

Immune responses in Lassa Fever survivors



Critical research for priority pathogens with epidemic potential

Online Meeting 18-01-2024

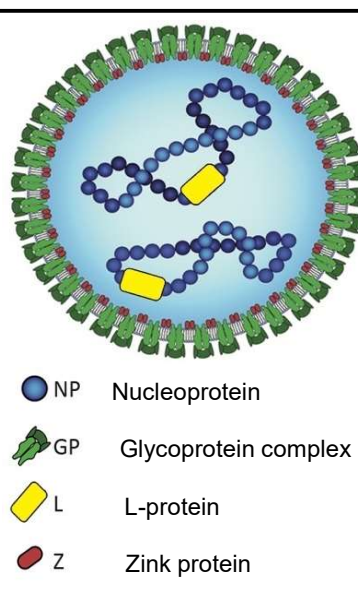
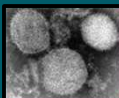
Lisa Oestereich, Department of Virology

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Lassa Virus as a prototype Arenavirus

Lassa virus and fever

- Ambisense RNA virus
- *Arenaviridae* family
- 2 Genome segments, 4 proteins



modified from Fedeli et al. 2018

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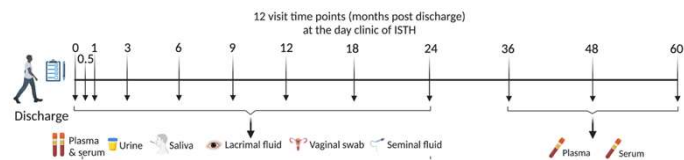
Lassa Virus

Lassa virus and fever

- Estimated 300.000 cases; 5.000 deaths annually
- Many asymptomatic cases; symptoms range from mild flu-like presentation to hemorrhage and fatal shock
- Endemic in West Africa, seasonal; especially in Nigeria
- 2019** Nigerian outbreak:
>810 cases and
>167 deaths
- 2023** 1270 laboratory-confirmed cases from 28 of 36 states; 18 % case fatality rate

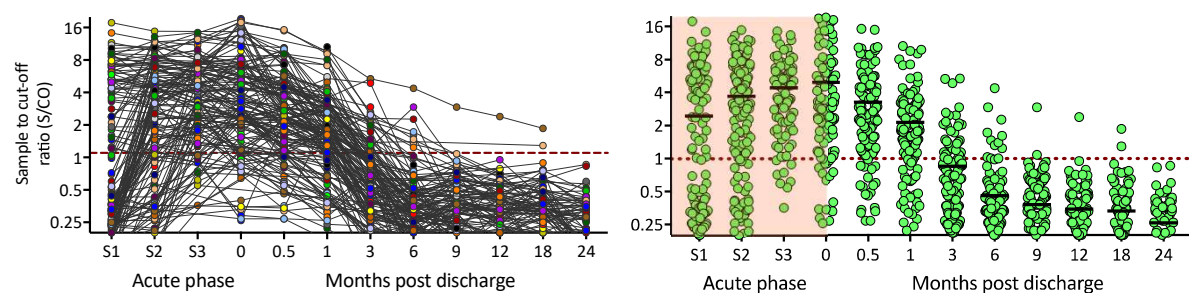
Clinical studies at ISTD, Nigeria

- Pathogenesis study linking laboratory data with clinical data
- Follow-up of Lassa Fever survivors

