

**THE ROLES OF SECRETED PROTEIN  
ACIDIC AND RICH IN CYSTIENE (SPARC)  
IN INTESTINAL INFLAMMATION,  
HEALING AND FIBROSIS.**

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**BSc(Hons.)**



**This thesis is presented for the degree of Doctor of Philosophy of  
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I declare that this thesis is my own account of my research and contains as its main content work which has not previously been submitted for a degree at any tertiary education institution.

.....  
(Yoke-Leng NG)

## **ABSTRACT**

Secreted Protein Acidic and Rich in Cysteine (SPARC) is a matricellular protein expressed during tissue repair and regulates cell proliferation and migration. It binds to, and interacts with collagen and regulates matrix metalloproteinase (MMP) expression. The aim was to determine if SPARC modifies intestinal inflammation, healing and fibrogenesis.

Intestinal disease was investigated using SPARC null (KO) and wild-type (WT) mice in which inflammation was induced by 3% dextran sodium sulphate (DSS) in the drinking water for 7 days. Inflammation was assessed endoscopically, clinically and histologically on days 7, 14, 21 and 35 after initiation of DSS treatment. Systemic and colonic cytokines and chemokines were quantitated by ELISA and CBA. Colon, mesenteric lymph node and spleen were analysed by flow cytometry and immunofluorescence for inflammatory cell infiltrates. To determine the effect of SPARC on the extracellular matrix (ECM) genes regulation, RNA from colonic tissue, and colonic myofibroblasts from WT and KO mice, were analysed by real time PCR for expression of ECM related genes.

KO animals had significantly lower endoscopic scores of inflammation, suffered less weight loss, diarrhoea, faecal blood and had lower spleen/body weight ratios compared to WT animals consistent with less colonic and systemic inflammation. WT mice had higher levels of histological inflammation and damage when compared to KO animals and in the majority of KO animals the colonic mucosa had completely regenerated by day 35 in contrast to the WT mice. Compared to WT mice, in KO animal colons at day 7 there was significantly less IL1 $\beta$  and MIG expression while TGF $\beta$ 1 levels were higher. Flow cytometric analysis identified a significantly greater percentage of FoxP3+ regulatory T cells in the spleen and draining lymph nodes of KO animals. KO mice also

had fewer of cells, such as CD68<sup>+</sup> macrophages and Ly6G<sup>+</sup> neutrophils, of the innate immune system infiltrating the inflamed colon.

Collagen (Col) 1 $\alpha$ 1, Col3 $\alpha$ 1 MMP13 and MMP3 expression levels were reduced in DSS-treated WT colons at day 7 and these were significantly lower than those observed in the KO colons. TIMP1 expression was significantly lower in KO mice at day 35 when healing was complete in this group. TIMP2 and TGF $\beta$ 1, TGF $\beta$ 3 were not different between the groups at any time point. The observation by others that collagen fiber diameters in KO colons were noted to be significant smaller than in WT animals suggesting that SPARC modifies the collagen bundling. Compared to unstimulated WT fibroblasts, KO cells had lower Col1 $\alpha$ 1 and Col3 $\alpha$ 1 expression. Stimulation with PMA reduced Col1 $\alpha$ 1 and Col3 $\alpha$ 1 and increased MMP13 and TIMP1 expression in all the isolated cells, but PMA had no effect on MMP3, TIMP2, TGF $\beta$ 1 and TGF $\beta$ 3 expression.

DSS induced less colonic and systemic inflammation in KO compared to WT mice and the inflammation appeared to resolve faster. This may be secondary to increased numbers of regulatory T cells and increased colonic TGF- $\beta$  levels which may inhibit effector cell activity, including cytokine and chemokine expression and aid in the more rapid resolution of inflammation and restoration of the intestinal mucosa. SPARC is able to modify tissue healing potentially through the regulation of collagen expression, bundling and its degradation by MMPs, which impacts on tissue turnover rate and thus delays healing. Hence, SPARC might represent a promising therapeutic target in clinical management of inflammatory bowel diseases (IBD).

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## CONFERENCE ABSTRACTS

**Ng Y.L., Kloplic B., Lloyd F, Greene W, Lawrance IC** “Combined Transforming Growth Factor- $\beta$ 1 and -3 Fibroblast stimulation modifies the Extracellular Matrix gene expression through Smad independant signalling”

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**Ng Y.L., Kloplic B., Lloyd F, Greene W, Lawrance IC** “SPARC attenuated Intestinal Inflammation in the DSS-treated Mouse?”

