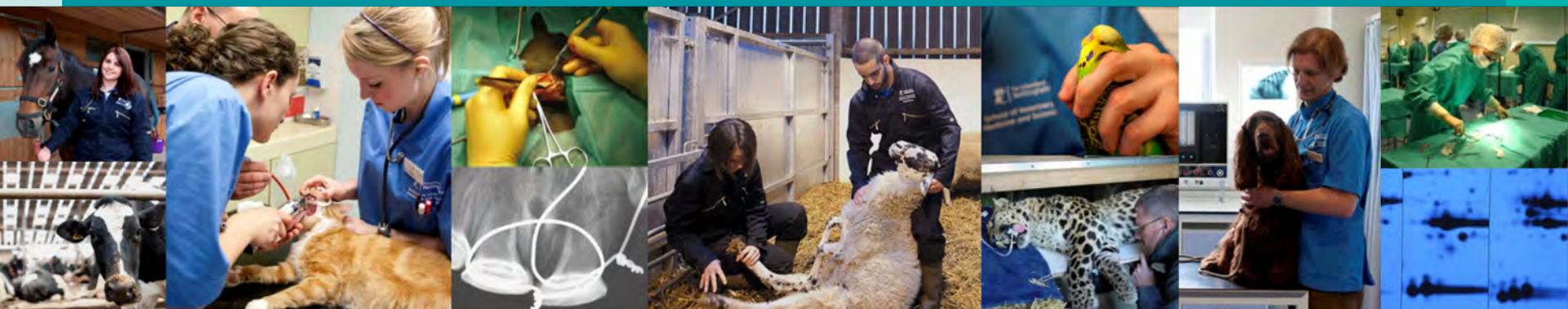




DISCOVERY OF ANTIBODIES TO *EIMERIA* INFECTIONS IN POULTRY USING NEXT GENERATION PHAGE DISPLAY

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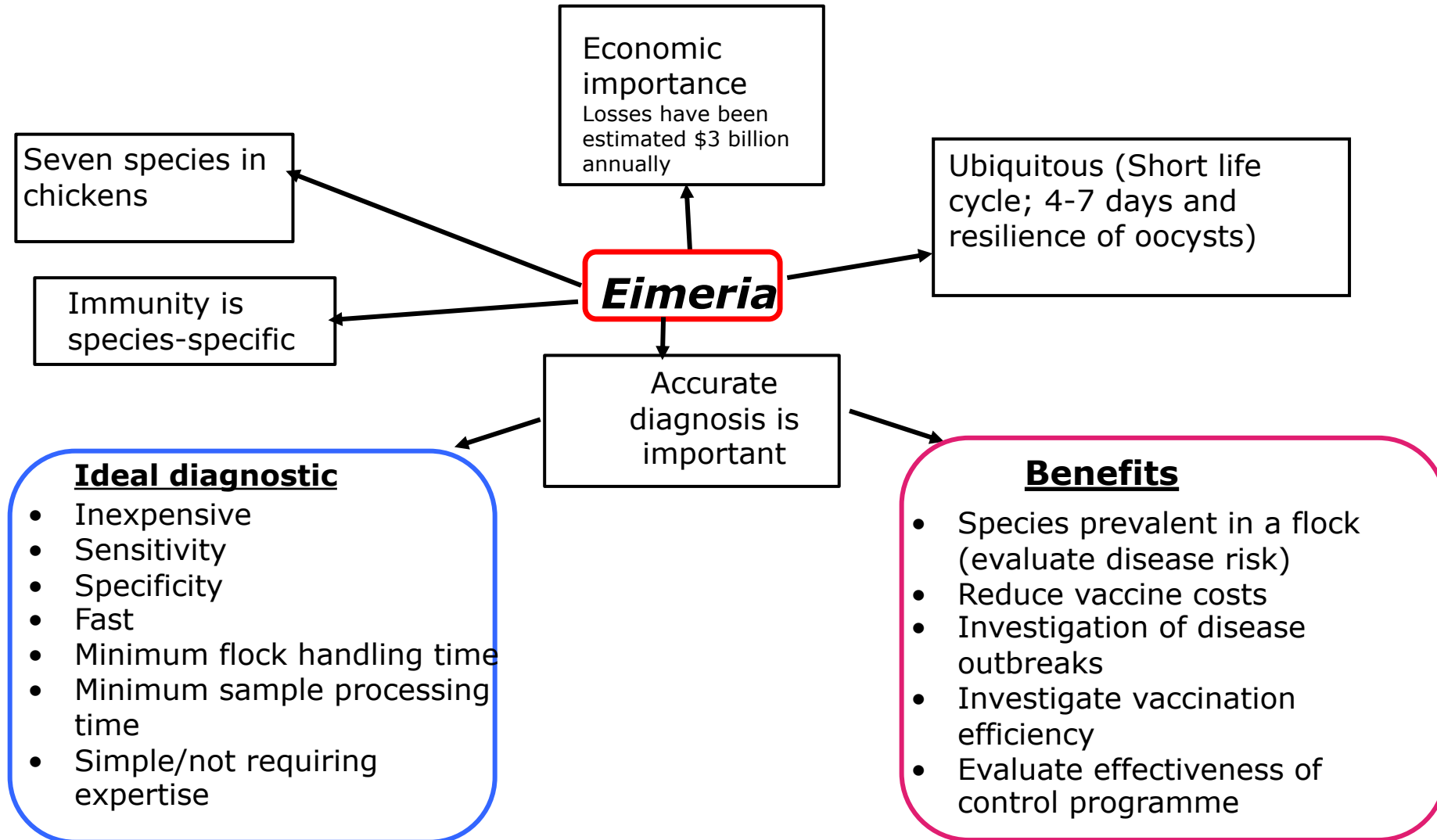


Outline

- Terms
- Introduction
- Hypothesis
- Aims and Objectives
- Methods
- Results
- Summary

- 1) Phagemid: Cloning vector that has plasmid and bacteriophage properties used in phage display
- 2) ScFv (Single chain fragment variable): fusion protein of the variable regions of the heavy and light chains of immunoglobulins. This protein retains the specificity of the original immunoglobulin, despite removal of the constant regions
- 3) HCDR3: Heavy chain complementarity determining region 3 ;part of immunoglobulin and most variable region

Introduction



Introduction

Phage display

Conventional phage display

- Handpicking of clones
- Traditional phage panning: Screen ~100 ligands by ELISA against a single target

- G. Smith (1985)
- Display peptides and proteins on phage
- Links genotype and phenotype through the phage coat protein
- Hugely successful for the development of monoclonal antibodies
- Starts with antibody library preparation
- Affinity selection technique

Next generation phage display

- Deep sequencing of genes
- NGPD: Screen 10,000's ligands against multiple targets

Hypothesis

- Next generation phage display will allow comparative screening of antibody binding events to identify antibodies to species-specific epitopes of *Eimeria* oocysts. Such ligands can then be used to develop immunoassays to differentiate infecting *Eimeria* species of poultry.

