



- 1. Roadmaps for the development of diagnostic tests and therapeutics for helminths**
- 2. Roadmaps for the development of candidate vaccines and control strategies for liver fluke and nematodes**
- 3. Roadmaps for the development of candidate vaccines, diagnostic tests and control strategies for FMD**
- 4. Roadmap for research to underpin the development of control strategies for ASF**

*SIRCAH Deliverable 3.4*

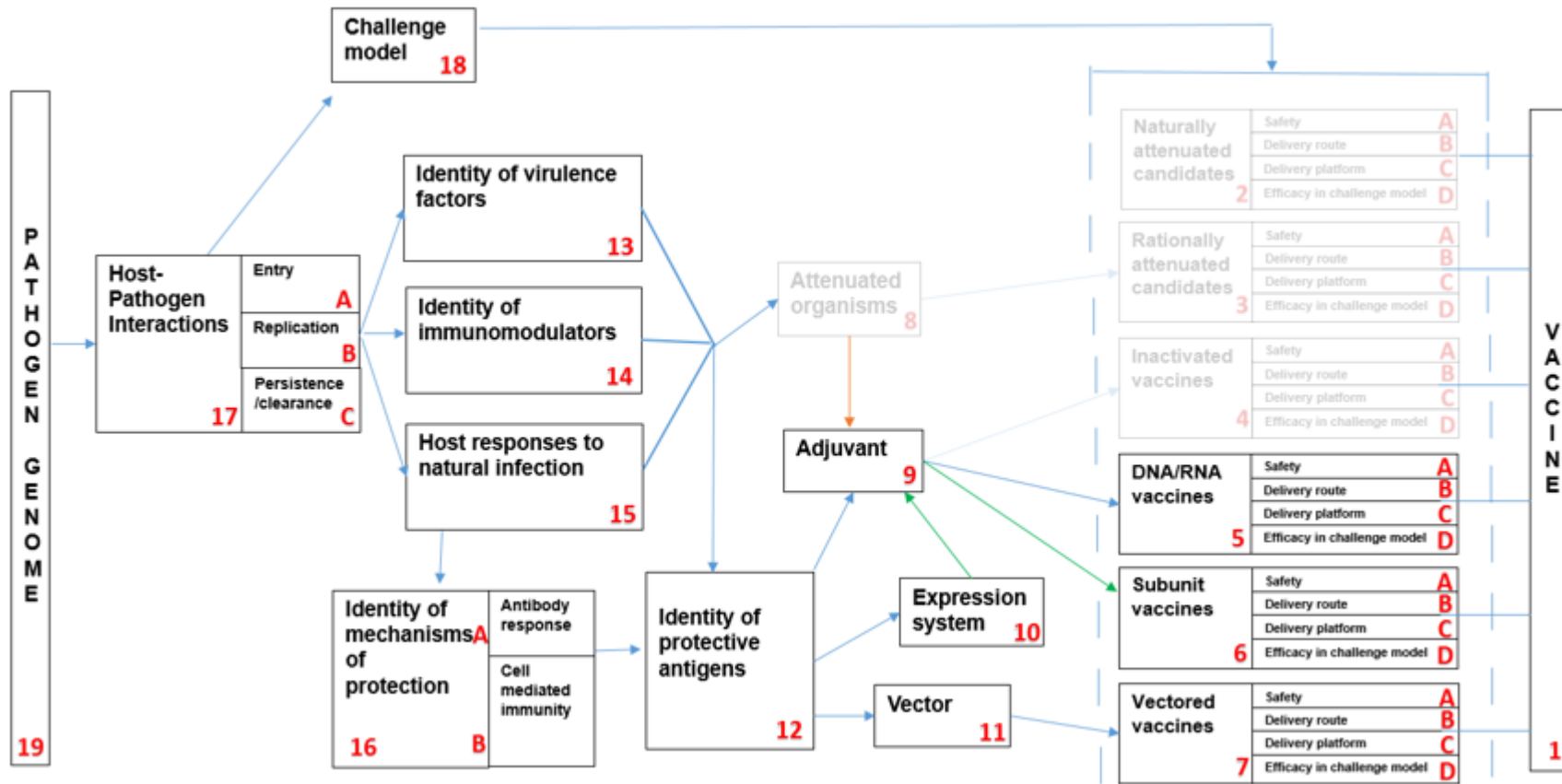
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**Interactive versions of the roadmaps in this report can be found at <https://roadmap.star-idaz.net>**



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## 2ai) Roadmap for the development of candidate vaccines for liver fluke



The roadmap for Trematode vaccine development has been developed by the Livestock Helminth Research Alliance (LiHRA; June 2019) with major contributions of Grace Mulcahy and Jozef Vercruyse.

