Testing Frozen-Spin HD Targets with Electron BEAMS

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- Developing a large acceptance transversely polarized proton target for the CLAS12 transversity program with electron beams.
- The HDice target became the prime candidate for CLAS12 transverse polarization experiments, due to its excellent performance with photon BEAMS, and Frozen-Spin properties.
- Electron beam induced polarization loss and related relaxation times, T₁, can be studied by tracking NMR signals under various beam and target conditions.















Quick test results: polarization lost in hours with unfavered beam-target conditions. (Raster rate, target temperature, spin-field alignment...)

 \Rightarrow Mechanisms for beam-induced depolarization:

- I. beam-heating
 - heat > partially polarized molecular electrons > interact with HD spins
 - solution: keep HD cold so that molecular electrons are 100% polarized and frozen
 ⇔ new target cells and new fast raster

II. Hyperfine mixing

- $\vec{\mu}(e)$ opposite to $\vec{\mu}(p)$ > polarized electrons mix and dilute H polarization
- solution: flip H spin against field, so that e and H polarizations are parallel

III. Radiation damage

- beam ionization > chemical changes that could bring HD out of frozen-spin state
- expected to be temperature dependent ⇔ needs detailed study







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Upgraded Injector Test Facility (UITF)















UITF Beamline (250keV – 9.7MeV)



Beam reastered at \sim 14 KHz with \sim 1 KHz repeating pattern.

HDice Beamline (9.7MeV)

HDice Target (1.1Tesla, 0.06K)





Rastered beam profile on a YAG viewer near the beam dump











- 10MeV electron beams were bent easily by multiple magnets along the beam path.
- Multiple scattering inside the HDice target smeared the beam further.
- Beam current determination below 1 nA was a challenge itself.
- The activities of nearby facilities in the shared building generated heavy noises electrically and mechanically.
 - \Rightarrow Large random and/or periodic NMR noises

(less in late night and weekends)

• Local temperature of HD varies as electrons passed through.











Results: NMR Signals





Proton Signal from PD, ~40% Polarization



Typical TE Signal from IBC, 170 scans, Ie=0pA





Proton Signal from IBC, ~40% Polarization, $I_{\rho} = 0 \text{ pA}.$ Proton Signal from IBC, ~40% Polarization, $I_e = 125 \text{ pA}.$

Periodic Noise Removal for X Channel



Down Absorption Signal of NMR Run # 73667103 Periodic Noise Baseline Raw Down Absorption Signal Treated Signal (Raw Signal - Periodic Baseline) Amplitude (V) Sign 1000 2000 3000 4000 5000 6000 7000 0 8000 Channels

Signal with periodical background

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Results: Polarization Tracking for Target #60 and #66 close



P vs. Time

P vs. Total Dose





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Results: CORE-RING-SKIN MODEL

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3 dose ranges (μC) (0<=D<20, 20<=D<60 and D>=60):

CORE (r<Rcore, uniform beam) RING (Rcore<=r<Rskin, shoulder beam) SKIN (Rskin<=r<RHD, no beam)

The polarization vs dose in Run 3B can be described as:

- 1. P=-1.39D+34 (0<=D<20, CORE+RING)
- 2. P=-0.124D+7.00 (21<=D<60, RING)
- 3. P=0 (D>=60, SKIN) (P=1.66 without SKIN subtraction).
- The relative portions of HD target in each region: CORE/total=0.756 RING/total=0.196 SKIN/total=0.048

 $\Rightarrow \text{RHD}=9.50\text{mm}, \text{Rcore}=8.26\text{mm}, \text{Rskin}=9.27\text{mm}.$ $\Rightarrow \text{The 3 HD volumes contributed NMR signals:}$ were 2.679cc, 0.695cc and 0.173cc for the L=12.5mm target.

The Run 3A were treated the same way.











Run 3A, TGT #60

Run 3B: Ring Polarization VS Ring Dose/Area



Run 3B, TGT #66

Target polarization lost (almost) linearly as Areal Dose accumulated.

$$\frac{dP}{d(\frac{Dose}{Area})} \cong 3.8\% \ ^{cm^2}/_{\mu C}$$

For both CORE and RING

(The difference between 3A and 3B at CORE region was caused by different raster patterns.)







• Starting with a model developed by C. Keith (SPIN 2016) and rewriting time in terms of the relative dose received during a beam condition,

$$P(t) = (P_0 - P_{TE})e^{-\frac{\alpha * D_{rel} * D_T * \beta}{I_e}} + P_{TE}$$

where $\beta = B^3 e^{-\frac{2\mu_e B}{kT}}$ and $t = \frac{D_{rel}}{I_e}$

the value of α can be found by fitting it with the expression.

