

The 3rd International Symposium on  
Microgenomics 2023

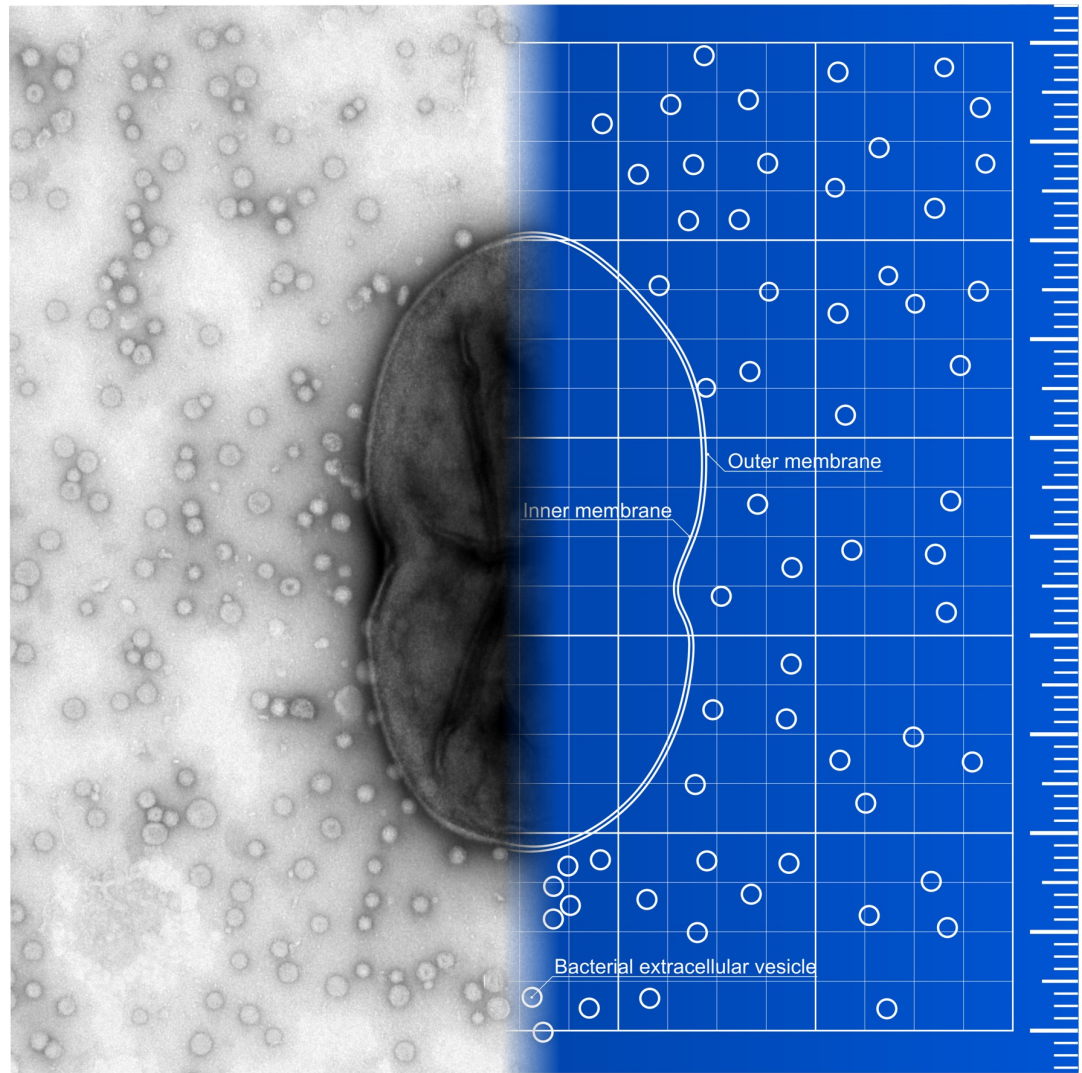
***Bacterial Extracellular  
Vesicles:  
Spheres of influence  
within and beyond the  
gut***

**Simon R Carding**

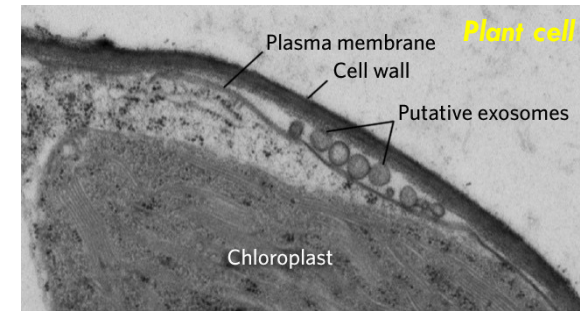
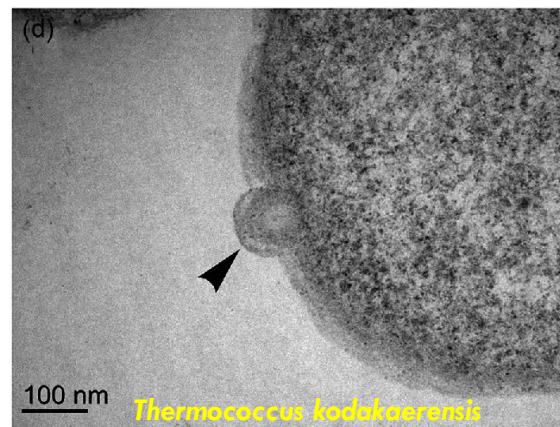
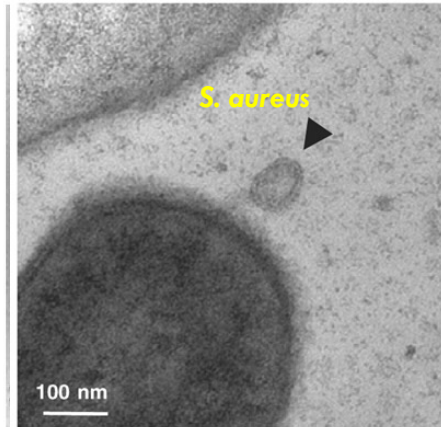
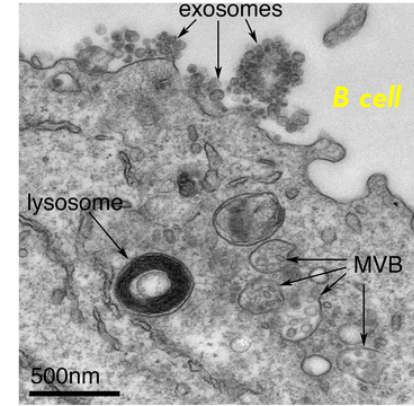
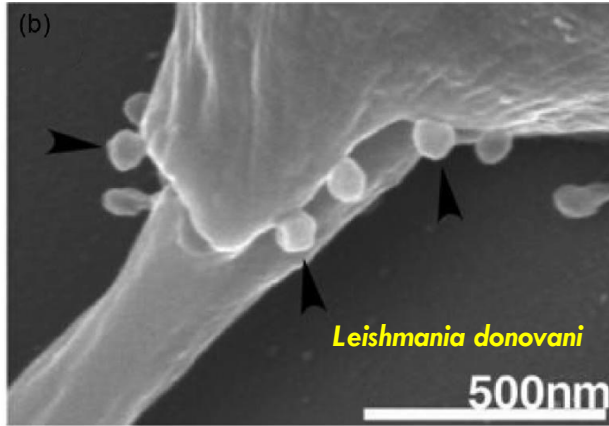
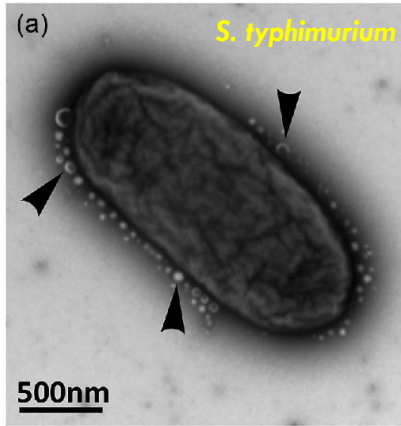
**Food, Microbiome, and Health  
Research Programme,  
The Quadram Institute**

**&**

**Norwich Medical School, UEA, UK**



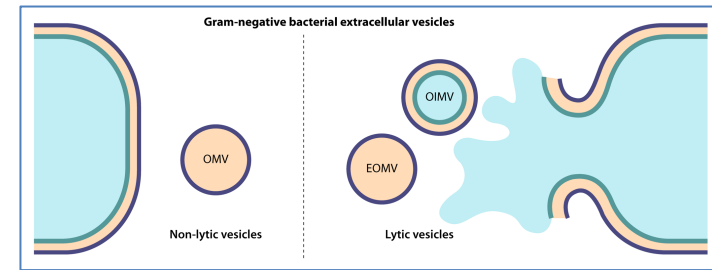
# Cells from all three domains of life produce EVs: Evolutionarily conserved mediators of intercellular communication



Gill et al., *FEMS Microbiol Rev* 2019; Lee et al., *Proteomics*, 2009; Edgar *BMC Biol.* 2016; Palmer, *ASPB*, 2019

# BEVs: a *bona fide* bacterial secretion system

- First described in 1960's by Bishop and Work for Gram negative bacteria and 30 years later for Gram positive bacteria
- Naturally shed by Gram negative bacteria during normal growth cycle by:
  - Budding from the outer membrane (OMVs)
  - Explosive lysis - outer inner MVs and outer MVs
  - Gram positive bacteria produce cytoplasmic MV
- Genetic basis for regulated production (stress response pathways)
- Selective, non-random, packaging of cytoplasmic and periplasmic contents

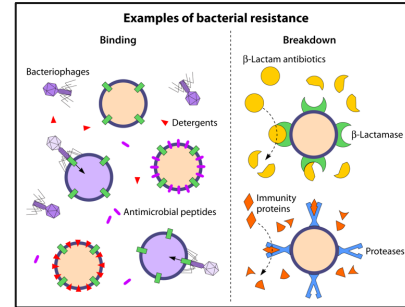


Juodeikis and Carding, 2022

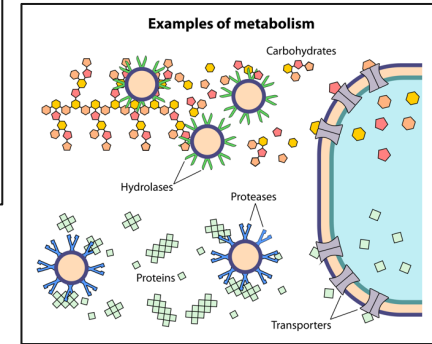


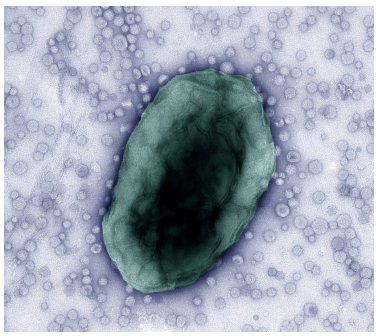
# BEVs: It's energetically costly so why bother?

