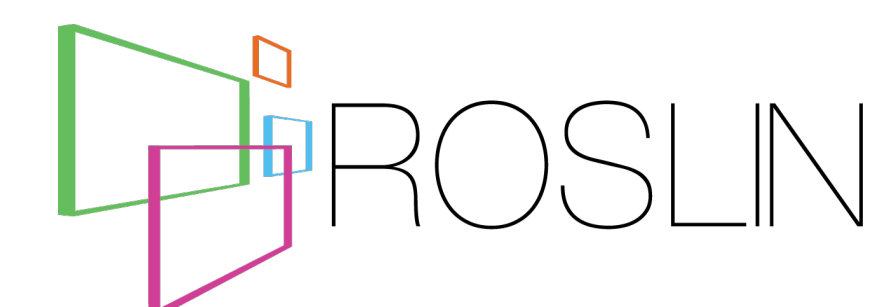




Smallholder dairy cattle epidemiology and health genetics in Tanzania



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INTRODUCTION

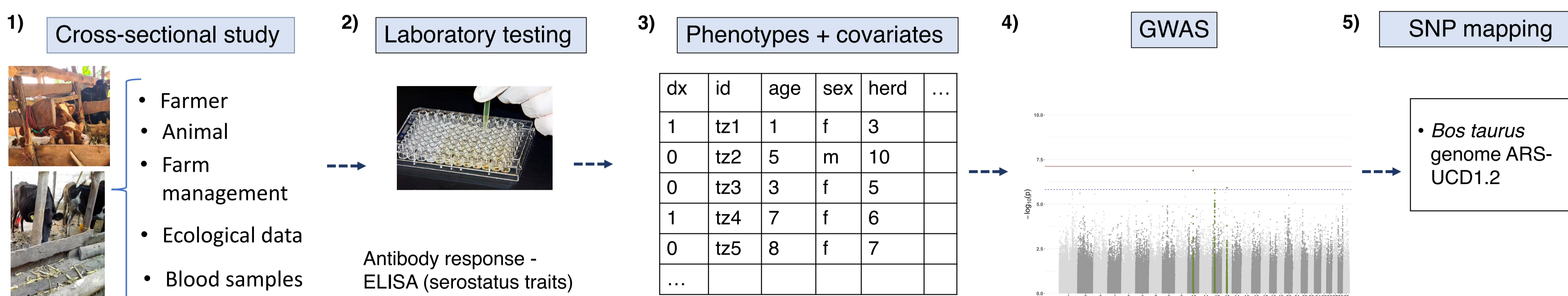
- High pathogen burdens threaten efficient livestock production in LMICs.
- Breeding disease resilient/tolerant dairy cattle could improve animal welfare, health and production while reducing livestock carbon emissions.
- Genetic component to susceptibility or tolerance of infectious diseases in cattle largely unquantified¹.

MATERIAL AND METHODS

- Crossbreed cattle (n = 2045) were genotyped and screened for six infectious diseases (**Fig 1A**) of production (and public health) importance across six dairy regions in Tanzania (**Fig 1B**).

- Genome-wide association studies² (GWAS) identified **41** SNP markers (**Fig 2**) linked to pathogen-specific serostatus while accounting for relatedness, population structure, and other environmental risk factors.

