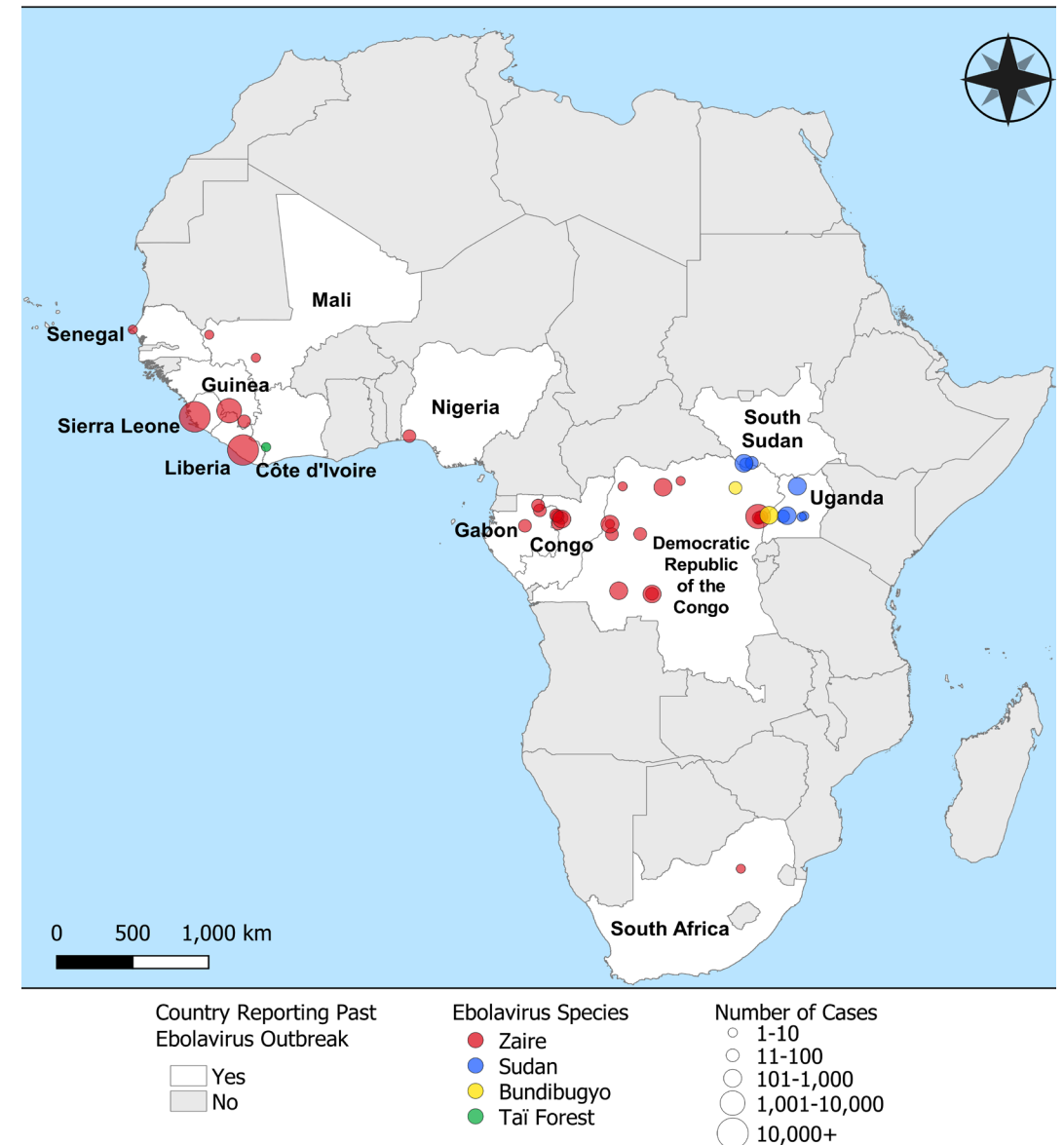


# Regulatory approval of Zabdeno/Mvabea

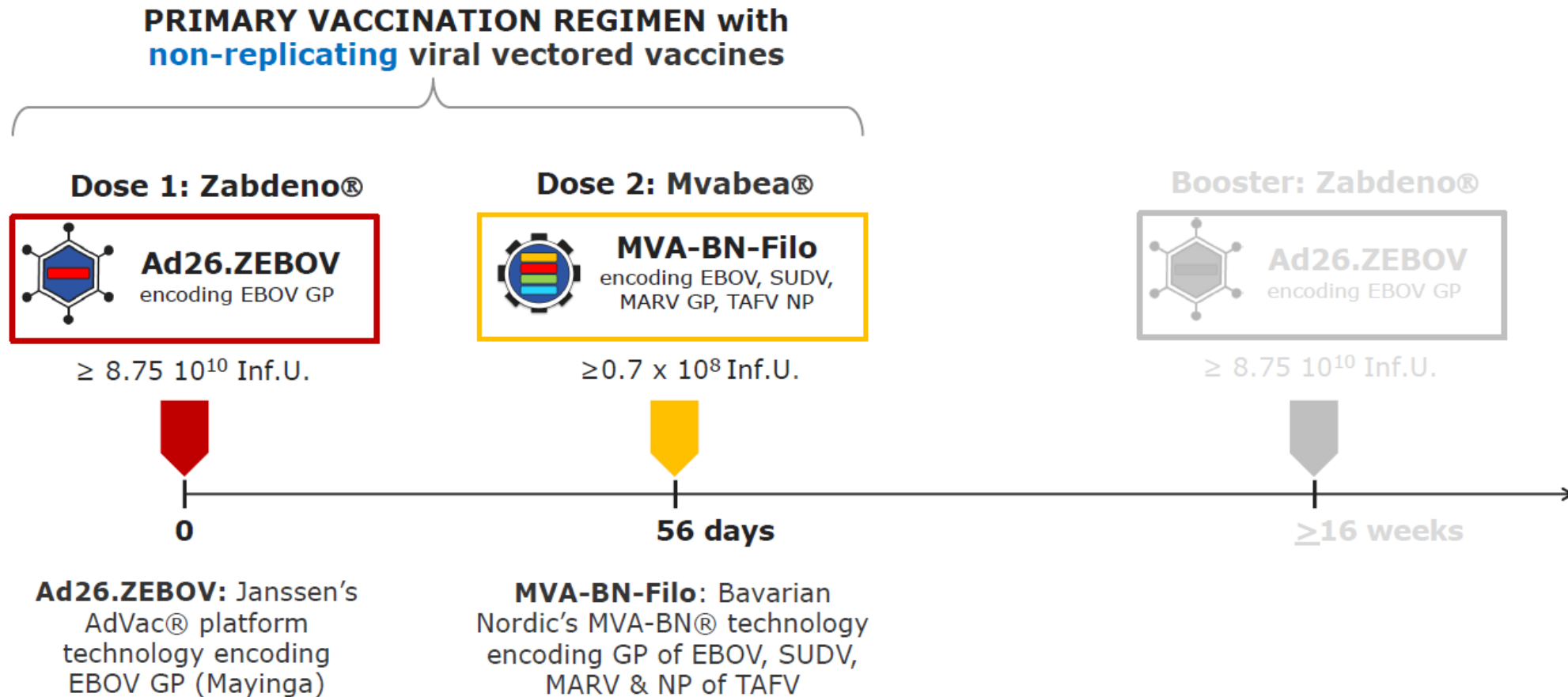
Jenny Hendriks  
Global Lead Clinical Immunology  
24nov25

