Numerically exact mimicking of quantum gas microscopy for interacting lattice fermions

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A numerical method is presented for reproducing fermionic quantum gas microscope experiments in equilibrium. By employing nested componentwise direct sampling of fermion pseudo-density matrices, as they arise naturally in determinantal quantum Monte Carlo (QMC) simulations, a stream of pseudo-snapshots of occupation numbers on large systems can be produced. There is a sign problem even when the conventional determinantal QMC algorithm can be made sign-problem free, and every pseudo-snapshot comes with a sign and a reweighting factor. Nonetheless, this “sampling sign problem” turns out to be weak and manageable in a large, relevant parameter regime. The method allows to compute distribution functions of arbitrary quantities defined in occupation number space and, from a practical point of view, facilitates the computation of complicated conditional correlation functions. While the projective measurements in quantum gas microscope experiments achieve direct sampling of occupation number states from the density matrix, the presented numerical method requires a Markov chain as an intermediate step and thus achieves only indirect sampling, but the full distribution of pseudo-snapshots after (signed) reweighting is identical to the distribution of snapshots from projective measurements.

I. INTRODUCTION

The Hubbard model is a highly simplified, yet paradigmatic model of materials with strong correlations which has found an accurate physical realization in cold atomic gases in optical lattices.$^8$ Its phase diagram is still poorly understood, which has led to an intense synergy of numerical approaches.$^{10,11}$

Remarkably, fermionic quantum gas microscopes$^{4–9}$ [see also Refs.$^{10,11}$ and references therein] with single-site and single-atom resolution give access to the full distribution function of occupation number states. This has allowed the direct measurement of two-point correlation functions$^{12,13}$ and of more unconventional quantities such as the full counting statistics (FCS) of macroscopic operators$^{14}$ or the non-local string order parameter characterizing spin-charge separation in 1D. Conditional correlation functions around dopants$^{15,16}$ and the analysis of patterns in the snapshots$^{17}$ have revealed polarons in the doped Hubbard model, in and out of equilibrium.$^{18}$ Furthermore, time-dependent measurements give access to transport properties.$^{19–22}$ In this context, comparison with numerical simulations is not only important for calibrating e.g. the temperature in cold atoms experiments, but quite generally for reliable benchmarking to prepare quantum simulators for parameter regimes where classical simulations are impossible.

Yet, for fermions in $d \geq 2$ dimensions, a numerically exact technique for mimicking such projective measurements of occupation number snapshots is still missing. A single hole in a system of infinitely strongly repulsive fermions ($t-J$ model) can be simulated with a worldline loop algorithm$^{20,21}$ without a sign problem and, more recently, worm algorithm Monte Carlo applied to the $t-J$ model has given unbiased results for spin configurations around a small number of dopant.$^{22–24}$ However, by definition, the $t-J$ model neglects doublon-hole fluctuations, and for the Fermi-Hubbard model at finite interaction path integral Monte Carlo simulations$^{25,26}$ are only possible in one dimension due to the fermionic sign problem which is extensive in the system size. We extend the determinantal QMC (DQMC) algorithm$^{33–35}$ by an inner loop, where Fock configurations are sampled directly, i.e. without autocorrelation time, from a fully tractable quasiprobability distribution. A common technique for obtaining a tractable joint probability distribution, which can be sampled directly, is to model it as a product of conditional distributions.$^{36}$ This idea is at the heart of autoregressive neural networks$^{37}$, the generation of natural images pixel by pixel$^{38}$ and recent algorithms for simulation and generative modelling of quantum systems.$^{39–41}$

Alternative DQMC approaches, summing all Fock states implicitly, exist for computing the FCS of quadratic operators$^{42}$ and all elements of the reduced density matrix on small probe areas.$^{43}$ The nested componentwise direct sampling technique presented here is more versatile in that pseudo-snapshots can be produced for probe areas as large as in current experiments with the proviso that a (mild) sign problem is manageable in experimentally relevant regimes. The resulting distribution of (a sufficiently large number of) pseudo-snapshots after reweighting is - within controllable statistical error - identical to the distribution of snapshots from projective measurements as generated in quantum gas microscope experiments.

We calculate (i) the joint FCS of the staggered spin and pseudo-spin magnetization of the Hubbard model at half filling, (ii) the distribution of the total number of holes and doubly occupied sites as a function of doping at high temperature, and (iii) the magnetization environment of...
a polaron, where we find qualitative agreement with a recent quantum Monte Carlo simulation for a single hole in the $t-J$ model\textsuperscript{23}. An apparent discrepancy between Refs.\textsuperscript{22,23} and the quantum gas microscope experiment of Ref.\textsuperscript{22}, which we can also reproduce qualitatively, can be pinpointed to a difference in doping regimes.

II. NESTED AND COMPONENTWISE DIRECT SAMPLING

We are considering the single-band Hubbard model

$$H = -t \sum_{\langle i,j \rangle, \sigma = \uparrow, \downarrow} (c_i^\dagger \sigma c_j \sigma + h.c.) + U \sum_i \hat{n}_{i,\uparrow} \hat{n}_{i,\downarrow} - \mu \sum_{i, \sigma} \hat{n}_{i, \sigma}$$

with the usual notation, and treat it within the DQMC framework\textsuperscript{33,35}. After a Trotter-Suzuki decomposition of the density operator $\hat{\rho} \sim \exp(-\beta H)$ at inverse temperature $\beta = 1/T$ into $N_r = \beta/\Delta \tau$ imaginary times slices and a Hubbard-Stratonovich (HS) transformation for decoupling the interactions by introducing a functional integral over HS fields $\{s\}$, the density operator reads\textsuperscript{19}

$$\hat{\rho} = \frac{1}{Z} \prod_{\{s\} \sigma = \uparrow, \downarrow} \left( w_{\sigma}^{\uparrow} \frac{e^{-\sum_i X_i^2 \sigma^\dagger(\{s\}) X_i^2 \sigma}}{w_{\sigma}^{\uparrow}} \right) \left( w_{\sigma}^{\downarrow} \frac{e^{-\sum_i X_i^2 \sigma^\dagger(\{s\}) X_i^2 \sigma}}{w_{\sigma}^{\downarrow}} \right)$$

$$\equiv \frac{1}{Z} \prod_{\{s\} \sigma = \uparrow, \downarrow} w_{\sigma}^{\uparrow} w_{\sigma}^{\downarrow} \hat{\rho}_{\{s\}}^{\sigma}.$$  (2a)  (2b)

