Monitoring ion-channel function in real-time through quantum decoherence

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In drug discovery, there is a clear and urgent need for detection of cell-membrane ion-channel operation with wide-field capability. Existing techniques are generally invasive or require specialized nanostructures. We show that quantum nanotechnology could provide a solution. The nitrogen-vacancy (NV) center in nanodiamond is of great interest as a single-atom quantum probe for nanoscale processes. However, until now nothing was known about the quantum behavior of a NV probe in a complex biological environment. We explore the quantum dynamics of a NV probe in proximity to the ion channel, lipid bilayer, and surrounding aqueous environment. Our theoretical results indicate that real-time detection of ion-channel operation at millisecond resolution is possible by directly monitoring the quantum decoherence of the NV probe. With the potential to scan and scale up to an array-based system, this conclusion may have wide-ranging implications for nanoscale biology and drug discovery.

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The cell membrane is a critical regulator of life. Its importance is reflected by the fact that the majority of drugs target membrane interactions (1). Ion channels allow for passive and selective diffusion of ions across the cell membrane (2), whereas ion pumps actively create and maintain the potential gradients across the membranes of living cells (3). To monitor the effect of new drugs and drug delivery mechanisms, a wide-field ion-channel monitoring capability is essential (4). However, there are significant challenges facing existing techniques stemming from the fact that membrane proteins, hosted in a lipid bilayer, require a complex environment to preserve their structural and functional integrity (1, 5–7). Patch clamp techniques are generally invasive, quantitatively inaccurate, and difficult to scale up (8, 9), whereas black lipid membranes (10, 11) often suffer from stability issues and can only host a limited number of membrane proteins.

Instead of altering the way ion channels and the lipid membrane are presented or even assembled for detection, our approach is to consider a unique and inherently noninvasive in situ detection method based on the quantum properties of a single-atom probe. The atomic probe is a single nitrogen-vacancy (NV) center in a nanodiamond crystal that is highly sensitive to magnetic fields and shows great promise as a magnetometer for nanobiosensing (12–19). The NV center in nanodiamond has already been used as a fluorescence marker in biological systems (20–24). However, up to now there has been no analysis of the effect of the biological environment on the quantum dynamics of the NV center—such considerations are critical to nanobiomagnetometry applications. We explore these issues in detail and, furthermore, show that the rate of quantum decoherence of the NV center is sufficiently sensitive to the flow of ions through the channel to allow real-time detection, over and above the myriad background effects. In this context, decoherence refers to the loss of quantum coherence between magnetic sublevels of the NV atomic system due to interactions with an environment. Such superpositions of quantum states are generally fleeting in nature due to interactions with the environment, and the degree and timescale over which such quantum coherence is lost can be measured precisely. The immediate consequence of the fragility of the quantum coherence phenomenon is that detecting the loss of quantum coherence (decoherence) in such a single-atom probe offers a unique monitor of biological function at the nanoscale (18, 19).

The NV probe (Fig. 1) consists of a diamond nanocrystal containing an NV defect at the end of an atomic force microscope tip, as recently demonstrated (16). For biological applications, a quantum probe must be submersible to be brought within nanometers of the sample structure, hence the NV system locked and protected in the ultrastable diamond matrix (Fig. L4) is the system of choice. The NV center alone offers the controllable, robust, and persistent quantum properties such room temperature nanosensing applications will demand, as well as zero toxicity in a biological environment (20–22). Theoretical proposals for the use of diamond nanocrystals as sensitive nanoscale magnetometers (12–14) have been followed by proof-of-principle experiments (15–17). However, such nanoscale magnetometers employ only a fraction of the quantum resources at hand and do not have the sensitivity to detect the minute magnetic moment fluctuations associated with ion-channel operation. In contrast, our results show that measuring the quantum decoherence of the NV induced by the ion flux provides a highly sensitive monitoring capability for the ion-channel problem, well beyond the limits of magnetometer time-averaged field sensitivity (19). To determine the sensitivity of the NV probe to the ion-channel signal, we describe the lipid membrane, embedded ion channels, and the immediate surroundings as a fluctuating electromagnetic environment and quantitatively assess each effect on the quantum coherence of the NV center. We consider the diffusion of nuclei, atoms, and molecules in the immediate surroundings of the nanocrystal and the extent to which each source will decohere the quantum state of the NV. We find that, over and above these background sources, the decoherence of the NV spin levels is highly sensitive to the ion flux through a single ion channel. Our theoretical findings demonstrate the potential of this approach to revolutionize the way ion channels and potentially other membrane-bound proteins or interacting species are characterized and measured, particularly when scaleup and scanning capabilities are considered.