# History of Ebolavirus outbreaks

- Zaire ebolavirus: 23 reported outbreaks since 1976
  - The most fatal Ebola virus.
  - 2014- 2016 outbreak in West Africa (largest Ebola outbreak to date with more than 28,600 cases)
  - 2018- 2020 outbreak in DRC (2<sup>nd</sup> largest outbreak)
- Two approved vaccines, one reactive and prophylactic (rVSVZEBOV, Merck) and one prophylactic (Ad26.ZEBOV, MVA- BN- Filo, Janssen)
- Efficacy data available for rVSV- ZEBOV, immunobridging data available for Ad26.ZEBOV, MVA- BN- Filo
- Binding antibodies strongly correlated with survival in NHP. Used in immunobridging to estimate vaccine efficacy of the **Ad26.ZEBOV, MVA- BN- Filo** regimen and is the basis for licensure in the EU

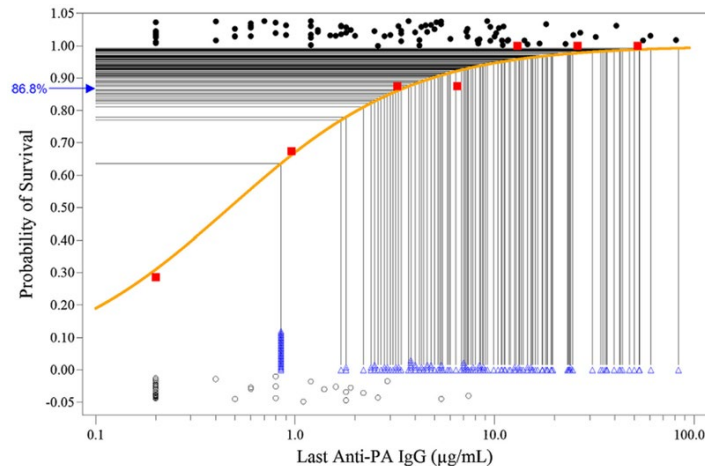


# The 2-dose primary regimen is administered approximately 8 weeks apart



# Why an immunobridging approach?

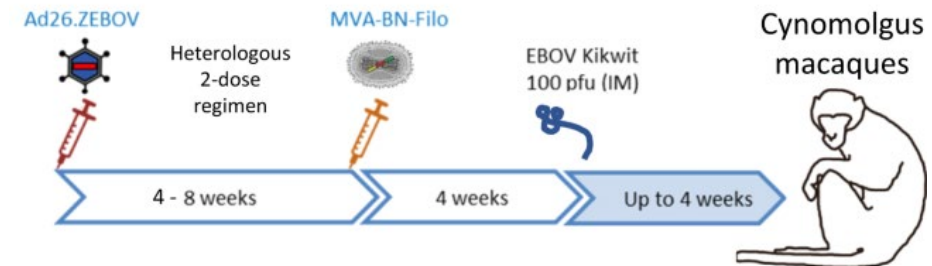
- Ebola outbreaks are not predictable in terms of location and size
  - Near impossible evaluation of a prophylactic vaccine efficacy
- In this situation and under the Animal Rule (FDA) / Conditional or Exceptional Circumstance authorization (EMA), licensure based on animal efficacy data bridged to human immunogenicity data may be acceptable
  - Confirmed at VRBPAC
- Methods based on a similar approach for Anthrax vaccine



# Selection and implementation of a bridging biomarker

## Major challenges

- Establish/use relevant animal model
- Identify immunological biomarkers to examine
- Statistical description of the immuno-bridging approach
- Bridging the immune response between species
- Interpretation for clinical relevance



Bridging immune responses in people to survival after Ebola virus challenge in a non-human primate model

- Establish immunological parameter that correlates with protection to Ebola challenge in a fully lethal non-human primate model
- Measure this immunological parameter following vaccination of people
- Infer clinical benefit of vaccination in people by comparison with protective immune responses in nonhuman primates

# Assumptions that pre-clinical endpoints can be surrogates for clinical endpoints



Pre-clinical disease model is a surrogate for human disease

Host susceptibility  
Challenge agent  
Exposure route  
Symptoms and manifestations



Pre-clinical model is a surrogate for vaccine responses in humans

Pre-clinical species immune system is comparable to humans  
Vaccine responses in pre-clinical species are similar to those in humans (PK,PD)



Associations of immune responses with protection in pre-clinical model is similar to those in humans