Methodological framework



RESULTS

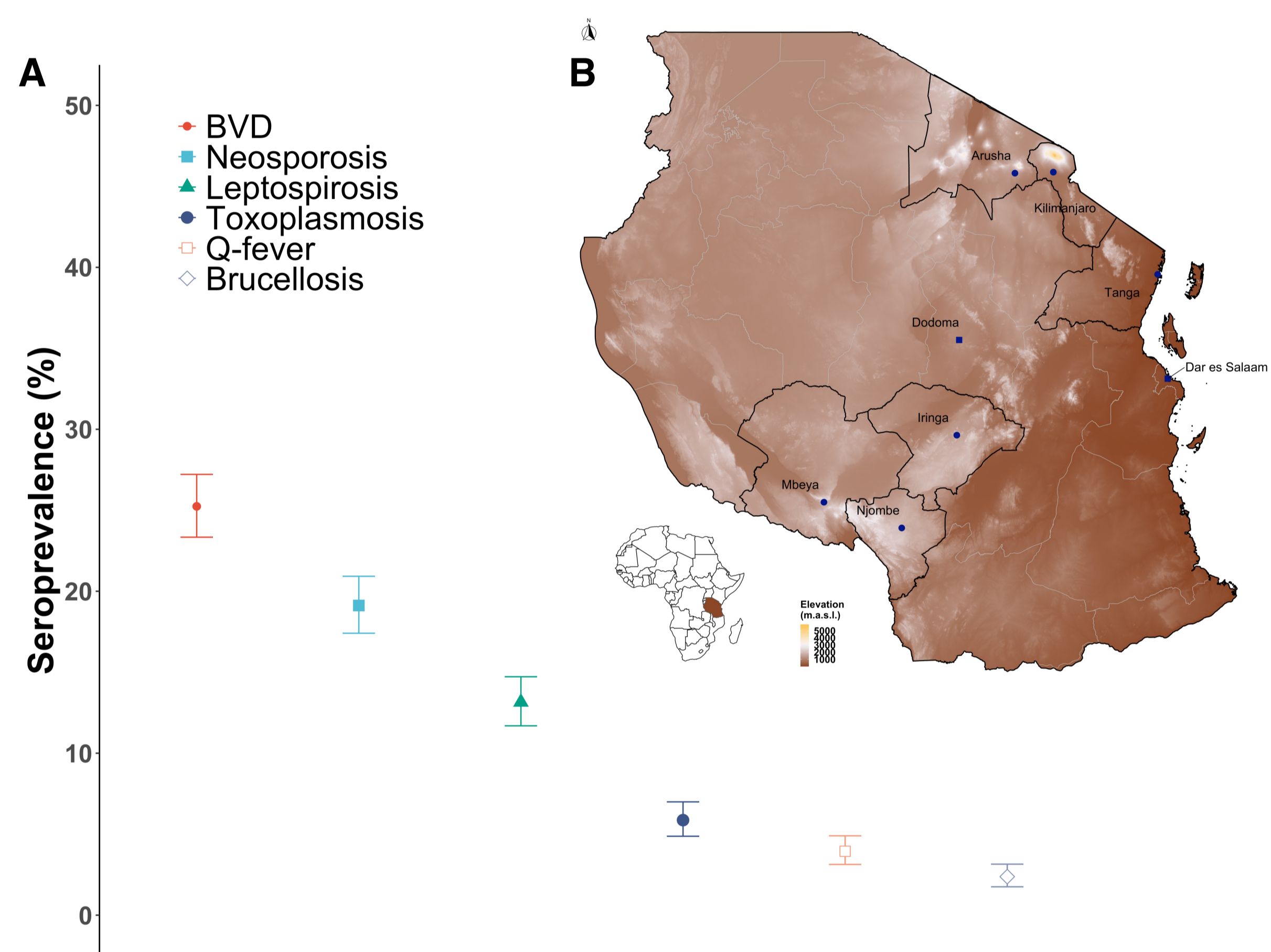


Fig 1. Sampling across six dairy regions in Tanzania in which seroprevalence for BVDV, *Neospora caninum*, *Leptospira interrogans* serovar Hardjo, *Toxoplasma gondii*, *Coxiella burnetii* (Q-fever) and *Brucella spp.* was variable and relatively high for most diseases.

DISCUSSION AND CONCLUSION

- Serostatus is heritable in this cattle population (h^2 , 0.05 – 0.46).
- Preliminary genome scans revealed loci potentially consistent with the polygenic nature of immunological response to viral, bacterial and parasitic diseases.
- Several loci mapped to genes in the cow genome involved in pathogenesis (**EXOC1**, **NTM**), immunological response (**TNFSF8**) and disease resistance (**FAM78B**)^{3,4,5}.

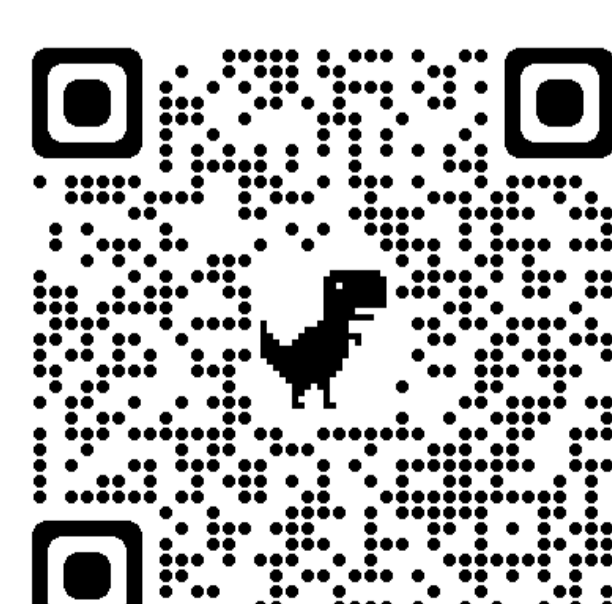
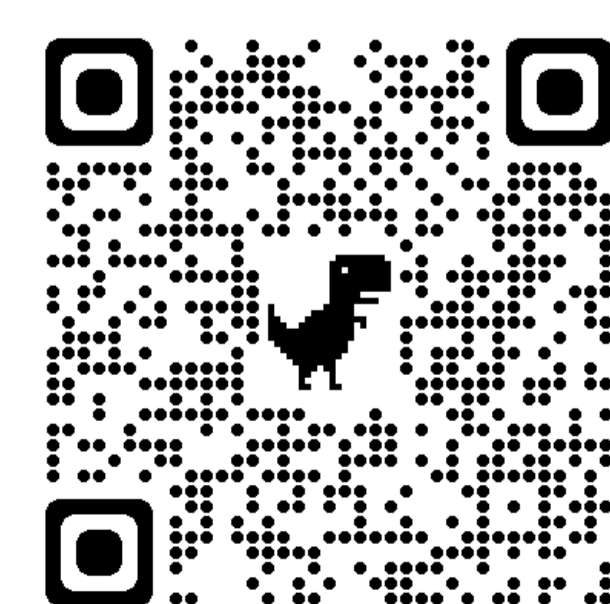
CONTACT

• Exciting new projects starting soon. Follow the QR codes for more information.

