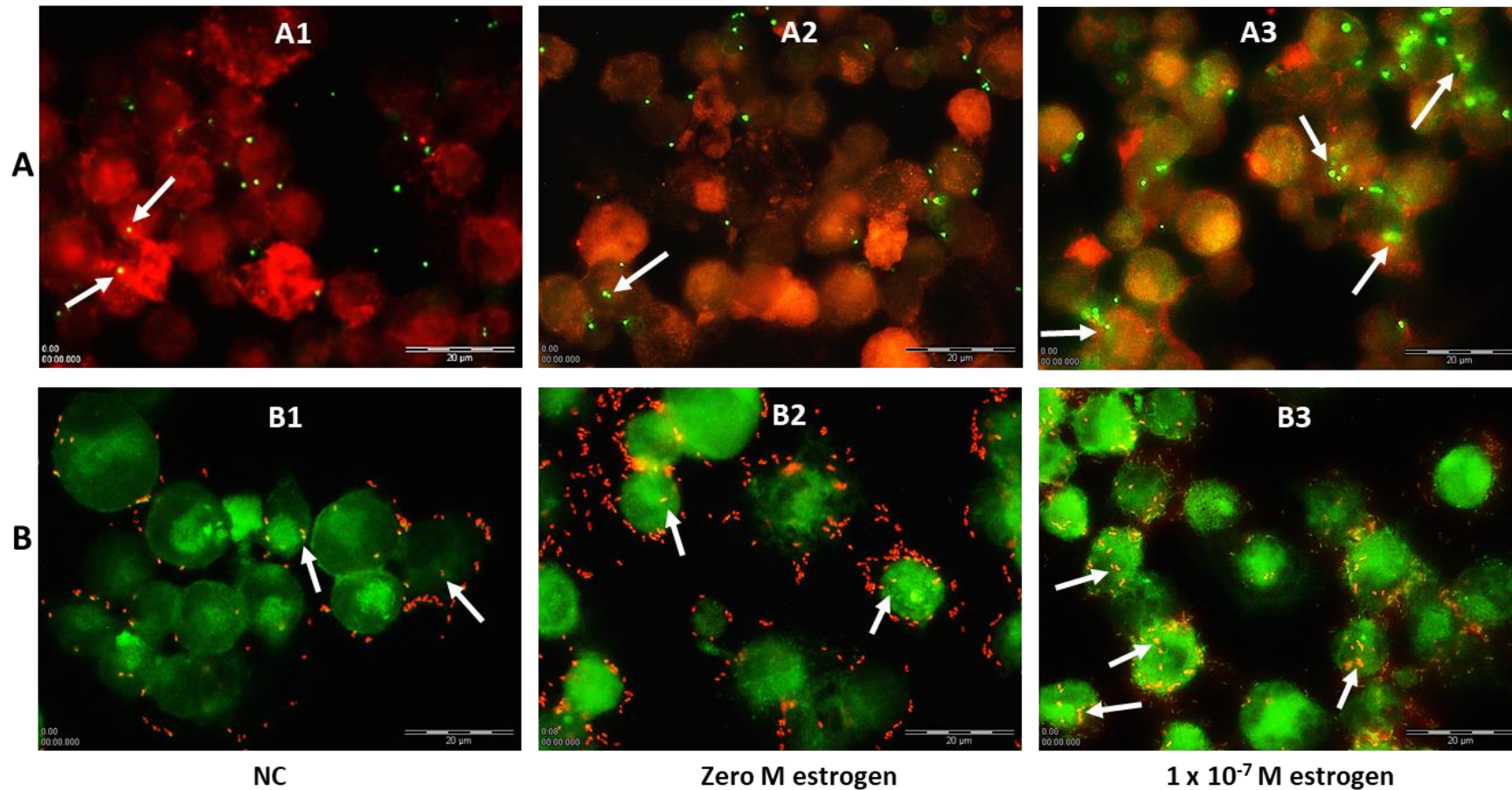


Figure 3.14. Effect of estrogen on the internalisation of GFP labelled-*S. aureus* (A) and mCherry-*P. aeruginosa* by U937 macrophages. U937-derived macrophages were treated with/without supraphysiological (1×10^{-7} M) estrogen supplementation for 24 hours prior to incubation with GFP labelled-*S. aureus* SH1000 (A) or mCherry labeled-PAO1 (B) for 3 hours. U937-derived macrophages were stained for phalloidin–tetramethylrhodamine B isothiocyanate conjugate from *Amanita phalloides* (A) or mouse anti-human CD18 monoclonal antibody followed by a goat anti-mouse IgG (H+L) superclonal secondary antibody, alexa fluor® 488 conjugate (B). The internalisation of *S. aureus* and *P. aeruginosa* by U937-derived macrophages (white arrows) was promoted by treatment with estrogen (A3 and B3) compared to negative control (NC) (A1 and B1) or macrophages lacking estrogen exposure (A2 and B2).

3.4.8.2 Z-Stack Analysis by Confocal Microscopy

To confirm the internalisation of bacteria by estrogen-treated U937 macrophages, Z-stacks were performed on phalloidin-stained phagocytes to visualise a series of layers through the interior of phagocytes and identify intracellular GFP-*S. aureus* colonies using a 63X objective on a Leica TSC SPE1000 confocal microscope. Z-stacks (example layer shown in (Figure 3.15) spanning the mid-intracellular region of estrogen-treated U937 macrophages (stained red) contained GFP-*S. aureus* colonies that fluoresced green, confirming the intracellular localisation of GFP-*S. aureus* inside (red) macrophages (Panel C). The size of Z-steps was set at 0.04 μm in this experiment.

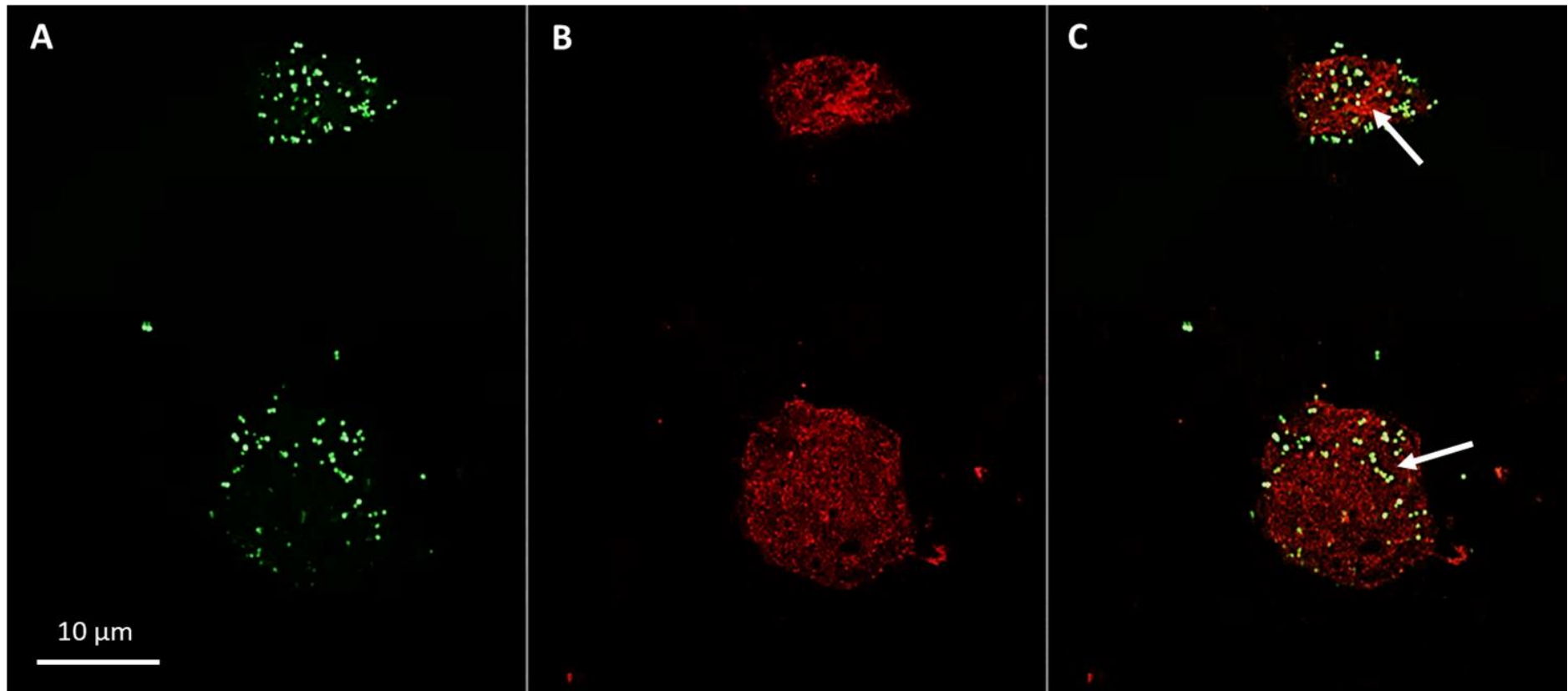


Figure 3.15. Z-Stack Demonstrating GFP-*S. aureus* Localisation in U937-Derived M0 macrophages Following Estrogen Supplementation. Bright green fluorescence was observed from GFP labeled *S. aureus* SH1000 (A). U937-derived M0 macrophages stained with phalloidin–tetramethylrhodamine B isothiocyanate conjugate from *Amanita phalloides* fluoresced red (B). The merged image confirmed several GFP-*S. aureus* colonies were located in the mid-intracellular region of estrogen-treated U937 M0 macrophages after 3 hours of host-pathogen interaction (C).

3.4.8.3 Scanning Electron Microscopy

In order to visualise and compare detailed features of the interactions between MRSA or PAO1 and macrophages, samples were visualised using a Supra 40VP scanning electron microscope as described in 2.2.11.2 (Figure 3.16). The moment of the MRSA and PAO1 uptake shows in the images (Panels B and C) as the estrogen treated macrophage sends out pseudopodia to engulf the invading bacteria. Surprisingly, the morphological response of macrophages toward PAO1 (Panel B) was distinct to and somewhat more pronounced than the response to MRSA (Panel C) in terms of the formation of membrane extensions, warranting further investigation. In concordance with these findings, PAO1 recovery was consistently higher than MRSA recovery in all host-pathogen experiments, supporting the notion of bacteria-specific interactions.

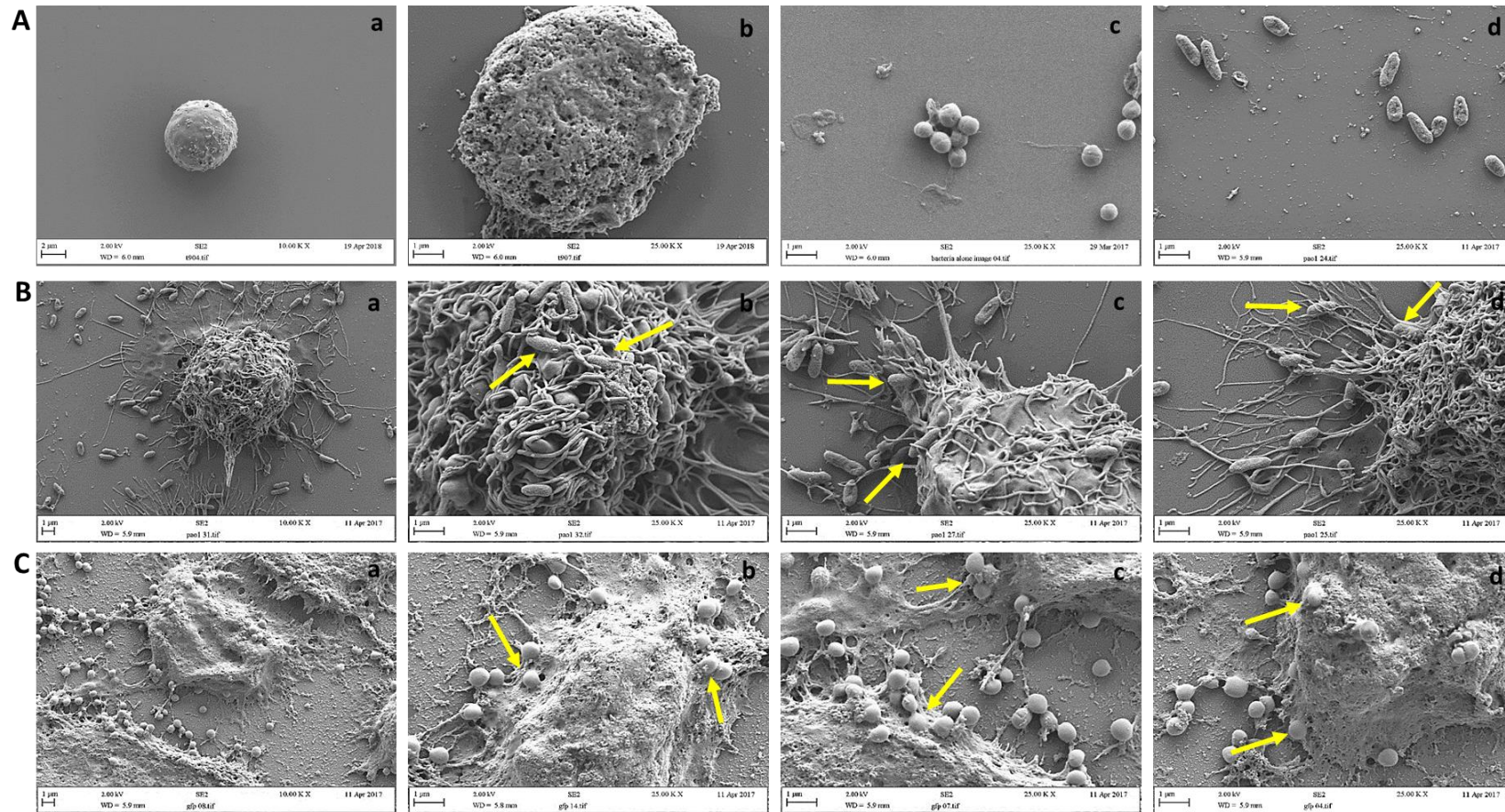


Figure 3.16. The Interaction of U937-derived M0 Macrophages with Gram Positive (MRSA11) and Gram Negative (PAO1) Bacteria Following Estrogen Supplementation. SEM images illustrating morphological changes in membrane extensions in estrogen-treated, U937-derived M0 macrophages after a 1h period of host-pathogen interaction with MRSA11 and PAO1. Control U937-derived M0 macrophages (Aa, Ab), MRSA11 (Ac) and PAO1 (Ad) were visualised before conducting the host-pathogen assay. Following the 1-hour period of host-pathogen interaction, estrogen-treated U937 M0 macrophages sent out morphologically distinct pseudopodial extensions to engage PAO1 (B) and MRSA (C) bacteria (yellow arrows).

3.5 Discussion

Wound infection is commonly quoted as a significant cause of delayed wound healing in the elderly and chronic wounds (Yotis, 1967). Chronic wounds are frequently colonised with a diverse range of Gram-positive and Gram-negative bacteria particularly *S. aureus* and *P. aeruginosa* (Dowd *et al.*, 2008). Studies of age-related impaired wound healing suggest that sex steroid hormones, particularly estrogen, may have a significant impact on the inflammatory response *in vivo*. Treatments with topical and systemic estrogen have been indicated to enhance the rate of acute wound healing in both elderly men, and particularly post-menopausal women, by reducing the inflammatory response (Gilliver *et al.*, 2007). Although estrogen has been shown to promote human acute wound healing in the elderly (Ashcroft *et al.*, 1999a), the effect of this hormone on bacterial clearance and wound infections has received little attention to date.

First, it was very important to develop and generate a successful *in vitro* model of monocyte-derived macrophages. PMA induced the differentiation of both U937 and human peripheral blood monocytes (HPBM) into M0 macrophages (Figure 3.2 and figure 3.3). This was confirmed via a high expression of the CD11c surface marker by macrophages, with monocytes lacking the marker. U937 and HPBM are known precursors of tissue macrophages and dendritic cells *in vitro* (Sintiprungrat *et al.*, 2010). PMA was described to induce the conversion of monocytes into macrophage-like cells *in vitro* (Martinez *et al.*, 2008; Rios de la Rosa *et al.*, 2017). Our results are supported by several reports as recent research reported that PMA-treated monocytes express high levels of CD11b, CD11c and protein kinase-C (PKC) (Sintiprungrat *et al.*, 2010). When treated with PMA, monocytes were shown to upregulate cell adhesion molecules including the beta-2-integrins (CD11a, CD11c, CD18 and CD11b) allowing the cells to attach to cell culture flask surfaces *in vitro*. PMA triggers calcium and phospholipid-dependent isoforms of PKC and encourages cyclic AMP metabolism, initiating maturation into a macrophage-like morphology (Luscinskas *et al.*, 1994).

A 3-hour incubation of both MRSA and *P. aeruginosa* with estrogen had no significant effect on their rate of growth. Therefore, any direct effect of estrogen on MRSA and PAO1 could be excluded during the host-pathogen assays in this study, with any changes in the number of bacteria instead arising solely due to the effect of estrogen on the interaction of macrophages with bacteria. These results are supported with many previous studies showing steroids, particularly estrogen and progesterone, have no effect on bacterial growth (Lev, 1959). However, some non-endogenous steroids have been reported to decrease the growth of many Gram-positive bacteria (Casas-Campillo *et al.*, 1961; Yotis, 1967), but the exposure to these steroids periods was for substantially longer periods than 3 hours.

The effect of phenol red on the host-pathogen interaction assays was investigated. Phenol red in the RPMI cell culture medium had no significant effect on MRSA and PAO1 recovery compared to RPMI medium lacking phenol red, regardless of the presence or absence of estrogen. Phenol red was shown to have very weak estrogenic effects on MCF-7 human breast cancer cells, with an affinity that was 0.001% that of 17 β -estradiol (Berthois *et al.*, 1986). Other studies reported that phenol red had no effect on estrogen-sensitive T47D breast cancer cells, primary immature rat pituitary cells or immature rat uterine cells (Welshons *et al.*, 1988). Thus, published findings are in concordance with the data in this study, demonstrating phenol red in the cell culture medium has negligible influence on host-pathogen interactions.

Multiplicity of infection (MOI) assays were then performed to investigate the effect of physiological estrogen (1×10^{-8} M) on phagocytosis over a range of starting bacterial inoculations (1×10^3 CFU, 1×10^4 CFU, 1×10^5 CFU and 1×10^6 CFU) for both MRSA or PAO1. The SERM tamoxifen, an estrogen-like molecule was found to enhance the bactericidal activity of neutrophils against *P. aeruginosa in vitro* and MRSA *in vivo* where (Corriden *et al.*, 2015) conducted *in vitro* host-pathogen assays at MOIs of 0.1. Findings in this study agree with published reports. The bacterial recovery of MRSA and PAO1 following estrogen (1×10^{-8} M) supplementation significantly increased as the MOI [bacteria: macrophage ratio] increased from 0.001 to 1. However, the bacterial recovery followed a similar pattern throughout,

regardless of the MOI ratio, with estrogen consistently increasing ($P<0.05$) bacterial clearance at all MOIs.

The effect of estrogen on the phagocytosis of the two major wound pathogens MRSA and *P. aeruginosa* by macrophages was investigated in this study. The results of the initial *in vitro* host-pathogen interaction assays (3.4.5.1) demonstrated novel data showing estrogen significantly ($P<0.05$) enhances the internalisation of both MRSA and *P. aeruginosa* in a dose-dependent manner following exposure to activated U937-derived M0 macrophages *in vitro*.

Li *et al.* (2000) extracted microglial cells from brain tissues of patients with Alzheimer disease. Similar to experimental procedures in this chapter, the macrophages were treated with 1×10^{-7} M estrogen for 24 hours and 48 hours prior to incubation with amyloid β -peptide (A β) for 24 hours. Estrogen was found to enhance the uptake of A β by brain macrophages (microglia) in a time- and dose-dependent manner. In order to confirm that estrogen enhances the phagocytic activity of microglia via a general engulfment mechanism, not specific to A β only, Li *et al.* (2000) re-conducted the same experiments replacing A β with *Escherichia coli*. Intriguingly, estrogen significantly increased the internalisation of *E.coli* in a concentration-dependent manner in comparison with the vehicle controls. These findings highlight the beneficial effects of estrogen on the phagocytic ability of microglia in the brain. However, other types of macrophages were not assessed and only *E.coli* was investigated so it remains unclear whether the effects of estrogen could be observed with varied bacterial species and strains, especially antibiotic-resistant strains.

Crompton *et al.* (2016) investigated the effect of estrogen on cutaneous murine wound healing following exposure to LPS derived from *Klebsiella pneumoniae*. Exposure of immune cells to LPS caused a substantial delay in wound healing. They then co-treated LPS-treated wounds with 17 β -estradiol, and interestingly, the delay in wound repair was reversed, with accelerated healing noted in mice following estrogen treatment (Crompton *et al.*, 2016). The results of this report were interesting but Crompton *et al.* (2016) only used locally applied *K. pneumoniae*-derived LPS in their model of infection. In order to generate realistic models of

infected wounds it is important to develop assays that infect wounds with live bacterial strains, not just bacterial-derived factors such as LPS. It is also important to consider both Gram-positive and Gram-negative models of infection. Moreover, *K. pneumoniae* is not a major wound pathogen compared to *S. aureus* and *P. aeruginosa*, which are the most frequently isolated bacterial species from infected wounds (Giacometti *et al.*, 2000; Gjødsbøl *et al.*, 2006; Kirketerp-Møller *et al.*, 2008; Rybtke *et al.*, 2015).

In vitro studies using cell lines provide only limited evidence to predict effective effects in humans. Thus, this study also confirmed findings in *ex vivo* primary human peripheral blood monocytes (HPBM) to establish a body of evidence using both a Gram-positive and a Gram-negative bacterium. Similar to results obtained with U937-derived macrophages *in vitro*, physiological and supraphysiological estrogen significantly increased ($P < 0.05$) the internalisation of both MRSA and PAO1 in comparison to the negative control.

Li *et al.* (2000) investigated the effect of estrogen on the uptake of fluorescent-*E.coli* by microglial cells using a fluorescence microplate reader, and showed estrogen enhanced the internalisation of *E.coli* in a concentration- and time-dependant manner. In line with this, and using similar experimental settings, data from the M0 macrophages-GFP *S. aureus* interaction assay strengthened the host-pathogen *in vitro* and *ex vivo* findings, indicating greater bacterial uptake following treatment with physiological and supraphysiological estrogen. Ashcroft and co-researchers reported that ovariectomized mice presented with high numbers of macrophages in injured tissues compared to wounds of normal healthy mice. Interestingly, in this thesis estrogen increased the number of macrophages that become active phagocytes following exposure to pathogens, thereby providing a justifiable reason for enhanced bacterial clearance described by Gilliver *et al.* (2007).

The effect of estrogen on the absolute killing of bacteria was investigated in the gentamicin protection assay. Gentamicin eradicates extracellular bacteria but it has no effect on intracellular bacteria due to poor ability to permeate the macrophage membrane (Waldbeser *et al.*, 1994; Hess *et al.*, 2004; Oelschlaeger, 2010; Hamad *et*

al., 2010). Interestingly, there was a significant decrease in the number of viable intracellular viable bacteria recovered following treatment of macrophages with physiological or supraphysiological estrogen, demonstrating estrogen promotes the killing of internalised pathogens, not just the uptake of bacteria by phagocytes.

Published research demonstrated that extended exposure of *ex vivo* macrophages to estrogen augments the expression of iNOS and the production of numerous cytokines, particularly IL-1 β , IL-12 and IL-6 following TLR4 activation by LPS *in vitro* (Deshpande *et al.*, 1997; Ghisletti *et al.*, 2005). The findings of this report combined with findings in Crompton *et al.* (2016) and Li *et al.* (2000) support the data in this Chapter, showing estrogen promotes the microbicidal activities of macrophages against bacteria.