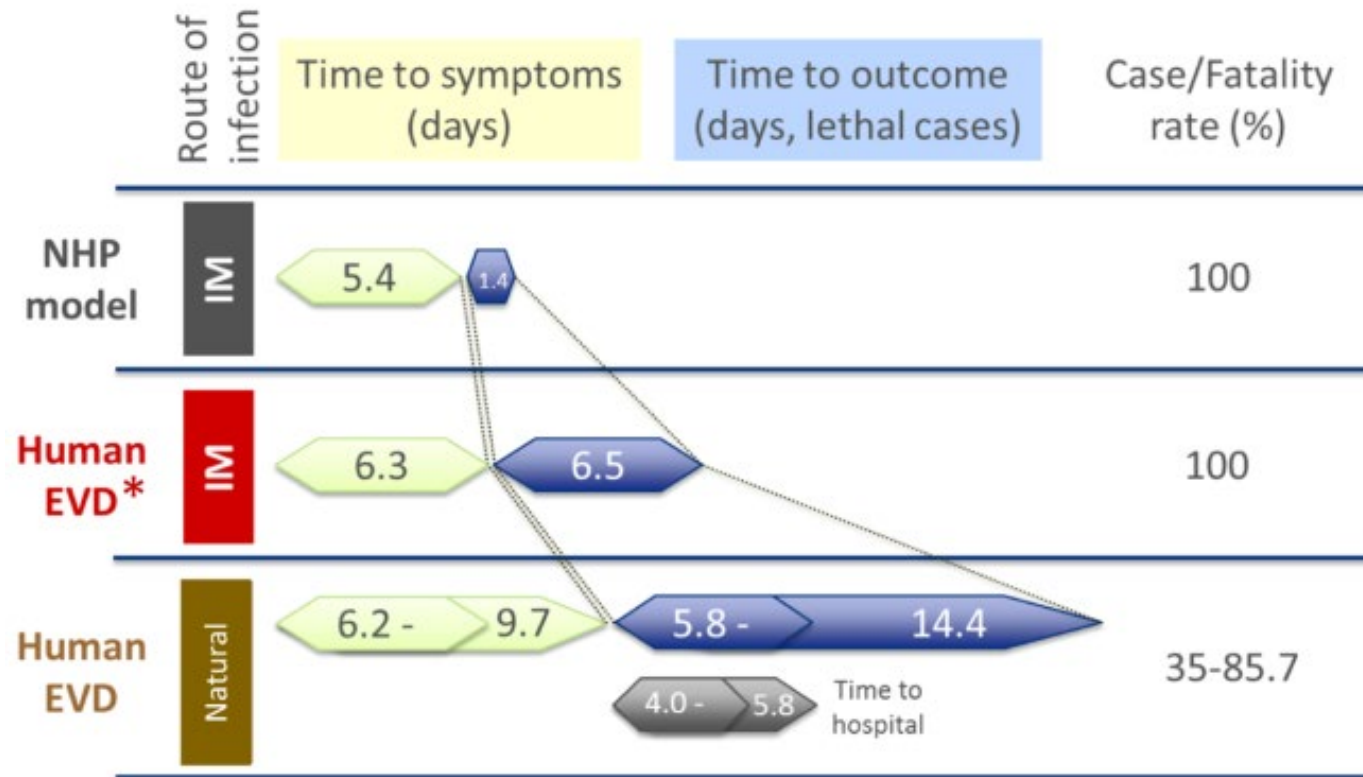


Supporting data can strengthen confidence in above assumptions

Natural history immune response data in humans

# Ebola: Pre-clinical data that led to the approval of Zabdeno/Mvabea

Pre-clinical disease model is a surrogate for human disease



Although the challenge model is more stringent, it is deemed appropriate to show reasonable likelihood of clinical benefit

# Ebola: Pre-clinical data that led to the approval of Zabdeno/Mvabea

Identifying relevant immune markers associated with protection from Ebola in NHP



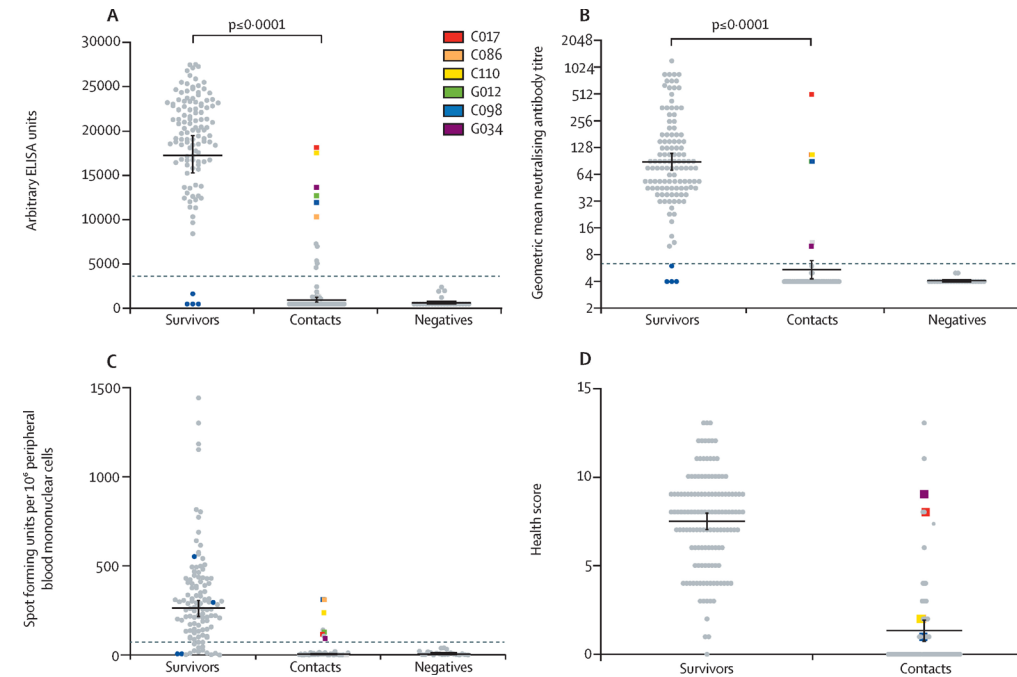
Selection based on:

