

Veterinary medicine

Serum miR-30 is a potential biomarker for hepatic copper accumulation in Labrador retrievers

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Background

- **Copper associated hepatitis in Labrador retrievers** is a hereditary disease with a complex genetic background and a female predisposition
- The disease is progressive:
 - Stage 1: copper build-up in the liver
 - Stage 2: Development of chronic hepatitis
 - Stage 3: Liver failure and liver cirrhosis
- Treatment outcome is highly dependent on early diagnosis
- **Diagnosis is often delayed** due to a late onset of clinical signs and because a **liver biopsy is needed** for diagnosis
- Liver biopsies are invasive for the patient and costly for the owner
- Treatment monitoring needs regular liver biopsies
- A **blood biomarker specific for hepatic copper levels** would simplify diagnosis and treatment monitoring
- MiRNAs are non-coding RNAs which regulate posttranscriptional gene expression and are released from tissues into the bloodstream
 - > Increasingly studied as **biomarkers for hepatic disease**
 - Potentially cheap and non-invasive biomarkers for diagnosis and disease monitoring



Figure 1 Liver cirrhosis due to copper associated hepatitis

Aim: To identify serum miRNAs associated with hepatic copper accumulation in Labrador retrievers

Results

- The miRNA array detected 186 (67.1%) from the 277 tested miRNAs
- From these 186 miRNAs, 6 were selected to be validated in the replication cohort
- miRNA miR-30b was significantly associated with hepatic copper levels in both the array cohort and the replication cohort

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p < 0.01

Materials and methods

- Study population selection:
- Labrador retrievers from a patient database with liver biopsies
- Control: Normal hepatic copper levels and no liver disease
 - Case: Elevated hepatic copper levels with and without hepatitis
 - All dogs: Serum stored at -80 °C available

Canine miRNA PCR array of 277 miRNAs in 5 cases and 5 controls

 Statistics: △△CT method with Student's t-test
Selection of upregulated miRNAs: >2-fold upregulated and p-value < 0.05
>4-fold upregulated



Figure 2 MiRNA 30b Gene Expression Values detected in the case (n=5) and control (n=5) group in the Canine miRNA PCR array



Replication of selected upregulated miRNAs in 18 cases and 13 controls

- Analysis of miRNA quantity: Quantitative Real-time PCR
- Statistics: \(\Delta\Delta\CT\) method with Mann-Whitney U test and Bonferroni correction for multiple testing, adjusted p-value <0.05 was considered significant

	Results array		Results replication cohort		
MiRNA	Fold change	p-value	Fold change	Unadjusted p-value	Adjusted p-value
miR-28	9.51	0.061	1.27	0.068	0.34
miR-30b	2.04	0.0090	2.17	0.00035	0.0018
miR-331	4.13	0.73	1.39	0.046	0.23
miR-411	10.58	0.041	Not detected	NA	NA
miR-502	6.02	0.0027	1.59	0.022	0.11
miR-551b	8.17	0.020	1.18	0.12	0.60

Table 1 Results from the six miRNAs that were either >2-fold upregulated with a p-value < 0.05 or were >4-fold upregulated in the cases compared to the controls in the miRNA array. Results are shown for the array and replication analysis.

Figure 3 MiRNA 30b Gene Expression Values detected in the case (n=18) and control (n=13) group in the replication cohort

Discussion

- Interestingly, MiR-30b hasn't been previously identified as a miRNA with a role in copper metabolism, this study indicates a previously unknown gene regulation function of this miRNA.
- A limitation of our study was the sample size and an overrepresentation of female cases due to limited patient availability for this study.

Conclusion: MiRNA miR-30b is a promising biomarker for hepatic copper levels in Labrador retrievers which warrants further research in a larger cohort to determine specificity and sensitivity for elevated hepatic copper levels in dogs.

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