

Department for Environment Food & Rural Affairs



National Centre for the Replacement Refinement & Reduction of Animals in Research

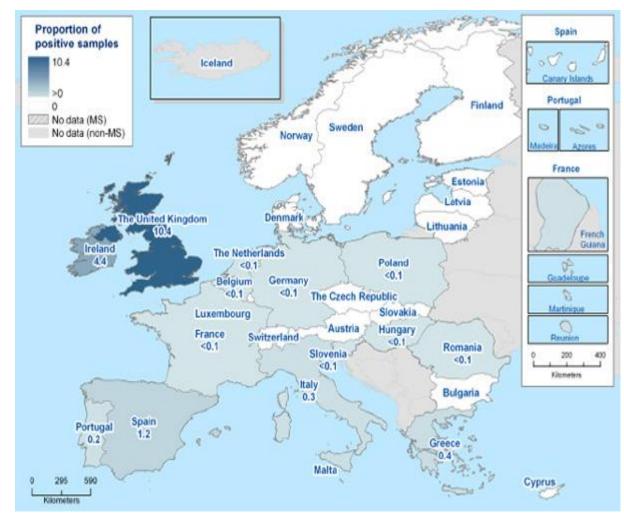
A bovine alveolus model to replace cattle in the study of host-pathogen interactions in bovine tuberculosis Diane Lee School of Veterinary Medicine University of Surrey, UK



- Mycobacterium bovis can infect almost all warm-blooded animals
- 1900
 - 40% of British cattle suspected to be infected
 - M. bovis responsible for ~15% of all human deaths from TB
- 1950s
 - Pasteurisation of milk and 'test-and-slaughter' introduced
- 1970s
 - 0.22% of tests positive for bTB
- 2016:
 - ~5%

Bovine tuberculosis (bTB)

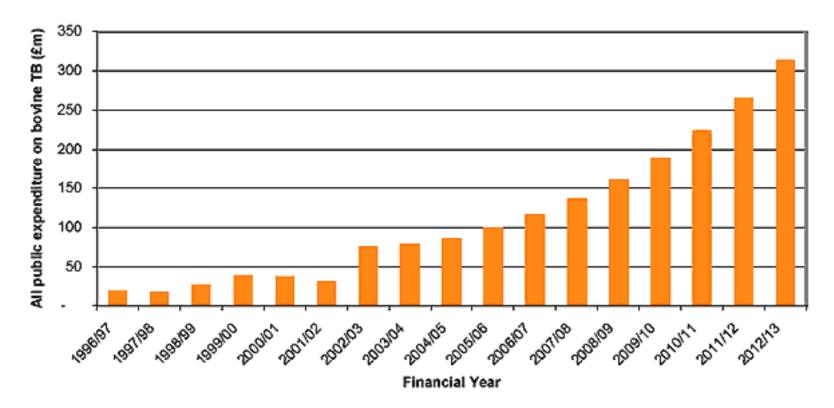




Proportion of existing cattle herds infected with or positive for *M. bovis* country based-data, 2012. EFSA Journal 2014

Bovine tuberculosis (bTB)





The Strategy for achieving Officially Bovine Tuberculosis Free status for England - 2014 https://www.gov.uk/government/uploads/system/uploads/attachment_data/file/300447/pb14088-bovine-tb-strategy-140328.pdf

https://www.gov.uk/government/news/tb-strategy-ahead-of-schedule-as-england-set-toapply-for-officially-tb-free-status-for-half-the-country "Dealing with Bovine TB in England ... required the culling of 28,000 cattle in 2015"

Bovine tuberculosis (bTB)



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ŧ	The 3Rs	Our science	Our resources	Funding	News	s & Blogs	Events	About us		
Home >	Home > Funding > Our funding schemes > Strategic awards > Replacing animal models of bovine tuberculosis									
Our funding schemes			Replacing animal models of bovine tuberculosis							
Project grants		lub								
Pilot study grants PhD Studentships David Sainsbury Fellowships		This c	This call is closed. Focus The current UK bovine tuberculosis (bTB) epidemic is one of the biggest challenges facing cattle farming in the UK with very serious economic and animal welfare consequences. The disease has spread from isolated pockets in the 1980s to cover large areas of the west and southwest of Britain, illustrating its increasing incidence.							
		in the U isolated								