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NV system is that, under optical excitation, the spin levels are distinguishable by a difference in fluorescence, hence spin-state readout is achieved by purely optical means (25, 26). These properties are fortuitous in the current context because mammalian ion-channel function is known to be insensitive to optical light (27). Because of the simple readout and control, the quantum properties of the NV system and the interaction with the immediate crystalline environment are well known (28, 29). The coherence time of the spin levels is very long even at room temperature: In type 1b nanocrystals $T_2 \sim 15 \mu$s (30), and, in isotopically engineered diamond, can be as long as 1.8 ms (17) with the use of a spin-echo microwave control sequence (Fig. 2C). More encouraging is that these times are predicted to be as high as 75 $\mu$s and 100 ms, respectively, with the application of optimal microwave pulse sequences (31). Nanocrystals of 5-nm diameter containing stable NV centers have recently been demonstrated (32).

Typical ion-channel species $K^+$, $Ca^{2+}$, $Na^+$, and nearby water molecules are electron spin paired, so any magnetic signal due to ion-channel operation will be primarily from the motion of nuclear spins. Ions and water molecules enter the channel in thermal equilibrium with random spin orientations and move through the channel over a microsecond timescale. The monitoring of ion-channel activity occurs via measurement of the contrast in probe behavior between the on and off states of the ion-channel. This protocol requires the dephasing due to ion-channel activity to be at least comparable to that due to the fluctuating background magnetic signal. We must therefore account for the decoherence due to the diffusion of water molecules, buffer molecules, saline components, as well as the transversal diffusion of lipid molecules in the cell membrane.

The $n$th nuclear spin with charge $q_n$, gyromagnetic ratio $\gamma_n$, velocity $v_n$, and spin vector $S_n$ interacts with the NV spin vector $P$ and gyromagnetic ratio $\gamma_P$ through the time-dependent dipole-dominated interaction

$$H_{\text{int}}(t) = \sum_{n=1}^{N} \kappa_{\text{tip}}^{(0)} \frac{\hat{P} \cdot \hat{S}_n(t)}{r_n^3(t)} - 3 \hat{P} \cdot \hat{S}_n(t) \frac{\hat{S}_n(t)}{r_n^3(t)}, \tag{1}$$

where $\kappa_{\text{tip}}^{(0)} = \frac{e}{2\hbar} \gamma_P \gamma_n$, are the probe-ion coupling strengths, and $r_n(t)$ is the time-dependent ion-probe separation. Additional Biot–Savart fields generated by the ion motion, both in the channel and the extracellular environment, are several orders of magnitude smaller than this dipole interaction and are neglected here. Any macroscopic fields due to intracellular ion currents are of nanotesla order and are effectively static over $T_2$ timescales. These effects will thus be suppressed by the spin-echo pulse sequence.

In Fig. 3, we show typical field traces at a probe height of 1–10 nm above the ion channel, generated by the ambient environment and the onset of ion-flow as the channel opens. The contribution to the net field at the NV probe position from the various background diffusion processes dominate the ion-channel signal in terms of their amplitude. Critically, because the magnetometer mode detects the field by acquiring phase over the coherence time of the NV center, both the ion-channel signal and background are well below the nanotesla Hz$^{-1/2}$ sensitivity limit of the magnetometer over the (~1 ns) self-correlated timescales of the environment. However, the effect of the various sources on the decoherence rate of the NV center are distinguishable because the amplitude-fluctuation frequency scales are very different, leading to remarkably different dephasing behavior.

