For improved pertussis vaccines a more detailed understanding of vaccine-induced immunity is required



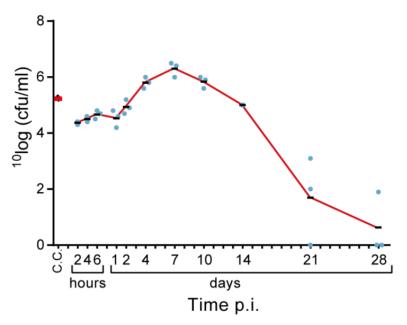


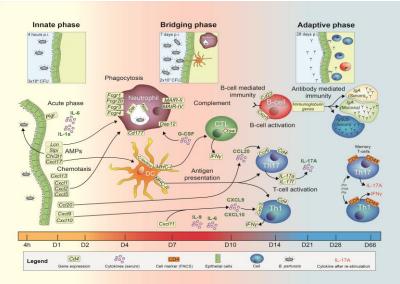
Molecular Signatures of the Evolving Immune Response in Mice following a *Bordetella pertussis* Infection

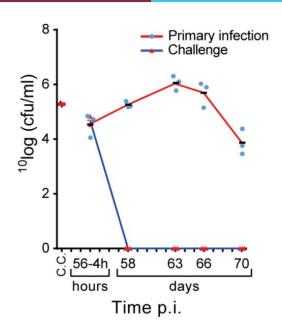
René H. M. Raeven^{1,2*}, Jolanda Brummelman³, Jeroen L. A. Pennings⁴, Olaf E. M. Nijst¹, Betsy Kuipers³, Laura E. R. Blok¹, Kina Helm³, Elly van Riet¹, Wim Jiskoot², Cecile A. C. M. van Els³, Wanda G. H. Han³, Gideon F. A. Kersten^{1,2}, Bernard Metz¹

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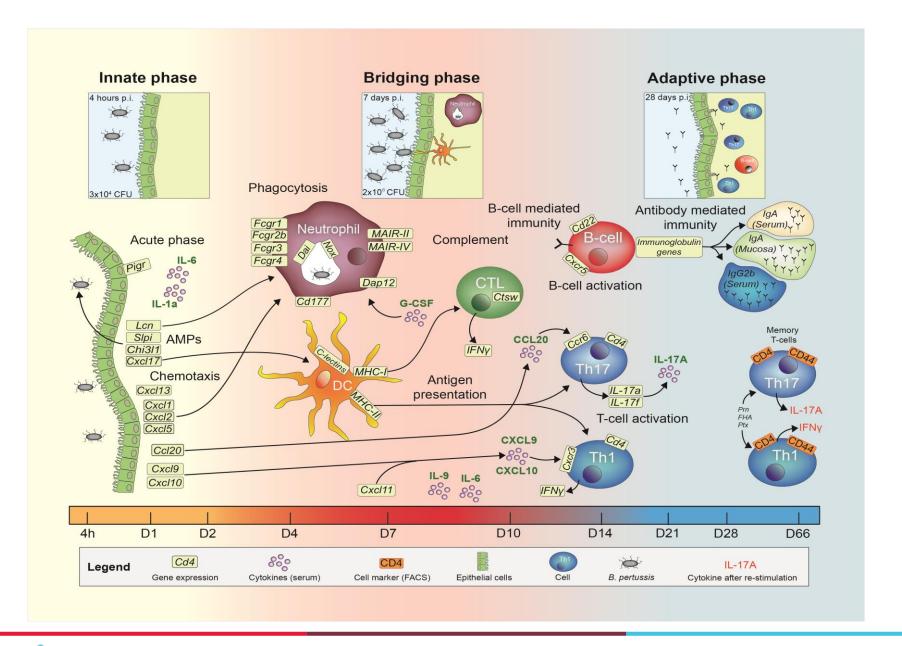
Sterilizing immunity

Different layers:

Transcriptomics
Proteomics
Cellular response
CFU

Kinetics







Which antigens should be presented?

and...

In what kind of package should they be wrapped?



Article

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Immunoproteomic Profiling of *Bordetella pertussis* Outer Membrane Vesicle Vaccine Reveals Broad and Balanced Humoral Immunogenicity

René H. M. Raeven,**,*,* Larissa van der Maas,* Wichard Tilstra,* Joost P. Uittenbogaard,* Tim H. E. Bindels,* Betsy Kuipers,* Arno van der Ark,* Jeroen L. A. Pennings,* Elly van Riet,* Wim Jiskoot,* Gideon F. A. Kersten,*,* and Bernard Metz*

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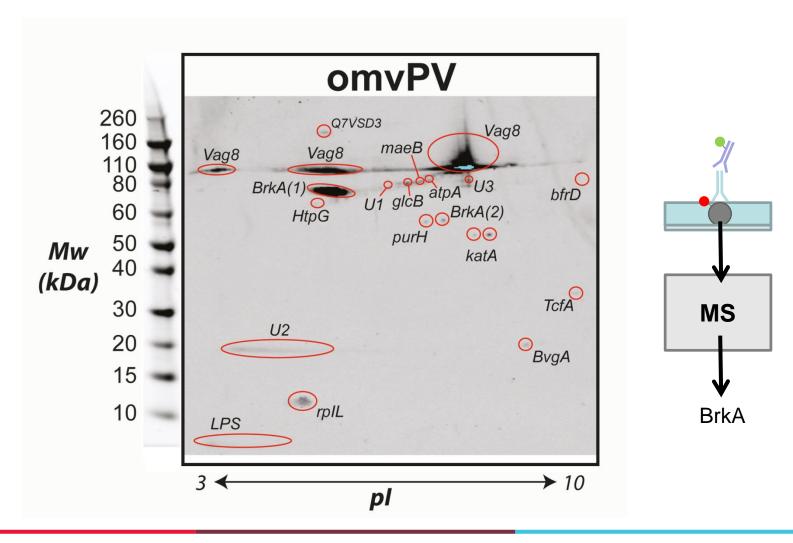


