

**Evidence on vaccine benefit from Human Infection
Models: insights and considerations**

Robert Read

University of Southampton

United Kingdom



Edward Jenner FRS
Physician and zoologist,
1749-1823

circa 1796

23 Subjects

Inoculated with Cowpox
pus

Subsequently injected with
variolous material

£10,000 grant from Royal
Society

Widespread vaccination
with cowpox begun 1840

Controlled Human Infection Models

Bacterial

Campylobacter

Cholera

E. Coli – (ETEC, EPEC, DAEC, EAaggEG)

Francisella tularensis

Gonorrhoea

H. Pylori

Salmonella Typhi and Paratyphi

Shigella

BCG

Bordetella pertussis

Streptococcus pneumoniae

Streptococcus pyogenes

Neisseria lactamica

Viral

Dengue

Influenza

RSV

Norovirus

Rhinovirus

Coronavirus

Rotavirus

Parasitic

Malaria

Giardia

Cryptosporidium

Leishmania

Rickettsial

Rocky Mountain

Spotted FEVER

Helminthic

Hookworm

Ethics and Informed Consent

- Risk of harm
 - Immunity, virulence/attenuation, antimicrobial activity, monitoring
- Quality and Value of Research
 - Benefit:risk
 - Professionalism and competence
- Justice
 - Self determination
 - Equality

Ethical Considerations: Controlled Human Infection

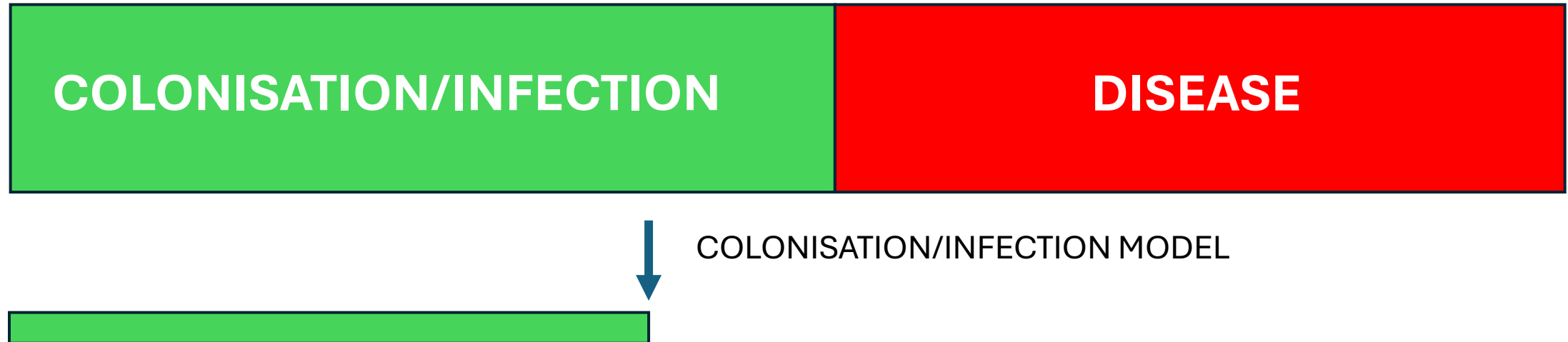
- **The objective can be justified**
- **The minimum adequate sample is used**
- **The challenge inoculum is the minimum required to provide a clear outcome measure**
- **Induced symptoms can be treated, and treatment is not withheld**

Controlled Human Infection models

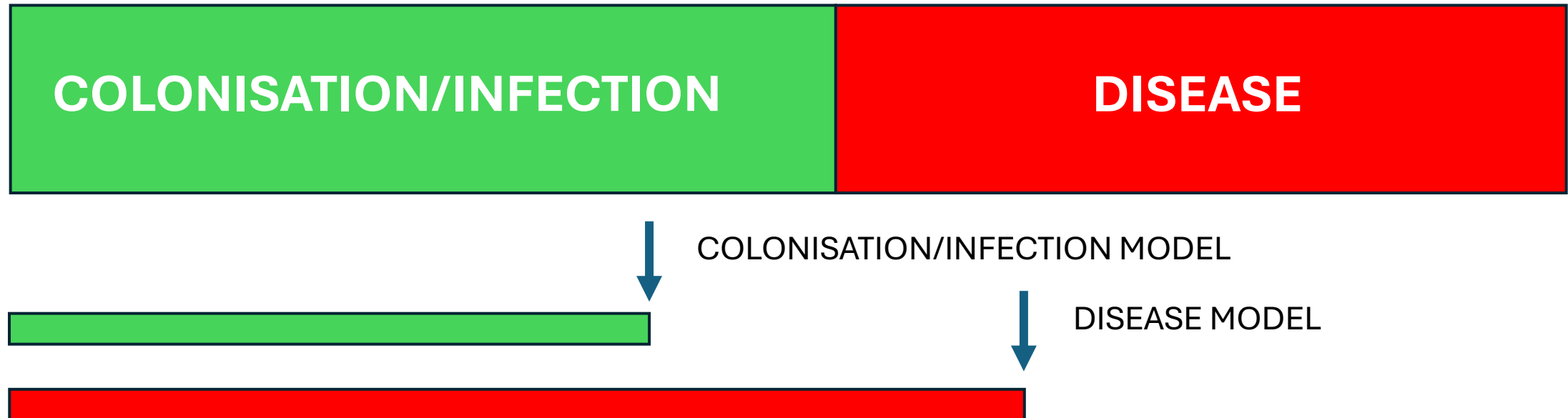
COLONISATION/INFECTION

DISEASE

Controlled Human Infection models



Controlled Human Infection models



Single-dose Live Oral Cholera Vaccine CVD 103-HgR Protects Against Human Experimental Infection With *Vibrio cholerae* O1 El Tor

Clinical and Bacteriologic Responses to Challenge

Clinical or Bacteriological Endpoint	All Vaccinees	Vaccine Day 10	Vaccine 3 mo	All Placebos	Vaccine Efficacy (95% CI) or P Value	
					Day 10	3 mo
Total challenged	68	35	33	66		
No. with severe cholera (%)	3 (4.4)	1 (2.9)	2 (6.1)	28 (42.4)	93.3% (56.2%–100%)	85.7% (46.2%–100%)
No. with moderate or severe cholera (%) ^a	6 (8.8)	2 (5.7)	4 (12.1)	39 (59.1)	90.3% (61.7%–100%)	79.5% (49.1%–100%)
No. with mild cholera ^b or worse (%)	20 (29.4)	5 (14.3)	15 (45.5)	61 (92.4)	84.5% (67.0%–100%)	50.8% (33.6%–66.8%)
Median diarrheal stool volume, mL (IQR)	0 (0–311)	0 (0–0)	178 (0–988)	4377 (1563–8087)	<.0001	<.0001
Median number of diarrheal stools (IQR)	0 (0–3.8)	0 (0–0)	2.0 (0–5.5)	22.5 (9.8–39.3)	<.0001	<.0001
Median peak <i>Vibrio cholerae</i> O1 count, CFU/mL (IQR)	4.2×10^1 (0– 3.5×10^5)	0 (0– 2.7×10^3)	9.7×10^4 (0– 1.7×10^7)	3.2×10^7 (6.1×10^6 – 1.2×10^8)	<.0001	<.0001
No. with fever (%) ^c	3 (4.4)	1 (2.9)	2 (6.1)	18 (27.3)	.0025	.0159
No. with nausea/vomiting (%)	16 (23.5)	9 (25.7)	7 (21.2)	38 (57.6)	.0032	.0006
No. with abdominal cramping (%)	18 (26.5)	8 (22.9)	10 (30.3)	45 (68.2)	<.0001	.0005
No. with malaise (%)	16 (23.5)	8 (22.9)	8 (24.2)	41 (62.1)	.0003	.0006

Chen WH, CID 2016

Abbreviations: CFU, colony-forming units; CI, confidence interval; IQR, interquartile range.

^a Moderate and severe cholera (study primary endpoint) is defined as ≥ 3 L and ≥ 5 L of cumulative diarrheal stool, respectively.

^b Mild diarrhea is defined as the passage of ≥ 2 unformed stools (grade 3–5) over a 48-hour period that equals or exceeds 200 mL or a single unformed stool of ≥ 300 mL and < 3 L total diarrhea.

^c Fever is defined as $\geq 38.0^\circ\text{C}$ (100.4°F).



Efficacy and immunogenicity of a Vi-tetanus toxoid conjugate vaccine in the prevention of typhoid fever using a controlled human infection model of *Salmonella* Typhi: a randomised controlled, phase 2b trial



Celina Jin, Malick M Gibani, Maria Moore, Helene B Juel, Elizabeth Jones, James Meiring, Victoria Harris, Jonathan Gardner, Anna Nebykova, Simon A Kerridge, Jennifer Hill, Helena Thomaidis-Brears, Christoph J Blohmke, Ly-Mee Yu, Brian Angus, Andrew J Pollard

Lancet 2017; 390: 2472-80
Published Online
September 28, 2017
[http://dx.doi.org/10.1016/S0140-6736\(17\)32333-9](http://dx.doi.org/10.1016/S0140-6736(17)32333-9)

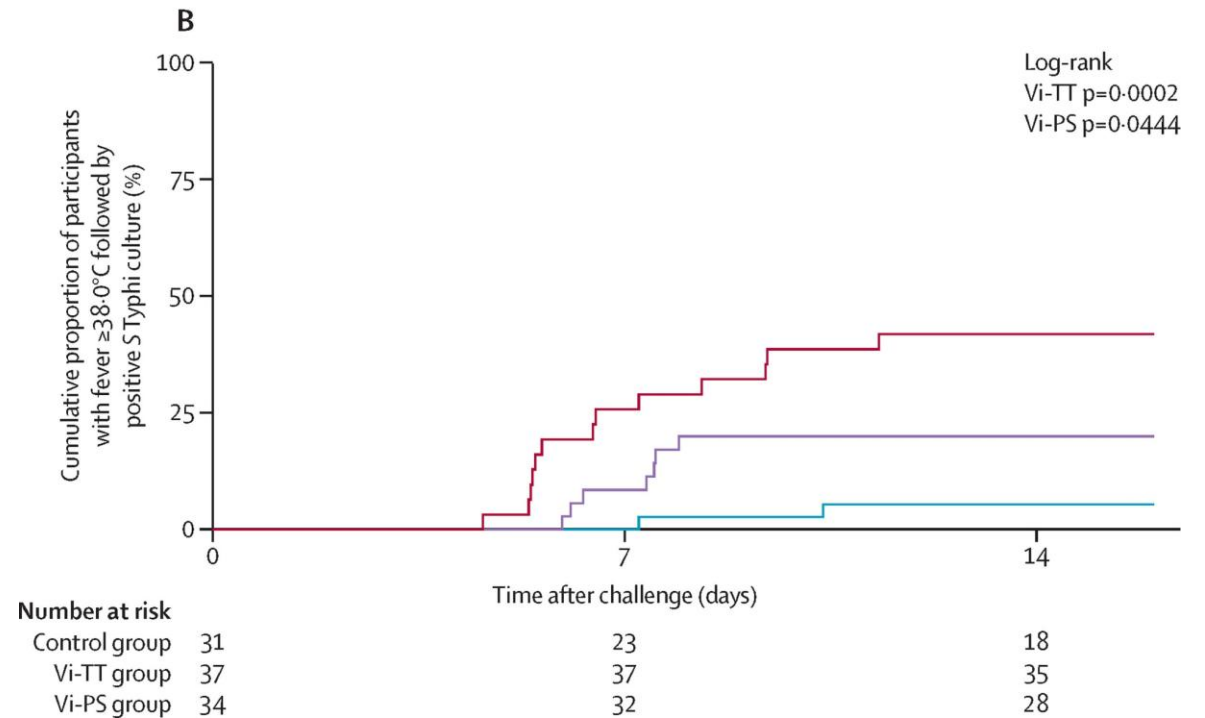
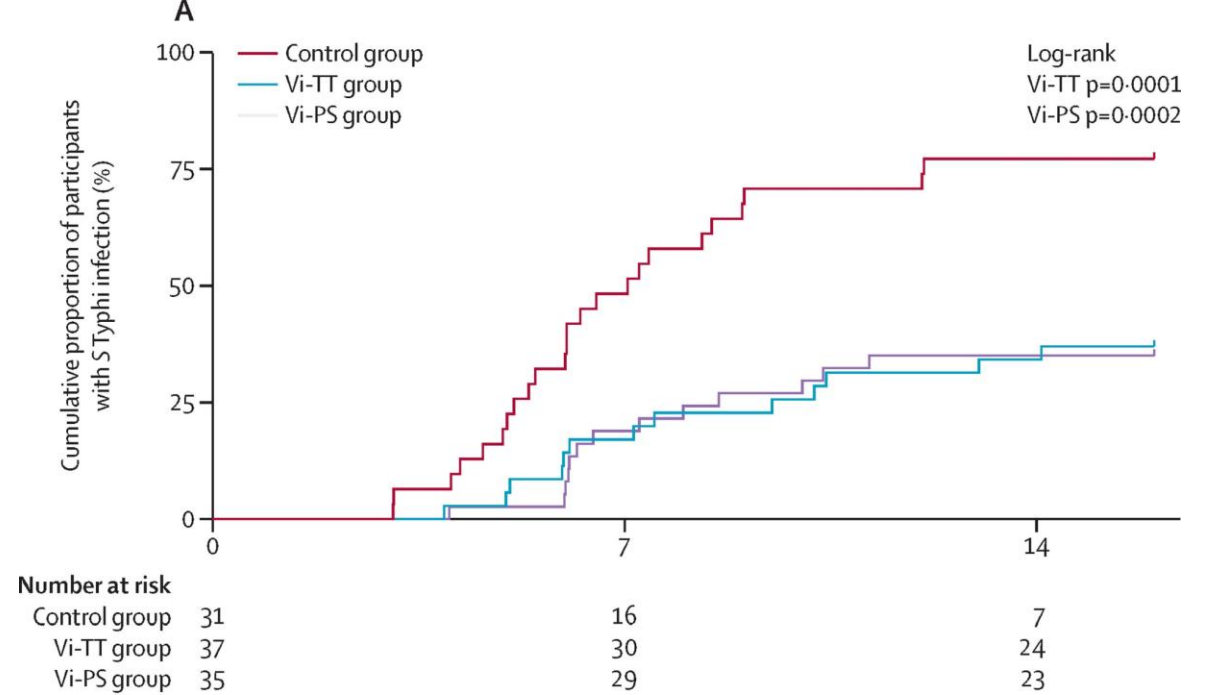
Summary
Background *Salmonella enterica* serovar Typhi (*S* Typhi) is responsible for an estimated 20 million infections and 200 000 deaths each year in resource poor regions of the world. Capsular Vi-polysaccharide-protein conjugate vaccines (Vi-conjugate vaccines) are immunogenic and can be used from infancy but there are no efficacy data for the leading candidate vaccine being considered for widespread use. To address this knowledge gap, we assessed the efficacy of a

Lin et al 2017
Lancet

	Control group (n=34)	Vi-TT group (n=41)	Vi-PS group (n=37)
Primary outcome			
Completed challenged	31	37	35
Total diagnosed (composite definition, clinical or microbiological typhoid diagnosis)	24/31 (77%)	13/37 (35%)	13/35 (37%)
Relative risk (95% CI)	..	0.45 (0.28-0.73)	0.48 (0.30-0.77)
Vaccine efficacy (%; 95% CI)	..	54.6% (26.8-71.8)	52.0% (23.2-70.0)
p value	..	0.0005	0.0010
Secondary outcomes			
Time to diagnosis (days)	6.0 (5.1-7.8)	6.5 (6.1-8.6)	7.2 (5.9-10.2)
Microbiological diagnosis	16/31 (52%)	12/37 (32%)	9/35 (26%)
Time to microbiological diagnosis (days)	6.0 (4.6-8.0)	6.3 (6.0-8.3)	6.1 (5.1-10.2)
Clinical diagnosis	8/31 (26%)	1/37 (3%)	4/35 (11%)
Time to clinical diagnosis (days)	6.8 (5.4-7.8)	10.4	8.5 (6.5-10.0)
Clinical outcomes			
Fever $\geq 37.5^{\circ}\text{C}$ (any duration)	20/31 (65%)	13/37 (35%)	18/35 (51%)
Fever $\geq 38.0^{\circ}\text{C}$ (any duration)	17/31 (55%)	6/37 (16%)	11/35 (31%)
Fever $\geq 38.5^{\circ}\text{C}$ (any duration)	14/31 (45%)	4/37 (11%)	9/35 (25%)
Time to first fever $\geq 38.0^{\circ}\text{C}$ (any duration; days)	7.2 (5.4-8.5)	10.4 (10.2-15.5)	7.5 (6.2-8.7)
Microbiological outcomes			
<i>S</i> Typhi bacteraemia	24/24 (100%)	13/13 (100%)	11/13 (85%)
Time to first positive blood culture (days)	6.1 (5.0-7.6)	6.5 (6.1-8.6)	6.1 (5.0-10.2)
Participants with positive <i>S</i> Typhi stool culture	22/31 (71%)	22/37 (59%)	21/35 (60%)
Diagnosed participants with positive <i>S</i> Typhi stool culture	19/24 (79%)	12/13 (92%)	10/13 (77%)
Median quantitative blood culture (CFU/mL; range)	0.4 (0.05-22.7)	0.075 (0.05-1.2)	0.1 (0.05-5.6)

Data are n, n/N (%), or median (IQR) unless otherwise stated. Vi-TT=Vi-tetanus toxoid conjugate vaccine. Vi-PS=Vi-polysaccharide vaccine. *S* Typhi=*Salmonella* Typhi. CFU=colony forming units.