- Strategic award between NC3Rs and Defra (2015)
- Replace cattle for the study of the pathogenesis of bTB by providing a tissue culture model with which to study fundamental events following infection of the bovine lung with virulent mycobacteria that can't be conducted currently *in vitro*



- Using human lung epithelium-endothelium submerged bilayer models:
 - Epithelial cells are hosts for *M. tuberculosis* and allow virulent bacteria to access the deeper tissues Birkness (1999)
 - Peripheral blood mononuclear cells (PBMCs) migrate from the basal layer to the apical layer during infection
 - Airway epithelial cells play an essential role in both innate and adaptive immune responses against *M. tuberculosis* PRR expression
- A bilayer of human pulmonary artery endothelial cells and human alveolar epithelial cells at an air-liquid interface on opposing sides of a Transwell developed to study invasive pulmonary aspergillosis
 - Professor William Hope, University of Liverpool (NCR3Rs grant, G0700599)



- Objective 1
 - Isolate and immortalise bovine type II alveolar epithelial (B2AE) cell line
- Objective 2
 - Assemble a bovine alveolus (boAlv) culture model using bovine cells (adaptation of the Hope model)

• Objective 3

 Demonstrate functional utility of the boAlv model by introducing PBMCs from cows expressing a strong or weak vaccine protection phenotype (VPP)



- Objective 1
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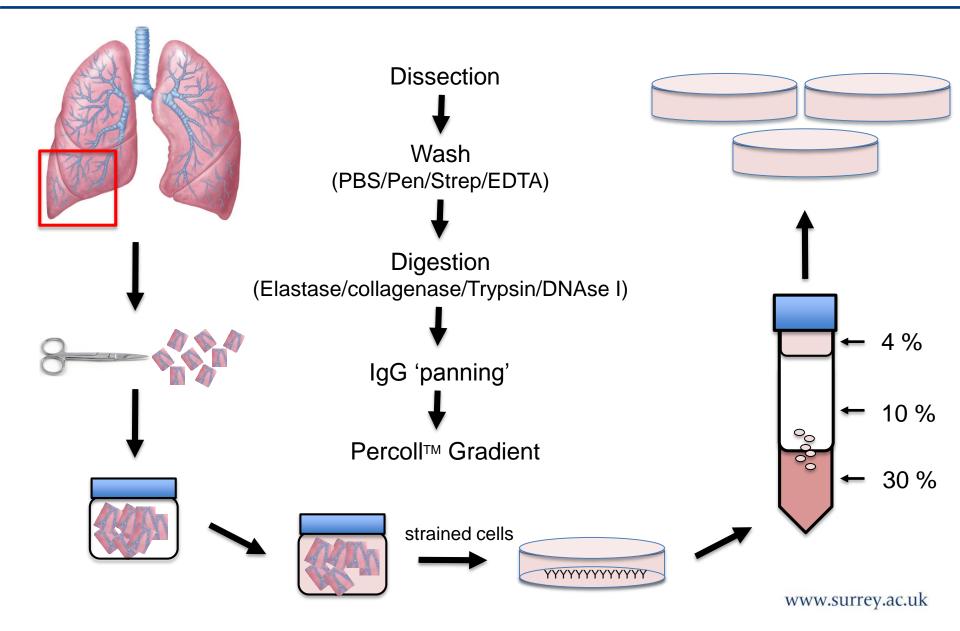


Objective 1 – Isolate and immortalise bovine type II alveolar epithelial (B2AE) cell line



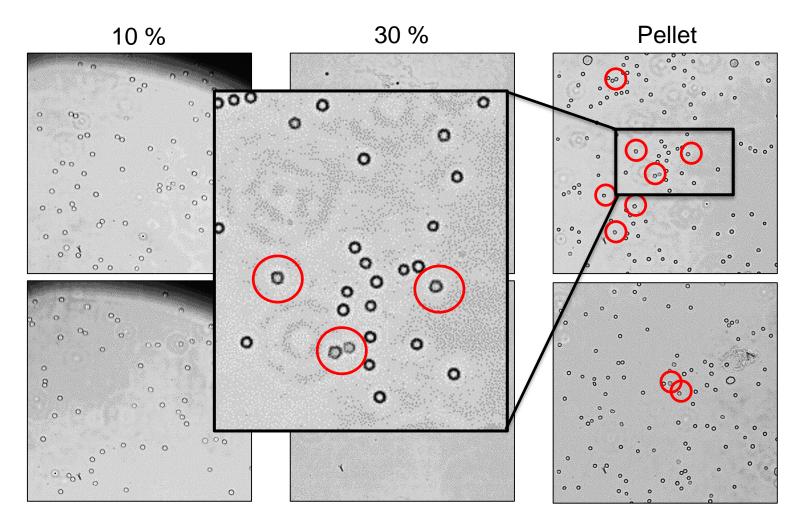
Objective 1 (i) Devise and implement an isolation strategy