In order to visualise the phagocytosis of *S. aureus* and *P. aeruginosa* with/without estrogen treatment, host-pathogen assays were conducted using fluorescent GFP *S. aureus* or mCherry *P. aeruginosa*. Visualisation of host-pathogen interactions was conducted via epifluorescent microscopy (Figure 3.14) and confocal microscopy (Figure 3.15). The findings demonstrated estrogen increased the internalisation of *S. aureus* and *P. aeruginosa* by macrophages, in concordance with the bacterial recovery data obtained from *in vitro* host-pathogen assays.

Images captured by SEM also confirmed increased host-pathogen interaction following treatment of macrophages with estrogen. It was noted that estrogen-treated macrophages had distinct morphological responses to MRSA and PAO1, in terms of pseudopodial formation. This finding was in concordance with data in the host-pathogen assays showing PAO1 recovery was consistently higher than MRSA recovery. The SEM images represent novel results that merit further investigation to determine the underlying mechanisms that might lead to pathogen-specific differences in pseudopodial formation. In order to better understand the biochemical processes occurring at the host-pathogen interface, future work could determine the effect of estrogen on physical and chemical changes in phagocytes. Fourier Transform Infra-Red (FTIR), spectroscopy and Raman spectroscopy can be used to determine changes in the functional groups on the cell surface of active phagocytes treated with estrogen.

The underlying mechanism by which estrogen affects bacterial phagocytosis has not been investigated yet. It is possible that estrogen promotes bacterial clearance by inducing genes or pathways involved in the processes of phagocytosis via activation of ER proteins. Future investigations will determine the precise ER pathways involved in estrogen-enhanced phagocytosis. Determining the mechanisms by which estrogen induces phagocytosis may lead to novel therapeutic strategies that combat wound infections in the elderly by modulating the host immune response. One potential avenue will be to use selective estrogen receptor modulators (SERMs) such as tamoxifen that act as agonists of ER proteins in a tissue-specific manner.

Tissue macrophages exist in different activation states: either pro-inflammatory (classically activated) by LPS or IFN- γ , referred to as M1 macrophages, or anti-inflammatory (alternatively activated) by IL-13 or IL-4, referred to as M2 macrophages (Aron-Wisnewsky *et al.*, 2009). Preliminary assays in this study were performed on PMA-differentiated M0 macrophages. Further work will investigate M1 and M2 macrophages to determine whether estrogen influences phagocytosis in an activation-dependent manner.

Images from the SEM (Figure 3.16) illustrated major morphological changes in the membrane of phagocytes (macrophages) following estrogen treatment, with the establishment of multiple pseudopodial structures suggesting estrogen induces significant alterations in the actin cytoskeleton of phagocytes. These findings suggested the downstream genes mediated by estrogen may include mediators of the actin cytoskeleton. Thus, further investigations could measure differences in activated/total levels of proteins involved in the regulation of the actin cytoskeleton, such as members of the Rho GTPase family.

Findings of this study are novel and support the preliminary work performed by Crompton *et al.* (2016). In summary, bacterial phagocytosis by macrophages was elevated following stimulation with physiological (typical of youth) and supraphysiological estrogen levels compared to estrogen deprivation and an absolute absence of estrogen. This suggests that the ageing process (age-related decline in estrogen levels) may increase the propensity and progression of wound

infections in the elderly, in line with the findings of Ashcroft *et al.* (1999a). The findings of this study suggest local and/or systemic hormone replacement therapy (HRT) might resolve or reduce the likelihood of Gram-positive and Gram-negative wound infections in the elderly.

**Chapter 4: Effect of Estrogen on
the Phagocytosis of MRSA and *P.*
aeruginosa by M1-like and M2-
like Macrophages**

4.1 Introduction

4.1.1 Macrophages Polarisation and Plasticity

The plasticity of macrophages is established by acquisition of distinct functional and morphological characteristics directed by particular tissues, and/or immunological and environmental stimuli (Mantovani *et al.*, 2005; Gordon and Martinez, 2010; Wynn *et al.*, 2013). Macrophages in tissues have several similar characteristics but they are an extremely heterogeneous population of cells in relation to their function and expression of surface markers (Figure 4.1) (Mackaness, 1964; Nathan *et al.*, 1983; Stein *et al.*, 1992; Mantovani *et al.*, 2005; Murray and Wynn, 2011; Murray *et al.*, 2014). Macrophages are key effectors of the innate immune system, providing an initial defence against foreign invaders such as bacteria (Duque and Descoteaux, 2015). They also play key roles in initiating and controlling the adaptive immune responses (Duque and Descoteaux, 2015). Macrophages are crucial in tissue repair, haemostasis, inflammation and remodelling (Mosser and Edwards, 2010). As a condition of a favourable resolution of inflammation, macrophages remove apoptotic leukocytes, and dysregulation in this particular macrophage role leads to chronic inflammatory and autoimmune diseases (Balhara and Gounni, 2012). Macrophages have multiple functions and characteristics (pro-inflammatory or anti-inflammatory) in immunity, depending on the type of stimulation and resulting functional phenotype (Martinez *et al.*, 2008). Macrophages were initially classified into two major phenotypes based on type 1 helper T-cell (Th1) and type 2 helper T-cell (Th2) polarisation (Hesse *et al.*, 2001; Puig-Kröger *et al.*, 2004; Yona *et al.*, 2013; Martinez and Gordon, 2014). Cytokines mainly released by Th1 cells, such as IFN- γ differentiate macrophages to pro-inflammatory M1 macrophages. However, cytokines released by Th2 cells such as IL-4 have been reported to dampen macrophage activation, resulting in M2 anti-inflammatory macrophages (Raes *et al.*, 2002; Kzhyshkowska *et al.*, 2006).

4.1.1.1 Classically Activated Macrophages

In response to cytokines such as IFN- γ , and/or bacterial components such as LPS, macrophages are polarized to a pro-inflammatory phenotype and are referred to as classically activated or M1 macrophages. Classical activation or M1 polarisation are terms that were first used during the 1960s when studies on mice reported macrophages with enhanced anti-bacterial activities towards infections with *Mycobacterium bovis bacillus Calmette-Guerin* (BCG) or *Listeria monocytogenes* (Mackness, 1964). The cytokine IFN- γ produced by TH1 cells was described as the main activator of antimicrobial and cytotoxic activities of macrophages (Mosmann *et al.*, 1986). Genetic deficiencies of IFN- γ and/or its receptors confirmed that macrophages polarisation to the M1 phenotype is essential for host defence against pathogenic infections in knockout mice models and in humans. Components of intracellular pathogens (e.g. LPS), TNF, GM-CSF and TLR4 ligands were also described to encourage M1 polarisation (Mantovani *et al.*, 2005).

M1 macrophages stimulate TH1 immunity by releasing high levels of IL-12 (Mantovani *et al.*, 2004), producing microbicide enzymes such as inducible NO synthase (iNOS) to destroy pathogenic invaders, secreting pro-inflammatory cytokines, such as, IL-1 β , IL-6, IL-12, IL-23, TNF- α , and releasing chemokines to attract immune cells to the infection site and MMPs (Mantovani *et al.*, 2004).

4.1.1.2 Alternatively Activated Macrophages

In 1990, a study reported for the first time the inhibition of M1 polarisation. This study revealed that the production of IFN- γ induced superoxide (O_2^-) by IL-4 was inhibited (Abramson and Gallin, 1990). Later on the 1990s, Gordon and colleagues described the term “alternatively activated macrophages (Stein *et al.*, 1992). It is commonly believed that M2 macrophages are involved in tissue repair, fibrosis and tumour progression (Wynn, 2004). They are identified by expression of surface markers, mainly the macrophage mannose receptor CD206 (Stein *et al.*, 1992) and CD163 (Zeyda *et al.*, 2007). Arginase (Arg-1) is an enzyme that is involved in the production of proline via conversion of arginine to ornithine (Martinez *et al.*, 2008). Proline has been linked with ECM formation and collagen deposition (Hesse *et al.*,

2001). M2 macrophages release anti-inflammatory cytokines, such as, IL-10, and are crucial in resolving chronic inflammation, stimulating angiogenesis and fibrogenesis and promoting tissue remodelling (Mosser and Edwards, 2008).

Depending on the stimuli that induces polarisation of macrophages to the M2 phenotype, M2 macrophages were divided into three subpopulations: M2a encouraged by stimulation with IL-4 and/or IL-13, M2b encouraged by TLR ligands, immune complexes (ICs), or IL-1R and M2c induced by IL-10/TGF- β (Mantovani *et al.*, 2004).

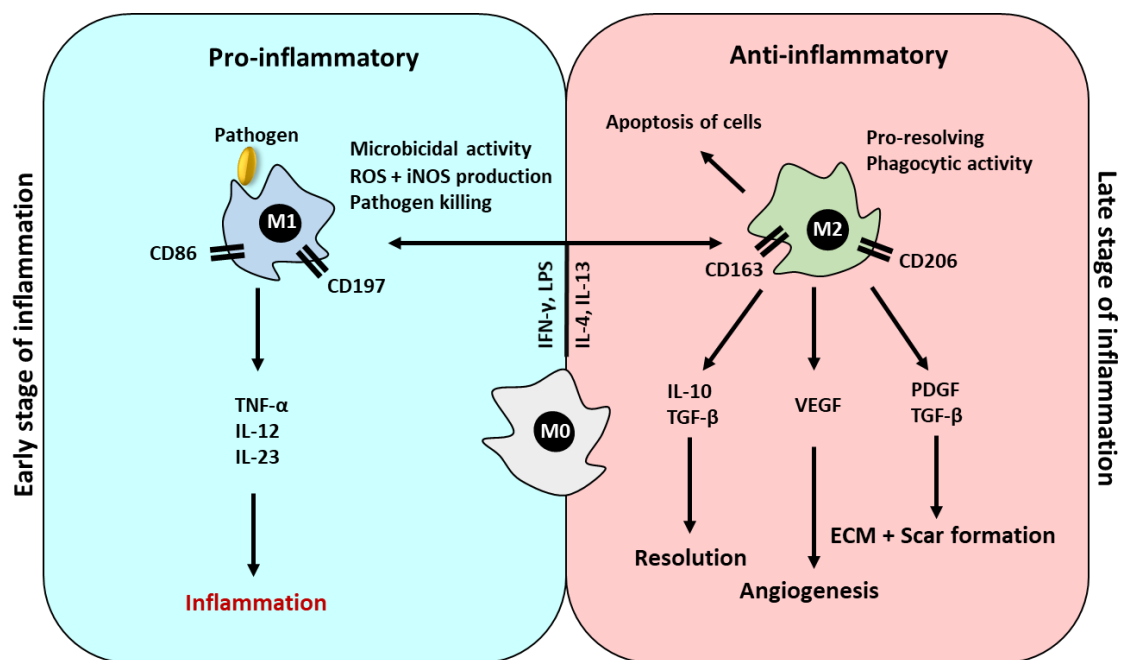


Figure 4.1. Plasticity of macrophages in wound healing. Following an injury, monocytes infiltrate to the wound site and become macrophages (M0). During the early stage of inflammation, macrophages switch to the M1 phenotype. M1 macrophages have a microbicidal activity and induce TH1 responses via release of IL-12. Close to the end of the inflammatory stage, the microenvironment of the wound changes and the process of clearing apoptotic cells initiates the polarisation of macrophages toward M2 macrophages. M2 cells play a key role in the resolution of inflammation and progression to subsequent stages of wound healing. Both M1 and M2 macrophages are key players in switching from inflammation to proliferation (angiogenesis, re-epithelialisation and matrix production). Figure drawn based on information in (Benoit *et al.*, 2008; Krzyszczyk *et al.*, 2018).

4.1.2 Macrophage Phenotypes and Plasticity in Wound Healing

The proliferation of primary monocytes is limited *in vitro* but evidence suggests skin macrophages can auto-renew *in vivo* (Davies *et al.*, 2013). In contrast, it remains unclear if macrophages are only recruited to the wound site when required and

removed after wound repair, or if macrophages deriving from monocytes can proliferate *in vivo* (Italiani and Boraschi, 2014).

Macrophages were initially classified into two major phenotypes referred to as M1 and M2 macrophages. M1 macrophages are commonly linked with pro-inflammatory events, while M2 macrophages are known to be anti-inflammatory and pro-regenerative. It is essential to understand that M1 and M2 macrophages are not separate types of cells, but they form a scale in which they take varying degrees of M1 or M2 properties. In addition, *in vivo* studies propose that macrophages are a heterogeneous group of cells all displaying a diversity of M1 and M2 characteristics (Martinez and Gordon, 2014; Murray *et al.*, 2014; Ogle *et al.*, 2016).

Distinct M1 and M2 macrophages classification is not applied when comparing wound healing macrophages *in vivo*. However, this classification is widely used *in vitro* when molecules inducing the polarisation are known and experimentally introduced to the system (Novak and Koh, 2013). The types and suggested roles of macrophages related with wound healing *in vitro* and *in vivo* are illustrated (figure 4.2).

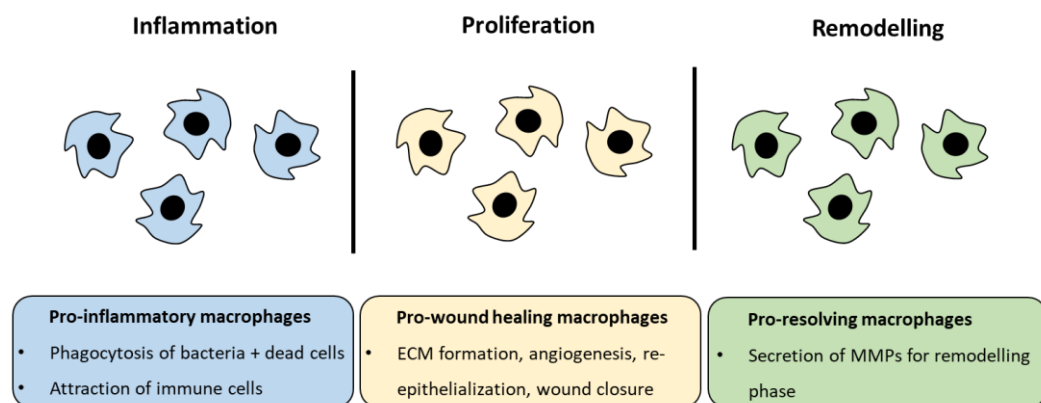


Figure 4.2. The role of macrophage phenotypes in wound repair. During haemostasis and inflammation, pro-inflammatory macrophages invade the wound area and start removing dead cells and bacteria. During the proliferative stage, pro-wound healing macrophages are present and release factors such VEGF, PDGF, TGF- β to aid angiogenesis, ECM and scar formation and re-epithelialization. During remodelling, pro-resolving macrophages are involved in strengthening the new skin barrier.

In line with the wound healing process, pro-inflammatory macrophages are dominant at the wound site after an injury, followed by pro-wound healing

macrophages that stimulate the growth of cells during the proliferative phase, and finally pro-resolving macrophages that dampen immune responses and stimulate the remodelling and maturation of collagen (Vannella and Wynn, 2017).

During wound healing, pro-inflammatory macrophages are a source of NO, ROS, IL-1, IL-6, and TNF- α . They release MMPs in order to disrupt the extracellular matrix and create a space for inflammatory cells to infiltrate the wound (Murray and Wynn, 2011). They also release important levels of growth factors such as PDGF, VEGF, insulin-like growth factor 1 (IGF-1) and TGF- β 1 which facilitate the proliferation of cells involved in angiogenesis and ECM deposition (Murray and Wynn, 2011). Pro-resolving macrophages dampen inflammation by upregulating IL-10 and produce MMPs to remodel and reinforce the ECM (Vannella and Wynn, 2017). The main roles of pro-resolving macrophages are to reinstate homeostasis and minimize fibrosis via apoptosis of myofibroblasts, and suppression T cell proliferation (Murray and Wynn, 2011).

Knowing that wound healing is a sequence of overlapping phases, all types of macrophages share many characteristics simultaneously. This is particularly true for pro-wound healing macrophages as they fluctuate between the early and late stages of wound repair, and therefore display similar features to pro-inflammatory and pro-resolving macrophages.

M1 macrophages are the dominant pro-inflammatory macrophages during the early pro-inflammatory stage of wound repair. They are phagocytic, and serve as cleaners of the wound site by removing foreign invaders and dead tissue. *In vitro*, M1 macrophages are induced by intracellular cytokines such as IFN- γ , and/or bacterial elements, such as LPS and peptidoglycans. M1 macrophages are distinguished *in vitro* by the expression of the chemokine receptor CCR7 (CD197) and high levels of the co-stimulatory molecules CD80 and CD86, resulting in efficient antigen presentation capacity (Mantovani *et al.*, 2004).

M2 macrophages play a regenerative role in wound healing. They are induced by IL-4 and IL-13 and characterised by a high expression of the mannose receptor (CD206) (Stein *et al.*, 1992), the haemoglobin scavenger receptor (CD163) (Zeyda *et*

al., 2007), IL-10, and TGF- β . M2 macrophages release only very low levels of pro-inflammatory mediators such as TNF- α , IL-12, and IL-8. In wounds M2 macrophages have been divided into three subtypes (M2a, M2b and M2c) relating to expression of different surface markers. The three subtypes are commonly identified and used *in vitro* to investigate M2 phenotype characteristics. However, distinct M2 macrophages classification is not widely applied when referring to wound healing *in vivo*. This is due to the heterogeneous populations present, which are generated from a variety of stimuli within wounds (Novak and Koh, 2013). Studies have reported that the wound healing M2 macrophages align with the well-defined subset of M2a (Ogle *et al.*, 2016). M2a macrophages are induced by IL-4/IL-13 and distinguished by the expression of CD206. They produce arginase-1, IGF-1, PDGF-BB and a variety of chemokines (CCL17, CCL18, CCL22) (Ogle *et al.*, 2016). They are crucial in the proliferative stage of wound healing by aiding ECM formation via production of collagen precursors and stimulation of fibroblasts. M2 macrophages also release high levels of PDGF, which is involved in angiogenesis (Vannella and Wynn, 2017).

Overall, macrophage classification is complex and not yet fully understood with uncertainty whether macrophage phenotypes are distinct, or even appropriate to *in vivo* models of wound healing (Martinez and Gordon, 2014). Although the mechanism is not understood, a balance between the M1 and M2 phenotypes is essential to preserve a homeostatic environment, and imbalance leads to disturbed inflammation.

Exogenous estrogen supplementation enhances healing in elderly people, indicating that the systemic and peripheral age-related decrease in endogenous estrogen levels impairs healing in both sexes (Ashcroft *et al.*, 1997b; Hardman and Ashcroft, 2005). Known for its anti-inflammatory properties, estrogen is reported to resolve excessive inflammation by directly effecting inflammatory responses of monocytes and macrophages (Ashcroft *et al.*, 2003; Guo and DiPietro, 2010). Estrogen has been reported to accelerate re-epithelialisation, promote angiogenesis, increase matrix deposition and wound contraction, dampen the inflammatory response and inhibit the release of pro-inflammatory cytokines (Ashcroft and Ashworth, 2003).

These findings suggest estrogen modulated both pro-inflammatory (M1) and anti-inflammatory / pro-regenerative (M2) macrophages during wound repair. Estrogen is also believed to be a key player in regulating immune responses against bacterial infections in wounds (Cutolo *et al.*, 2004). Data in the previous chapter showed that estrogen significantly enhanced the phagocytosis of MRSA and *P. aeruginosa* by classical M0 macrophages, in a concentration-dependant manner when above levels described as estrogen deprivation. Using models of macrophage polarisation into M1 and M2 phenotypes, the effect of estrogen on the phagocytosis of MRSA and *P. aeruginosa* by M1-like and M2-like macrophages was investigated.

4.2 Aim and Objectives

4.2.1 Aim

To investigate the effect of estrogen (age-related changes in estrogen concentrations) on the phagocytic function of both pro-inflammatory (M1) and anti-inflammatory (M2) macrophages using *in vitro* and *ex vivo* models of host-pathogen interactions.

4.2.2 Objectives

- Generate an *in vitro* model of HPBM and U937 monocyte differentiation into M1-like and M2-like macrophages.
- Compare the phagocytic function of M1 and M2 macrophages under conditions of physiological and supraphysiological levels of estrogen with conditions of estrogen deprivation or an absolute absence of estrogen.

- Compare the effect of age-related changes in estrogen levels on the phagocytosis of Gram-positive and Gram-negative bacteria by M1-like and M2-like macrophages.

4.3 Methods

4.3.1 Flow Cytometry

U937 and HPBM were differentiated into M1-like and M2-like macrophages *in vitro* (Figure 4.3) as described in section 2.2.5. The expression of surface markers specific for M1 (CD197) and M2 (CD206) macrophages (relative MFI and % fluorescence) on M1- and M2-like cells were compared with untreated M0 control macrophages.

4.3.2 *In vitro* and *Ex Vivo* Models of Host-Pathogen Interactions

U937 and HPBM M1-like/M2-like macrophages were adhered to wells and treated with different concentrations (1×10^{-7} M, 1×10^{-8} M, 1×10^{-9} M and zero M) of estrogen in antibiotic-free medium for 24 hours to represent the models of age-related changes in estrogen levels described in Sections 2.2.2 and 2.2.8.2. Cells were incubated with MRSA or *P. aeruginosa* (1×10^4 CFU) for 3 hours at 37 °C and 5% CO₂ prior to counting bacterial colonies (CFU/mL) following the methods described in Section 2.2.8.2.

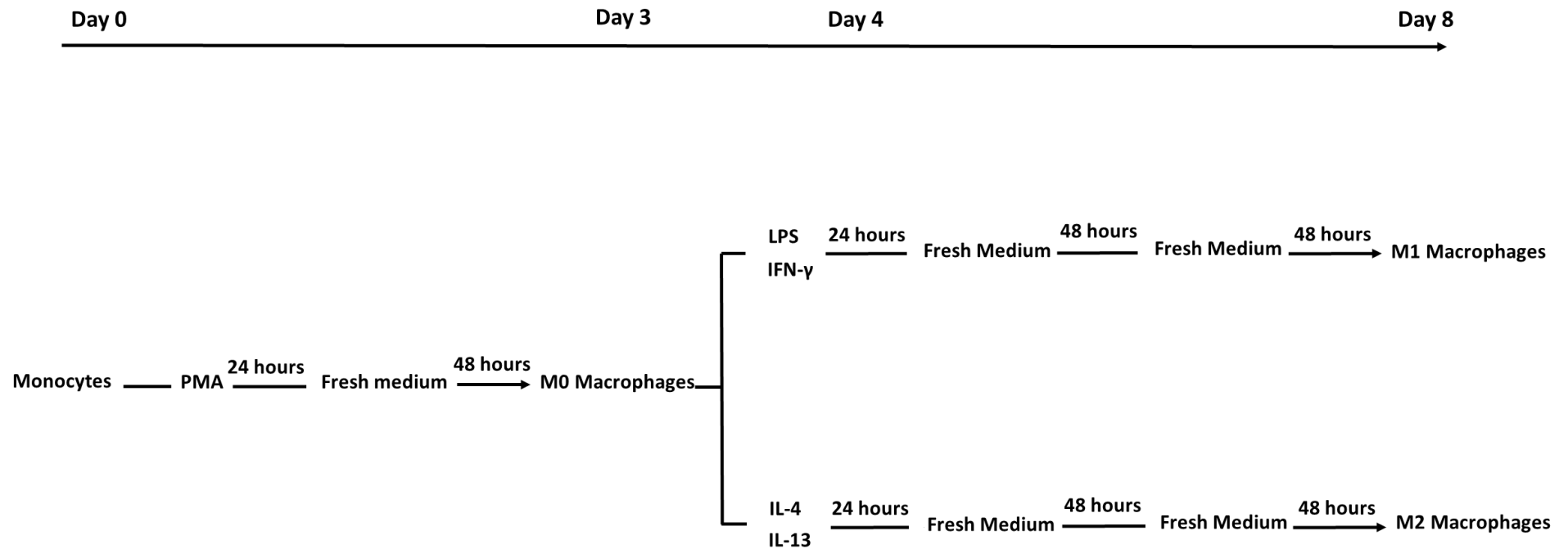


Figure 4.3. Experimental Generation of M1 and M2 Macrophages. U937 and CD14⁺ HPBM were differentiated into adherent M0 macrophages using PMA for 24 hours (with medium being changed after 24 hours). Further polarisation into classically activated (M1) or alternatively activated (M2) macrophages was performed using LPS + IFN-γ or IL-4 + IL-13 for 24 hours, respectively. The medium was changed every other day to generate resting M1-like and M2-like macrophages.