Here, formally, we have $\exp(-X^2\sigma) \equiv \prod_{k=1}^{N_s} B_k^\sigma$, where $B_k^\sigma = e^{-\Delta \tau V_k^\sigma(\{s\})}$ is the matrix representation of the single-particle propagator for spin $\sigma$ of the potential and kinetic part after HS transformation\textsuperscript{33}. Note that we use the conventional HS transformation\textsuperscript{33} in which the discrete auxiliary fields couple to the $S^z$-component of the electron spin, $S_i^z = \hat{n}_{i,\uparrow} - \hat{n}_{i,\downarrow}$, which, as will be discussed below, is crucial. After integrating out the fermionic degrees of freedom, $w_{\sigma}^{\uparrow} = \det \left(1 + e^{-X^2 \sigma(\{s\})}\right) = Z(\{s\})$ is the contribution of the spin component $\sigma$ to the Monte Carlo weight of the HS field configuration $\{s\}$, which can also be interpreted as the partition sum of the non-interacting fermion system $\hat{\rho}_{\{s\}}^\sigma$. Since the kinetic and potential matrices in the matrix product leading to Eq. (2) do not commute, the resulting matrix $e^{-X^2 \sigma(\{s\})}$ is not Hermitian and (except in 1D) not all diagonal matrix elements of $\hat{\rho}_{\{s\}}$, are semi-positive-definite; hence, $\hat{\rho}_{\{s\}}$ is termed a pseudo-density matrix, while the total $\hat{\rho}$ in Eq. (2) is a true density matrix.

The structure of the density matrix in Eq. (2) suggests a nested sampling approach, in which the HS fields of the pseudo-density matrices $\hat{\rho}_{\{s\}}$ are sampled using the Markov chain of the conventional determinantal QMC algorithm, while the occupation numbers can be sampled directly, i.e. without autocorrelation time, from each free-fermion pseudo-density matrix $\hat{\rho}_{\{s\}}^\sigma$ given that for fixed $\{s\}$ their distribution function and all its marginals can be calculated efficiently.

From the chain rule of basic probability theory every probability distribution can be decomposed into a chain of conditional probabilities

$$p_{\{s\}}(n_1, n_2, \ldots, n_D) = \prod_{k=1}^{D} p_{\{s\}}(n_k | n_{k-1}, n_{k-2}, \ldots, n_1)$$

(3)

where some ordering of the random variables $n_1, n_2, \ldots, n_D$ is implied. A sample from the joint distribution is then generated by traversing the chain as

$$p_{\{s\}}(n_1) \xrightarrow{n_1 \sim p_{\{s\}}(n_1)} p_{\{s\}}(n_2 | n_1) \xrightarrow{n_2 \sim p_{\{s\}}(n_2)} \ldots$$

where $n_k \sim p_{\{s\}}(n_k)$ denotes sampling variable $n_k$ from $p_{\{s\}}(n_k | n_{k-1}, n_{k-2}, \ldots)$ and the sampled value is "inserted" into the next conditional probability along the chain [see Fig. 1(a)].

Below we discuss how to calculate the conditional quasiprobabilities in Eq. (3) for a free fermion pseudo-density matrix $\hat{\rho}_{\{s\}}^\sigma$. Note that for fixed HS field configuration, the pseudo-density matrices for spin up and
down are statistically independent. Per HS sample, spin up and down occupancies are sampled independently and then combined into full pseudo-snapshots, see Fig. 1(b). Henceforth, we drop the subscripts \{s\} and \(\sigma\) for notational convenience.

### A. Direct sampling in the grand canonical ensemble

In the atomic microscopy, the quantity of central interest is the quasiprobability to find a given snapshot of the fermion occupation:

\[
p(n_1, n_2, \ldots, n_D) = \text{Tr}(\rho \Pi_{n_1} \Pi_{n_2} \cdots \Pi_{n_D}),
\]

\[
p(n_1, n_2, \ldots, n_k) = (-)^{n_1+n_2+\cdots+n_k} \det \left( G_{ij} \right),
\]

where \(G_{ij} \equiv G_{ij}^\sigma(\{s\}) = \langle c_i, \sigma c_j^\dagger | \{s\} \rangle \) is the equal-time single-particle Green’s function of spin species \(\sigma\) for a given HS field configuration \(\{s\}\) at a randomly chosen imaginary time slice. Eq. (5) is proven in appendix A.

We now decompose the high-dimensional quasiprobability distribution Eq. (4) into a chain of conditional quasiprobability distributions, which by definition can be computed as

\[
p(n_{k+1} | n_1, n_2 \cdots n_k) = \frac{p(n_1, n_2 \cdots n_k, n_{k+1})}{p(n_1, n_2, \cdots n_k)}.
\]

Inserting Eq. (5) and using the determinant formula for block matrices we find:

\[
p(0 | n_1, n_2 \cdots n_k) = G_{k+1,k+1} - \Delta G_{k+1},
\]

\[
p(1 | n_1, n_2 \cdots n_k) = 1 - G_{k+1,k+1} + \Delta G_{k+1},
\]

with the “correction term”

\[
\Delta G_{k+1} = \sum_{i=1}^k G_{k+1,i}(G_{K,K} - N_{K,K})^{-1} G_{i,k+1},
\]

where \(K = \{1, 2, \ldots, k\}\) denotes the ordered set of site indices, \(G_{k,K}\) is the corresponding submatrix of the
Green’s function, and $N_{K,K} = \text{diag}(n_1, n_2, ..., n_k)$ is a diagonal matrix whose entries are the sampled occupation numbers on the sites $K$.

If the correction term $\Delta G_{k+1}$ were zero, then the conditional probability $p(n_{k+1}|n_1, n_2, ...) = 0$ would be simply given by the diagonal element $G_{k+1,k+1}$ of the Green’s function and be independent of the other occupation numbers. Therefore the correction term is crucial for inter-site correlations. While traversing the chain of conditional probabilities, the block structure of the matrix whose inverse is required in Eq. (8) can be exploited recursively such that no calculation of a determinant or matrix inverse from scratch is necessary (see appendix B).

B. Reweighting

As said earlier, the pseudo-density operator $\hat{\rho}(s)$ is not Hermitian and not all conditional quasiprobabilities in Eq. (7) are non-negative. Therefore, we rewrite them as

$$p(n_k) = \text{sign}(p(n_k)) \frac{|p(n_k)|}{N_k} \times N_k,$$

with the shorthand notation $p(n_k) \equiv p(n_k|n_1, n_2, ..., n_{k-1})$. The sampling for component $k$ is then carried out using the valid probability distribution $q(n_k) \equiv \frac{|p(n_k)|}{N_k}$ with normalization $N_k = |p(n_k = 0)| + |p(n_k = 1)|$ (see Fig. 1(a)).