To understand this effect, we need to consider the full quantum evolution of the NV probe. In the midst of this environment, the probe’s quantum state, described by the density matrix $\rho(t)$, evolves according to the Liouville equation,

$$\frac{d}{dt} \rho(t) = -\frac{i}{\hbar} [H(t), \rho(t)] - \frac{1}{\tau} (\rho(t) - H(t)), \tag{2}$$

where $\rho(t)$ is the incoherent thermal average over all possible unitary evolutions of the entire system, as described by the full Hamiltonian, $H = H_{\text{int}} + H_{\text{g}}$.
$H_{bg}$, where $H_{bg}$ is the Hamiltonian of the NV system, and $H_{int}$ describes the interaction of the NV system with the background environment (e.g., diffusion of ortho spin water species and ions in solution) and any intrinsic coupling to the local crystal environment. The self-evolution of the background system is described by $H_{bg}$, which, in the present methodology, is used to obtain the noise spectra of the various background processes. We note that the following analysis assumes dephasing to be the dominant decoherence channel in the system. We ignore relaxation processes because all magnetic fields considered are at least 4 orders of magnitude less than 2.88 GHz, and are hence unable to flip the probe spin over relevant timescales. Phonon excitation in the diamond crystal may also be neglected (17). Before moving onto the numerical simulations, we consider some important features of the problem.

The decoherence rate of the NV center is governed by the accumulated phase variance during the control cycle. Maximal dephasing due to a fluctuating field will occur at the cross-over point between the fast and slow fluctuation regimes (19). A measure of this point is the dimensionless ratio $\Gamma \equiv f_e/\tau_r \sigma_B$, where $\tau_r = 1/f_e$ is the correlation time of the fluctuating signal, with cross-over at $\Gamma \sim 1$.

In what follows, we consider exclusively the behavior of a sodium ion channel. Sodium-23 has an effective abundance of 100% and a nuclear magnetic moment of $\mu_{Na} = 2.22 \mu_N$, where $\mu_N = 5.05 \times 10^{-27}$ JT$^{-1}$ is the nuclear magneton. Other magnetically active ion-channel species include potassium-39, having a 93.1% abundance and a nuclear magnetic moment of 0.39 $\mu_N$, and chlorine-35, having a 75.4% abundance and a nuclear magnetic moment of 0.82 $\mu_N$, ensuring sodium ions will interact most strongly with the NV center. We can estimate the field standard deviation $\sigma_B^2$ due to the random nuclear spin of ions and bound water molecules moving through an ion channel (ic) as

$$\sigma_B^2 \sim \frac{1}{4\pi \hbar} \sqrt{N_{Na} \mu_{Na}^2 + N_{H_2O} \mu_{H_2O}^2}$$

where $N_{Na}$ and $N_{H_2O}$ are the average numbers of sodium ions and water molecules inside the channel. By modeling the channel as a cylinder with typical sodium channel influx/outflux rates (33), we may numerically calculate the rms fluctuation strength of the ion-channel magnetic field, $\sigma_B^2$, as a function of the probe standoff distance $h_p$, as shown in Fig. 3B. The fluctuation rate is defined by the rate at which ions move through the channel and is independent of whether the ions are moving into or out of the cell. However, in typical physiological processes (neuronal firing for example), the sodium flux will be inward. Ion flux rates give an effective fluctuation rate of $f_e^p \sim 3$ MHz (33). For probe-channel separations of 2–8 nm, values of $\sigma$ range from 0.4 to 3.4 GHz (Fig. 3C). Thus, the ion-channel flow hovers near the cross-over point, with an induced dephasing rate of $\Gamma_e \sim 10^{-4}–10^4$ Hz.

At these separation distances, the presence of the diamond surface is expected to have a negligible effect on the ion-channel dynamics. To reduce the decoherence effects of surface spins on the NV, the surface may be terminated with H or OH moieties. This surface termination essentially replaces the electron spins associated with the sp2 hybridized orbitals, with weaker nuclear spins (32). The ions in the channel, also being nuclear spins, couple very weakly to their surrounding environment. We may estimate their coupling to the diamond surface as $f_e^p \sim \frac{\mu_{Na}^2}{4\pi \hbar} \mu_B^2 h_p^2 / L^2$, where $L$ is the typical distance between adjacent ions in the channel. Additionally, we may approximate the ratio between the magnetic force on the ions due to the surface spins and the electric force due to adjacent ions as $f_B/F_e = (3\pi\epsilon_0)\mu_B^2 h_p^2 / (\hbar c^2 \Delta r^2) \sim 10^{-14}$, where $\Delta r$ is the typical distance between adjacent ions in the channel. Similarly, the ratio between the magnetic force due to the NV spin and the electric force is $f_B/F_e = (3\pi\epsilon_0)\mu_B\mu_B\mu_B h_p^2 / (\hbar c^2 \Delta r^2) \sim 10^{-12}$, thus we expect the presence of the probe to be truly noninvasive.