4.4 Results

4.4.1 Differentiation of Monocytes into M1 and M2 Macrophages

The expression of CD197 and CD206, which are surface markers for M1 (Mantovani *et al.*, 2004) and M2 (Stein *et al.*, 1992) macrophages respectively, were examined by flow cytometry (Figure 4.4). The findings confirmed the successful conversion of U937 M0 macrophages into CD197⁺ M1 and CD206⁺ M2 macrophages. The U937 macrophages treated with LPS/IFN- γ were 94.7% CD197⁺ and 0.4% CD206⁺, whereas the U937 macrophages treated with IL-4/IL-13 were 6.6% CD197⁺ and 79.8% CD206⁺. The MFI for M1 surface marker CD197 was significantly ($P<0.01$) higher in U937 macrophages treated with LPS/IFN- γ compared to untreated U937 macrophages or U937 macrophages treated with IL-4/IL-13. In contrast, the MFI for the M2 surface marker CD206 was significantly ($P<0.01$) higher in U937 macrophages treated with IL-4/IL-13 compared to untreated U937 macrophages or macrophages treated with LPS/IFN- γ .

Similarly, the findings confirmed the successful conversion of HPBM-derived macrophages into CD197⁺ M1 and CD206⁺ M2 macrophages (Figure 4.5). There was substantial expression of the M1 surface marker CD197 (67.2%) by HPBM-derived macrophages treated with LPS/IFN- γ , whereas CD197 was almost absent in HPBM-derived M0 macrophages (1%) and at very low levels (5.7%) in HPBM-derived macrophages treated with IL-4/IL-13. The M2 surface marker CD206 was substantially expressed (73.4%) by HPBM-derived macrophages treated with IL-4/IL-13 whereas CD206 was almost absent from HPBM-derived M0 macrophages (0.1%) and HPBM-derived macrophages treated with LPS/IFN- γ (0.2%). The MFI for CD197 from HPBM-derived macrophages treated with LPS/IFN- γ was significantly ($P<0.01$) greater than from HPBM-derived M0 macrophages or HPBM-derived macrophages treated with IL-4/IL-13. In contrast, the MFI for CD206 from HPBM-derived macrophages treated with IL-4/IL-13 was significantly ($P<0.01$) higher than from HPBM-derived M0 macrophages or HPBM-derived macrophages treated with LPS/IFN- γ .

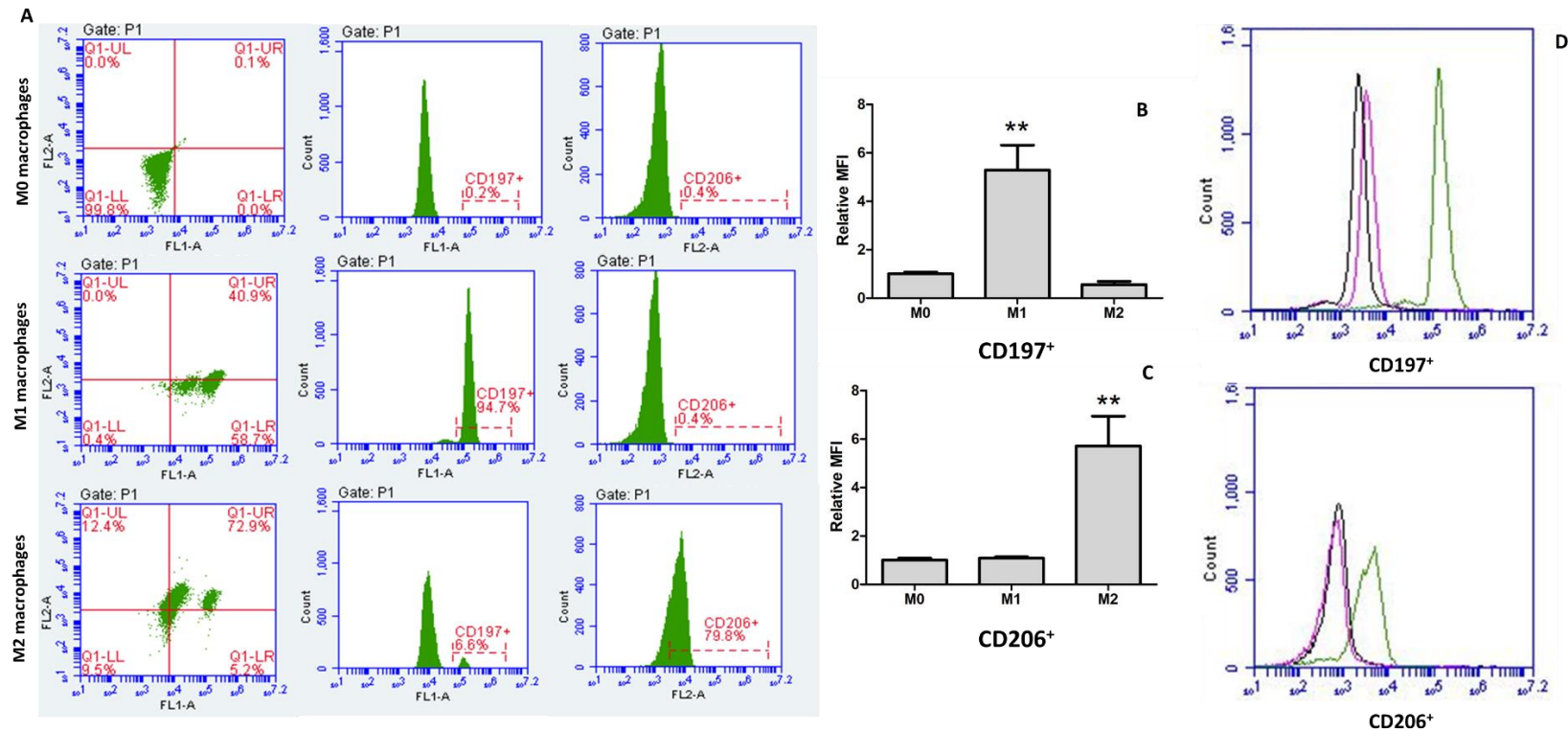


Figure 4.4. Differentiation of U937 M0 macrophages onto M1/M2 macrophages. The polarisation of U937 M0 macrophages into M1-like and M2-like macrophages was confirmed by through detection of CD197 and CD206 surface markers by flow cytometry respectively. M0 macrophages did not express CD197 or CD206 markers (0.2% and 0.4% respectively). U937 macrophages treated with LPS/IFN- γ expressed high levels of CD197 (94.7%) but almost lacked CD206 (0.4%). U937 macrophages treated with IL-4/IL-13 expressed CD206 (79.8%) but very low levels of CD197 (6.6%) (A). Three distinct populations of U937 macrophages were identified (D) with significantly higher MFI for CD197 from M1 macrophages compared to M0 and M2 macrophages (B), and a significantly higher MFI for CD206 from M2 macrophages compared to M0 and M1 macrophages (C). Data represent averages of $n = 3$ experiments. ** Indicates significant difference (One-way ANOVA) in MFI ($P < 0.01$) compared to M0 macrophages. Error bars represent the standard error of the mean (StEM).

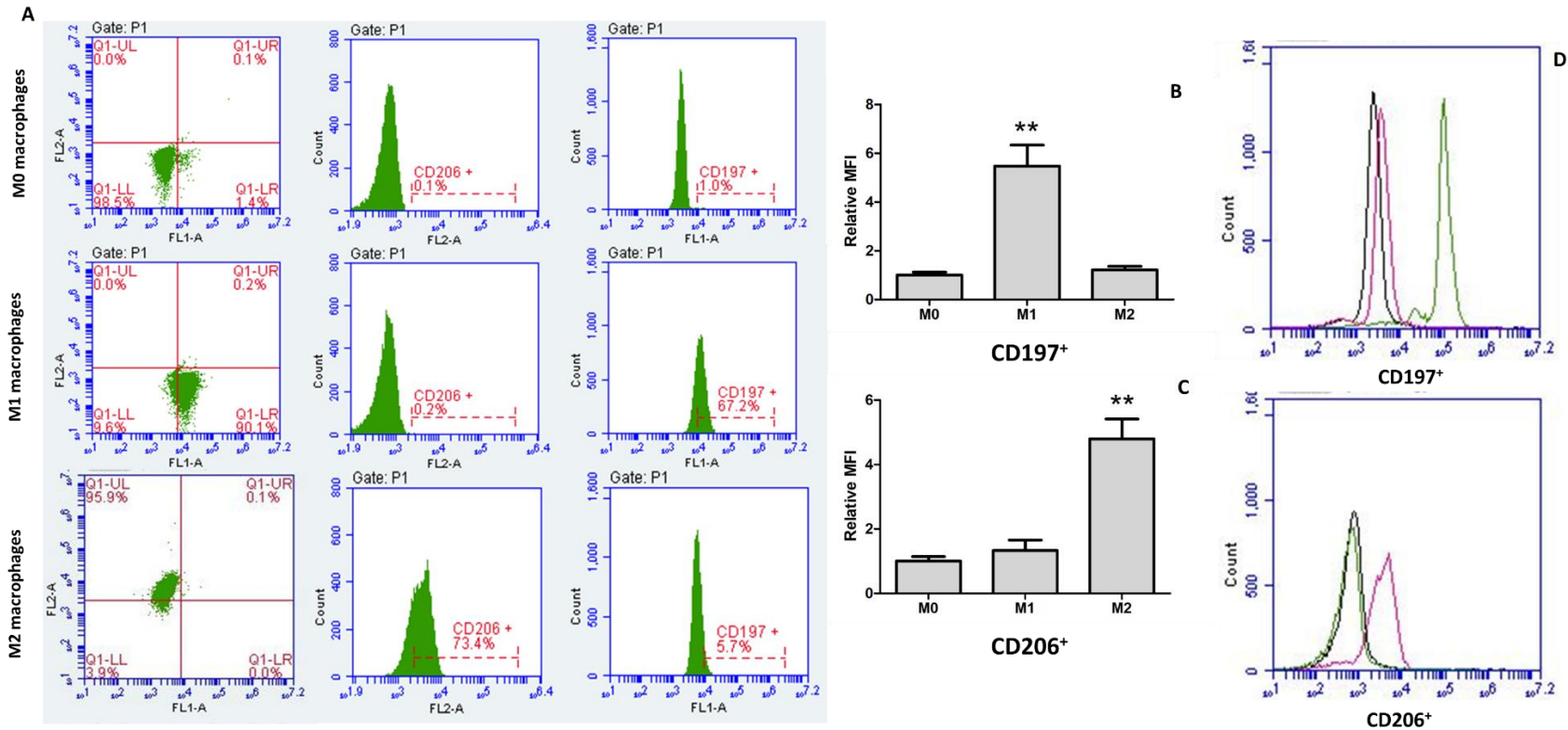


Figure 4.5. Polarisation of HPBM M0 macrophages into M1/M2 macrophages. The polarisation of HPBM M0 macrophages into M1 and M2 macrophages was confirmed via detection of CD197 and CD206 surface markers by flow cytometry respectively. PMA-differentiated M0 macrophages almost lacked both CD197 (1%) and CD206 (0.1%). In contrast, macrophages treated with LPS/IFN- γ expressed CD197 (67.2%) but almost lacked CD206 (0.2%). Macrophages treated with IL-4/IL-13 expressed CD206 (73.4%) but very low levels of CD197 (5.7%) (A). Three distinct populations of HBPM macrophages were detected (D) with significantly higher MFI for CD197 from M1 macrophages compared to M0 and M2 macrophages (B), and significantly higher MFI for CD206 from M2 macrophages compared to M0 and M1 macrophages. Data represent an average of $n = 3$ experiments. ** Indicates significant difference (One-way ANOVA) in MFI ($P < 0.01$) compared to M0 macrophages. Error bars represent the standard error of the mean (StEM).

4.4.2 Effect of Age-Related Changes in Estrogen Levels on *In vitro* Phagocytosis of MRSA and *P. aeruginosa* by U937-derived M1 and M2 Macrophages

The effect of age-related changes in estrogen levels on the clearance of MRSA and *P. aeruginosa* by U937-derived M1 and M2 macrophages was investigated (Figure 4.6). Compared to the negative control, conditions representing an absolute absence of estrogen (zero M) and estrogen deprivation (1×10^{-9} M) had no significant effect ($P > 0.05$) on the clearance of MRSA or PAO1 by both U937-derived M1 and M2 macrophages.

In contrast, physiological and supraphysiological levels of estrogen significantly ($P < 0.01$) reduced the recovery of MRSA and *P. aeruginosa* by both U937-derived M1 and M2 macrophages in a dose-dependent manner compared to the negative control. After a 3-hour incubation with U937-derived M1 macrophages, the MRSA recovery was significantly ($P < 0.05$) reduced to 1586 CFU/mL and 1476 CFU/mL following treatment with physiological and supraphysiological levels of estrogen respectively, compared to the negative control (2117 CFU/mL). Although somewhat less effective, after a 3-hour incubation with U937-derived M2 macrophages with physiological or supraphysiological levels of estrogen still significantly ($P < 0.05$) reduced the recovery of MRSA to 2927 CFU/mL and 2533 CFU/mL respectively compared to the negative control (3538 CFU/ml).

Similarly, treating U937-derived M1 macrophages with physiological or supraphysiological levels of estrogen significantly ($P < 0.05$) reduced PAO1 recovery to 2405 CFU/mL and 1920 CFU/mL respectively compared to the negative control (2660 CFU/mL). Although somewhat less effective, treating U937-derived M2 macrophages with physiological or supraphysiological levels of estrogen still significantly ($P < 0.05$) reduced the recovery of PAO1 to 3052 CFU/mL and 2594 CFU/mL respectively compared to the negative control (3790 CFU/mL).

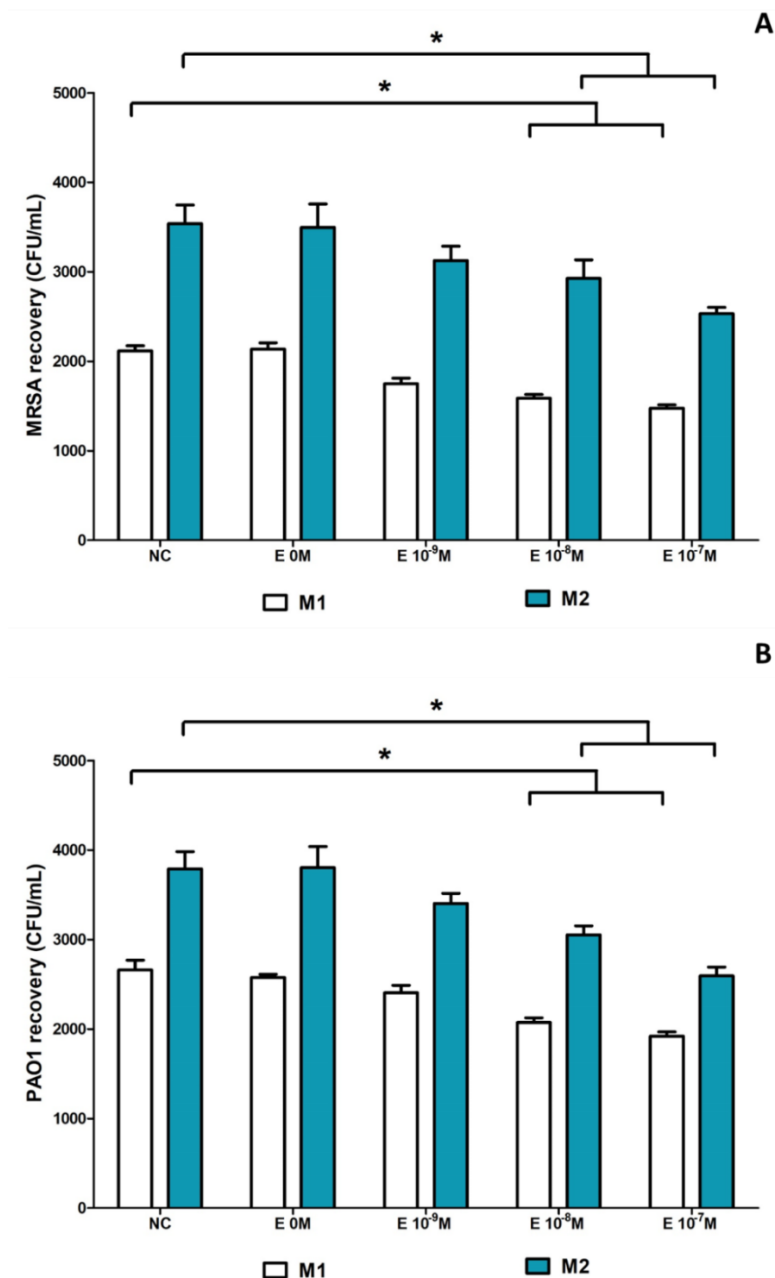


Figure 4.6. Estrogen enhances the phagocytosis of MRSA (A) and PAO1 (B) by U937-derived M1/M2 macrophages *in vitro*. The recovery of MRSA (A) and PAO1 (B) after incubation with U937-derived M1/M2 macrophages were significantly (*: $P < 0.05$, **: $P < 0.01$) decreased following treatment with physiological (1×10^{-8} M) or supraphysiological (1×10^{-7} M) levels of estrogen compared to the negative control. U937-derived M1 macrophages internalised significantly ($P < 0.05$) more MRSA/PAO1 respectively than U937-derived M2 macrophages. Data is mean \pm StEM, $n = 10$.

4.4.3 Effect of Age-Related Changes in Estrogen Levels on the Phagocytosis of MRSA and *P. aeruginosa* by *Ex Vivo* M1 and M2 Human Macrophages

The effect of estrogen on the phagocytosis of MRSA and *P. aeruginosa* by *ex vivo* HPBM-derived M1/M2 macrophages was investigated (Figure 4.7). After a 3-hour

incubation of MRSA or *P. aeruginosa* with HPBM-derived M1/M2 macrophages, the number of recovered MRSA and *P. aeruginosa* under conditions representing an absolute absence of estrogen (zero M) and estrogen deprivation (1×10^{-9} M) were not significantly different ($P > 0.05$) to the negative control.

Physiological and supraphysiological levels of estrogen significantly ($P < 0.05$) reduced MRSA and *P. aeruginosa* recovery by HPBM-derived M1 and M2 macrophages compared to the negative control. After a 3-hour incubation with HPBM-derived M1 macrophages, the MRSA recovery significantly ($P < 0.05$) decreased from 2825 CFU/mL in the negative control to 2023 CFU/mL and 1343 CFU/mL following treatment with physiological and supraphysiological levels of estrogen respectively. Although significantly less effective than M1 macrophages, treatment of HPBM-derived M2 macrophages with physiological or supraphysiological levels of estrogen still significantly ($P < 0.05$) reduced the recovery of MRSA to 2745 CFU/mL and 2437 CFU/mL respectively compared to the negative control (3635 CFU/mL).

Similarly, treating HPBM-derived M1 macrophages with physiological or supraphysiological levels of estrogen significantly ($P < 0.05$) reduced *P. aeruginosa* recovery to 2430 CFU/mL and 2083 CFU/mL respectively compared to the negative control (3352 CFU/mL). Although somewhat less effective, treating HPBM-derived M2 macrophages with physiological or supraphysiological levels of estrogen still significantly ($P < 0.05$) reduced the recovery of *P. aeruginosa* to 3062 CFU/mL and 2339 CFU/mL respectively compared to the negative control (4052 CFU/mL).

Intriguingly, the MRSA and *P. aeruginosa* uptake was significantly ($P < 0.05$) greater with both U937-derived and HPBM-derived M1 macrophages compared to M2 macrophages.

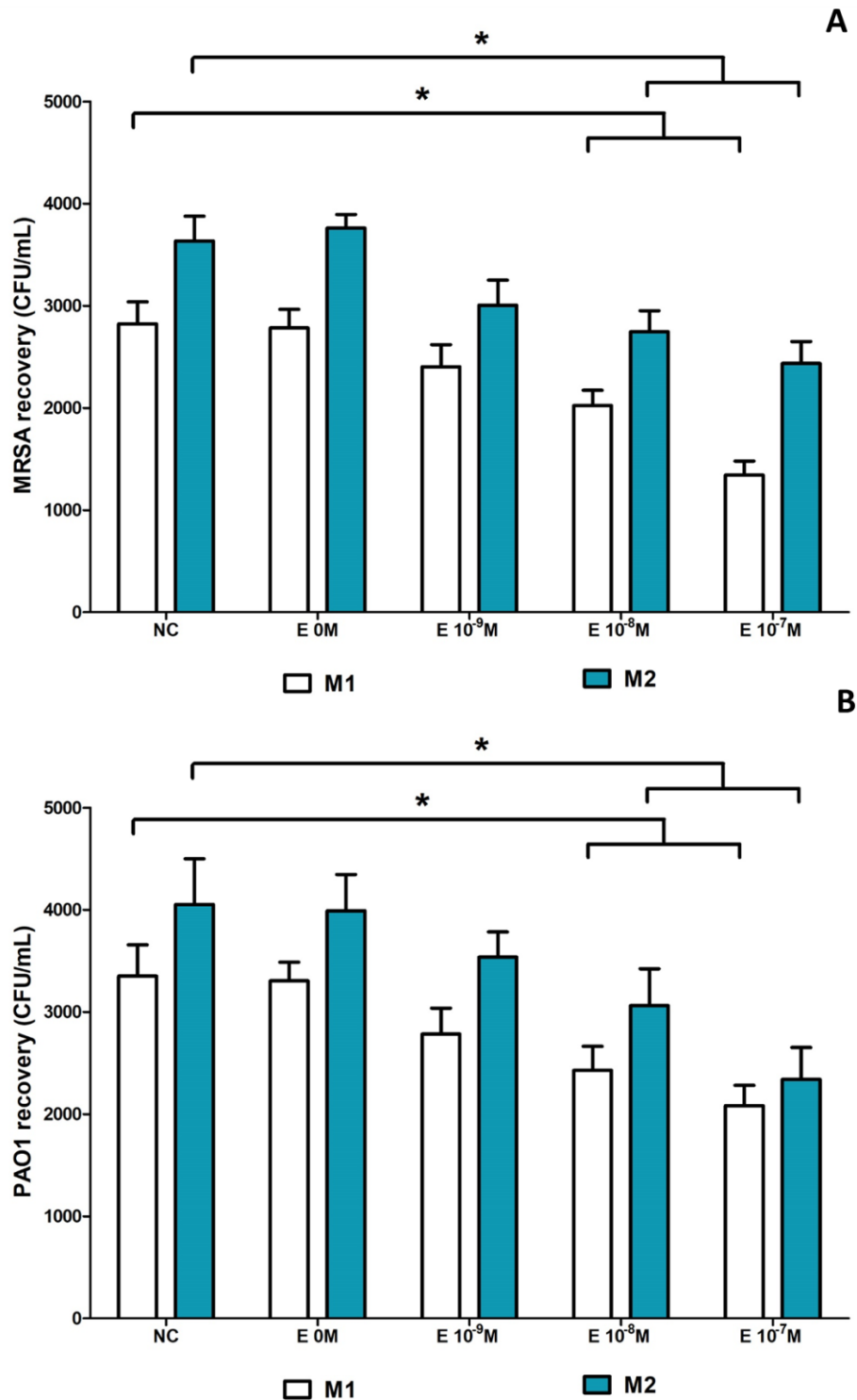


Figure 4.7. Estrogen improves the phagocytosis of MRSA (A) and PAO1 (B) by *ex vivo* HPBM-derived M1/M2 macrophages. The recovery of MRSA (A) and *P. aeruginosa* (B) after incubation with HPBM-derived M1/M2 macrophages were significantly (*: $P < 0.05$, **: $P < 0.01$) decreased following treatment with physiological (1×10^{-8} M) or supraphysiological (1×10^{-7} M) levels of estrogen compared to the negative control. HPBM-derived M1 macrophages internalised significantly ($P < 0.05$) more MRSA/PAO1 respectively than HPBM-derived M2 macrophages. Data is mean \pm StEM, $n = 10$.