• Prof Mark Bronsvort (PI)

• Dr Enrique Hernandez

• CTLGH



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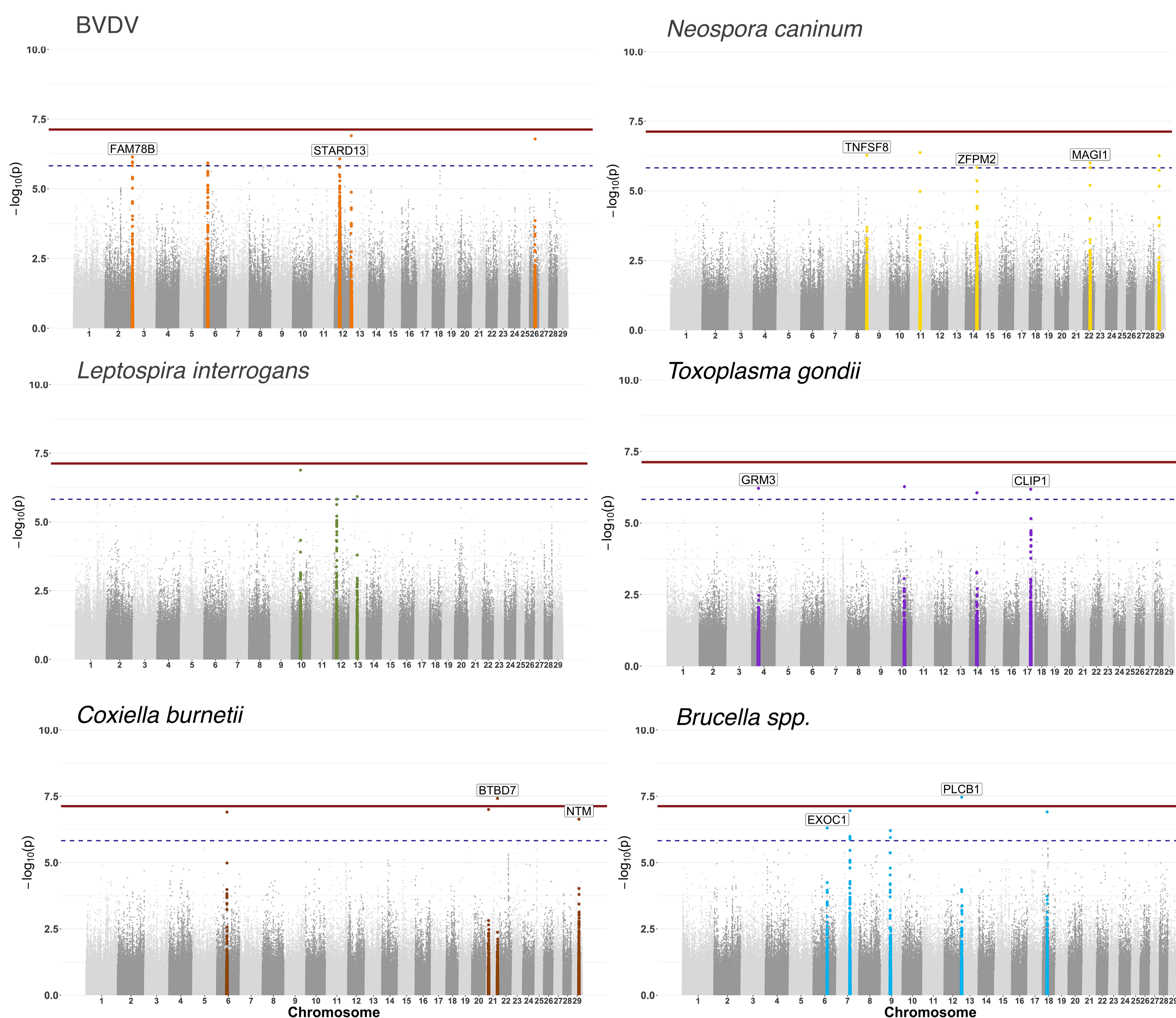


Fig 2. Manhattan plots show regions (colour-coded dots) in the *Bos taurus* genome with strong association to serostatus for a given pathogen. Several SNP markers mapped to several annotated regions.

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