TOWARDS AN INTEGRATED ANALYSIS OF CRYPTORCHIDISM IN GERMAN SHEEP POODLE

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BACKGROUND

Cryptorchidism is a disease, in which one or both testes do not descend into the scrotum properly¹. It affects up to 4% of men² and 10% of male pups³. Amongst other disorders it is associated with a 13.6 times higher risk for the development of malignant neoplasms and is correlated with an impaired





spermatogenesis⁴.

Despite its associated health risks and accompanying economic damage, resulting from surgery and losses in breeding¹, studies on canine cryptorchidism and its causes are relatively rare.

Figure 1: German Sheep Poodle⁵

OBJECTIVE

- Setting up a relational database with known cryptorchidism-associated genetic variants based on the CryptoGene database⁶
- Sequencing associated genes using the MinION to identify new variants and develop a genetic test



CONCLUSION

A database containing 74 genetic variants in dogs, humans and mice was set up. Only 14.9% of all entries are found in dogs. PCR based amplification of INSL3 and four exons of RXFP2 was established.

OUTLOOK

The bioinformatics and experimental studies serve as starting points in preparation to identify variants that could elucidate genetic etiology of canine cryptorchidism.



Figure 3: Database entry distributions of a) data types, b) genes and c) species



Figure 4: Amplicon agarose gel

After an update in spring 2019 the database now contains 364 entries. Amplification of several additional gene sections, all of which represent promising candidate genes for canine cryptorchidism, is under establishment for downstream Nanopore sequencing.



Figure 5: Entry comparision before (green) and after the update (beige) of a) all data and b) dog associated data

Figure 2: Objective and outlook of the study (portrayed stage is boxed)

electrophoresis of INSL3 and exons 6-8 and exon 10 of RXFP2

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