Table 2: Primary and secondary outcomes

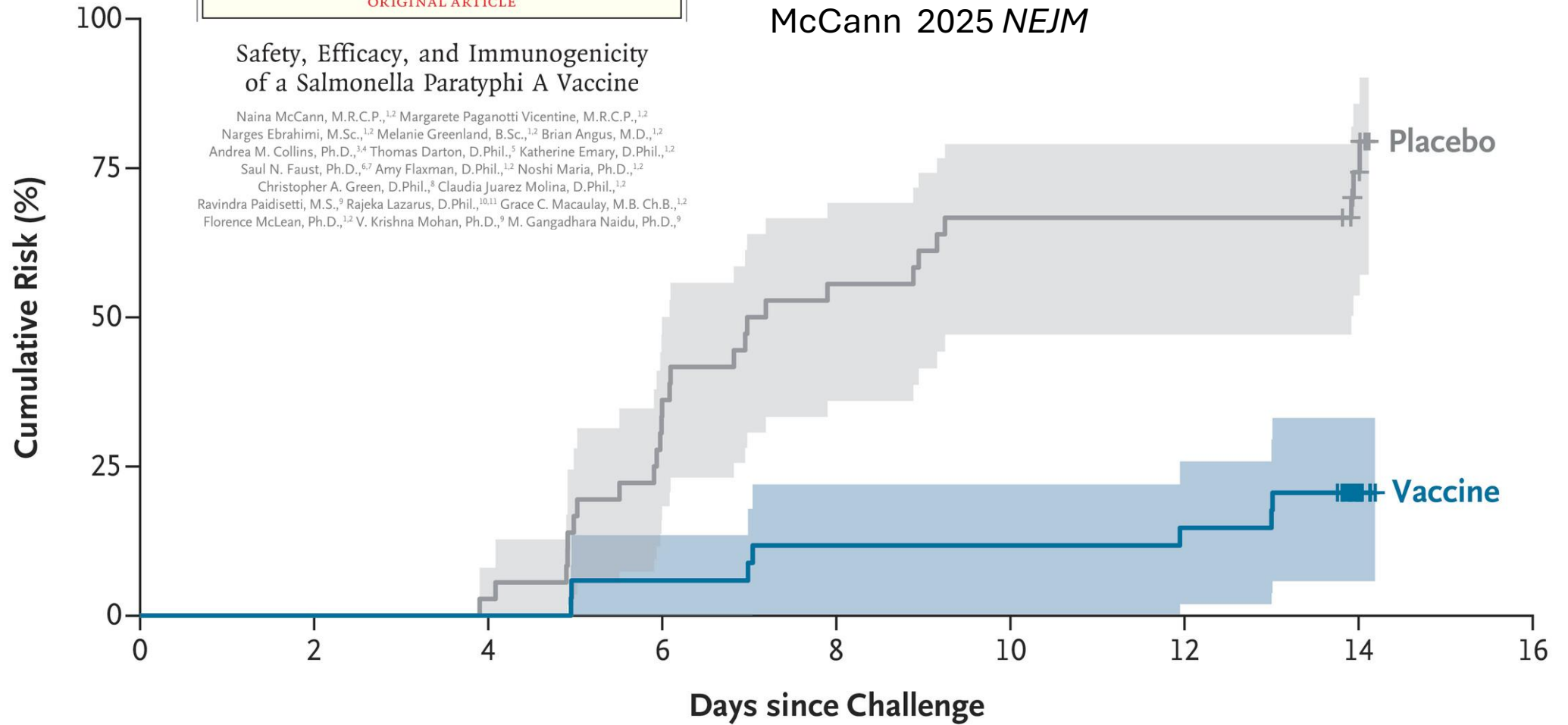


ORIGINAL ARTICLE

McCann 2025 *NEJM*

Safety, Efficacy, and Immunogenicity of a Salmonella Paratyphi A Vaccine

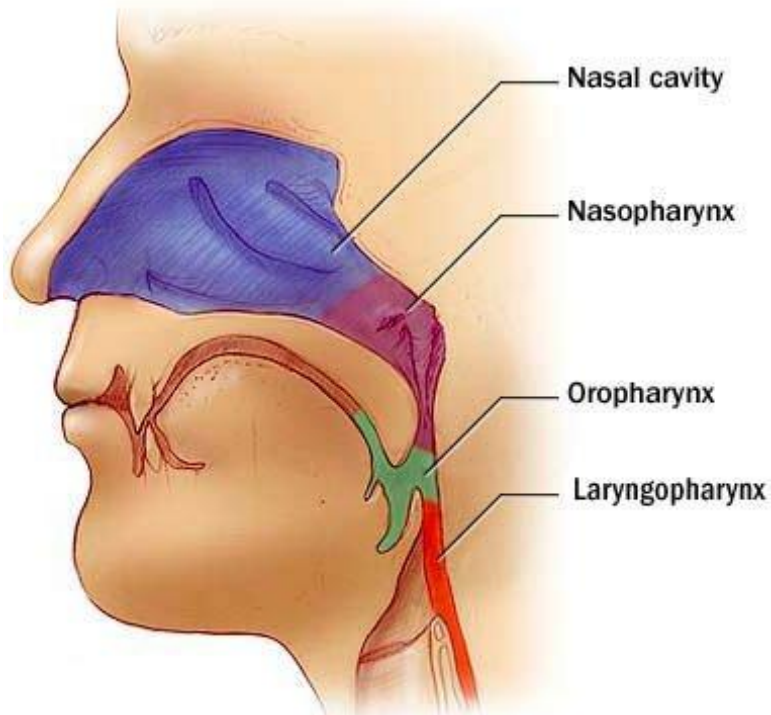
Naina McCann, M.R.C.P.,^{1,2} Margarete Paganotti Vicentine, M.R.C.P.,^{1,2}
 Narges Ebrahimi, M.Sc.,^{1,2} Melanie Greenland, B.Sc.,^{1,2} Brian Angus, M.D.,^{1,2}
 Andrea M. Collins, Ph.D.,^{3,4} Thomas Darton, D.Phil.,⁵ Katherine Emary, D.Phil.,^{1,2}
 Saul N. Faust, Ph.D.,^{6,7} Amy Flaxman, D.Phil.,^{1,2} Noshi Maria, Ph.D.,^{1,2}
 Christopher A. Green, D.Phil.,⁸ Claudia Juarez Molina, D.Phil.,^{1,2}
 Ravindra Paidiseti, M.S.,⁹ Rajeka Lazarus, D.Phil.,^{10,11} Grace C. Macaulay, M.B. Ch.B.,^{1,2}
 Florence McLean, Ph.D.,^{1,2} V. Krishna Mohan, Ph.D.,⁹ M. Gangadhara Naidu, Ph.D.,⁹



No. at Risk

Placebo	36	36	35	23	16	12	12	6	0
Vaccine	34	34	34	32	30	30	29	8	0

The nasopharynx...the root of all....



- Pneumonia
- Otitis Media
- Sinusitis
- Pharyngitis
- Chronic Bronchial Sepsis
- Meningitis
- Staphylococcal sepsis

Haemophilus influenzae
Streptococcus pneumoniae
Bordetella pertussis
Moraxella catarrhalis

Neisseria meningitidis
Streptococcus pyogenes
Staphylococcus aureus
Streptococcus agalactiae

Mycobacterium spp.
Fusobacterium spp.
Klebsiella spp
Corynebacterium spp.

Experimental inoculation of *S.pneumoniae* 6B following 13 valent PCV

Collins et al *AJRCCM* 2015

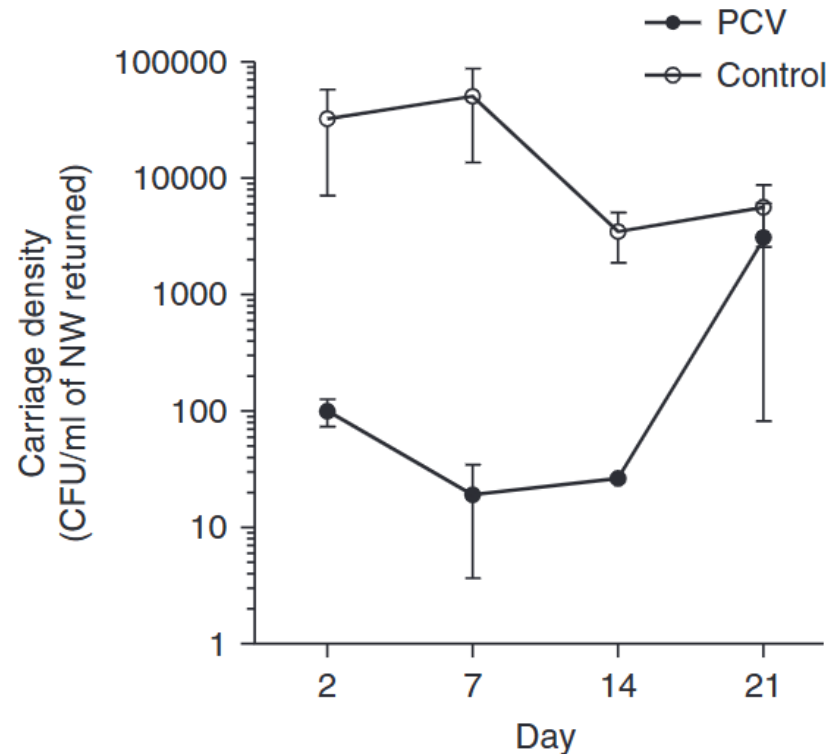


Table 3. 6B Colonization Density According to Vaccination Group at Each Time Point

Day	Density (CFU/ml) Geometric Mean \pm SD		Ratio (95% CI)	P Value
	PCV (No. Colonized at Each Time Point)	Control (No. Colonized at Each Time Point)		
Day 2	99 \pm 53 (4)	33,694 \pm 120,812 (21)	0.17 (0.04–0.65)	0.0099
Day 7	19 \pm 31 (4)	50,517 \pm 169,172 (21)	0.02 (0.00–0.14)	<0.0001
Day 14	26* (1)	3,476 \pm 6,962 (19)	1.94 (0.09–43.90) [†]	0.6767 [†]
Day 21	3,085 \pm 4,247 (2)	5,623 \pm 11,855 (15)		

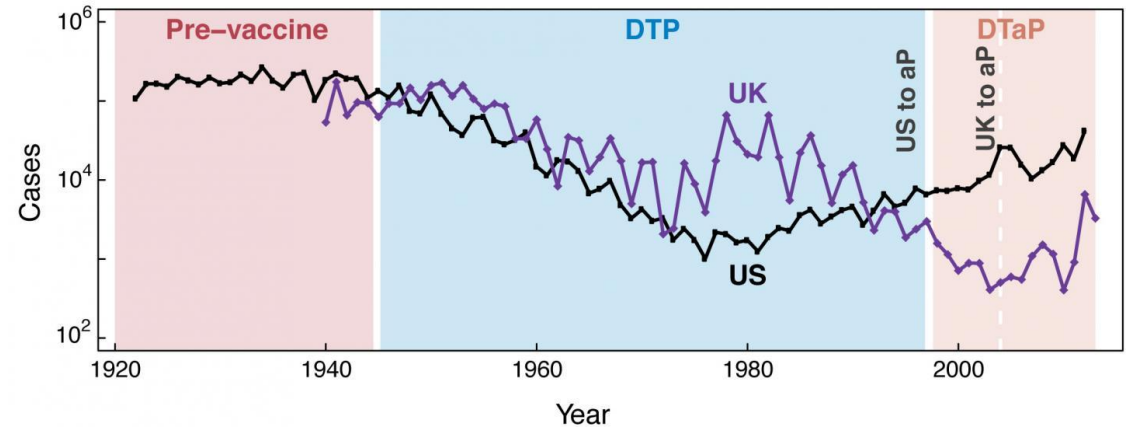
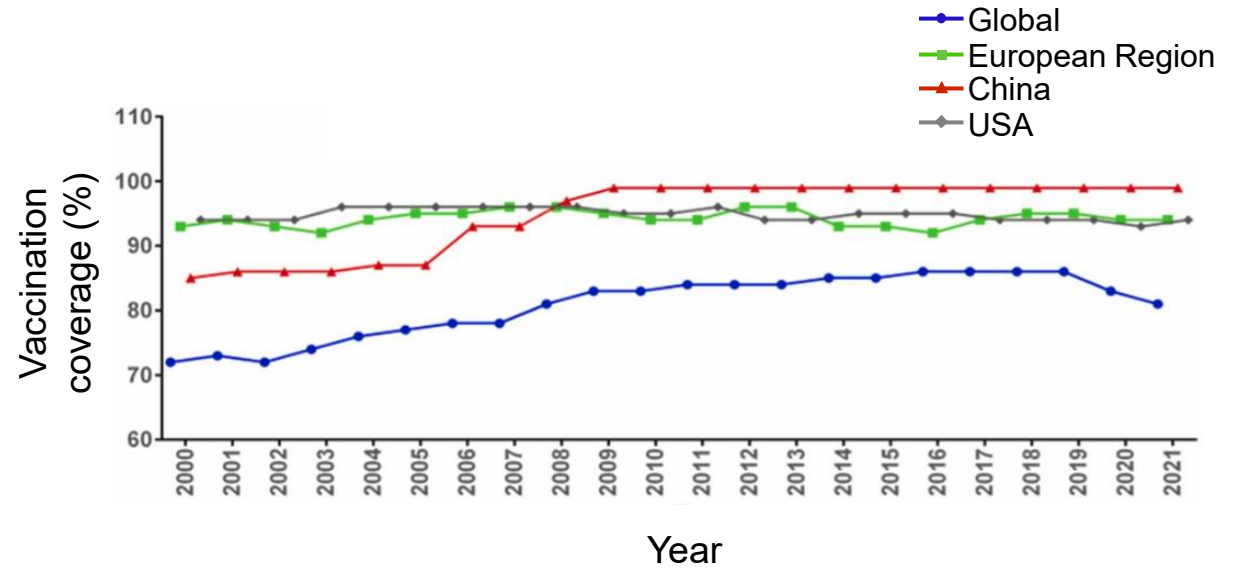
Definition of abbreviations: CI = confidence interval; PCV = pneumococcal conjugate vaccine. Density is only reported in participants who are colonized with 6B.