- Known immune responses in Ebola survivors
- Known immune responses elicited by the vaccine platform

For this vaccine/challenge model:

- Binding antibodies by GP ELISA
- Neutralizing antibodies by psVNA
- Specific T cells by IFN- $\gamma$  ELISpot

*(not considered as limited/no data available: wt VNA; ICS; detailed characterization of Abs)*



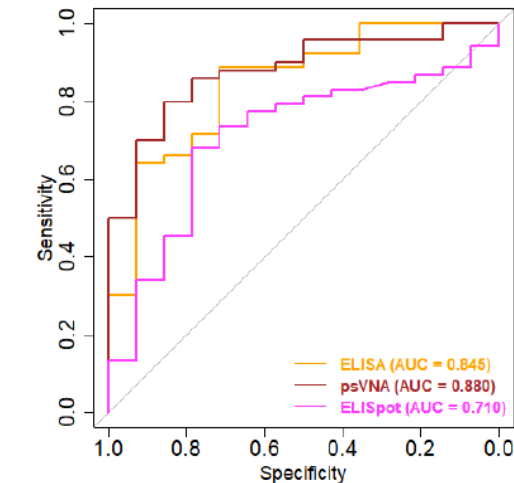
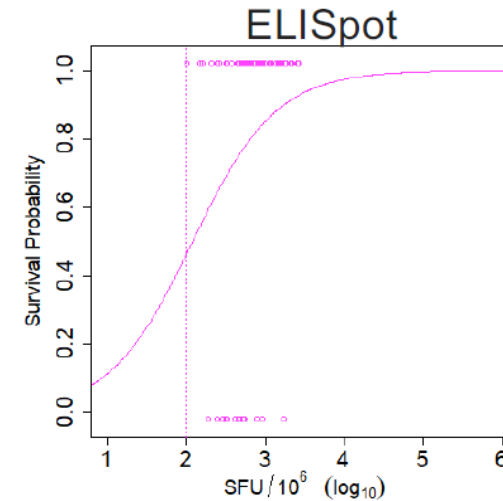
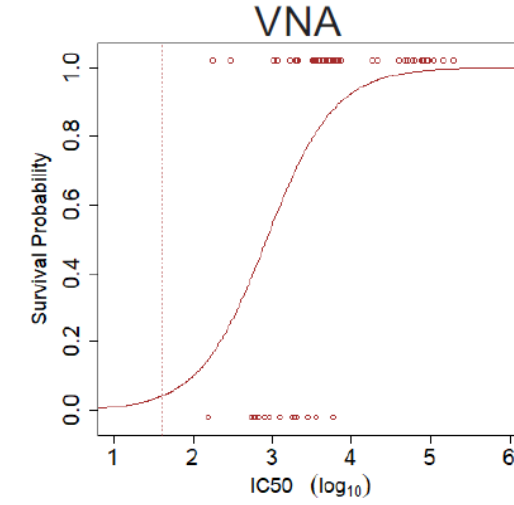
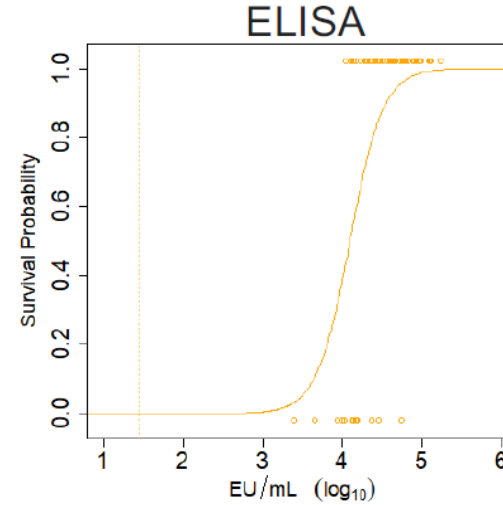