4.5 Discussion

Plasticity of macrophages is a crucial process for an effective inflammatory phase during wound healing. Macrophages ability to continuously switch from a phenotype to another was demonstrated in mice (Thomas *et al.*, 1992; Erwig *et al.*, 1998). Experiments in this study were conducted using pro-inflammatory (M1-like) and anti-inflammatory (M2-like) macrophages. *In vitro* M1 macrophages can be induced by differentiation with IFN- γ and/or LPS, and distinguished through the expression of the surface markers CD197 and/or CD86 (Mantovani *et al.*, 2004). In contrast, M2 macrophages can be induced by differentiation with IL-4 and/or IL-13, and distinguished through the expression of the macrophage mannose receptors, CD206 (Stein *et al.*, 1992; Daigneault *et al.*, 2010) and/or CD163 (Zeyda *et al.*, 2007). Successful polarisation of U937-derived, and HPBM-derived M0 macrophages into M1-like and M2-like macrophages was achieved in this study, with high expression of CD197 confirming polarisation into M1-like macrophages by IFN- γ /LPS, and high CD206 expression confirming polarisation into M2-like macrophages by IL-4/IL-13.

The effect of estrogen on the phagocytosis of MRSA and *P. aeruginosa* by U937-derived M1-like and M2-like macrophages was investigated *in vitro*. In line with findings of chapter 3, the results from this chapter demonstrated that the physiological and supraphysiological levels of estrogen enhanced the clearance of both MRSA and PAO1 in a dose-dependent manner in both U937-derived M1 and M2 macrophages. Interestingly, the bacterial recovery followed a similar pattern throughout, regardless of the macrophage phenotype, with estrogen consistently increasing ($P < 0.05$) the phagocytosis of bacteria. These results are supported by the findings of Crompton *et al.* (2016), when estrogen was found to significantly enhance the rate of wound repair in mice in a *K. pneumonia* LPS model of infection, and by Li *et al.* (2000) when a 48-hour stimulation with estrogen was found to enhance the uptake of amyloid β -peptide and *E. coli* by macrophages extracted from human brain tissues (microglial cells). Cutolo *et al.* (2004) also described estrogen as a key regulator of immune responses against bacterial infections in macrophages as estrogen was found to enhance the expression of growth and proliferation markers in macrophages in patients with autoimmune diseases.

It is believed that during early stages of wound healing, M1 macrophages are highly phagocytic, and serve as cleaners of the wound site by removing foreign invaders and dead tissue (Stein *et al.*, 1992) whereas M2 macrophages have anti-inflammatory properties and play a more regenerative role as the inflammatory response phase subsides (Mantovani *et al.*, 2004). In line with this, M1 macrophages were described to be highly phagocytic *in vitro*, and produce microbicide enzymes such as iNOS, essential for pathogen destruction. They also release many pro-inflammatory cytokines, such as, IL-1 β , IL-6, IL-12, IL-23 and TNF- α (Mantovani *et al.*, 2004). Werner and Grose (2003) reported an up-regulation of these cytokines during the inflammatory phase of wound repair. In addition, research indicated that genetic deficiencies of IFN- γ and/or its receptors confirmed that macrophages polarisation to the M1 phenotype is essential for host defence against pathogenic infections in knockout mice models and in humans. Components of intracellular pathogens (e.g. LPS), TNF, GM-CSF and TLR4 ligands were also described to encourage M1 polarisation (Mantovani *et al.*, 2005). Collectively, data from the host-pathogen assays support these reports, with significantly enhanced clearance of MRSA and *P. aeruginosa* by M1 macrophages compared to M2 macrophages or M0 macrophages (results in chapter 3).

The assays were conducted first using the immortal cell line “U937 monocytes”. *In vitro* results would provide an insufficient evidence to predict effects *in vivo*. To avoid this limitation, primary monocytes derived from human peripheral blood were used to conduct an *ex vivo* model of host-pathogen interactions. Unsurprisingly, the findings were consistent with the *in vitro* assays. The bacterial recovery was shown to decrease with increasing estrogen concentrations with both HPBM-derived M1 and M2 macrophages, with a significantly greater uptake of bacteria with M1 macrophages compared to M2 macrophages. This suggests enhanced phagocytosis is a feature of M1 macrophages, regardless of their source. In support of these results, Deshpande *et al.* (1997) reported that estrogen causes an increase in the production of iNOS by murine macrophages as well as many other cytokines, particularly, IL-1 β , IL-12 and IL-6. This is due to TLR4 activation with LPS in murine macrophages. In contrast, other studies reported many anti-

inflammatory effects of estrogens on macrophages *in vitro*. Indeed, a 2-hour estrogen treatment of macrophages (RAW 264.7 cell line) *in vitro* caused a substantial diminution in the expression of inflammatory mediators, mainly NF- κ B, subsequent to TLR4 activation (Ghisletti *et al.*, 2005). Long-time (chronic) exposure to estrogen was found to promote the alternative activation of macrophages and to dampen the IFN- γ /LPS actions on human macrophages (Campbell *et al.*, 2014; Toniolo *et al.*, 2015). In this study, a 24-hour incubation with estrogen was shown to increase the rate of bacterial phagocytosis in all macrophages regardless the macrophage subtype. Further investigations are warranted to determine whether estrogen have an effect on the polarisation of macrophages *in vitro*. One application will be to measure the expression of the M1-like and M2-like specific markers (e.g. CD197 and CD206 respectively) before and after treatment of macrophages with estrogen.

Human trials have confirmed that estrogen supplementation accelerates healing by reducing inflammatory response in the elderly (Deshpande *et al.*, 1997; Srivastava *et al.*, 1999) but further *in vivo* investigations are needed to demonstrate the effect of estrogen on bacterial clearance in elderly subjects. Investigations could also be conducted using existing animal models of age-related impaired healing (Deshpande *et al.*, 1997; Ghisletti *et al.*, 2005) to confirm *in vivo* effects of estrogen on the phagocytic function of M1 and M2 macrophages.

Billiau and Vandekerckhove (1991) reported that bacterial clearance in wounds is reliant on good regulation of TNF- α . Interestingly, estradiol was shown to suppress the production of TNF- α in murine macrophages (Deshpande *et al.*, 1997; Srivastava *et al.*, 1999). The work presented in this chapter suggests that TNF- α could be associated with the mechanism of estrogen-enhanced phagocytosis in M1 macrophages. Further research is warranted to investigate the change in TNF- α levels and other pro-inflammatory cytokines (e.g. MIF) involved in the estrogen-increased phagocytosis in macrophages.

Compared to results reported in chapter 3, M1-like macrophages were shown to be more phagocytic than M0-like macrophages regardless of the source of macrophage. Further work investigating mechanisms of estrogen-enhanced

phagocytosis in this thesis will therefore be conducted using pro-inflammatory M1-like macrophages instead of M0-like or M2-like macrophages. Investigations were conducted (chapter 5) to determine the ER pathways linked with the estrogen-mediated enhancement of phagocytosis through the use of: selective estrogen receptor modulators (SERMs) (e.g. Tamoxifen), selective estrogen receptor degraders (SERDs) (e.g. Fulvestrant), selective ER- α and/or ER- β agonists (e.g. PPT, DPN) to determine the exact ER activation profile involved in the estrogen-enhanced phagocytosis. In order to investigate the mechanism by which estrogen enhanced bacterial clearance in M1 macrophages, the changes in the levels of proteins that regulate the actin cytoskeleton dynamics including Rho GTPase proteins, and focal adhesion proteins on the cell membrane of estrogen-treated M1 macrophages are presented (detailed in chapter 6).

In summary, these findings confirm for the first time that estrogen deprivation inhibits phagocytosis by both M1-like and M2-like human macrophages whereas physiological/supraphysiological levels of estrogen stimulate the phagocytic function of M2 but particularly M1 human macrophages against Gram-positive and Gram-negative wound pathogens. Together, the results suggest topical or systemic estrogen supplementation may potentially resolve or reduce the likelihood of wound infections in the elderly.

**Chapter 5: Estrogen Enhances the
Phagocytosis of MRSA and *P.*
aeruginosa by Human M1
Macrophages via Binding to ER- α**

5.1 Introduction

5.1.1 Estrogen Receptors

Estrogen signals principally through the estrogen receptor (ER) proteins that are part of the nuclear receptor (NR) family. ERs display a structure that is typical of the NR family, containing six domains (A-F) (Kuiper *et al.*, 1998; Klinge, 2000; Begam *et al.*, 2017) (Figure 5.1).

ER-alpha (ER- α) was first discovered in 1958 and cloned in 1985 (Jensen, 1962; Kuiper *et al.*, 1998). However, ER-beta (ER- β) was first characterised in 1996 in rat prostate and ovary. Later in 1998, a human ER- β was identified (Mosselman *et al.*, 1996; Kuiper *et al.*, 1998). Both ERs are located in various tissues (Campbell *et al.*, 2010). ER- α is highly expressed in ovaries, cancer breast tissues and reproductive tissues (Kuiper *et al.*, 1997; Ali and Coombes, 2000; Campbell *et al.*, 2010). ER- β is mainly found in peripheral tissues such as bone, heart and hypothalamus, but it can also be found in healthy breast tissue, the prostate and testis (Kuiper *et al.*, 1997; Ali and Coombes, 2000; Campbell *et al.*, 2010). ERs are expressed in the skin, suggesting that estrogen regulates skin functions and development (Ashworth, 2005). Both ER- α and ER- β have been identified in the epidermis of young and old women, with isoforms observed in epithelial cells of the hair follicles (Ashworth, 2005).

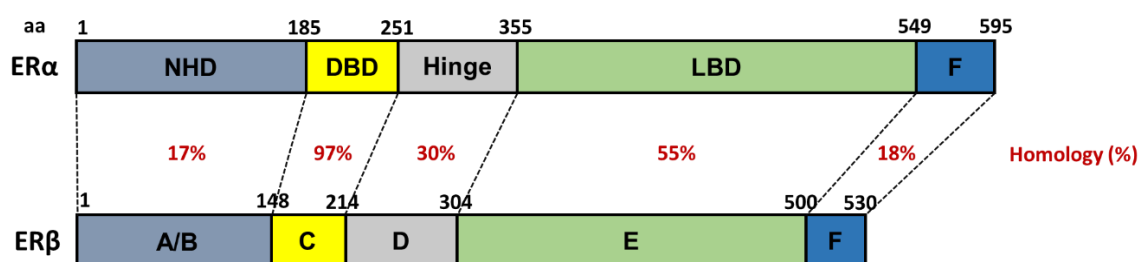


Figure 5.1. Schematic representation comparing the structure of human ER- α with ER- β . Homology between domains (A-F) is represented as percentage (%) similarity. NHD = N-terminal homology domain, DBD = DNA-binding domain, LBD = ligand-binding domain. Figure adapted from (Webb *et al.*, 1999; Klinge, 2000; Begam *et al.*, 2017).

Estrogen signals predominantly by binding to inactive ERs in the nucleus of the cell (Klinge, 2000). When estrogen binds to ERs, they become activated and dimerize (Klinge, 2000). Each ER has a DNA-binding domain (DBD) which binds to the target

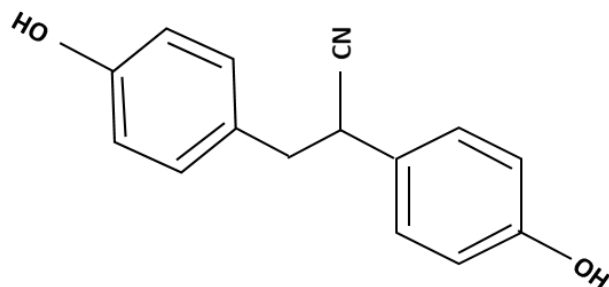
gene at a site called the estrogen response element (ERE) (Kuiper *et al.*, 1998; Klinge, 2000). In cells expressing a single ER subtype, homodimers of ER- α or ER- β are formed (Kuiper *et al.*, 1998). In cells that express both ER subtypes, a heterodimer containing one ER- α and one ER- β may form (Kuiper *et al.*, 1998). ERs heterodimers and ER- α homodimers bind to DNA with a similar affinity. However, ER- β homodimers bind to DNA with a lower affinity (Kuiper *et al.*, 1998). DNA-bound homodimers and heterodimers bind to steroid receptor coactivator-1 and trigger gene transcription (Marino *et al.*, 2006).

Both ER- α and ER- β enhance acute wound repair, however, their functions are different; ER- α is described to control and regulate inflammation, however, ER- β is linked with the modulation of wound healing (Campbell *et al.*, 2010). While ER- α and ER- β express a 97% homology in the DBD region, they only have 55% homology in the LBD region (Barkhem *et al.*, 1998; Webb *et al.*, 1999; Klinge, 2000). This made possible the design of artificial estrogens such as SERMs with specific binding affinity to one isoform over another.

5.1.2 Estrogen Receptor Agonists

Estrogen receptor agonists are molecules that induce one or both ER subtypes in order to partially or fully induce a pharmacological effect (Sun *et al.*, 1999). Selective agonists of ER- α and ER- β are extensively used to investigate the distinct functions of the ERs in several human diseases. Propyl pyrazole triol (PPT) and Diarylpropionitrile (DPN) are the most frequently used selective ERs agonists. PPT is a selective ER- α agonist (Thammacharoen *et al.*, 2009) that has 410-fold more selectivity for ER- α than ER- β (Thammacharoen *et al.*, 2009). In 1999, PPT was used for the first time to investigate the role of ER- α in several tissues (Sun *et al.*, 1999). DPN is a selective ER- β agonist (Minutolo *et al.*, 2011) that has 70-fold more selectivity for ER- β than ER- α (Meyers *et al.*, 2001). DPN was first used in 1999 in a purpose to develop new ligands selective for ER- β (Sun *et al.*, 1999; Meyers *et al.*, 2001) (Figure 5.2).

A



B

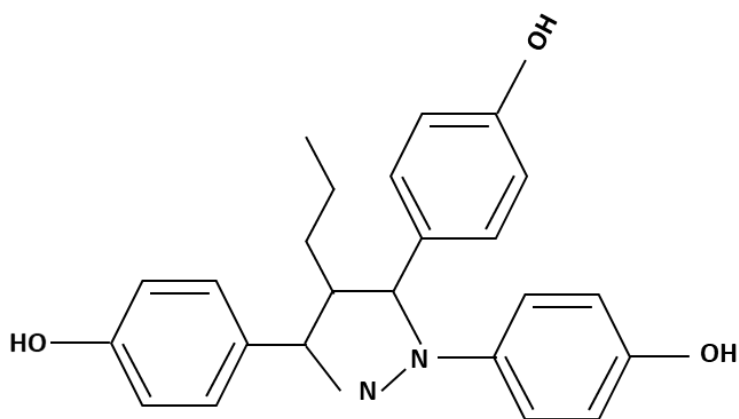


Figure 5.2. Structure of DPN (A) and PPT (B).

5.1.3 Estrogen Receptor Antagonists

Estrogen receptor antagonists are compounds characterised to have high affinity to bind to one or both of the ER subtypes, without the potential of inducing the normal pharmacological effects typically mediated by the receptors (Wakeling *et al.*, 1991; Barkhem *et al.*, 1998). ER antagonists compete with estrogen to bind to the specific receptor site (Morris and Wakeling, 2002; Osborne *et al.*, 2004). Therefore, the physiological effect produced in response to estrogen will be dampened in the presence of ER antagonists. ER antagonists have been used clinically to control fertility or treat various cancers (Morris and Wakeling, 2002; Osborne *et al.*, 2004).

Theophylline, 8-[(benzylthio)methyl]-7,8-dimethylxanthine (TPBM) is a potent inhibitor of ER- α via blocking the binding of ER- α to the ERE DNA (Mao *et al.*, 2008). TPBM is highly selective for ER- α compared to ER- β , with a greater selectivity for blocking gene transcription mediated by ER- α relative to ER- β (Mao *et al.*, 2008). TPBM has been used to investigate the role of ER- α in many biological processes and disorders including cancer (Mann, 2008; Plant and Zeleznik, 2014).

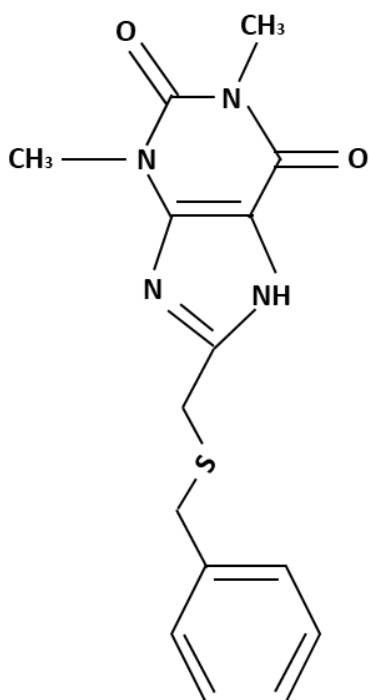


Figure 5.3. Structure of TPBM.

5.1.4 Selective Estrogen Receptor Modulators (SERMs)

Selective estrogen receptor modulators (SERMs) are ER-interacting molecules that have the ability to bind the ERs and act as agonists in specific tissues whilst acting as antagonists in different tissues (Cho and Nuttall, 2001). SERMs have been used clinically to promote the beneficial effects of estrogen in target tissues whilst avoiding/decreasing detrimental effects of estrogen in other non-target tissues such as increasing the risk of breast cancer in breast tissues (Mirkin and Pickar, 2015). SERMs can be differentiated from ER agonists/antagonists due to their opposing effects in tissues (Park and Jordan, 2002). SERMs are structurally different to estrogens but their chemical structures allow them to bind to the LBD of ERs (Brzozowski *et al.*, 1997).

Tamoxifen (Figure 5.4) is one of the most documented and frequently used SERMs. It is known to have estrogenic effects in numerous peripheral tissues such as bone, but is anti-estrogenic in the breast tissue and therefore used extensively in breast cancer research (Furr and Jordan, 1984; Morris and Wakeling, 2002; Park and Jordan, 2002; Mirkin and Pickar, 2015). Tamoxifen was discovered and reported by the Food and Drug Administration (FDA) in 1977 (Park and Jordan, 2002; Jordan, 2006; Mirkin and Pickar, 2015; Quirke, 2017). Tamoxifen binds to both ERs, and its effects depend on the cell and tissue types. It is anti-estrogenic in the breast, and is therefore commonly used to prevent and/or treat breast cancer in postmenopausal and premenopausal females (Zidan *et al.*, 2004; Quirke, 2017). Tamoxifen has also been reported to maintain the density of bone in rats and humans (Jordan *et al.*, 1987; Zidan *et al.*, 2004). However, it has multiple side effects and is frequently linked with endometrial cancer due to its estrogenic effects in the uterus (Kedar *et al.*, 1994).

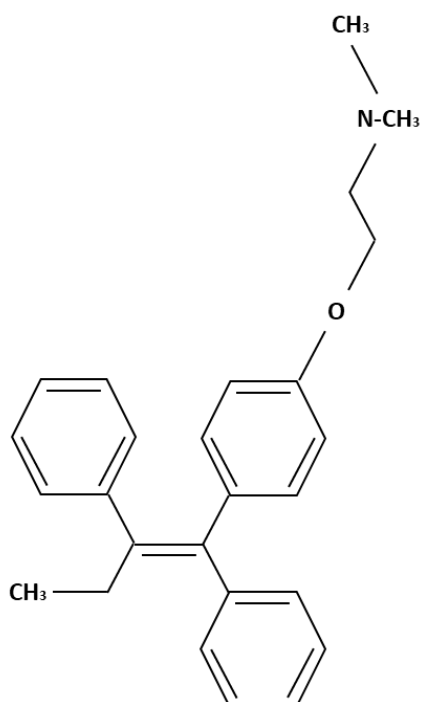


Figure 5.4. Structure of Tamoxifen.

5.1.5 Selective Estrogen Receptor Degradors (SERDs)

Selective estrogen receptor degraders (SERDs) are molecules that have high affinity to bind ERs but lead to their degradation, leading to reduced levels of functional

ERs (Lee *et al.*, 2017). Along with SERMs and aromatase inhibitors, SERDs are mainly used to treat breast cancer (Lee *et al.*, 2017). Fulvestrant, the most commonly used SERD, was officially discovered in 2002 (Lee *et al.*, 2017). In November 2016, other SERDs such as elacestrant and brilanestrant were developed (Lai and Crews, 2017). Fulvestrant is an analogue and competitive inhibitor of 17 β -estradiol (Osborne *et al.*, 2004), with a high selective binding affinity (89% that of estrogen) to the ER (Morris and Wakeling, 2002). The binding of fulvestrant to the ER weakens the receptor dimerization and blocks the nuclear localisation of ER (Morris and Wakeling, 2002; Osborne *et al.*, 2004). In addition, the fulvestrant-ER complex is unstable, which leads to its rapid degradation leading to enhanced cellular ER deprivation compared with estrogen or other SERMs (e.g. tamoxifen) (Morris and Wakeling, 2002). The reduced levels of functional ER protein does not cause any decrease in the levels of ER mRNA. Therefore, after binding to, blocking and accelerating the destruction of the ER protein, fulvestrant inhibits estrogen signalling (Morris and Wakeling, 2002; Osborne *et al.*, 2004).

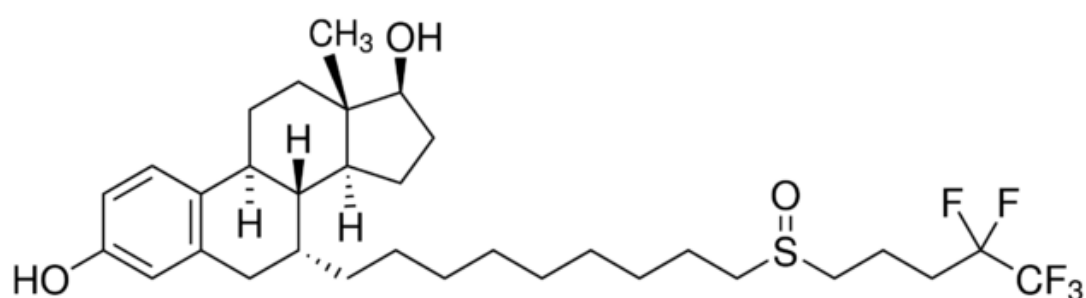


Figure 5.5. Structure of Fulvestrant.

5.1.6 Estrogen Receptors and Wound Infection

ER- α and ER- β are both expressed by inflammatory cells, including neutrophils and macrophages (Ashcroft *et al.*, 2003). More widely, ER- α is expressed in human skin, fibroblasts and the adipose tissue (Kuiper *et al.*, 1997; Kuiper *et al.*, 1998; Campbell *et al.*, 2010; Campbell *et al.*, 2014). ER- β is expressed in the dermis, epidermis, fibroblasts, endothelial cells and muscle cells (Thornton *et al.*, 2003).