*Only one observation, and SD is unavailable.

[†]Days 14 and 21 were combined to generate a stable estimate.

Pertussis

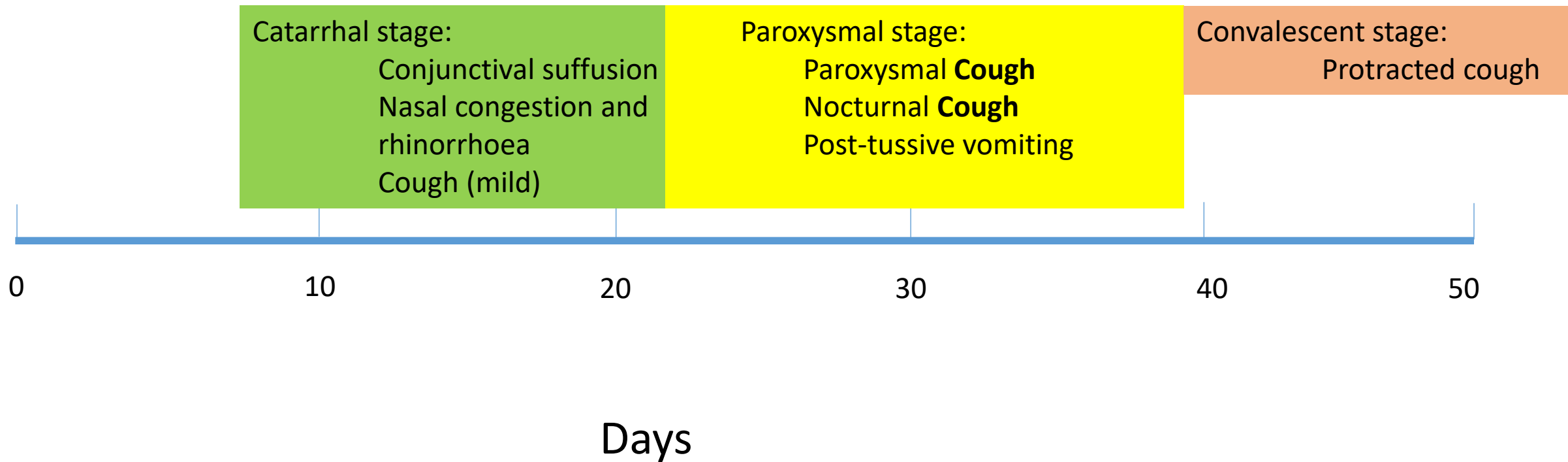
- Resurgence of Pertussis disease despite high vaccine coverage.
- ~ 24.1 million cases and ~ 160,700 deaths annually.
- Occurring despite high vaccine coverage.



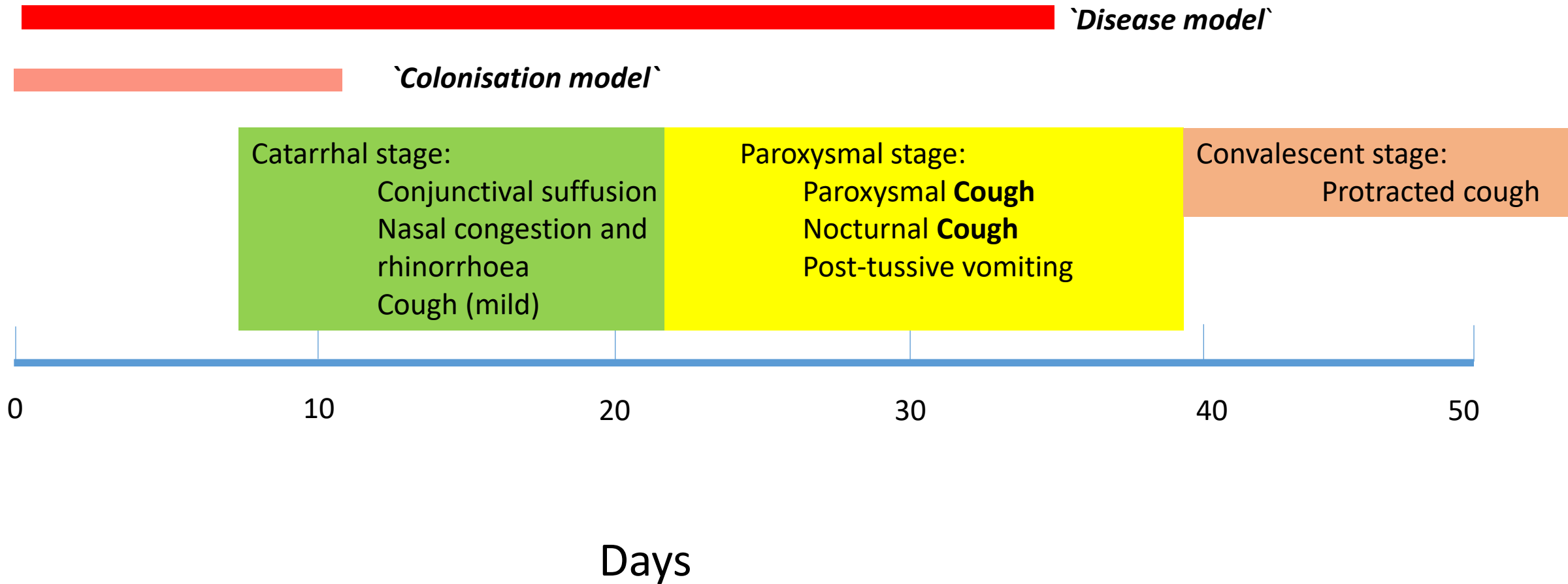
Althouse and Scarpino, *BMC Med*, 2015.
 Yeung et al, *Lancet Infect Dis*, 2017.
 Liu et al, *J Infect*, 2024.



Clinical Pertussis



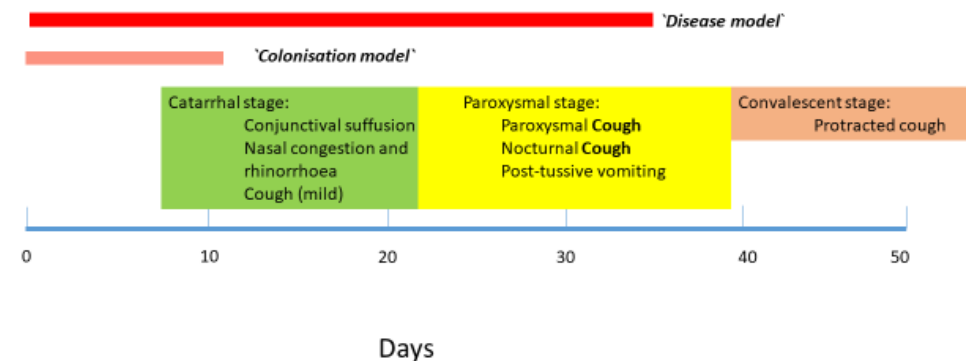
Controlled Human Infection



Disease model

ADVANTAGES	DISADVANTAGES
Clear Clinical and Regulatory Relevance	
Paroxysmal cough is easily detected and measured	
Correlate micro/immunology with clinical endpoints	
Study the full/mature disease process	
Interventions measured against clinical endpoints	

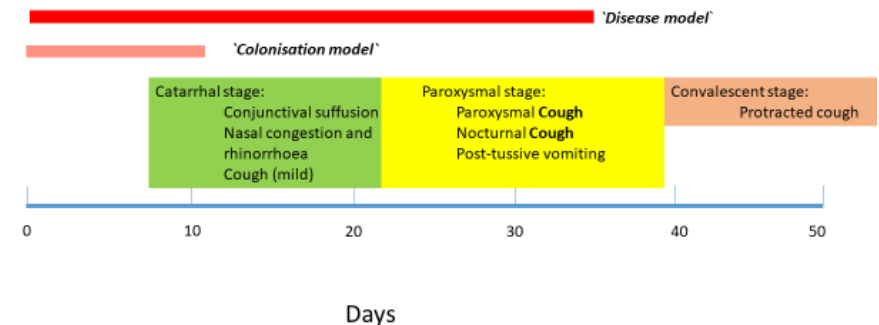
Controlled Human Infection



Disease model

ADVANTAGES	DISADVANTAGES
Clear Clinical and Regulatory Relevance	Unable to quantify risk
Paroxysmal cough is easily detected and measured	Unable to reverse disease after paroxysmal cough onset
Correlate micro/immunology with clinical endpoints	Secondary cases
Study the full/mature disease process	Risk to nursing staff
Interventions measured against clinical endpoints	Reputational risk

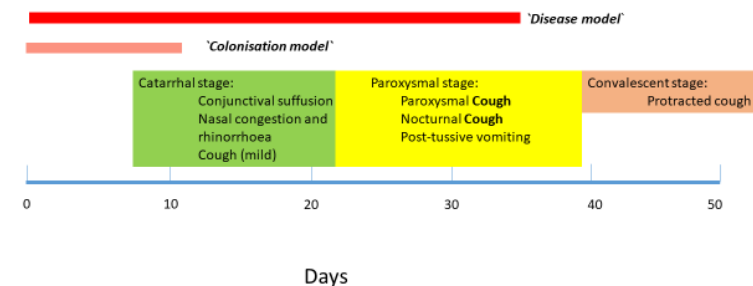
Controlled Human Infection



Colonisation model

ADVANTAGES	DISADVANTAGES
Safe	
Detection of bacteria is an objective endpoint	
Probably less infection control issues	
May be the commonest manifestation of wild infection	
Colonisation is pre-requisite to disease so Vaccine/Challenge studies would inform herd protection estimates	

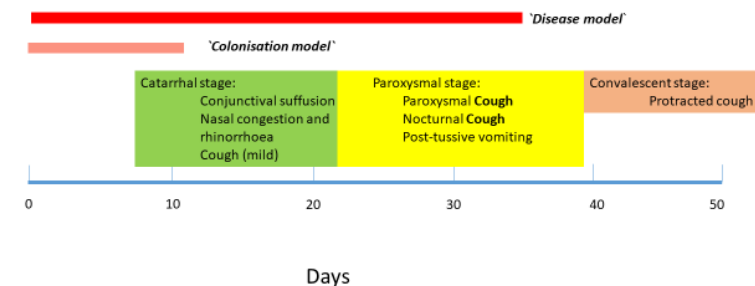
Controlled Human Infection



Colonisation model

ADVANTAGES	DISADVANTAGES
Safe	<i>Bordetella pertussis</i> is difficult to detect in asymptomatic people
Detection of bacteria is an objective endpoint	The site of colonisation is unknown
Probably less infection control issues	Duration of carriage is unknown
May be the commonest manifestation of wild infection	Detection of <i>B.pertussis</i> in an asymptomatic person may not imply active biological interaction
Colonisation is pre-requisite to disease so Vaccine/Challenge studies would inform herd protection estimates	Different carriage states – density??

Controlled Human Infection



University of Southampton

Bp Controlled Human Infection Model



- *Primum non nocere*
- Asymptomatic nasopharyngeal colonisation
- *Bordetella pertussis* strain B1917
- Clearance with azithromycin

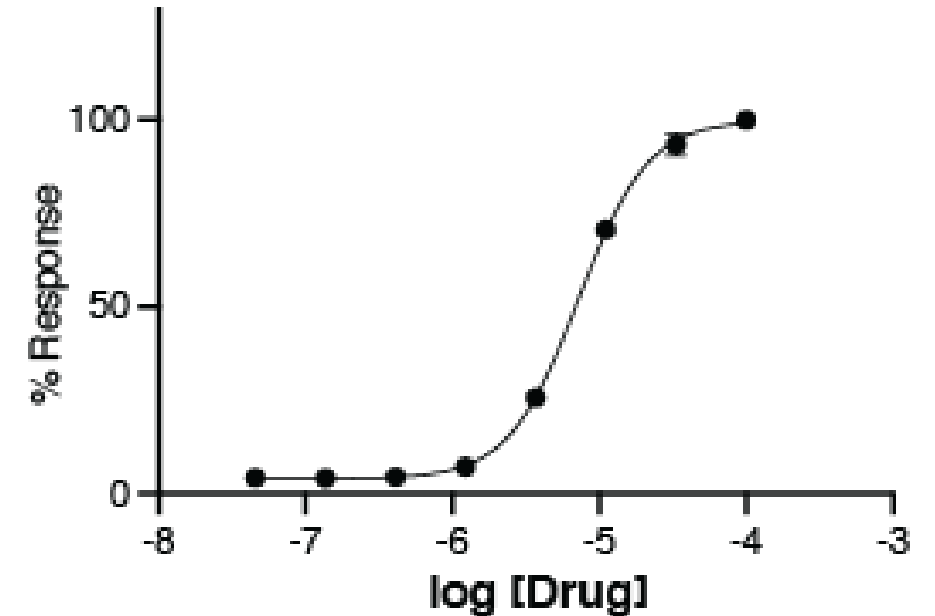
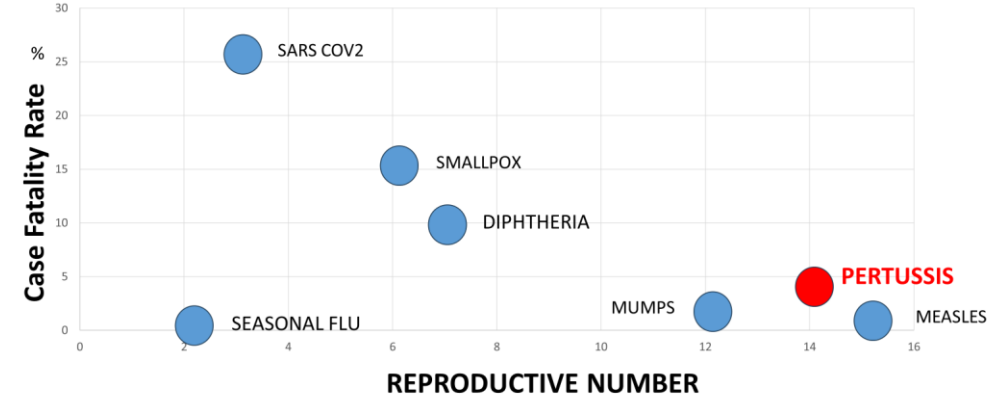
- Potential utilities:
 - Biomarkers associated with protection
 - Platform for vaccine testing



Dose and Administration

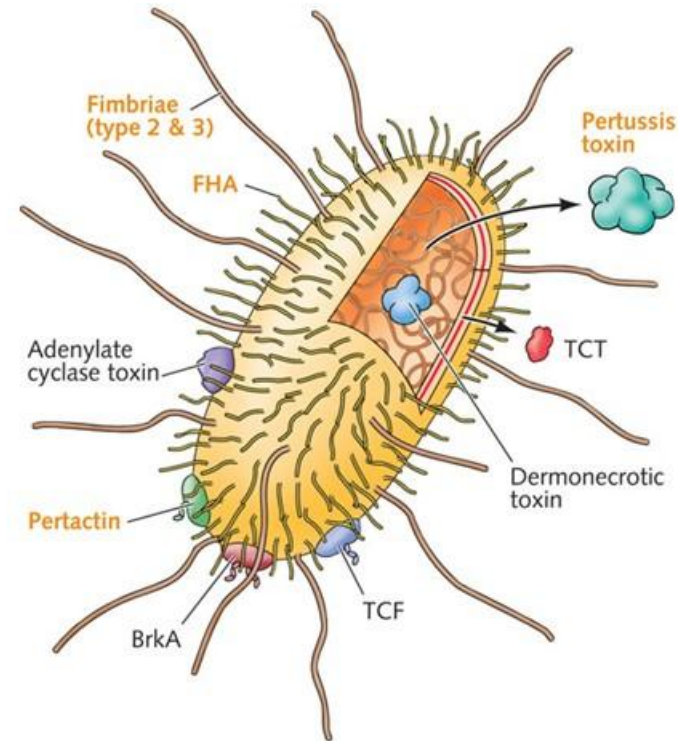
- *B.pertussis* is highly infectious
- Human-adapted infection
- Natural infection requires $<10\text{cfu}$
- Aerosol transmission