Having sampled the entire chain of conditional probabilities for both spin components, the generated snapshot is associated with a (signed) reweighting factor $R = R^a R^d$ where

$$R^a = \text{sign}(w^a(s)) \prod_{k=1}^{D} \left( \text{sign}(p(s) (n^a_k)) N^a_k \right),$$

and all quantities $O(n) = O(n_1, n_2, ..., n_D)$ that are evaluated on $M$ generated snapshots $n_i^{(1)}$ need to be reweighted as

$$\langle O \rangle = \frac{\sum_{i=1}^{M} R_i O(n_i^1)}{\sum_{i=1}^{M} R_i}.$$

The joint pseudo-probability $p(n_1, n_2, ...) = 0$ can be factored in arbitrary order into components. However, the reweighted distribution $q(n_1, n_2, ...)$ and thus the magnitude and sign of the reweighting factor depends on the chosen order leaving room for optimization in a given HS sample.

The severity of the sign problem is basis dependent: We find that it strongly depends on the single-particle basis for sampling and the chosen HS transformation. Sampling in the $S_z^-$ or $S_z^+$-basis of the electron spin (or in the $S_z$-basis in momentum space) leads to a very strong sign (phase) problem. Choosing the HS transformation that couples to the electron charge density rather than the electron spin also gives a very severe sign problem.

Even at half filling, where the DQMC algorithm can be made sign-problem free for the purpose of computing expectation values, there is a sampling sign problem, with a non-uniform dependence on $U/t$. The average sign diminishes as $U/t$ increases from $U/t = 0$ and reaches a minimum at an intermediate value $(U/t)_{\text{MIC}} \approx 4 - 7$, where a metal-to-insulator crossover occurs. For $U/t > (U/t)_{\text{MIC}}$ the sign problem again gradually becomes much less severe. The dependence of the sampling sign problem on temperature, interaction strength, and doping is presented in Figs. C.1, C.2, and C.3 in Appendix C. The average sampling sign of pseudo-snapshots for the parameters in all figures of the main text is $(\bar{s}) \geq 0.75$ with $(\bar{s})$ defined in Eq. (C2).

In all simulations, we use a Trotter discretization of $\Delta t = 0.02$ and generate around 20 pseudo-snapshots per HS sample on equidistant imaginary time slices. The simulation code has been verified by comparing with exact diagonalization results (see appendix D).

III. APPLICATIONS

A. Joint full counting statistics (FCS)

At half filling, the Hubbard model has an enlarged $(SU(2) \times SU(2))/Z_2 = SO(4)$ symmetry, which is the combination of spin-rotational and particle-hole symmetry and is generated by two commuting sets of angular momentum operators describing the total spin and total pseudo-spin of the system, respectively. As the temperature is lowered, domains form with an order parameter of the same symmetry, which, apart from fluctuating in length, can rotate on an $SO(4)$ sphere between antiferromagnetic, $s$-wave pairing and charge-density wave correlations, as one goes from one domain to a neighbouring domain. A joint histogram of the operators representing different components of the order parameter should reflect that they are projections of the same vector along different directions in order parameter space. Fig. 2 shows the joint distribution of the projections of the staggered magnetization $M_{\text{stag}} = \sum_{i=1}^{N_x} (-1)^i (\hat{n}_{i,\uparrow} - \hat{n}_{i,\downarrow})$ and the staggered pseudo-spin $Q_{\text{stag}} = \sum_{i=1}^{N_x} (-1)^i (\hat{n}_{i,\uparrow} + \hat{n}_{i,\downarrow} - 1)$ on a square probe area of size $N_A = L^2_A$, as obtained from re-weighted pseudo-snapshots. As $U/t$ increases [Fig. 2 (a-c)], the suppression of charge fluctuations manifests itself in the narrowing of the pseudo-spin distribution. The even-odd effect visible in the joint distributions is due to the fact that an even number of sites can only accommodate an even magnetization of spin-$\frac{1}{2}$ (pseudo-spin) moments.

Note that $P(M_{\text{stag}}^{\uparrow}, Q_{\text{stag}}^{\uparrow})$ could also be obtained using the generating function approach of Ref. [22]. This is not true for the FCS of the number of doublons $N_d = \sum_{i=1}^{N_x} \hat{n}_{i,\uparrow} \hat{n}_{i,\downarrow}$ and the total number of holes $N_h = \sum_{i=1}^{N_x} (1 - \hat{n}_{i,\uparrow}) (1 - \hat{n}_{i,\downarrow})$ since these operators are non-
quadratic in fermionic operators. We find that e.g. for $U/t = 12$ and $\beta = 4$, the FCS of $N_h$ and $N_d$ as a function of doping are accurately modelled by (shifted) binomial distributions (see appendix E).

B. Three-point spin-charge correlator

A three-point spin-charge correlator in the reference frame of the hole\cite{fig3} was calculated for a single hole in the $t-J$ model at zero temperature with DMRG and trial wave functions\cite{Blomquist2019} and at finite (high and low) temperature using worm algorithm QMFT\cite{Hosur2018}. Presumably the same correlator was measured experimentally in Ref.\cite{Blomquist2019} for the Hubbard model at large interactions. For the purpose of meaningful comparison between the Hubbard and $t-J$ model we calculate the slightly modified correlator:

$$C_{id}(r) = \frac{\langle \mathcal{P}_r^h S_r^z S_{r+\frac{\delta}{2}}^z S_{r+\frac{\delta}{2}}^2 \rangle}{\langle \mathcal{P}_0^h \rangle}$$

where $S_r^z = \hat{n}_{r+\delta/2} - \hat{n}_{r-\delta/2}$, and the projector

$$\mathcal{P}_r^h = \hat{n}_{r_0}^h \hat{n}_{r_0+\delta/2}^\dagger \hat{n}_{r_0-\delta/2}^\dagger \hat{n}_{r_0+\delta/2} - \hat{n}_{r_0}^\dagger$$

with $\hat{n}_{r}^h = (1 - \hat{n}_{r+\delta/2}) (1 - \hat{n}_{r-\delta/2})$ and $\hat{n}_{r}^\dagger = (1 - \hat{n}_{r+\delta/2}) \hat{n}_{r-\delta/2} + (1 - \hat{n}_{r-\delta/2}) \hat{n}_{r+\delta/2}$ selects configurations with an empty site at position $r_0$ surrounded from eight sides by spin-only states, which serves to exclude nearest- and next-nearest neighbour double-hole pairs from the statistics. The conditional correlation functions in the reference frame of the hole, Eq.\cite{fig3}, are implemented straightforwardly by applying a filter to the pseudo-snapshots, whereas implementing such higher order correlation functions using a generalized form of Wick’s theorem would require separate coding for each specific correlator thus hampering quick experimentation (although this would give better statistics since all Fock state are summed implicitly).

