# Polymorphic Immune Molecules and Their Functional Diversity Across Species

# Program 3 - 5 November 2025

Monday November 3 <sup>rd</sup>	13:00	Natasja de Groot MHC Diversity in Great Ape Species: Evolution and
13:00 Registration open	13:30	Host-Pathogen Interactions Thomas Goeury Markov chain modelling of MHC sequences in 38 primates species reveals complex evolutionary events Pascale Gerbault Contrasting patterns of nucleotide diversity along MHC class I genes A, B and C in bonobos, chimpanzees and humans
<ul><li>18:00 Paul Norman Introduction</li><li>18:15 John van Rood Prize Lecture:</li><li>John Trowsdale FMedSoc</li></ul>	13:45	
Immune Defence and Polymorphism  19:00 Welcome drinks and dinner	14.00	
Tuesday November 4th	14:00	Richard Natarajan A quantitative characterisation of the extent of homopolymer and repeat region polymorphism within the MHC
09:00 Louise Boyle	14:15	Break
Peptides Controlling the Orchestration of Immune	14:45	John Hammond
Responses	14.40	Combining Genomics and Single Cell Sequencing
09:30 Griffin Kutler Dodd		to Understand the Evolution of MHC and NK Cell
Chimpanzees and gorillas lack an MHC-A peptide-		Receptors in Non-Model Species
binding specificity common in humans	15:15	Avik Bhattacharya
09:45 Malcom Sim		High-resolution investigation of the LILR gene
High-throughput discovery of novel HLA-C bound		complex unveils extensive allele and allotype
peptide ligands for NK cell receptor KIR2DS4		diversity
10:00 Jim Kaufman	15:30	Jamie Winternitz
A tissue-specific helminth-inducible class II system in		High-resolution genotyping of complex immune gene
birds (and mammals?)		families in model and non-model species using hybrid
10:15 Break		target capture and long-read sequencing
10:45 Jacek Radwan	15:45	Susan Ott
MHC Genes in the Red Queen Race 11:00 Joanna Różańska-Wróbel		The characterization and annotation of the highly
		diverse genomic IG regions in rhesus macaques
Interplay between vertebrate adaptive immunity and	16:00	Lightning talks (x6)
bacterial infectivity genes: Bank vole MHC versus	19:00	Drinks and dinner
Borrelia afzelii OspC	20:30	Erik Rozemuller
11:15 Javier Avalos Nunez		A Royal Dessert
Searching for signatures of natural selection within		
the regions encoding NK cell receptors 11:30 Tobias Lenz		
II:30 TODIAS LENZ		

T-cell repertoire diversity analysis in a small teleost as a natural model system for adaptive immunity

11:45 Lunch

# Wednesday November 5th

# 09:00 Josephine Pemberton FRS

Maintenance of MHC Class II Variation in a Wild Sheep

09:30 Dorothea Heimeier

MHC diversity in the New Zealand southern right whale (Eubalaena australis; Tohorā nō Aotearoa)

09:45 Johannes Dijkstra

Extreme allelic MHC class I diversification in bony fish: evolutionary and structural perspectives of the salmonid example

10:00 Tony De Tomaso

Exceptional diversity of allorecognition receptors in a non-vertebrate chordate reveal principles of innate allelic discrimination and convergent evolution of mammalian immunity

#### 10:15 Break

#### 10:45 Anna Savage

MHC Polymorphism and Disease Susceptibility in Wild Amphibian and Reptile Populations

11:00 Magdalena Migalska

Major Histocompatibility Complex in Salamanders: annotation pitfalls and evolution of complex immune regions in giant genomes

11:15 Dimitri Monos

The complete haplotypic characterization of the swine MHC using ONT adaptive sampling and sequencing. Optimizing the swine xenotransplantation model

11:30 Saghar Kaabinejadian Canine MHC Class I Molecules and their Human Homologs

### 11:45 Lunch

#### 13:00 Alexander Mentzer

HLA, Hepatitis B and Viral Evasion: Unravelling Vaccine Immunity in Bangladeshi Infants

13:30 Keyi Zheng

Investigating the Role of Chromosome 14 IGH Locus in COVID -19 Vaccine Immunogenicity and Infection

13:45 Kristen Wade

Novel multiple sclerosis MHC risk loci impact expression of cellular homeostasis , innate immune and  $\,$ 

14:00 Katherine Kichula

Dual effects of maternal killer cell immunoglobulinlike receptor genotype on preeclampsia risk in Ugandan women

#### 14:15 Break

# 14:45 Ed Hollox

Copy Number Variation of Immunity Genes in the Age of Biobanks

15:15 Gabriela Sato Paes

Population-Specific Diversity of KIR2DS1 and KIR2DS4 Activating Receptors: Allelic Variation and Copy Number Patterns Across Global Ancestries

15:30 Joanna Malukiewicz

Organization, Diversity, and Copy Number Variation in the Major histocompatibility complex (MHC) region of Brazilian Marmosets (Callithrix)

15:45 Jesse Bruijnesteijn

Structural diversity and allelic polymorphism in the LILR gene cluster of humans and rhesus macaques

16:00 Closing Remarks

19:00 Drinks and dinner

## Lightning Talks

Jonathan Beaumont Clinton Azuure Christopher Thorpe Daniel Alvarez-Sirvent Dominic Barker Jonathan Downing