Objective 1 (i) Devise and implement an isolation strategy – results 1

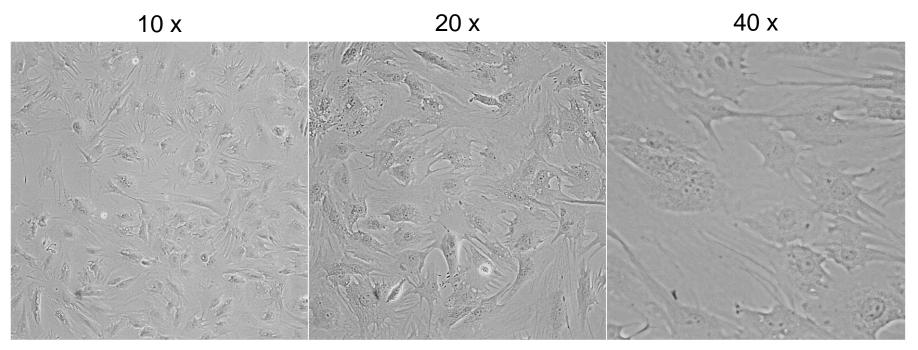




Day 1 post-isolation

Objective 1 (i) Devise and implement an isolation strategy – results 2

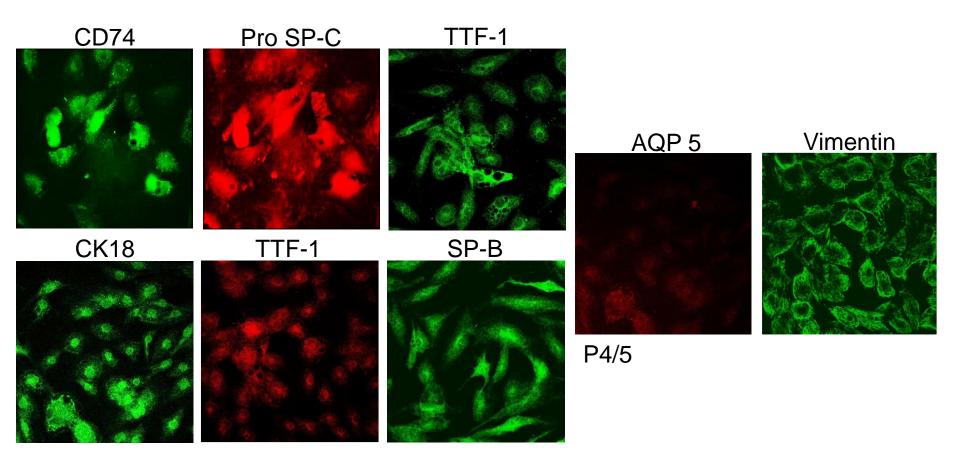




p2

Objective 1 (i) Devise and implement an isolation strategy – results 3

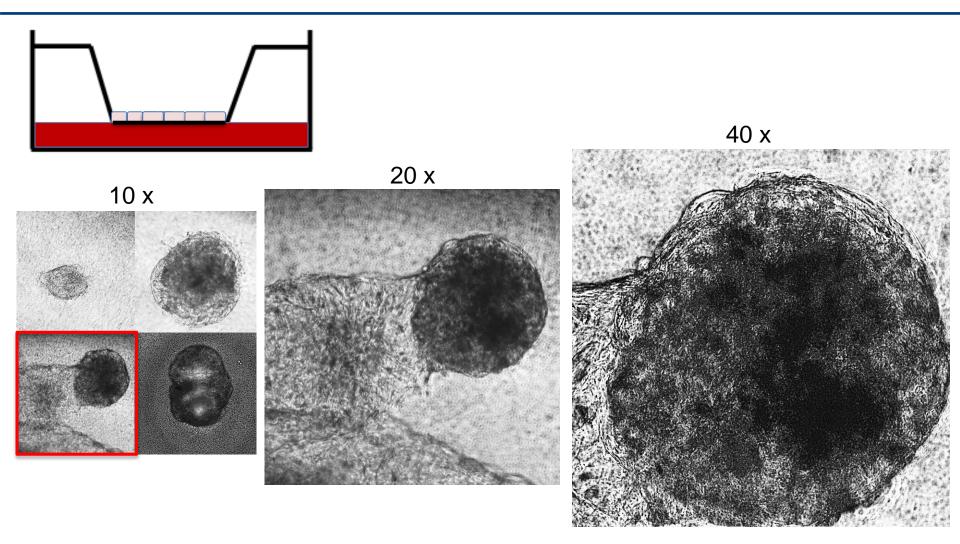




p2

Objective 1 (i) Devise and implement an isolation strategy – results 4 - ALI cultures