Fig. 3 shows overall qualitative agreement of $C_{id}(r)$ for our data for the Hubbard model and that of Ref.\cite{Blomquist2019} for the $t-J$ model, with some notable differences in $C_{s^2}$ and $C_2$ in the immediate vicinity of the hole. A careful comparison of the $t-J$ model with the Hubbard model would require a renormalization of all correlators in the former by a polynomial in $t/U$\cite{Blomquist2019} (although certain qualitative model differences may not be captured perturbatively)\cite{Blomquist2019}.

There are qualitative differences to the experimental data of Ref.\cite{Blomquist2019}, which were already noted in Ref.\cite{Blomquist2019}. The data of Ref.\cite{Blomquist2019} can be reproduced by nested component-wise sampling, pointing, however, to a very different conclusion: namely, that magnetic polarons may have disappeared for the relatively high doping level of Ref.\cite{Blomquist2019}. This is illustrated in appendix E.

IV. CONCLUSION

In conclusion, the presented method for generating pseudo-snapshots allows both theorist to take part in the exploration of fermionic quantum microscopy and experimentalists to use numerical simulations in a more versatile way\cite{Blomquist2019}. The data analysis of pseudo-snapshots is quite analogous to that of experimental snapshots generated by projective measurements except that a signed reweighting factor needs to be taken into account. While it is not meaningful to compare individual pseudo-snapshots with actual experimental snapshots, the full distribution of pseudo-snapshots after reweighting is identical to the distribution of snapshots from projective measurements. The difference is that quantum gas microscope experiments achieve direct sampling of occupation number states from the density matrix, whereas the method presented here relies on indirect sampling from the overall interacting fermion density matrix (we can achieve direct sampling only at the level of the constituent free fermion density matrices). Arbitrary quantities can be evaluated...
on the reweighted pseudo-snapshots, including those that cannot be feasibly expressed as expectation values of operators. A case in point is the FCS of macroscopic operators which are higher than quadratic in fermionic operators; there, the generating function method of Ref. does not apply, and evaluation as a sum of projectors onto Fock states would require a number of terms which is exponential in the number of sites.

Our nested componentwise direct sampling method is generic to all fermionic Monte Carlo methods that are based on the “free fermion decomposition” and is easily adapted to projector DQMC for accessing zero-temperature properties where Slater determinants rather than thermal free-fermion pseudo-density matrices are sampled directly in the inner loop of the Markov chain. For Hubbard models around intermediate interaction strength $U/t \approx (U/t)_{\text{MIC}}$ further improvements are required to reduce the sampling sign problem. There, the interest lies in potentially observing non-Gaussian fluctuations or characterizing attraction and spin-correlations between dopants in the crossover from a polaronic metal to a Fermi liquid. While a general solution of the sampling sign problem is unlikely, it remains to be investigated how more general HS descriptions and representations of the electron operator which were successful at eliminating the sign problem of the Monte Carlo weights, affect the sampling sign problem and whether snapshots in another single-particle basis can be generated.

The total computing time spent on the generation of Figs. 1-3 amounts to the equivalent of approximately $3 \times 10^4$ CPU hours on an Intel(R) Core(TM) i5-6300U CPU with 2.40GHz clock cycle.

V. ACKNOWLEDGMENTS

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Appendix A: Inductive proof of Eq. (5) in the main text

Let

\[ G_{i_a,J_b}^{(0)} = \langle \hat{c}_{i_a} \hat{c}_{j_b}^\dagger \rangle_0 \]  

(A1)

be the single-particle Green’s function of a free fermion system, and \( \alpha, \beta \in \{1, \ldots, D\} \). First we prove the well-known fact that Wick’s theorem for higher-order correlation functions can be expressed in the compact determinant form

\[ \langle \left( \hat{c}_{i_1} \hat{c}_{j_1}^\dagger \right) \cdots \left( \hat{c}_{i_n} \hat{c}_{j_n}^\dagger \right) \rangle_0 = \det \left( G_{i_i,j_j}^{(0)} \right)_{i,j = \{i_1, i_2, \ldots, i_n\} \land \{j_1, j_2, \ldots, j_n\}} \equiv \left[ G_{i_j}^{(0)} \right]_{I,J} \]  

(A2)

We use the convention that \([A]_{I,J}\) refers to the determinant of the submatrix of \(A\) whose row index (column index) runs in the set \(I (J)\), which is the “inclusive” definition of the minor.

The proof goes by induction. The case \(n = 1\) is true by virtue of the definition (A1). In the induction step, we use Wick’s theorem writing all non-vanishing contractions for a product of \(n + 1\) pairs of fermionic operators as

\[ \langle \left( \hat{c}_{i_1} \hat{c}_{j_1}^\dagger \right) \cdots \left( \hat{c}_{i_n+1} \hat{c}_{j_{n+1}}^\dagger \right) \rangle_0 = -G_{i_1,j_{n+1}}^{(0)} \langle \left( \hat{c}_{i_{n+1}} \hat{c}_{j_1}^\dagger \right) \left( \hat{c}_{i_2} \hat{c}_{j_2}^\dagger \right) \cdots \left( \hat{c}_{i_n} \hat{c}_{j_n}^\dagger \right) \rangle_0 \]

\[ - \sum_{k=2}^{n-1} G_{i_k,j_{n+1}}^{(0)} \langle \left( \hat{c}_{i_1} \hat{c}_{j_1}^\dagger \right) \cdots \left( \hat{c}_{i_{k-1}} \hat{c}_{j_{k-1}}^\dagger \right) \left( \hat{c}_{i_k} \hat{c}_{j_k}^\dagger \right) \cdots \left( \hat{c}_{i_n} \hat{c}_{j_n}^\dagger \right) \rangle_0 \]

\[ - G_{i_{n+1},j_1}^{(0)} \langle \left( \hat{c}_{i_1} \hat{c}_{j_1}^\dagger \right) \cdots \left( \hat{c}_{i_{n-1}} \hat{c}_{j_{n-1}}^\dagger \right) \left( \hat{c}_{i_n} \hat{c}_{j_n}^\dagger \right) \rangle_0 \]

\[ + G_{i_{n+1},j_{n+1}}^{(0)} \langle \left( \hat{c}_{i_1} \hat{c}_{j_1}^\dagger \right) \cdots \left( \hat{c}_{i_n} \hat{c}_{j_n}^\dagger \right) \rangle_0 \].