Aim

- To isolate antibodies to oocysts to develop in-field species-specific diagnostic assays for *Eimeria* infection

Using Next generation phage display technology [NGPD]

Objectives

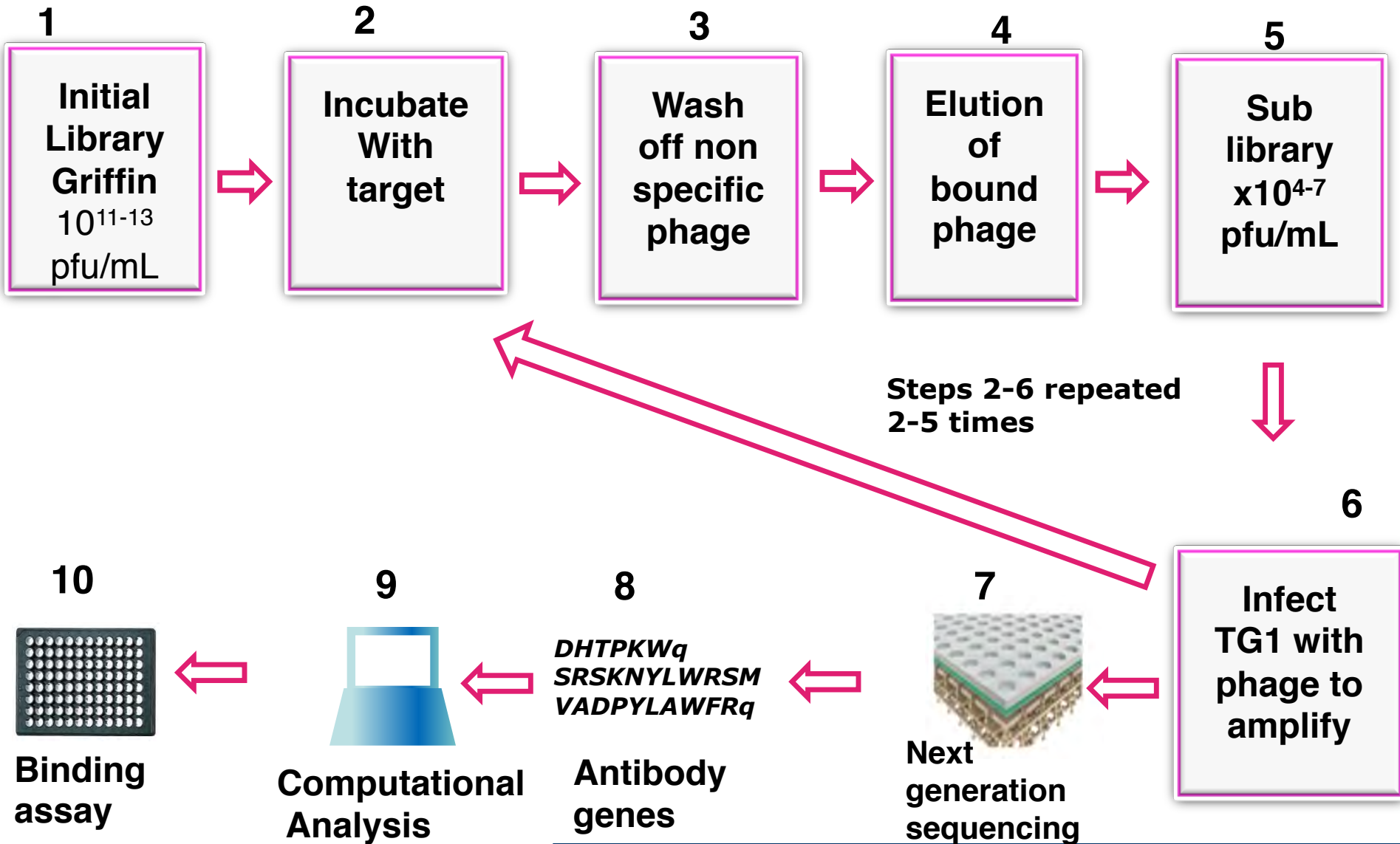
- To carry out antibody-phage display panning against oocysts
 - Isolate antibodies against oocysts of all seven *Eimeria* spp
 - To screen the antibodies isolated via ELISA
- To compare traditional phage panning with Next generation phage display (NGPD)

Methods(NGPD)



The University of
Nottingham

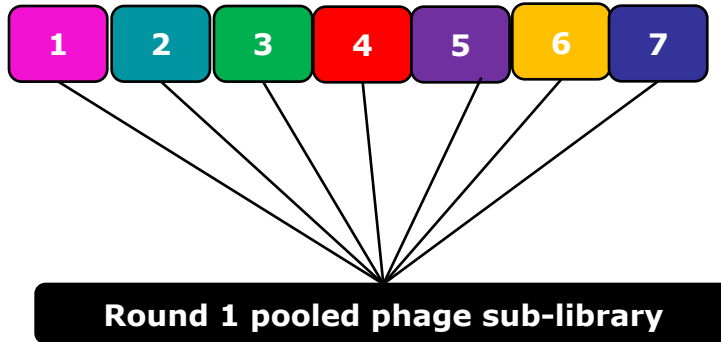
UNITED KINGDOM • CHINA • MALAYSIA



- The project began with oocysts
 - Sourcing of oocysts (MSD Animal health)
 - Counting of oocysts (7.0×10^5 - 6.60×10^6 /ml)
- Griffin Phage library grown 10^{12} PFU/ml
- Target antigen Oocysts 10^4
- Panning experiments with seven species with Griffin library
 - Five rounds of panning, 5 phage and Griffin library(41 samples)