Estrogen is key molecule that enhances wound healing in both genders and these actions appear to be mediated via the ERs (Ashcroft *et al.*, 1999a). Age-related

estrogen deprivation is frequently linked with delayed wound healing and might contribute to the development of chronic wounds in the elderly (Ashcroft and Ashworth., 2003). In particular, polymorphisms of the ER- β gene are associated with venous ulceration and a pronounced inflammatory response (Ashworth, 2005). ER- α has been implicated in reducing the inflammatory response in wounds by dampening levels of pro-inflammatory cytokines such as macrophage migration inhibitory factor (MIF) (Campbell *et al.*, 2010). In contrast, ER- α activation by PPT has been shown to considerably reduce early wound strength in models of ovariectomized mice compared to untreated ovariectomized mice, whereas activation of ER- β with DPN had no effect on early wound strength (Gál *et al.*, 2010). Although the involvement of ER subtypes in the wound healing process is documented, there is a lack in knowledge regarding the precise roles of ER- α and ER- β in the clearance of bacteria from infected wounds.

Ashcroft *et al.* (1999a) showed that estrogen replacement therapy accelerates wound repair in postmenopausal females. Furthermore, this study has showed estrogen enhances the clearance of the typical wound pathogens MRSA and *P. aeruginosa* (chapter 3 and 4). However, Anderson *et al.* (2004) reported that estrogen supplementation in postmenopausal females is associated with a higher risk of breast cancer and heart diseases. Thus, rather than using systemic estrogen supplementation, the use of appropriate ERs agonists/antagonists, SERDs and particularly SERMs could be beneficial in treatment of age related impaired healing and infected wounds.

5.2 Aim and Objectives

5.2.1 Aim

To investigate the involvement of ERs in mediating the beneficial effects of estrogen on promoting phagocytosis by human M1 macrophages using *in vitro* and *ex vivo* models of host-pathogen interactions.

5.2.2 Objectives

- Determine the effect of ERs agonists and antagonists, SERDs and SERMs on the phagocytosis of Gram-positive (MRSA) and Gram-negative (*P. aeruginosa*) bacteria by U937- and HPBM-derived M1 macrophages.

5.3 Methods

5.3.1 Estrogen Receptors Stimulation/Blockade

In order to test the effect of ERs agonists/antagonists, SERMs and SERDs on the MRSA and *P. aeruginosa* clearance, *in vitro* and *ex vivo* host-pathogen assays were conducted as described in section 2.2.12.

5.4 Results

Similar to previous results, supraphysiological estrogen levels significantly decreased ($P < 0.01$) MRSA and *P. aeruginosa* recovery compared to the negative control and the absence of estrogen (Figure 5.6). Treatment with the ER agonist tamoxifen mirrored the effects of estrogen, significantly ($P < 0.01$) decreasing the bacterial recovery of both MRSA and *P. aeruginosa*. In contrast, treatment with the ER degrader fulvestrant blocked the effects of supraphysiological levels of estrogen on the phagocytosis of MRSA and *P. aeruginosa*, with no significant difference in bacterial recovery ($P > 0.05$) detected compared to the negative control and the absence of estrogen. Of interest, ER- α agonism with PPT significantly reduced ($P < 0.01$) the recovery of both MRSA and *P. aeruginosa* in fashion similar to supraphysiological estrogen levels when compared to the negative control and the absence of estrogen. In contrast, ER- β stimulation with DPN did not significantly ($P > 0.05$) affect the uptake of MRSA and *P. aeruginosa* by M1 macrophage compared to the negative control or the absence of estrogen. Intriguingly, the selective ER- α antagonist, TPBM, reversed the effect of estrogen and PPT on the phagocytosis of MRSA and *P. aeruginosa*, with no significant difference ($P > 0.05$) in bacterial recovery detected compared to the negative control or the absence of estrogen.

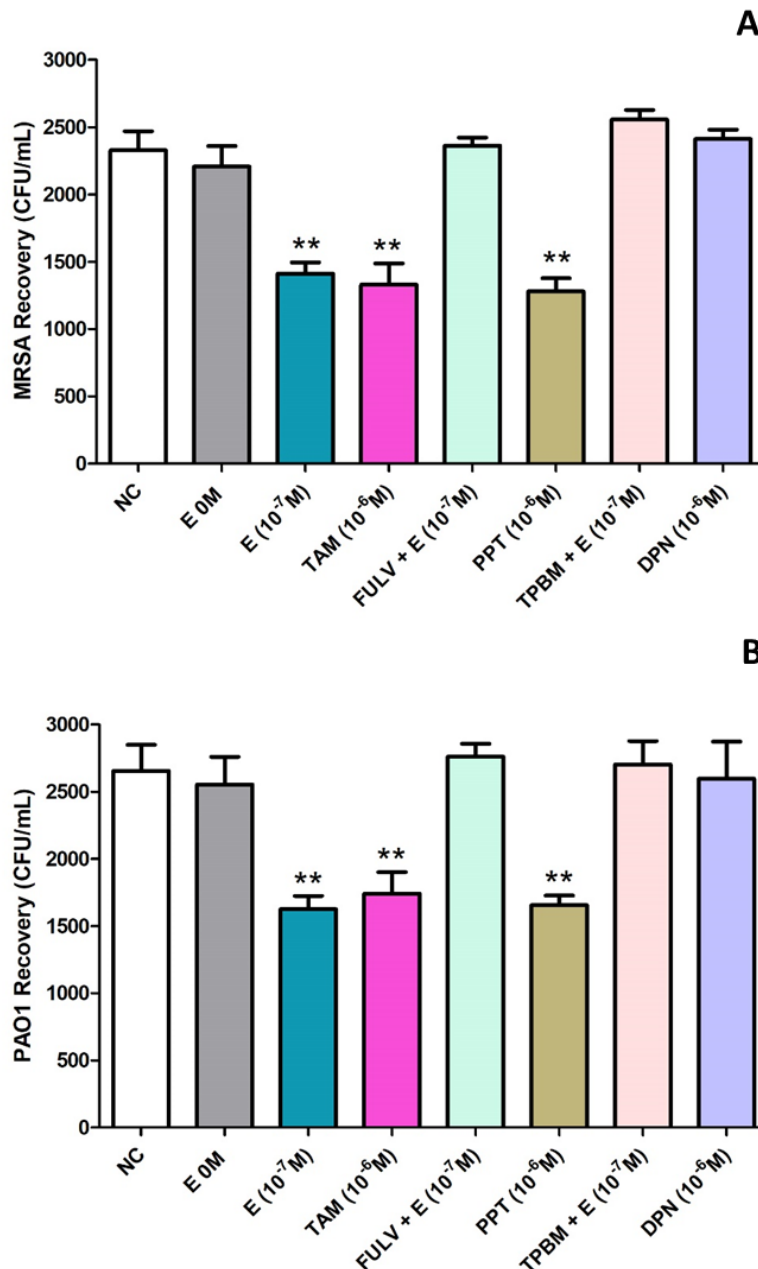


Figure 5.6. Estrogen, tamoxifen and the selective ER- α agonist, PPT, enhances the phagocytosis of MRSA (A) and *P. aeruginosa* (B) by U937 M1 macrophages. Estrogen, tamoxifen and PPT significantly decreased the MRSA (A) and *P. aeruginosa* (B) recovery compared to the negative control or the absence of estrogen (E 0M). Fulvestrant and TPBM significantly blocked the effects of estrogen on the phagocytosis of both MRSA and *P. aeruginosa*, with no significant difference in bacterial recovery detected compared to the negative control or the absence of estrogen. DPN did not significantly affect bacterial uptake compared to the negative control or the absence of estrogen. Data represent an average of $n = 6$ experiments. ** Indicates significant difference ($P < 0.01$). Error bars represent the standard error of the mean (StEM).

The *in vitro* host-pathogen assays were repeated using *ex vivo* primary HPBM-derived M1 macrophages in order to confirm these initial findings (Figure 5.7). The

results were similar to those obtained with U937-derived M1 macrophages, with estrogen, tamoxifen and PPT significantly ($P < 0.01$) reducing MRSA and *P. aeruginosa* recovery compared to the negative control and the absence of estrogen. Again, fulvestrant and ER- α antagonism with TPBM blocked the effects of estrogen whereas ER- β agonism with DPN had no significant ($P > 0.05$) effect on the phagocytosis of MRSA and *P. aeruginosa* compared to the negative control or the absence of estrogen.

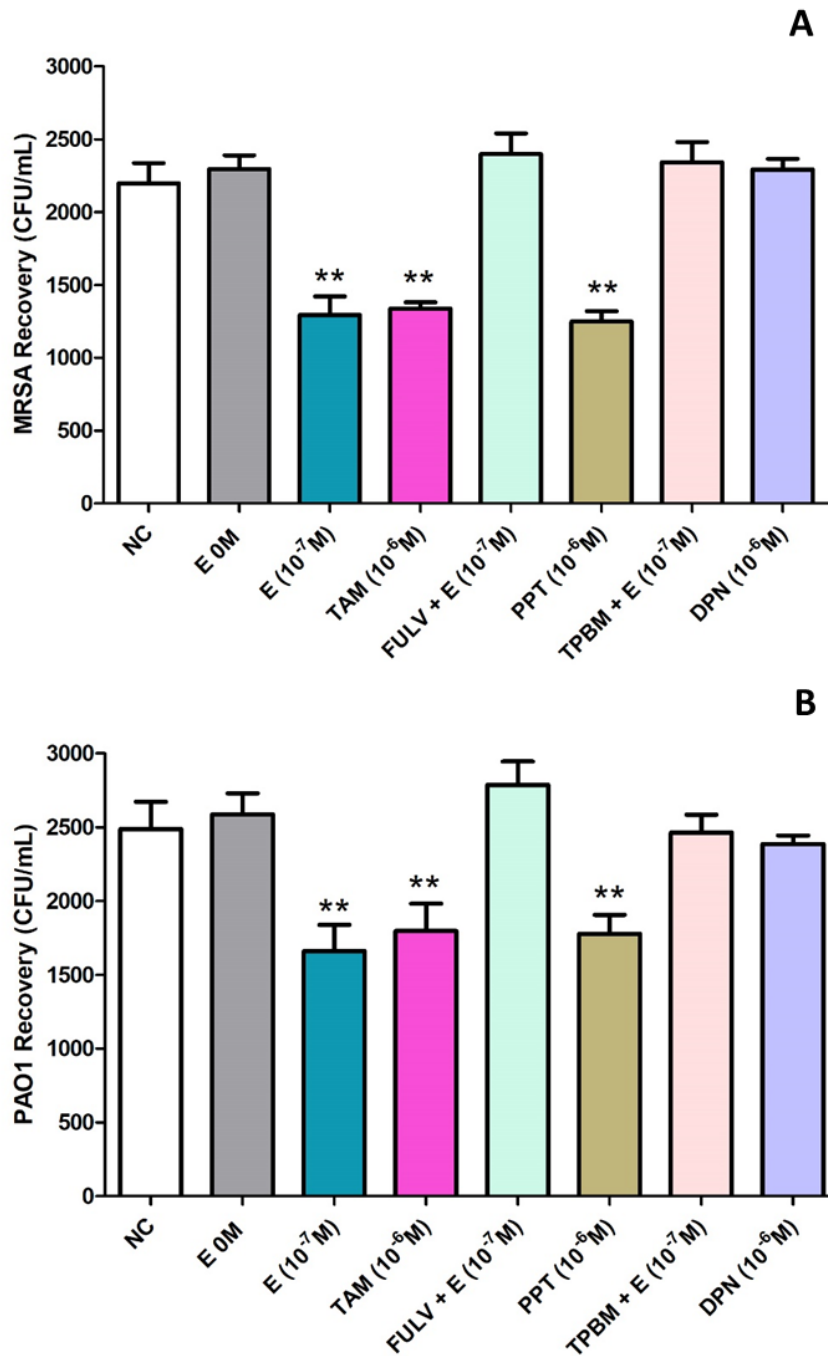


Figure 5.7. Estrogen, tamoxifen and the selective ER- α agonist, PPT, enhance MRSA (A) and *P. aeruginosa* (B) uptake by HPBM M1 macrophages. Estrogen, tamoxifen and PPT significantly decreased the recovery of MRSA (A) and *P. aeruginosa* (B) compared to the negative control and the absence of estrogen (E 0M). Fulvestrant and TPBM significantly blocked the effects of estrogen on the phagocytosis of both MRSA and *P. aeruginosa*, with no significant difference in bacterial recovery detected compared to the negative control or the absence of estrogen. DPN did not significantly affect bacterial uptake compared to the negative control or the absence of estrogen. Data represent an average of $n = 6$ experiments. ** Indicates significant difference ($P < 0.01$). Error bars represent the standard error of the mean (StEM).

5.5 Discussion

The age-related change in estrogen levels is linked with skin degeneration and impaired wound healing (Ashcroft *et al.*, 1999b). Estrogen is frequently linked with pathological wounds (Ashcroft and Ashworth., 2003) and a case-cohort study reported that elderly (over 65 years old) patients treated with estrogen were 30-40% less likely to develop a venous leg ulcer than untreated patients (Margolis *et al.*, 2002).. Moreover, estrogen deprivation diminishes the phagocytic capability of neutrophils and macrophages, and increases the risk of infection and a postponement in healing (Ashcroft and Ashworth., 2003).

Due to their possession of nuclear ERs, the function of inflammatory cells such as macrophages are known to be influenced directly by estrogen (Weusten *et al.*, 1986; Suenaga *et al.*, 1996; Suenaga *et al.*, 1998) through its binding to ER- α and/or ER- β (Kuiper *et al.*, 1998; Klinge, 2000). SERMs and SERDs are ER-interacting molecules that are able to bind to the ERs and achieve normal mechanisms of estrogen signalling to function as agonists or antagonists of estrogen in a tissue-specific manner (Cho and Nuttall, 2001). SERMs and SERDs are compounds typically used to treat several ER-mediated pathologies (e.g. breast cancer) due to their tissue-specific responses (Mirkin and Pickar, 2015; Lee *et al.*, 2017). Estrogen stimulated the phagocytic function of human macrophages against MRSA and *P. aeruginosa* (Chapter 3 and 4). This study used SERMs and SERDs to investigate which ERs are involved in the estrogen-mediated stimulation of bacterial clearance by M1-like macrophages.

An accumulating body of evidence demonstrates that tamoxifen acts as ER antagonist in breast tissue, but is agonist in the uterus and peripheral tissues such as bone (Fisher *et al.*, 2005; DeMichele *et al.*, 2008). Tamoxifen enhanced the phagocytosis of MRSA and *P. aeruginosa* by human M1 macrophages suggesting tamoxifen was acting in a similar manner to estrogen. In support of this, tamoxifen was shown to enhance neutrophils chemotaxis and phagocytosis functions, boost human neutrophil bactericidal ability against a range of bacteria *in vitro* and

improve clearance of the leading nosocomial antibiotic-resistant bacteria, MRSA, *in vivo* (Corriden *et al.*, 2015).

Tamoxifen has been shown to accelerate wound healing in ovariectomized mice, suggesting that it can potentially be used to treat impaired wound repair in the elderly (Hardman *et al.*, 2007). This study adds weight to this theory by showing tamoxifen may be beneficial in the treatment of infected wounds in the elderly. Pickar *et al.* (2010) have detailed the beneficial effects of SERMs on females health after menopause, SERMs such as tamoxifen are more beneficial during wound healing when they are applied locally on wounds rather than orally. A new group of SERMs, such as Ophena, are now developed to mimic the positive biological and tissue specific estrogen functions but not the undesirable side effects such as the risk of breast and uterine cancers (Pickar *et al.*, 2010).

Fulvestrant is a compound structurally different to SERMs that due to its high binding affinity to the ERs, binds in competition with estrogen to the ERs (Morris and Wakeling, 2002). When bound to ERs, fulvestrant deteriorates the receptor dimerization and blocks nuclear localisation of the ERs (Morris and Wakeling, 2002; Osborne *et al.*, 2004). The findings showed that, unlike treatment with estrogen or tamoxifen, pre-treatment of M1 macrophages with fulvestrant blocked stimulation of bacterial clearance by estrogen. This provided additional evidence that estrogen may be acting through the ERs to promote phagocytosis and that disruption of the ERs prevents the beneficial effect of estrogen on bacterial clearance. Collectively, data generated from the use of tamoxifen and fulvestrant provide evidence that estrogen is enhancing the phagocytosis of bacteria, by pro-inflammatory M1 macrophages via binding and activation of the ERs.

It has been reported that the anti-proliferative effects of fulvestrant on cells in breast are due to the degradation of the ER- α protein (Long and Nephew, 2006). Interestingly, it has been reported that ER- α is the key receptor in regulating the thickness of epidermis during wound healing (Stumpf *et al.*, 1974; Bidmon *et al.*, 1990). Toutain *et al.* (2009) reported that ER- α is also the main isoform involved in promoting the beneficial effects of 17 β -estradiol in a model of skin necrosis. On the other hand, ER- β was described to be the key receptor in the protection against

skin damage resulted from exposure to UV (Chang *et al.*, 2010). Both ER subtypes are expressed on fibroblasts and inflammatory cells such as neutrophils and macrophages (Campbell *et al.*, 2010). ER- α has been noted to control and regulate inflammation, whereas ER- β is mainly linked with modulating the later phases of wound healing (Campbell *et al.*, 2010). This study used two major agonists of ER- α (PPT) and ER- β (DPN) to investigate the effect of activating ER isoforms on the phagocytosis of MRSA and *P. aeruginosa*. PPT is an ER- α agonist that is 410-fold more selective for ER- α than ER- β whereas DPN is an ER- β agonist with a 70-fold higher selectivity for ER- β than ER- α (Campbell *et al.*, 2010). Intriguingly, selective activation of ER- α with PPT mirrored the effects of estrogen by significantly ($P < 0.01$) increasing the phagocytosis of both MRSA and *P. aeruginosa* by U937-derived and HPBM-derived M1 macrophages. In contrast, stimulation with the ER- β agonist DPN failed to improve the phagocytosis of either MRSA or *P. aeruginosa* by macrophages. Of interest, blocking the ER- α with TPBM overturned the effect of PPT on the bacterial clearance. The results of this study are in concordance with published findings implicating ER- α as the key mediator of the inflammatory response, with activation of ER- α in M1 macrophages with PPT increasing bacterial clearance whereas activation of ER- β by DPN had no effect on the phagocytosis of MRSA and PAO1. Moreover, blockade of ER- α with TPBM reversed the effect of PPT. Collectively, these novel findings suggest that estrogen stimulates promotion of phagocytosis by human macrophages through ER- α rather than ER- β .

Tamoxifen is estrogenic in the uterus, therefore, this molecule is frequently linked with a high risk of uterine cancer (Jordan and Morrow, 1999; Bergman *et al.*, 2000). Another commonly used SERM, called raloxifene, is in contrast estrogenic in bone (Heaney and Draper, 1997) and ovaries (Neven *et al.*, 2002), but displays anti-estrogenic effects in the uterus (Bryant *et al.*, 1996). Hardman and Ashcroft (2008) indicated that estrogen, tamoxifen and raloxifene all accelerated wound healing in ovariectomised mice. Future research could investigate a range of SERMs including raloxifene to determine the most effective mediator(s) of bacterial clearance in macrophages. This might potentially help detecting the most promising topical SERM to treat wound infections in the elderly.

In summary, novel findings reported in this chapter provide clear evidence that ER- α is the ER isoform through which estrogen stimulates phagocytosis by U937- and HPBM-derived M1 macrophages. Selective targeting of ER- α with agents such as PPT may potentially lead to the development of therapeutic approaches to combat wound infections in the elderly.

Chapter 6: Estrogen Enhances the Phagocytic Activity of Human Macrophages via Regulation of Actin Cytoskeleton Proteins

6.1 Introduction

Macrophages are immune cells characterised by the ability to engulf and digest bacteria, damaged cells and any foreign substance that does not express on its surface normal host-cell specific proteins via a process known as phagocytosis (Duque and Descoteaux, 2015). Phagocytosis is a vital process for protecting the host against foreign dangers such as bacterial invasion (Desjardins *et al.*, 2005). This process needs to be performed efficiently because unsuccessful or imperfect clearance of bacteria can have detrimental circumstances on the host, and is linked with the progression of several pathological conditions such as chronic inflammation, wound infections and several autoimmune diseases (Engelich *et al.*, 2001). Attracted to sites of infection via chemotaxis, phagocytes such as macrophages recognize and attach to microorganisms. This results in actin polymerisation within the phagocyte, re-arrangement of the actin cytoskeleton to form pseudopodia followed by engulfment of the microorganism to form a phagosome and subsequent phagolysosome formation on fusion with a lysosome (Freeman and Grinstein, 2016; Garcia-Gomez *et al.*, 2016). Finally, the destruction of bacteria is mediated by lysosomal enzymes and ROS within the phagolysosome (Pauwels *et al.*, 2017). Successful engulfment of pathogens requires continuous reorganization of the phagocyte cell membrane that involves dynamic rearrangement of the actin cytoskeleton. The control of these mechanical changes on the cell membrane during cell movement and pathogen engulfment involves Rho GTPase proteins that regulate the polymerisation of actin filaments (Etienne-Manneville and Hall, 2002; Groves *et al.*, 2008).

In this study it has been shown that estrogen enhances the phagocytosis of both MRSA and *P. aeruginosa* by human macrophages. The interaction between macrophages and bacteria were visualised by SEM (Figure 3.16), demonstrating estrogen promotes the formation of pseudopodia by macrophages during the uptake of both MRSA and *P. aeruginosa*. These findings suggest estrogen may be regulating the levels or activation of one or more proteins involved in the regulation of the actin cytoskeleton during phagocytosis.

6.1.1 The Actin Cytoskeleton and Phagocytosis:

The process of engulfing a foreign particles, phagocytosis, is a crucial process for the wide diversity of organisms. From simple unicellular organisms that use phagocytosis for nutrition, to complex ones in which phagocytic cells play a fundamental role in the immune response. Phagocytosis is a step-wise process in which particles are first recognized and bind to the cell surface, followed by internalisation and formation of membrane-bound vacuoles known as phagosomes (Jilkine, 2009). After a target particle is recognised the phagocytic process requires a dynamic assemblage of the actin cytoskeleton to promote the formation of pseudopodia (Alberts *et al.*, 2008; Freeman and Grinstein, 2016; Garcia-Gomez *et al.*, 2016; Pollard *et al.*, 2016; Pauwels *et al.*, 2017).

The actin cytoskeleton is constructed by a wide range of proteins (Insall and Machesky, 2009). Actin is an abundant protein found in all cells and it exists in two different forms; globular monomers named G-actin, and polymeric chains of the two G-actin subunits, F-actin and filamentous actin (Holmes *et al.*, 1990). The F-actin filaments are structurally polarised (Wegner, 1976). The negative (known as “pointed”) end holds a subunit with an open ATP binding site, whereas the positive (known as “barbed”) end is open to adjacent G-actin monomers (Wegner, 1976). The polymerisation/depolymerisation of actin is a continuous constant process (Wegner, 1976) during which, the energy generated from ATP conversion to ADP stimulates the accumulation of G-actin monomers at the barbed end before polymerisation to F-actin filaments (Wegner, 1976; Alberts *et al.*, 2008) (Figure 6.1). After the complete establishment of F-actin filaments, actin-binding proteins add an extra layer of control over the actin cytoskeleton network. For instance, actin severing proteins, such as profilin and cofilin, can bind the actin polymers and alter the filaments length and strength (T. D. Pollard and Borisy, 2003), and increase/decrease the amount of barbed ends by cleaving actin filaments (Devreotes and Horwitz, 2015).

Capping proteins can also bind to the termination of the filaments regulating their attachment and detachment (Isenberg *et al.*, 1980). Additionally, F-actin filaments

bind to integrin and myosin II filaments producing focal adhesive complexes, which allow cells to slide past each other and connect with the extracellular matrix and other adjacent cells via the cytoskeleton network, therefore allowing cell-cell and cell-matrix exchange (Burrige *et al.*, 1988; Peskin *et al.*, 1993; Pollard and Borisy., 2003; Alberts *et al.*, 2008).