Reproductive Number/ Case Fatality Rate



Selected organism

- GMP manufactured by Q Biologicals, Belgium
- B1917, characterised by *ptxP3-ptxA1-prn2-fim3-2, fim2-1* MLVA27, PFGE BpSR11
- Expresses
 - Pertactin
 - Pertussis Toxin
 - Filamentous haemagglutinin
- Representative of current isolates in Europe

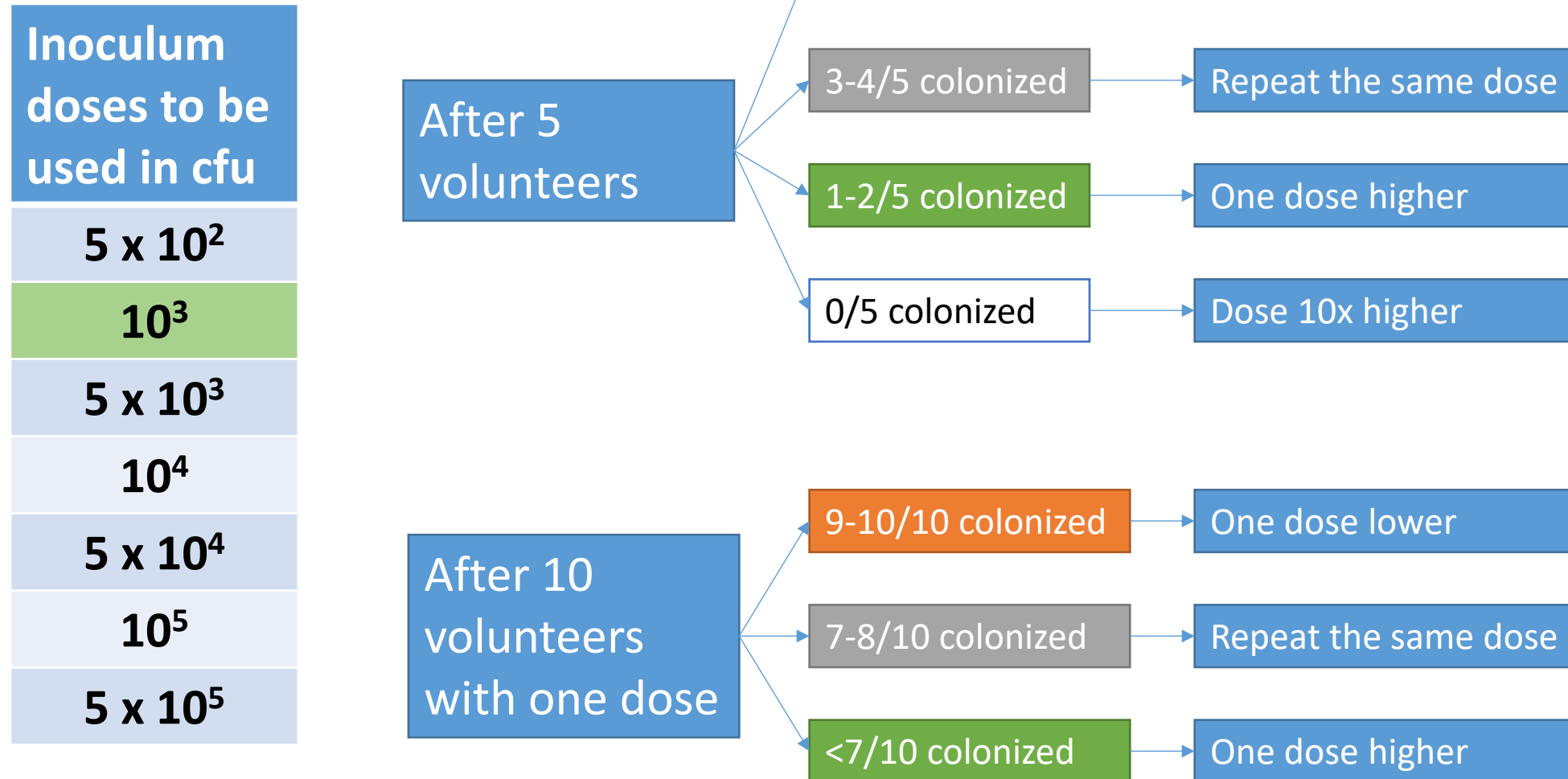


PERISCOPE Phase A

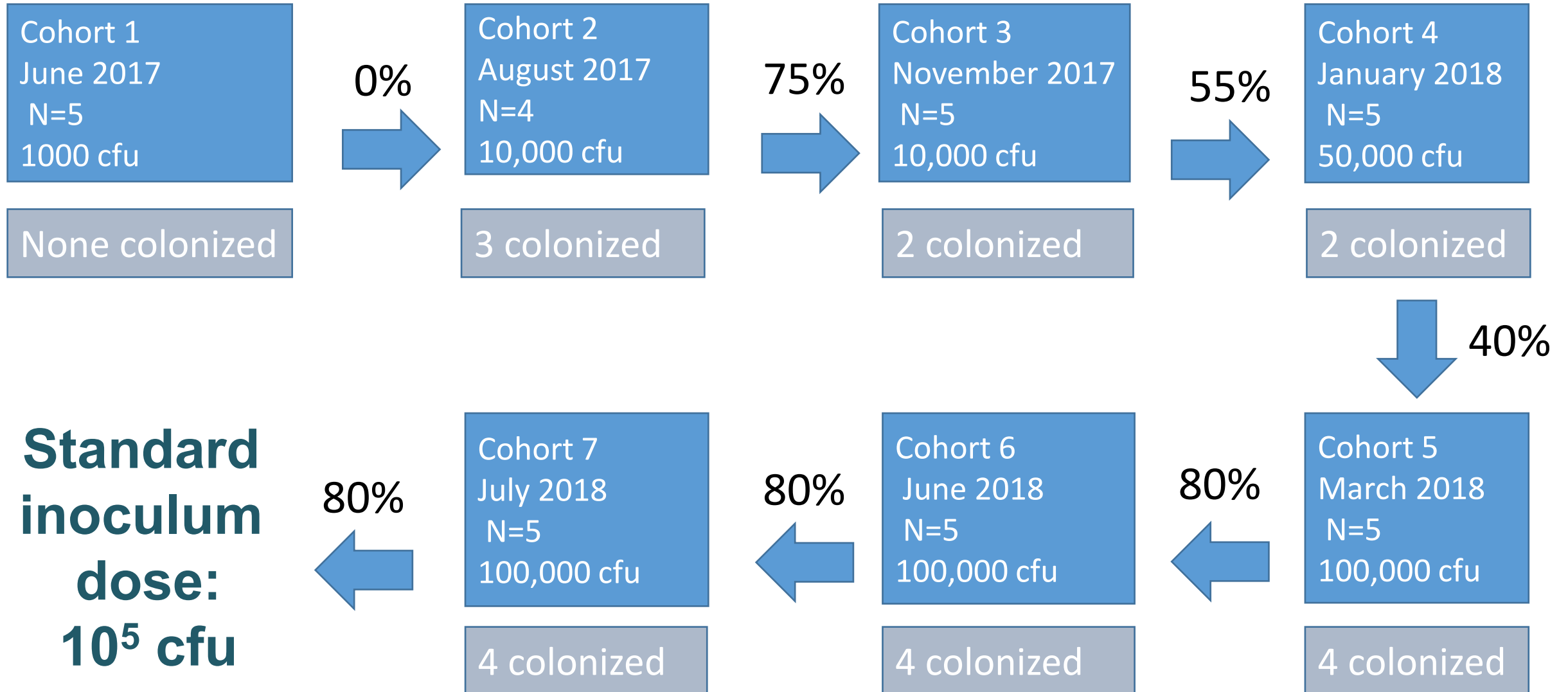


- Dose ranging study : selected for anti-PT IgG < 20 IU/mL
- Inpatient model : 16-day admission period
- Completed in 2019, n=34

To determine the dose of the standard inoculum: Dose escalation



Time line dose escalation





Phase A – main findings

- Standard inoculum dose 10^5 CFU
- Colonisation fraction 0.8 (12 of 15 participants colonised)
- Safe
- Azithromycin effective in clearing colonisation
- No shedding detected
- Seroconversion in some colonised participants

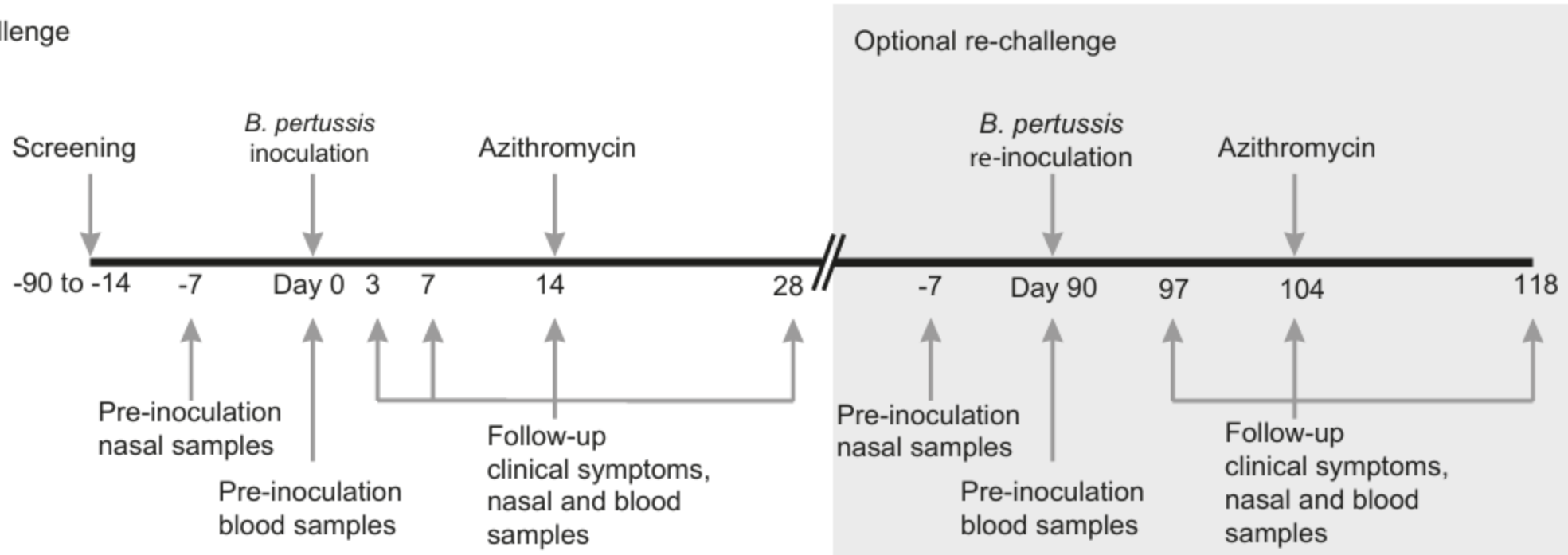
De Graaf H. et al *Clinical Infectious Disease* 2000

Diagnostics – Phase A



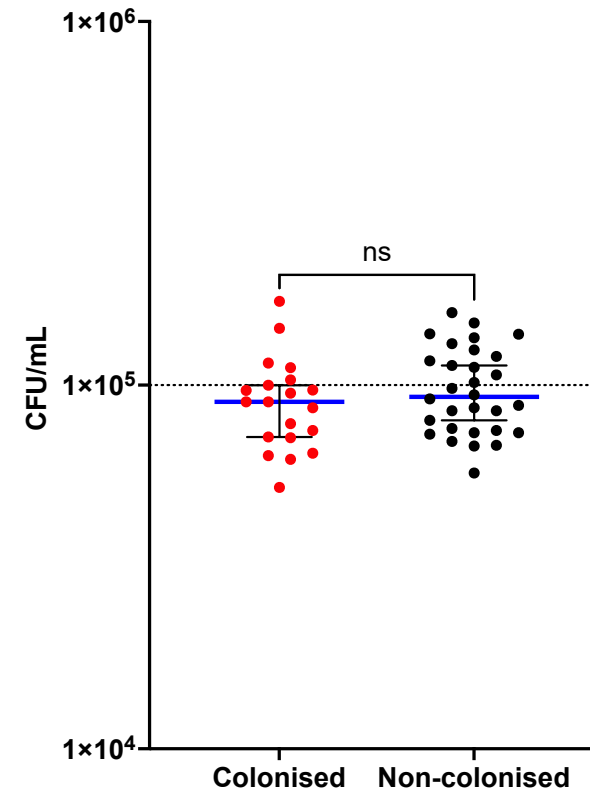
- Nasal wash most sensitive sampling technique
- Nasal wash culture and PCR equivalent
- Pernasal swab- culture 36% sensitive versus PCR 77%
- PCR of throat swabs detected 36% of all PCR positive samples (for pernasal swab this was 54%, and nasal wash 94%)