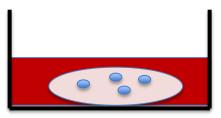


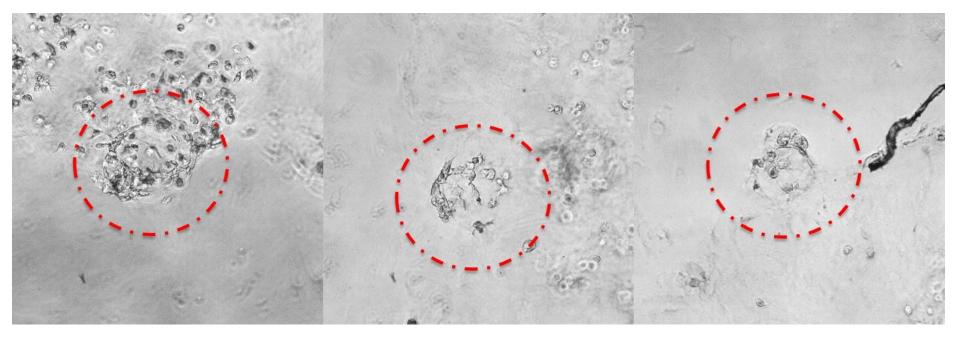


5 days on MG-coated Transwell-Clear 12 mm inserts; p5

Objective 1 (i) Devise and implement an isolation strategy – results 5 - 3D ECM



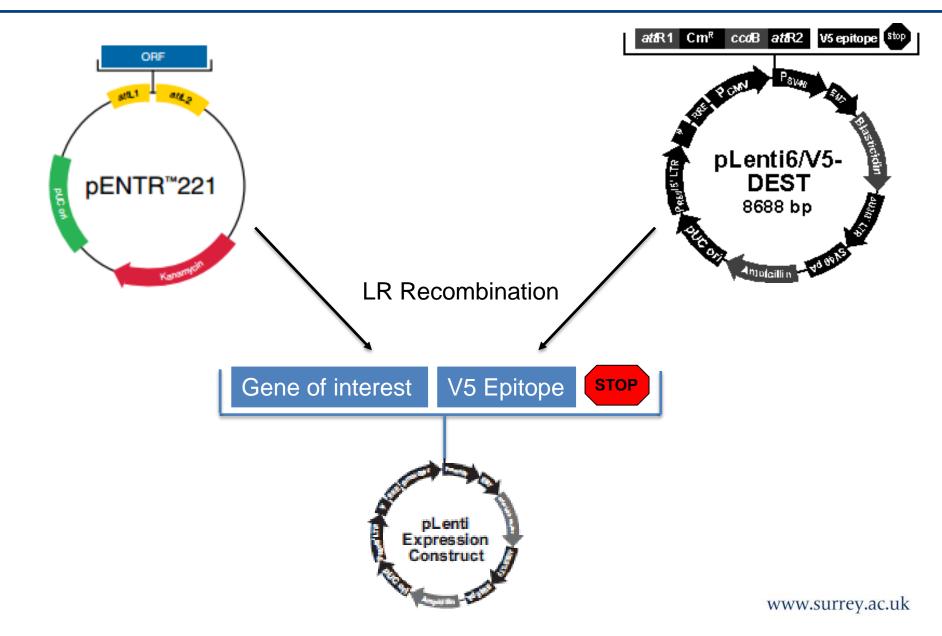




10 days in 3D MG, submerged in SAGM; 10 x, p6

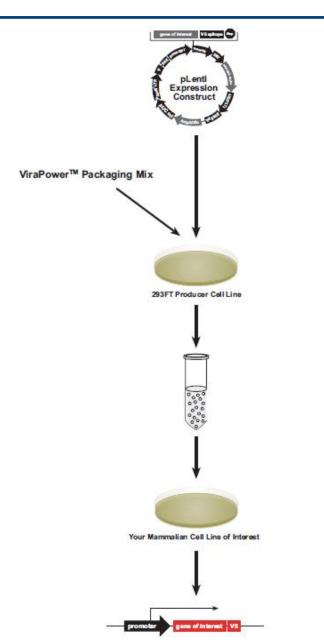
Objective 1 (ii) Immortalise isolated cells using Bmi1 and hTERT lentiviral constructs





Objective 1 (ii) Immortalise isolated cells using Bmi1 and hTERT lentiviral constructs





Co-transfect 293FT producer cell line with expression construct and packaging mix

Harvest viral supernatant and determine titre