# Ebola: Pre-clinical data that led to the approval of Zabdeno/Mvabea

Single immunological correlates of protection in NHP

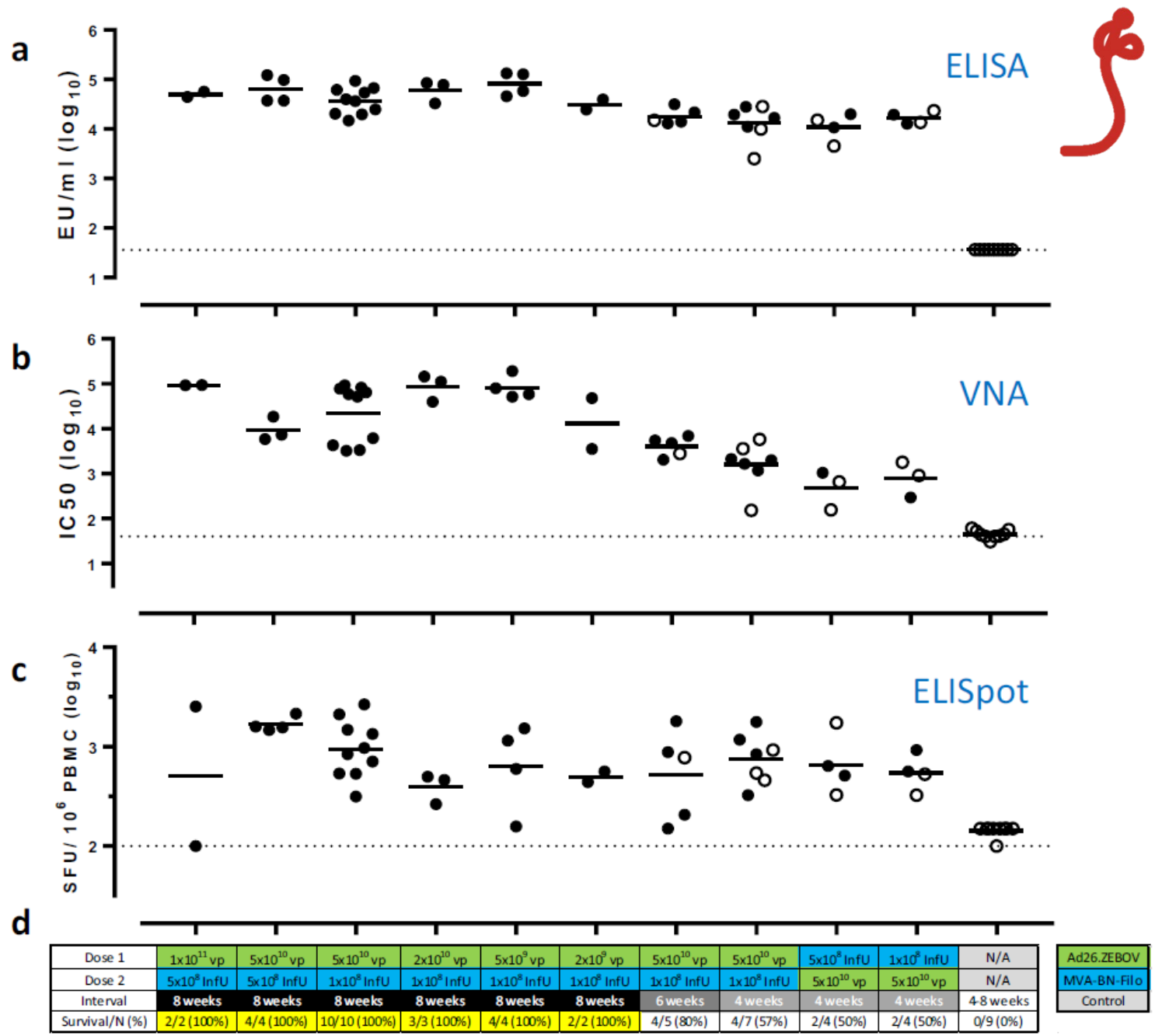
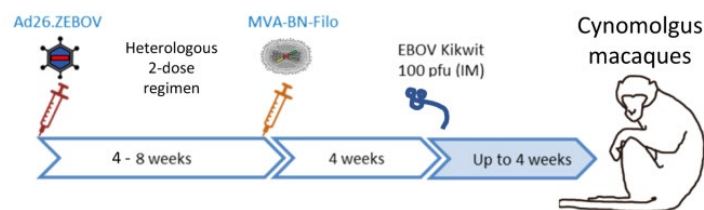


- Binding antibodies, neutralizing Abs and T cells are significantly correlated to challenge outcome in NHP
- ELISA and VNA are comparable for discriminatory power
- Lower discriminatory capacity with T cells



# Ebola: Pre-clinical data that led to the approval of Zabdeno/Mvabea

## Refining the curve

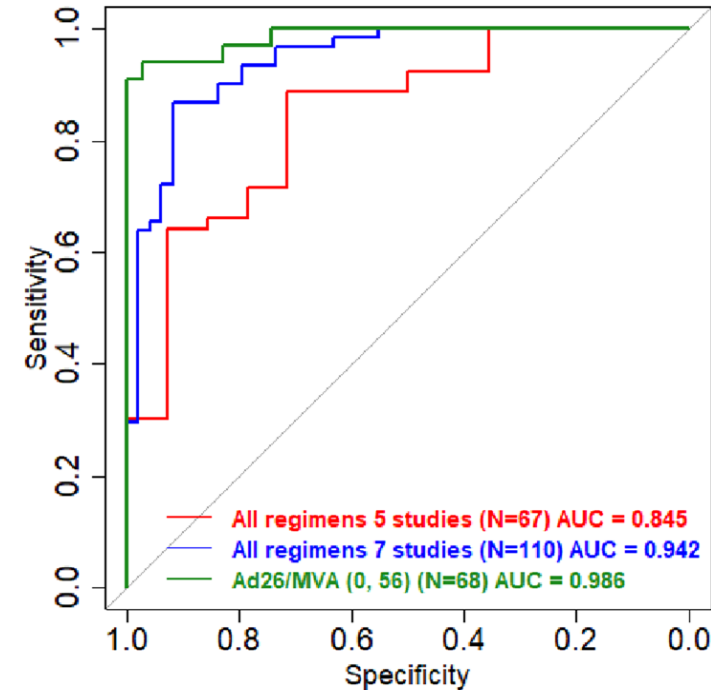
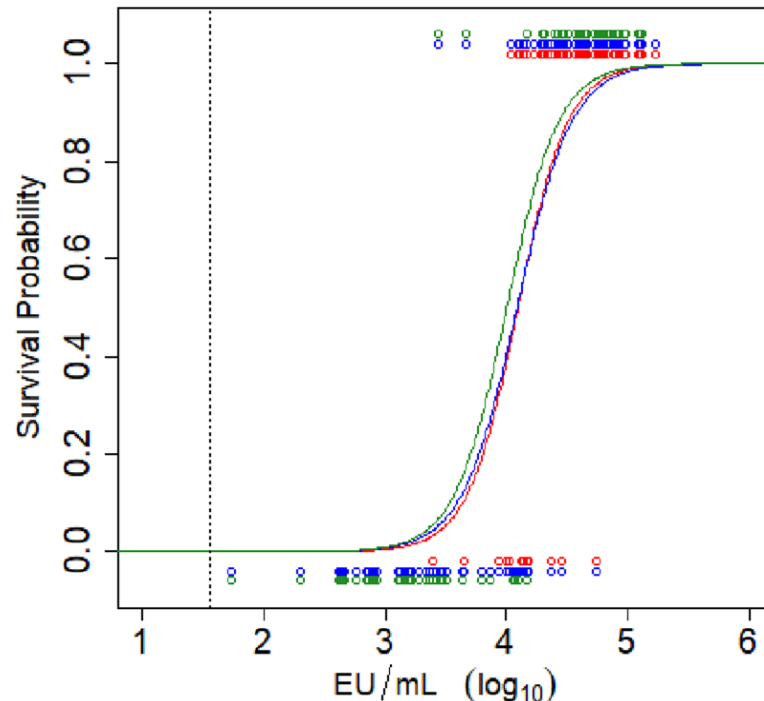