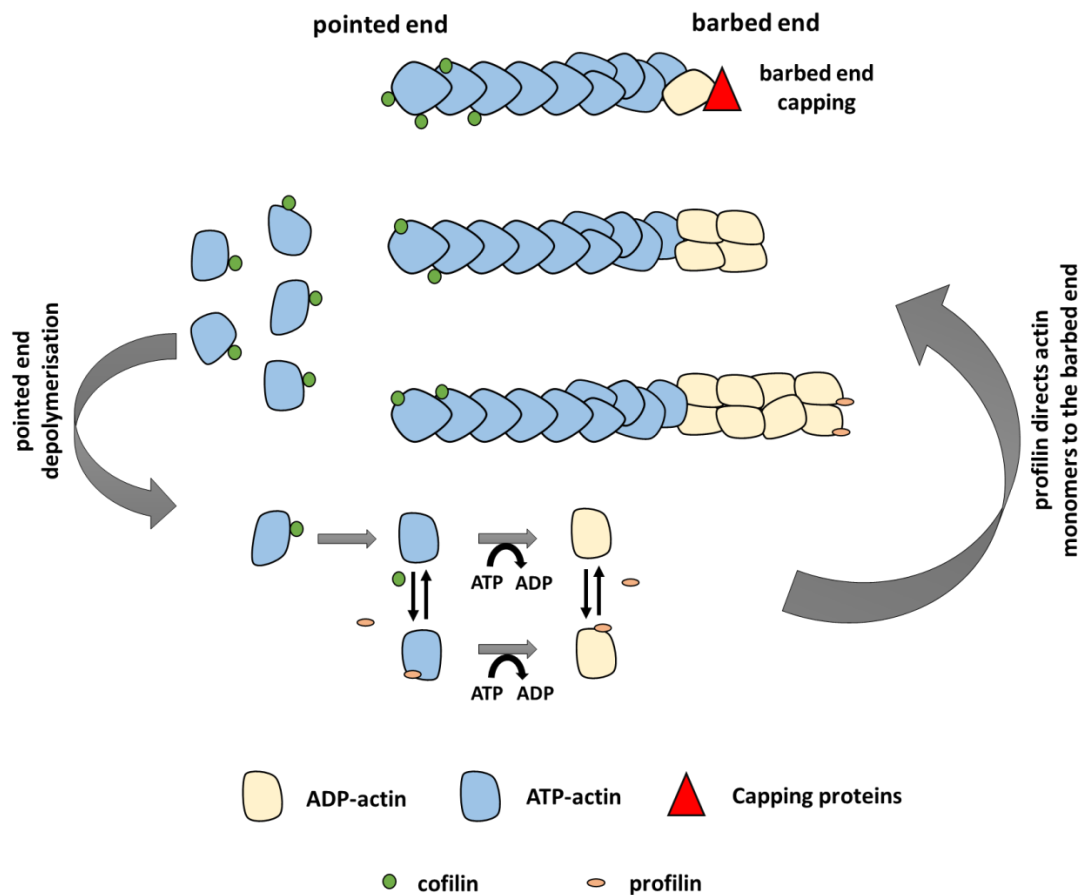


Figure 6.1. Actin cycle. F-actin filaments are separated due to Cofilin binding at the pointed end. Profilin binds to F-actin filaments at the barbed end before capping proteins are attached, allowing the attachment and detachment of filaments. Figure drawn based on details of the actin cycle described by Brown (2016).

6.1.2 Regulation of the Cytoskeleton Organisation by Rho GTPases

Estrogen was shown to promote the phagocytosis of the two major wound pathogens MRSA and *P. aeruginosa* (Chapters 3, 4 and 5). However, the link between estrogen and regulators of the actin cytoskeleton such as Rho GTPases during phagocytosis has not been investigated to date.

The organised movement of membranes and actin cytoskeleton are coordinated in phagocytosis by a family of proteins known as Rho GTPase (Etienne-Manneville and Hall, 2002). The small GTPases of the Rho-family (Rho-GTPases) are members of the Ras superfamily of small GTPases. The Rho-GTPase family includes 20 members distributed into 8 subfamilies (Figure 6.2) (Etienne-Manneville and Hall, 2002). While RhoA, Cdc42 and Rac1 are the most highly conserved and well-studied members in eukaryotic species, the 17 other members of the Rho GTPase family are less studied and their functions and effects have not been fully investigated and understood yet (Etienne-Manneville and Hall, 2002). Rac1, Cdc42 and RhoA are best known for their functions in regulating the actin cytoskeleton (Nobes and Hall, 1995; Machesky and Hall, 1996; A. Hall, 1998; Machesky and Insall, 1999; Etienne-Manneville and Hall, 2002). Working on 2D tissue-culture fibroblast models, (Ridley and Hall, 1992; Nobes and Hall, 1995) it has been reported that RhoA was mainly involved in rearranging actin and myosin filaments to form new stress fibres, however, Cdc42 and Rac1 induced actin polymerisation causing the formation of lamellipodia and filopodia. RhoA, Cdc42 and Rac1 have all emerged as master regulators of cell polarity and contractility, progression of the G1 cell cycle, gene transcription and microtubule dynamics (Etienne-Manneville and Hall, 2002). In addition to regulating the actin cytoskeleton dynamics, Rho GTPases have been linked with many biological functions such as cytoskeletal regulation, phagocytosis, cell polarity establishment, cell proliferation and motility, reactive oxygen species (ROS) production, and tumorigenesis (Etienne-Manneville and Hall, 2002; Jaffe and Hall, 2005).

Proteins of the Rho GTPase family are key regulators of the polymerisation of actin filaments resulting in cytoskeleton reorganisation (Etienne-Manneville and Hall, 2002). These proteins are parts of the GTP-binding proteins (G-proteins) family, which are conserved in eukaryotic organisms from yeast and amoeba to mammals (Etienne-Manneville and Hall, 2002). G-proteins contain a GTP-binding domain and function as molecular switches (Raftopoulou and Hall, 2004). In order to become active, the protein must bind to GTP causing its hydrolysis (Raftopoulou and Hall, 2004). The process of GTP hydrolysis is the origin of the name GTPase. As a

consequence of GTP hydrolysis to GDP, the protein changes structurally and converts into an inactive form (Raftopoulou and Hall, 2004). The Rho GTPase proteins receive signals from cell surface receptors, and interact indirectly with actin, via binding to effector molecules, which causes rearrangement of the cytoskeleton and enable phagocytosis (Raftopoulou and Hall, 2004).

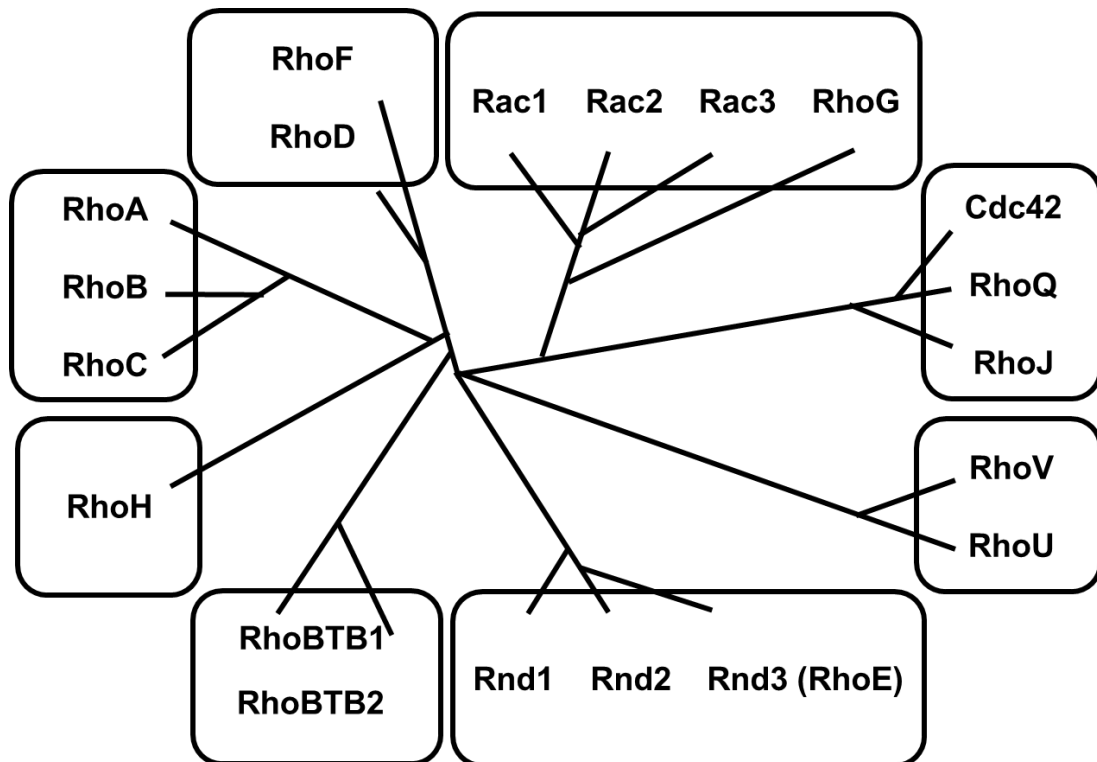


Figure 6.2. Phylogenetic tree representation of the Rho GTPases family. Rho GTPases family comprises 20 proteins divided into eight major groups according to sequence and phylogenetic similarities: Rho, Rac, Cdc42, RhoD/F, Rnd, RhoU/V, RhoH and RhoBTB. Figure based on published details of Rho GTPases (Aspenström *et al.*, 2007; Heasman and Ridley, 2008).

6.1.2.1 Cdc42

Cdc42 is an important Rho GTPase protein that controls signalling pathways regulating many cell functions essentially phagocytosis, cell cycle development and cell migration (Ridley *et al.*, 1992; Kozma *et al.*, 1995; Qadir *et al.*, 2015). Cdc42 is a key regulator of the cytoskeleton dynamics, which is essential for the phagocytosis process (Ridley *et al.*, 1992; Kozma *et al.*, 1995). In order to elucidate the functions of Cdc42 in humans, researchers studied the dominant negative (DN) and the constitutively active (CA) forms of Cdc42 (Van Aelst and D’Souza-Schorey, 1997), and demonstrated that cdc42 is potentially required in the initiation of cell signals

crucial for the regulation of actin cytoskeleton (Van Aelst and D'Souza-Schorey, 1997; Johnson, 1999). The Cdc42-related proteins stimulate the formation of filopodia, through binding to and activation of an effector protein known as "Wiskott Aldrich Syndrome Protein (WASP)" via Cdc42- and Rac-interacting and binding domain (CRIB) (Johnson, 1999; Rohatgi *et al.*, 1999; Carlier *et al.*, 2003). This leads to the recruitment of G-actin and a specific actin protein called the "Arp2/3 complex", which bind to the existing actin filaments resulting in a new filament (Rohatgi *et al.*, 1999; Carlier *et al.*, 2003).

Similar to WASP, other proteins such as the tyrosine p21-activated kinases (PAK1 and PAK2) and focal adhesive kinase (FAK), attach to Cdc42 via the CRIB domain and initiate the cytoskeleton reorganisation leading to membrane deformation, essential for phagocytosis (Hofmann *et al.*, 2004).

Cdc42 treadmills continuously and reversibly from an active GTP-bound form to an inactive GDP-bound form. This balance is initiated and controlled by guanine nucleotide exchange factors (GEFs) which encourage the conversion of GDP-bound molecules to unbound GTP (Van Aelst and D'Souza-Schorey, 1997). GTPase activating proteins (GAPs) promote GTP hydrolysis to GDP (Lamarche and Hall, 1994), whereas guanine nucleotide dissociation inhibitors (GDIs) block the detachment of the nucleotide from Cdc42 and result in deactivation of the protein (Olofsson, 1999; Valdés-Mora *et al.*, 2009).

6.1.2.2 Rac1

The GTPase Rac1 is one of the most documented and well-studied members of the Rho family (Bishop and HALL, 2000; Etienne-Manneville and Hall, 2002). Rac1 is a member of the Rac subfamily that comprises RhoG, Rac1b, Rac2, and Rac3 (Aspenström *et al.*, 2007; Heasman and Ridley, 2008). The Rac proteins all share more than 80% sequence homology and differ principally in the C-terminal region (Wennerberg, 2004). Rac1 is known for being a master regulator of the actin cytoskeleton organisation. Rac1 participates in the initial step of pseudopodia formation (Etienne-Manneville and Hall, 2002) during phagocytosis, but is also involved in many other cell functions including cell killing mediation via ROS, cell

differentiation, cell adhesion, apoptosis, signalling transcription factors and gene expression (Jaffe and Hall, 2005). The activity of Rac1 has been extensively studied in recent years, with findings showing active Rac1 induces the formation of membrane extensions called lamellipodium (Ridley *et al.*, 1992). In addition, RAC1 is essential in moderating several signalling pathways linked with cell growth and cell division (Bid *et al.*, 2013). Similar to Cdc42, the activities of Rac1 are mediated via interaction with specific effectors such as the protein kinase PAK that changes from an inactive to active state after interacting directly with GTP-Rac1 (Bid *et al.*, 2013).

6.1.2.3 RhoG

RhoG is a member of the Rac subgroup of the Rho GTPase family that was reported for the first time in 1992 (Vincent *et al.*, 1992). It shares 72% sequence identity with Rac1 but is characterised by its incapability to bind to proteins such as PAKs (Leung *et al.*, 1996; Gauthier-Rouviere *et al.*, 1998). Similar to most Rho GTPase proteins, RhoG is involved in a various cell signalling mechanisms including the regulation of actin cytoskeleton (Katoh *et al.*, 2006), phagocytosis (Prieto-Sánchez *et al.*, 2006), gene transcription (Vigorito *et al.*, 2003) and the regulation of NADPH oxidase in neutrophils (Condliffe *et al.*, 2006).

Both Rac1 and RhoG are able to signal downstream effectors after binding to GTP and become inactive when bound to GDP. Three protein families interact with Rac1 or RhoG to control the GTP/GDP balance: 1) GEFs are involved in facilitating the conversion of GDP for GTP, leading to the promotion of RhoA/RhoG-mediated signalling (Van Aelst and D'Souza-Schorey, 1997); 2) GAPs promote hydrolysis of GTP to GDP, thereby ending RhoA/RhoG-mediated signalling (Lamarche and Hall, 1994); 3) GDIs inhibit the GDP dissociation and lock the G protein in its inactive state (Olofsson, 1999).

It has been reported that the activation of RhoG provokes Rac1- and Cdc42-like activities such as lamellipodia and filopodia formation (Gauthier-Rouviere *et al.*, 1998). Activated Rac1 and/or RhoG are believed to be coupled with numerous downstream effectors, leading to the control of several signalling pathways. The mechanism by which RhoG signals is not fully understood. However, a detailed

signalling pathway of RhoG leading to activation of Rac proteins, via GEFs, has been described. This mechanism of RhoG-dependent Rac signalling was reported to encourage the phagocytosis of apoptotic cells in *C. elegans* (Kato and Negishi, 2003; Nakaya *et al.*, 2006).

6.1.2.4 RhoA

RhoA is a small protein member of the Rho GTPase family. Like the other GTPases, RhoA acts as a molecular switch to regulate signalling pathways that mediate the actin cytoskeleton dynamics and organisation (Hall, 1998). RhoA was initially reported to be a key player in the regulation of the cytoskeletal dynamics, and the formation of actin stress fibres in fibroblasts stimulated with growth factors (Ridley and Hall, 1992). It was later shown by Hackam and Grinstein in 1997 that RhoA is an essential requirement in phagocytosis. Olson *et al.* (1995) also demonstrated that RhoA regulates the growth of cells and triggers cell cycle progression via the phase G1.

The majority of downstream targets of RhoA are serine-threonine kinases, including ROCK1 which binds to RhoA in a GTP-dependant manner (Leung *et al.*, 1995). RhoA was also demonstrated to form actin stress fibres and focal adhesion complexes via binding to the ROCK kinase, an isoform of Rho kinase. Another Rho kinase known as Rho kinase-alpha or ROK was also shown to bind to RhoA in a GTP-dependent manner, leading to the assembly of actin filaments and the cytoskeleton organisation (Leung *et al.*, 1996).

The interaction between estrogen and Rac1 in cardiovascular disease has been documented, with estrogen downregulating the expression and activity of Rac1 in monocytes and vascular smooth muscle cells and resulting in a decrease in the production of ROS in the cardiovascular system (Laufs *et al.*, 2003). Estrogen enhances the phagocytic activity of rat peritoneal macrophages (Csaba, 2017) but inhibits monocyte recruitment and adhesion in the cardiovascular system (Friedrich *et al.*, 2006)

6.1.3 Focal Adhesion Kinase (FAK)

The focal adhesion kinase (FAK) was first discovered in the early 1990s (Kanner *et al.*, 1990; Schaller *et al.*, 1992) as a new tyrosine kinase that does not possess a receptor but is expressed in several tissues (Hanks *et al.*, 1992). FAK is highly conserved among different species with human FAK sharing high percentage homology with mice (97%) and zebrafish (90%) (Schaller, 2010). It has been reported that FAK plays a key role in integrin signalling which regulates phagocytosis, cell adhesion and cell survival (Schlaepfer *et al.*, 2004). The gene protein tyrosine kinase 2 (*PTK2*) encodes for FAK (André and Beckerandre, 1993). It has been demonstrated that blockage of FAK in breast cancer cells reduce their metastatic ability by decreasing cell mobility (Chan *et al.*, 2009).

FAK is expressed in all cell types except some blood cells, and is naturally localised in structures called focal adhesions. Focal adhesions are sites where the cytoplasmic cytoskeleton interact with the ECM (Chrzanowska-Wodnicka and Burridge, 1996). When FAK tyrosine kinase is activated, intracellular signalling pathways involved in cell migration are initiated (Hanks *et al.*, 1992; Schaller *et al.*, 1992). Estrogen is a known regulator of cytoskeletal constituents (Sapino *et al.*, 1986), mainly actin fibres via the control of actin dynamism and organisation (Giretti *et al.*, 2008; Flamini *et al.*, 2009). Estrogen also regulates cell morphology and cell interaction with the ECM, a key process in phagocytosis (Giretti *et al.*, 2008; Flamini *et al.*, 2009; Flamini *et al.*, 2011). Flamini *et al.* (2011) investigated the effect of estrogen on the remodelling of actin cytoskeleton in the Ishikawa endometrial adenocarcinoma cell line and endometrial stromal cells (ESC). This study showed remodelling estrogen promotes FAK phosphorylation in Ishikawa and ESC cells in a dose-dependent manner, resulting in an enhancement of actin and membrane remodelling, and phagocytosis.

6.2 Aim and Objectives

6.2.1 Aim

To investigate the effect of estrogen on regulating protein levels of Rac1, RhoA, Cdc42, RhoG and FAK in U937-derived M1 macrophages.

6.2.2 Objectives

To compare protein levels of Rac1, RhoA, Cdc42, RhoG and FAK in U937-derived M1 macrophages treated with/without supraphysiological levels of estrogen for 24 hours against BSA-treated negative control.

6.3 Methods

6.3.1 Generation of U937-derived M1 Macrophages

U937 monocytes were differentiated into M0 macrophages and polarised into M1-like macrophages using methods described in Section 2.2.5.

6.3.2 Immunoblotting

U937-derived M1 macrophages were treated with 1×10^{-7} M BSA, zero M estrogen (17β -estradiol) or 1×10^{-7} M estrogen for 24 hours. Macrophages were retrieved and lysed prior to extracting and quantifying proteins following methods described in Section 2.2.13.1. Immunoblotting assays were conducted as described in Section 2.2.13.2 to quantify levels of Rac1, RhoA, Cdc42, RhoG and FAK with protein normalisation conducted against the housekeeping gene β -tubulin.

6.4 Results

6.4.1 Estrogen Enhances Rac1 by U937-Derived M1 Macrophages

The effect of estrogen on levels of the GTPase Rac1 by U937-derived M1 macrophages was investigated (Figure 6.3). Estrogen had no significant effect ($P>0.05$) on levels of β -tubulin in U937-derived M1 macrophages compared to untreated macrophages (zero M estrogen) and the negative control (Figure 6.3.A and C). However, estrogen supplementation (Figure 6.3. B and C) significantly ($P<0.01$) increased relative Rac1 levels (1.9-fold change) compared to the negative control (NC). There was no significant difference ($P>0.05$) between Rac1 levels in the NC and untreated macrophages that lacked estrogen supplementation.

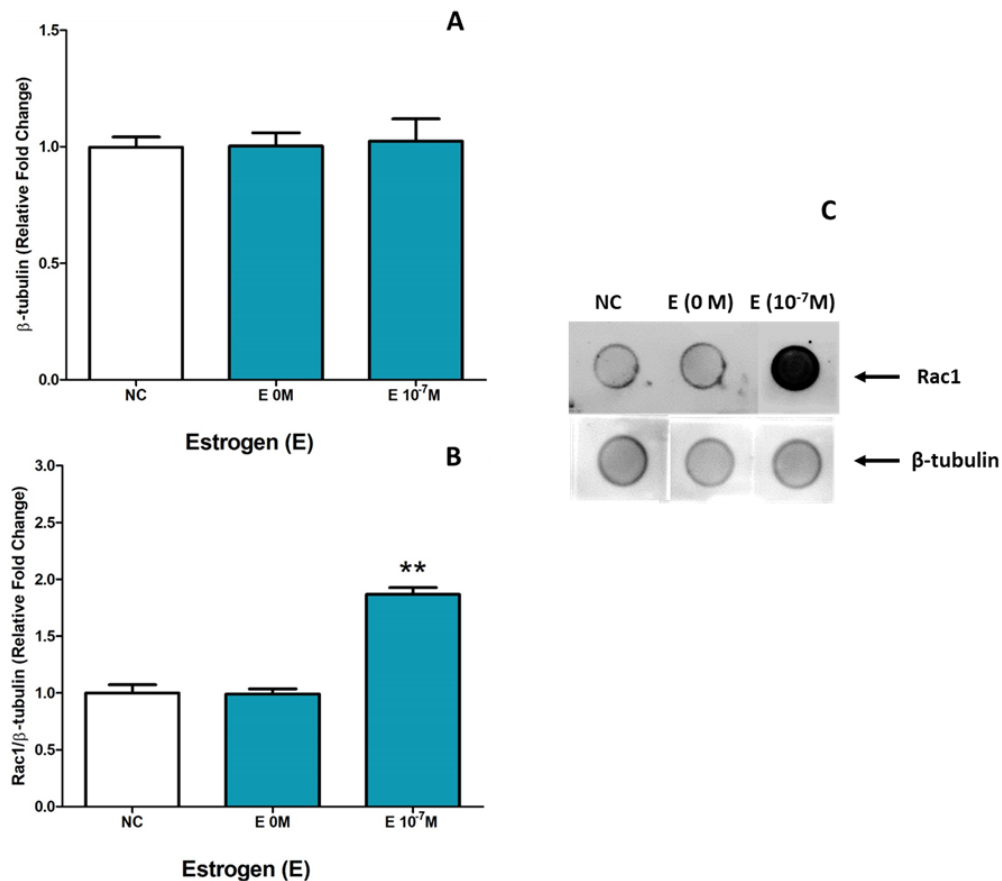


Figure 6.3. Rac1 levels in U937-derived M1 macrophages increase following exposure to supraphysiological estrogen. Rac1 levels were assessed by immunoblotting with β -tubulin used as loading control. Estrogen (E) had no effect on the β -tubulin levels in macrophages ($P>0.05$) (A and C), but significantly ($P<0.01$) increased Rac1 levels compared to the negative control (NC) and untreated (E 0M) macrophages (B and C). Results represent an average of $n = 12$ experimental repeats. ** Indicates significant difference in protein levels

(fold-change) compared to the NC ($P < 0.01$). Error bars represent the standard error of the mean (StEM).

6.4.2 Estrogen Decreases RhoA in U937-Derived M1 Macrophages

Estrogen had no significant ($P > 0.05$) effect on the housekeeping protein β -tubulin (Figure 6.4.A and C) but significantly ($P < 0.01$) reduced relative levels of RhoA (0.7-fold change) in U937-derived M1 macrophages compared to the negative control (NC). There was no significant ($P > 0.05$) difference in RhoA levels between the NC and untreated macrophages lacking estrogen supplementation (Figure 6.4.B and C).