## Lead Summary 1

**Title:** Vaccine giving long-lasting protection against infection by *F. hepatica*

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

A vaccine giving long-lasting protection with two initial injections followed by annual boosting, ideally in both cattle and sheep

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

Need for frequent boosting where hidden antigens are used  
The parasite modulates the hosts immune responses  
Unknown sources of variability in extent of protection in trials with identical protocols  
Need to communicate the level of efficacy likely to be delivered by fluke vaccines – will not be likely to match the 100% or near expected for many bacterial and viral vaccines

### Solution Routes

*What approaches could/should be taken to address the research question?*

Slow release of purified antigens from an adjuvant that drives the immune response in the correct direction.

Meta-analysis of the results of previous trials to understand the sources of variability  
Continued basic immunological studies to understand fluke-host interactions and fluke-induced immunomodulation  
Adjuvants capable of modulating the response to infection in a protective direction  
Use of a replicating vector to reduce number of boosting doses required  
Nucleic-acid based vaccines based including those based on self-amplifying RNA technology  
A combination of different candidates (subunit/nucleic acid based/peptide/vectored) in a Prime-boost approach  
Multivalent vaccines incorporating pathogens generating strong Th1-responses and counter-acting *F. hepatica* induced immunoregulation

### Dependencies

*What else needs to be done before we can solve this need?*

Difficulties in agreeing measures of protection – e.g. fluke burden, egg output, liver damage, specific pathology  
Modelling how much protection is required for an effective and commercial vaccine

More in-depth knowledge about host-pathogen relationships and the nature of immune response that is capable of being protective

### State of the Art

#### *Existing knowledge including successes and failures*

Protection of ruminants can be achieved using a number of antigens, both purified and recombinant, derived from *F. hepatica* E/S molecules. However, this is inconsistent.

Possible reasons for this inconsistency include:

Genetic variation in fluke strains used for challenge  
Differences in fine-specificity or other parameters of the host response  
Contribution of parasite glycans to protective and non-protective responses  
Host genetic background factors

Unknown unknowns!

### Projects

#### *What activities are planned or underway?*

PARAGONE, and EU-funded H2020 project involving vaccine development for liver fluke as well as for a range of other livestock parasites, will finish in 2019

A Science-Foundation Ireland project on basic immunology and applied vaccine science will finish in 2020

An SFI Research Professorship to J.P. Dalton at NUIG will commence in 2019

A number of national projects on fluke immunology and fluke vaccines are underway in Spain

## Lead Summary 5

**Title:** Development of nucleic acid-based vaccines

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

Generation of a protective immune response against *F.hepatica*

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

DNA vaccine approach hasn't worked as well in ruminants as in rodent models.

Anticipated regulatory challenges for this approach

### Solution Routes

*What approaches could/should be taken to address the research question?*

DNA vaccine

Self-amplifying RNA technology

Efficacy trials of existing candidates in experimental and field challenge studies

### Dependencies

*What else needs to be done before we can solve this need?*

A challenge model,

Definition of minimal acceptable efficacy

Identity of protective antigens

Identity of suitable molecular adjuvants

### State of the Art

*Existing knowledge including successes and failures*

Very limited evidence in the target species

### Projects

*What activities are planned or underway?*

No major projects are highlighted

## Lead Summary 6

**Title:** Development of subunit/peptide vaccine

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

Generation of a protective immune response against *F.hepatica* using parasite peptides/glycopeptides giving protection for up to a year.

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

Lack of a natural boosting effect from field exposure if hidden antigens are used

Immunomodulation by field infection

Obtaining protective immune responses in the range of MHC haplotypes represented in target populations, given the limited T-cell epitopes likely to be represented in a peptide-based vaccine

Obtaining recombinants that mimic the protective capacity of native proteins in the case of a sub-unit vaccine

### Solution Routes

*What approaches could/should be taken to address the research question?*

Identify antigens that are physiologically important for the parasite's development and or survival – secreted or hidden.

Identify of secreted antigens that the parasite uses to modulate host responses

Produce effective recombinants and/or peptides that induce protective, but not decoy or deleterious responses

### Dependencies

*What else needs to be done before we can solve this need?*

A challenge model,

Definition of minimal acceptable efficacy

Identity of adjuvants to give long lasting protective immune responses.

Identity of suitable expression vectors for large scale expression.