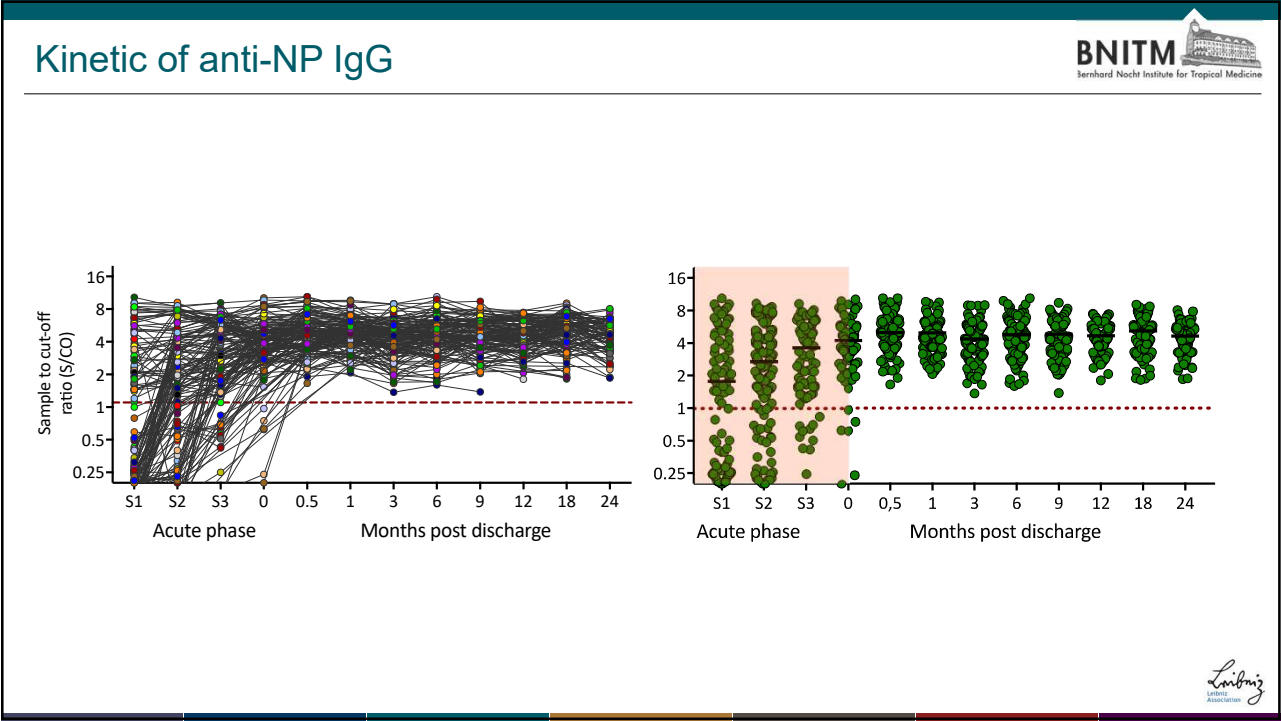


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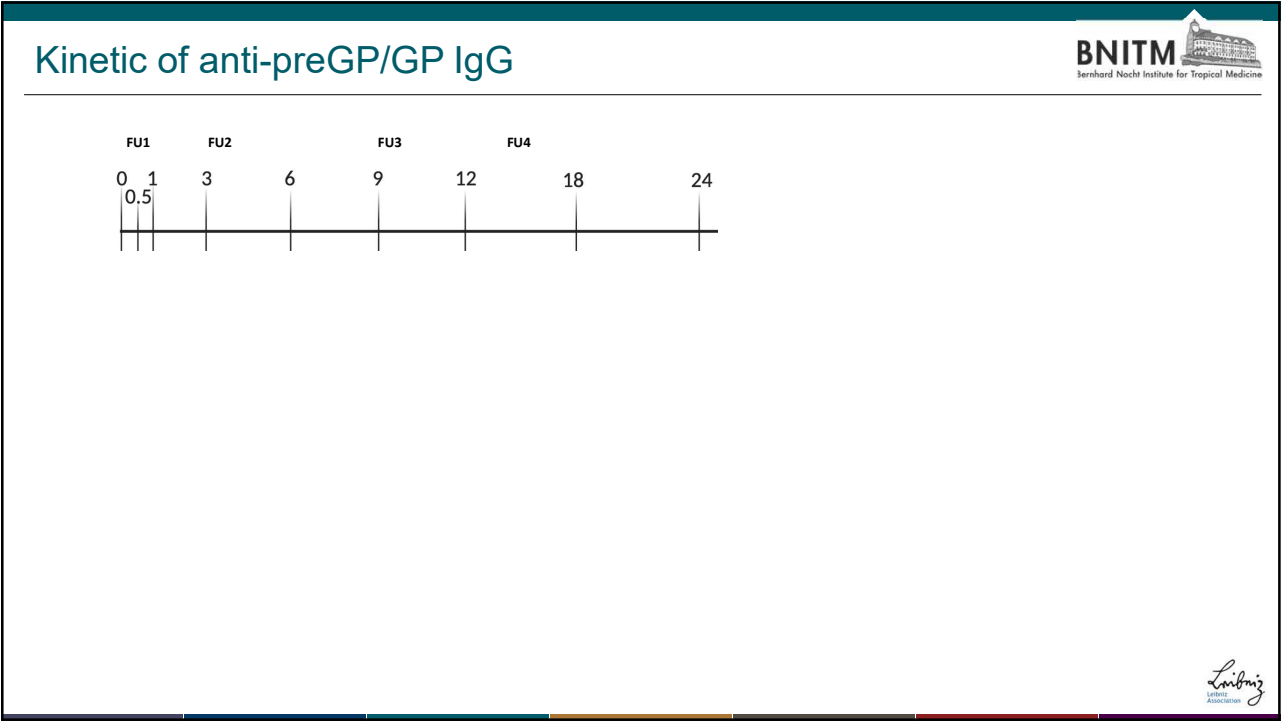
Kinetic of anti-NP IgM



