reproduced by Mod 1, (see Fig. 4). This behavior is maintained with the capture of the carrier by the environment as described by $D_{\rm l}[\rho]$. Effects of the carrier wavefunction dephasing caused by thermal noise and described by $D_{\rm 2}[\rho]$, are small for tunneling $(N \leq 3)$, but introduce significant increase of F by factor 2 in the hopping transfer regime at $N > 3$.

As seen from Fig.5, the double -strand Mod 2 reproduces only a general trend of the N-dependence of the measured hole transfer velocity $F(N)$. The matter is that when solving the Lindblad equation for Mod 2 only one particular set of the model parameter values was used by us. We are planning to test other parameter sets available from the literature and check their better relevance for reproducing data of Giese experiment

References

Each base pair is considered as a structureless site numbered as $n = 0, 1...$, N, N+1, in a single -stranded electron model with the Hamiltonian :

Dissipative dynamics of a hole transfer in single- and doublestrand electronic model of DNA

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IV. Results

I. Introduction

V. Conclusion

We conclude that a drastic change in the hole transfer mechanism from tunneling (fast decrease of F at $N \leq 3$), to hopping (nearly constant F at N > 3) is accurately

Num erical solutions of the Lindblad equation in the qubit representation are found using the open source LindbladMPO code, which simulates elements of a quantum algorithm on a classical com puter [3]. The calculated results are compared to available experim ental data [2].

$$
\mathbf{u} = \mathbf{v} \cdot \mathbf{v}
$$

II. Simplified single-strand electronic model (Mod1) of DNA oligonucleotide.

 $\left| \bullet \right|$ g=0.01

Fig. 5. Obtained from Mod 2 (marked in yellow) the dimen sionless hole transfer velocity $F(N)$ as a function of the length of the bridge $(A-T)_{N}$ is presented without dephasing, i.e. $g=0$ in $D_2[\rho]$. The curve marked in blue is the same as in Fig. 4

of a hole transitions between neighboring sites.

The energy level diagram with the model param eters used is shown in Figure 3.

Two physical sources, and the corresponding operators $\left\langle D_{\text{l}}[\rho]\right\rangle$ and $D_2[\rho]$ in the Lindblad equation, give rise to the carrier dissipative dynamics: (1) capture of the carrier by the environment, resulting in the escape of the hole from the DNA, (2) the loss of the carrier wavefunction coherence under the influence of stochastic fluctuations of the

> Fig. 4. Obtained from Mod1 the dim en sion less hole tran sfer velocity *F*(*N*) as a function of the length of the bridge $(A-T)_N$ is presented without dephasing, i.e. $g=0$ in $D_2[\rho]$ (marked in green) and in cluding dephasing, $q=0.3$ meV (m arked in blue).

Our goal is to check whether both structurally distinct DNA models, discussed earlier in [1] and denoted here as Mod1 and Mod2, enable to reproduce accurately two different regimes of a hole motion of a hole carrier, namely, coherent tunneling in short DNA oligonucleotides and incoherent hopping in longer ones, in accordance with results of experimental measurements of the rate of charge transfer along a DNA oligonucleotide as a function of its length N.

> Fig. 1. Fragment of DNA with two pairs of nitrogenous bases: aden ine thym ine (A-T) and guanine - cytosine (G-C). The overlapping π -orbitals of the carbon and nitrogen atom s of the nitrogenous bases form the "paths" for charge movement along DNA.

a) From Fermi to Pauli operators: $1 \quad \bigg\{ \quad n-1$ 0 $j=0$, *n*-1 \bigcap 1 \bigcap *n* $z \sim^+$ z^+ $\sqrt{1-z^2}$ *n* $\left(\prod_{j} O_j | O_n \right)$, $\left(\prod_{j} O_j | O_n \right)$ *j* =0 \bigcup \bigcup *a a* ^σ ^σ ^σ ^σ -1 $\qquad \qquad$ $\qquad \qquad$ \qquad \qquad + α^+ $\sqrt{|\mathbf{u}|^2}$ α^2 $= 0$) $j =$ $\left(\begin{array}{cc} n-1 \\ \overline{1\end{array}\right)$ + $\left(\begin{array}{cc} n-1 \\ \overline{1\end{array}\right)$ $\rightarrow \left(\prod_{j=0} \sigma_j^z \right) \sigma_n^+, \quad a_n^+ \rightarrow \left(\prod_{j=0} \sigma_j^z \right)$