# Ebola: Pre-clinical data that led to the approval of Zabdeno/Mvabea

## Refinement of ELISA logistic model by vaccine dose down studies



- Initial data set was collated from 5 studies that tested different regimens in addition to the selected clinical regimen: Dose, lots, directionality and spacing
- Negligible contribution of other covariates to discriminatory capacity of the model
- Dose down studies were conducted, to generate more data in the lower end of the curve
- These studies improve model with relation to discriminatory capacity between survivors and non-survivors

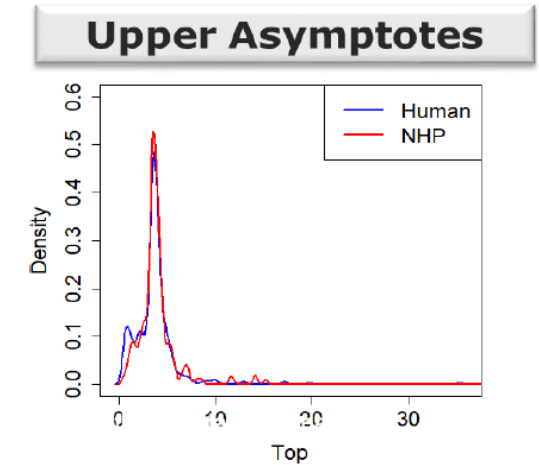
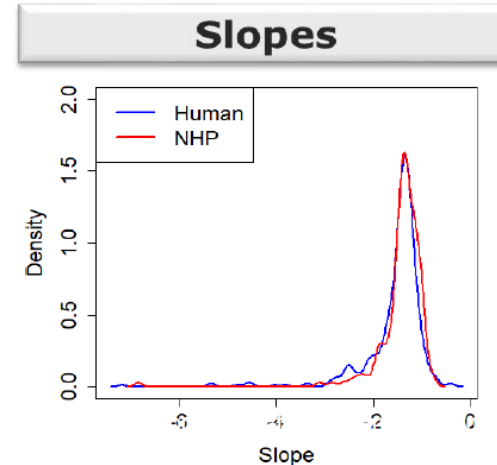


# Ebola: Pre-clinical data that led to the approval of Zabdeno/Mvabea

Measuring the immune response for NHP and human samples



- Bridge the human immunogenicity results; One Lab, One Assay
- Transfer to CRO, to increase control over assay performance and resourcing
- FANG ELISA parallelism investigation: Suitability of using the same assay for both species
- Secondary Ab cross-reactive experiment
- (Sponsor-specific) validation of ELISA for human serum
- (Sponsor-specific) validation of ELISA for NHP serum
- (re-)measured all NHP samples in validated NHP ELISA to establish logistic regression curve

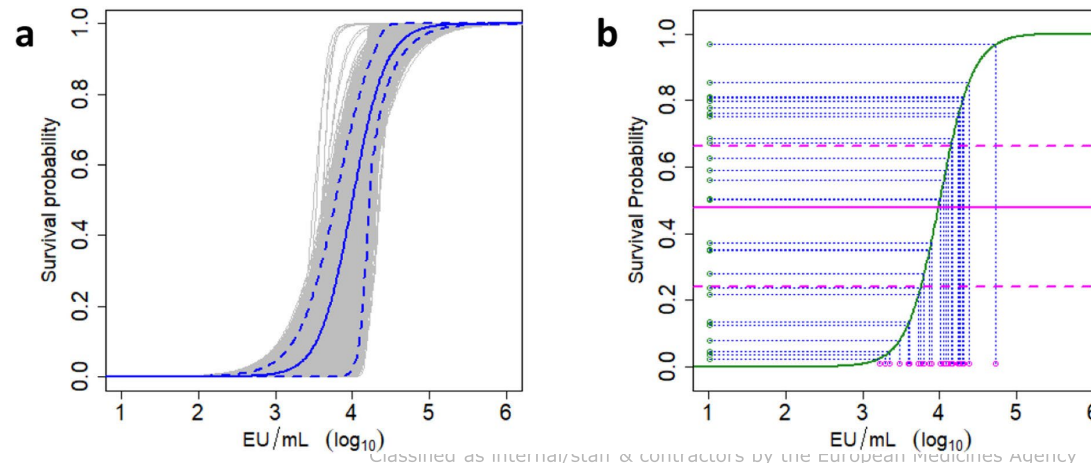


# Ebola: Pre-clinical data that led to the approval of Zabdeno/Mvabea

FDA- and EMA-approved immunobridging strategy: inferring clinical benefit



- Use curve to calculate mean survival probability and 95% CI in humans using double-bootstrap method
- Clinical benefit demonstrated if lower limit of 95% CI is above pre-specified success criterion of 20% (FDA- and EMA- approved)
- NHP model more stringent than EVD in humans:
  - Model can provide evidence for clinical benefit
  - Model cannot quantify vaccine effectiveness, as 1:1 translation will likely underestimate clinical benefit
- Quantification of the actual clinical effectiveness must be determined in a field study

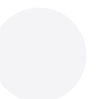


# Ebola: Pre-clinical data that led to the approval of Zabdeno/Mvabea

## Immuno-bridging Statistical Analysis



- **Primary analysis on all adults** (18- 50 years) from Ph2 and Ph3 studies. N=1550
- **Pre- specified sensitivity analyses** to investigate the robustness of the primary analysis:
  - Analysis stratified per baseline EBOV GP FANG ELISA level to assess the impact of pre- existing EBOV GP binding antibody levels
  - Analysis excluding the subjects of the Sierra Leone study (EBL3001)
- **Demographic sub- analyses** stratified by:
  - age
  - Sex
  - Race
  - geographic region, including East Africa, West Africa, Europe and USA.
- **Post- hoc analysis** for further insight to the primary analysis:
  - Analysis stratified by country
  - Analysis of entire adult population stratified by country
  - Analysis stratified by special populations (elderly [ $>50$  years], HIV- infected adults, pediatrics)