Biopanning strategy

Round 1



1-7 represent the seven species of *Eimeria*

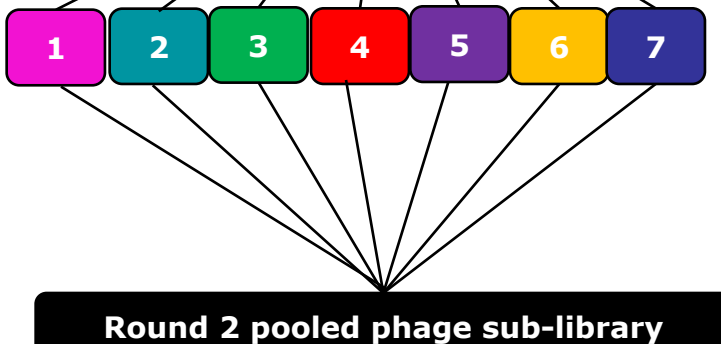
1st round of panning;
Antibody library was
panned against
Individual species of
Eimeria

Elution by PH shift

A total of 5 rounds of
biopanning was carried
out

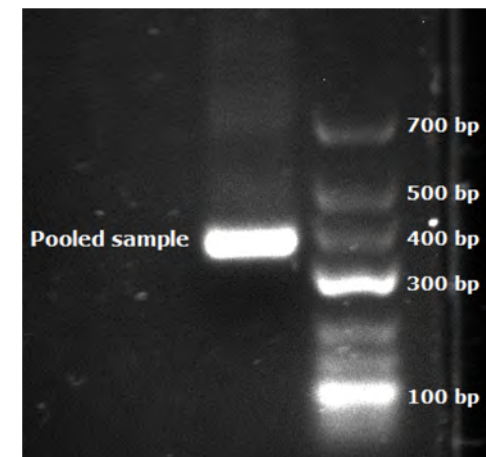
The HCDR3 region
was amplified

Round 2



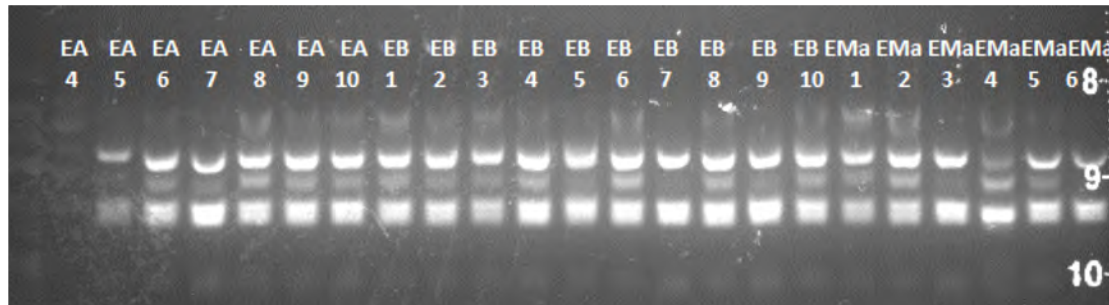
Barcode
identifiers

PCR
amplicon



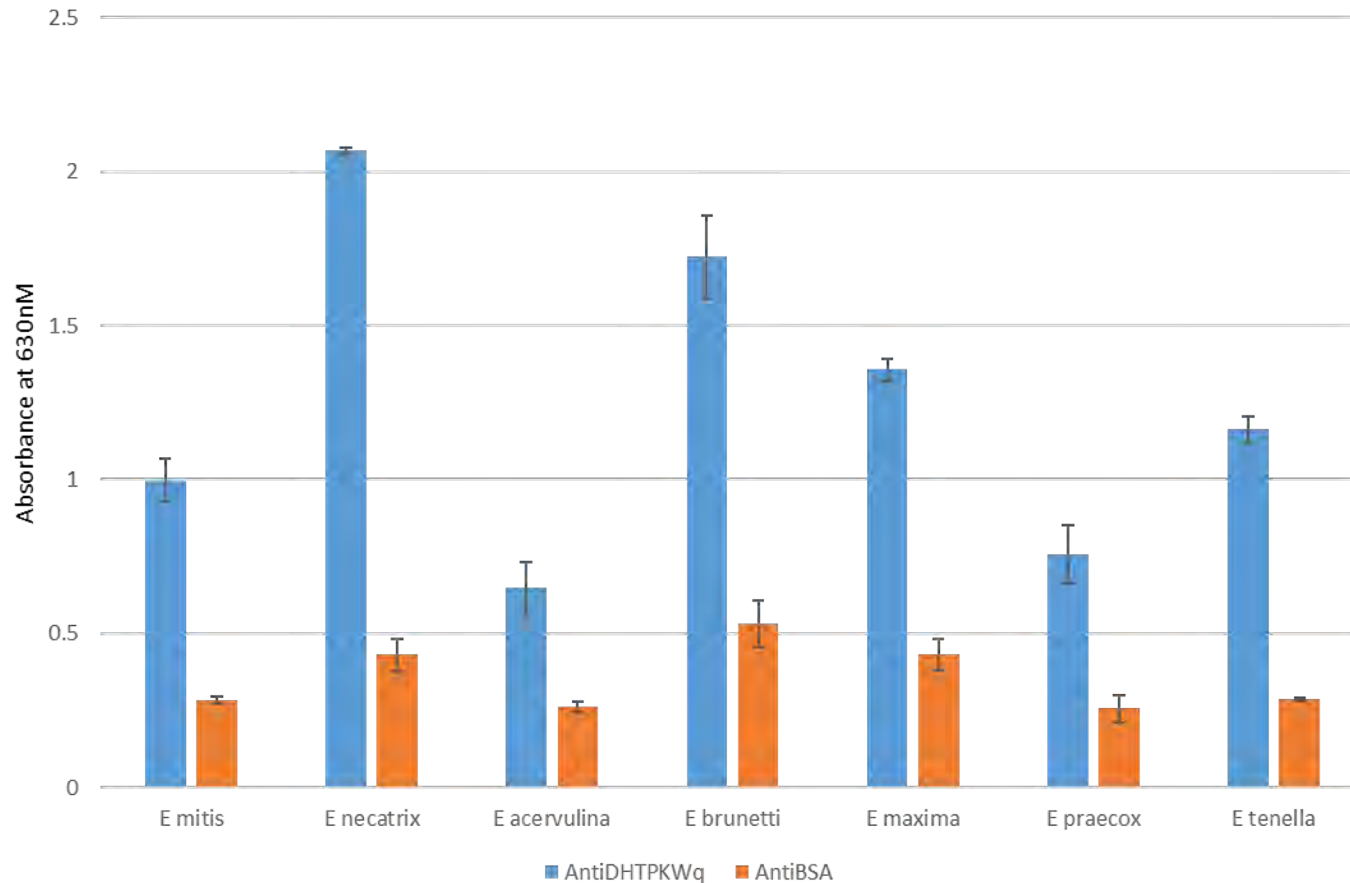
While waiting for NGPD data...