The Jordan-Wigner transformation defines the Ham iltonian in the qubit representation : $\frac{1}{2}\left(\sigma_{0}^{z}+\sigma_{N+1}^{z}\right) +\sum_{n=0}V_{n,n+1}\left(\sigma_{n}^{+}\sigma_{n+1}^{-}+\sigma_{n+1}^{+}\sigma_{n}^{-}\right)$ *N* $Z \leftarrow Z$ $H = \frac{\Delta}{2} (\sigma_0^z + \sigma_{N+1}^z) + \sum V_{n,n+1} (\sigma_n^+ \sigma_{n+1}^- + \sigma_{n+1}^+ \sigma_n^-)$ Δ $=\frac{\Delta}{2}\Big(\sigma_{0}^{z}+\sigma_{N+1}^{z}\Big)+\sum V_{n,n+1}\Big(\sigma_{n}^{+}\sigma_{n+1}^{-}+$

1. Cun iberti G., Maciá E., Rodríguez A., Röm er R.A. Tight-Bin din g Modeling of Ch arge Migration in DNA Devices. In : Ch akraborty, T. (eds) Ch arge Migration in DNA. Springer, Ch .1. P. 1-20 . 2007. 2. Giese B., Am audrut J., Köhler A.K., Sporm ann M., Wessely S. Direct observation of hole transfer through DNA by hopping between adenine bases and by tunneling. Nature. 2001. V. 412. P. 318-320 . 3. Lan da H. and Misguich G. Nonlocal correlations in noisy m ultiqubit system s sim ulated using m atrix product operators. SciPost Phys. Core. 2023. V. 6. P. 037-38.

d) Num erical solutions of the Lindblad equation are obtained using the Lindblad MPO Solver program [3].

Formulation of the problem and theoretical approach

The transfer of a hole carrier in a fragment of artificial DNA (oligonucleotide) is investigated by considering the "hole carrier in DNA + dynamic environment" as an open quantum system. DNA structure is approximated either by (Mod1) a simplified single-strand model or (Mod2) the more realistic double-strand one. Time evolution of the system density operator $\rho(t)$ is described by the Lindblad equation that is solved numerically.

> In the fram ew ork of a more realistic double-stranded electronic model Mod2 (not specified here for brevity) the hole transfer velocity $F(N)$ have been also obtained withing the same approach based on the Lin dblad equation and compared to experim ental data [2] (see Fig.5).

environment.

Giese et al. [2] laid the foundations for measuring a charge transfer in oligonucleotides - homogeneous and regular DNA sequences. Here, a quantum models, Mod1 and Mod2, for describing results obtained in these experiments are explore.

Fig. 2. Schem atic view on DNA oligonucleotide in the Giese experim ent [2, 3]. Donor and acceptor centers, each form ed by (G-C) base pairs, are highlighted and join ed by the $(A-T)_N$ bridge along which a hole carrier transfer.

$$
H = \sum_{n=0}^{N+1} \varepsilon_n a_n^+ a_n + \sum_{n=0}^{N} V_{n,n+1} \left(a_n^+ a_{n+1} + a_{n+1}^+ a_n \right)
$$

of a set of operators of creation (annihilation) of a set

where $a_n^+(a_n)$ are the operators of creation (annihilation) of a spinless fermion, \mathcal{E}_n is the energy level of the hole carrier at site *n*, and $V_{n,n+1}$ are the amplitudes

Fig. 3. Energy level diagram . \ket{D} and \ket{A} – states of a hole carrier on the donor $(n=0)$ and acceptor $(n = N+1)$, $\varphi_n(x) = 1 \div N$ – state of the charge carrier on the n-th (A-T)-pair of the bridge.

III. Method of calculating the charge transfer rate.

n

=

 $N+1$ \int \sum \int $n,n+1$ \int σ $n+1$ \int σ $n+1$ σ $n+1$

c) Selection of Mod1 param eters:

b) The charge transfer evolution is described by the Lindblad equation:
\n
$$
\frac{\partial \rho}{\partial t} = -\frac{i}{\hbar} [H, \rho] + D_1 [\rho] + D_2 [\rho],
$$
\n
$$
D_1 [\rho] = \sum_{n=0,1,N+1} \gamma_n \left(\sigma_n^+ \rho \sigma_n^- - \frac{1}{2} \{ \sigma_n^+ \sigma_n^- , \rho \} \right), \qquad D_2 [\rho] = g \sum_{n=1}^N \left(\sigma_n^z \rho \sigma_n^z - \rho \right)
$$

$$
\Delta = \varepsilon_{D/A} = -150 \text{meV}; \qquad V_{01} = V_{N,N+1} = V = 90 \text{meV};
$$

\n
$$
V_{n,n+1} = V_B = 30 \text{meV}; \qquad n = 1,...N-1
$$

\n
$$
\gamma_0 = 0.1 \text{meV}, \qquad \gamma_1 = 10 \text{meV}, \qquad \gamma_{N+1} = 40 \text{meV}, \qquad g = 0.3 \text{meV}
$$