Challenge

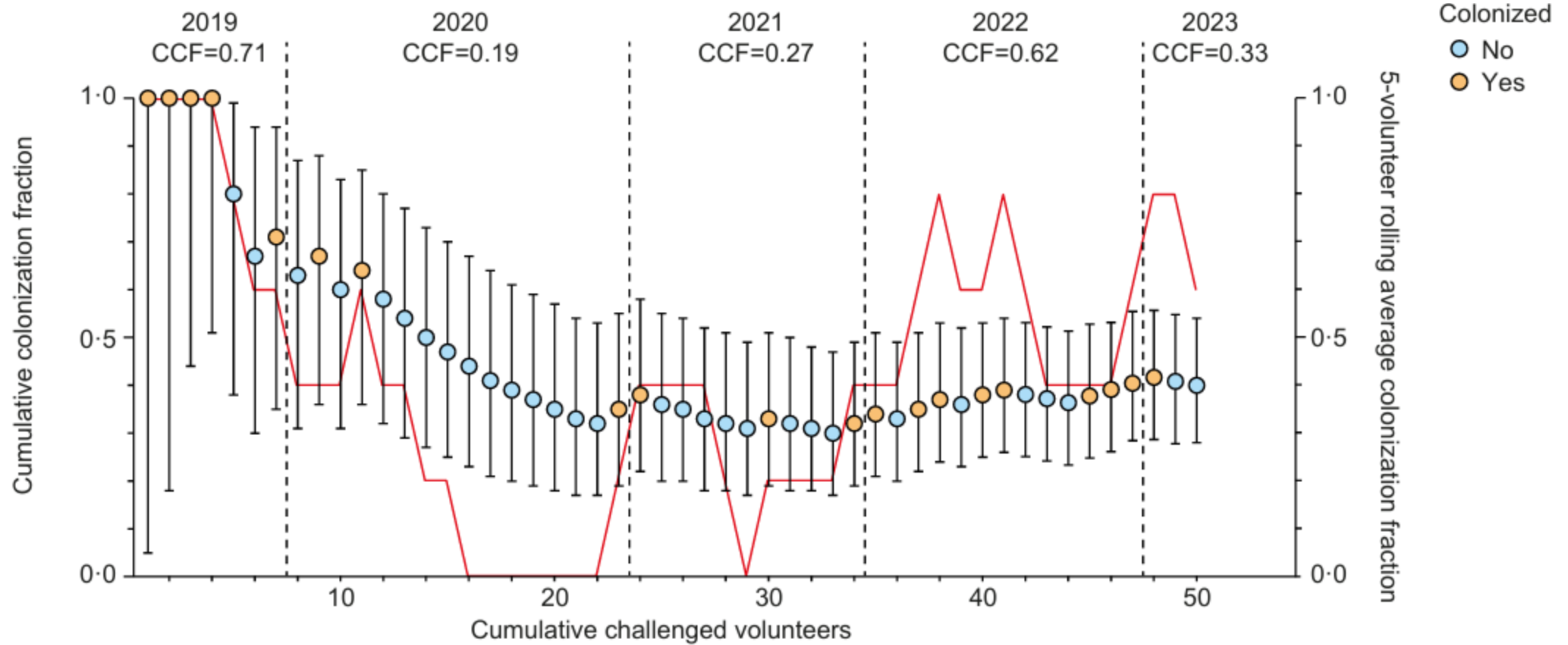


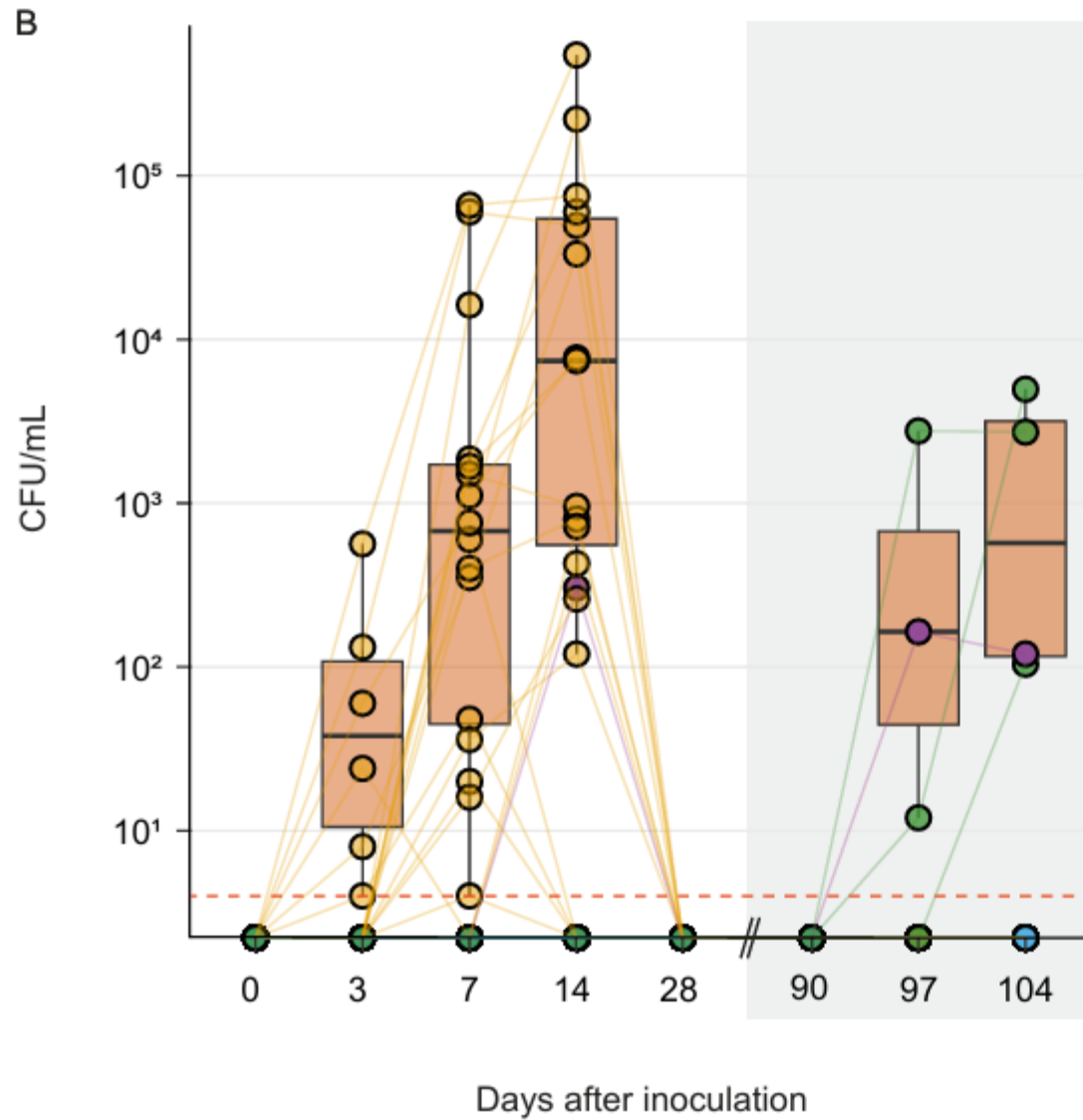
Primary challenge: Colonisation

- **20 of 50** Challenge volunteers colonised
- Colonisation fraction = 0.4
- No difference in inoculum
- 14 contact volunteers enrolled
 - 6 corresponding Challenge volunteer colonised
 - No transmission detected

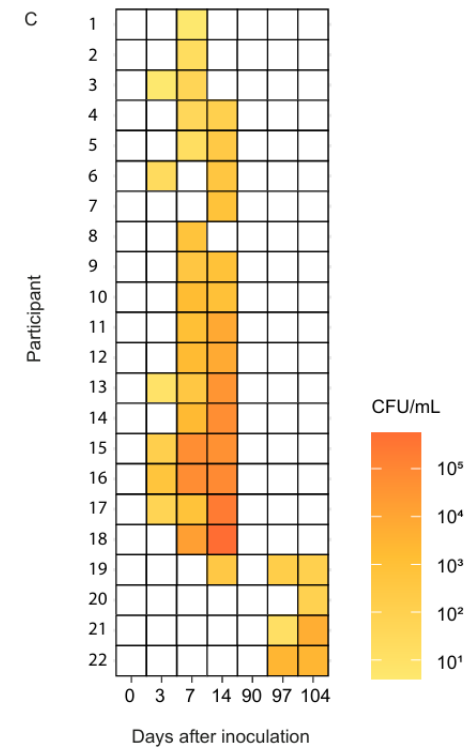


A



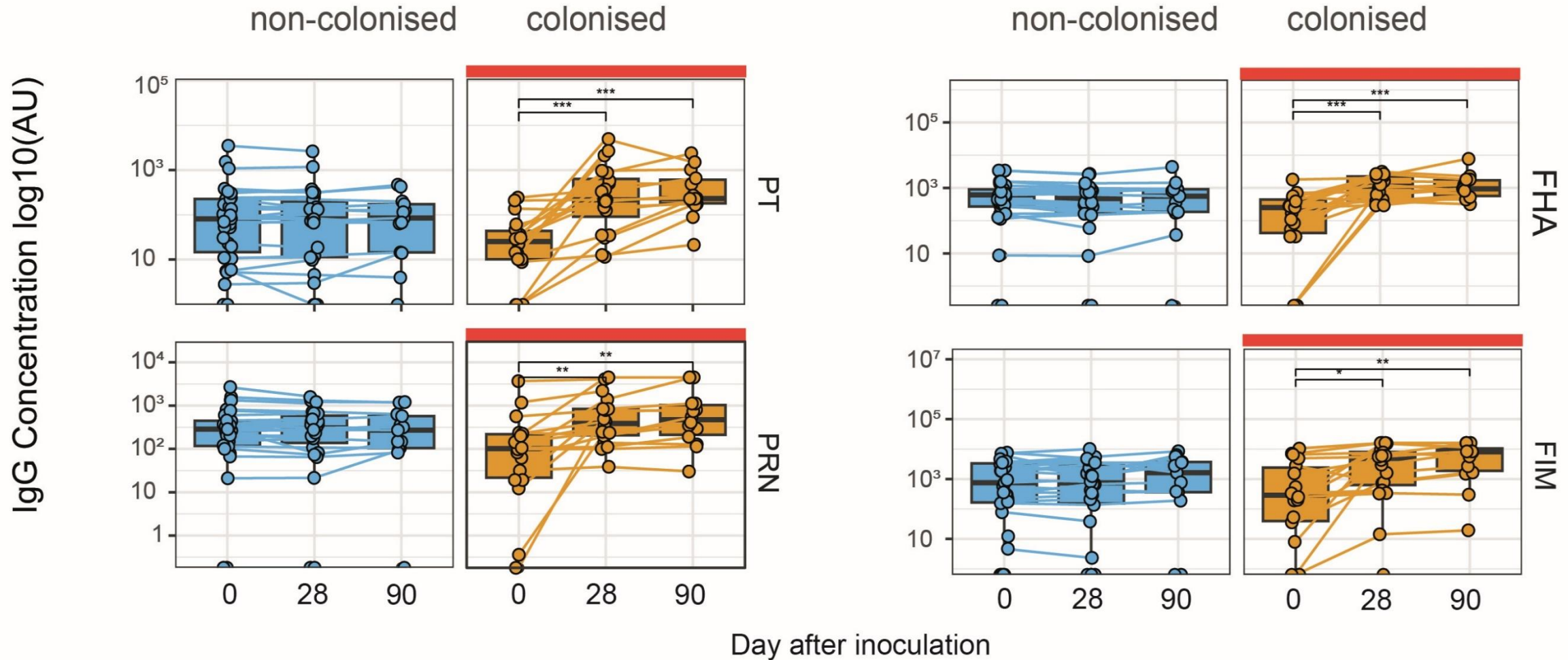


First challenge	Re-challenge
Colonized	Colonized
Colonized	Non-colonized
Non-colonized	Colonized
Non-colonized	Non-colonized

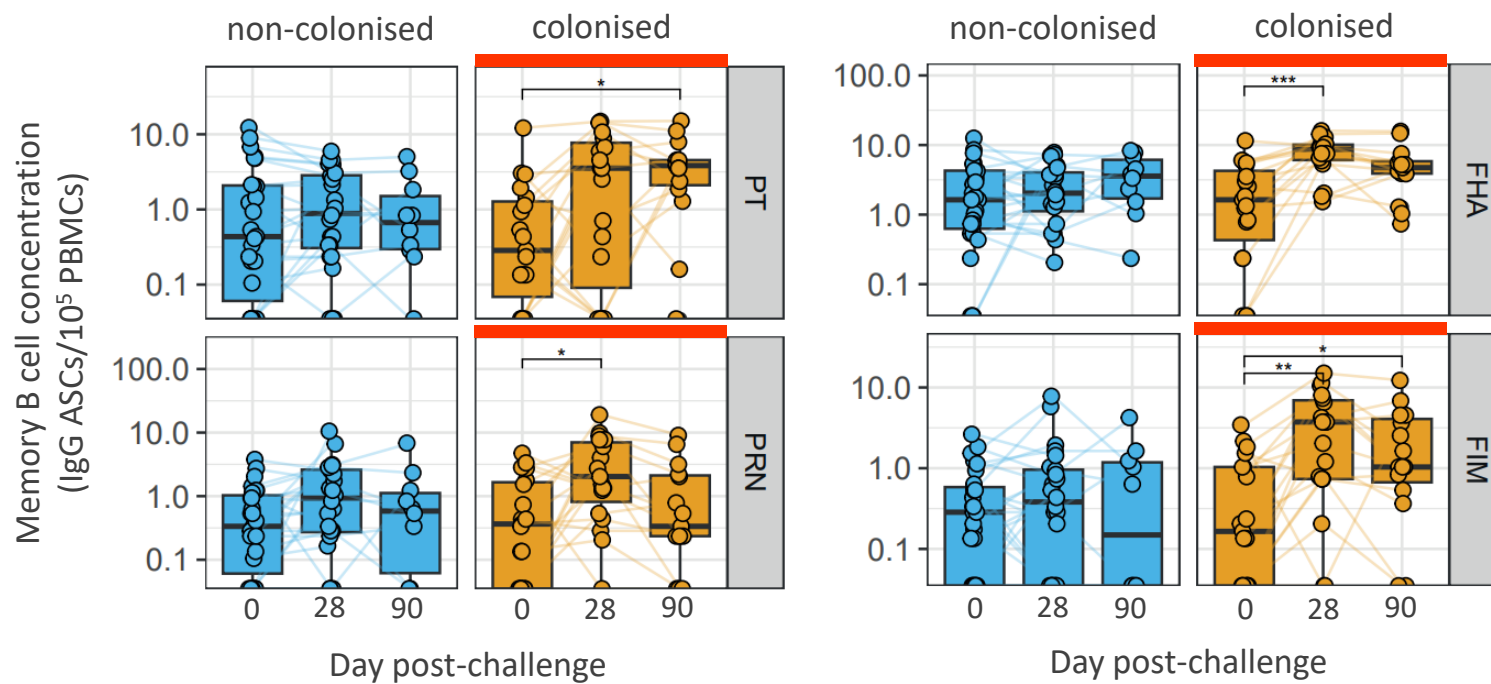


Does experimental infection induce an immune response?

Colonisation is an immunising event - IgG



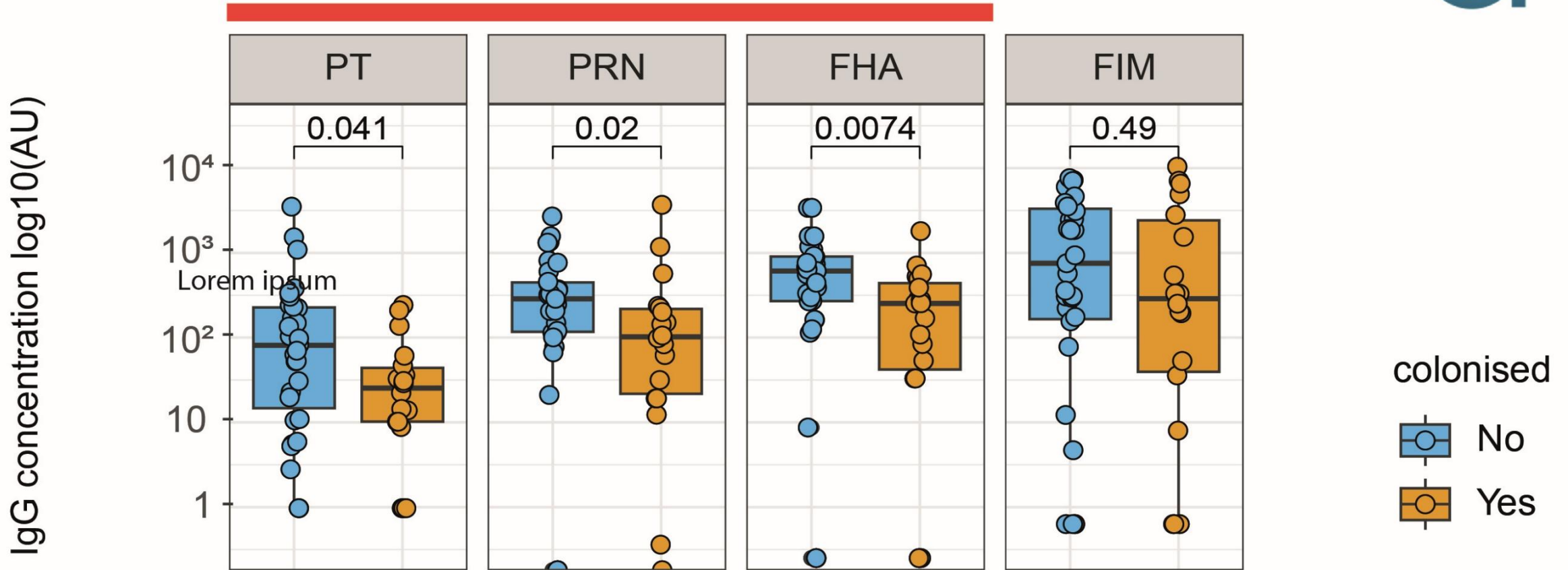
Colonisation induces Bp-specific memory B cell responses