Identity of protective antigens

### State of the Art

*Existing knowledge including successes and failures*

Vaccine trials are under way evaluating a range of antigens.

Excretory products have been trialled – including cathepsin L2, GST, HDM, PRx, LAP

### Projects

*What activities are planned or underway?*

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## Lead Summary 7

**Title:** Development of a vectored vaccine

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

Using a replicating organism to give enhanced exposure to protective antigens

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

If several antigens are needed to give optimal protection the vector may need to have a relatively large genome.

That the expressed antigen has the correct conformation

That revaccination doesn't result in rapid elimination of the vector

### Solution Routes

*What approaches could/should be taken to address the research question?*

Development of a range of genetically modified organisms expressing protective antigens of one or more parasite species either as a secreted entity or as a surface molecule

### Dependencies

*What else needs to be done before we can solve this need?*

Identification of an effective and acceptable vector system

Interdisciplinary collaboration to achieve this

### State of the Art

*Existing knowledge including successes and failures*

There have been no studies to date on vector-expressed vaccines for *Fasciola*

### Projects

*What activities are planned or underway?*

None on this specific topic, currently



## Lead Summary 9

**Title:** Conventional and molecular Adjuvants

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

Establish the identity of suitable adjuvants for conventional (an effective antigen delivery system) and/or nucleic acid-based vaccines including an effective delivery system.

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

There is comparatively little empirical knowledge on the mechanism of action of available adjuvants  
In the case of newer adjuvants, many are within the IP portfolio of animal health companies  
There is a need to investment in the development of “smart adjuvants” capable of targeting the qualitative aspects of immune responses

### Solution Routes

*What approaches could/should be taken to address the research question?*

Trial different delivery platforms

Work with animal health companies to trial proprietary adjuvants with vaccine candidates

Establish the immune response and protection levels generated using various combinations of antigens and adjuvants

### Dependencies

*What else needs to be done before we can solve this need?*

Precise identity of protective immune response – Th1/Th2.  
Precise Identity of protective antigens.

### State of the Art

*Existing knowledge including successes and failures*

The analysis of a large number of trials carried out in both cattle and sheep by a number of groups independently has shown that it is possible to deliver protection of the order of 40-50% reduction in fluke burden. However, for unknown reasons, these results are variable, and some trials with identical protocols have not delivered protection.

### Projects

*What activities are planned or underway?*

It will be very important to make efforts to understand the reasons for this variability through a metaanalysis of previous studies

## Lead Summary 10

**Title:** Expression system for subunit vaccine

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

Development of an affordable stable expression system for large-scale production of the recombinant proteins/glycoproteins.

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

That the expressed antigens have the correct conformation and glycosylation  
Generation of stable genetically modified organisms allowing largescale production

### Solution Routes

*What approaches could/should be taken to address the research question?*

Various cell line expression systems – bacteria; parasite or plant cell lines  
Collaboration with commercial entities who have proprietary expression systems  
Certain plant protein expression systems are capable of mimicking helminth protein glycosylation systems

### Dependencies

*What else needs to be done before we can solve this need?*

Identity of the protective antigens  
Collaboration between groups expert in a variety of expression systems  
Detailed knowledge of the importance of parasite glycans in host-parasite responses

### State of the Art

*Existing knowledge including successes and failures*

Quill A has been used in a number of the vaccine trials in which efficacy has been demonstrated

### Projects

*What activities are planned or underway?*

Adjuvant studies are included in most of the projects underway at present

## Lead Summary 11

**Title:** Vector identification for Vectored vaccine

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

Using a replicating organism to express the protective antigens generating a stronger longer-lasting immune response

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

Immune response against the vector

### Solution Routes

*What approaches could/should be taken to address the research question?*

Explore a range of possible vectors – large viruses (Herpes viruses, endogenous retroviruses), protozoa (*Trypanosoma theileri*, *Eimeria*), bacteria (replication deficient such as aroA mutants)

### Dependencies

*What else needs to be done before we can solve this need?*

Identity of the protective antigens  
Successful antigen expression

### State of the Art

*Existing knowledge including successes and failures*

No specific trials with Fasciola antigens as yet.

### Projects

*What activities are planned or underway?*

None planned at present

## Lead Summary 12

**Title:** Identity of Protective antigens

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

To identify antigens against which protective immune responses can be generated

The parasite must be doing something which allows it to feed and evade the host responses as it migrates through the tissues – can these be used as immunogens

Alternatively there may be factors which could be targeted such as gut antigens

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

Generation of genetically modified parasites lacking “virulence” factors or immunomodulators or use of RNAi technology

### Solution Routes

*What approaches could/should be taken to address the research question?*

Reverse vaccinology to identify possible protective antigens.