- 10 single clones were picked from each species (*round 5 panning*)
- Similar restriction enzyme digest pattern

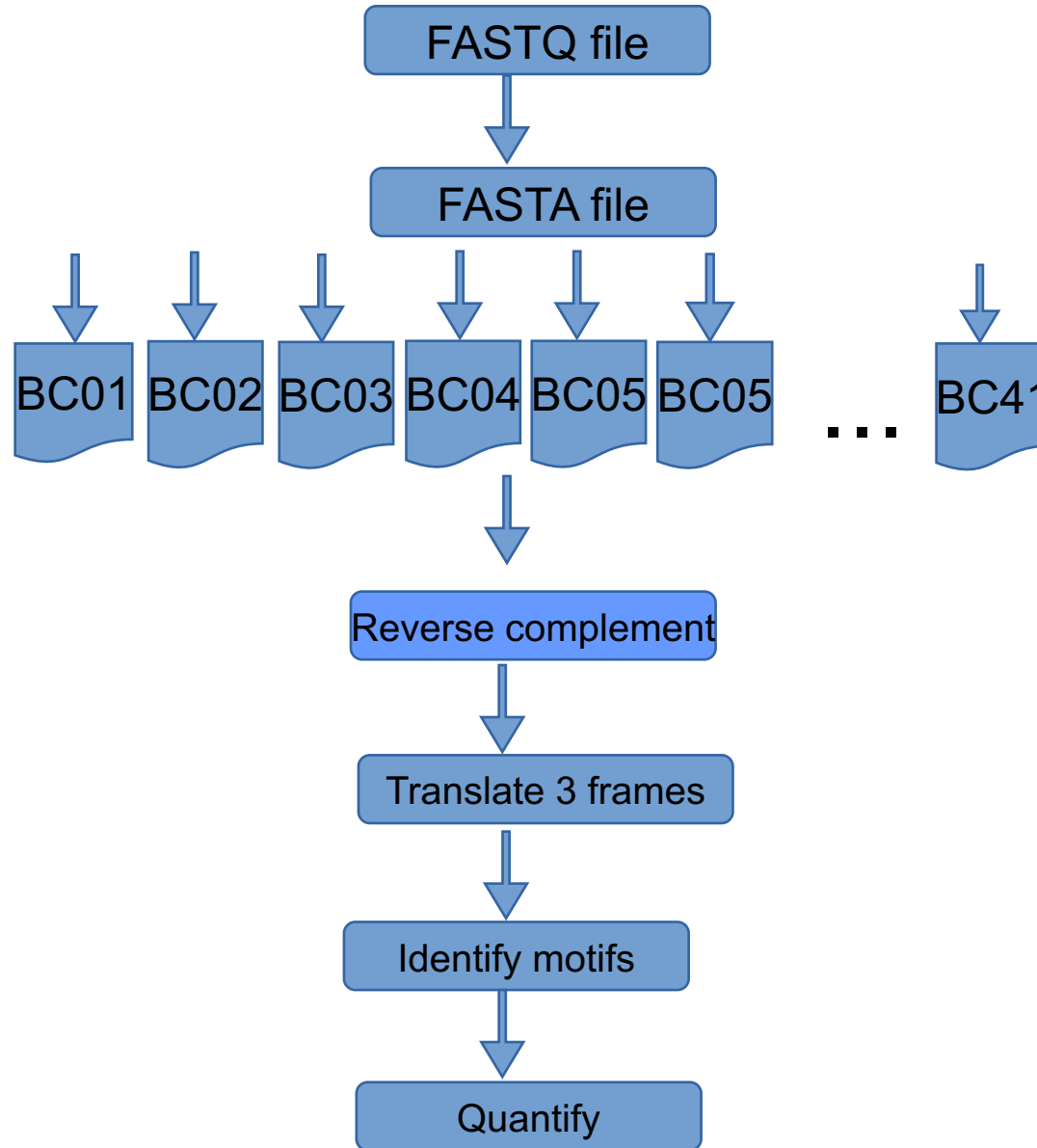


- 1/3rd (23 clones) randomly picked clones were Sanger sequenced DHTPKWq
- Screened via ELISA

Traditionally picked scFv DHTPKWq with AntiBSA phage ELISA



NGPD data analysis



Results

Criteria for selection

- Z scores (ratio and frequency of binding). They define relative enrichment
- Z score > 5
- At least 5 reads
- Enriched in at least 2 panning rounds
- Enriched in selection Phase not only growth