Non-colonised versus colonised – serum IgG



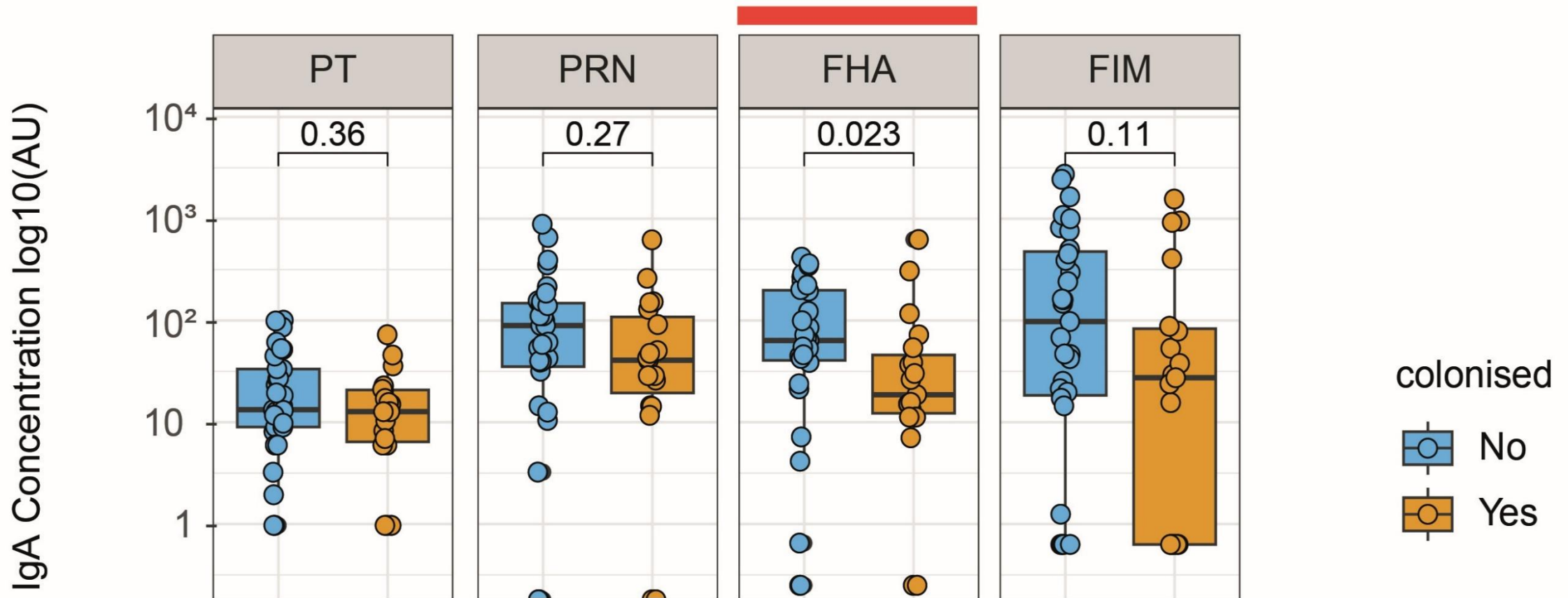
Baseline



Non-colonised versus colonised – serum IgA



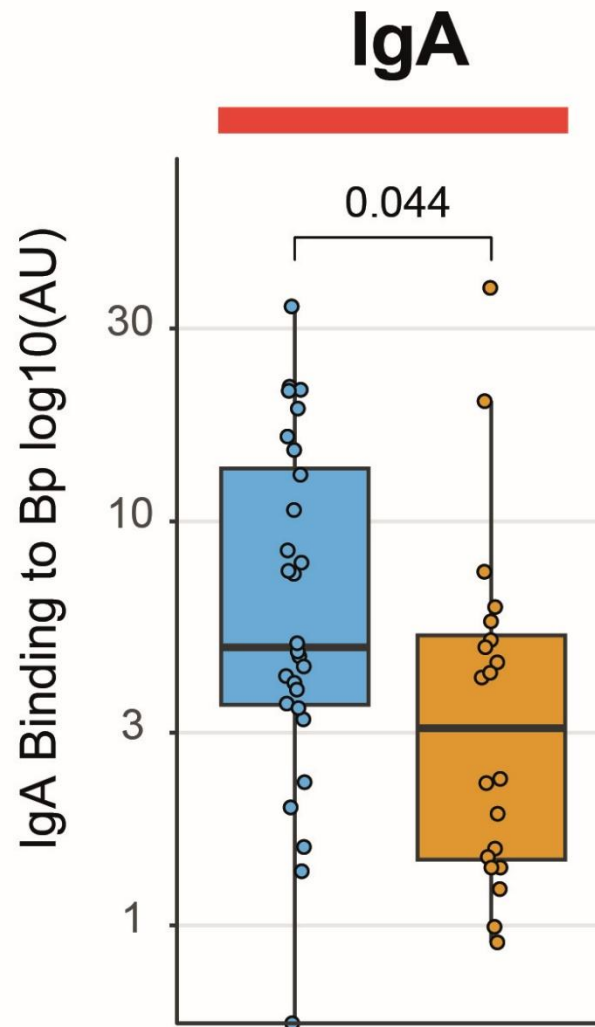
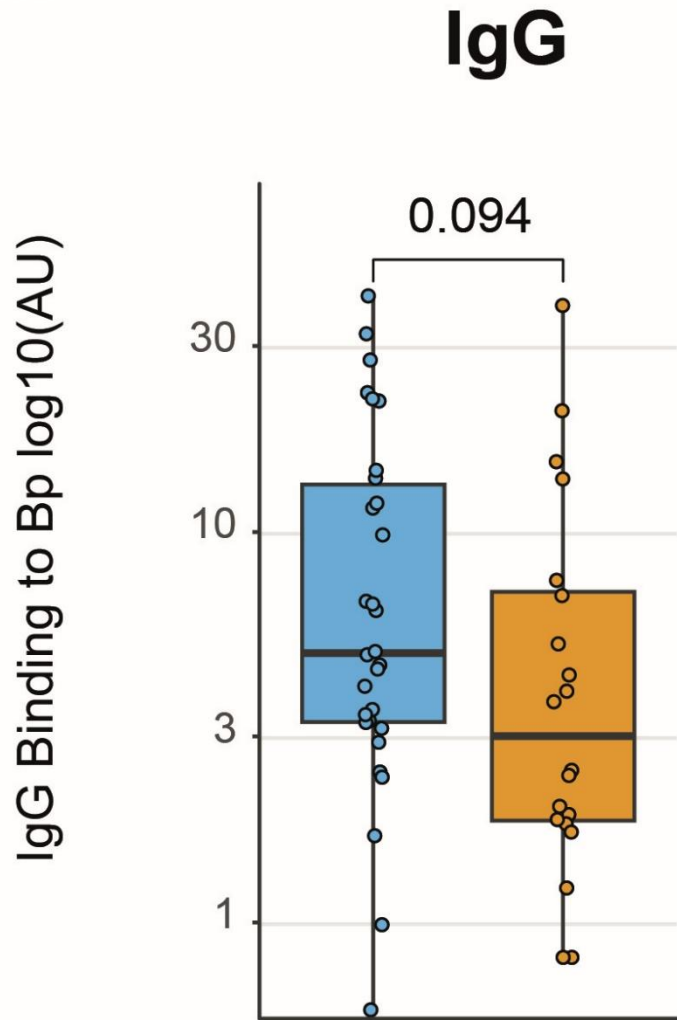
Baseline



Non-colonised versus colonised – serum Bp binding



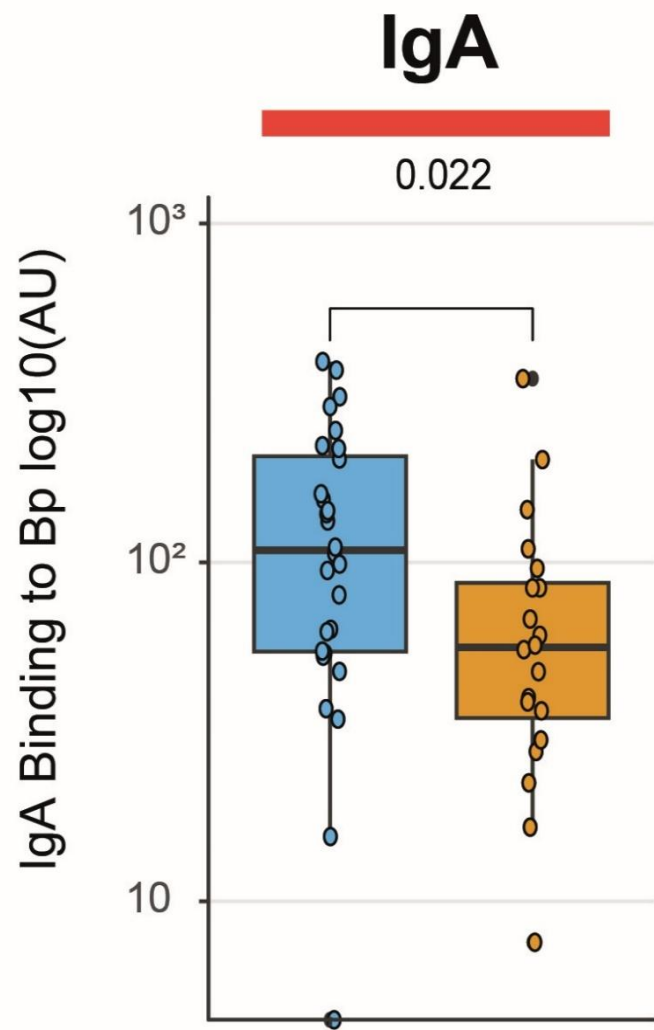
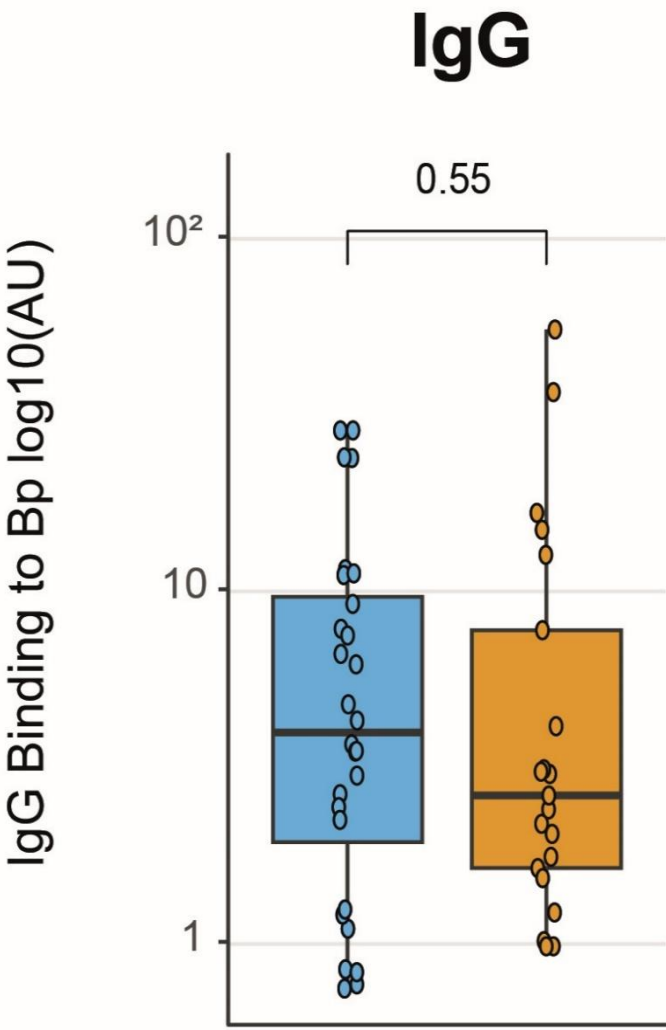
Baseline



colonised

- No
- Yes

Non-colonised versus colonised — mucosal fluid Bp binding

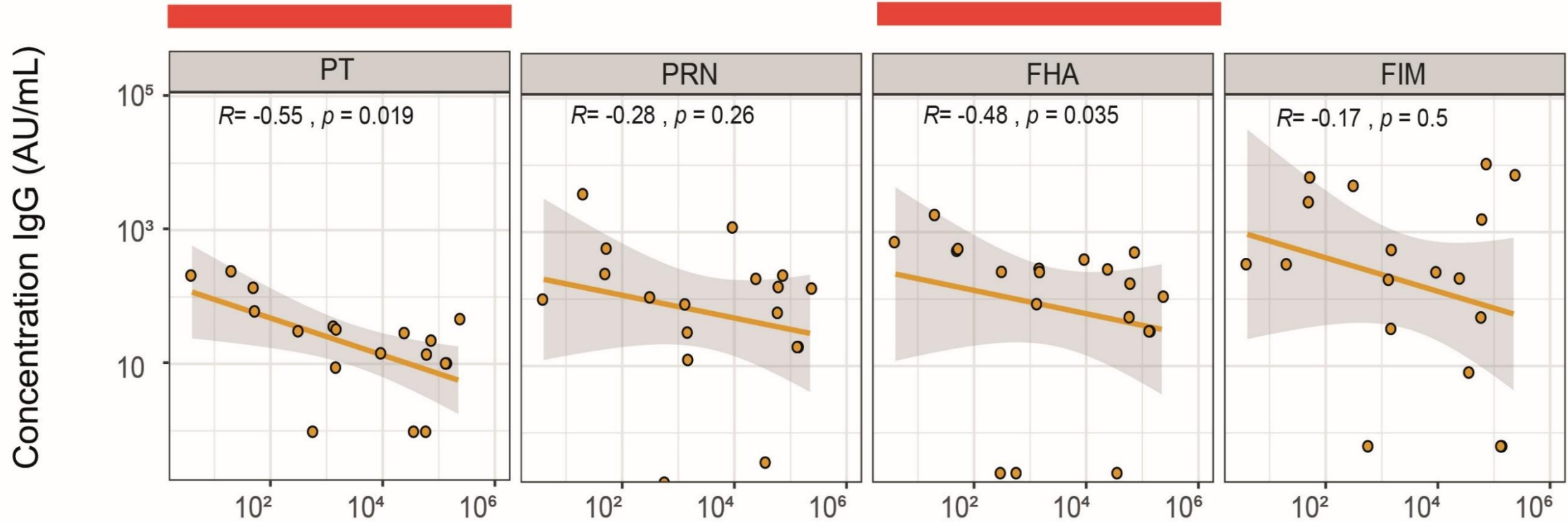


colonised

- No
- Yes

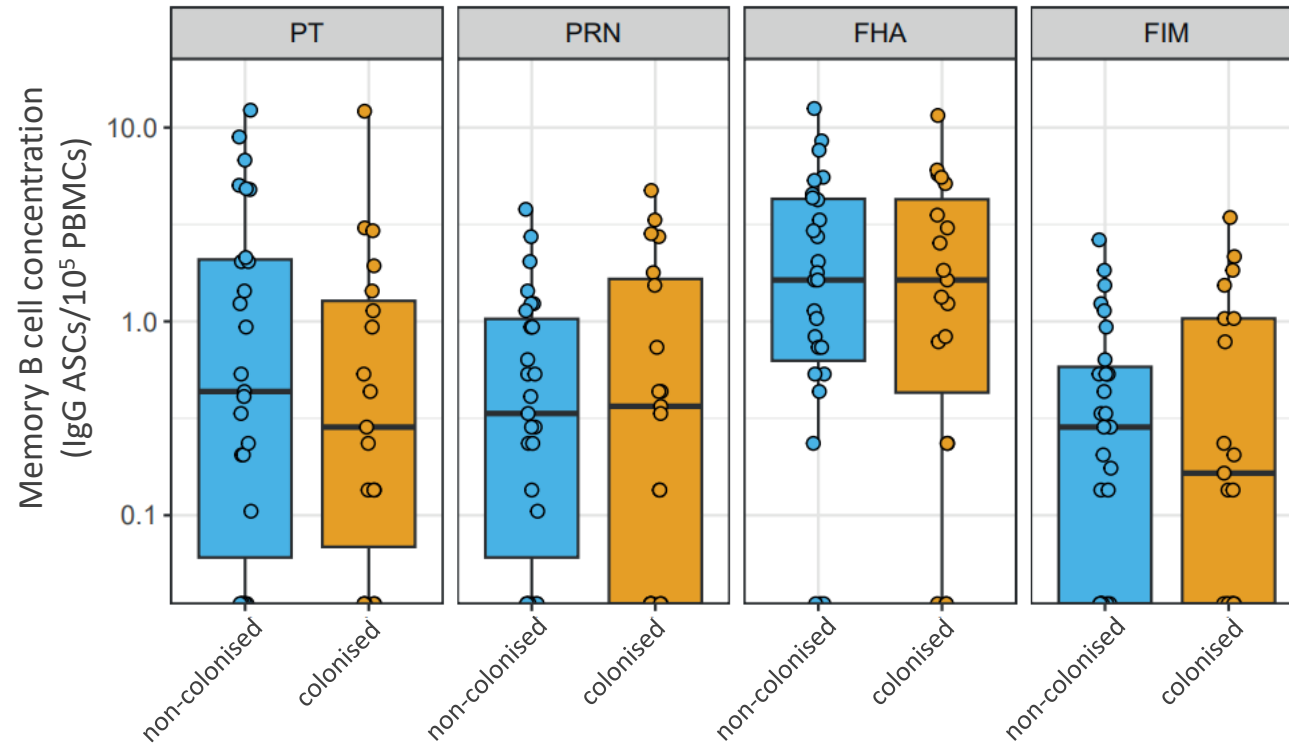
Control of colonisation density - Serum IgG