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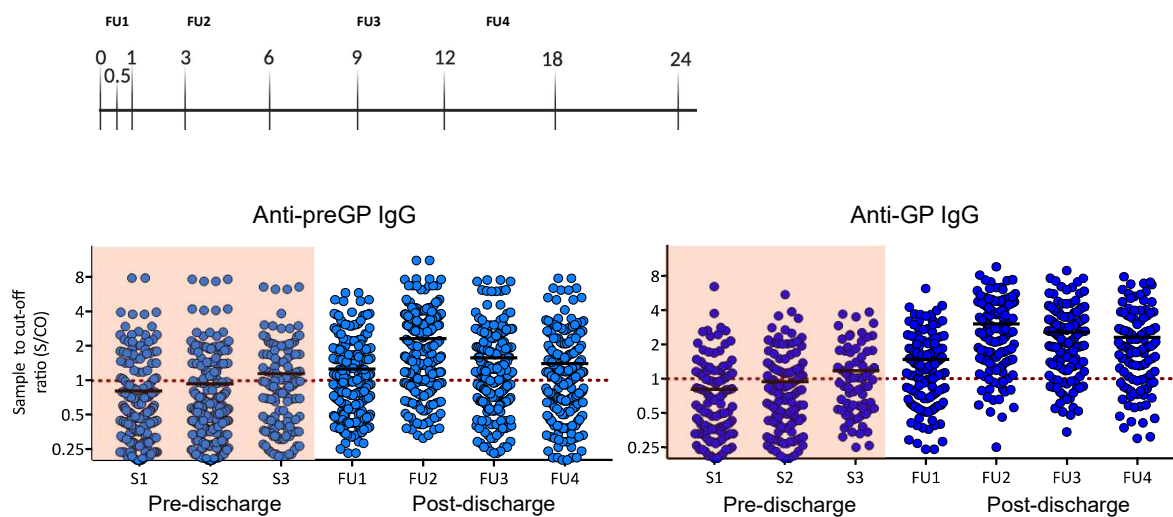