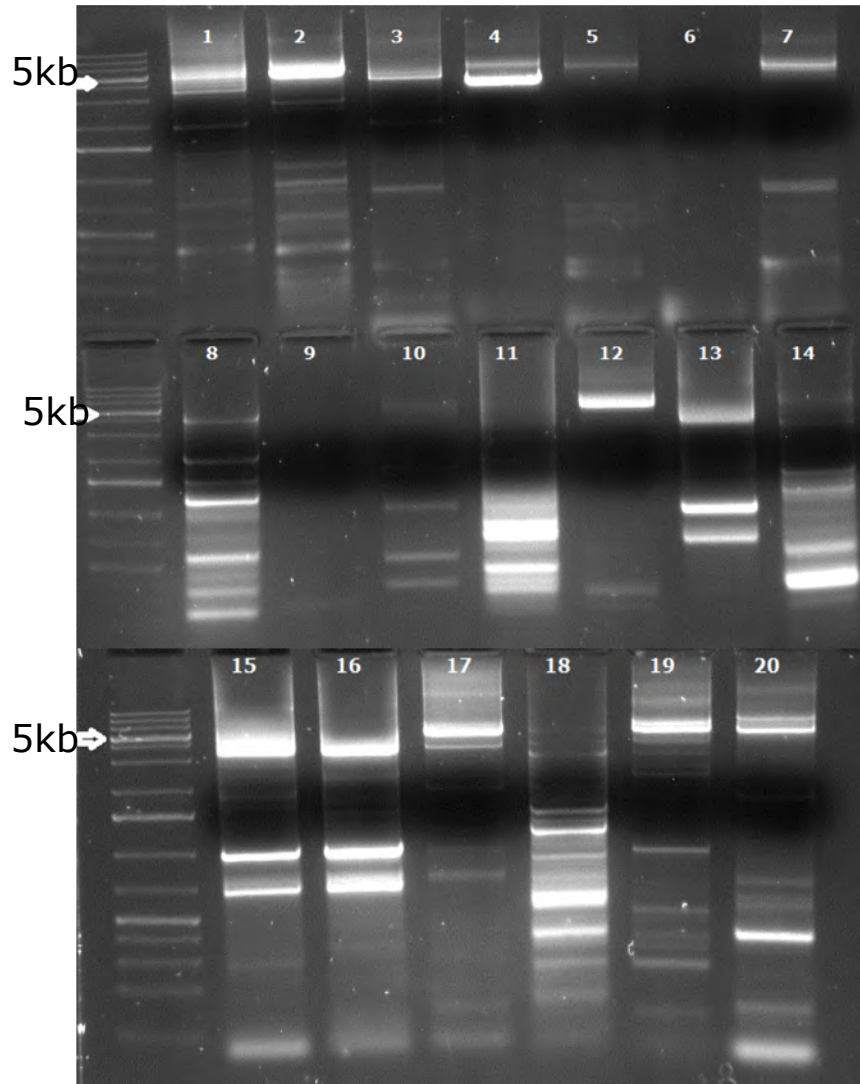
Selected scFv

- 1 AWTVNQTTNY
- 2 DGKPPSG
- 3 DRSYTACDTSK
- 4 DKSSqG
- 5 EDRqCKHG
- 6 ESSqMNS
- 7 VAADFDY
- 8 YIRYTSPV
- 9 STQHP
- 10 SSDAP
- 11 SPRLPFDY
- 12 QTGMN
- 13 LMVYKH
- 14 KDVGVVVPK
- 15 IHRMVYL
- 16 HPMLGGLQGK
- 17 SRSKNYLWRSM
- 18 RDLTqDMWR
- 19 TTALP
- 20 SAqHP

From the DNA sequence, inverse PCR primers were designed to recover that phagemid from the panning round with the highest frequency of the clone



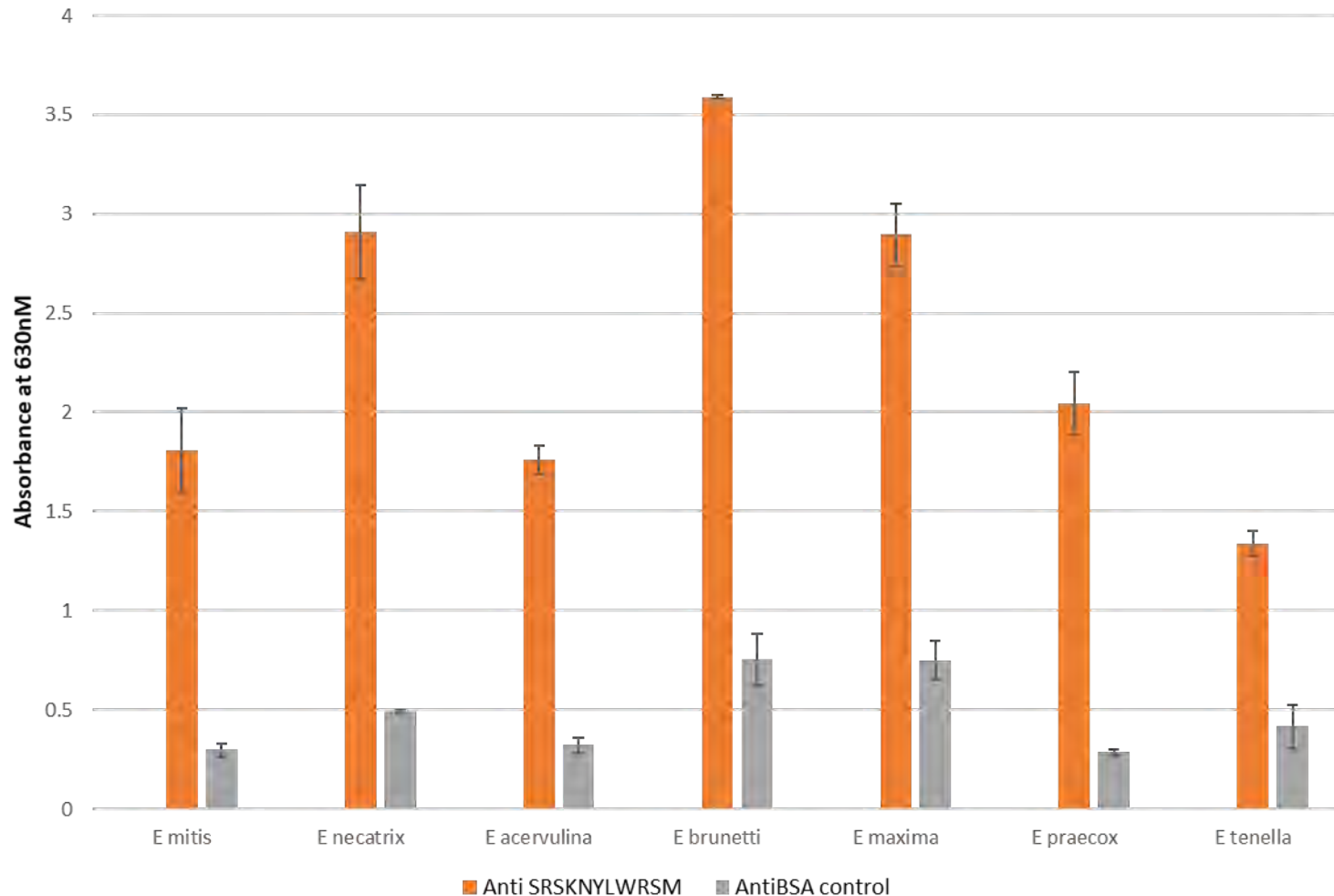
Inverse PCR (phagemid rescues)



