The Role of *Mycobacterium avium* ss *paratuberculosis* (MAP) in patients with Crohn's Disease.

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ABSTRACT

The cause of chronic inflammation in the gut of subjects with Crohn's Disease (CD) is unclear; however most would agree that 3 interacting factors are critical to mucosal inflammation: genetic susceptibility, enteric microflora and the host immune system. The most controversial theory is whether or not a particular microbe(s) infects and maintains in intestinal tissues, resulting in chronic inflammation. The most discussed microbe is *Mycobacterium avium* ss *paratuberculosis* (MAP). The aim of this thesis was to better document the natural history of MAP infection in subjects with Crohn's disease, IBS, "non-Crohn's colitis" and normal subjects, correlating clinical status with parameters relating to MAP, to the mucosal T cell response and to the genetic susceptibility gene CARD15/NOD2.

MAP is an obligate intracellular pathogen, which causes chronic inflammation in the intestine of many species, including primates. MAP was first identified as the causative agent in Johnes Disease, a gastrointestinal disease in ruminants and primates. The infection of domestic livestock with MAP is now widespread, which has increased the risk of transmission of MAP to humans via milk products. The involvement of MAP in human patients with Crohn's disease (CD) has been difficult to prove, due to the difficulties in isolating and detecting this organism.

By using nested PCR (Polymerase Chain Reaction) on DNA extracted from fresh human intestinal mucosal biopsy samples the presence of MAP can be verified. Bacteria contain unique insertion elements, which play a role in virulence, pathogenicity and antibiotic resistance. Within MAP a unique 1.4 kb insertion element was identified, IS900. By using DNA primers for PCR within this unique insertion element we were able to identify MAP in tissue. There was found to be no statistical significance among the disease groups tested.

The next objective was to determine whether the presence of MAP in gut biopsies was associated with a different cytokine secretion profile as observed in organ and whole blood culture, using an ELISA assay. Significantly higher levels of TNF- α

were found in culture supernatants from organ culture for Crohn's Disease when compared to ulcerative colitis (p<0.05), irritable bowel (p<0.01) and controls (p<0.0001). When TNF- α levels were correlated with the presence of MAP, significantly greater concentrations were only found in MAP positive Crohn's Disease patients (p<0.05). In whole blood culture significantly higher levels of IL-4 (p<0.05) and IL-2 (p<0.05) were found in MAP positive Crohn's Disease patients compared to MAP negative CD, which is consistent with a skewed Th2 immune response. This data provided the first evidence of an abnormal macrophage handling of MAP and an altered T cell function linked to MAP infection.

Finally, the co-existence of MAP infection and NOD2/CARD15 mutation status was investigated. SNP analysis of the three most common NOD2/CARD15 variants: missense mutations R702W (2104C \rightarrow T, SNP8) and G908R (2722G \rightarrow C, SNP12) and the frameshift mutation 1007fs (3020insC, SNP13); was performed on trial participants and compared to MAP status to determine if genetic susceptibility to CD predisposes them to MAP infection. Analysis of CD patients NOD2 status and cytokine profile found no correlation. Given that there is no clear link between NOD2 gene mutation, presence of MAP and cytokine secretion in our CD patients the suggestion that defective handling of MAP is due to NOD2/CARD15 gene mutations is not relevant to our group.

We can conclude that

- 1. MAP is not essential for CD,
- 2. That MAP is present in IBD and non-IBD patients,
- 3. There is a defect in the cellular handling of MAP in CD and
- MAP in CD has the capacity to enhance drive a Th2 response, which in turn down regulates the protective Th1 response and enhances mucosal permeability which leads to increased inflammation.

CHAPTER ONE: Literature Review.

1.1	Crohn's Disease	1
1.2	Pathogenesis of Crohn's Disease	2
1.3	Mycobacterium avium ss paratuberculosis – MAP	3
1.4	Immune response in Crohn's Disease	14
1.5	CARD15/NOD2 gene mutations	16
1.6	Hypothesis	19

CHAPTER TWO: MAP Detection and Culture from Mucosal Biopsies.

2.1	Introduction	21
2.2	Materials and Methods	23
2.2.1	Mucosal Biopsy Specimen Collection	23
2.2.2	DNA Extraction from Mucosal Biopsies	24
2.2.3	IS900 + pDIL60 Plasmid Preparation	26
2.2.4	IS900 Nested PCR	27
2.2.5	DNA Sequencing	30
2.2.6	Culture of Mucosal Biopsy Specimens	33
2.2.7	Analysis of Sera by Recombinant p35 and p36	34
2.3	Results	35
2.3.1	IS900 PCR Results	35
2.3.2	IS900 Detection in Biopsy Specimens	37
2.3.3	Mucosal Biopsy Culture	37

2.4	Discussion	40
2.3.4	Analysis of Sera by p35 and p36 Antigens	38

CHAPTER THREE: Pattern of Cytokine Secretion in Crohn's Disease Patients

3.1	Introduction	43
3.2	Materials and Methods	44
3.2.1	Organ Culture	44
3.2.2	Preparation of MAP Antigen for Culture Stimulation	45
3.2.3	Whole Blood Culture	46
3.2.4	Whole Blood Culture and IL-4 Cytokine ELISA	47
3.2.5	Cytokine ELISA – IL-2, IFN- γ , IL-10, IL-2 and TNF- α	52
3.2.6	Statistical Analysis	55
3.3	Results	56
3.3.1	Organ Culture	56
3.3.2	Cytokine Secretion from Organ Culture	57
3.3.3	Whole Blood Culture	60
3.3.4	Cytokine Secretion in Whole Blood Culture	60
3.3.5	Effect of MAP Antigen on Cytokine Secretion	62
3.4	Discussion	65

CHAPTER FOUR: Does the Presence of MAP Alter the Cytokine Secretion

Profile.

4.1	Introduction	68
4.2	Materials and Methods	69
4.2.1	Whole Blood Culture and IL-4 ELISA	70
4.2.2	Statistical Analysis	75
4.3	Results	76
4.3.1	IL-4 Secretion in Whole Blood Culture	76
4.3.2	IL-2 Secretion from Whole Blood Culture	78
4.3.3	IFN- γ and TNF- α Secretion in whole blood culture	78
4.3.4	Effect of MAP Antigen on Cytokine Secretion	78
4.3.5	Cytokine secretion in organ culture in relation to MAP status	80
4.4	Discussion	83

CHAPTER FIVE: Analysis of CARD15/NOD2 Polymorphisms by Real-Time PCR.

5.1	Introduction	
5.2	Materials and Methods	90
5.2.1	Preparation of DNA	90
5.2.2	CARD15/NOD2 SNP Analysis	93
5.2.3	Allelic Discrimination Analysis Procedure	
	using ABI Prism7900HT Sequence Detection System	95
5.2.4	Statistical Analysis	98

5.3	Results	100
5.3.1	CARD15/NOD2 Mutation Analysis	100
5.3.2	CARD15/NOD2 SNPs and MAP Status	103
5.3.3	CARD15/NOD2 and Cytokine Analysis	105
5.4	Discussion	111
CHAF	PTER SIX: FINAL DISCUSSION	114
REFE	RENCES	122
APPE	NDICES	153
1.	Solutions	153
2.	Published papers	162