Transduce mammalian cells and select for clones

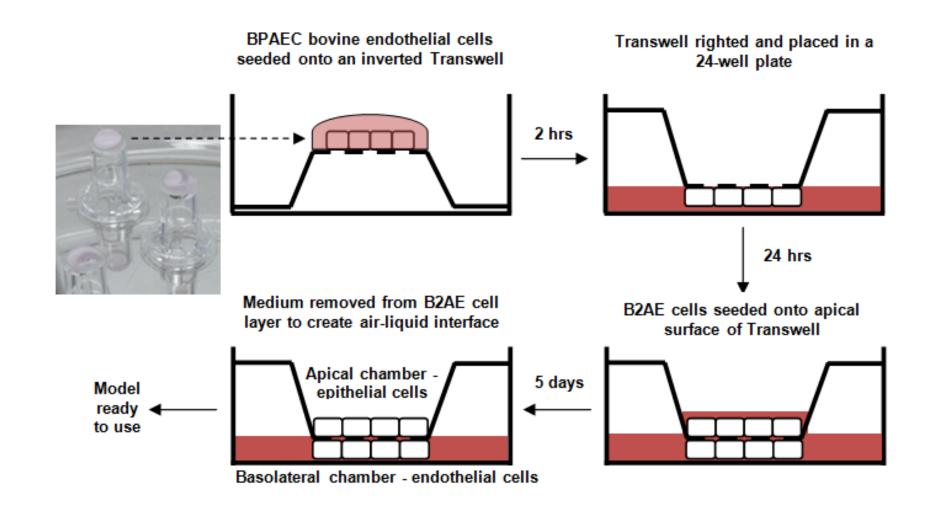


Criteria		Status		
Digest of expression construct	(At miniprep and midiprep stages)	Digest verified at both stages		
Sequencing (external)	(CMV and V5 C-term)	Sequence verified using diagnostic PCR and Sanger sequencing		
Sequencing (internal)	(two sets of primers designed for hTERT)	Sequence verified using diagnostic PCR and Sanger sequencing		
Blasticidin selection	(at titration and transduction stage)	Titrations obtained for both lentiviral preparations: > 2 x 10 ⁷ TU/mL		



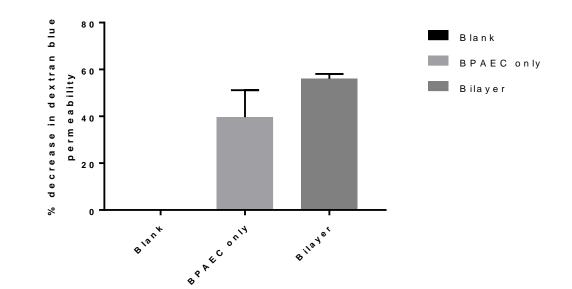
Objective 2 - Assemble a bovine alveolus (boAlv) bilayer culture model using bovine cells