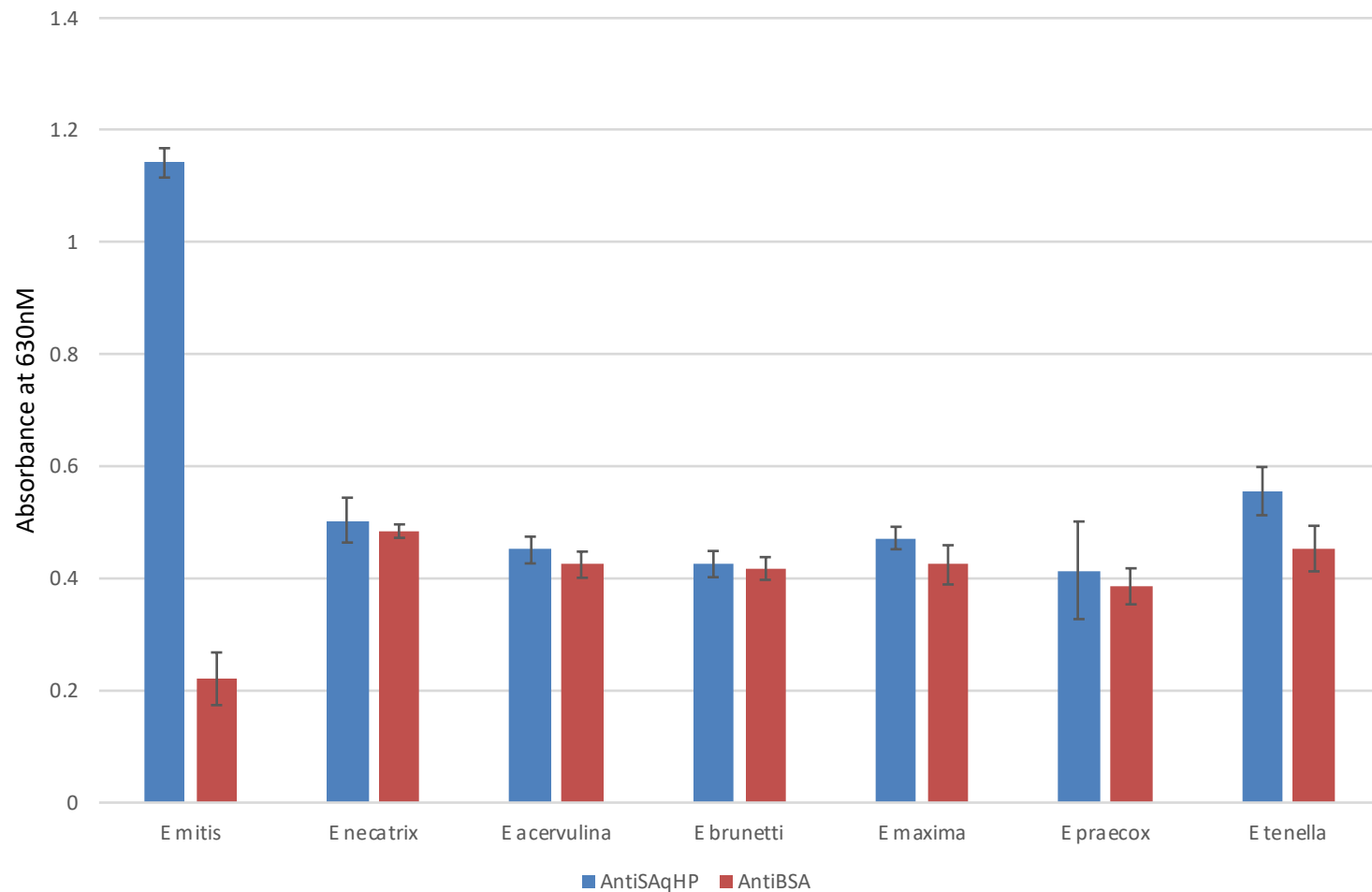
- 1 AWTVNQTTNY
- 2 DGKPPSG
- 3 DKSSqGT
- 4 DRSYTACDTSK
- 5 EDRqCKHG
- 6 ESSqMNS **failed**
- 7 VAADPFDY
- 8 YIRYTSPV
- 9 STQHP **failed**
- 10 SSDAP **failed**
- 11 SPRLPFDY **failed**
- 12 QTGMN
- 13 LMVYKH
- 14 KDVGVPK **failed**
- 15 IHRMVYL
- 16 HPMLGGLQGK
- 17 SRSKNYLWRSM
- 18 RDLTqDMWR **failed**
- 19 TTALP
- 20 SAqHP