- Garbage disposal
- Re-modelling
- Building fences and self protection
- Export and transport
- Scavenging
- Good neighbors, community development and altruism
- Détente and stabilising cooperative relationships with the host



Juodeikis and Carding, 2022, *MMBR*





# *Bacteroides thetaiotaomicron* (Bt): A Model Human GIT Commensal

Obligate anaerobe, non-endospore forming, motile/ immotile:

6.26 MB genome encoding 4776 proteins, ~90% for binding/import of polysaccharides

Wide environmental distribution: Constituent of multiple biomes inc. animal microbiomes

Represents ~10-25% of all anaerobes in the human colonic microbiota

Normally resident at the mucosal interface in the lower GIT

# *Bacteroides* BEVs - Questions of Interest

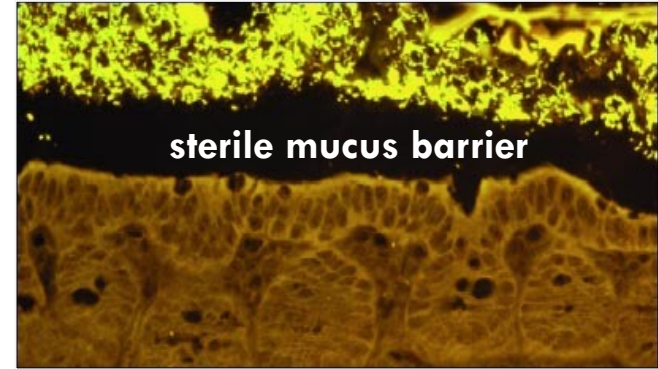
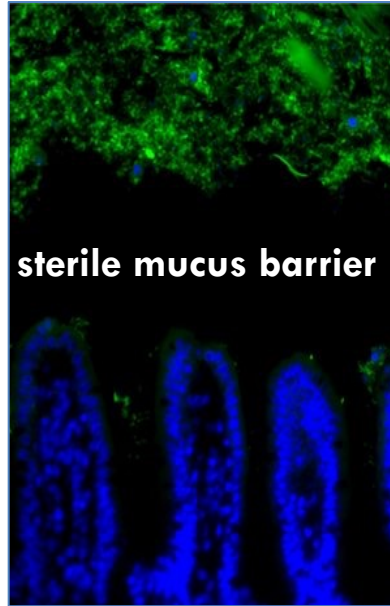
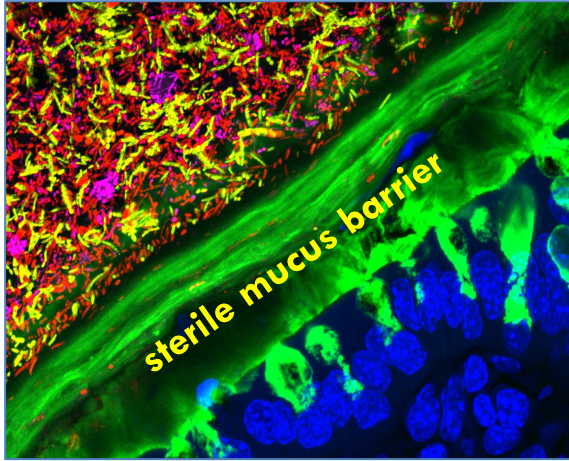
- Production
- Distribution
- Function
- Exploitation

## Context

Defining the mechanism and pathways by which gut bacteria communicate with the host to promote health

# How do gut microbes communicate with their host?

Vaishnava et al. 2009. *J. Physiol. Pharm.*



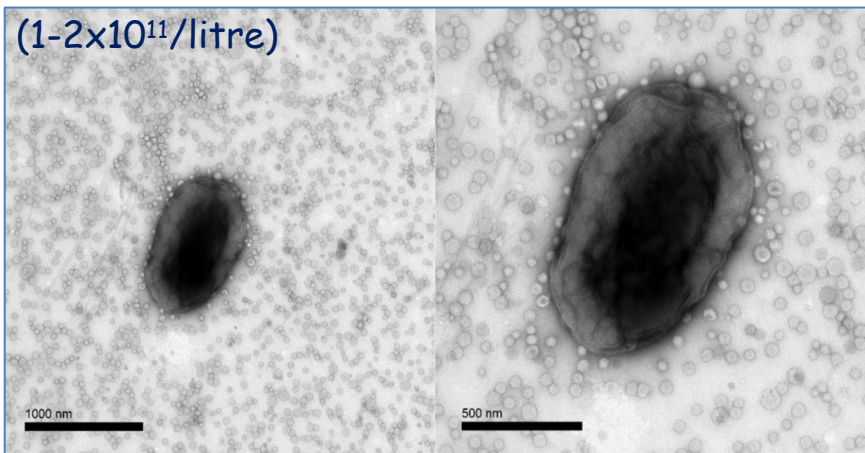
Mucosal barrier crossing by signals is required for microbe-host cell crosstalk

Kamphuis et al. 2017. *Scientific Reports*

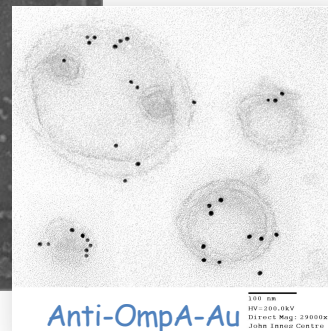
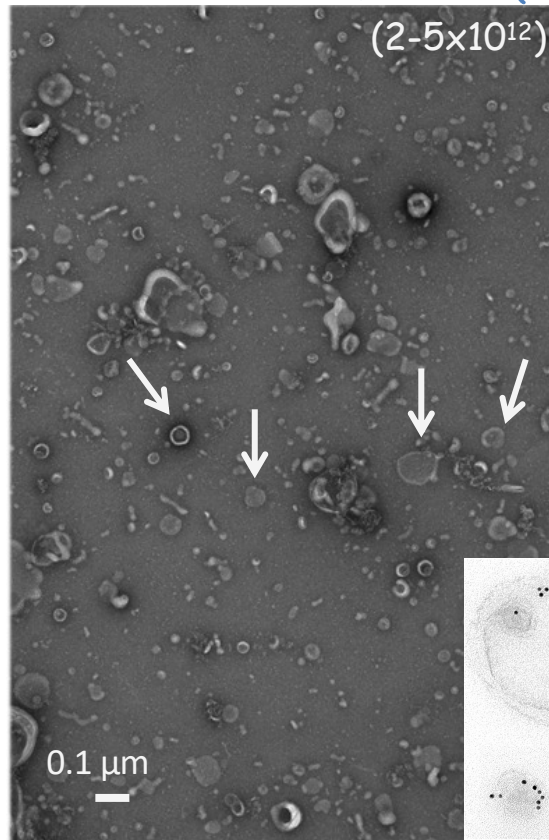
HOW?

- ❖ Small Diffusible Molecules
  - ❖ Signalling Molecules & Hormones
  - ❖ SCFAs
- ❖ Nanovesicles (BEVs)

# Bt BEVs: Abundant & naturally produced in the GIT

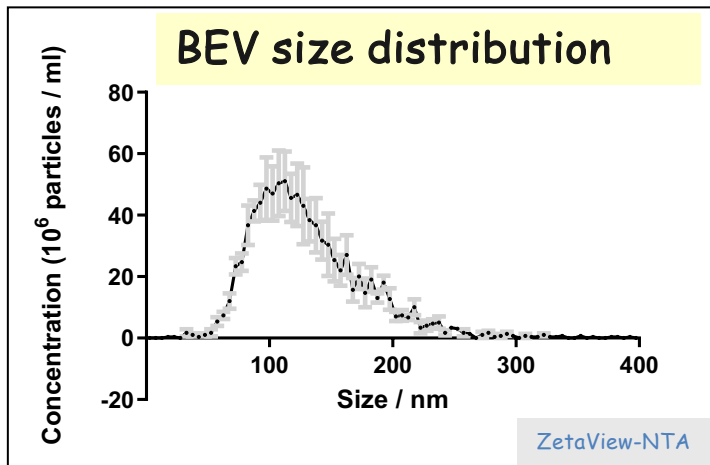


Bt mono-colonised *Germfree* mice (caecum)



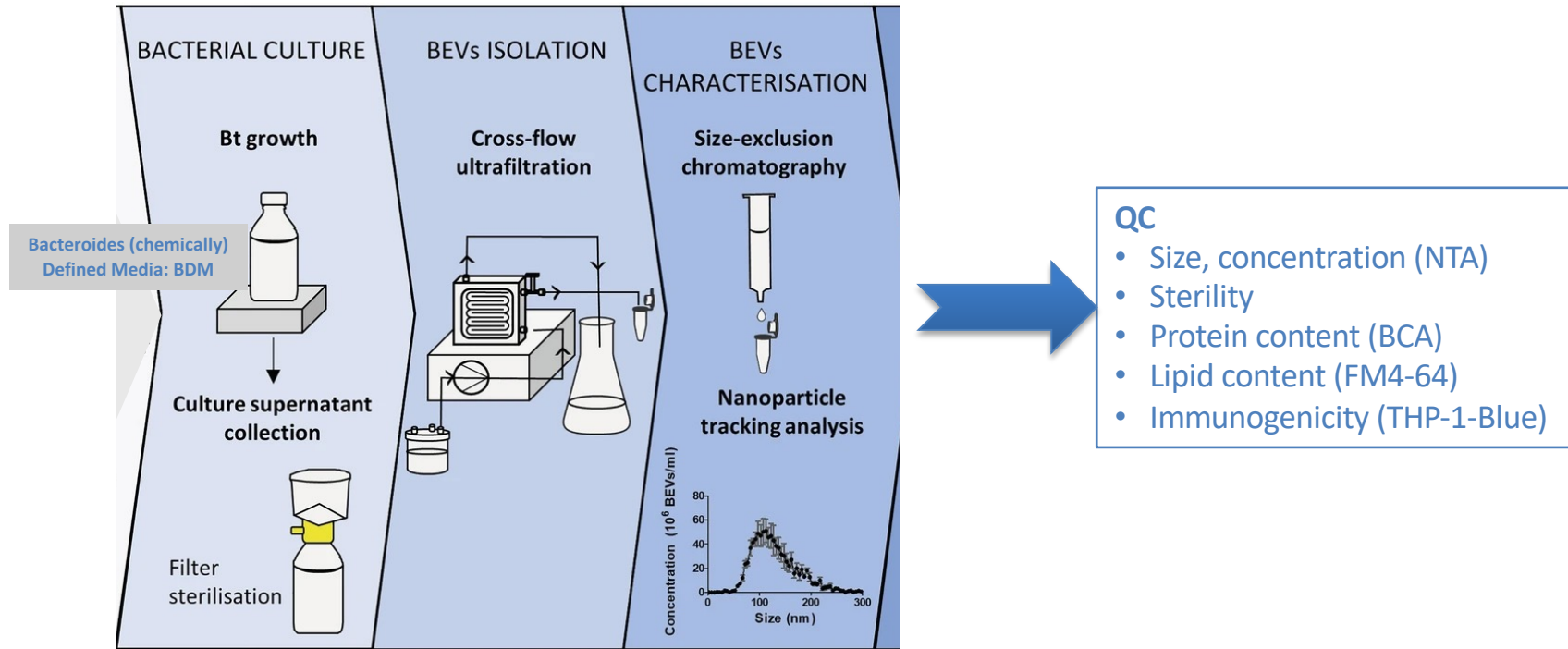
R. Stentz, K. Cross

Ian Brown





# BEV isolation & characterisation



# Bt BEVs *in vivo* enriched in metabolic enzymes

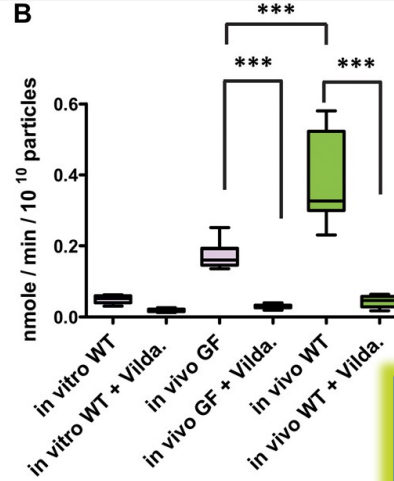
## BEVs + Ala-Pro-pNa (DPP IV)