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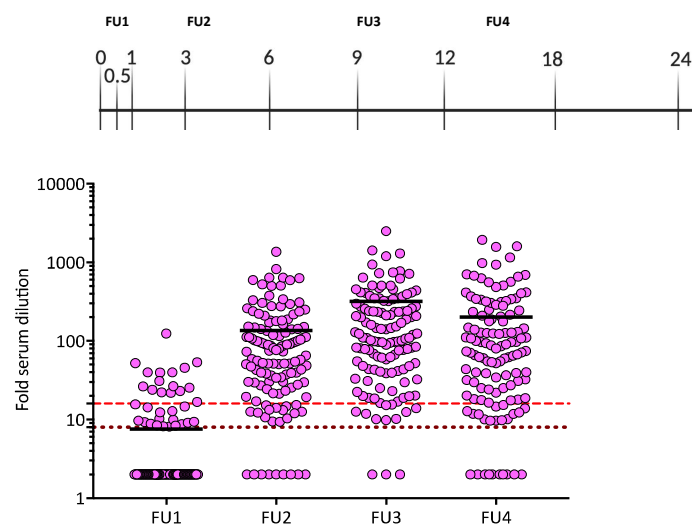
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Kinetic of anti-preGP/GP IgG

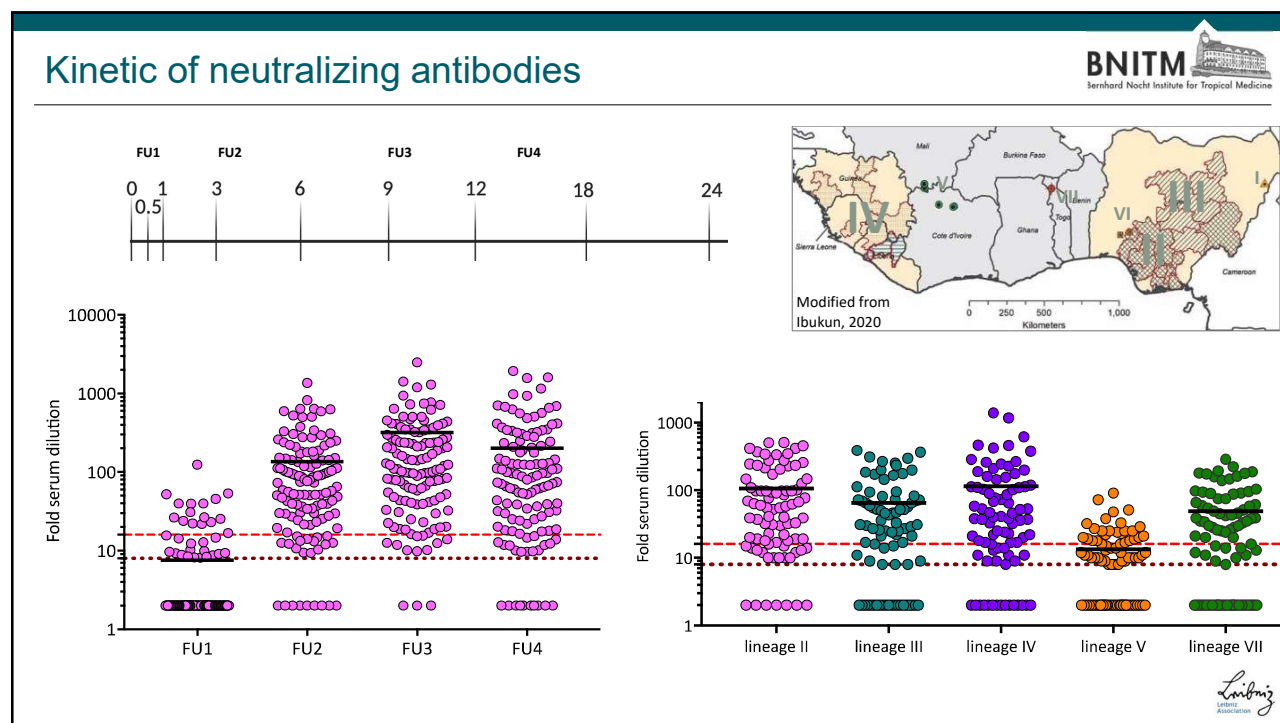


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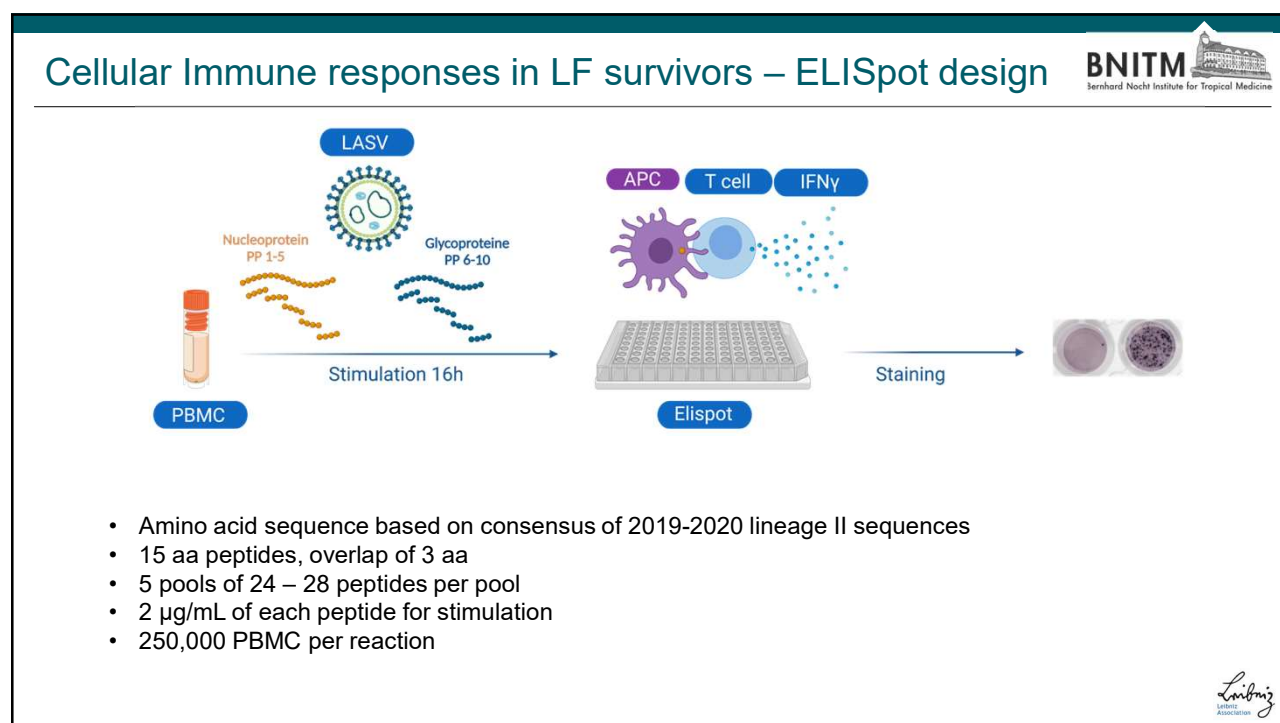
Kinetic of neutralizing antibodies