We now consider the dephasing effects of the various sources of background magnetic fields. The first source of background noise is the fields arising from the motion of the water molecules and ions throughout the aqueous solution. Due to the nuclear spins of the hydrogen atoms, liquid water consists of a mixture of spin neutral (para) and spin-1 (ortho) molecules. The equilibrium ratio of ortho to para molecules (OP ratio) is 3:1 (34), making 75% of water molecules magnetically active. In biological conditions, dissolved ions occur in concentrations 2–3 orders of magnitude above this number density and are ignored here (they are important however for calculations of the induced Stark shift; see below). The rms strength of the field due to the aqueous solution is $\sigma_{H_2O}^B = g_1 \mu_B \mu_B \hbar / 2 \sqrt{\hbar \rho_{H_2O}}$, giving a fluctuation rate of $f_e^{H_2O} \sim \frac{\mu_B^2 \mu_B^2 h_p^2}{\hbar c^2 \Delta r^2} \sim 10^{-10}$ Hz and corresponding dephasing envelope $\varphi_{H_2O} = e^{-\Gamma_e^{H_2O} f_e^{H_2O}}$.

An additional source of background dephasing is the lipid molecules comprising the cell membrane. Assuming magnetic contributions from hydrogen nuclei in the lipid molecules, lateral diffusion in the membrane gives rise to a fluctuating B field, with a characteristic frequency related to the diffusion rate. Atomic hydrogen densities in the membrane are $n_H = 3 \times 10^{26}$ m$^{-3}$. At room temperature, the populations of the spin states of hydrogen will be equal, thus the rms field strength is given by $\sigma_h \sim g_B \mu_B \mu_B \hbar / 4 \sqrt{\pi g_0 n_H}$, where $g_0$ is the response factor of the hydrogen spin system. The strength of the fluctuating field due to the lipid bilayer is of the order of $10^{-7}$ T (Fig. 3A). The diffusion constant for lateral Brownian motion of lipid molecules in lipid bilayers is $D_L = 2 \times 10^{-15}$ m$^2$ s$^{-1}$ (35), giving a fluctuation of...
frequency of $f_2^2 \sim 125$ Hz and $\Theta_0 \sim 10^{-4}$ (Fig. 3C). At this frequency, any quasi-static field effects will be predominantly suppressed by the spin-echo refocusing. The leading-order (gradient-channel) dephasing rate is given by (19) $\Gamma_{1L,2} = \frac{1}{2\sqrt{\Delta^2 + \frac{g_0^2}{\gamma_0^2}}} f_1^2 + \Theta_{1L} f_1^2$, giving rise to dephasing rates of the order $\Gamma_{1L} \sim 100$ Hz, with corresponding dephasing envelope $\mathcal{D}_1(t) = e^{-\Gamma_{1L} t}$.

The electric fields associated with the dissolved ions also interact with the NV center via the ground-state Stark effect. The coefficient for the frequency shift as a function of the electric field applied along the dominant ($z$) axis is given by $R_{3D} = 3.5 \times 10^{-3}$ Hz in V m$^{-1}$ (36). Fluctuations in the electric field may be related to an effective magnetic field via $B_{eff} = B_{3D}/\gamma P_{\rho}$, which may be used in an analysis similar to that above. An analysis using Debye–Hückel theory (37) shows charge fluctuations of an ionic solution in a spherical region $\Lambda$ of radius $R$ behave as $\langle Q_i^2 \rangle = D_E k_B T (1 + x R) e^{-\gamma R} [R \cosh(\gamma R) - \sinh(\gamma R)]$, where $D_E$ is the diffusion coefficient of the electrolyte, and $\gamma$ is the inverse Debye length ($\lambda_D$): $\lambda_D = 1/\kappa = 1.3$ nm for biological conditions. Although this analysis applies to a region $\Lambda$ embedded in an infinite bulk electrolyte system, simulation results discussed below show very good agreement when applied to the system considered here. The electric field variance may be obtained from $\langle Q_i^2 \rangle$, giving $\sigma_{E} = \sqrt{\langle E^2 \rangle} - \langle E \rangle^2 \sim 10^{-1}$ V m$^{-1}$, as a function of $\rho_{\rho}$. Relaxation times for electric field fluctuations are $\tau_{rel} = \rho_{\rho} e_{rms}$ (38), where $\rho_{\rho}$ is the resistivity of the electrolyte, giving $f_2^2 \sim 1/\tau_{rel} = 1.4$ GHz under biological conditions. Given the relatively low strength (Fig. 3A) and short relaxation time of the effective Stark-induced magnetic field fluctuations ($\Theta \sim 10^{5}$) (Fig. 3B), we expect the charge fluctuations associated with ions in solution to have little effect on the evolution of the probe.