Cell Reports  
Article

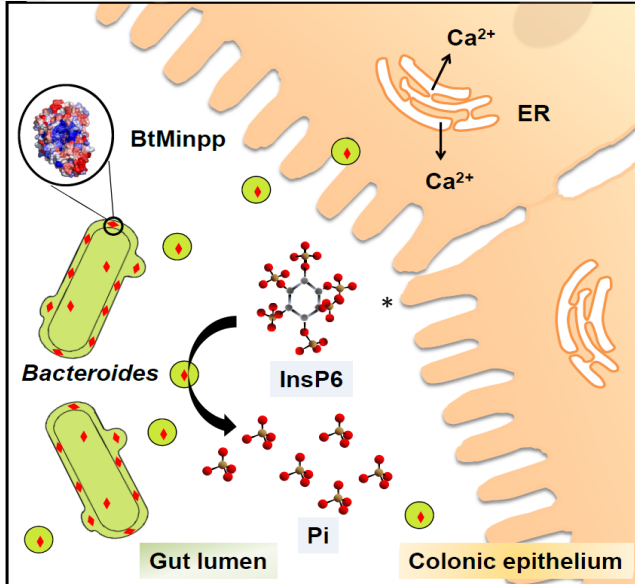
Open ACCESS

**A Bacterial Homolog of a Eukaryotic Inositol Phosphate Signaling Enzyme Mediates Cross-kingdom Dialog in the Mammalian Gut**

Régis Stentz,<sup>1</sup> Samantha Osborne,<sup>1</sup> Nikki Hom,<sup>1</sup> Arthur W.H. Li,<sup>2</sup> Isabelle Hautefort,<sup>1</sup> Roy Bongaerts,<sup>1</sup> Marine Rouyer,<sup>1</sup> Paul Bailey,<sup>2</sup> Stephen B. Shears,<sup>2</sup> Andrew M. Hemmings,<sup>2,3</sup> Charles A. Brearley,<sup>2,4</sup> and Simon R. Carding<sup>1,4,\*</sup>



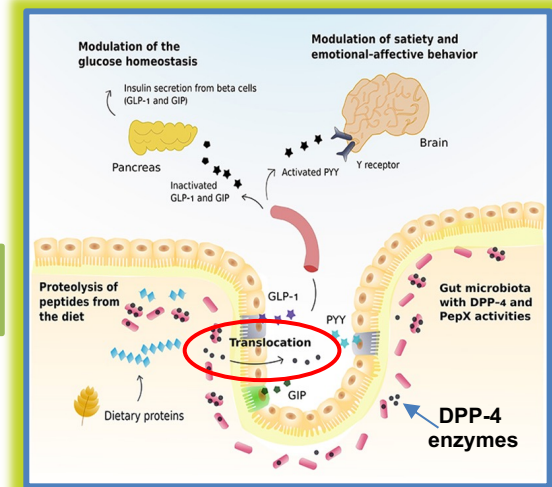
\*0.33μM Vildagliptin (DPP IV inhib.)



Human GLP-1 metabolism (QMUL)

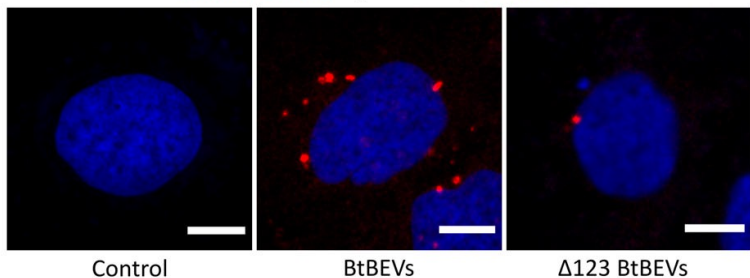
Recombinant DPP IV degrades:

- ✓ **GLP-1**
- ✓ **Substance P**

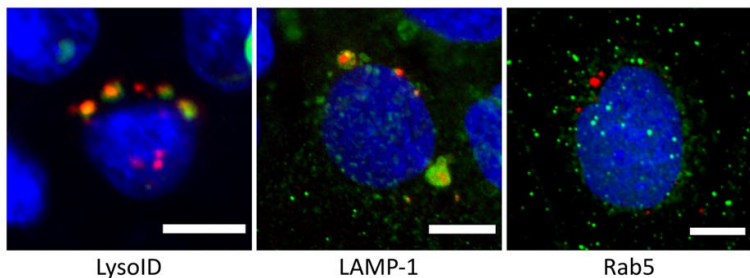


# Bt BEVs *in vivo* express high levels of BtuG proteins that deliver cobalamin/vit B12 to host cells

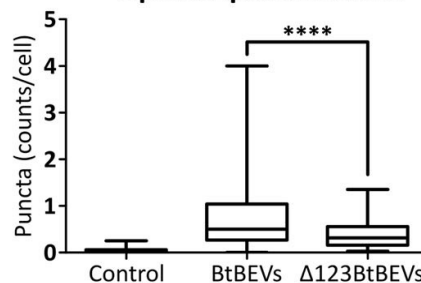
(a) **fb12-BtBEV uptake by Caco-2 cells**



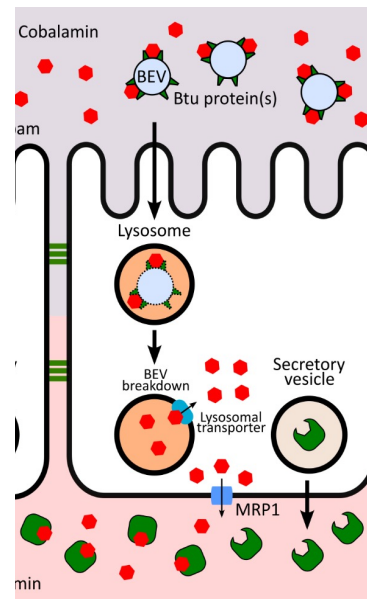
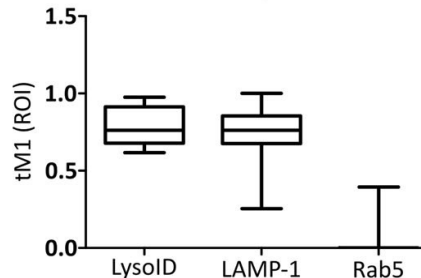
(c) **fb12-BtBEV localisation in Caco-2 cells**



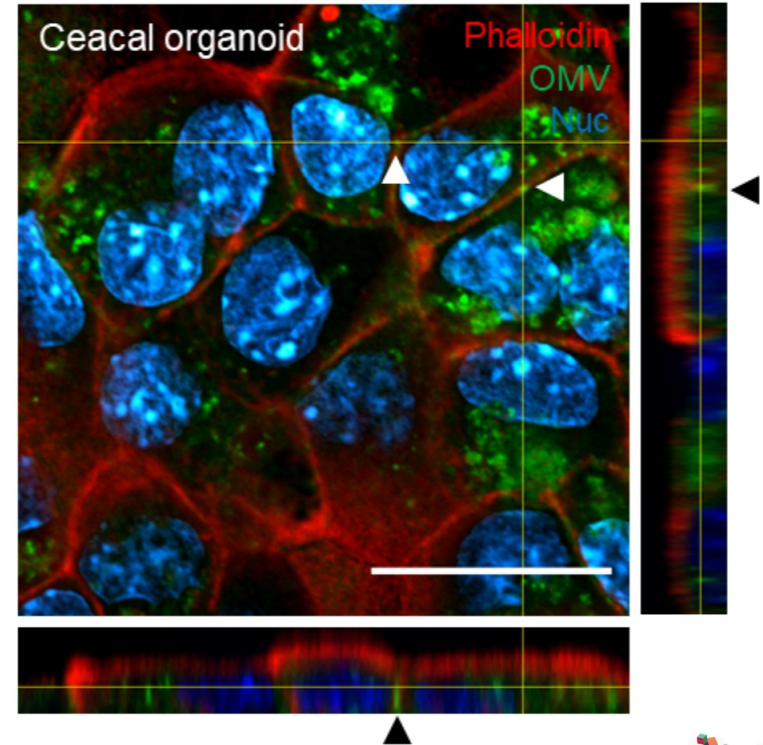
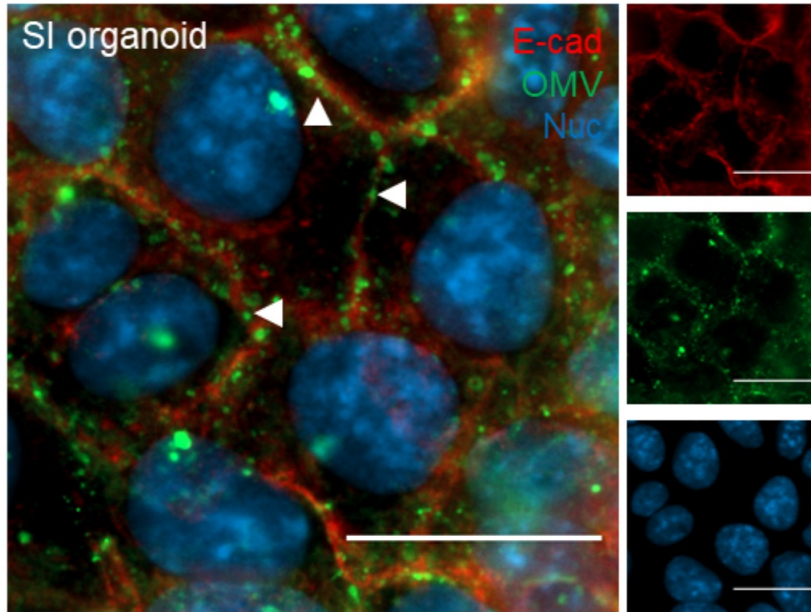
(b) **Uptake quantification**



(d) **Localisation quantification**



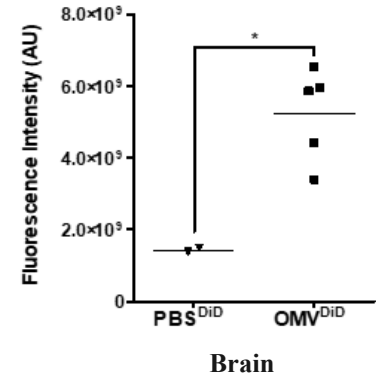
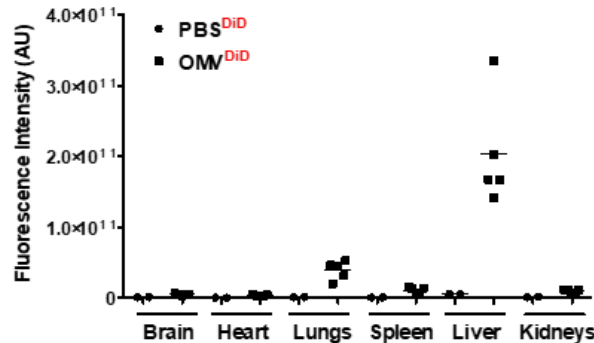
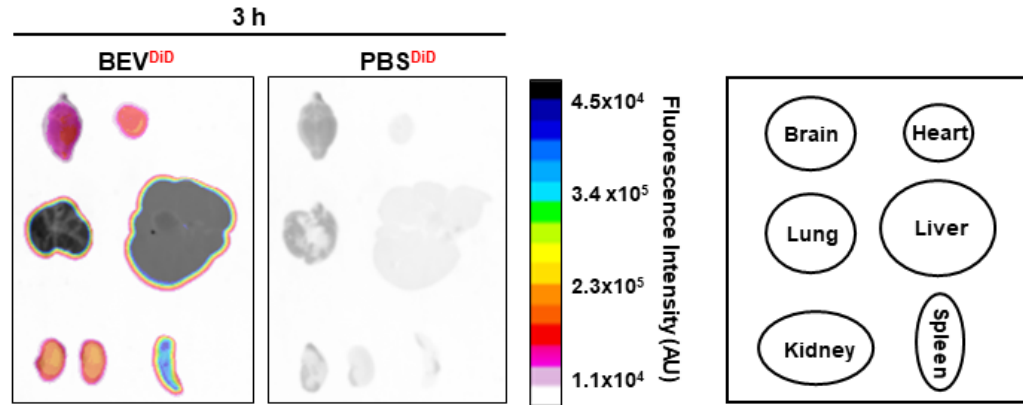
# Bt BEVs access and cross the host GIT



