## LEADING ARTICLE

# Microbes, immunoregulation, and the gut

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Two distinct, but rapidly converging, areas of research (the hygiene hypothesis and the study of probiotic/prebiotic effects) have emphasised the need to understand, and ultimately to manipulate, our physiological interactions with commensal flora, and with other transient but harmless organisms from the environment that affect immunoregulatory circuits. The story began with allergic disorders but now inflammatory bowel disease is increasingly involved.

# EARLY FORMULATIONS OF THE HYGIENE HYPOTHESIS

The hygiene hypothesis was first proposed in the late 1980s to explain the rise in allergic conditions (reviewed by Rook and colleagues<sup>1</sup>). The incidence of these disorders in the USA and Europe increased from the late 19th century, and appears to have doubled in some decades, particularly during the 1960s and 1970s. Epidemiological correlations with the modern way of life prompted the assumption that modern hygiene was reducing contact with pathogens that prime T helper 1 (Th1) responses. At that time it was believed that this would result in a compensatory increase in T helper 2 (Th2) activity that characterises allergic disorders. This concept, requiring Th1 inducing infections to control Th2 mediated allergic conditions, arose because of the remarkable compartmentalisation of medical knowledge. Readers of this journal, aware of the simultaneous increase in several Th1 mediated disorders such as Crohn's disease, type 1 diabetes, and multiple sclerosis, will be sceptical. Indeed, the incidences of allergic disorders (Th2) and of type 1 diabetes (Th1) correlate closely both within Europe and outside.2

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#### IMMUNOREGULATORY DISORDERS

The unifying hypothesis that can explain the simultaneous increase in autoimmunity and inflammatory bowel disease (IBD) (Th1 mediated) and allergies (Th2 mediated) is that modern living conditions can lead to defective maturation of regulatory T cells ( $T_{reg}$ ) and regulatory antigen presenting cells (APC<sub>reg</sub>). Therefore, rather than Th1/Th2 balance, the crucial factor is likely to be the effector T cells ( $T_{effector}$ )/ $T_{reg}$  balance. In the absence of optimal levels of immunoregulation, the individual may develop a Th1 or a Th2 mediated inflammatory disorder, depending on his/her own particular

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Th1/Th2 bias, immunological history, and genetic background. The argument that a similar lack of  $T_{reg}$  activity could underlie the increases in such diverse disorders as allergies, IBD, and autoimmunity is given added weight by the observation that mice or humans who have genetic defects of the transcription factor Foxp3, which is required for some  $T_{reg}$  functions, have a syndrome that includes components of all of these disease types. Thus diminished immunoregulation can lead to inappropriate immune responses to allergens, gut contents, or self (discussed by Rook and colleagues<sup>1</sup>).

#### T<sub>REG</sub> DEFECTS IN CHRONIC INFLAMMATORY DISORDERS

If this reinterpretation of the hygiene hypothesis is correct, the increase in human immunoregulatory disorders is at least partly attributable to defective T<sub>reg</sub> activity. Evidence to confirm this hypothesis has come from studies of allergic disorders,<sup>3</sup> multiple sclerosis,<sup>4</sup> autoimmune polyglandular syndromes,5 and cow's milk intolerance.6 It is likely to be true for IBD too, though more difficult to prove. The intestine is always in a state of controlled inflammation, and T cells of the regulatory phenotype are abundant in the guts of patients with IBD.7 Nevertheless, data from animal models of IBD suggest that the problem is likely to be an immunoregulatory one,8 and there is evidence that there is defective induction of oral tolerance in IBD patients.9 Moreover, they have exaggerated responses to bowel flora<sup>10</sup> which also appear to be the disease triggering antigens in animal models.8

# MICROBIAL EXPOSURE AND IMMUNOREGULATION

How does this  $T_{reg}$  orientated concept relate to the original hygiene hypothesis, and why would microbial exposure affect maturation of regulatory pathways? To answer these questions we must first establish what we mean by hygiene.

One interpretation of the word "hygiene" in this context, mostly promoted by the media, assumes that the critical factor is domestic hygiene (bathing, soaps, detergents, antibacterial kitchen cutting boards, etc). However, a comprehensive recent report has shown that the development of these practices in the home does not correlate with the observed changes in the occurrence of immunoregulatory disorders.<sup>11</sup>

Abbreviations: IBD, inflammatory bowel disease; Th1, Th2, T helper 1, 2; T<sub>effector</sub>, effector T cells, whether Th1 or Th2; T<sub>reg</sub>, regulatory T cell; APC, APC<sub>reg</sub>, antigen presenting cell/regulatory antigen presenting cell; DC, dendritic cell; TLR, toll-like receptor; KO, gene knockout; IL, interleukin; TGF- $\beta$ , transforming growth factor  $\beta$ 

A second view is that the critical change is the decreased frequency of infections due to pathogenic organisms. When the data available from the Centres for Disease Control and Prevention for the incidence of some infections are plotted against time, the graphs suggest that some of the decreases did occur during the critical period 1960-1985 when some chronic inflammatory disorders were doubling every decade.12 However, more detailed analysis of European data reveals that most of the changes in exposure to pathogens took place long before the crucial period.<sup>11</sup> In addition, there is strong epidemiological evidence to suggest that certain pathogens, such as childhood viruses and respiratory infections, cause an increase rather than a decrease in the incidence of allergic disorders.13 Interestingly, despite the detrimental effect of infections, this study still identified protective effects of being sent to day care, keeping pets, and living on a farm.<sup>13</sup> The latter has been a consistently robust observation, and the protective effect of exposing children to cowsheds is well documented.14 If childhood infections do not protect and home hygiene does not correlate, what might be the protective factors associated with pets, farms, and day care centres?