(A3)

The minus-sign, e.g. in the second line of Eq. (A3), comes from the permutation of \(\hat{c}_{i_{n+1}}\) with \(\hat{c}_{j_{n+1}}\). Using the induction hypothesis Eq. (A2), valid for \(n\) pairs of fermionic operators, the remaining correlators in Eq. (A3) can be expressed as determinants

\[ \langle \left( \hat{c}_{i_1} \hat{c}_{j_1}^\dagger \right) \cdots \left( \hat{c}_{i_{n+1}} \hat{c}_{j_{n+1}}^\dagger \right) \rangle_0 = -G_{i_1,j_{n+1}}^{(0)} \cdot \det \left( G_{i_{n+1},j_1}^{(0)} G_{i_{n+1},j_2}^{(0)} \cdots G_{i_{n+1},j_{n-1}}^{(0)} \right) 
\]

\[ - \sum_{k=2}^{n-1} G_{i_k,j_{n+1}}^{(0)} \cdot \det \left( G_{i_{n+1},j_k}^{(0)} G_{i_{n+1},j_{k+1}}^{(0)} \cdots G_{i_{n+1},j_{n-1}}^{(0)} \right) 
\]

\[ - G_{i_{n+1},j_1}^{(0)} \cdot \det \left( G_{i_1,j_1}^{(0)} G_{i_1,j_2}^{(0)} \cdots G_{i_1,j_{n-1}}^{(0)} \right) 
\]

\[ + G_{i_{n+1},j_{n+1}}^{(0)} \cdot \det \left( G_{i_1,j_1}^{(0)} G_{i_1,j_2}^{(0)} \cdots G_{i_1,j_{n-1}}^{(0)} \right) \].

(A4)

By rearranging rows in (A4), it can be concluded that (A4) is the expansion of the \((n + 1) \times (n + 1)\) determinant

\[ \det \left( G_{i_a,j_b}^{(0)} \right)_{a,b = \{i_1, i_2, \ldots, i_{n+1}\} \land \{j_1, j_2, \ldots, j_{n+1}\}} \]  

(A5)
accompanying the single-particle Green’s function \(G_{d}^{(0)}\) in Eq. (A4) we need to perform \((n - k)\) row exchanges which results in a factor \((-1)^{n-k}\) such that the total sign is \((-1)^{n-k+1}\), which is identical to the alternating factor \((-1)^{n+1+k}\) coming from Laplace’s formula.

Eq. (5) differs from Eq. (A2) in that instead of pairings \((\hat{c}_{i} \hat{c}_{j}^\dagger)\) there are projectors of the form

\[
\Pi_k = n_{k} \hat{c}_{jk} \hat{c}_{jk} + (1 - n_{k}) \hat{c}_{jk} \hat{c}_{jk}. \tag{A6}
\]

Depending on whether \(n_{k} = 0\) or \(n_{k} = 1\), only one of either terms in (A6) applies. Let us replace for the moment only the \(k\)-th pairing \((\hat{c}_{i} \hat{c}_{j}^\dagger)\) in Eq. (A2) by \(\Pi_k\). If \(n_{k} = 0\), the resulting expression is covered by Eq. (A2). The case \(n_{k} = 1\) is different since \(\hat{c}_{jk}\) is to the right of \(\hat{c}_{jk}\). Using \(\hat{c}_{jk} \hat{c}_{jk} = 1 - \hat{c}_{jk} \hat{c}_{jk}\), one obtains

\[
\langle (\hat{c}_{i} \hat{c}_{j}^\dagger) \cdots \Pi_k \cdots (\hat{c}_{i} \hat{c}_{j}^\dagger)\rangle_0 = (-1)^{n_{k}} n_{k} \left[ \det (G_{I\setminus jk, J\setminus jk}^{(0)}) - \det (G_{I, J}^{(0)}) \right]. \tag{A7}
\]

To make the connection with Eq. (5) we consider the matrix

\[
\left( G^{(0)} \right)_{ij} = \left( G^{(0)} \right)_{ij} - n_{k} \delta_{i,jk} \delta_{j,jk} \tag{A8}
\]

and develop its determinant with respect to the \(k\)-th column:

\[
\det (\bar{G}) = \sum_{l=1}^{n} (-1)^{l+k} \left( G^{(0)} \right)_{i_l,j_k} \left[ G^{(0)} \right]_{I\setminus i_l, J\setminus j_k} \tag{A9}
\]

\[
= \sum_{l=1}^{n} (-1)^{l+k} \left( G^{(0)} \right)_{i_l,j_k} \left[ G^{(0)} \right]_{I\setminus i_l, J\setminus j_k} - n_{k} (-1)^{2k} \left[ G^{(0)} \right]_{I\setminus j_k, J\setminus j_k}. \tag{A10}
\]

Here, angular braces \((\cdot)\) denote matrix elements and square brackets \([\cdot]\) denote the inclusive definition of the minor. Note that the minors \([G^{(0)}]_{I\setminus j_k, J\setminus j_k} = [G^{(0)}]_{I\setminus i_l, J\setminus j_k}\) since \(G^{(0)}\) and \(\bar{G}^{(0)}\) only differ in the element \((j_k, j_k)\) which is excluded from the minors. Now, one can recognize in the first sum Eq. (A10) the Laplace expansion of \(G^{(0)}\) with respect to the \(k\)-th column and undo it again to recover \(\det (\bar{G}^{(0)})\):

\[
\det (\bar{G}^{(0)}) = \det \left( G^{(0)} \right) - n_{k} \left[ G^{(0)} \right]_{I\setminus j_k, J\setminus j_k}. \tag{A11}
\]

Eq. (A11) is identical to Eq. (A7), which proves that

\[
\langle (\hat{c}_{i} \hat{c}_{j}^\dagger) \cdots \Pi_k \cdots (\hat{c}_{i} \hat{c}_{j}^\dagger)\rangle_0 = \det \begin{pmatrix}
G^{(0)}_{i_1,j_1} & G^{(0)}_{i_1,j_2} & \cdots & G^{(0)}_{i_1,j_n} \\
G^{(0)}_{i_2,j_1} & G^{(0)}_{i_2,j_2} & \cdots & G^{(0)}_{i_2,j_n} \\
\vdots & \vdots & \ddots & \vdots \\
G^{(0)}_{i_n,j_1} & G^{(0)}_{i_n,j_2} & \cdots & G^{(0)}_{i_n,j_n}
\end{pmatrix} - n_{k} \left[ G^{(0)} \right]_{I\setminus j_k, J\setminus j_k}. \tag{A12}
\]

Repeatedly replacing each pairing \((\hat{c}_{i} \hat{c}_{j}^\dagger)\) in Eq. (A2) by a projector \(\Pi_l\) of the form (A6) and repeating the derivation from (A8) to (A12), with a Laplace expansion carried out with respect to the \(l\)-th column, completes the proof of Eq. (5).

