

Target Analysis

Water NMR Analysis

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Outline

1 Water NMR Fit

2 Water Calibration Constant

Update From Last Week

- I discovered that when converting the water NMR signal from mV to μV I was applying an extra power of 10
- I also found that if I first use Minuit to find initial parameters
- Then fit the NMR data using ROOT (which uses MIGARD algorithm) gives better fit results
- In addition to fit residuals, X lock in channel was also checked

NMR Water X-Y Signals

6,189 sweeps

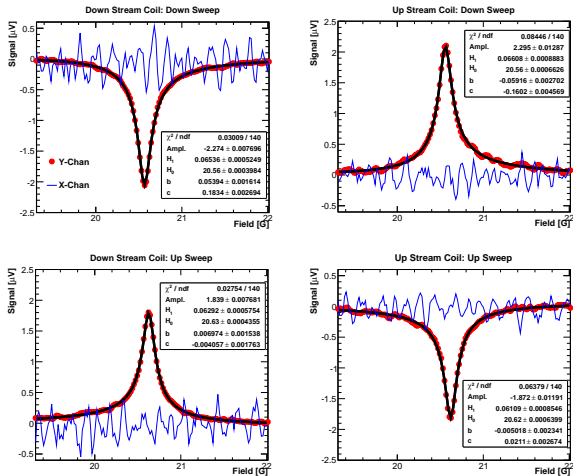


Figure: Presented are the sweep up and sweep down signals for the downstream and upstream coils. The Y lock in channel is

NMR Water Fit Residuals

6,189 sweeps

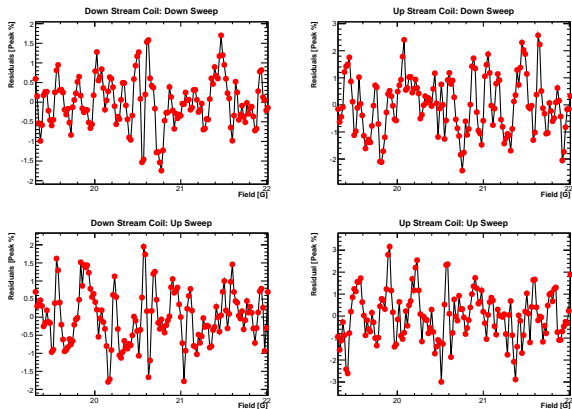


Figure: Presented are the sweep up and sweep down fit residuals for the downstream and upstream coils. The fit residual is defined as $100(\text{data} - \text{fit})/\text{peak}$.

NMR Water Coil Signals

6,189 sweeps

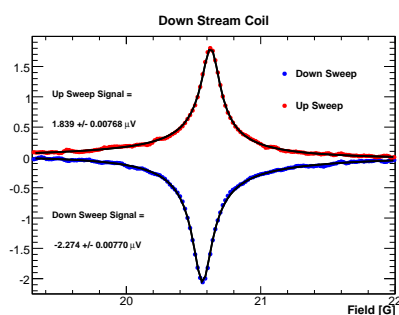


Figure: Presented are the sweep up and sweep down signals for the downstream coil. The error associated with the amplitude is statistical only.

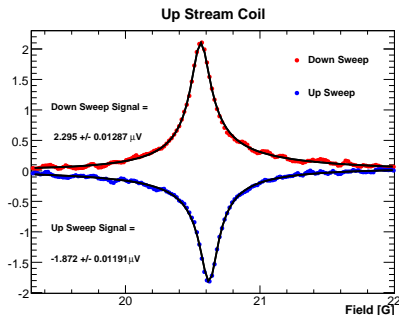


Figure: Presented are the sweep up and sweep down signals for the upstream coil. The error associated with the amplitude is statistical only.

Water NMR Fit: Summary/Comments

- Since the upstream and downstream coil measurements are in agreement we can combine like sweeps via:

$$\bar{S} = \frac{\sum_i w_i |S_i|}{\sum_i w_i}, \quad \sigma_{\bar{S}} = \sqrt{\frac{1}{\sum_i w_i}}, \quad w = \frac{1}{\sigma_S^2}$$

- Where S is the NMR signal and σ_S is the statistical uncertainty on the signal

Table: Up and down sweep signals and stat. errors from up and down stream coils and the combined coil results for 6,189 sweeps.