### "Contact with "old friends" is greatly diminished in rich countries but increased on farms, in cowsheds, and through contact with pets"

The answer might lie in certain relatively harmless microorganisms (including helminths, saprophytic mycobacteria, and lactobacilli) that have been present throughout mammalian evolution. We have called this the "old friends" hypothesis.1 Contact with "old friends" is greatly diminished in rich countries but increased on farms, in cowsheds, and through contact with pets. A number of reports have provided evidence for this interpretation. Allergic disorders are less frequent in individuals with helminth infections, and atopic sensitisation increases after treatment of intestinal helminths.15 Similarly, there are less lactobacilli in the guts of children with allergies,16 and a preliminary clinical study suggests that high doses of lactobacilli may inhibit development of atopic eczema in genetically high risk children.<sup>17</sup> Finally, the saprophytic mycobacterium M vaccae, originally isolated, as its name suggests, from a cow shed, potently drives maturation of T<sub>reg</sub> that will treat pre-existing allergy in a mouse model18 and has given encouraging results in clinical trials in allergic disorders.<sup>19 20</sup>

# THE MECHANISM BEHIND THE ''OLD FRIENDS'' HYPOTHESIS

We suggest that because of our long evolutionary association with these organisms, they are recognised by the innate immune system as harmless or, in the case of some helminths, treated as "friends" because a response would merely lead to immunopathology (fig 1). Therefore, rather than priming aggressive immune responses, these organisms prime immunoregulation.<sup>1</sup> They do it by inducing an unusual pattern of maturation of dendritic cells  $(DC)^{21\ 22}$  such that these retain the ability to drive  $T_{reg}$ . This effect requires the innate immune system. Toll-like receptor 2 (TLR2) may be involved for helminths<sup>21</sup> and TLR9 for lactobacilli.<sup>23</sup> It is interesting that polymorphisms of NOD2 (an intracellular receptor for bacterial peptidoglycan) are linked to increased susceptibility to both Crohn's disease<sup>24</sup> and asthma.<sup>25</sup> Thus an extension of the "old friends" mechanism suggests that in an environment that less actively primes  $T_{\rm reg}$  activity, immunoregulatory disorders will occur first in those individuals whose innate immune systems are least efficient at driving T<sub>reg</sub>.<sup>26</sup>

### BYSTANDER AND SPECIFIC IMMUNOREGULATION

The increased DC<sub>reg</sub> and T<sub>reg</sub> induced by "old friends" then lead to two immunoregulatory mechanisms mediated in part by release of interleukin (IL)-10 and transforming growth factor  $\beta$  (TGF- $\beta$ ). Firstly, continuing exposure to "old friends" will cause continuous background activation of T<sub>reg</sub> specific for the "old friends" themselves, resulting in a constant background of bystander suppression.

''Continuing exposure to ''old friends'' will cause continuous background activation of  $T_{\rm reg}$  specific for the ''old friends'' themselves, resulting in a constant background of bystander suppression''

This mechanism has been elegantly demonstrated in a model of colitis.<sup>27</sup> Secondly,  $DC_{reg}$  inevitably sample self, gut contents, and allergens and so induce  $T_{reg}$  specific for the target antigens of the three groups of chronic inflammatory disorder. These mechanisms may be aborted when there are legitimate "danger" signals. For example,  $T_{reg}$  function can be turned off by appropriate "danger signals" in vitro.<sup>28</sup>

### "OLD FRIENDS", THE GUT, AND PROBIOTICS

"Old friends" provide a conceptual link between the increase in allergic disorders, which triggered the formulation of the hygiene hypothesis, and the simultaneous increase in IBD, as similar organisms may be involved. For example, Weinstock *et al* have reviewed evidence that diminished exposure to helminths is a critical factor in the increase in IBD. They also have encouraging clinical results using oral delivery of the ova of *Trichuris suis*, which transiently colonises the human intestine.<sup>29</sup> Similarly, there have been several preliminary studies of the efficacy of bacterial probiotics, usually derived from lactobacillus strains, for inducing or maintaining remission in IBD (reviewed by Sartor<sup>30</sup>). This issue was discussed at a recent meeting of the International Scientific Association for Probiotics and Prebiotics (ISAPP; http:// www.isapp.net/).