**Appendix B: Exploitation of block matrix structure**

Eq. (5) implies that the expressions for the joint quasiprobability distributions of successive numbers of components are related by a block matrix structure. Using the formula for the determinant of a block matrix and noticing that
Given that $X_{k-1}$ is itself a block matrix

$$X_{k-1} = \begin{pmatrix} X_{k-2} & G(x_1, x_{k-1}) \\ \cdot & \cdot & \cdot \\ G(x_{k-2}, x_1) G(x_{k-2}, x_2) \cdots G(x_{k-2}, x_{k-2}) & G(x_{k-2}, x_{k-1}) & \cdots & G(x_{k-2}, x_{k-1}) \\ \cdot & \cdot & \cdot \\ G(x_{k-1}, x_1) G(x_{k-1}, x_2) \cdots G(x_{k-1}, x_{k-2}) & G(x_{k-1}, x_{k-1}) & \cdots & G(x_{k-1}, x_{k-1}) \end{pmatrix}$$

with $G(x_{k-1}, x_{k-1})$ just a number and assuming that the inverse $X_{k-2}^{-1}$ is already known, one can make use of the formula for the inversion of a block matrix to compute the inverse of $X_{k-1}$ in an economical way. We define

$$g \equiv G(x_{k-1}, x_{k-1}) - \sum_{i,j=1}^{k-2} G(x_{k-1}, x_i) [X_{k-2}^{-1}]_{i,j} G(x_j, x_{k-1})$$

and recognize that

$$g = \frac{p(x_1, x_2, \ldots, x_{k-1})}{p(x_1, x_2, \ldots, x_{k-2})} = p(x_{k-1}|x_{k-2}, \ldots, x_1)$$

which means that we have computed $g$ already previously when sampling the $(k-1)$-th component.

Using the formula for the inverse of a block matrix

$$X_{k-1}^{-1} = \begin{pmatrix} X_{k-2}^{-1} + g^{-1} \bar{u} \otimes \bar{v}^T & -g^{-1} \bar{u} \\ -g^{-1} \bar{v}^T & g^{-1} \end{pmatrix}$$

where

$$[\bar{u}]_i = \sum_{j=1}^{k-2} [X_{k-2}^{-1}]_{i,j} G(x_j, x_{k-1})$$

and

$$[\bar{v}^T]_j = \sum_{i=1}^{k-2} [X_{k-2}^{-1}]_{i,j} G(x_{k-1}, x_i)$$

and

$$[\bar{u} \otimes \bar{v}^T]_{ij} = [\bar{u}]_i [\bar{v}^T]_j.$$
Appendix C: Sign problem for nested componentwise direct sampling at and away from half filling

The severity of the conventional sign problem in the determinantal QMC algorithm is measured by the average sign of the Monte Carlo weight

$$\langle s \rangle = \frac{1}{N_{\text{HS samples}}} \sum_{\{s\}} \text{sign}(w^\uparrow_{\{s\}} w^\downarrow_{\{s\}}),$$

where the sum is over auxiliary field configurations of the Hubbard-Stratonovich samples. To quantify the “sampling sign problem” in the nested componentwise sampling algorithm we introduce the average sign of a snapshot, \(\langle \tilde{s} \rangle\), which is given by

$$\langle \tilde{s} \rangle = \frac{1}{M} \sum_{\{s\}} \text{sign}(w^\uparrow_{\{s\}} w^\downarrow_{\{s\}}) \sum_{i=1}^{N_{\text{snapshots per HS}}} \prod_{\sigma=\uparrow,\downarrow} \prod_{k=1}^{D} \text{sign}(p_{\{s\}}^{(i)}(n^\sigma_k)),$$

where \(M = N_{\text{HS samples}} \times N_{\text{snapshots per HS}}\) is the total number of generated snapshots. Here, it is understood that snapshots drawn from the same HS sample, the number of which is \(N_{\text{snapshots per HS}}\), are multiplied by the sign of the corresponding Monte Carlo weight \(w^\uparrow_{\{s\}} w^\downarrow_{\{s\}}\); in case that there is already a conventional sign problem at the level of the determinantal QMC algorithm. Note that within one HS sample, \(\{s\}\), snapshots for spin-\(\uparrow\) and spin-\(\downarrow\) can be paired up arbitrarily into a full snapshot due to the statistical independence of the two spin species.

Furthermore, one can define the average (unsigned) reweighting factor

$$R_{\text{av.}} = \frac{1}{M} \sum_{\{s\}} \sum_{i=1}^{N_{\text{snapshots per HS}}} \prod_{\sigma=\uparrow,\downarrow} \prod_{k=1}^{D} N_{\sigma,(i)}^{\sigma,(s)}$$

FIG. C.1. Sampling sign problem at half filling. The probe area is the total \(L \times L\) square system \((L = L_A)\). (a) Average sign of a pseudo-snapshot. (b) Average (unsigned) reweighting factor (full lines) and average maximum reweighting factor (dashed lines). The red dashed-dotted line indicates a rough estimate of the threshold of maximum reweighting factors below which numerically exact results can be obtained with modest computational effort.
FIG. C.2. Conventional sign problem and sampling sign problem away from half filling: dependence on $U/t$. (a,c) Average sign ($\langle s \rangle$, full lines) and average sampling sign ($\langle \tilde{s} \rangle$, dashed lines); (b,d) average reweighting factor (full lines) and average maximum reweighting factor (dotted lines). The inverse temperature $\beta$ is in units of $1/t$. The probe area is equal to the system size with $L_A = L = 8$.