# Biodistribution of DiD- labelled Bt BEVs

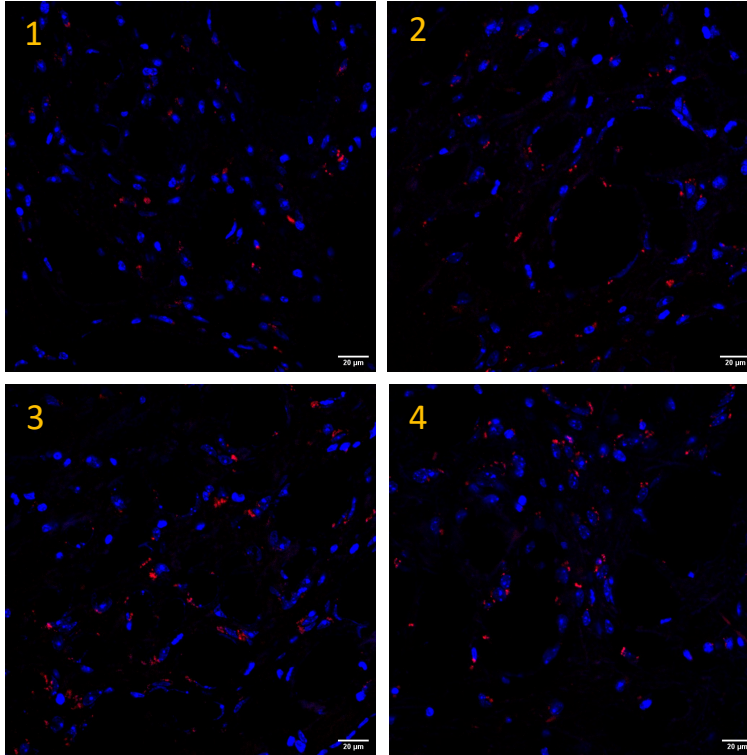
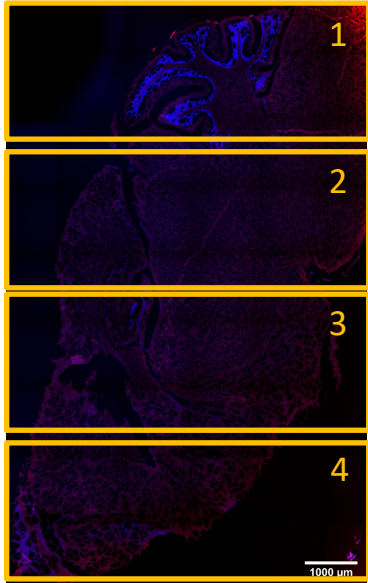
## Post oral administration to adult SPF mice

*In vivo* Xtreme multi-modal optical and x-ray small animal imaging system (Bruker) with a back-illuminated 4MP CCD detector



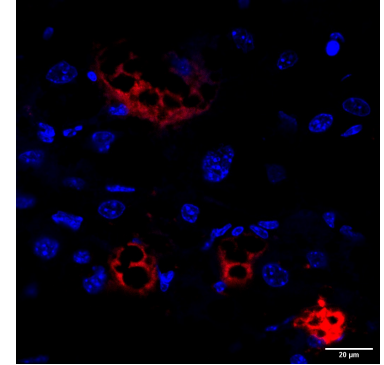
# Localisation of DiD-Bt BEVs to the Liver & CNS

R hemisphere

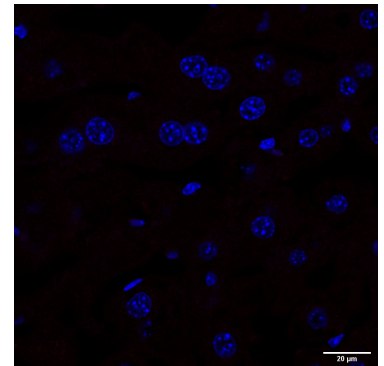


Median lobe

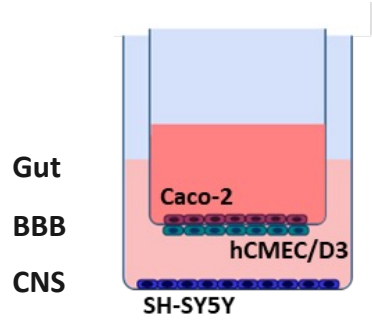
+BEVs



+PBS

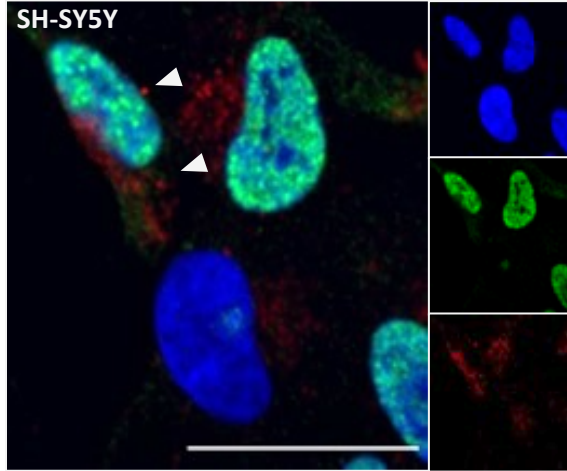


# A simplistic *in vitro* three cell model of the gut-blood-brain axis

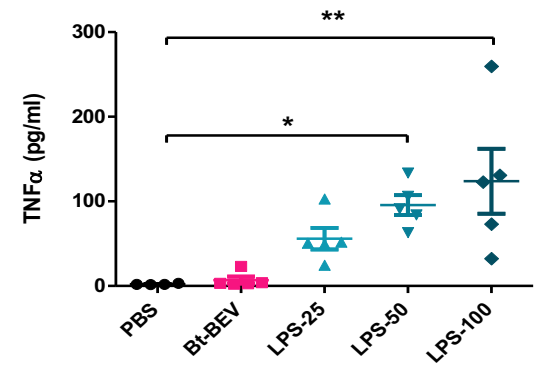
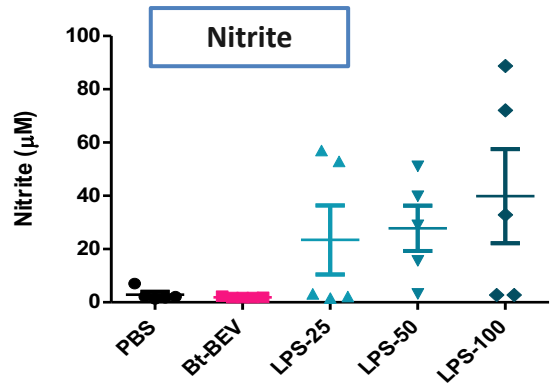
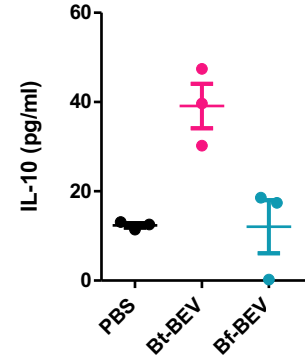


Nuclei  
PCNA  
DiD-BEVs

+ Bt-BEV



IL-10

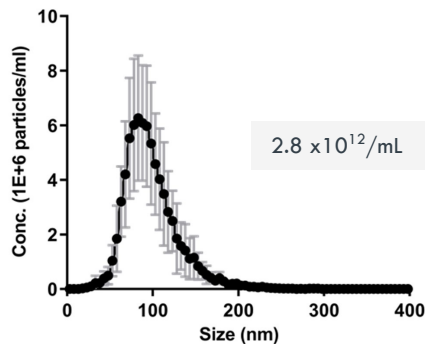
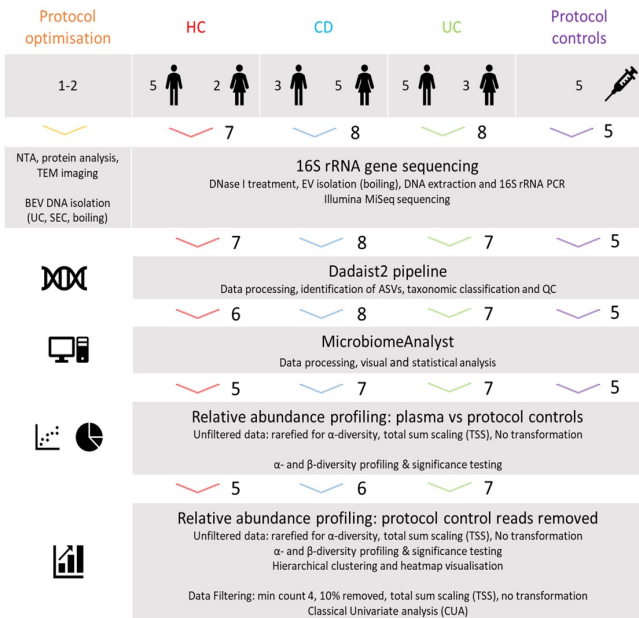


TNFα

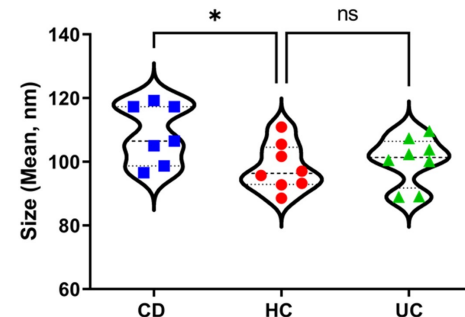
# Plasma of healthy individuals contains BEVs

Article  
**The Origin of Plasma-Derived Bacterial Extracellular Vesicles in Healthy Individuals and Patients with Inflammatory Bowel Disease: A Pilot Study**

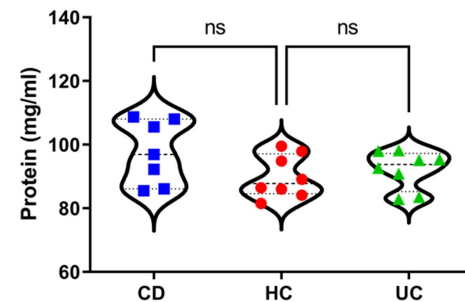
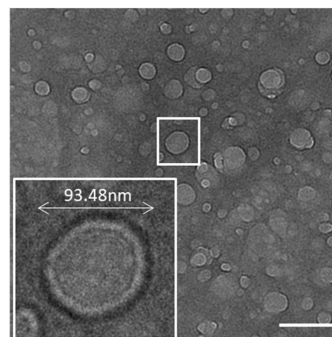
Emily Jones <sup>1</sup>, Régis Stentz <sup>1</sup>, Andrea Telatin <sup>1</sup>, George M. Savva <sup>1</sup>, Catherine Booth <sup>2</sup>, David Baker <sup>1</sup>, Steven Rudder <sup>1</sup>, Siella C. Knight <sup>1</sup>, Alistair Noble <sup>3,4</sup> and Simon R. Carding <sup>1,4,\*</sup>