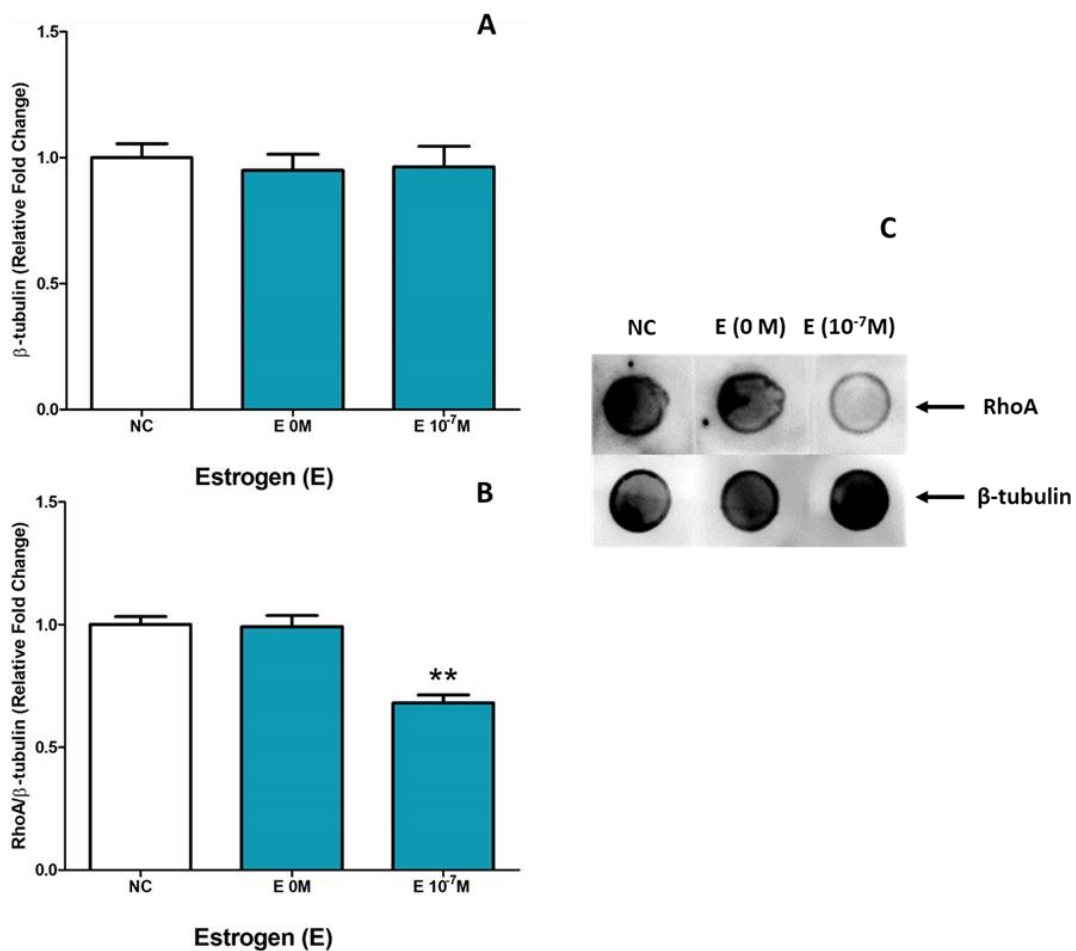


Figure 6.4. RhoA levels in U937-derived M1 macrophages reduce following treatment with supraphysiological estrogen. RhoA levels were measured by immunoblotting using β -tubulin as the housekeeping protein. Estrogen (E) had no significant ($P > 0.05$) effect on levels of β -tubulin (A and C) but significantly ($P < 0.01$) decreased RhoA levels compared to the negative control (NC) and untreated (E 0M) macrophages (B and C). Results represent an average of $n = 12$ experimental repeats. ** Indicates significant difference ($P < 0.01$) in protein levels (fold-change) compared to the NC. Error bars represent the standard error of the mean (StEM).

6.4.3 Estrogen Increases Levels of Cdc42 in U937-Derived M1 Macrophages

Estrogen had no significant ($P>0.05$) effect on levels of the housekeeping protein β -tubulin (Figure 6.5.A and B). In comparison to the negative control (NC), the absence of estrogen in untreated macrophages had no significant ($P>0.05$) effect on Cdc42 levels (1.0-fold change). However, estrogen induced a significant ($P< 0.01$) increase in levels of Cdc42 (2.3-fold change) compared to the NC (Figure 6.5.B and C).

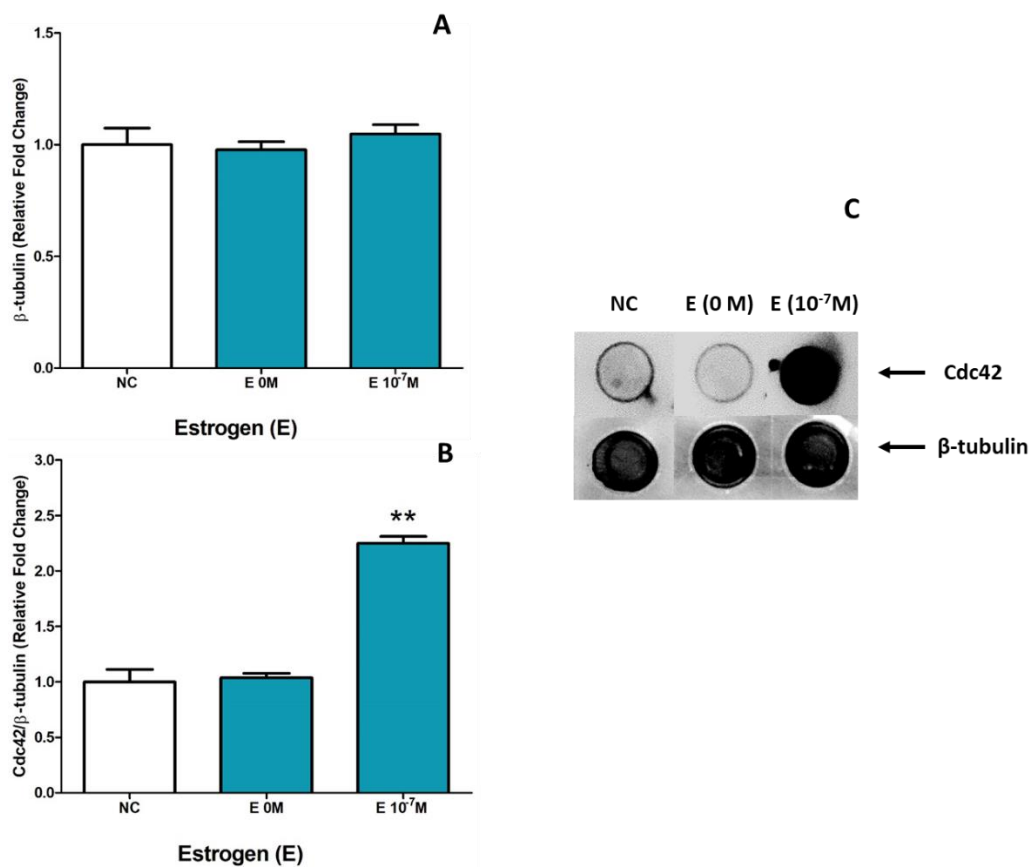


Figure 6.5. Cdc42 levels in U937-derived M1 macrophages increase following exposure to supraphysiological estrogen. Cdc42 levels were assessed by immunoblotting with β -tubulin used as loading control. Estrogen (E) had no effect on the β -tubulin levels in macrophages ($P>0.05$) (A and C), but significantly ($P<0.01$) increased Cdc42 levels compared to the negative control (NC) and untreated (E 0M) macrophages (B and C). Results represent an average of $n = 12$ experimental repeats. ** Indicates significant difference in protein levels (fold-change) compared to the NC ($P<0.01$). Error bars represent the standard error of the mean (StEM).

6.4.4 Estrogen Increases Levels of RhoG in U937-Derived M1 Macrophages

The effect of estrogen on levels of RhoG in U937-derived M1 macrophages was investigated (Figure 6.6). Estrogen had no significant effect on levels of β -tubulin in

U937-derived M1 macrophages, compared to negative control (NC) and untreated macrophages (E 0M) (Figure 6.6.A and C). Compared to the NC, the level of RhoG was not significantly ($P>0.05$) altered in untreated (E 0M) macrophages (1.1-fold change). However, estrogen significantly ($P< 0.01$) increased relative levels of RhoG (2.3-fold change) in comparison to the NC (Figure 6.6.B and C).

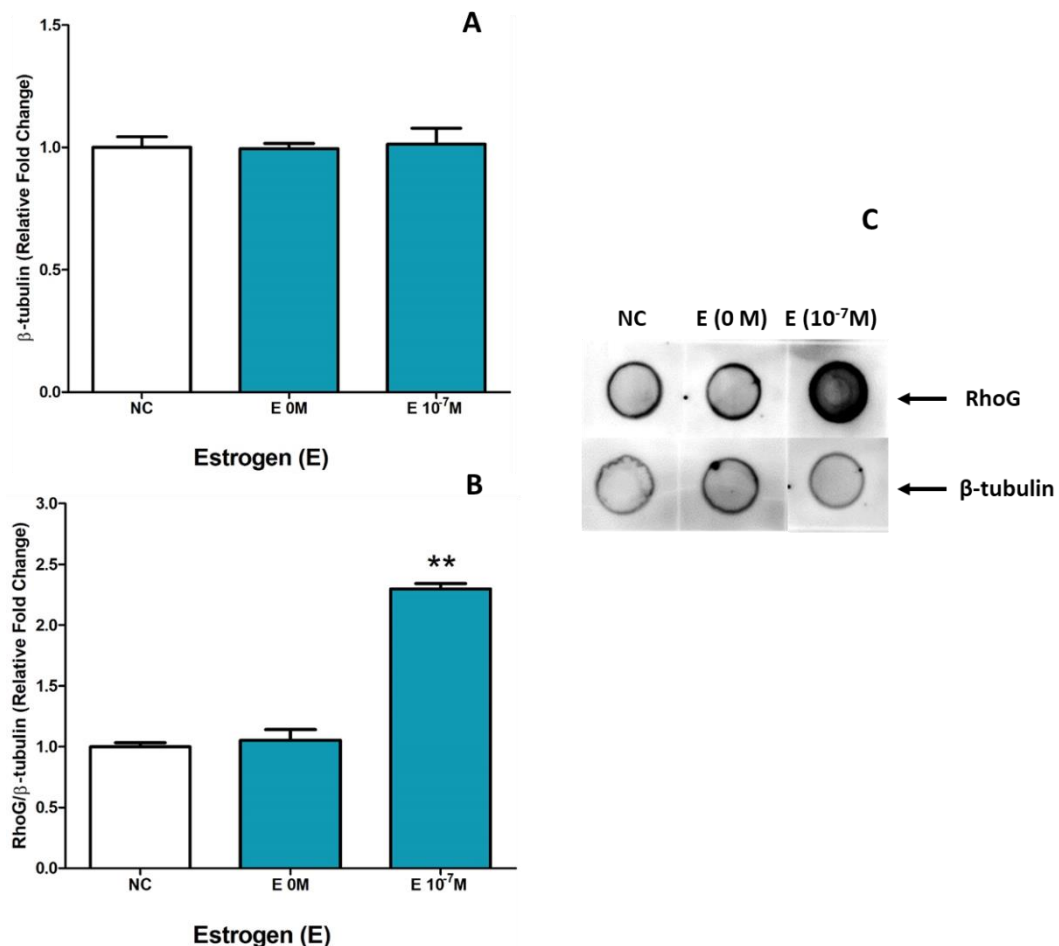


Figure 6.6. RhoG levels in U937-derived M1 macrophages increase following treatment with supraphysiological estrogen. RhoG levels were assessed by immunoblotting with β -tubulin used as loading control. Estrogen (E) had no effect on the β -tubulin levels in macrophages ($P>0.05$) (A and C), but significantly ($P<0.01$) increased RhoG levels compared to the negative control (NC) and untreated (E 0M) macrophages (B and C). Results represent an average of $n = 12$ experimental repeats. ** Indicates significant difference in protein levels (fold-change) compared to the NC ($P<0.01$). Error bars represent the standard error of the mean (StEM).

6.4.5 Estrogen Increases Levels of Focal Adhesion Kinase (FAK) in U937-Derived M1 Macrophages

Levels of the housekeeping protein β -tubulin did not significantly change ($P>0.05$) following treatment of U937-derived M1 macrophages with supraphysiological

estrogen (Figure 6.7.A and C). Levels of FAK in untreated macrophages (1.0-fold change) were not significantly ($P>0.05$) different to levels found in the negative control (NC). Intriguingly, relative levels of FAK significantly increased ($P<0.01$) following treatment with estrogen (2.3-fold change) compared to the NC (Figure 6.6.B and C).

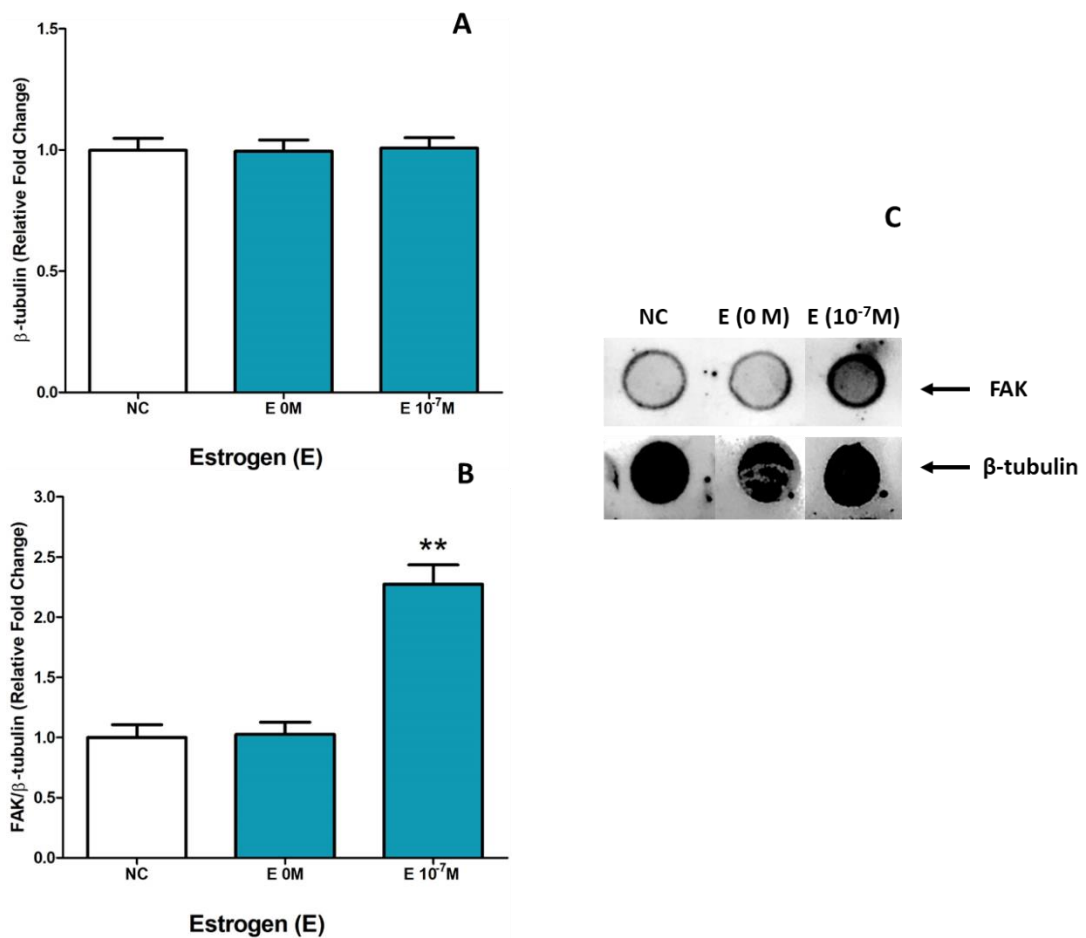


Figure 6.7. FAK levels in U937-derived M1 macrophages increase following treatment with supraphysiological estrogen. FAK levels were assessed by immunoblotting with β -tubulin used as loading control. Estrogen (E) had no effect on the β -tubulin levels in macrophages ($P>0.05$) (A and C), but significantly ($P<0.01$) increased FAK levels compared to the negative control (NC) and untreated (E 0M) macrophages (B and C). Results represent an average of $n = 12$ experimental repeats. ** Indicates significant difference in protein levels (fold-change) compared to the NC ($P<0.01$). Error bars represent the standard error of the mean (StEM).

6.5 Discussion

Estrogen promotes the phagocytosis of both MRSA and *P. aeruginosa* by human macrophages through specific binding to ER- α . (Chapter 5). Moreover, data from the SEM (Chapter 3) demonstrated major changes in the morphology of phagocytic macrophages following estrogen supplementation, with the formation of pseudopodia suggesting estrogen induces substantial changes in the actin cytoskeleton. In order to determine whether Estrogen stimulates the phagocytic activity of macrophages through changes in the actin cytoskeleton, this study measured the protein levels of four major Rho GTPase proteins (Cdc42, Rac1, RhoG and RhoA) and the focal adhesion kinase (FAK) in U937-derived M1 macrophages in the presence/absence of estrogen for 24 hours.

Estrogen significantly increased the level of FAK in M1 macrophages. FAK has been shown to be an important regulator of phagocytosis (Tzircotis *et al.*, 2011). Tzircotis *et al.* (2011) transfected macrophages with RNAi against FAK and found that macrophages lacking FAK presented a 65% decrease in the number of phagocytes produced with Fc γ R- and CR3-mediated phagocytosis. Activation of FAK was also associated with provoking Rac1- and Cdc42-like activities, such as lamellipodia and filopodia formation (Gauthier-Rouviere *et al.*, 1998), but not RhoA activities (Tzircotis *et al.*, 2011). It has been reported that fibroblasts lacking FAK have impaired cell motility. Importantly, FAK is an important regulator of the pathogen-killing functions of neutrophils (Kasorn *et al.*, 2009). FAK signalling is linked with the organisation of integrin-based adhesion sites and the formation of lamellipodia in migrating cells, such as neutrophils and macrophages (Kasorn *et al.*, 2009). Moreover, FAK deficiency in macrophages reduces cell adhesion, and is associated with failure to form lamellipodia and cell migration to sites of infection and inflammation *in vivo* (Owen *et al.*, 2007).

Estrogen is a known regulator of cytoskeletal constituents, mainly actin fibres (Giretti *et al.*, 2008; Flamini *et al.*, 2009) via the control of actin dynamism and organisation. Estrogen also regulates cell morphology and interaction with the ECM, a key process in phagocytosis (Giretti *et al.*, 2008; Flamini *et al.*, 2009). Flamini *et al.*

(2011) investigated the effect of estrogen on the remodelling of actin cytoskeleton and on regulating cell mobility in an Ishikawa cell line and ESC. They showed estrogen encourages FAK phosphorylation in Ishikawa and ESC cells in a dose-dependent manner, resulting in an enhancement of cell mobility, actin and membrane remodelling (Flamini *et al.*, 2011).

Importantly, estrogen has been shown to stimulate the production of membrane ruffles and pseudopodia in breast cancer cells via activation of ER- α in concordance with the findings (Chapter 3) of this study (Song *et al.*, 2002). Simoncini *et al.* (2006) confirmed that estrogen encourages the migration of endothelial cells via phosphorylation and activation of FAK. Song and Santen (2006) indicated that estrogen enhances the formation of filopodia and lamellipodia with localisation of ER- α in the plasma membrane. Stimulation of cancer cells with estrogen induces cell migration and motility via interaction of FAK with ER- α and other factors including phosphatidylinositol 3 kinase and p60 Src tyrosine kinase (c-Src) (Li *et al.*, 2010). Given estrogen stimulated phagocytosis via ER- α (Chapter 3) and increased levels of FAK further investigations in this area are now warranted to confirm a causal link between elevated FAK levels and enhanced phagocytosis in estrogen supplemented M1 macrophages.

In this study, macrophage levels of Rac1 significantly increased following treatment with estrogen for 24 hours compared to the negative control or macrophages lacking estrogen supplementation. Rac1 is crucial for dynamic rearrangement of the actin cytoskeleton that is essential for membrane ruffling and the development of focal complexes that are needed for filopodia and lamellipodia formation (Ridley *et al.*, 1992; Allen *et al.*, 1997). Rac1 regulates the organisation of actin cytoskeleton and cell adhesion in murine macrophages *in vitro* (Allen *et al.*, 1997). Upregulation of active Rac1 have been detected in phagocytic cups (Hoppe and Swanson, 2004) while its inhibition was found to reduce the phagocytosis of both C3bi- and IgG-opsonised molecules (Cox *et al.*, 1997). It is noteworthy that estrogen-like molecules such as daidzein have been shown to have an effect on the regulation of Rac1 during the phagocytic process. Daidzein, a soy isoflavone extract, was found to stimulate the production of estrogen in trophoblast cells (Richter *et al.*, 2009).

Moreover, diadzein enhances the macrophage phagocytic capability via upregulation of the Rac1 activity (Yen *et al.*, 2014).

In another report, it has been shown that the estrogen metabolite (2-ME) blocks the RhoA/ROCK1 pathway in human smooth muscle cells through downregulation of RhoA (Rigassi *et al.*, 2015). It has also been reported that the upregulation of RhoA inhibits the process of phagocytosis in macrophages (Tosello-Tramont *et al.*, 2003; Colucci-Guyon *et al.*, 2005; Nakaya *et al.*, 2006). Results in this study have demonstrated that treatment of macrophages with estrogen for 24 hours significantly decreased protein levels of RhoA compared to the negative (BSA) negative control and macrophages cultured in the absence of estrogen. Whether estrogen stimulates phagocytosis by dampening the RhoA/ROCK1 pathway warrants further investigation but accumulating evidence has indicated that upregulation of RhoA inhibits the process of phagocytosis in macrophages (Tosello-Tramont *et al.*, 2003; Colucci-Guyon *et al.*, 2005; Nakaya *et al.*, 2006).

It is worth pointing out that many other reports support the findings of this study and show a harmonised balance of Rac1 and RhoA is essential for a successful phagocytosis. Research investigations on *Candida elegans* in mammals have reported that Rac1 and RhoA, are antagonistically implicated in the modulation of cytoskeleton reorganisation during the process of phagocytosis, as their signalling oppose each other (Ohta *et al.*, 2006; Freeman and Grinstein, 2014). To date, their exact functions are not fully clear, but recent studies using mutants of Rac1 and RhoA specified that upregulation of Rac1 stimulates phagocytes to engulf apoptotic cells (Akakura *et al.*, 2004; Nakaya *et al.*, 2008) whereas RhoA had an opposite effect (Kim *et al.*, 2017). Kim *et al.* (2017) investigated the role of Rac1 and RhoA in the clearance of apoptotic cells and showed a dynamic balance between the two proteins was needed to regulate efferocytosis, with Rac1 upregulated and RhoA downregulated during pseudopodial formation.

RhoG is associated with the regulation of actin cytoskeleton dynamics (Kato *et al.*, 2006), phagocytosis (Prieto-Sánchez *et al.*, 2006), gene transcription (Vigorito *et al.*, 2003) and the regulation of NADPH oxidase in neutrophils (Condliffe *et al.*, 2006). Data in this chapter demonstrates for the first time that treatment of macrophages

with estrogen causes an increase in RhoG levels leading to enhanced phagocytosis of bacteria. This result supports the findings of Tzircotis *et al.* (2011) that demonstrated that RhoG stimulates phagocytosis of IgG-opsonised (for Fc γ R) or C3bi-opsonised (for CR3) sheep red blood cells (RBCs) in mouse macrophages. However, the findings in this chapter are the first to show that estrogen stimulates RhoG in human M1 macrophages. Therefore, elevation of RhoG levels could potentially be a novel and effective therapeutic approach for the treatment of wound. However, further investigations are needed to provide a confirmation of this hypothesis.