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**Ng Y.L., Kloplic B., Lloyd F, Greene W, Lawrance IC** “A role for SPARC in modifying inflammation and healing in the DSS murine model of colitis”

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**Ng Y.L., Kloplic B., Lloyd F, Fu S.K., Lawrance I.C.** “SPARC modifies colonic tissue healing and inflammation by regulating collagen and MMP expression”

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**Ng Y.L., Kloplic B., Lloyd F, Greene W., Lawrance I.C.** “SPARC enhances intestinal inflammation in dextran sodium sulphate-induced murine colitis”

Poster presented at 7<sup>th</sup> Congress of European Crohn’s and Colitis Organisation – February 2012; Barcelona, Spain

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## ABBREVIATIONS

The following abbreviations are used throughout this thesis:

$\alpha$ -SMA	$\alpha$ -smooth muscle actin
AF	Alexa fluor®
APC	Antigen presenting cells
BCP	Bromochloropropane
BSA	Bovine serum albumin
CCL28	Mucosal epithelial chemokine
CD	Cluster of differentiation
cDNA	Complementary deoxyribonucleic acid
COX	Cyclooxygenase
CrD	Crohn's disease
CTGF	Connective tissue growth factor
DC	Dendritic cell
DMEM	Dulbecco's Modified Eagle Medium
DMSO	Dimethyl sulphoxide
DSS	Dextran sodium sulphate
DTT	Dithiothreitol
ECM	Extracellular matrix
EDTA	Ethylene diamine tetra acetic acid
FBS	Foetal bovine serum
FSC	Forward scatter
FITC	Fluorescein isothiocyanate
FoxP3	Forkhead box P3
GALT	Gut-associated lymphoid tissue
GIT	Gastrointestinal tract
H&E	Haemotoxylin and eosin
HBSS	Hank's balanced salt solution
HCl	Hydrogen chloride
HEPES	N-2-Hydroxyethylpiperazine-N'-2-ethanesulfonic acid
IBD	Inflammatory bowel diseases
IEL	Intraepithelial lymphocytes
IFN	Interferon
Ig	Immunoglobulin
IL	Interleukin
ILF	Isolated lymphoid follicles
KCl	Potassium chloride

KHCO <sub>3</sub>	Potassium hydrogen carbonate
KO	Knockout
LAP	latency-associated peptide
LP	Lamina propria
LPMC	Lamina propria mononuclear cell
LPS	lipopolysaccharide
LTBP	Latent TGF- $\beta$ binding protein
MAdCAM	Mucosal vascular addressin cellular adhesion molecules
MgCl	Magnesium chloride
M cell	Specialised epithelial cells in GLT
MCP-1	monocyte chemotactic protein-1, CCL2
MEICS	Murine endoscopic score of colitis severity
MHC	Major histocompatibility complex
MIG	Monokines induced by IFN- $\gamma$ , CXCL9
MIP	Macrophage inflammatory proteins
MLN	Mesenteric lymph node
MMP	Matrix metalloproteinase
Na <sub>2</sub> CO <sub>3</sub>	Sodium carbonate
NaCl	Sodium chloride
NaN <sub>3</sub>	Sodium azide
NaOH	Sodium hydroxide
NF	Nuclear factor
NH <sub>4</sub> Cl	Ammonium Chloride
NOD	Nucleotide oligomerisation domain
OD	Optical density
Oligo	Oligonucleotide
PAI-1	Type-1 plasminogen activator inhibitor
PBS	Phosphate buffered saline
PCR	Polymerase chain reaction
PE	Phycoerythrin
PRR	Pattern recognition receptors
PMA	Phorbol 12-myristate 13-acetate
PMN	polymorphonuclear leucocytes
PDGF	platelet derived growth factor
PG	Prostaglandin
PP	Peyer's Patches
PSF	Penicillin /Streptomycin/Fungizone

RANTES	Regulated on activation normal T cell expressed and excreted, CCL5
RNase	Ribonuclease
RNasin	Ribonuclease inhibitor
ROR	Retinoic-acid-receptor-related orphan receptors
RT	Room temperature
SEM	Standard error of mean
SPARC	Secreted Protein Acidic and Rich in Cystiene
SSC	Side scatter
STAT	Signal transducer and activators of transcription
TBS	Tris-buffered saline
TBST	TBS-Tween
TGF	Transforming growth factor
Th cell	T helper cell
TIMP	Tissue inhibitors of metalloproteinases
TLR	Toll-like receptors
TNBS	trinitrobenzenesulfonic acid
TNF	Tumour necrosis factor
tPA	tissue type plasminogen activator
Treg cell	Regulatory T cell
TREM-1	Triggering Receptor Expressed on Myeloid cells-1
UC	Ulcerative colitis
uPA	Urokinase type plasminogen activator
UV	Ultra-violet
VEGF	Vascular endothelial growth factor
WT	Wild type