(a)

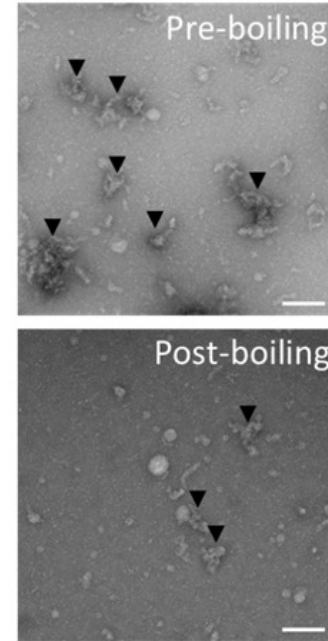
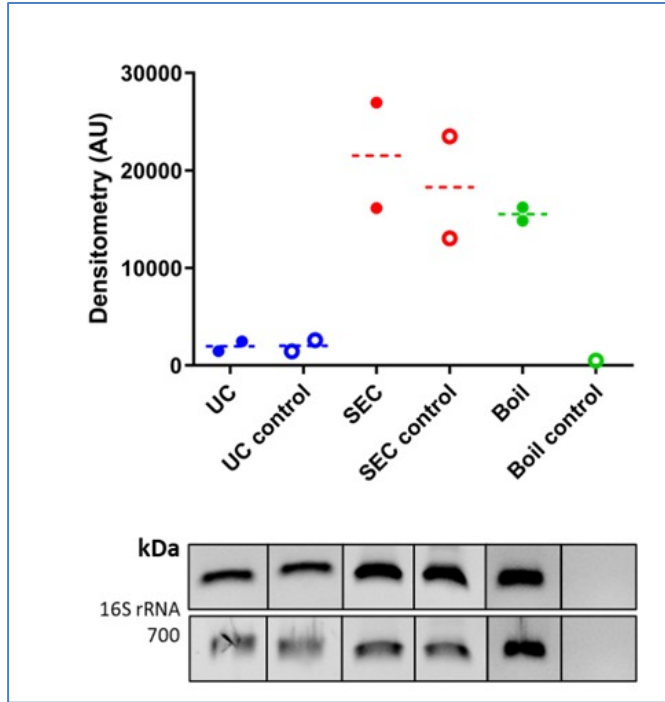


(b)





# Optimising bacterial extracellular DNA isolation from plasma BEVs



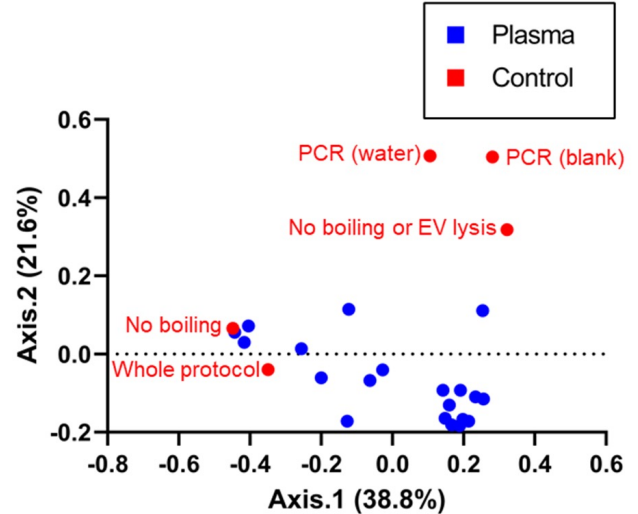
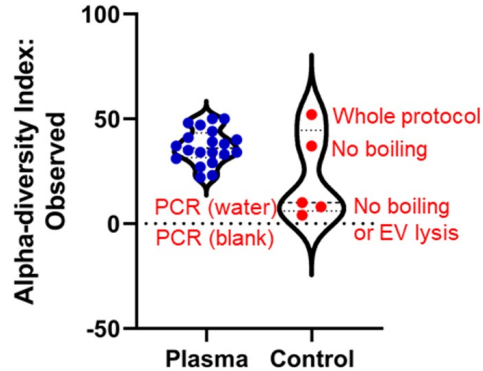
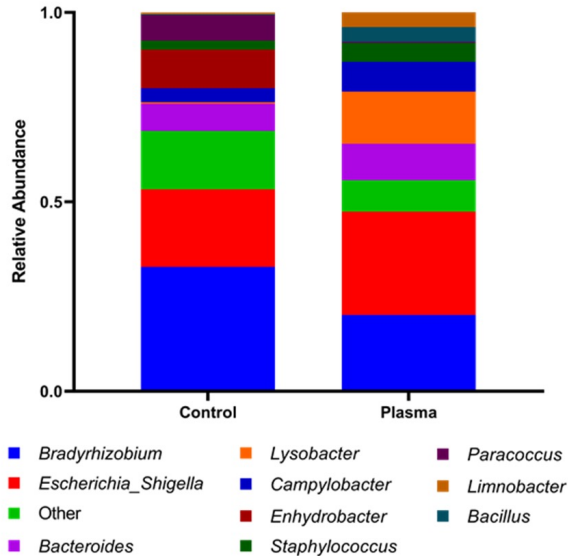
UC = ultracentrifugation (150,000 × *g*, 2 h at 4°C)

SEC = size exclusion chromatography (qEV/35nm series column)

Boil = boiling (100°C for 40 min + centrifugation at 13,000 × *g*, 30 mins at 4°C)