# ''We hypothesise that in the context of IBD, the property that matters most is the ability to drive $T_{\rm reg}{}^{\prime\prime}$

Some of the effects of probiotics are beginning to be understood at a molecular level and involve striking mechanisms, including competition for ecological niches within the gut, inhibition of signalling via nuclear factor  $\kappa$ B, direct antimicrobial effects of secreted components, modulation of apoptosis, and activation of macrophages that take part in driving epithelial repair (partly reviewed by Ghosh and colleagues<sup>31</sup>). However, we hypothesise that in the context of IBD, the property that matters most is the ability to drive T<sub>reg</sub>.

### **PROBIOTICS AND TREG**

There is evidence that some probiotics can induce  $T_{reg}$ . Orally administered *Lactobacillus casei* reduced skin inflammation due to contact sensitivity in animals sensitised to dinitrofluorobenzene.<sup>32</sup> This finding cannot be attributed to the local gut specific effects of *L casei* but rather appears to require CD4+ T cells and is likely to have been mediated by  $T_{reg}$ . Moreover, some strains of lactobacillus mature DC so that they release little TNF- $\alpha$  or IL-12 but maintain their ability to release IL-10. This might facilitate induction of  $T_{reg}$ .<sup>33 34</sup> In IL-10 gene knockout (KO) mice, probiotics that attenuate the colitis to which these animals are susceptible downregulate Th1 cytokines while maintaining TGF- $\beta$ .<sup>35</sup> Both oral and subcutaneous administration promote this effect.<sup>36</sup> This activity of lactobacilli via the subcutaneous route protects

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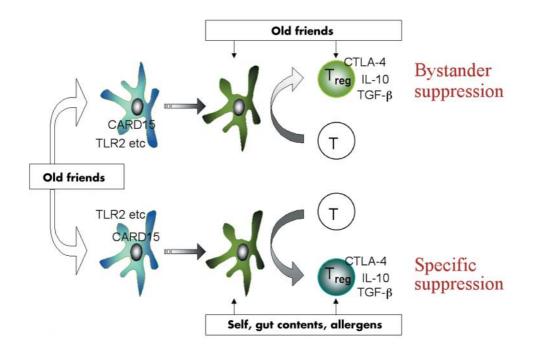


Figure 1 A diagram of the hypothesised modes of action of "old friends" and immunoregulatory probiotics. The "old friends", which have been present throughout mammalian evolutionary history, are recognised as harmless by the innate immune system and so cause antigen presenting cells (APC) to mature into regulatory APC (APC<sub>reg</sub>) that drive regulatory T cell ( $T_{reg}$ ). Some of these  $T_{reg}$  will recognise the "old friends" themselves and so provide continuous background bystander regulation. However, APC<sub>reg</sub> will also process and present epitopes from self, allergens, and gut contents, and so drive specific immunoregulation. These mechanisms can be silenced in the presence of appropriate danger signals. In the absence of "old friends", both specific and bystander regulation will be defective. IL-10, interleukin 10; TGF-β, transforming growth factor β; TLR2, toll-like receptor 2.

not only against colitis in IL-10 KO mice but also against collagen arthritis, a mainly Th1 mediated model of autoimmunity.36

#### "The gut may be the major site for T<sub>reg</sub> induction even when the probiotic is given subcutaneously'

The fact that probiotics work in models of colitis and arthritis, whether given orally or subcutaneously, is evidence that the important function in this context is not a gut specific one. Once generated, T<sub>regs</sub> can travel to other tissues. It is interesting however that the gut may be the major site for T<sub>reg</sub> induction even when the probiotic is given subcutaneously. Antigens containing bacterial polysaccharides<sup>37 38</sup> or whole organisms,<sup>39</sup> even when given parenterally, may evoke a pattern of response, detected as IgA secreting cells in peripheral blood, that mimics the response to mucosal immunisation. It seems likely that this is due to cross reactivity with antigens already experienced by the gut associated lymphoid tissue. Moreover, induction of IgA in the gut is heavily dependent on TGF- $\beta$  which is also closely involved in the maturation of T<sub>reg</sub>.<sup>40</sup>

#### CONCLUSIONS

The "old friends" hypothesis has evolved into a concept similar to that which lies behind attempts to modulate disease by altering the bowel flora. The strong parallels point to the following conclusions. Firstly, we suggest that the strains (whether "old friends" or probiotics; whether bacteria or helminths) used for clinical trials in disorders of immunoregulation (allergies, IBD, autoimmunity) ought to be those which can be shown to drive  $T_{reg}$ . Secondly, the particular organism used might need to be tailored to the individual patient. The "old friends" mechanism implies that, in rich countries, we live in an environment that does not efficiently prime T<sub>reg</sub> activity. This might precipitate immunoregulatory disorders most frequently in those individuals whose innate immune systems have genetic polymorphisms that further reduce the efficiency of  $T_{\text{reg}}\xspace$  induction.  $^{26}$  Clearly, an immunotherapeutic that requires TLR2 to drive T<sub>reg</sub> will not work in an individual in whom this receptor is nonfunctional.

Ultimately, the answer must lie in more rigorous proof that diminishing exposure to these  $T_{\rm reg}$  inducing organisms is a factor in the increase in immunoregulatory disorders. If confirmed, we will be able to devise subtle changes to our lifestyles that reconstitute this exposure by the oral route.

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Topics: 177; 191; 196