**12 were
recovered**

Rescued scFv AntiSRSKNYLWRSM with AntiBSA control phage ELISA

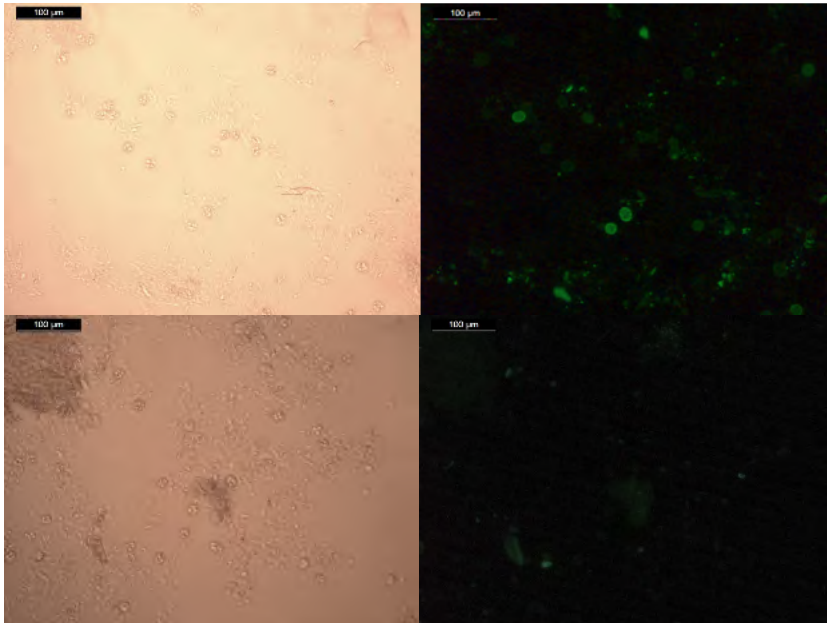
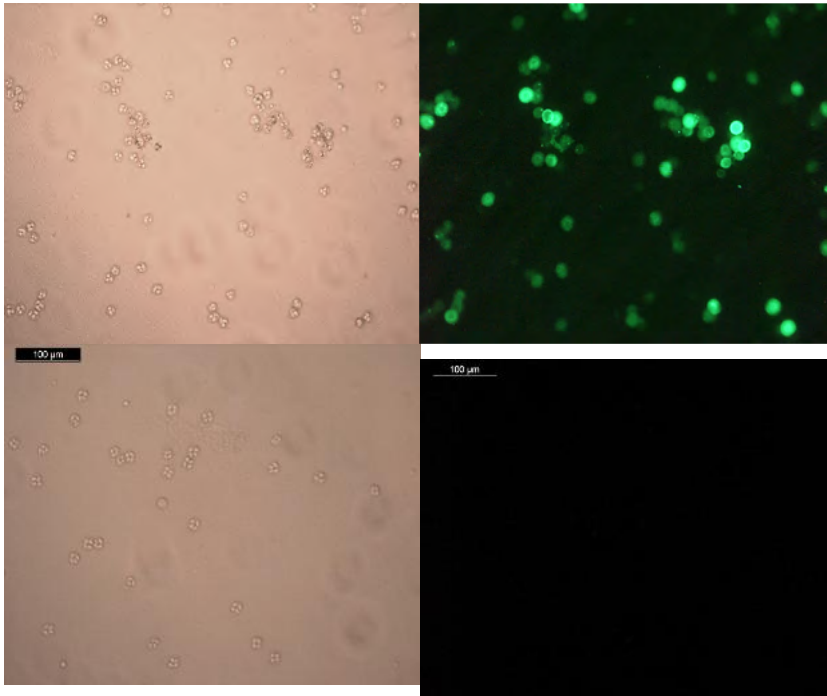


Rescued scFv AntiSAqHP with AntiBSA control phage ELISA



AntiSRSKNYLWRSM phage

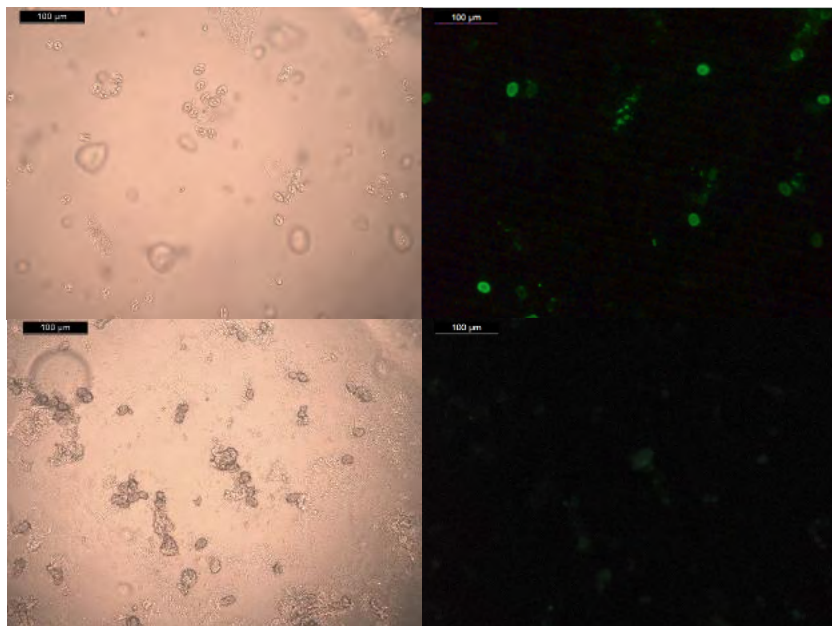
Eimeria mitis brightfield and
fluorescent images above
and BSA phage brightfield
and fluorescent control
below



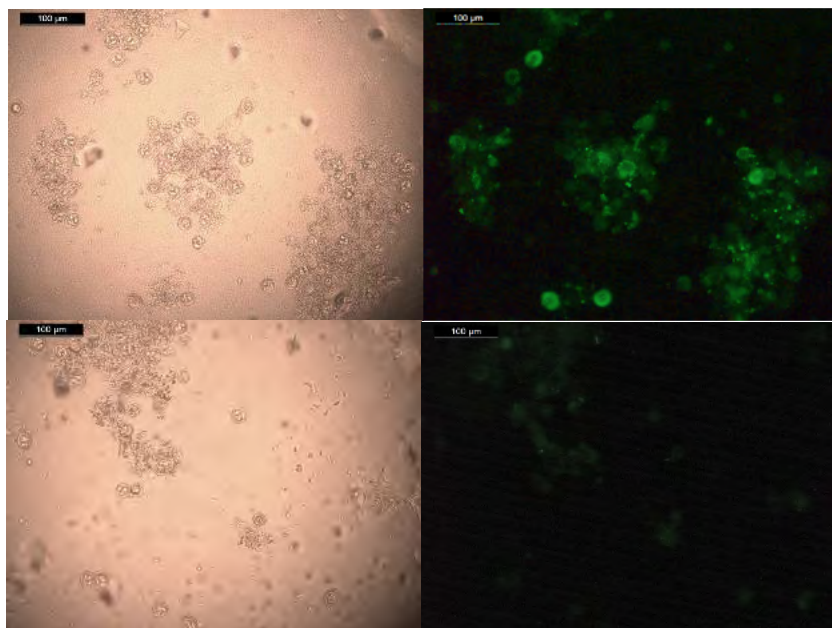
Eimeria necatrix
brightfield and fluorescent
images above and BSA
phage brightfield and
fluorescent control below