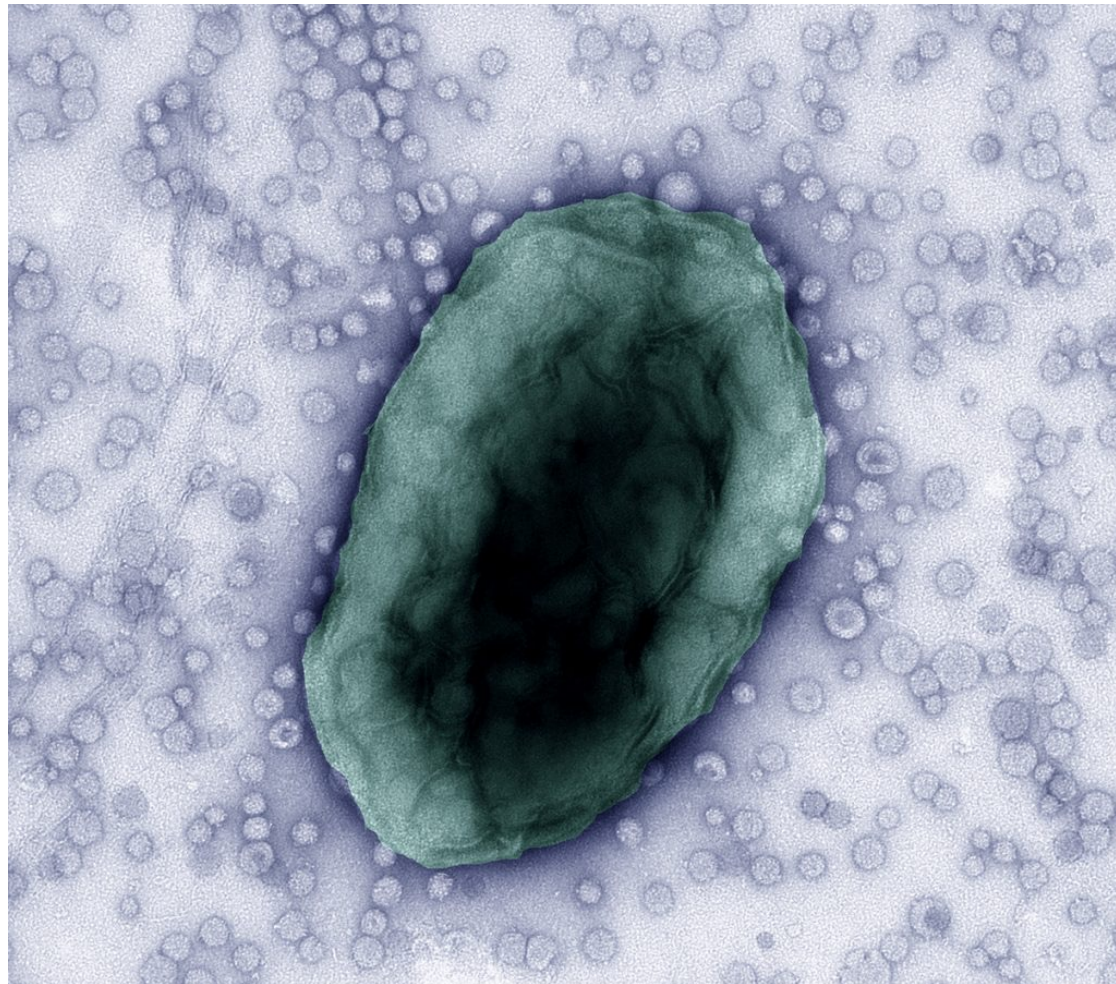
2.9-3.1 ng/ml DNA

# Identifying sources of contaminating DNA (kit-omes)

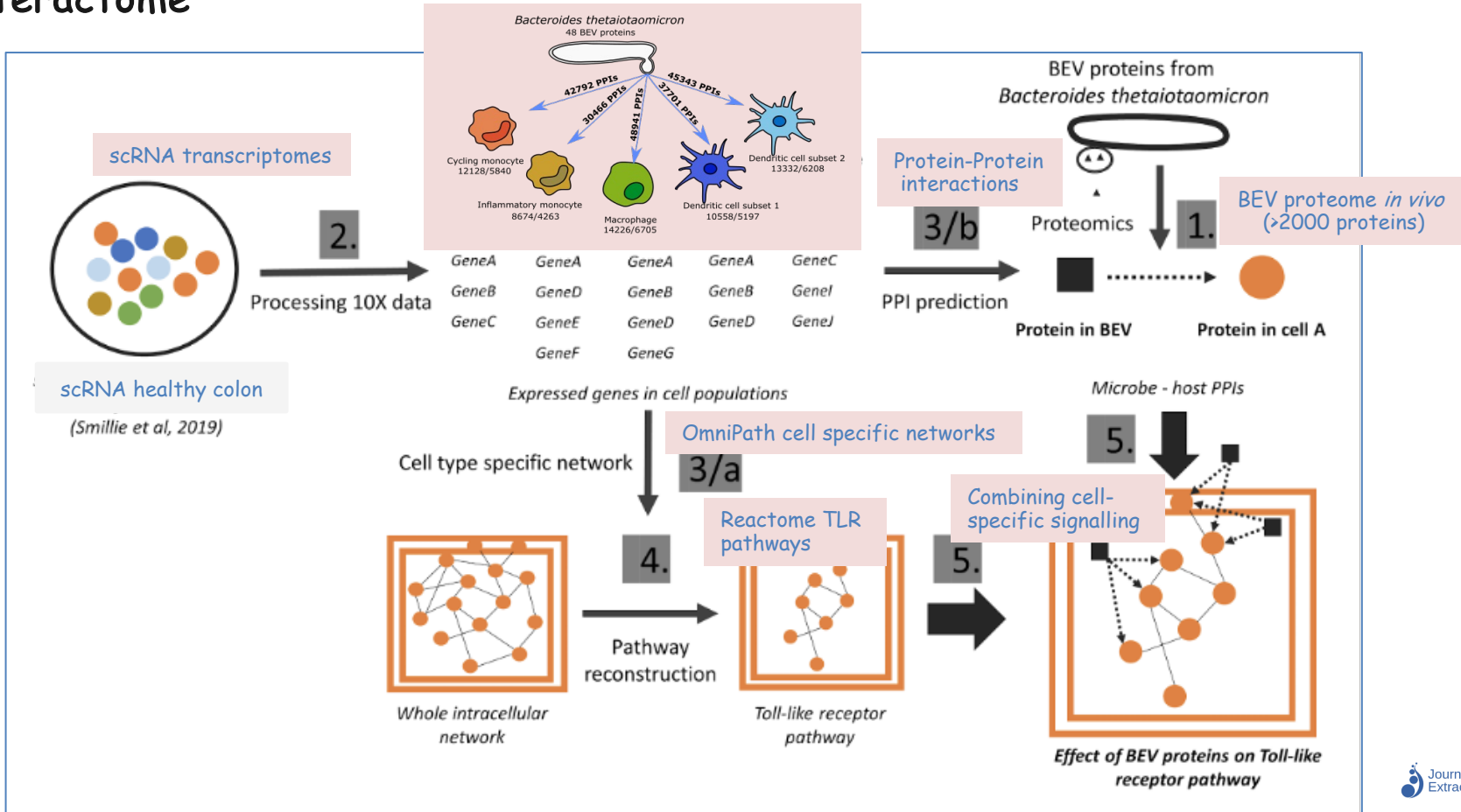


- DNase I treatment of plasma to eliminate extracellular DNA
- Major source of contamination is post-EV lysis – buffers and kits
- Sequence based filtering to remove protocol control reads >20% prevalence
  - ~15,000 Amplicon Sequence Variants
  - $743 \pm 586$  average reads/sample

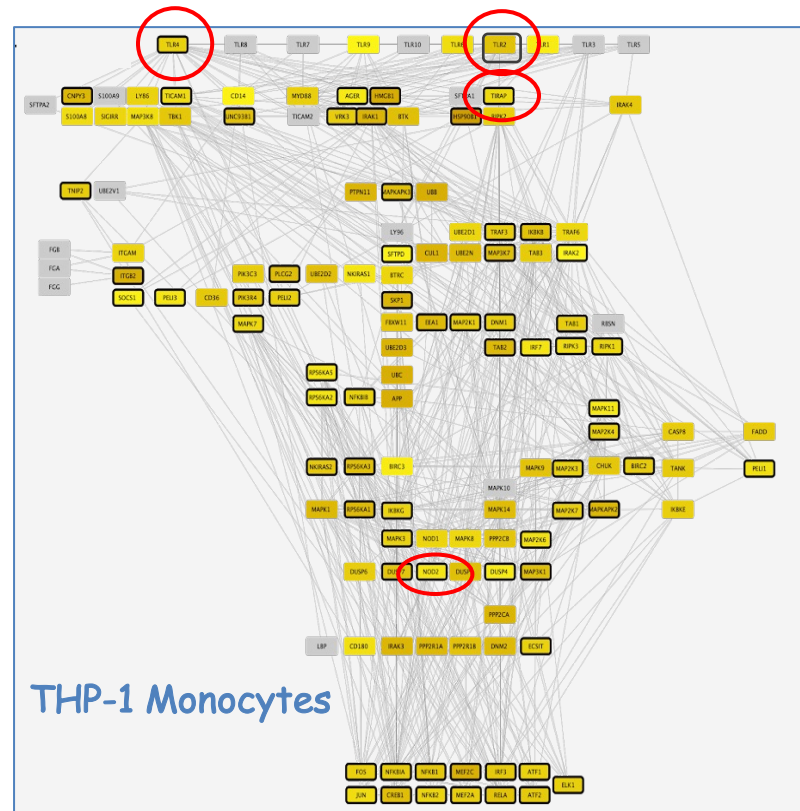
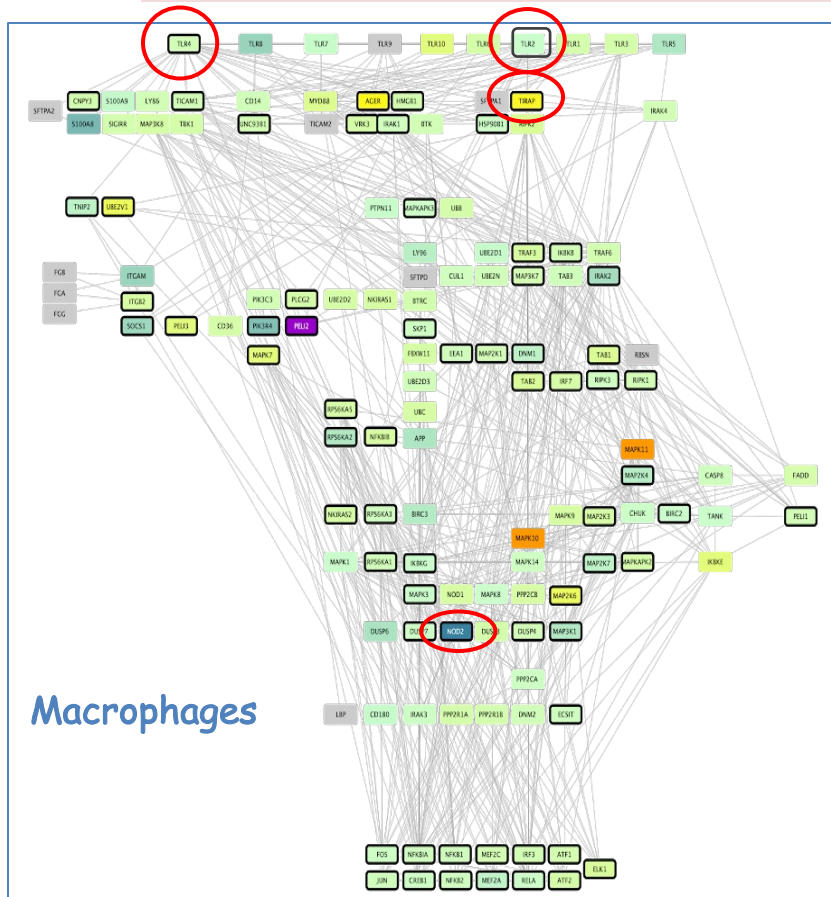
Mediators and  
outcome of BEV  
interactions  
with immune  
cells



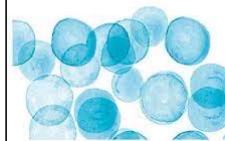
# A computational workflow to define the BEV-host cell Interactome



# TLR Pathway Analysis identifies specific PRRs as a BEV target in human macrophages

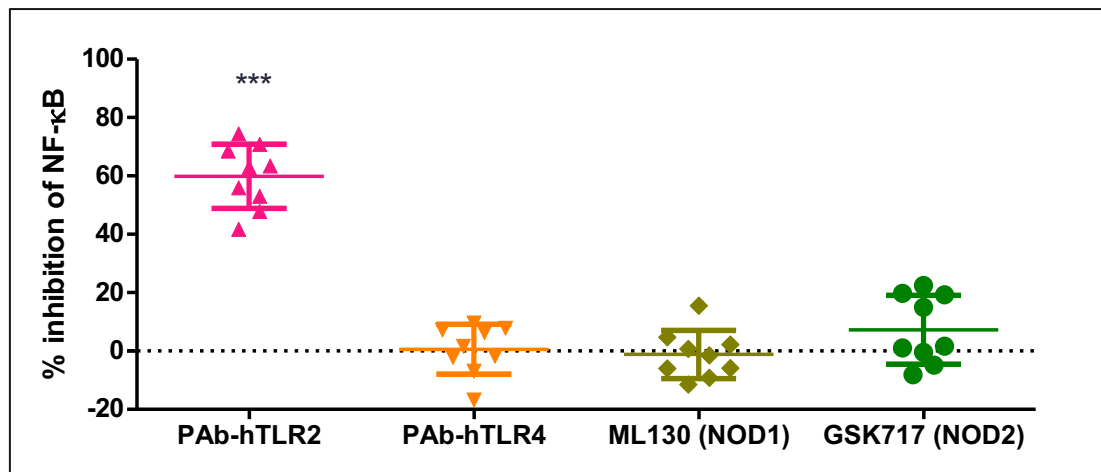
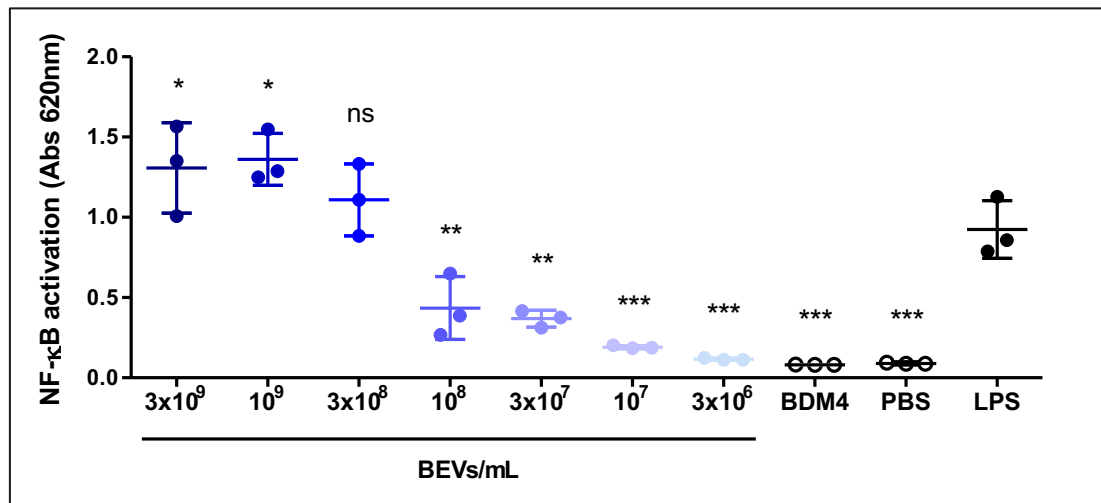


Macrophages: TLR2, TLR4 & NOD2 are candidates for mediating interactions with BEV proteins