• Once α is determined for each condition (both core and ring), a global alpha found from the weighted average of all individual values.







Core Simulation Using Separate Condition Alphas (Top) Core Simulation Using Global Alpha (Bottom)













Ring Simulation Using Separate Condition Alphas (Bottom) Ring Simulation Using Global Alpha (Bottom)



Run 3A, TGT #60





Run 3B: Measured Polarization of the Ring Section



Run 3B, TGT #66





T_{HD} Calibration Curve from HDice Run 2A & 2B



• Temperature calibration for T_{HD} required for beam on NMR measurements.

(Run 2A and 2B, 2 TE targets, $T_1^H < 1s$)

 From Curie's Law <> The product of NMR Signals from beam-off conditions and the HD temperature were calculated across the Run 3A run period:

$$S_{(beam-on)} * T_{HD(beam-on)} = S_{(beam-off)} * T_{HD(beam-off)}$$

 The HD temperature at various beam currents is determined from the relationship above:

$$T_{HD(beam-on)} = \frac{S_{(beam-off)} * T_{HD(beam-off)}}{S_{(beam-on)}}$$









• The model was used to determine the scaling factor between the HDice T_1 values and the T_1 value from the Mano and Honig experiment.

$$\frac{T_1^{H_I}}{T_1^{H_{II}}} = \frac{\left(\frac{1}{(B_I = 1.04T)^3}\right) \exp\left(\frac{2\mu_e(B_I = 1.04T)}{k(T_I = 0.15K)}\right)}{\left(\frac{1}{(B_{II} = 0.28T)^3}\right) \exp\left(\frac{2\mu_e(B_{II} = 0.28T)}{k(T_{II} = 4.2K)}\right)} = 10.30$$

- The scaling factor was determined to be an order of magnitude, which isn't enough to account for the difference between the results.
- The remaining difference could be due to a higher effective dose during the Mano and Honig experiment, due to the production of secondaries in the relatively thick walls of their cryostat.











The same model used to scale Mano and Honig's T_1 values can be used to project the relaxation times for ideal conditions.

If the HD temperature and field were maintained at 120mK and 1.25 T, the T_1 values increase by:



Potential T_1 enhancements, are quite large (~10⁴ times larger), but the beam-off T_1 dominated by another polarization loss mechanism. This imposes a limit on the gains in the beam-on T_1 that might be achieved with more ideal conditions.









- During the initial $\sim 1.9 \mu C/cm^2$ for both targets, the polarization loss was insignificant.
- By 6µC/cm² the polarization of the each section dropped to 1/e of their initial values with complete polarization loss at ~10µC/cm².
- Beam-on spin-relaxation rates (T₁) dropped with dose. Beam-off T₁ values followed similar trend, but about 10x larger.
- With both Run 3A and 3B data, it became evident that at a fixed dose, T₁ also drops with increasing current.
- The *eHD* experiments showed significantly better performance than the Mano and Honig experiment.
- Improvements to Beam On T_{HD} and B could greatly improve the T1, but the additional Beam Off depolarization mechanism limits the potential gains.
- 2 PhD theses produced:
 - Kevin Wei, Univ. of Connecticut, 2021.

"The Response of Polarized Protons in Solid Hydrogen-Deuteride(HD) to Electron Beams"

Thomas O'Connell, Univ. of Connecticut, 2022.

"Measurements of Electron Beam Induced Spin Relaxation in Frozen Spin Hydrogen Deuteride HD"





Φ^{ice} Extra: Relaxation Time (T₁) Values in Runs 3A & 3B closs

• The polarization relaxation times (T_1) could be determined by fitting the polarization values of each of the beam on/off conditions to the following equation:

$$P(t) = (P_0 - P_{TE})e^{-t/T_1} + P_{TE}$$

• A difference of an order of magnitude seen between the T_1 values from beam-on and beam-off conditions.





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$\mathbf{\Phi}^{ice}$ Extra: Relaxation Time (T₁) Values in Runs 3A & 3B closs



Run 3A and 3B: All Beam-On Core T1 Values

100 10 9 83pA T1 (Days) 125pA 167pA 250pA 0.1 0.01 0 2 6 8 10 Dose/Area (µC/cm²)

Runs 3A and 3B: Relaxation Times VS Average Beam Current at an Accumulated





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Extra: Beam Profile at Target, viewer, and along axis close