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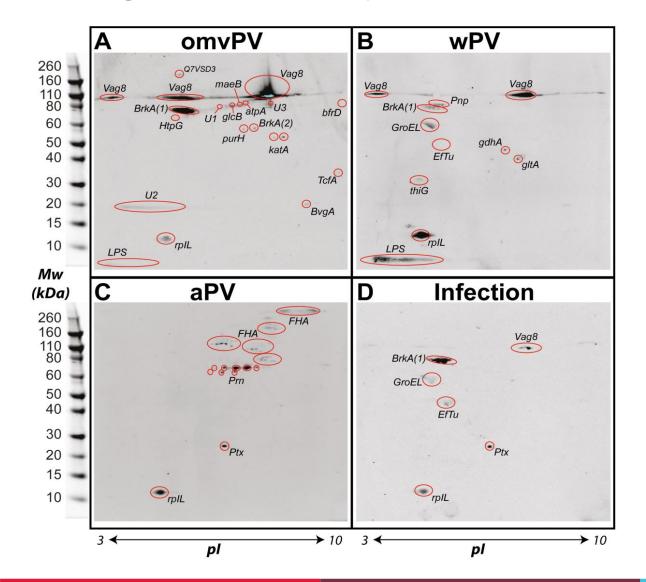
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2D-immunoblotting

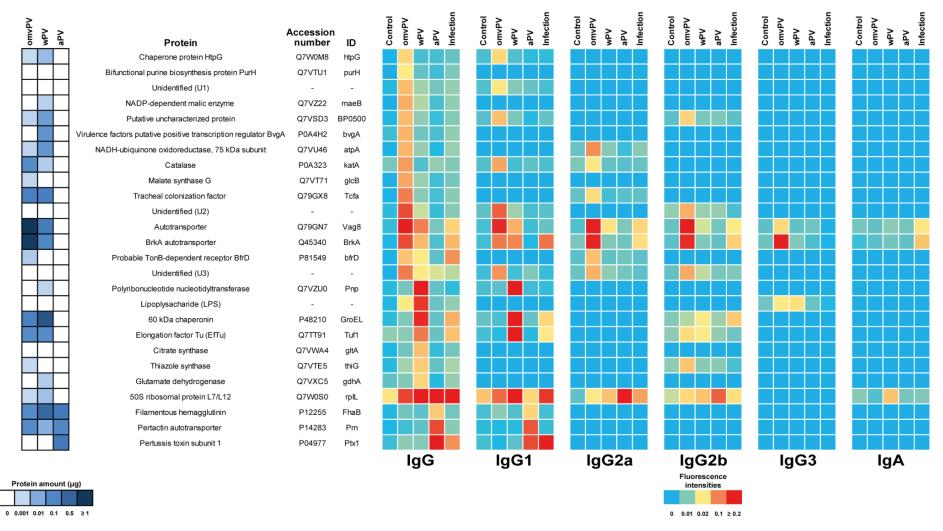


Antigen specificity



	# Abs
omvPV	16
wPV	9
aPV	3
Infection	5

Antibody profiles

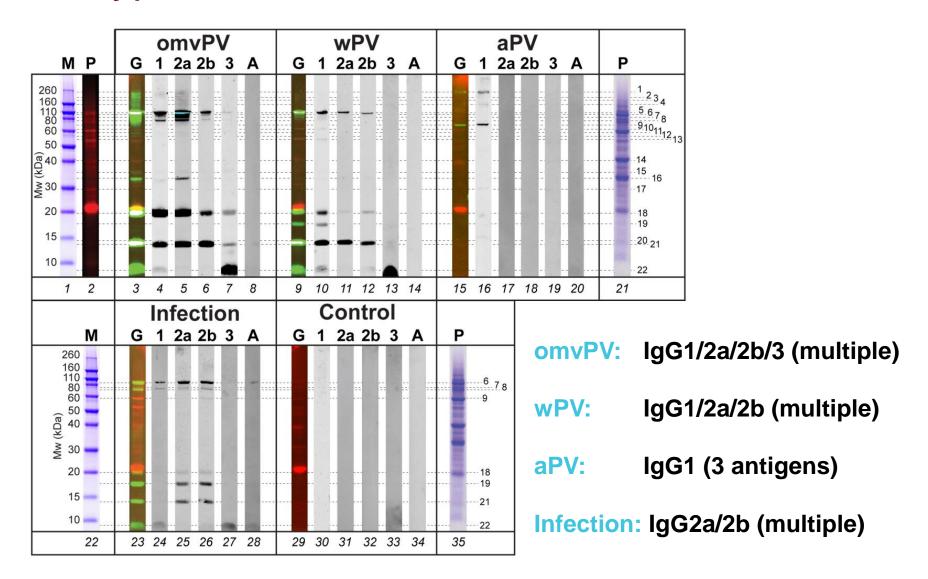


Protein concentration LC-MS

IgG (n=3), subclasses and IgA (n=1), warping with Delta2D software



Isotype and subclass distribution





Lung response

<u>Transcriptomics:</u> 558 genes

- Up (red): 446

- Down (green): 112

Functional analysis:

GO-BP terms **KEGG** pathways

Result:

Chronological cascade of immunological process

 $p \le 0.001$; FR ≥ 1.5