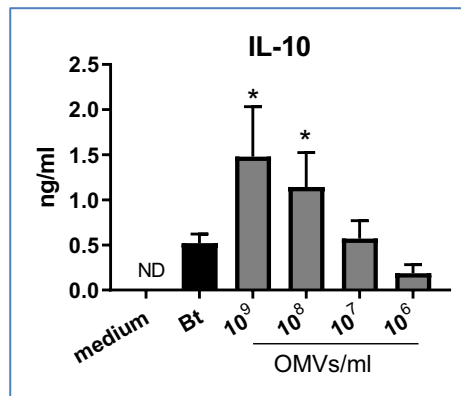
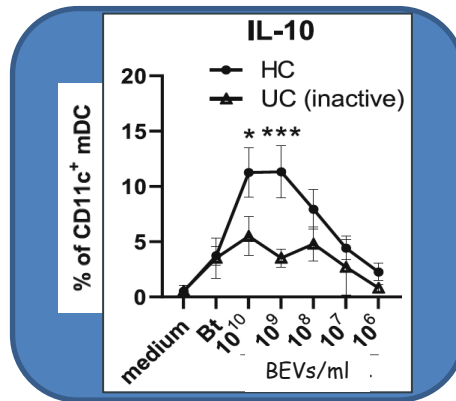
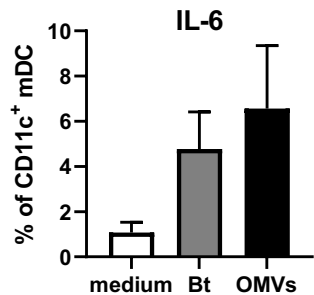
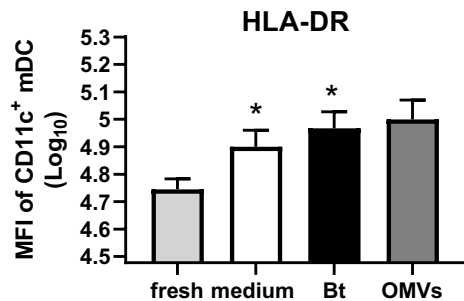
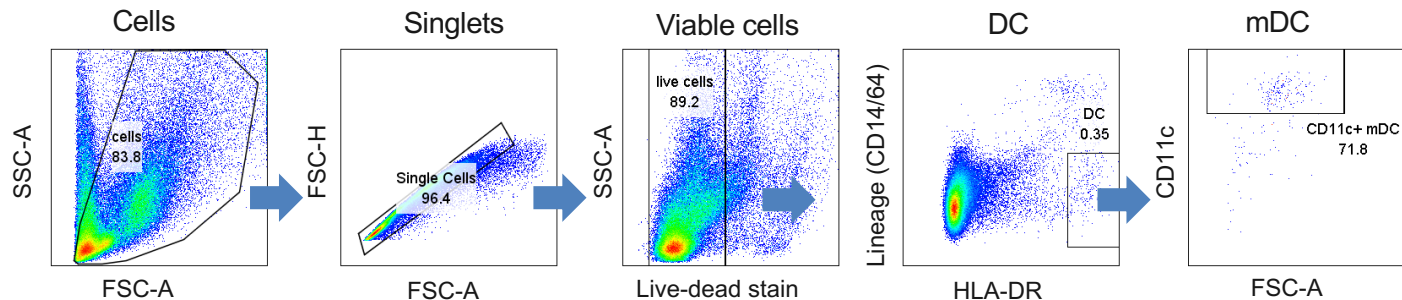


# TLR2-mediated activation of THP-1 Blue\* cells by Bt BEVs

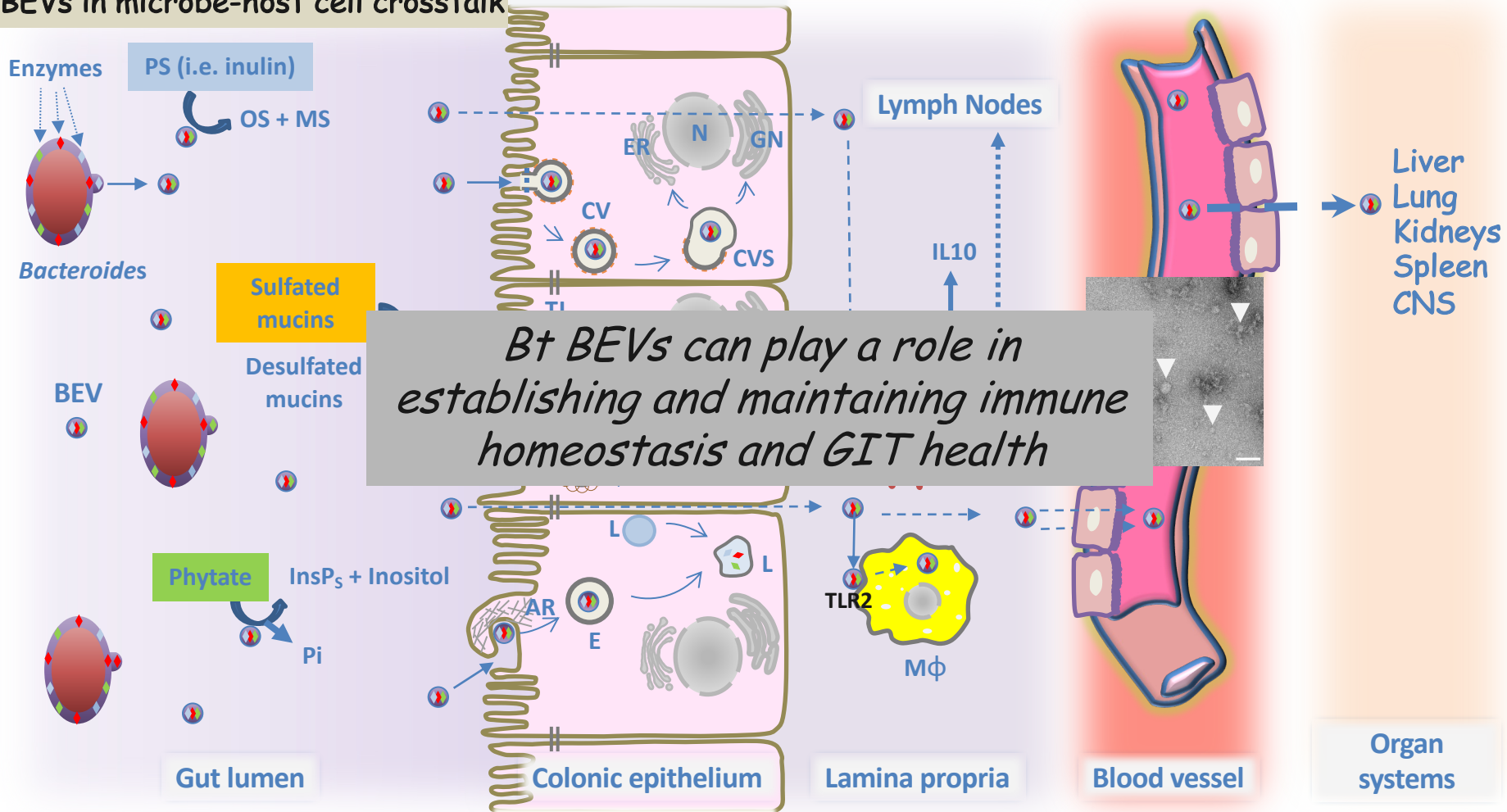
\* NF- $\kappa$ B-inducible secreted alkaline phosphatase (SEAP) reporter gene



# Bt BEVs stimulate production of immunoregulatory IL-10 from mDCs of healthy individuals but not those of IBD patients



# BEVs in microbe-host cell crosstalk





# TRANSLATION

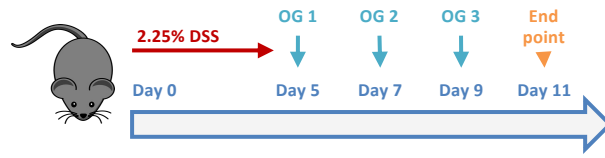
## Native BEVs

- Limiting/controlling inflammation – IBD (Fonseca et al., 2022, Front. Micro.)
- Cancer therapy
- Adjuvant – boosting natural immunity - Ageing

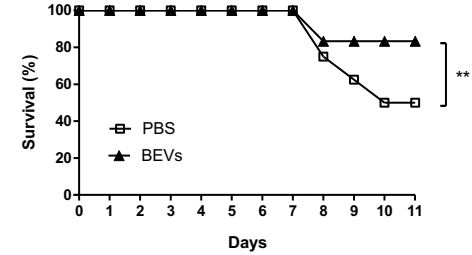
## Engineered BEVs

- Delivery of biologics
  - BEV-Hu.KGF2 for oral treatment of acute colitis (Carvalho et al., 2019, JEV)
- Mucosal vaccines
  - BEV-*Yersinia pestis* (pneumonic plague) oral/i.n. vaccine (Carvalho et al., 2019 Clin. Exp. Immunol.)
  - BEV-IAV (influenza virus) i.n. vaccine (Carvalho et al., 20219, JEV)
  - BEV-SARS-CoV-2 i.n. vaccine

A.

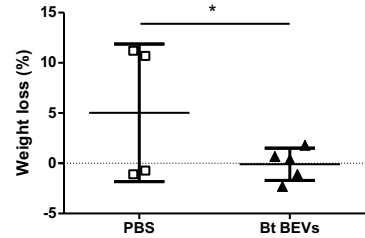


B.

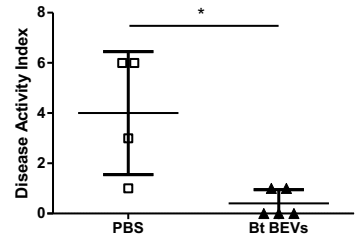


**Anti-inflammatory effect of Bt BEVs in a pre-clinical model of colitis**

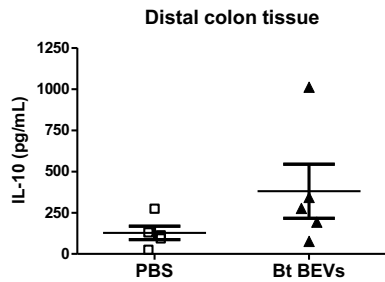
C.



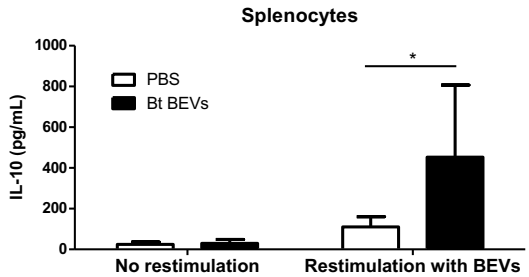
D.



