

# Longitudinal T<sub>2</sub> relaxometry to monitor repeated cycles of DSS inducing a chronically relapsing inflammation

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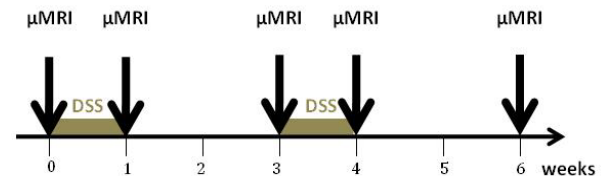
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## Introduction

Most experimental animal models of inflammatory bowel diseases (IBD) fail to accurately reflect the chronically relapsing inflammation underlying the complications of human Crohn's disease. This study investigated whether repeated cycles of dextran sulfate sodium salt (DSS) adequately reflect the effects of chronic transmural healing compared to the acute murine (DSS) colitis model with recovery. Longitudinal T<sub>2</sub> relaxometry at 9.4T was used to monitor the transmural changes during a 2 cycle treatment with DSS.

## Materials & Methods

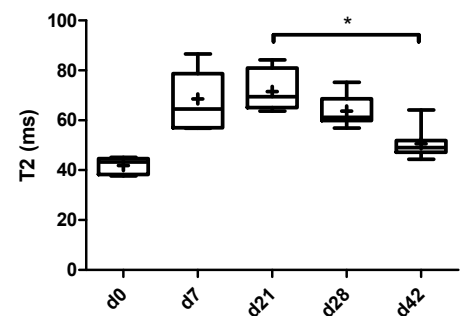
DSS colitis was induced in 6 week-old C57BL6/J mice: a first group (n=8) received 2 cycles of 7 days of DSS followed by 2 weeks of normal drinking water. For histological evaluation, qRT-PCR and macroscopic scoring of the colon, a second group (n=4) received 7 days of DSS followed by a 5 weeks recovery period and third control group (n=3) received normal drinking water only. Mice from the 2 cycle group were monitored longitudinally (day 0, 7, 21, 28 and 42) on a 9.4T MRI system (Bruker Biospin; horizontal bore, 20 cm) equipped with an actively shielded gradient insert (1200mT/m) and using a 3.5 cm volume resonator (Rapid Biomedical). T<sub>2</sub> weighted images and T<sub>2</sub> maps of the distal colon were recorded (T<sub>2</sub>w: RARE; TE<sub>eff</sub>=52ms, TR=4800ms, FOV=3x3cm, matrix= 256x256; slice thickness = 1mm T<sub>2</sub>map:TE=10-100ms, matrix 128x128). Maps were calculated using a plug-in for ImageJ [1]. Regions of interest delineating the colon wall were manually drawn on the T<sub>2</sub>w images, acquired with the same geometry as the T<sub>2</sub> map, and then copied to the T<sub>2</sub> map to determine mean T<sub>2</sub> values. After scanning and euthanasia, the distal colon was harvested for histology and qRT-PCR. Collagen deposition was quantified with Martius-Scarlett-Blue staining. To compare mean T<sub>2</sub> values at different time points within the 2 cycle group a non-parametric paired (Friedman) test was used with Dunns multiple comparison correction.



## Results

The T<sub>2</sub> after 2 cycles with recovery (day 42) showed a significant decrease compared to the 1 cycle with recovery (21 days) time point (respectively, T<sub>2</sub> = 72±8ms and 51±6ms; p<0.05). T<sub>2</sub> values at day 42 also tended to be increased vs. baseline levels (day 0: T<sub>2</sub> = 42±3ms) although not statistically significant with the non-parametric Dunns test.

Mice with 2 DSS cycles had a significant shorter colon (p<0.001), higher colon weight (p=0.001) and higher colon weight/length ratio (p<0.001) vs. 1 cycle mice with recovery and control mice. The colon of the 2 cycle group had a significantly higher macroscopic score than the 1 cycle group (p<0.001). Histology showed a trend towards a higher collagen content in the 2 cycle model compared to the 1 cycle model and a significantly higher value compared to control mice (p=0.046). No significant difference was observed in TGFβ, IL13, IL13Ra1 and IL13Ra2 expression. TNFα was significantly higher in the 1 cycle model vs. 2 cycle model and controls (p=0.002).



## Discussion and Conclusion

Based on macroscopic scoring of the colon and histology the chronic repeated cycle DSS model with relapse and remission is clearly different from the acute DSS model with recovery. The in vivo T<sub>2</sub> relaxometry also showed significant differences between the first and second cycle of DSS making it a promising non-invasive assessment method in IBD. A more detailed statistical analysis relying on the full T<sub>2</sub> histogram at each time point could more accurately reveal these differences. This model opens perspectives to study the effects of healing and fibrosis in Crohn's disease.

## References

(1) D. Prodanov, MRI Processor, ImageJ Documentation Wiki, 2009, [http://imagejdocu.tudor.lu/doku.php?id=plugin:filter:mri\\_processor:start](http://imagejdocu.tudor.lu/doku.php?id=plugin:filter:mri_processor:start)