Baseline

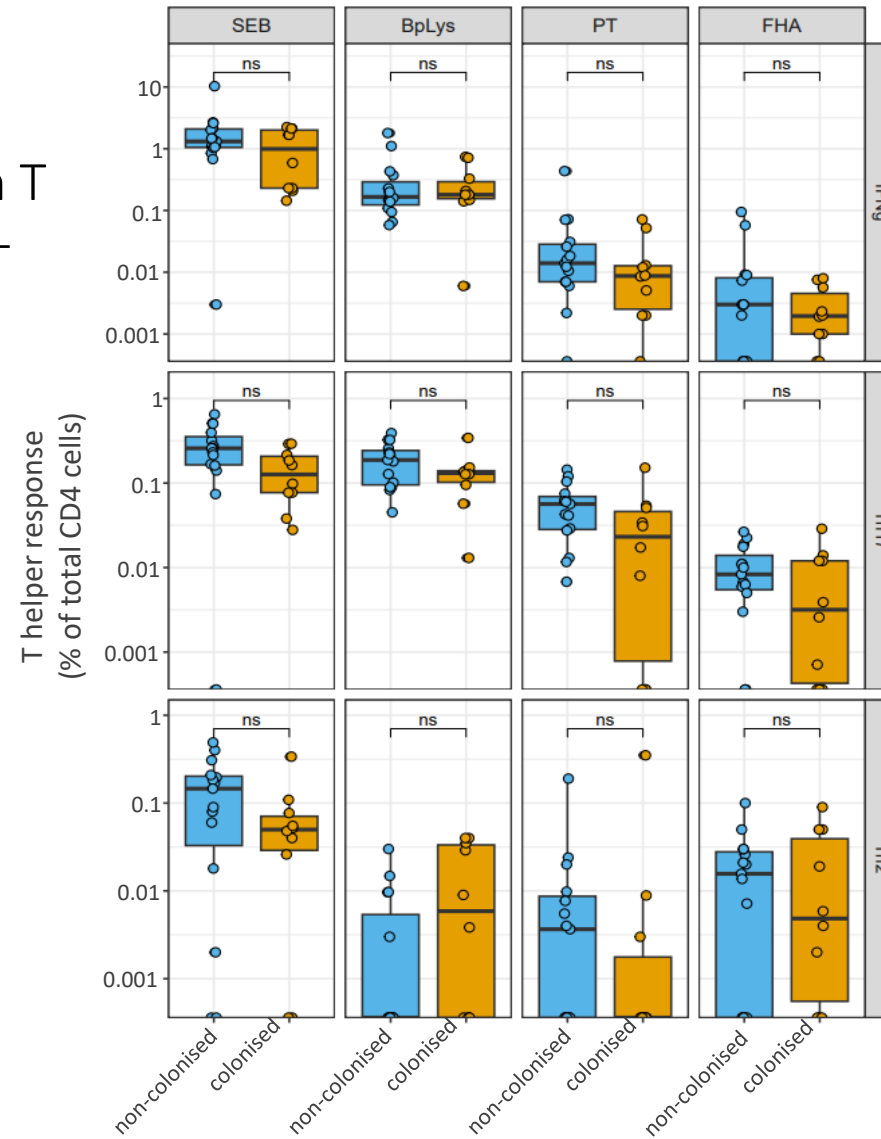


Total CFU in period 0 - 14 days after challenge

Baseline memory B cell concentrations

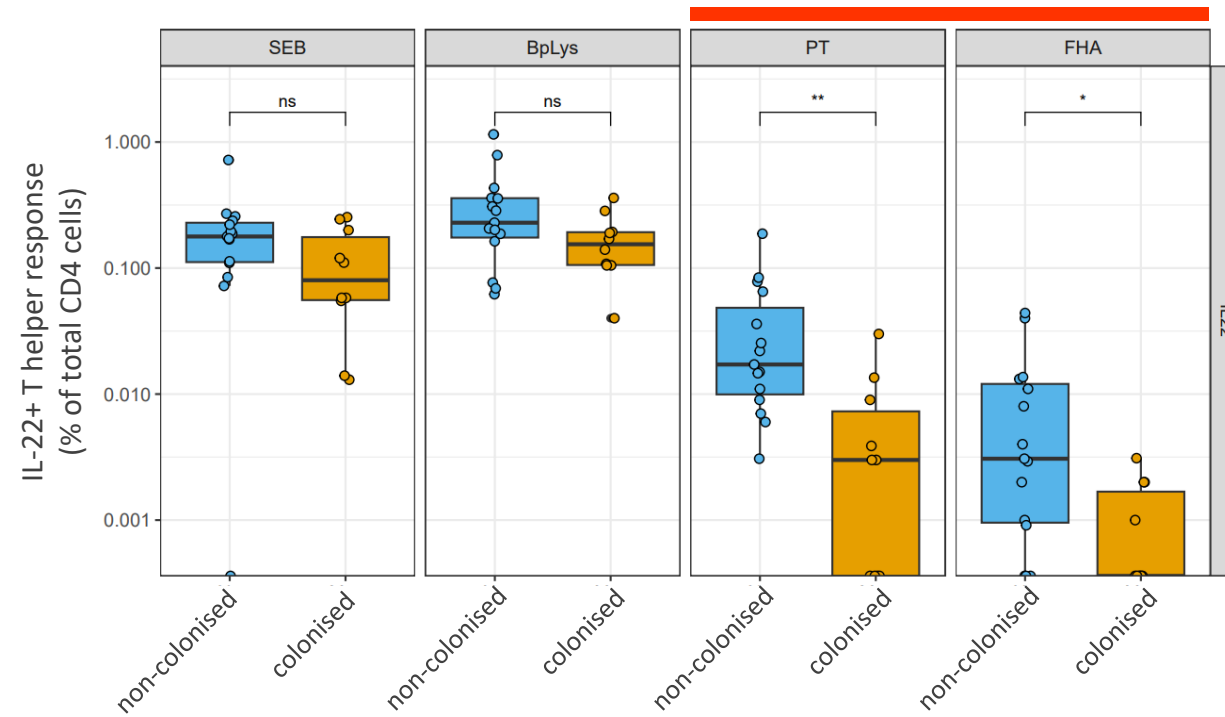


Baseline comparison between T helper cell responses in non-colonised and colonised volunteers



Data is mock subtracted

Baseline IL-22+ T helper cell responses



Overall Summary – PERISCOPE Human Challenge Program

- A dose of 10^5 cfu BP1917 induces:
 - In selected participants (PT \leq 20)
 - Colonisation Fraction 0.8
 - In unselected participants
 - Colonisation Fraction 0.4-0.56

Overall Summary – PERISCOPE Human Challenge Program

- Minor symptoms occur and are tolerated, even in non-colonised
- Nasal washing is the most sensitive microbiological sampling technique
- Colonisation is detected in most by day 7 and density peaks at Day 14
- Colonisation clears spontaneously but this may take weeks
- Azithromycin clears infection by 48 hours in most cases
- There is no environmental shedding, and transmission has not been observed
- The model can be safely conducted in an outpatient environment
- Colonisation induces a `protected` phenotype

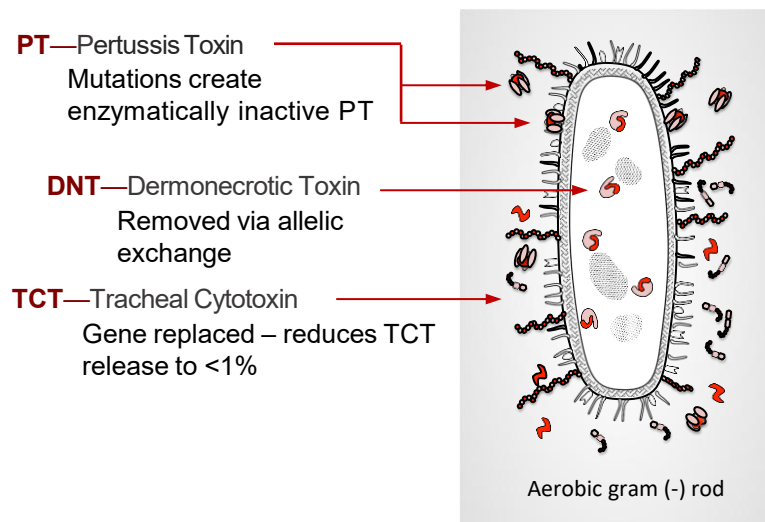
Overall Summary – PERISCOPE Human Challenge Programme

- **Protection against colonisation is associated with:**
 - Serum IgG anti-PT, anti-PRN, anti-FHA
 - Serum IgA anti-FHA, and IgA binding to Bp
 - Nasal mucosal lining fluid IgA binding to Bp
 - IL22-expressing T helper cell responses to PT and FHA
- **Control of colonisation density is associated with:**
 - Serum IgG anti-PT and anti-FHA
 - Nasal mucosal lining fluid IgG binding to Bp
 - Bp-specific Memory B cell responses
- **Colonisation induces a `protective` immunophenotype**

BPZE1 Vaccine

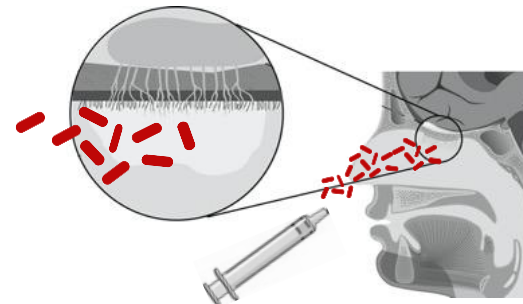
Live attenuated vaccine

- *B. pertussis* Tohama I strain
- Genetically modified with 3 toxins inactivated or removed



Mucosal delivery

- Lyophilised formulation
- Nasally administered by mucosal atomization device



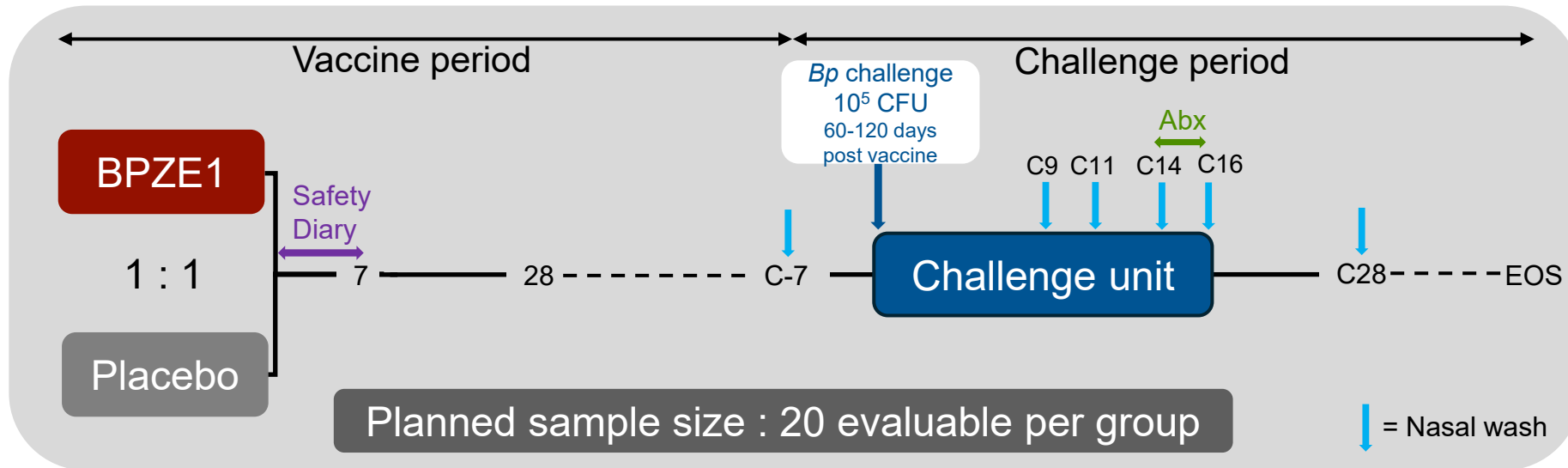
Characteristics

- Acceptable safety profile
- Transient colonisation
- Protection against rechallenge with BPZE1
- Immunogenic
 - Mucosal s-IgA
 - Serum IgG and IgA
- Functional
 - Serum bactericidal activity against PRN+ and PRN- *B. pertussis*
 - Pertussis toxin neutralisation assay

Jahnmatz et al 2020, Creech et al 2022, Keech et al 2023

BPZE1 Vaccine-challenge study

- BPZE1 impact on asymptomatic colonisation
- Anti-PT IgG < 20 IU/mL; Anti-PRN IgG < 30 IU/mL
- Primary endpoint: no colonisation in nasal wash cultures on Days 9, 11 and 14 post virulent *Bp* challenge



Efficacy, immunogenicity, and safety of the live attenuated nasal pertussis vaccine, BPZE1, in the UK: a randomised, placebo-controlled, phase 2b trial using a controlled human infection model with virulent *Bordetella pertussis*

Diane Gbesemete, Mahesh N Ramasamy, Muktar Ibrahim, Alison R Hill, Lucy Raad, Daniela M Ferreira, Jonathan Gray, Adam P Dale, Joy R Laver, Tjeren Coutinho, Saul N Faust, Thomas A N Reed, Gavin Ballhage, Lisa Westfeldt, Wei Long, Camille Loch, Vivek Samal, Peter Goldstein, Ken Solovey, Keith Rubin, Stephanie Novello, Robert C Reed

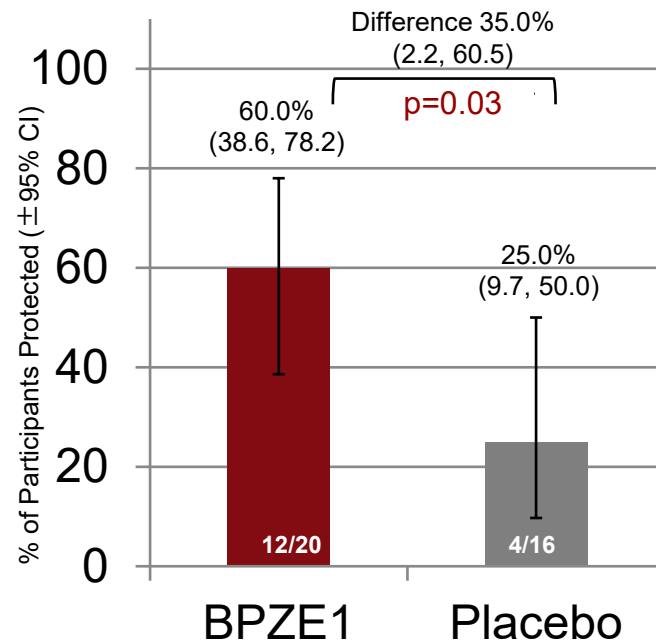
Summary
Background Pertussis is a severe respiratory disease caused by *Bordetella pertussis*. Although vaccines prevent disease for a limited duration, they do not prevent infection and transmission. We aimed to assess the safety and efficacy of BPZE1 at preventing or substantially reducing colonisation by virulent *B pertussis* using a robust controlled human infection model.