Establish the identity and role of putative “virulence” factors and immunomodulators by generating genetically modified parasites.

**Establish which parasite genes are being expressed at different stages of feeding and their role**

### Dependencies

*What else needs to be done before we can solve this need?*

Improved understanding of the mechanisms of protection

- (i) The role of antibody classes in protection
- (ii) The role of cell-mediated immune responses
- (iii) Understanding the immuno-regulatory aspects of the target parasite

Transcriptomic studies of the host response

### State of the Art

*Existing knowledge including successes and failures*

Native fatty acid binding proteins induced 55% protection and cathepsin L1 42-69% protection in cattle.

Native leucine aminopeptidase(LAP) induced 89% protection in sheep.

Native glutathione S transferase – showed variable results in cattle and sheep  
Recombinant Schistosoma mansoni 14 antigen has shown positive results in some experiments.

F.hepatica rLAP shown positive response in sheep

Recombinant versions of Cl1 and CL3 have demonstrated protection in cattle

### Projects

*What activities are planned or underway?*

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## Lead Summary 13

**Title:** Identity of parasite virulence factors

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

To identify the factors that aid parasite migration and feeding. If the host could be immunised against these would it result in the parasite being unable to feed.

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

Understanding how the parasite interacts with and moves across the host intestine

### Solution Routes

*What approaches could/should be taken to address the research question?*

Studies on the immune responses of naturally exposed animals to establish the direction of immunomodulation  
Establish which parasite genes are being expressed at different stages and their role in parasite migration/feeding

### Dependencies

*What else needs to be done before we can solve this need?*

An improved understanding of Host-parasite interaction

### State of the Art

*Existing knowledge including successes and failures*

### Projects

*What activities are planned or underway?*

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## Lead Summary 14

**Title:** Identity of immunomodulators

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

To identify how the parasite modulates the host immune responses. If the host could be immunised against these would it result in the parasite being unable to migrate and be eliminated

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

### Solution Routes

*What approaches could/should be taken to address the research question?*

Establish which parasite genes are being expressed at different stages and their role in parasite migration/feeding  
Establish the roles of Cathepsins, peroxiredoxin and helminth defence molecules in immunomodulation

### Dependencies

*What else needs to be done before we can solve this need?*

An improved understanding of Host-parasite interaction

### State of the Art

*Existing knowledge including successes and failures*

We can describe the ways in which the response of infected animals is altered, and how it affects the response to other pathogens  
Modulation involves suppression of Th1 responses, induction of Th2/Treg responses, and polarisation of macrophages towards the alternatively-activated pathway.

### Projects

*What activities are planned or underway?*

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## Lead Summary 15

**Title:** Host response to natural infection

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

Identify what the host is responding to and its role in protection or possibly as a decoy dominant antigen which is preventing resistance developing

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

Establish the protective mechanisms required to kill juvenile and adult flukes  
Use NGS and bioinformatic tools to map the pathways up- and down-regulated following infection

As animals do not become immon-protective une to re-infection, these must be n

### Solution Routes

*What approaches could/should be taken to address the research question?*

Establish the dynamics of host immune responses under field conditions.  
Study the response of "Resistant" animals compared to those that are highly susceptible  
Transfer studies to establish the role of antibody and immune cell populations  
Studies on the immune responses of naturally exposed animals to establish the direction of immunomodulation

NGS studies of local and systemic immune responses in target species  
NGS studies of animals after protective-vaccination

### Dependencies

*What else needs to be done before we can solve this need?*

We are ready now to address these steps

### State of the Art

*Existing knowledge including successes and failures*

There are recent publications on the ovine systemic and hepatic responses to infection

### Projects

*What activities are planned or underway?*

A study of the bovine systemic transcriptomic response has been completed and ready for publication. A study of the transcriptomic changes in hepatic lymph nodes of sheep following challenge infection is underway



## Lead Summary 16

**Title:** Identify Mechanisms of protection against *F. hepatica*

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

Studying the transcriptomic response of ruminants to the acute and chronic stages of infection  
Identifying gene expression pathways involved in immunoregulation and whether these are open to manipulation  
Identify how animals that are considered more resistant are controlling/containing the infection. What are the roles of the various antibody classes and cellular responses.

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

The role of parasite glycans in the immune response, in immunomodulation and in the early stages of host-parasite infection

### Solution Routes

*What approaches could/should be taken to address the research question?*

Study the response of "Resistant" animals compared to those that are highly susceptible  
Role of Molecular pattern recognition (interaction with TLRs) in determining the immune response.