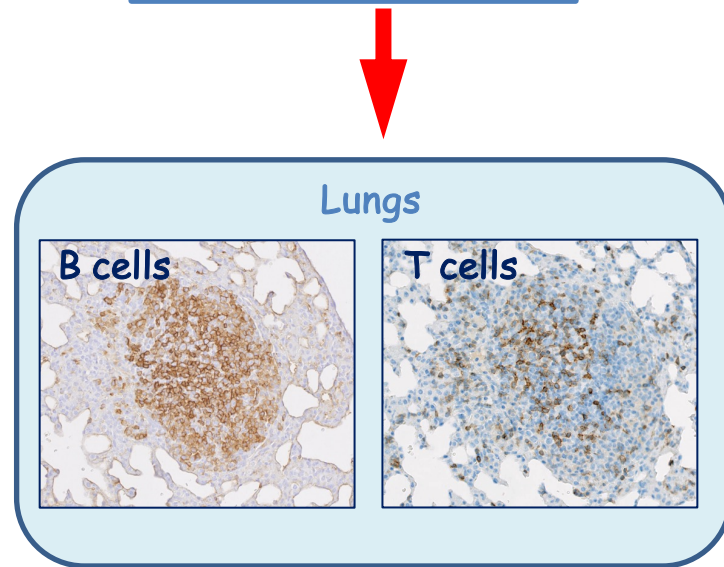
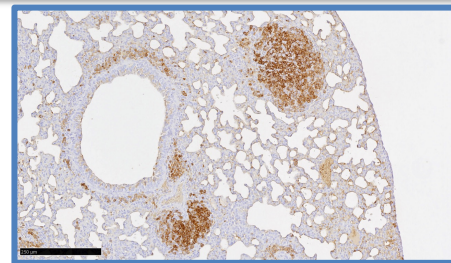
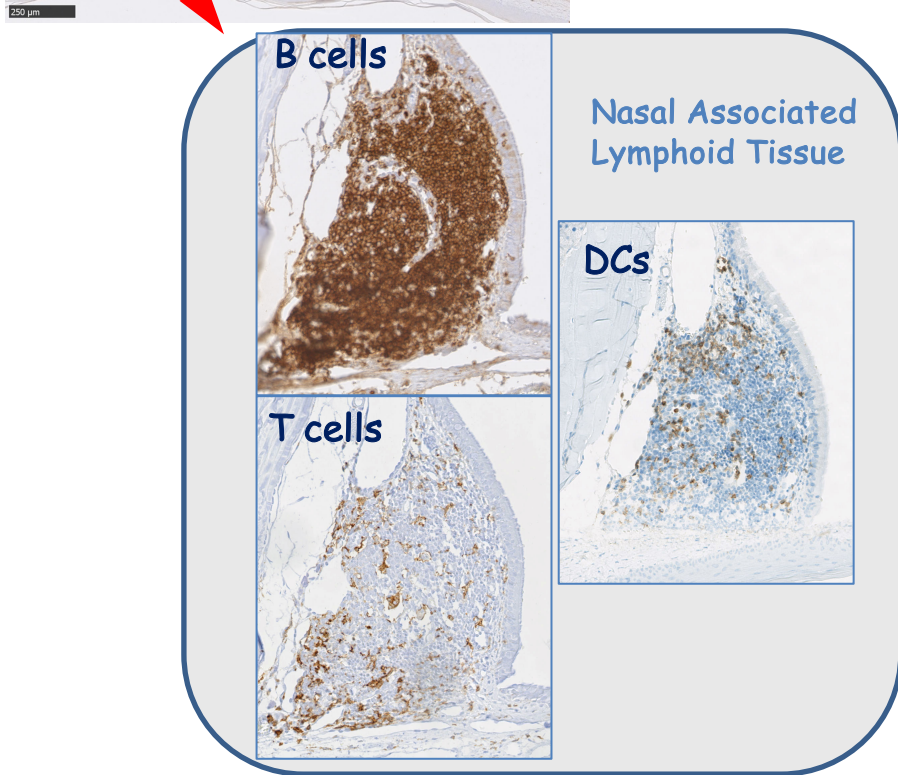
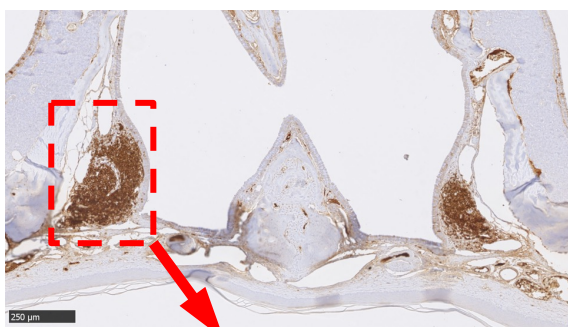
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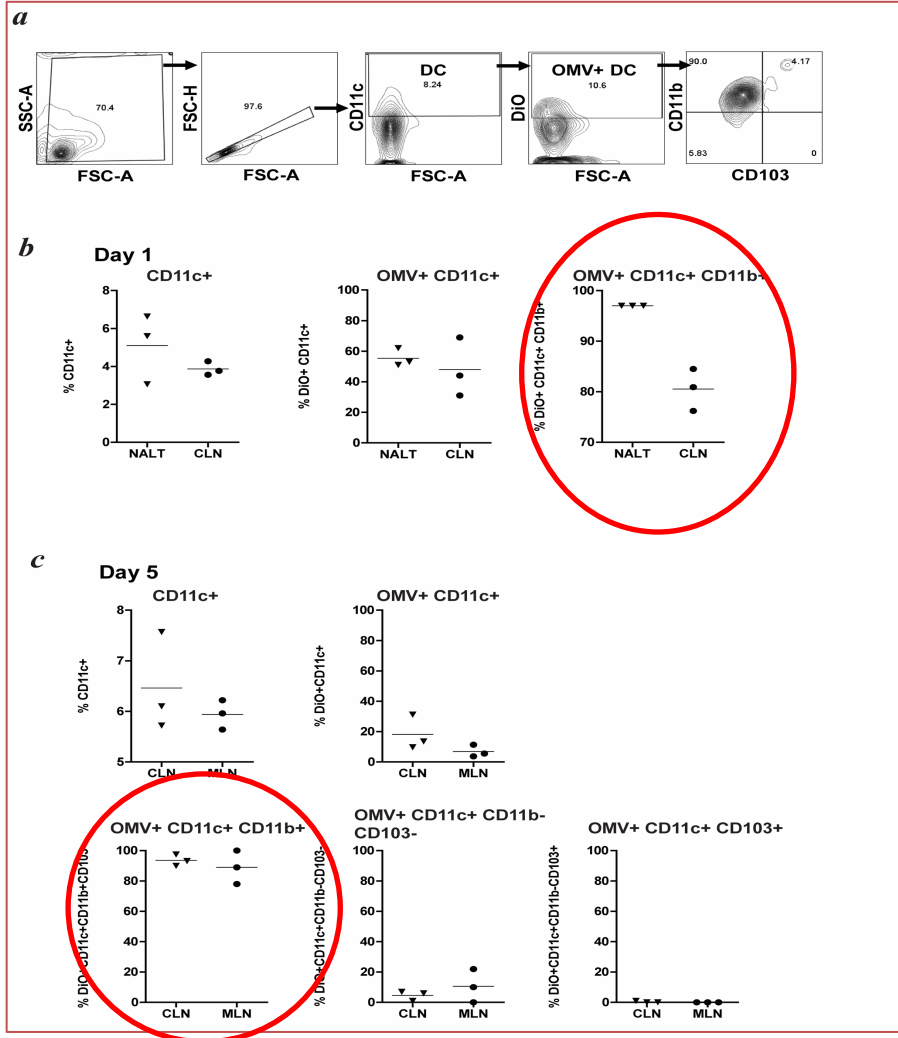
F.



# I.N. delivered native BEVs-mediate immune potentiation in the Upper and Lower RT



Rapid (<24h)  
 acquisition of native  
 Bt BEVs by  
 pulmonary-associated  
 DCs and trafficking  
 to cervical and  
 mesenteric lymph  
 nodes (CLN and MLN)  
 after intranasal  
 administration



# Bioengineered Bt BEVs for mucosal delivery of vaccines for respiratory pathogens



## Pneumonic plague

Use of bioengineered human commensal gut bacteria-derived microvesicles for mucosal plague vaccine delivery and immunization  
Carvalho AL., et al. 2019. 196:287-304.

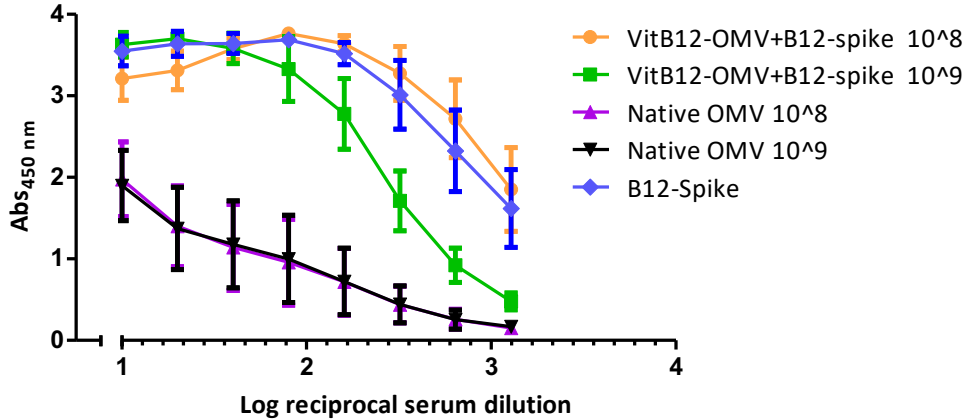


## Influenza

Bioengineering human gut commensal bacteria derived outer membrane vesicles for the delivery of biologics to the gastrointestinal and respiratory tract  
Carvalho AL., et al. 2019. 8:1632100.

# A multivalent Bt BEV SARS-CoV-2 mucosal vaccine

## Serum anti-Spike IgG



Drugtargetreview.com

## Plug-and-Play System

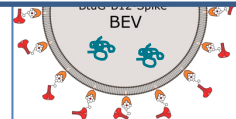
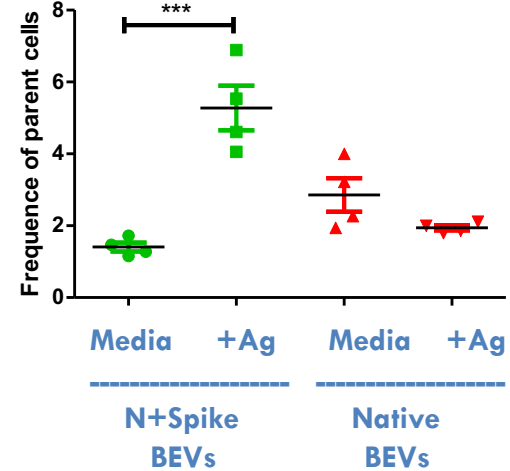
Based upon using the VitB12-B12-R (BtuG) high affinity covalent interactions to "label" BEVs

Isolation/purification rec. Spike proteins (HEK/CHO cells) + vitB12



Spike

## IFN $\gamma$ <sup>+</sup> N-specific CD8 T cells



# Summary: Reevaluating the Biological Role(s) of BEVs

Findings to date challenge established orthodoxies concerning mechanisms of symbiosis in the GI-tract

Roles for bacteria-generated microvesicles and BEVs in:

- Nutrition (e.g., phytate and vitamin metabolism)
- Inter-kingdom communication
- Host cell physiology - adaptation & immune homeostasis
- Ecology of the intestinal microbiota
- Susceptibility vs. resistance to infection (AMR)

# Exploiting BEVs and BEV Technology

## Native BEVs for:

- Biomarkers (Blood, SNF...)
- Immune-potential
- Adjuvant therapy
- Boosting natural immunity:
  - Immunocompromised
  - Immunosenescence
  - Infection - viral/bacteria
  - Cancer

## Engineered BEVs for mucosal & systemic delivery:

- Vaccines - human & animal
- Enzymes (replacement therapy)
- Antimicrobials (bacteriocins, anti-virals)
- Cytokines, hormones, neurotransmitters, QS mols
- Anti-tumor agents - angiogenesis inhibitors
- Neutralising antibodies (scFv, sdAb)
- Metabolites, anti-metabolites



# Acknowledgement



Régis Stentz  
Ana Carvalho  
Ariadna Clopés  
Anne Verkerk  
Khoon-S Kok  
Nikki Horn  
Udo Wegman  
Emily Jones

Aimée Parker  
Catherine Booth  
Sonia Fonseca  
Jo Brooks  
Nadezha Gicheva  
Rokas Juodeikis  
Amisha Modasia  
Anne VerKerk

Dimitris Latousakis  
Arlaine Brion  
Andy Goldson



Tamas Korcsmaros  
Leila Gul  
Dezso Modos

Matthew Madgwick  
John Thomas  
Paddi Sudhakar

Martin Warren



Mark Smales

Maria Stanley, Emi Nemoto-Smith, Dave Beal,  
James Budge, Mingzhi Liang, Ian Brown,  
Kevin Howland, Phoebe Lee



THE UNIVERSITY  
of LIVERPOOL

James Stewart  
Eleanor Bentley



Universität  
Zürich<sup>UZH</sup>

Anjar Kipar



The  
University  
Of  
Sheffield.

Colin Bingle  
Lynn Bingle



Imperial College  
London

Stella Knight  
Alistair Noble  
Lydia Durrant



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*Thank You*