FIG. C.3. Conventional sign problem and sampling sign problem away from half filling: dependence on inverse temperature $\beta t$. System size $L_A = L = 8$. 
and the average maximum reweighting factor, averaged over independent Markov chains,

\[ R_{\text{av. max}} = \frac{1}{N_{\text{Markov chains}}} \sum_{m=1}^{N_{\text{Markov chains}}} \max_{\text{snapshots } i} (|R_i|). \]  

(C4)

The average maximum reweighting \( R_{\text{av. max}} \) is the most relevant indicator of the severity of the sign problem as it quantifies the "inflation" of value of an individual snapshot. Because the reweighting factor can fluctuate over several orders of magnitude, the average sign alone is not sufficient for a characterization of the "sampling sign problem". How the sampling sign and the reweighting factor depend on doping, inverse temperature \( \beta t \) and interaction strength \( U/t \) is shown in Figs. C.1, C.2, and C.3.

The snapshots generated in quantum gas microscope experiments originate from independent experimental runs whereas the pseudo-snapshots of the nested componentwise sampling are affected by the autocorrelation time inherent in the sampling of Hubbard-Stratonovich field configurations via the standard determinantal QMC algorithm. For large \( U/t \), pseudo-snapshots taken from the same HS sample (and at the same imaginary time slice), differ mostly in the positions of holes and doubly occupied sites, while the spin background stays largely fixed. Additionally, pseudo-snapshots come with a sign and reweighting factor due to the non-Hermiticity of the pseudo-density operator within each HS sample. Therefore, experimental snapshots and pseudo-snapshots are not directly comparable. An important question is how many pseudo-snapshots \( M \) are required to obtain a comparable precision of measurement quantities as from \( M^{(\text{exp})} \) experimental snapshots. Taking into account the average maximum reweighting factor, the effective number of pseudo-snapshots \( M' \) that is equivalent to \( M^{(\text{exp})} \) can be roughly estimated as

\[ M' = M/(\tau_{\text{DQMC}} \times R_{\text{av. max.}}) \leftrightarrow M^{(\text{exp})}. \]  

(C5)

Here, \( \tau_{\text{DQMC}} \) is some measure of the autocorrelation of the Markov chain generated by the standard determinantal QMC algorithm.

The "sampling sign" deteriorates exponentially with the probe area \( N_A = L_A^2 \). Yet, an extent of the probe area \( L_A = \xi(T/t, U/t, L) \), where \( \xi \) is the correlation length, is often sufficient for meaningful simulations.

Appendix D: Code verification

For benchmarking purposes an irregular model instance of the Hubbard model on five sites (see inset in Fig. D.1(b)) is chosen

\[ H = - \sum_{\langle i,j \rangle, \sigma = \uparrow, \downarrow} t_{ij} \sigma \left( c_{i,\uparrow}^\dagger c_{j,\sigma} + h.c. \right) + U \sum_{i=1}^{5} n_{i,\uparrow} n_{i,\downarrow} - \sum_{\sigma = \uparrow, \downarrow} \mu_{\sigma} \sum_{i=1}^{5} n_{i,\sigma}, \]  

(D1)

which breaks translational, point group and spin rotational symmetry. The hopping matrix is identical for both spin species and reads

\[
[t^\sigma]_{i,j} = t \begin{pmatrix}
0 & 0.7 & 1.1 & 0 & 0.8 \\
0.7 & 0 & 1.05 & 0.9 & 1.2 \\
1.1 & 1.05 & 0 & 1.0 & 0 \\
0 & 0.9 & 1.0 & 0 & 0 \\
0.8 & 1.2 & 0 & 0 & 0
\end{pmatrix},
\]  

(D2)

The onsite repulsion is \( U/t = 4 \) and the chemical potential for spin up and down is \( \mu_{\uparrow}/t = 1.5 \) and \( \mu_{\downarrow}/t = 1.8 \), respectively; the inverse temperature is \( \beta t = 4 \) discretized into \( N_r = 256 \) Trotter time slices with \( \Delta t = \beta t/N_r = 1/64 \). Fig. D.1 compares the probabilities \( P(s) \) of all microstates \( s \) in the occupation number basis with results from exact diagonalization. An enlarged view of Fig. D.1(a) is shown in Fig. D.1(b). The occupation number state \( \vec{n} = [n_{1,\uparrow}, \ldots n_{5,\uparrow}; n_{1,\downarrow} \ldots n_{5,\downarrow}] \) with \( n_{i,\sigma} = n_{\sigma} \in \{0,1\} \) is interpreted as a bitstring and is represented by the integer \( s = \sum_{n=0}^{N_{\text{dim}}-1} n_{\sigma} 2^n \). With the knowledge of the eigenenergies \( E_i \) and eigenvectors \( |\phi_i\rangle \) of the Hamiltonian \( H \) the probability of an occupation number state \( s \) is

\[ P(s) = \sum_{i=1}^{\text{dim}(H)} e^{-\beta E_i} \frac{|\langle s|\phi_i\rangle|^2}{Z}. \]  

(D3)
with $Z = \sum_{i=1}^{\text{dim}(H)} e^{-\beta E_i}$. In order to demonstrate that the nested componentwise direct sampling method yields consistent microscopic states also for much larger system sizes, Fig. D.2 shows the probabilities of spin-only states on the eight sites around an isolated, mobile hole for a system of $10 \times 10$ lattice sites at small doping. Only isolated holes, i.e., those surrounded exclusively by singly occupied sites, are considered (see Eq. (13) of the main text). With the labelling of sites around a hole as shown in the right panel of Fig. D.2, the spin environment is characterized by the state $\vec{\sigma} = [\sigma_0, \sigma_1, \ldots, \sigma_7]$ where $\sigma_p = 1$ if there is a spin-$\uparrow$ at position $p$ around the hole and $\sigma_p = 0$ if it is a spin-$\downarrow$.

With this convention, the state index in Fig. D.2 is given as $s = \sum_{p=0}^{7} \sigma_p 2^p$. Fig. D.2 shows that there is a hierarchy of groups of states (right panel of Fig. D.2). Furthermore, states related by symmetry, which are grouped together in coloured boxes in Fig. D.2, appear with approximately the same probability. This is not a built-in feature of the nested componentwise sampling algorithm and thus provides strong evidence that all relevant occupation number states are sampled with their correct probabilities.

**Appendix E: FCS of number of doublons and holes**

Fig. E.1 displays the doping dependence of the FCS of the number of doubly occupied sites $N_d = \sum_{i=1}^{N_A} n_{i,\uparrow} n_{i,\downarrow}$ and holes $N_h = \sum_{i=1}^{N_A} (1 - n_{i,\uparrow})(1 - n_{i,\downarrow})$ for $U/t = 12, \beta t = 4$ on an $L \times L$ system with $L = L_A = 8$. For small doping values, the FCS are well described by binomial distributions (dashed and dotted lines) which are generated by populating lattice sites independently with doublons (holes) with probability $p_d = \langle n_{i,\uparrow} n_{i,\downarrow} \rangle$ ($p_h = \langle (1 - n_{i,\uparrow})(1 - n_{i,\downarrow}) \rangle$).