Cell types involved in the various stages of infection, including Th1/Th2 balance in "resistant" and highly susceptible animals  
Identification of specific protective epitopes within putative vaccine antigens

### Dependencies

*What else needs to be done before we can solve this need?*

A better understanding of the host-parasite interaction.

### State of the Art

*Existing knowledge including successes and failures*

In cattle a Th1 response with high titres of IgG2 and low titres of IgG1 has been correlated with protection.  
Conversely, infected animals produce specific IgG1 and almost no specific IgG2, and are not protected  
A proportion of the antibody response generated in natural infection may be directed at non-protective, "decoy" antigens

### Projects

*What activities are planned or underway?*

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## Lead Summary 17

**Title:** Host-parasite interactions

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

How the host-parasite interact allowing or otherwise parasite migration, feeding and reproduction

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

How is the very early stage of infection, across the intestinal mucosa, orchestrated

### Solution Routes

*What approaches could/should be taken to address the research question?*

Establish which parasite genes are being expressed at different stages of feeding and their role in parasite feeding.  
Establish the degree of genetic variation there is in the parasite populations and the impact of these variations on the host-parasite interaction.  
Role of epithelial activation in initiating the immune responses.  
Establish the importance of the different interaction of tegumental Ags and ES Ags with TLR4 and dendritic cells.  
Establish the role of a range of Th2 cytokines, especially IL-25, IL-33 and TSLP  
Identify the parasite proteins/Ags that modulate dendritic cell maturation  
Establish the role of basophils early in infection  
Establish the role of Antibody-dependent Cell cytotoxicity

Are macrophages and eosinophils targets of immunomodulation

### Dependencies

*What else needs to be done before we can solve this need?*

Parasite genome sequence - annotated

### State of the Art

*Existing knowledge including successes and failures*

In Schistosoma infected mice after an early phase of antigen specific cellular proliferation and IFN $\gamma$  synthesis there is a shift to a Th2 profile with IL-4 and IL-13 production predominating and a decrease in cellular responses. By the time infection becomes patent the response is dominated by IL-10 and TGF $\beta$  and IgG1 predominates.

Initial recognition of the metacercariae takes place in the GIT.

The tegument of NEJs is primarily composed on oligomannose and core fucosylated truncated N-glycans. Tegumental Ag induce anergic T-cells via dendritic cells

In both cattle and sheep IgG2 are linked with the expression of resistance or protection

### Projects

*What activities are planned or underway?*

Parasite glycan characterisation

Fluke genome projects

ERC grant

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## Lead Summary 18

**Title:** Challenge model

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

What are acceptable challenge models for screening of vaccine candidates

Will we obtain different results with bolus as compared with trickle infections?

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

Is there a difference in the vaccine response to trickle (field) and bolus challenges

Can we predict differences in vaccine protection based on fluke strain

Is there a non-challenge proxy for estimating protective capacity

### Solution Routes

*What approaches could/should be taken to address the research question?*

Meta-analysis of previous trials used in an attempt to determine and standardise the optimum vaccine challenge model

### Dependencies

*What else needs to be done before we can solve this need?*

What would regulators accept in relation to minimal efficacy

Understand the contribution of fluke genetic variability on protection following vaccination

### State of the Art

*Existing knowledge including successes and failures*

Protective effect has been observed using both trickle, and bolus, challenge models to date. The same is true for non-protective trials

### Projects

*What activities are planned or underway?*

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## Lead Summary [*Number*]

**Title:** Bystander Immunoregulation

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

What effect does Fasciola infection have on the diagnosis and control of other livestock pathogens?  
How can this relationship be utilised to enhance disease control in a holistic fashion

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

Funding for such studies which have a less obvious “end-product” than more applied vaccine studies

### Solution Routes

*What approaches could/should be taken to address the research question?*

Clearly communicate the utility of such studies

### Dependencies

*What else needs to be done before we can solve this need?*

Incorporate sophisticated epidemiological modelling studies along with immunological insights into co-infection

### State of the Art

### *Existing knowledge including successes and failures*

*F. hepatica* infection has been documented as decreasing the responsiveness of cattle to diagnostic tests for bovine tuberculosis, but also decreasing bacterial load in co-infected cattle  
The epidemiological significance of these effects needs to be dissected and understood.  
Some evidence is beginning to be compiled concerning interaction between liver fluke and other pathogens including *Mycobacterium paratuberculosis*, and *Calicophoron daubneyi*

### Projects

*What activities are planned or underway?*

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