# Ebola: Pre-clinical data that led to the approval of Zabdeno/Mvabea

Immuno-bridging Statistical Analysis



Final Im m u n o - bridging (PPS) Final Prim ary analysis	
Per Protocol Immunogenicity Analysis Set	Ad26.ZEBOV, MVA-BN-Filo (0,56)
N	1550
Mean Predicted Survival Probability (95.7% CI)	57.3% ( <b>41.2%</b> ; 71.0%)

The 95.7% CI lower limit of **41.2%** passes the pre- specified success criterion of 20%

- Evidence for clinical benefit successfully dem onstrated at final analysis
- In view of the stringency of the model, the point estimate cannot be used for absolute quantification of vaccine efficacy in humans.
- Estimate of clinical effectiveness and duration of protection to be demonstrated in subsequent studies

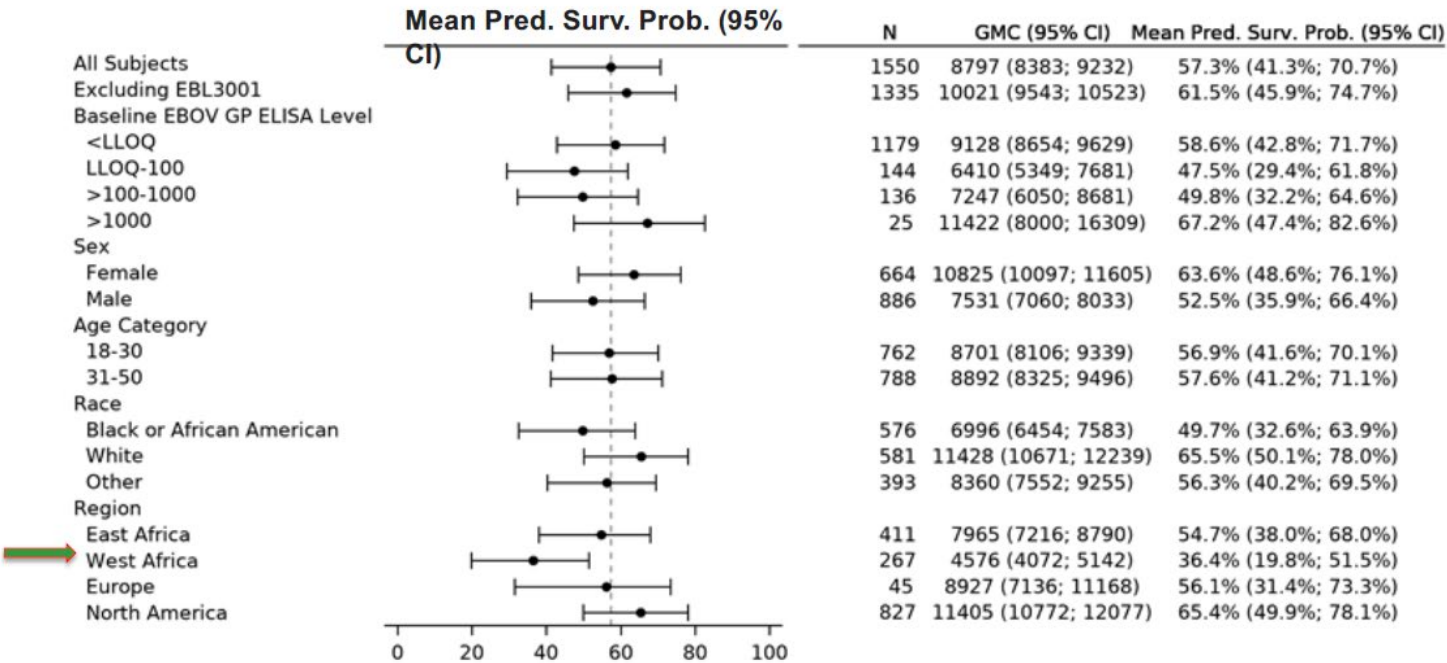


# Ebola: Pre-clinical data that led to the approval of Zabdeno/Mvabea

Immuno-bridging Statistical Analysis



Final Immuno-bridging (PPS)  
Subgroup analysis



- Observed clinical benefit not driven by inclusion of potentially pre- exposed Sierra Leone subjects.
- Sensitivity analyses fully consistent with primary analysis.
  - Trend to lower prediction in West- Africa, influenced by lower immune responses in Sierra Leone.



# Several questions remain

- Translation of animal model vaccine efficacy to protection from human EVD: how can we assess effectiveness?
- Bridging to other vaccines where efficacy has been shown such as ERVEBO, how to approach a possible immuno- bridge?
- Mechanism of Action:
  - binding Abs are a suitable bridging biomarker, but protection is assessed in the vaccinated animal, with all immunological parameters that may contribute to protection
- Protection is assessed at peak immunogenicity response; how does this predict for long term protection?
  - Longevity of the immune response and memory responses are established; how do they contribute to protection?

