28.1 A Handheld 50pM-Sensitivity Micro-NMR CMOS Platform with B-Field Stabilization for Multi-Type Biological/Chemical Assays

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Point-of-use (PoU) biological/chemical assays are aimed to transform bulky laboratory instruments into easy-to-use lab-on-a-chip platforms, bringing down the cost, size, and sample-use by orders of magnitude [1,2]. Micro-Nuclear Magnetic Resonance (NMR) is a trail-blazing tool for *target* pinpointing, by utilizing functionalized magnetic nanoparticles (MNPs) as the *probe* [3]. Screening by micro-NMR is repeatable, versatile and low-cost as it is label- and washing-free for the samples, and immobilization-free for the electrodes. Herein, a high-sensitivity micro-NMR CMOS platform with magnetic (B)-field stabilization and thermal management is reported (Fig. 28.1.1). This handheld tool unifies multi-type assays (target detection, protein state analysis, and solvent-polymer dynamics), and is suitable for healthcare, food industry, and colloidal applications.

Micro-NMR relaxometry detects the spin-spin relaxation time (T_2) by extracting the echoes envelopes from the response of the non-zero spin nuclei (i.e., 'H). The nuclei, under magnetization with a static magnetic field (B_0), absorb orthogonal RF exciting magnetic field (B_1) at the Larmor frequency, $f_1 = \gamma B_0$ (v: gyromagnetic ratio), and precess about the direction of magnetization at f_1 even after the cessation of the excitation. In an existing micro-NMR system [3], frequency deviation of the local oscillator (LO) from f_1 induces improper frequency excitation, paralyzing the operation. Confounded by the thermal instability of the portable magnet (B_0 =0.46T, T.C.=-1200ppm/K), LO tracking is essential to safeguard the system against environmental changes.

Our micro-NMR platform (Fig. 28.1.2) is tailored with B_0 -field stabilization and thermal management to enhance the robustness and simplify the hardware. The dynamic B_1 -field transduction is based on a spiral coil driven by a transmitter (TX)/receiver (RX) together with a matching capacitor $C_{\rm M}$, to excite/obtain the magnetic signal to/from the droplet samples (2.5µL) normal to the chip surface. The TX is based on a tapped inverter-chain power amplifier (PA), measured 31.6% power efficiency, to deliver programmable pulse sequences pertaining to the LO. The RX features a multi-stage low-noise amplifier (LNA) for high RX sensitivity (down to 1nV/ $\sqrt{\rm hz}$ input-referred-noise), and a dynamic-bandwidth lowpass filter for fast recovery from saturation after excitation pulses. The B_0 -field sensor and calibrator manage the lateral B_0 -field together with a current driver, which injects a calibration current to the magnet (75mT/A) stabilizing the bulk magnetization on the nuclei. The spiral coil also serves as a heater allowing thermal profiling of the samples. The thermal-induced error on the B_0 -field sensor and calibrator, and the hotness of the samples, are monitored by a BJT temperature sensor.

To sense the lateral B_0 -field normal to the chip surface, a *current-mode* 4-folded vertical hall sensor (VHS) arranged in a Wheatstone bridge is employed (Fig. 28.1.3). Each VHS element is composed by an n-well as the substrate and three n-diffusions as contacts [4]. P-diffusions are embedded between the n-diffusions to avert current flowing at the surface, soothing the 1/f noise. To achieve sub-nA sensitivity, the VHS readout circuit (Fig. 28.1.3) is based on a low-noise TIA. Current-spinning and chopper are applied reducing the 1/f noise corner by >5,000×. Switches S_{1-8} control the flows of the current and reset the capacitors C_F . Small switches (280 Ω each) can exacerbate the impedance of the TIA, ($R_{\text{in,TIA}}$ =210 Ω) if there is current passing through the switches connected between the core OTA of the TIA and VHS (i.e., S_{5-6}). To address this, S_{7-8} are managed to guide the current passing through the negative feedback path, nullifying the impact of resistances of S_{5-6} on the TIA. Thanks to this switching scheme, $R_{\text{in,TIA}}$ is suppressed by 84%, absorbing ~21% more current into the TIA than the general approach [5].