**Discussion**

We now turn to the problem of noninvasively resolving the location of a sodium ion channel in a lipid bilayer membrane. When the channel is closed, the dephasing is the result of the background activity and is defined by $\mathcal{D}_{off} = D_{H2O} \mathcal{D}_{E} \mathcal{D}_{1L,2}$. When the channel is open, the dephasing envelope is defined by $\mathcal{D}_{on} = D_{off} \mathcal{D}_{ic}$. Maximum contrast will be achieved by optimizing the spin-echo interrogation time, $\tau$, to ensure $\mathcal{D}_{off} = -\mathcal{D}_{on}$ is maximal. Thus, in the vicinity of an open channel at the point of optimal contrast, $\tau \approx T_2/2$, we expect an ensemble ground-state population of $P_{\rho}(\frac{T}{2}) = \frac{1}{2} \{1 + \mathcal{D}_{on}(\frac{T}{2})\} = 0.61$, and $P_{\rho}(\frac{T}{2}) = \frac{1}{2} \{1 + \mathcal{D}_{off}(\frac{T}{2})\} = 0.93$ otherwise. By scanning over an open ion channel and monitoring the probe via repeated measurements of the spin state, we may build up a population ensemble for each lateral point in the sample. The signal-to-noise ratio improves with the dwell time at each point. Fig. 4 shows simulated scans of a sodium channel with corresponding image acquisition times of 4, 40, and 400 s. It should be noted here that the spatial resolution available with this technique is beyond that achievable by magnetic field measurements alone, because for large $\Theta$, $\Delta P \propto B^2 \propto h_{\rho}^2$.

We may employ similar techniques to temporally resolve a sodium ion-channel switch-on event. By monitoring a single point, we may build up a measurement record, $J$. In an experimental situation, measurement frequency has an upper limit of $f_m = (\tau_m + \tau_{2p})^{-1}$, where $\tau_m \approx 900$ ns is the time required for photon collection, and $\tau_{2p}$ is the time required for all three microwave pulses. A tradeoff exists between the increased dephasing due to longer interrogation times and the corresponding reduction in measurement frequency. Interrogation times are limited by the intrinsic $T_2$ time of the crystal. A second tradeoff exists between the variance of a given set of $N_i$ consecutive measurements and the temporal resolution of the probe. For the monitoring of a switching event, the spin state may be inferred with increased confidence by performing a running average over a larger number of data points, $N_i$. However, increasing $N_i$ will lead to a longer time lag before a definitive result is obtained. The uncertainty in the ion-channel state goes as $\Delta P \sim (\sqrt{N_i})^{-1}$, where $N_i$ is the number of points included in the dynamic averaging. We must take sufficient $N_i$ to ensure that $\Delta P < \Delta P_r (\tau_{2p}, T_2) = P_{off} - P_{on}$. The temporal resolution depends on the width of the dynamic average and is given by $\tau \approx N_i (\tau_m + \tau_{2p})$, giving the relationship $\tau = f_m (\tau_m + \tau_{2p})^{5/2}$. We wish to minimize this function with respect to $\tau$ for a given standoff ($h_{\rho}$) and $T_2$ time.

In reality, not all crystals are manufactured with equal $T_2$ times. An important question is therefore, for a given $T_2$, what is the best temporal resolution we may hope to achieve? Fig. 5B shows the optimal temporal resolution as a function of $T_2$. It can be seen that $\delta \tau$ improves monotonically with $T_2$, until $T_2$ exceeds the dephasing time due the fluctuating background fields (Fig. 5A). Beyond this point, no advantage is found from extending $T_2$.