In this study, levels of Cdc42 in M1 macrophages following treatment with estrogen significantly increased. Resveratrol, a plant-derived polyphenol found in grapes and associated products such as wine, has both anti-estrogenic effects in breast cancer and but is believed to be estrogenic in peripheral tissues such as bone (Bhat *et al.*, 2001; Tou, 2015). Azios *et al.* (2007) investigated the effect of resveratrol in estrogen-responsive breast cancer, and found that resveratrol decreased the activity of Cdc42 and Rac1 whereas a combination of resveratrol with estrogen resulted in a significant upregulation of both Cdc42 and Rac1 in breast cancer cells. Cdc42 is linked with the organisation of the actin cytoskeleton, membrane ruffling and formation of filopodia in fibroblasts and macrophages (Kozma *et al.*, 1995; Cox *et al.*, 1997). Research indicates Cdc42 is crucial for successful phagocytosis of *Salmonella* in COS-1 cells (L.-M. Chen *et al.*, 1996; Cox *et al.*, 1997). In addition, the injection of macrophages with activated Cdc42 results in the formation of long and indistinct filopodia *in vitro* (Allen *et al.*, 1997). Cdc42 is an essential requirement for Fc receptor-mediated phagocytosis and for the formation of membrane ruffling mediated by diverse receptors in macrophages (Allen *et al.*, 1997).

In summary, estrogen increased levels of Rac1, Cdc42, RhoG and FAK, but reduced levels of RhoA in U937-derived M1 macrophages. Further interrogation of pathways involved in the actin cytoskeleton could potentially reveal other mediators of estrogen-induced phagocytosis. Whilst results presented in this chapter provided exciting and novel links between estrogen and proteins of the cytoskeleton during the process of phagocytosis, further investigations are warranted to determine the

effect of estrogen on relative amounts of proteins in their activated (phosphorylated) state during the process of phagocytosis. Western blotting, kinase activity assays (to measure FAK activation), and intracellular flow cytometry would be very robust methods of measuring activated protein levels. Inhibitors of phagocytosis pathways could also be exploited to further understand the mechanisms by which estrogen induces phagocytosis. In addition, further work is required to determine whether the changes in protein levels are mirrored by concomitant alterations in gene expression. In order to determine the effect of ER- α activation on downstream gene expression in macrophages, RNA sequencing (RNAseq) and real time polymerase chain reaction (qPCR) analysis could be adopted. Understanding the precise mechanisms by which estrogen promotes phagocytosis might reveal key downstream targets to interrogate further and may ultimately lead to the development of novel therapies to treat wound infections in the elderly with high efficacy and minimal side effects.

Chapter 7: Discussion and Future Work

7.1 Discussion

Chronic wounds are difficult to treat and often become colonised with bacteria leading to substantial mortality and morbidity in the elderly. Impaired wounds can become colonised with a varied range of pathogens including *S. aureus* and *P. aeruginosa* (Gjødsbøl *et al.*, 2006; Kirketerp-Møller *et al.*, 2008; Rybtke *et al.*, 2015). Treatment of bacterial infections in wounds requires aggressive medication, and the use of last-generation antibiotics (Beasley and Hirst, 2004). Thus, the treatment of wound infections in the elderly represents a big challenge for the healthcare services (Beasley and Hirst, 2004). The propensity for wound infections increases in the elderly, in part due to the delay in wound repair (Cooper *et al.*, 2015). Estrogen is a key regulator of the wound healing process and estrogen deprivation in the elderly is linked with pathological wound healing (Ashcroft *et al.*, 1999a; Margolis *et al.*, 2002). Indeed, impaired healing in the elderly is mediated by changes in steroid hormones, particularly declining levels of estrogen with increasing age (Ashcroft *et al.*, 1999a; Ashcroft *et al.*, 2003; M. J. Hardman and G. S. Ashcroft, 2008). Estrogen deficiency causes a significant delay in wound repair, with topical and systemic estrogen replacement reversing this delay (Ashcroft *et al.*, 1997a; Ashcroft *et al.*, 1999a). Furthermore, estrogen supplementation promotes age-related wound healing in both elderly males and females (Ashcroft *et al.*, 1999a; Guo and DiPietro, 2010). Delayed wound healing in the elderly is associated with a delayed, but prolonged and excessive inflammatory response, with dysregulated expression of pro-inflammatory cytokines and proteolytic mediators. The key role of estrogen in regulating inflammation has been extensively reviewed (Gilliver *et al.*, 2007; Guo and DiPietro, 2010), with estrogen supplementation decreasing the inflammatory response in elderly subjects of both genders (Ashcroft *et al.*, 1999a). However, despite the pronounced inflammatory response, evidence suggests declining levels of estrogen with increasing age can result in an impaired ability of inflammatory cells to eliminate bacteria from infected wounds (Emori *et al.*, 1991; Thomas, 2001). The possible role estrogen plays in macrophage-mediated bacterial clearance has not been fully characterised to date. In order to elucidate the effect of age-related changes in estrogen levels on the clearance of bacteria by human macrophages,

this study conducted host-pathogen interaction assays to investigate the phagocytosis of two major wound pathogens, MRSA and *P. aeruginosa* by *in vitro* U937-derived macrophages and *ex vivo* human peripheral blood monocyte (HPBM)-derived macrophages. Investigations were performed under conditions that model estrogen levels found in the elderly (estrogen deprivation), young adults (referred to as physiological estrogen in this thesis) and following exogenous estrogen supplementation.

The conversion of monocytes into macrophage-like cells *in vitro* was an essential requirement to conduct all the host-pathogen assays in this study. PMA is reported to induce the differentiation of monocytes into macrophages *in vitro* (Martinez *et al.*, 2006; Rios de la Rosa *et al.*, 2017) via upregulation of cell adhesion molecules such as beta-2-integrins (CD11a, CD11c, CD18 and CD11b) (Luscinskas *et al.*, 1994), enabling cells to adhere to cell culture flasks *in vitro*. The differentiation of monocytes into macrophages was assessed in this study via the detection of the CD11c surface marker by flow cytometry. The high expression of CD11c in PMA-differentiated macrophages, compared with monocytes incubated in the absence of PMA, confirmed the successful transformation of U937 monocytes and HPBMs into macrophage-like cells *in vitro*.

Macrophage stimulation with supraphysiological levels of estrogen increases the phagocytic function of human brain macrophages (microglial cells) to take up amyloid β -peptide and *E. coli* (Li *et al.*, 2000). Evidence has indicated that estrogen increases bacterial clearance in an *in vitro* LPS model of *K. pneumoniae* infection in acute wounds Crompton *et al.* (2016). In concordance with these reports, novel findings presented in this thesis (Chapter 3) have demonstrated that physiological (typical of youth) and supraphysiological levels of estrogen promote the engulfment of both Gram-positive and Gram-negative live bacteria by both human U937-derived macrophages and HPBM-derived macrophages. Further, *in vitro* host-pathogen assays involving the uptake of fluorescent bacteria (GFP-*S. aureus*) by U937-derived macrophages validated these findings, with significantly increased intracellular fluorescence following treatment of phagocytes with estrogen. Due to the poor ability of the antibiotic gentamicin to permeate the macrophage

membrane, a gentamicin protection assay was performed to determine the number of viable bacteria inside the phagocytes following the killing of extracellular bacteria using gentamicin (Hamad *et al.*, 2010; Rios de la Rosa *et al.*, 2017). Novel findings from this assay indicated that physiological (typical of youth) and supraphysiological levels of estrogen promote not just the internalisation, but also the killing, of both MRSA and *P. aeruginosa* in a dose-dependent manner. Further evidence captured by fluorescent microscopy (epifluorescent and confocal microscopes) indicated a higher uptake of both GFP-*S. aureus* and mCherry-*P. aeruginosa* by estrogen-treated macrophages in comparison to estrogen-deprived macrophages, with Z-stack analysis confirming co-localisation of the bacteria within macrophages. Images captured by SEM provided additional evidence of enhanced host-pathogen interactions following treatment of macrophages with estrogen, with macrophages showing pathogen-specific morphological changes in response to MRSA and *P. aeruginosa*. Macrophages generated longer membrane extensions (pseudopodia) in response to *P. aeruginosa* compared to MRSA. This pathogen-specific difference is intriguing since the clearance of MRSA was consistently higher than the uptake of *P. aeruginosa* ($P < 0.05$) regardless of the experimental conditions. The lower bacterial uptake and longer pseudopodia produced by macrophages following interaction with *P. aeruginosa*, may be due to *P. aeruginosa*'s being a motile bacterium (O'Toole and Kolter, 1998) and therefore phagocytes may have reduced success when stretching out further to capture a moving target due to increased demands on cytoskeleton reorganisation.

Known for its anti-inflammatory properties, estrogen reduces excessive and prolonged inflammation by directly affecting the inflammatory response of monocytes, macrophages and neutrophils (Ashcroft *et al.*, 2003; Guo and DiPietro, 2010). Estrogen dampens the inflammatory response via inhibition of pro-inflammatory cytokines production by tissue macrophages (Ashcroft and Ashworth., 2003). However, estrogen also stimulates tissue regeneration and remodelling following the resolution of inflammation (Ashcroft *et al.*, 1999b; Ashcroft and Ashworth, 2003; Ashcroft *et al.*, 2003). Collectively, these reports suggest that estrogen modulates both pro-inflammatory (M1) and pro-regenerative

(M2) macrophages during wound repair. The effect of estrogen on the phagocytosis of MRSA and *P. aeruginosa* by U937-derived and HPBM-derived M1/M2 macrophages was investigated in Chapter 4. High expression of CD197 confirmed the polarisation of U937-derived and HPBM-derived M0 macrophages into M1-like macrophages by IFN- γ /LPS, whilst high CD206 expression confirmed polarisation into M2-like macrophages by IL-4/IL-13 (Stossi *et al.*, 2012; Rios de la Rosa *et al.*, 2017). Consistent with the data in Chapter 3, physiological (typical of youth) and supraphysiological levels of estrogen were found to promote the phagocytosis of MRSA and *P. aeruginosa* in a dose-dependent manner by both M1 and M2 macrophages compared to estrogen deprivation and the absolute absence of estrogen. However, the phagocytosis of bacteria by M1 macrophages was significantly ($P < 0.05$) higher than that with M2 or M0 macrophages, regardless of whether the macrophages were derived from U937s or HPBMs. This data demonstrated the enhanced phagocytosis of bacteria by M1 macrophages which is in agreement with published findings (Mantovani *et al.*, 2004; Benoit *et al.*, 2008; Murray and Wynn, 2011; Krzyszczyk *et al.*, 2018).

Further work assessing potential mechanisms of estrogen-enhanced phagocytosis was subsequently conducted using pro-inflammatory M1-like macrophages instead of M0-like or M2-like macrophages. In order to elucidate the importance of the ERs in mediating the estrogen-enhanced phagocytosis of bacteria by human M1 macrophages, host-pathogen interaction assays were conducted using the SERM tamoxifen and the SERD fulvestrant. Tamoxifen and fulvestrant are mainly used to treat breast cancer (Cole *et al.*, 1971; Cummings *et al.*, 1999) owing to their antiestrogenic effects in breast tissue (Love *et al.*, 1992; Black *et al.*, 1994). Tamoxifen acts as an ER antagonist in the breast, but is agonist in peripheral tissues such as the bone and the uterus (Love *et al.*, 1992; Black *et al.*, 1994). Interestingly, stimulation of macrophages with tamoxifen mirrored the effects of estrogen by significantly ($P < 0.01$) increasing the phagocytosis of both MRSA and *P. aeruginosa* by U937-derived and HPBM-derived M1 macrophages. These results are supported by the findings of Corriden *et al.* (2015) which showed tamoxifen improves the phagocytic function of human neutrophils against a range of bacteria, including

MRSA. Hardman *et al.* (2007) also reported that tamoxifen accelerates wound healing in ovariectomised mice by exhibiting estrogenic effects, principally by reducing the inflammatory response. Collectively, these findings suggest that tamoxifen could potentially be a beneficial therapeutic strategy to treat infected wounds in the elderly.

Degradation of the ER by fulvestrant blocked the beneficial effects of estrogen on the phagocytosis of MRSA and *P. aeruginosa* by U937-derived and HPBM-derived M1 macrophages. Compared with estrogen, fulvestrant has a higher binding affinity to the ER (Morris and Wakeling, 2002), and when fulvestrant is coupled with the ER, the nuclear localisation of the ER is blocked leading to its degradation (Morris and Wakeling, 2002; Osborne *et al.*, 2004).

Together, these data from the use of tamoxifen and fulvestrant provide evidence that estrogen is promoting phagocytosis via the binding and activation of the ER. Consequently, the role of the two estrogen receptor subtypes, ER- α and ER- β , in this process was investigated using ER isoform-specific agonists and antagonists. In a similar manner to estrogen and tamoxifen, selective activation of ER- α with PPT increased the phagocytosis of both MRSA and *P. aeruginosa* by M1 macrophages. However, treatment with the ER- β agonist DPN failed to promote the phagocytosis of either MRSA or *P. aeruginosa*. Intriguingly, blockade of ER- α with TPBM reversed the effect of PPT on the bacterial clearance. These novel findings collectively indicate that estrogen induces the phagocytosis of bacteria through activation of ER- α . This supports published findings showing that ER- α regulates inflammation, with ER- β modulating the later phases of wound healing (Campbell *et al.*, 2010). Toutain *et al.* (2009) identified ER- α as the main isoform involved in promoting the beneficial effects of 17 β -estradiol in a model of skin necrosis. ER- α has also been shown to be the predominant ER involved in regulating the thickness of epidermis during wound healing (Stumpf *et al.*, 1974; Bidmon *et al.*, 1990). In summary, the findings described in Chapter 5 provide an indication that ER- α is the ER isoform through which estrogen promotes phagocytosis by macrophages. Selective targeting of ER- α with agents such as PPT may potentially lead to the development of therapeutic strategies to combat wound infections in the elderly.

Images from the SEM (Figure 3.16) illustrated major morphological changes in the membrane of phagocytes (macrophages) following estrogen treatment, with the establishment of multiple pseudopodial structures suggesting estrogen induces significant alterations in the actin cytoskeleton of the phagocytes. These findings suggested that the downstream genes mediated by estrogen may include mediators of the actin cytoskeleton.

In chapter 6, protein levels of four major Rho GTPase proteins (Cdc42, Rac1, RhoG and RhoA) and the focal adhesion kinase (FAK) were measured in M1 macrophages following treatment with/without estrogen.

FAK is a key molecule involved in regulating the actin cytoskeleton during phagocytosis (Cote-Vélez *et al.*, 2001; Tzircotis *et al.*, 2011). Immunoblotting analysis of FAK levels in this thesis indicated that estrogen supplementation significantly induced the levels of FAK in M1 macrophages. The importance of FAK in phagocytosis was highlighted by a study showing that *in vitro* transfection of macrophages with RNAi against FAK caused a 65% fall in the number of phagocytes during FcγR- and CR3-mediated phagocytosis (Tzircotis *et al.*, 2011). The pathogen-killing activities of neutrophils and macrophages were mediated by FAK through the formation of lamellipodial structures (Kasorn *et al.*, 2009). In contrast, FAK deficiency in macrophages is associated with a reduction in cell adhesion, and a failure in the formation of lamellipodia (Owen *et al.*, 2007). These findings suggest that FAK is a potential therapeutic target that may (at least in part) mirror the beneficial effects of estrogen on bacterial clearance by macrophages.

Whilst estrogen increased the levels of Rac1 in M1 macrophages, an opposite effect on RhoA was observed. This was interesting in light of studies showing that Rac1 enhanced the phagocytosis of apoptotic cells (Akakura *et al.*, 2004; Nakaya *et al.*, 2008) whereas RhoA was shown to dampen the process (Kim *et al.*, 2017). These reports also highlighted the need of a dynamic balance between the two proteins to regulate the apoptosis process, with concurrent upregulation of Rac1 and downregulation of RhoA required for engulfment of apoptotic cells. Similarly, other studies have also shown the opposing nature of Rac1 and RhoA during cytoskeleton reorganisation in *Candida elegans* (Ohta *et al.*, 2006; Freeman and

Grinstein, 2014). The concomitant increase of Rac1 and FAK as inducers of phagocytosis is consistent with the fact that Rac1 is downstream from FAK in the regulation of actin dynamics (Kallergi *et al.*, 2003; Papakonstanti *et al.*, 2003; Siesser and Hanks, 2006; Kallergi *et al.*, 2007). These findings suggest that manipulating the balance of Rac1 and RhoA in phagocytes of the elderly may be an additional therapeutic strategy to increase the appetite of macrophages to engulf bacteria, particularly in individuals where hormone replacement therapy (HRT) is contraindicated.

Similar to Rac1, RhoG is linked with the regulation of the actin cytoskeleton (Kato *et al.*, 2006), phagocytosis (Prieto-Sánchez *et al.*, 2006), gene transcription (Vigorito *et al.*, 2003) and the regulation of NADPH oxidase in neutrophils (Condliffe *et al.*, 2006). Novel data in this study demonstrate that stimulation of macrophages with estrogen results in an increase in RhoG levels and enhanced phagocytosis. This result is in concordance with the findings of Tzircotis *et al.* (2011) that showed RhoG stimulates phagocytosis of IgG-opsonised (for FcγR) or C3bi-opsonised (for CR3) sheep red blood cells (RBCs) in mouse macrophages. However, the findings in this thesis are the first to show that estrogen stimulates RhoG in human M1 macrophages. Thus, promotion of RhoG may be an effective and extremely novel therapeutic strategy for infected wounds since it may recapitulate (at least in part) the beneficial effect of estrogen on bacterial clearance. However, further investigations are required to confirm this hypothesis.

Activation of Cdc42 is required for the regulation of the actin cytoskeleton dynamics, the formation of membrane ruffling and the formation of filopodia, a crucial requirement for effective phagocytosis in macrophages (Kozma *et al.*, 1995; Cox *et al.*, 1997). Immunoblotting analysis in this study showed an increase in the levels of Cdc42 in M1 macrophages following supplementation with estrogen. The concurrent increased levels of both Cdc42 and FAK in M1 macrophages are consistent with Cdc42 being downstream of FAK in the signalling cascade that promotes pseudopodia formation by phagocytes (Schaller, 2010). The increased levels of Cdc42 agrees with the findings of Allen *et al.* (1997) who showed that injection of macrophages with active Cdc42 resulted in the formation of long and

indistinct filopodia *in vitro*. Thus, further investigations are warranted to determine whether the longer pseudopodia produced by macrophages in this study, following interaction with *P. aeruginosa* compared with MRSA could be associated with higher levels of Cdc42. Such work may lead to novel therapeutics designed to be more effective against problematic Gram-negative/motile bacteria such as *P. aeruginosa*.

7.2 Future Work

- In order to reinforce the findings of this study, *in vitro* and *ex vivo* investigations using other phagocytes such as neutrophils (e.g. HL-60 cell line or *ex vivo* neutrophils) are warranted. One development of this would be to determine the effect of estrogen on the formation of neutrophil extracellular trap (NET) formation during host-pathogen interaction assays. This would help build a wider picture of the effects of estrogen on the clearance of wound pathogens.
- Bacterial growth in chronic wounds has been reported to take the form of irregularly shaped micro-colonies that can comprise hundreds of bacteria known as biofilms (Bjarnsholt *et al.*, 2008; Rybtke *et al.*, 2015). Host pathogen assays in this study investigated the effect of estrogen on bacterial clearance by macrophages. However, only single (planktonic) bacterial pathogens were utilised throughout the study. Replicating the *in vitro* and *ex vivo* experiments using biofilms of MRSA and *P. aeruginosa* would improve our *in vitro* model of chronic wound infections.
- Unpublished research in the Ashworth laboratory has shown that androgens, such as testosterone and dihydrotestosterone (DHT) inhibit the phagocytosis of MRSA in U937-derived macrophages. Indeed, the literature suggests that testosterone has opposing effects to estrogen (Gilliver *et al.*, 2007). This creates many opportunities and directions for extending the

scope of this research, linking various sex steroid hormones with phagocytosis in wounds. Future work could investigate the effect of other steroid hormones and steroid hormone precursors such as dehydroepiandrosterone (DHEA) and dehydroepiandrosterone-sulfate (DHEA-S) on the phagocytosis of bacteria by various human phagocytes. Moreover, local conversion of testosterone to estrogen could be investigated using aromatase inhibitors and conversion of testosterone to DHT could be investigated using 5 α -reductase inhibitors.

- The findings of this study were obtained from host-pathogen assays conducted using *in vitro* and *ex vivo* macrophages. Repeating these experiments *in vivo* would provide robust evidence to confirm the beneficial effects of estrogen on promoting bacterial clearance during wound infections. One application would be to use mouse models that have been used previously to study age-related impaired healing (Ashcroft *et al.*, 1997b; Hardman *et al.*, 2007). *In vivo* investigations in animal models will be needed to develop potential therapeutic approaches for the treatment of wound infections before conducting human clinical trials involving chronic wounds in the elderly.
- The SEM data in Chapter 3 (Figure 3.16) illustrated distinct morphological changes occurring in macrophages when incubated with MRSA and *P. aeruginosa*. Future research is needed to determine the physical and biochemical effects induced by estrogen on pathogen-specific interactions with host cells. Fourier Transform Infra-Red (FTIR) Spectroscopy and Raman Spectroscopy could be used to determine changes in the functional groups on the membrane of host cells and cell wall of bacteria during host-pathogen interactions.

- Tamoxifen enhanced the phagocytic activity of macrophages against Gram-positive and Gram-negative bacteria (Chapter 5). Hardman and Ashcroft (2008) indicated that estrogen, tamoxifen and raloxifen all improve wound repair in ovariectomised mice. Future work could investigate a range of SERMs, including raloxifen, to determine the most effective mediator(s) of bacterial clearance. This could help identify the most promising topical SERM(s) to treat wound infections in the elderly.
- Several proteins involved in regulating the actin cytoskeleton were found to be mediated by estrogen (Chapter 6). Protein levels were measured by simple immunoblotting due to time restrictions, but these initial findings should now be confirmed by an additional technique, such as Western blotting. Moreover, Western blot analysis of phosphorylation (activation) patterns of these proteins at different time-points before and after stimulation with estrogen would add depth to current findings and potentially reveal further details on the mechanisms of estrogen-mediated phagocytosis. Intracellular flow cytometry could be used to explore these proteins in phagocyte populations, and kinase activity assays could be adopted to measure FAK activity. In addition, further investigations are required to determine whether the changes in protein levels are mirrored by concomitant alterations in gene expression. In order to determine the effect of ER- α activation on downstream gene expression in macrophages, RNAseq and qPCR analysis could be adopted. Future work would also be to replicate the findings using agonist/antagonist treatments to stimulate/block the activity of the proteins investigated in chapter 6. Inhibitors of phagocytosis could also be exploited to further understand the mechanisms by which estrogen induces phagocytosis.

7.3 Conclusion

In conclusion, this thesis delivers a body of experimental evidence demonstrating that estrogen directly enhances the phagocytosis of the two major wound

pathogens, MRSA and *P. aeruginosa*, by human macrophages via activation of ER- α and possible transcription of genes involved in regulating the actin cytoskeleton of phagocytes. When translating the findings to the context of wound infections, the data suggest that estrogen may promote the clearance of wound infections during youth but this protection is lost as estrogen levels decline with increasing age, resulting in increased propensity and progression of wound infections in the elderly. Understanding the precise mechanisms by which estrogen promotes phagocytosis would assist in identifying downstream targets to develop effective novel therapies for the treatment of wound infections with minimal side effects.

Based on the findings in this thesis, novel dressings that provide estrogen supplementation or selective activation of ER- α and/or specific targeting of proteins that regulate the actin cytoskeleton of phagocytes may provide effective therapeutic options for infected wounds in the elderly and warrant further investigation.

Chapter 8: References

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