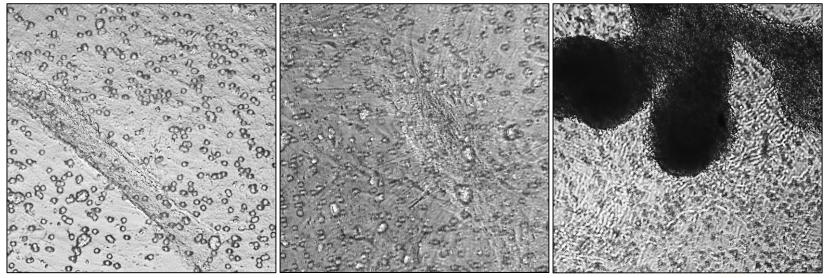




Objective 2 - Assemble the boAlv culture model – Results 1

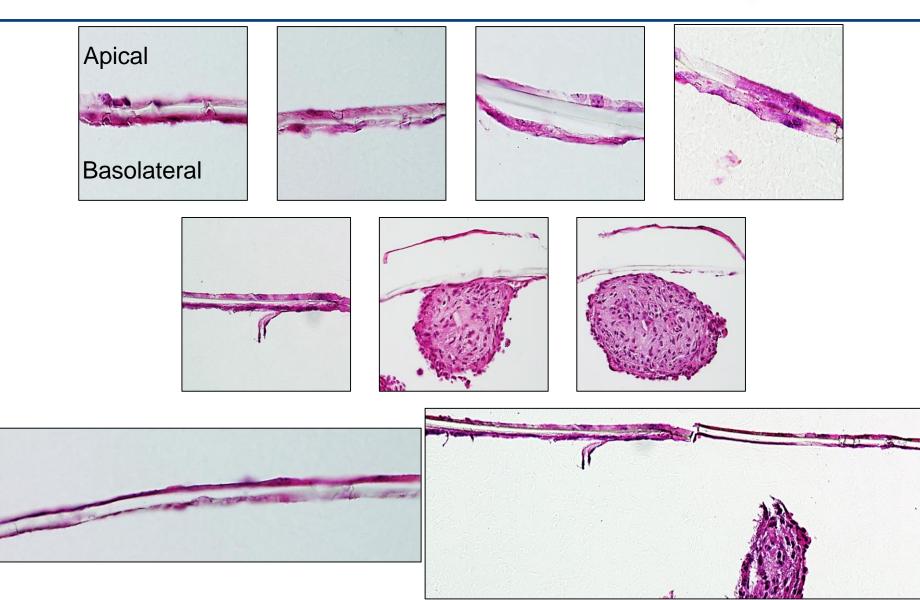






Objective 2 - Assemble the boAlv culture model – Results 2





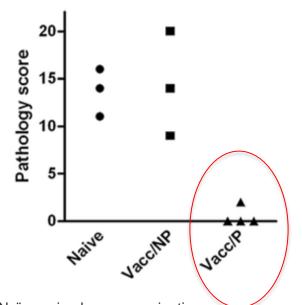


- Perform fresh isolation of ATII cells from bovine lung and immortalise (B2AE cell line)
- Characterize clones in parallel with WT ATII
 - Bmi1/hTERT expression (qRT-PCR, WB)
 - Growth curve
 - Karyotype
 - ATII markers SP-B, SP-C, TFF1, CK18, CD74 (qRT-PCR and/or IF)
- Optimise bilayer model
 - seeding densities
 - culture period



- Evaluate at least 4 possible scenarios associated with vaccine protection, related to the speed and activity of the host response and the pathogen behaviour within the alveolus:
 - PBMCs from strong VPP animals restrict better the growth of mycobacteria;
 - PBMCs from strong VPP animals restrict better the migration of mycobacteria through the epithelium -> endothelium;
 - PBMCs from strong VPP animals migrate more efficiently and in greater numbers through the endothelium->epithelium;
 - PBMCs from strong VPP animals express and stimulate a greater ratio of IL-22/17

Distribution of vaccinated cattle into those that express a strong and weak VPP



Naïve animals = no vaccination Vacc/NP = vaccinated calves that were not protected (weak VPP) Vacc/P = vaccinated calves that were protected (strong VPP) Taken from: Bhuju S et al. PLoS Pathog 8(12): e1003077.





Prof. Mark Chambers Dr Javier Salguero-Bodes Prof. Graham Stewart Prof. Martin Vordermeier

Ms Abbe Martyn Mr John Cooper

Ms Ella May Mr Duncan Grainger Mrs Gillian Wallis

Prof. William Hope Ms Clara Negri



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