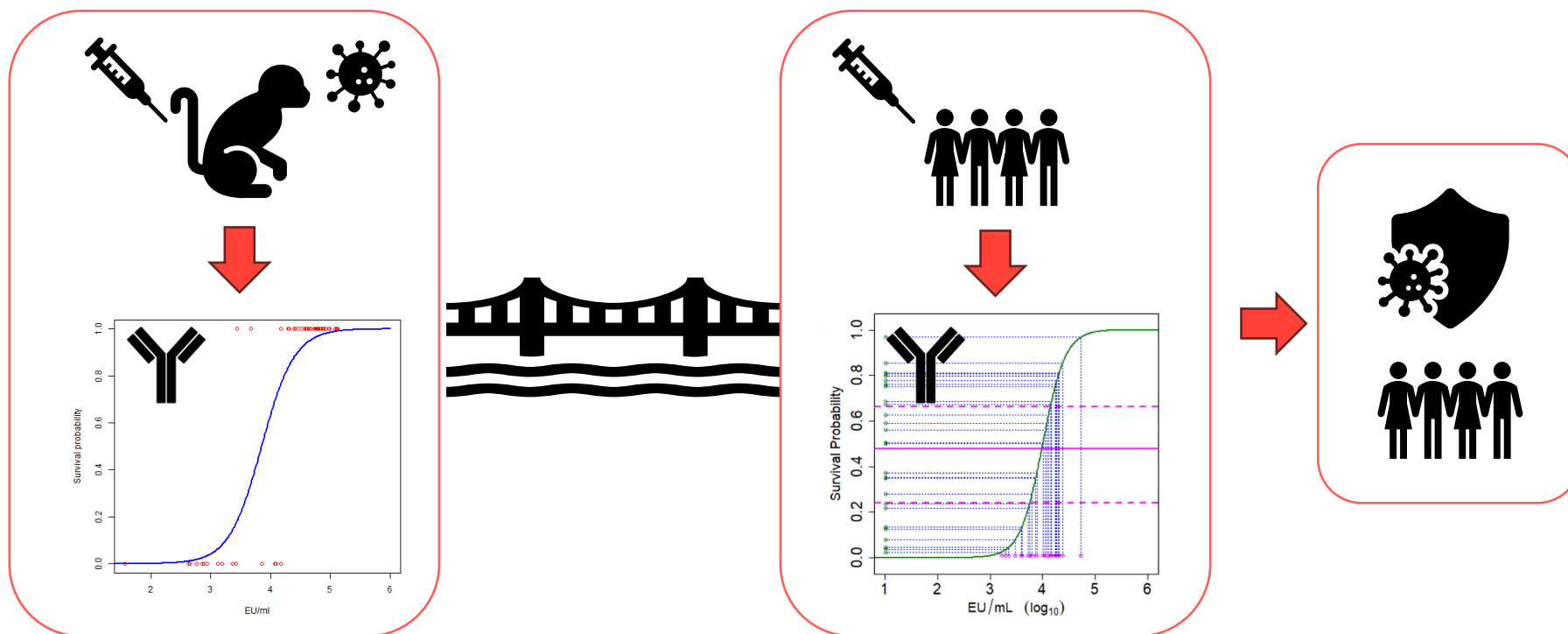
# CEPI

# Back up

# Alternative pathway to vaccine licensure: Immuno-bridging



Vaccine immune responses associated with protection in NHP provide a model to bridge to human vaccine immune responses and estimate the likelihood of clinical benefit



Current vaccine licenses based on immuno-bridging\*:

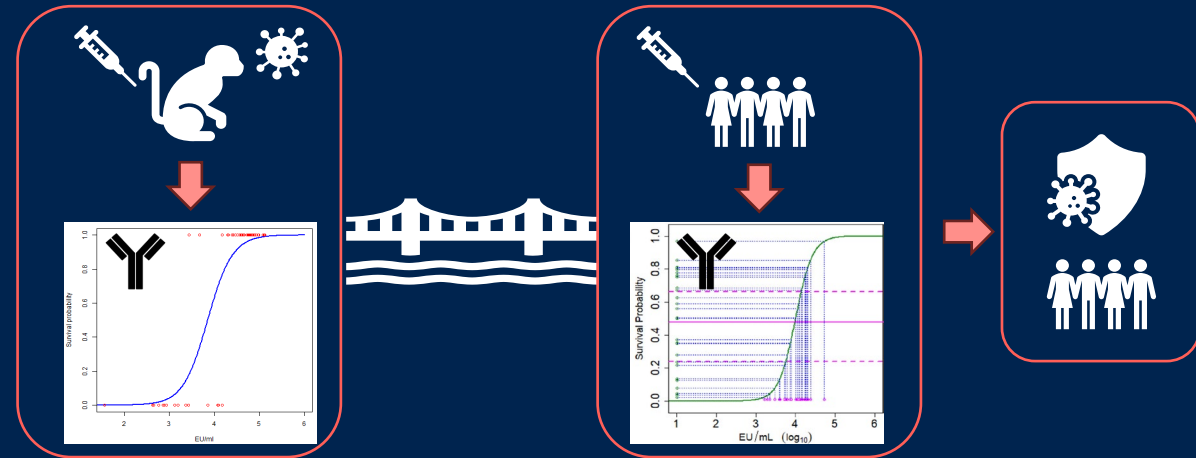
Anthrax : Biothrax and CYFENDUS

Ebola Zaire : Zabdeno/Mvabea

Chikungunya: IXCHIQ and VIMKUNYA

# Summary

- Immuno-bridging is an approach to vaccine licensure when conventional vaccine licensure is not possible; focus should lie on bridging the gaps
- Ensure argumentation around feasibility of efficacy studies
- Study FDA and EMA guidance and case studies
- Ensure that pre-clinical disease model is a surrogate for human disease and well characterized
- Examine associations of immune responses with protection in pre-clinical model and relate to human immune responses
- Supporting data can strengthen confidence in immuno-bridge: Natural history immune response data in humans
- Prepare to meet quality requirements for assays
- Define success criteria with uncertainty/confidence in mind
- Clinical data package may be a lot lighter than conventional late stage vaccine development packages



# Setting success criteria



Requires a well thought out strategy and thorough knowledge of the elements required



Not about target thresholds or point estimates, it is about the lower bound



->Confidence vs uncertainty



Variation at NHP and human side, each with a lower bound



Passive transfer: indicates mechanism, using human antibodies to protect NHP, set threshold of protection-> xx% of vaccinees to exceed threshold (lower bound 95%CI)



Logistic regression: mechanism unknown, relate NHP protective responses to human responses, set target xx% probability of protection (lower bound 95%CI)