A plot of $\delta \tau$ as a function of $\tau$ is given in Fig. 5C for standoffs of 2–6 nm. Solid lines depict the resolution that maybe achieved with $T_2 = 300 \mu s$. Dashed lines represent the resolution that may be achieved by extending $T_2$ beyond the dephasing times of background fields. We see that $\delta \tau$ diverges as $\tau \to T_2$ and is optimal for $\tau \to 1/T_2$.

As an example of monitoring of ion-channel behavior, we consider a crystal with a $T_2$ time of 300 $\mu s$ at a standoff of 3 nm. Fig. 5C tells us that an optimal temporal resolution of

![Fig. 5. Temporal characteristics of the measurement protocol. (A) Dephasing rates due to the sources of magnetic field plotted as a function of probe standoff, $h_{\rho}$. (B) Optimum temporal resolution as a function of crystal $T_2$ times for $h_{\rho} = 2$–6 nm. (C) Temporal resolution as a function of interrogation time, $\tau$, for separations of 2–7 nm. The limits corresponding to $T_2 = 15 \mu s$ and $T_2 = 300 \mu s$ are shown as vertical dashed lines.](https://www.pnas.org/cgi/doi/10.1073/pnas.1002562107)
\[ \delta t \approx 1.1 \text{ ms} \] may be achieved by choosing \( \tau \approx 100 \mu \text{s} \). This interrogation time suggests an optimal running average will employ \( N_s = \delta t (r + \tau_m) \approx 11 \) data points. Fig. 6A shows a simulated detection of a sodium ion-channel switch-on event using \( N_s = 20, 50, \) and 100 points. The effect of increasing \( N_s \) is shown to give poorer temporal resolution but also produces a lower variance in the signal, which may be necessary if there is little contrast between \( P_{\text{off}} \) and \( P_{\text{on}} \). Conversely, decreasing \( N_s \) results in an improvement to the temporal resolution but leads to a larger signal variation.

We now consider an ion-channel switching between states after an average waiting time of 5 ms (200 Hz) (Fig. 6B). To ensure the condition \( \delta \Phi < \Delta \Phi \) is satisfied, we perform the analysis using \( N_s = 20 \), giving a resolution of \( \delta t \approx 2 \text{ ms} \). The blue curve shows the response of the NV population to changes in the ion-channel state. Fourier transforms of the measurement record, \( \mathcal{F}(\mathcal{J}) \), are shown in Fig. 6C–E). The switching dynamics are clearly resolvable for heights less than 6 nm. The dominant spectral frequency is 100 Hz which is half the 200 Hz switching rate as expected. Beyond 6 nm, the contrast between \( P_{\text{off}} \) and \( P_{\text{on}} \) is too small to be resolvable due to the \( T_2 \) limited temporal resolution, as given in Fig. 5B. This contrast may be improved via the manufacturing of nanocrystals with improved \( T_2 \) times, allowing for longer interrogation times (dashed curves, Fig. 5C).

With regard to scaleup to a wide-field imaging capability, beyond the obvious extrinsic scaling of the number of single channel detection elements (in conjunction with microconfocal arrays), we consider an intrinsic scaleup strategy using many NV centers in a bulk diamond probe (39), with photons collected in a pixel arrangement. Because the activity of adjacent ion channels is correlated by the micrometer scale activity of the membrane, the fluorescence of adjacent NV centers will likewise be correlated, thus wide-field detection will occur via a fluorescence contrast across the pixel. Implementation of this scheme involves a random distribution of NV centers in a bulk diamond crystal. The highest reported NV density is \( 2.8 \times 10^{24} \text{ m}^{-3} \) (40), giving typical NV–NV couplings of \(<10 \text{ MHz} \) which are strong enough to introduce significant additional decoherence. We seek a compromise between increased population contrast and increased decoherence rates due to higher NV densities, \( n_{\text{nv}} \), given by \( \Gamma_{\text{nv}} \sim \frac{\delta t}{r_{\text{eff}}^2 r_{\text{ff}}} \) (19).