Attributed to the prodigious nominal B_0 -field, a typical TIA can be saturated and fail to sense the tiny B_0 -field variation (3.75mT). To solve it, a nominal B_0 -field compensator made by a passive switched-capacitor network (Fig. 28.1.3) nullifies the nominal B_0 -field entering into the TIA.

Before the micro-NMR assay, the VHS reads B_0 and responds to the current driver (Fig. 28.1.4). B_0 may shift away from its nominal value due to the environmental changes (e.g., temperature and sample-to-magnet position). Thus, untracked f_L can be easily off-center from the LO frequency $f_{\rm OSC}$ (BW=16.7kHz), Here, by modulating the magnet according to an updated B_0 (sensitivity: 4.12V/T), f_L is reset to $f_{\rm OSC}$. Also, with signal-averaging performed in the frequency domain to suppress the background noise, the calibration improves the B_0 -field stability by 13× (from 2 to 0.15mT) at 0.46T (f_L =19.6MHz). Under the synergy of micro-NMR and VHS, the stabilized f_L inspires the use of a simple crystal oscillator as the LO that measures low phase noise (-116dBc/Hz at 1kHz offset) at very low power (79 μ W).

Human Immunoglobulin G (IgG), which protects the body from infections, can be quantified by utilizing Protein A coated water-soluble MNPs (i.e., Fe₂O₃) based on their T_2 . T_2 of the sample is shortened commensurate with the amount of IgG upon nanoparticles agglomeration, enabling quantification of IgG down to 5nM (Fig. 28.1.5). The specificity of micro-NMR assay is evinced with the addition of Chicken Immunoglobulin Y (IgY), which does not conjugate with Protein A. The negligible change of T_2 (<2%) validates the selectivity of the assay. The versatility of the platform is manifested with DNA detection apt for life-threatening bacteria screening. With a pair of probe-decorated MNPs, the platform quantifies the synthesized DNA derived from *Enterococcus faecalis*, with a detection limit down to 50pM in 2.5µL samples (125amol). By varying the MNP concentration, the detection range is impelled to 125nM. The response to single-nucleotide polymorphism is indistinguishable to T_2 baseline (<4%), substantiating that single-base mismatch DNA can be differentiated.

Probing the molecular structure can digest the protein state for food quality inspection. Protein β -lactoglobulin (β -LG) denatures and aggregates irreversibly after heating to >60°C [6]. This state transformation can be embodied by measuring \mathcal{T}_2 of the samples attributed to the dissimilar interaction between the water molecules and protein at different states and sizes (Fig. 28.1.6). For the colloidal industry, Poly(N-isopropylacrylamide) (PNIPAM) is widely used as advanced sensor and drug delivery carrier [7]. It is a colloidal polymer that exhibits a temperature-induced reversible volume phase transition in water, affects the local environment on solvent confinement and thus \mathcal{T}_2 of the solvent. By duty-cycling the heater (coil), PNIPAM undergoes a volume phase transition above 33°C, resulting in \mathcal{T}_2 decrement of the solvent.