According to the distribution $B_{n,p}(x) = \binom{n}{x} p^x (1-p)^{n-x}$ with $n = N_A$ and $p \in \{p_d, p_h\}$. In the strongly doped case (lower panel), the distribution of the number of holes can be modelled accurately by a “shifted” binomial distribution (dashed-dotted line), which is obtained by populating the lattice with holes arising from doublon-hole fluctuations according to a binomial distribution with parameter $p = p_d$ (since the number of doublon-hole pairs is approximately equal to the number of doublons) which is then shifted so that the distribution is centered about its mean. Of course, for a large probe area and away from a critical point, the distribution of holes (doublons) should approach a Gaussian distribution which could be characterized by computing the mean and variance directly.
FIG. D.2. (a) Probabilities of spin-only states around an empty site as obtained from reweighted pseudo-snapshots in the reference frame of the hole. Parameters: $U/t = 12$, $\mu/t = -2$, $\beta t = 2.5$, $L \times L$ square lattice with $L = 8$. For this system size, the average number of holes and doubles $\langle N_h \rangle \approx 1.50$ and $\langle N_d \rangle \approx 1.31$ so that the average number of excess holes is $\langle N_h \rangle - \langle N_d \rangle \approx 0.19$. With 56 independent Markov chains a total number of $70 \times 10^6$ occupation number samples of the entire system was obtained. In the reference frame of the hole, those states around the hole position are filtered out which do not contain charge fluctuations (“spin-only states”). Per Hubbard-Stratonovich configuration, 200 direct sampling steps were performed during the Monte Carlo sweeps at different imaginary time slices. Trotter discretization $\Delta \tau t = 0.02$. (b) Illustration of the most important spin-only states around an isolated hole which are marked in panel (a). Dashed lines delineate different antiferromagnetic domains.

FIG. E.1. FCS of the total number of doubly occupied sites $\langle N_d \rangle$ and holes $\langle N_h \rangle$, obtained from reweighted pseudo-snapshots for different levels of hole doping (top and bottom panel). Dashed and dotted lines are binomial distributions; the dashed-dotted line is a “shifted” binomial distribution. Parameters: $U/t = 12$, $\beta t = 4$, $L = 8$.

Appendix F: Additional data for the polaron problem

This section illustrates the significance of numerical simulations for exploring the parameter space. In agreement with Ref. 29 for the $t - J$ model and the results of Fig. 3 for the Hubbard model, one can identify the joint condition $C_{d=2}(r) = 0 < 0$ and $C_{d=2}(r) = 1 > 0$ as a characteristic correlation pattern of a magnetic polaron. However, $C_{d=2}(r) = 0 < 0$ by itself occurs also for larger doping values where the antiferromagnetic spin background is too diluted to host polarons. Fig. F.1 juxtaposes the evolution of the three-point correlation function in the reference
frame of a hole

\[ C_{|d|=2}(|r|) = \frac{\langle n_{c_0}^{(h)} S^z_{c_0 + r + \frac{d}{2}} S^z_{c_0 + r - \frac{d}{2}} \rangle}{\langle n_{c_0}^{(h)} \rangle}. \] (F1)

as a function of chemical potential \( \mu/t \) for \( L = 4 \) (c) and \( L = 6 \) (e), respectively, with the distribution of the number of holes (b,g) and doubly occupied sites (a,f). Unlike in Eq. (12) of the main text, for the study of the hole environment in Fig. [F.1] no projection operator restricting the sites around the hole to be singly occupied has been applied prior to evaluating the correlation function. The data points connected by a thick red line are taken from Fig. 4(c) of Ref.\(^{19}\). From \( \mu/t = -4.0 \) to \( \mu/t = -5.5 \) the correlator \( C_{|d|=2}(|r|) \) changes markedly which leads to the conclusion that Fig. 4 of Ref.\(^{19}\) is consistent with the disappearance rather than the presence of magnetic polarons, which is due to the relatively high level of doping chosen in Ref.\(^{19}\). This picture is supported by visually comparing the two randomly selected pseudo-snapshots for \( \mu/t = -4.0 \) versus \( \mu/t = -5.0 \) in Fig. [F.1](d). However, note that the pseudo-snapshots should not be taken at face value since they come with a sign and a reweighting factor. Fig. [F.2] shows the FCS of the number of doubly occupied sites and holes for the same parameters as in Fig. 5 of the main text, indicating that only a small number of excess holes is present in the pseudo-snapshots.

FIG. F.1. The three-point spin-charge correlation function \( C_{|d|=2}(|r|) \) of Eq. (F1) computed on a 4 \( \times \) 4 system (a-c) as well as a 6 \( \times \) 6 system (d-g) with periodic boundary conditions and with parameters \( t/t = 14, \beta t = 2.5, \mu = -4.0, -4.5, \ldots, -6.5 \). The dashed magenta line in (c) [denoted \( \text{ conditional } C_2 \)] shows the correlation function \( C_{|d|=2}(|r|) \) calculated for \( \mu/t = -4.0 \) according to Eq. (12) in the main text, in which only those snapshots are counted where the hole is surrounded on eight sites by spin-only states (i.e. it is not adjacent to a doublon or a hole and can be regarded as an isolated dopant). The experimental data of Ref.\(^{19}\) are also shown for comparison. There, the system size is approximately 4 \( \times \) 6 to 6 \( \times \) 6 lattice sites (slightly inhomogeneous with open boundary conditions), interactions \( U/t \approx 14 \) and stated temperature \( T = 1.4J \) which corresponds to \( T/t \approx 0.4 \) (i.e. \( \beta t = 2.5 \)) for \( J = 4t^2/U \). In the quantum gas microscope experiment, there are on average 1.95(1) dopants present in each experimental realization (doublon-doped instead of hole-doped)\(^{19}\). For our setting on a 4 \( \times \) 4 system at \( \mu/t = -4.0 \), on the other hand, the average number of holes (hole-doped) is \( \langle N_h \rangle = 0.72 \), which is small enough for measuring the spin-environment of an isolated dopant. (d) Randomly chosen pseudo-snapshots for \( \mu/t = -4 \) (left) and \( \mu/t = -5 \) (right).
FIG. F.2. FCS of the number of doubly occupied \( N_d \) and empty sites \( N_h \) for the same parameters as in Fig. 3 of the main text, which are \( L = 10, U/t = 14, \mu/t = -3.0, \beta t = 2.5 \).