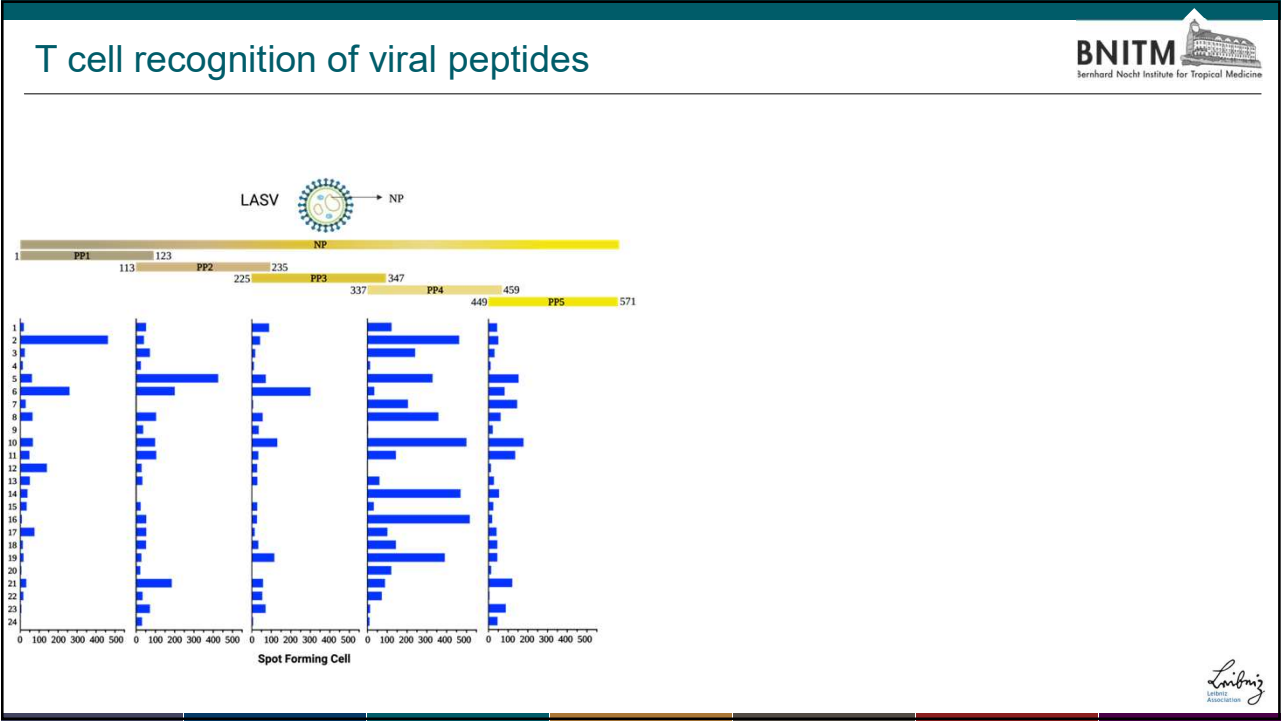
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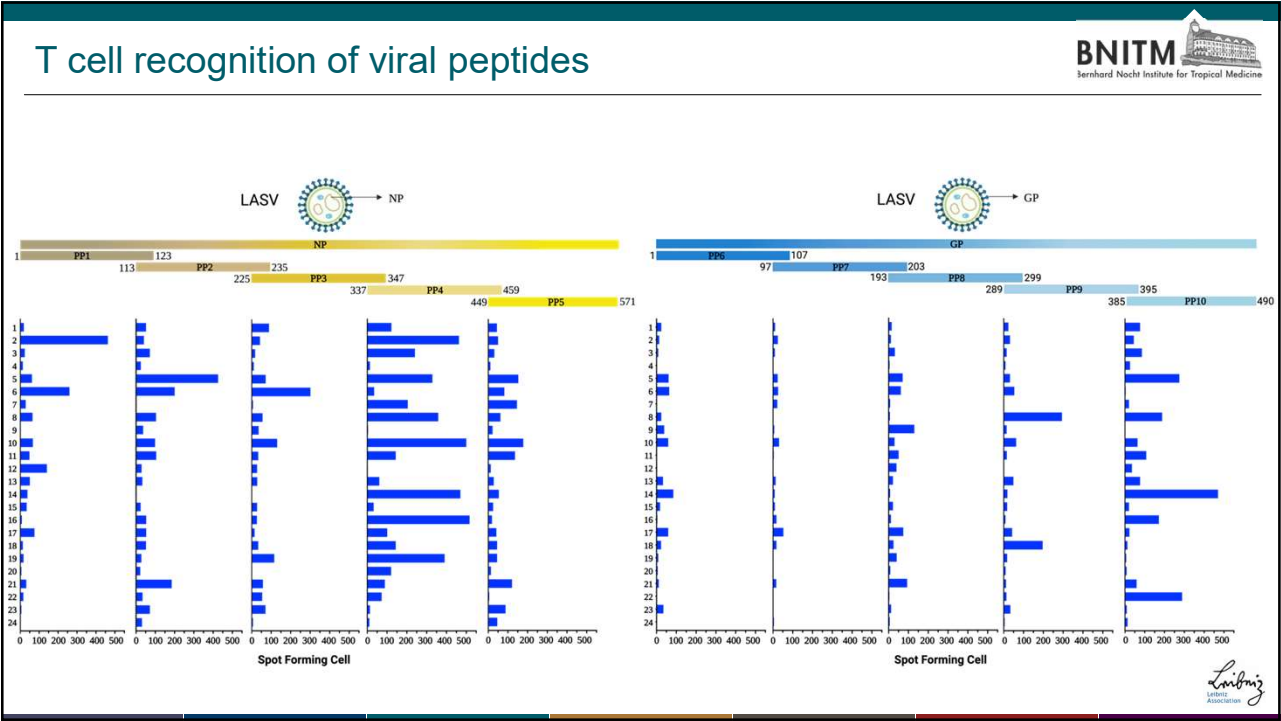
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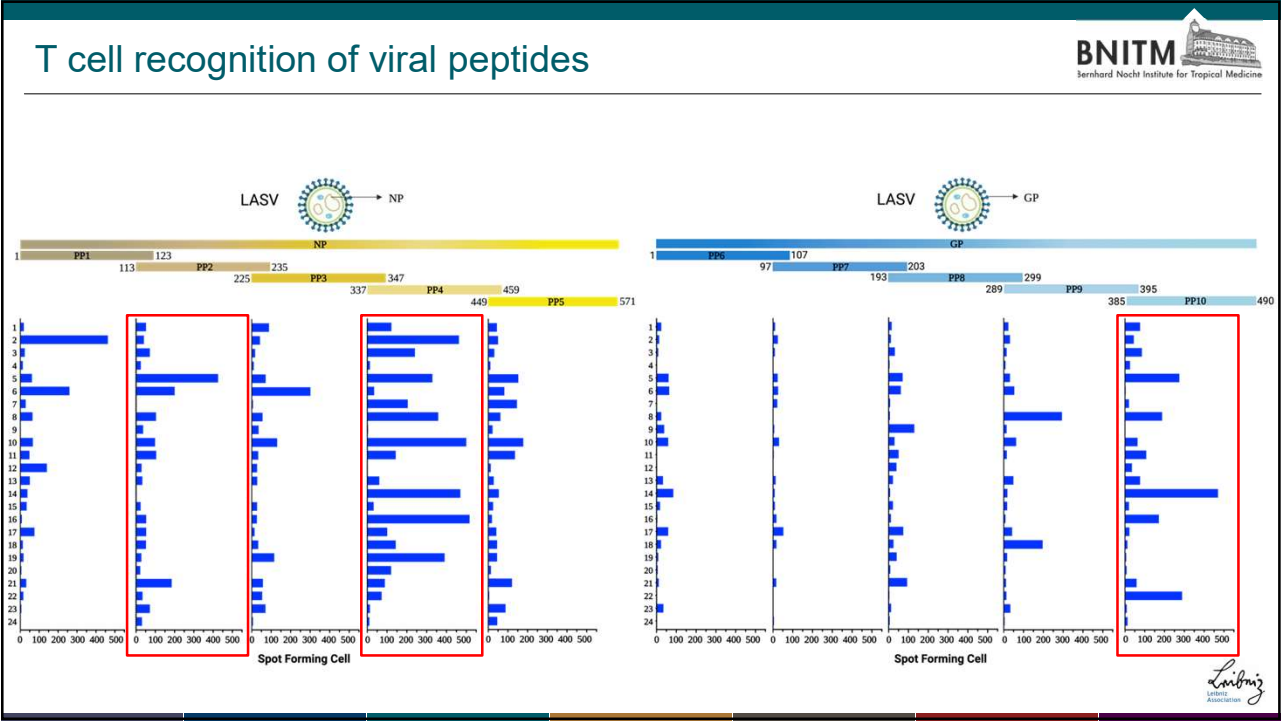
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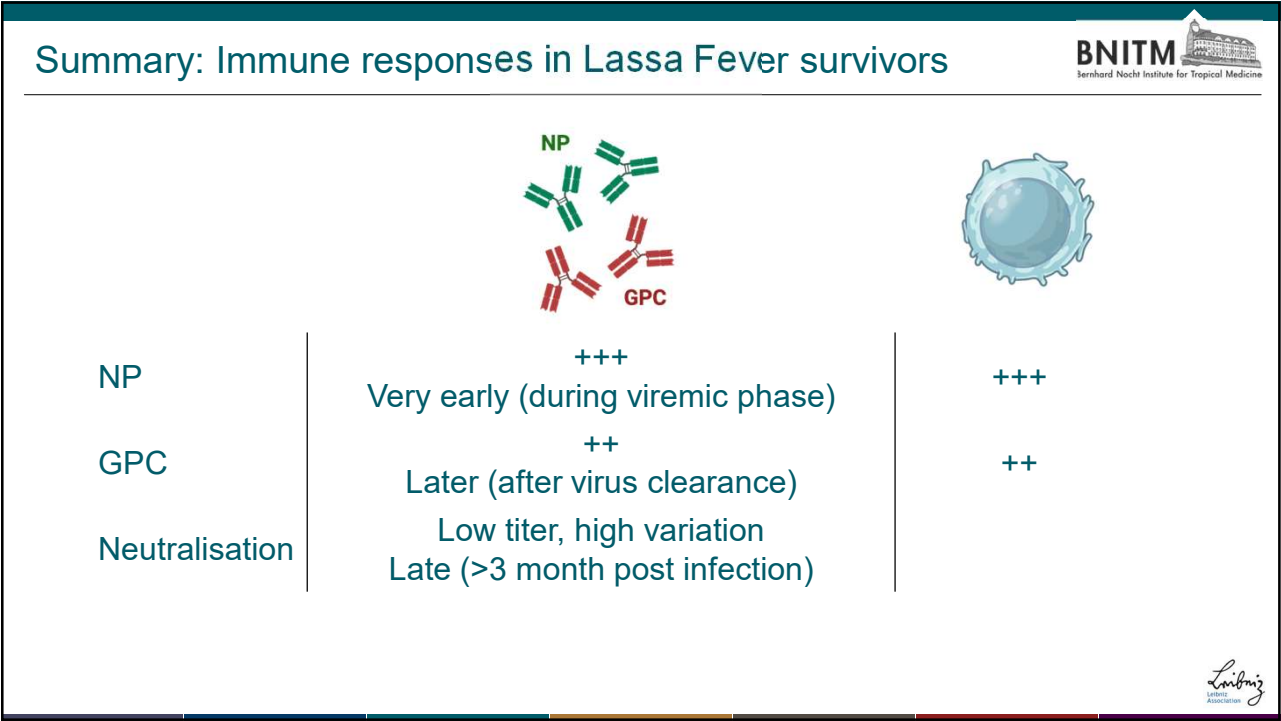
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Challenges



- ❖ Most vaccines target contain GPC as an antigen
- ❖ High genetic variability of LASV strains; vaccines based on “old” lab strain
 - ❖ Crossprotection?
- ❖ Low neutralizing antibody titer and overall delayed and lower antigenicity of GPC compared to NP
- ❖ More and stronger T cells epitopes in NP compared to GPC



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Arenaviruses beyond LASV




- ❖ Mopeia and Morogoro virus infection of mice or NHP induces protective immunity (LASV challenge)
- ❖ No/low cross-reactivity of T cells between different Old World Arenaviruses
- ❖ Limited cross-reactivity of antibodies between old World Arenaviruses
- ❖ No recognition of New World Arenaviruses with Lassa Fever survivor plasma



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