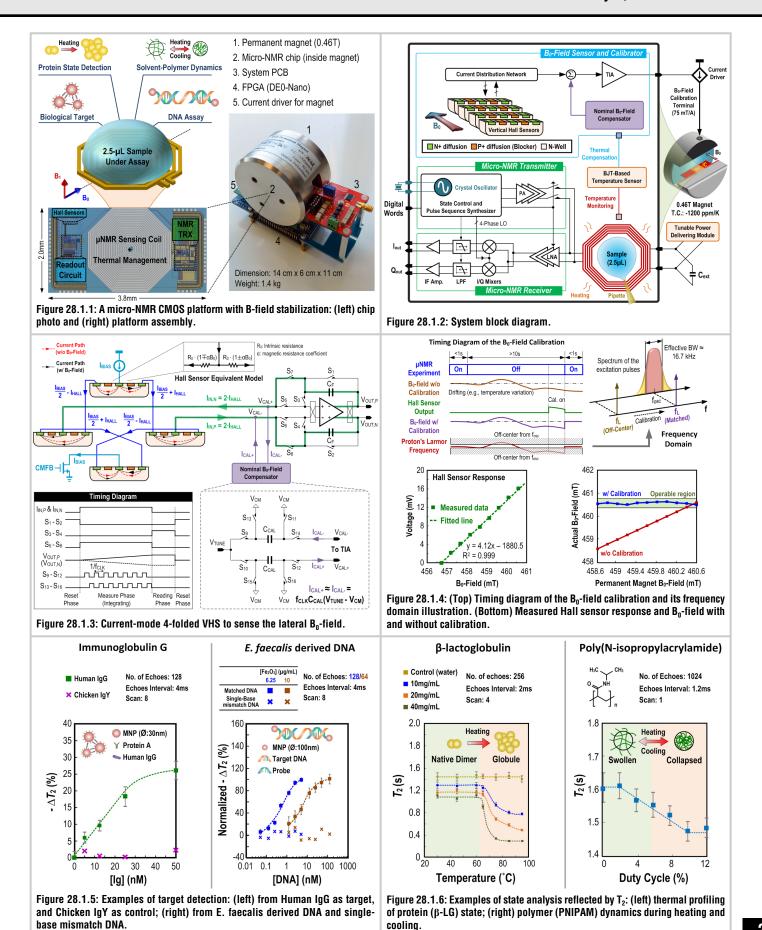
Benchmarking with the recent PoU tools (Fig. 28.1.7), this work supports multitype assays in one unified platform, while achieving high sensitivity and selectivity for DNA, as well as other proteins targeting capability in tiny sample with functionalized MNPs. The platform consumes $120\times$ less samples, and is $96\times$ lighter, $175\times$ smaller, and $16\times$ cheaper than a commercial product (Bruker mq-20 [8]).

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ISSCC 2016 PAPER CONTINUATIONS

į	This Work	PH. Kuo et al., ISSCC'15	KH. Lee et al., ISSCC'12	N. Sun et al., ISSCC'10	B. Jang et al., ISSCC'09
Specificity	Target detection Solvent-Polymer dynamics	Target detection	Target detection	Target detection	Target detection
TargetLabeling	3. Protein state analysis Label-free	Label-free	Label-free	Label-free	Cy3-label
Demo Target	68 base E. faecalis derived DNA	NT-ProBNP &	21 base H5N1 virus	hCG cancer marker	18 base DNA
Detection Limit	50 pM	TNF-alpha	100 pM	5000 pM	125 pM
Sample Handling	(DNA)		(DNA)	(Cancermarker)	(DNA)
Limit Sample Handling	2.5 µL	-	-	5.0 µL	
Physics	NMR relaxometry + Thermal management	Magnetic- sensing	Capacitance- sensing	NMR relaxometry	Fluorescent- sensing
Post-Fabrication Necessity	No (immobilization free)	Probe (antibody) immobilization	Probe (DNA) immobilization on Au electrodes	No (immobilization free)	Fiber-optical faceplate
External Part	Portable magnet	No	No	Portable magnet	Lightsource
LO Generation	Crystal oscillator+ B₀-field calibration	-	-	Off-chip	-
Robustness to	Robust to	Robust to	Vulnerable to	Vulnerable to	Vulnerable to
environments	temperature & sample position variations	temperature variation	bias current variation	B-field variation	background noise
CMOST ech.	0.18 μm	0.35 µm	0.35 µm	0.18 µm	0.35 µm
Chip Area	7.6 mm²	8.9 mm ²	20.0 mm ²	11.3 mm ²	9.0 mm ²
Figure 28 1 7	: Benchmark with o	ther CMOS-	hased Poll	systems	