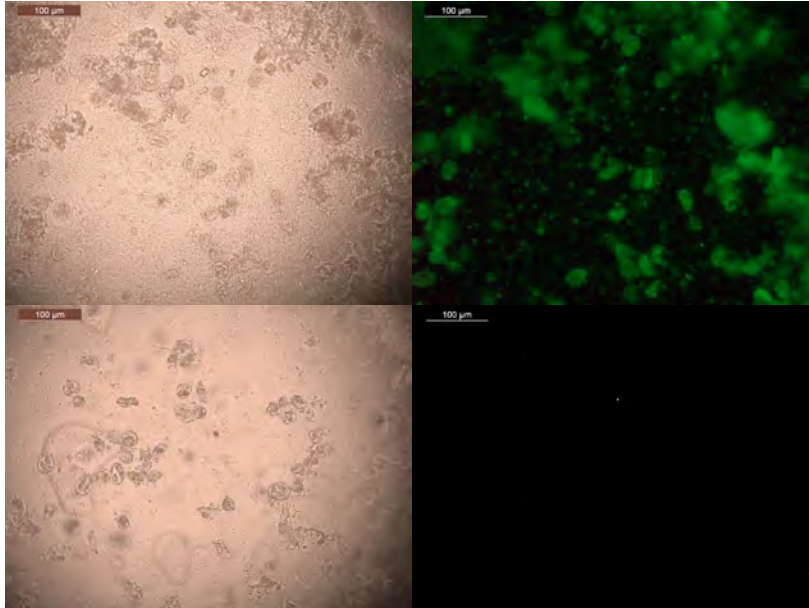


Eimeria acervulina brightfield and
fluorescent images above and BSA
phage brightfield and fluorescent
control below

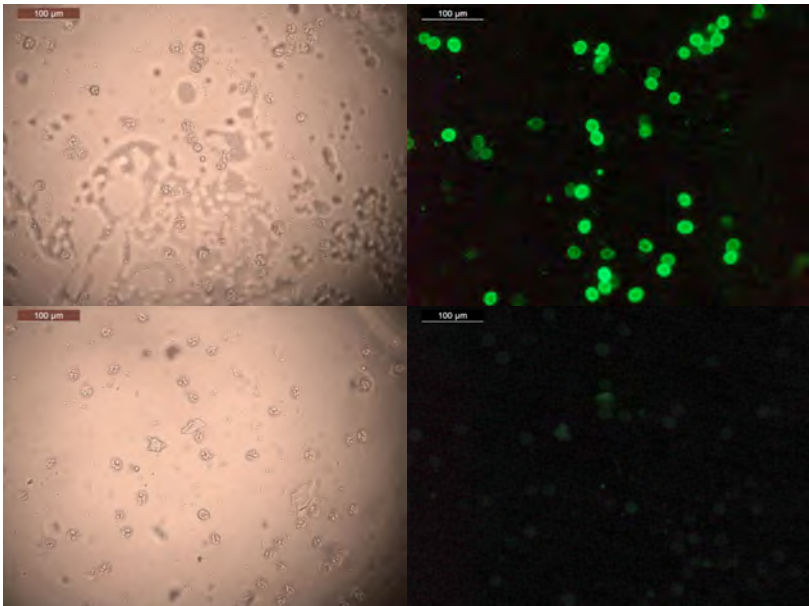


Eimeria brunetti brightfield and
fluorescent images above and BSA
phage brightfield and fluorescent
control below

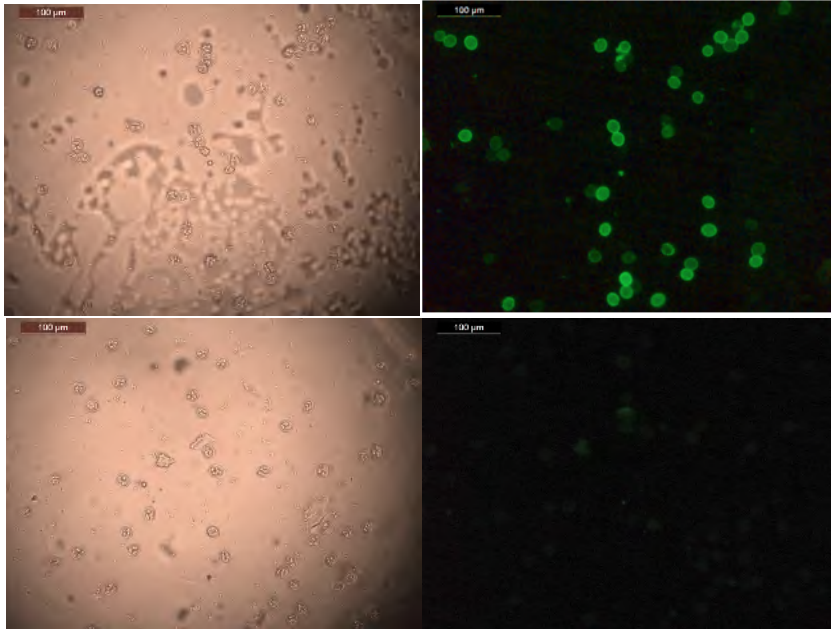




Eimeria maxima brightfield
and fluorescent images
above and BSA phage
brightfield and fluorescent
control below



Eimeria praecox brightfield
and fluorescent images above
and BSA phage brightfield and
fluorescent control below



Eimeria tenella brightfield
and fluorescent images
above and BSA phage
brightfield and fluorescent
control below

Summary

Objective 1

Isolate antibodies against oocysts of all seven *Eimeria* species

Objective 2

To compare traditional phage panning with Next generation phage display (NGPD)

- We isolated two antibodies against all seven *Eimeria* species
- We isolated an antibody against *Eimeria mitis*
- More antibodies are currently in development
- We found NGPD better because true binders can easily be missed by phage that have growth advantage

Acknowledgements

- Kevin Gough (University of Nottingham)
- Richard Emes (University of Nottingham)
- Ben Maddison (ADAS Biotechnology)
- Jon Owen
- Keith Bishop
- Claire Baker
- Rob Workman
- Rose Reader

Thank
you

University of Agriculture Makurdi



4/1/19