Methods This randomised, placebo-controlled, phase 2b trial was conducted at University Hospital Southampton and

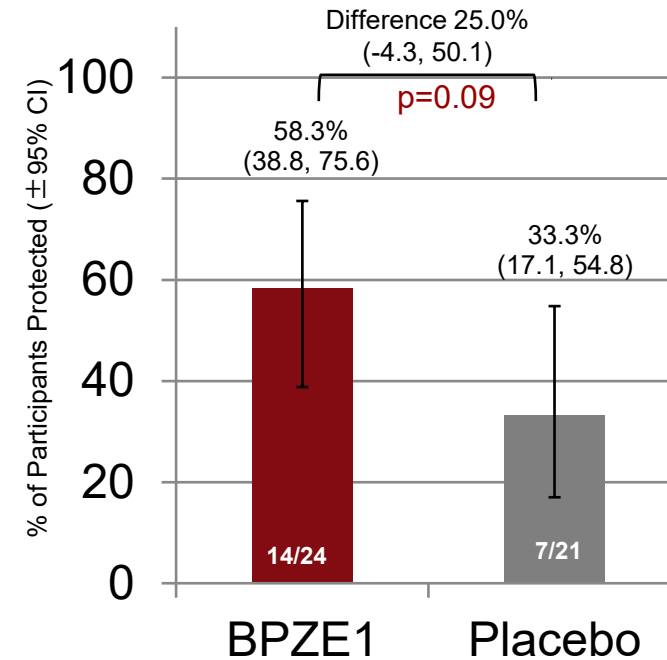


Colonisation

Per-Protocol Adequate Challenge Inoculum

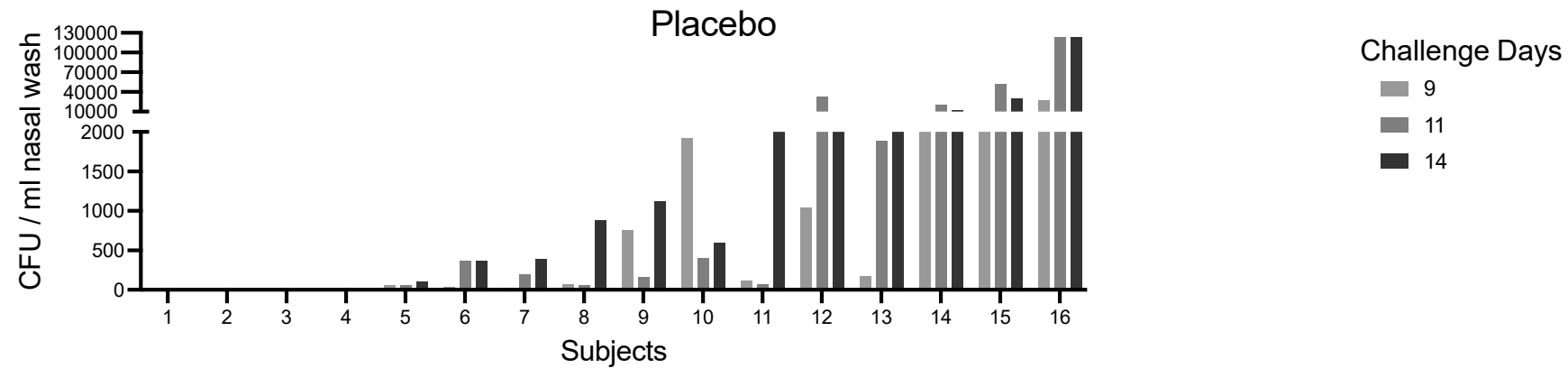
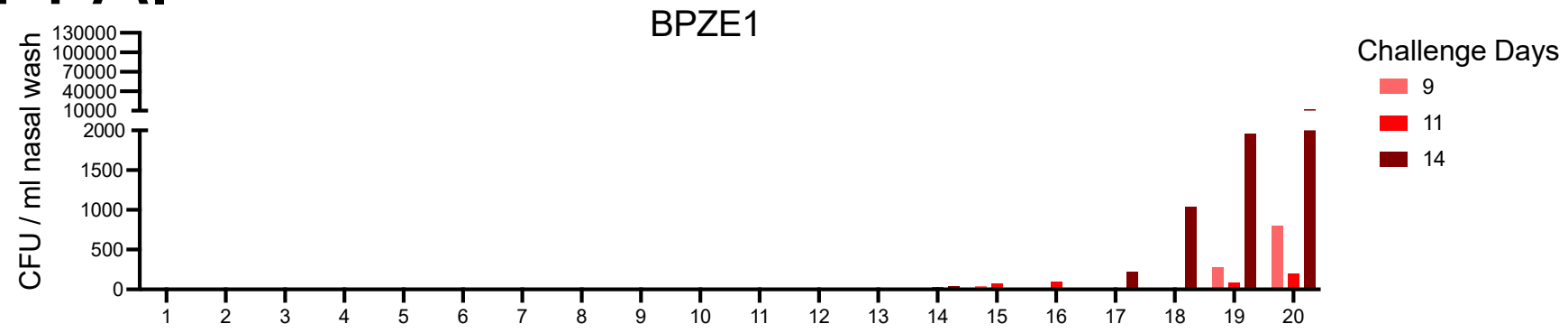


Modified Intent-to-Treat



p-values using Chi-square test

By-subject Colonisation Density - PPAI



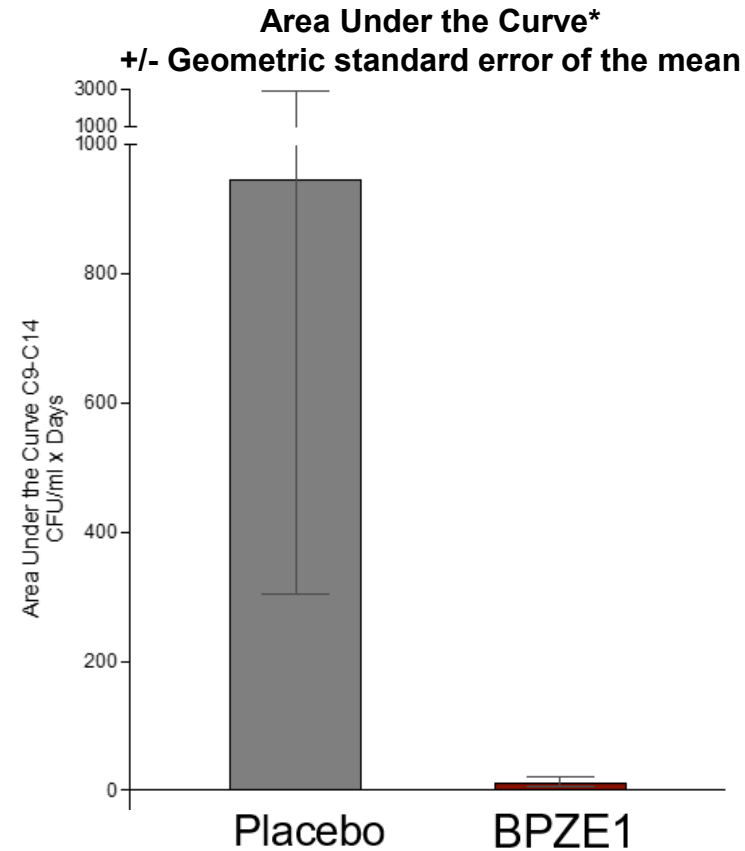
B. pertussis burden

Per-protocol Adequate Inoculum

- BPZE1-vaccinated participants had a **98.7%** reduction vs placebo

Modified Intent-to-Treat

- BPZE1-vaccinated participants had a **97.1%** reduction vs placebo

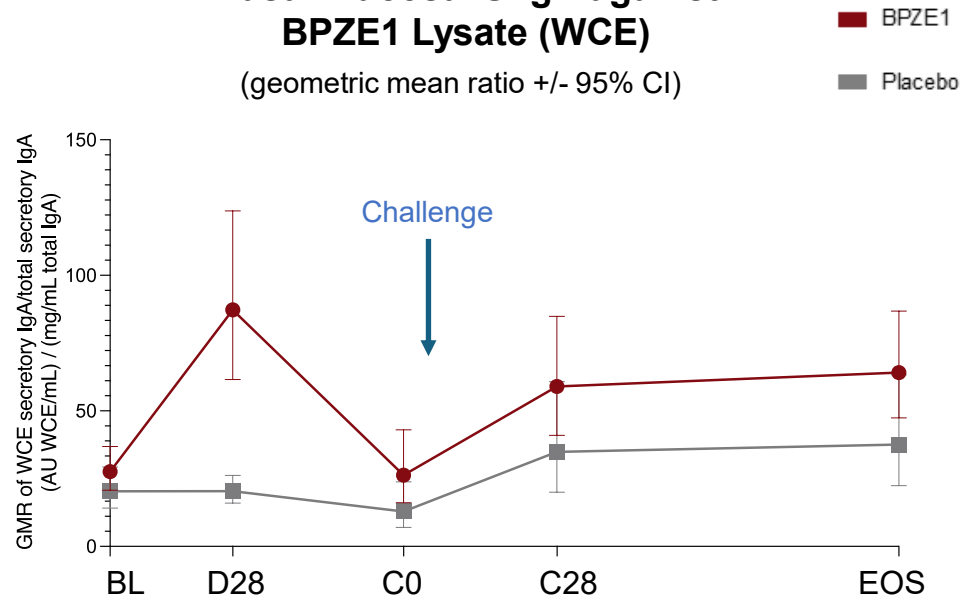


*Geometric means of individual AUCs from C9 through C14 using the trapezoidal rule

Immunogenicity

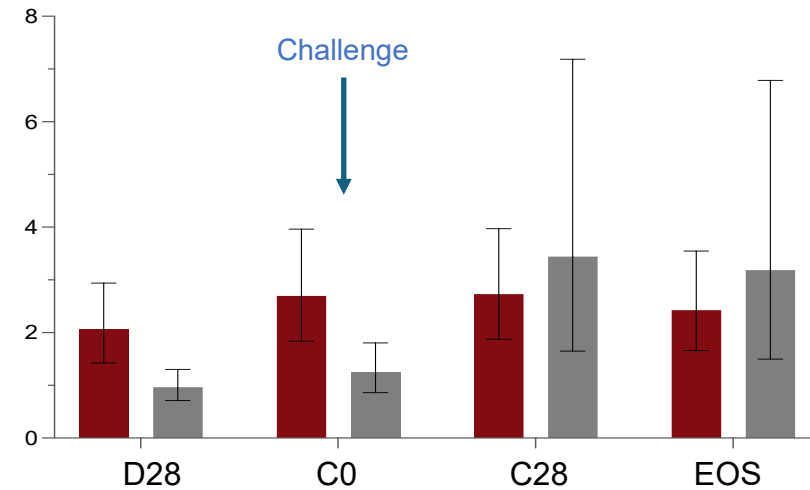
Nasal Mucosal S-IgA against BPZE1 Lysate (WCE)

(geometric mean ratio +/- 95% CI)



Serum IgG Anti-Pertussis Toxin

(geometric mean fold rise from baseline +/- 95% CI)



- Similar trends in S-IgA antibodies against PRN, FHA, FIM, PT

BL = baseline; D28 = Day 28 post vaccine; C0 = pre-challenge; C28 = Day 28 post challenge; EOS = End of study; GMR = geometric mean ratio; IgG = immunoglobulin G; mL= millilitre; WCE=whole cell extract

Immune correlates

(Hill A, Dale A. et al manuscript submitted)

- Vaccination with BPZE1 induced circulating CD4⁺ T cells of Th17 and Th22 effector phenotype.
- These responses correlated positively with BPZE1-induced SIgA responses
- In BPZE1 recipients who developed breakthrough infection following virulent *B. pertussis* challenge, higher vaccine-induced SIgA titres were significantly associated with reduced colonisation density (but serum IgA and IgG titres were not)
- A Th17/Th22-SIgA immune axis as a key feature of BPZE1 vaccination and highlight SIgA as a candidate correlate of reduced *B. pertussis* colonisation in humans.



Thanks to the volunteers, the teams in Southampton and Radboud, and our partners within PERISCOPE

Southampton University, UK: **Nursing staff, Clinical Research Facility lab, CHIG Technician team, Muktar Ibrahim, Alison R. Hill, Diane Gbesetmete, Robert C. Read**

RadboudUMC, The Netherlands: **Dimitri A. Diavatopoulos, Janeri Fröberg, Hans de Graaf**

RIVM, The Netherlands: **Annemarie Buisman, Cecile Van Els, Guy A.M. Berbers**

UK Health Security Agency, UK: **Andrew Gorringe**

IAVI, USA: **Kent E. Kester**

ULB, Belgium: **Véronique Corbière, Françoise Mascart**

PERISCOPE has received funding from the Innovative Medicines Initiative 2 Joint Undertaking under grant agreement No 115910. This Joint Undertaking receives support from the European Union's Horizon 2020 research and innovation programme and EFPIA and BMGF.





Robert Read
 Lucy Raud
 Tyween Coutinho
 Alison Hill
 Jon Guy
 Jay Laver
 Adam Dale
 Muhammad Ahmed
 Muktar Ibrahim
 Hans de Graaf



Stephanie Noviello
 Camille Locht
 Peter Goldstein
 Andre Morias Campos
 Theresa Creighton
 Lisa Weissfeld
 Wei Lang
 Keith Rubin
 Kevin Solovay



Maheshi Ramasamy
 Daniela Ferreira
 Sophia Hawkins
 Carla Solorzano Gonzalez
 Jasmin Kinch
 Hannah Robinson
 Bruno Rocha de Macedo
 Juliette Cotton
 Conor Whelan

Thank you

Study participants and site staff



UK Health
 Security
 Agency

Mary Matheson
 Hannah Cuthbertson



Radhika Mhatre
 Angela Williamson
 Diana Dimov
 Maria Rios
 Ken Almedilla
 Simphewe Ndlovu



Bea Selby
 Liga Krauge
 Devi Nair-Apadoo
 Stephen Saich
 and the CRF PM team

Patricia Acosta Carrazco
 Alice Sebastien
 Marta Adamczyk
 Liza Shiner-Clark
 Caroline Grabau
 and the CRF Nursing team

Laura Presland
 Sarah Horswill
 Tim Whitbread
 and the Vaccine Hub team

Thomas Reed
 Thomas Durham
 Simone Paulson
 and the CRF Fellows

Gavin Babbage
 Eloise Summerton
 and the CRF lab team