For ion-channel operation correlated across each pixel, the total population contrast \( \Delta \Phi \) between off and on states is obtained by averaging the local NV state population change \( \Delta \Phi (\tau) = P_{\text{off}}(r_{\text{ff}}, r_{\text{eff}}, \tau) - P_{\text{on}}(r_{\text{ff}}, r_{\text{eff}}, \tau) \) over all NV positions \( r_{\text{ff}} \) and orientations, and ion-channel positions \( r_{\text{eff}} \) and species, and maximizing with respect to \( \tau \). As an example, consider a crystal with \( n_{\text{nv}} = 10^{24} \text{ m}^{-3} \) whose surface is brought within 3 nm of the cell membrane containing a sodium and potassium ion-channel densities of \( \sim 2 \times 10^{15} \text{ m}^{-2} \) (41). Higher densities will yield better results, however, these have not been realized experimentally as yet, and electron spins in residual nitrogen will begin to induce NV spin flips. We expect ion-channel activity to be correlated across pixel areas of \( 1 \times 1 \mu \text{m} \), so the population contrast between on and off states is \( \Delta \Phi \approx 15 \). At these densities, the optimal interrogation time is \( \tau \sim 0.8 \mu \text{s} \), yielding an improvement in the temporal resolution by a factor of 10,000, opening up the potential for single-shot measurements of ion-channel activity across each pixel.

We have carried out an extensive analysis of the quantum dynamics of a NV diamond probe in the cell-membrane environment and determined the theoretical sensitivity for the detection, monitoring, and imaging of single ion-channel function through quantum decoherence. Using current demonstrated technology, a temporal resolution in the 1–10 ms range is possible, with spatial resolution at the nanometer level. With the scope for scaleup and unique scanning modes, this fundamentally different detection mode has the potential to revolutionize the characterization of ion-channel action, and possibly other membrane proteins, with important implications for molecular biology and drug discovery.

Materials and Methods

Geometry. All numerical calculations were performed with Monte Carlo methods using MATLAB, and conducted on a rectangular slab. Reflective boundary conditions are imposed at the top and bottom, representing the diamond crystal and lipid bilayer, respectively. Periodic boundary conditions are imposed on the sides of the volume. The lower boundary also hosts ion channels with the surface density given in the main text. The NV center is placed on a variable height above the diamond crystal boundary.

Ion Channel. We assume the waiting time between ion ejections from the channel to follow a Poissonian distribution about a mean rate of 3 MHz. Each ion will couple to the NV spin via Eq. 1, with \( t_w(\tau) \) describing the spatial separation between the NV spin and the dipoles in the ion channel. Upon exiting the channel, the timescales associated with the ion motion will change to that described by the self-diffusion rate of nuclear spins in the electrolyte. Because the addition of ions to the electrolyte at these frequencies is slow compared to the characteristic Brownian motion of the electrolyte, any ions exiting the channel will rapidly diffuse away. Thus for simplicity, ions are terminated upon exiting the channel, making the nuclear dipole concentration near the channel opening equal to that of the bulk.

Magnetic Field of the Electrolyte. For the purpose of modeling the fluctuating field due to self-diffusive behavior of the electrolyte, we do not distinguish between different species of nuclear dipole, and instead consider a weighted-average nuclear dipole moment as defined by the concentrations of water molecules and dissolved ions. The coupling of each ion to the NV center is again described by Eq. 1, however, couplings between ions, or to the

\[ \delta \Phi \sim \frac{\Delta \Phi}{\delta t} \]
mean field, are ignored because the associated timescales are much slower than those of their diffusive motion. Because of this weak coupling, each ion is assumed to not flip during its interaction with the NV center. Each of the \( r_s(t) \) then describes a random walk of a dipole of random but fixed orientation, as defined by the self-diffusion coefficient of the electrolyte.

**Magnetic Field of the Lipid Bilayer.** Lipid diffusive motion is modeled in a similar fashion to the bulk nuclear motion using the diffusion rate given in the main text, however, the lipids are confined to the two-dimensional plane comprising the lower boundary of the system geometry.


**Electric Field of the Electrolyte.** Regions of nonneutral charge density, together with their correlation lengths are found using Monte Carlo numerical techniques outlined in ref. 37. The rms electric field felt by the NV center may then be found by integrating over the charge distribution in the electrolyte.

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