Signal	Amplitude [μV]	Error [μV]	Error [%]
$S_u^{(1)}$	-1.872	0.0119	0.636
$S_u^{(2)}$	1.839	0.00768	0.418
$S_d^{(1)}$	2.295	0.0129	0.562
$S_d^{(2)}$	-2.274	0.0077	0.337
\bar{S}_u	1.8487	0.00645	0.349
\bar{S}_d	2.2795	0.00661	0.290

Water NMR Fit: Summary/Comments

- Appears to be **little signal** if any in **X** channel (failed to fit NMR X signal)
- Residuals around **1.5%** of amplitude value
- Statistical errors on NMR signals **< 1%** on all sweeps and coils
- Up and down sweeps in **both coils** are in **agreement**, combine to one sweep up and sweep down measurement
- Need to determine a systematic error

Water Calibration Constant (1)

The signal induced in the pick-up coils by the rotating spins at resonance are proportional to the macroscopic magnetization of the sample:

$$S = kM = k\mu nP$$

- S : amplitude of the lock in amplifier signal
- k : proportionality constant, taking into account gains, losses in the cables etc...
- μ : magnetic moment of the sample
- n : density of spins
- P : polarization of the sample

Water Calibration Constant (2)

The ratio of the ^3He NMR signal height S_{He} to the water NMR signal height S_w can be written as:

$$\frac{S_{He}}{S_w} = \frac{G_{He}}{G_w} \frac{\mu_{He}}{\mu_p} \frac{n_{He}}{n_p} \frac{\Phi_{He}}{\Phi_w} \frac{P_{He}}{P_w}$$

- S_{He} : amplitude of helium signal
- S_w : amplitude of water signal
- G_{He} : gain of the pick-up coil preamplifier in helium configuration
- G_w : gain of the pick-up coil preamplifier in water configuration
- μ_{He} : magnetic moment of ^3He nucleus
- μ_p : magnetic moment of the proton in water
- n_{He} : density of ^3He nuclei in the helium target
- n_p : density of protons in the water target
- Φ_{He} : total magnetic flux of helium cell through pick-up coils
- Φ_w : total magnetic flux of water cell through pick-up coils
- P_{He} : polarization of the helium target
- P_w : polarization of the water sample

Water Calibration Constant: P_w

- The water polarization can be computed using [Boltzmann statistics](#)

$$P_w = \text{Tanh} \left(\frac{\mu_p H_0}{k_B T_w} \right)$$

- with k_B being the Boltzmann constant and T_w being the temperature of the water cell
- However, we can use the [Bloch equations](#) to numerically calculate the water polarization
- There are [two](#) sources of error that arise when using the Bloch equations
 - The [start](#) and [end](#) points of the integration (H-field)
 - Sweep range $H = 18 - 24$ G
 - Vary start and stop points by 2 G
 - The uncertainty in T_w
 - Still new to analyze

Water Calibration Constant: P_w Uncertainty Results

Table: Results of varying integration limits on water polarization.

H_i [G]	H_f [G]	Up Sweep Value $\times 10^{-9}$	Down Sweep Value $\times 10^{-9}$
18	24	6.59397	7.7827
18	26	6.59397	7.9373
16	24	6.47505	7.7827

Table: Uncertainty results from Bloch equation.

Parameter	Value	Units	Uncertainty Source [%]
Up sweep P_w	6.59397	-	1.8
Down sweep P_w	7.7827	-	2.0

Water Calibration Constant (3)

$$P_{He} = c_w \frac{S_{He}}{n_{He} \Phi_{He}}$$

$$c_w = \frac{1}{S_w} \frac{G_w}{G_{He}} \frac{\mu_p}{\mu_{He}} n_p \Phi_w P_w$$

Table: Parameters used to compute the water constant. Highlighted parameters still need to be calculated.

Parameter	Description	Value	Units	Uncertainty [%]	Source
S_w	Sweep Up	1.849	μV	0.349	Fit
	Sweep Down	2.280	μV	0.290	Fit
G_w	Gain of pick-up coil pre-amp. for water cell	20	-	-	-
G_{He}	Gain of pick-up coil pre-amp. for 3He cell	1	-	-	-
μ_p / μ_{He}	-	1.3127	-	neg.	-
n_p	at 22°C	2482	amg	0.1	see M. Romalis thesis
Φ_w	Magnetic flux through pick-up coil	-	cm^2	-	-
P_w	Sweep Up	6.59397	-	1.8	Model of Bloch Eqs.
	Sweep Down	7.7827	-	2.0	Model of Bloch Eqs.

Summary and To Do

- Get water cell temperature
- Fit ^3He NMR data
- Look at how acceptance holes affect asymmetries
- Begin systematic error on BB cuts (All or subset of runs?)