



A Numerically Exact Approach to Quantum Impurity Problems in Realistic Lattice Geometries

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We review a controlled numerical approach to quantum impurity problems in realistic geometries, consisting of exactly mapping the complete lattice Hamiltonian onto an equivalent one dimensional system through a unitary transformation. The resulting dimensional and entanglement reduction allows one to study the quantum many-body problem on arbitrary *d*-dimensional lattices using the density matrix renormalization group (DMRG) method. The real-space resolution allows one to position the impurities at the boundary or bulk of the sample and to study screening effects due to edge or surface modes. We describe how to generalize this approach to multi-impurity problems, discuss applications and possible extensions.

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1. INTRODUCTION

Research in engineering and fabrication of magnetic nano-structures has been, and continues to be, a fast growing field [1–5]. Their usefulness can potentially be applied to data storage, spintronics, sensors, magnetic imaging, quantum computing, and more [6]. Moreover, magnetic adatoms have been regarded as the building blocks of quantum nanostructures that can be assembled on crystal surfaces through single-atom manipulation with a scanning tunneling microscope (STM). These can serve as quantum simulators for studying magnetism with excellent real-space and energy resolution [7–9]. In the same spirit as cold-atom setups, this technique can unveil magnetic order and excitations in different geometries [10–13]. The surface not only supports the spin centers, but also plays a crucial role in stabilizing magnetic order [14–17]. Fundamental to all of these considerations is how individual spins interact with each other. Exchange interactions between impurities may have different physical origins: they can either arise from direct exchange for nearest-neighbor adsorption sites or indirect substrate-mediated coupling that asymptotically decay as a power-law with increasing separation between the impurities (Ruderman-Kittel-Kasuya-Yosida or RKKY interaction) [18–20].

The physics of a single magnetic impurity is well understood in terms of the Kondo problem: the magnetic moment is either partially or totally screened by the spin of the conduction electrons forming a collective many-body state with the Fermi sea [21]. In essence, the impurity ends up acting as a scattering center. However, in reality it is an extended object with a complex internal structure, usually referred-to as the hybridization cloud or "Kondo cloud," which is centered at the impurity and decays with a characteristic range R_K [22–24].

In order to understand how multiple impurities can realize magnetic order, it is first necessary to determine the effective interactions mediated by the conduction electrons in the substrate. A good point to start is by considering two Kondo impurities interacting locally with free fermions in the bulk via an antiferromagnetic exchange coupling J_K [25]:

$$H = H_{\text{band}} + J_K \left(\vec{S}_1 \cdot \vec{s}_{r_1} + \vec{S}_2 \cdot \vec{s}_{r_2} \right). \tag{1}$$

where H_{band} is the lattice Hamiltonian for non-interacting electrons, parameterized by a hopping *t* and \vec{s}_{r_i} represents the conduction electron's spin at the impurity's coordinate r_i , for impurities i = 1, 2. The same form of the Hamiltonian can be generalized to multiple impurities.

Historically, the standard approach is to assume that the impurities see each-other via an effective RKKY interaction, or indirect-exchange. It was first introduced in 1954 by Rudderman and Kittel to explain the coupling of nuclear magnetic moments [19]. Kasuya realized the same method could be applied to localized d or f electrons in a metallic host [20] and further work was carried on by Yosida [18]. The problem can be thought of as one impurity interacting with the conduction electrons which then "carry" this interaction to the next impurity. The effective Hamiltonian in second order perturbation theory is written as:

$$H_{eff} = J_{RKKY}(\mathbf{R})\mathbf{S}_1 \cdot \mathbf{S}_2, \qquad (2)$$

where $J_{RKKY}(\mathbf{R}) = J_K^2 \chi(\mathbf{R})$, is a distance dependent function and $\chi(\mathbf{R})$ is the Fourier transform of the non-interacting static susceptibility, or Lindhard function, which varies with distance, dimensionality and fermion density. The corresponding expression which is often cited in the literature is derived from assuming a uniform electron gas with a quadratic dispersion $E(k) \sim k^2$ [26] and its asymptotic behavior at long distances ($k_F R \gg 1$) and in *d* dimensions is given by:

$$\chi(R) \sim \frac{\sin\left(2k_F R + \pi \, d/2\right)}{R^d}.$$

As multiple impurities are included, the problem is complicated by the fact that there are now competing energy scales. This perturbative approach ignores many important features such as the many-body physics that can lead to a Kondo ground state. Therefore, as we shall see, these arguments alone can not accurately describe multiple impurity physics at low temperatures.

As perturbative approaches fail for strongly interacting systems, numerical methods become essential for a better understanding of experimental results. In this review, we describe in detail a completely unbiased and controlled numerical approach to solve quantum impurity problems in *d*-dimensional lattices [27–29]. The method relies on an exact canonical transformation in which a Lanczos recursion allows one to recast the complete lattice Hamiltonian as a tri-diagonal matrix that in turn can be associated to an equivalent one dimensional tight-binding problem. This procedure can be recognized as a generalization of Wilson's numerical renormalization approach

[30] and Haydock's recursion method [31–34]. After the noninteracting part of the problem has been reduced to a chain, the many-body physics is introduced in the form of a Kondo or Anderson impurity. The resulting low-dimensional problem can efficiently be solved using the density matrix renormalization group (DMRG) [35–38] method. We generalize these concepts to the multi-impurity case and discuss improvements to optimize the calculations in certain particular cases.

2. MAPPING AND DIMENSIONAL REDUCTION

2.1. Single Impurity

2.1.1. Lanczos Transformation

Solving for the ground state in an interacting lattice problem in dimensions greater than one is quite a challenging problem. Some impurity solvers include quantum Monte Carlo (QMC) [39–43] and the numerical renormalization group (NRG) [30, 44]. However, these methods have limitations, such as capturing the finite details of a lattice, or running into the well-known sign-problem (modern diagramatic QMC techniques somewhat overcome these issues). Fortunately, onedimensional systems can be solved efficiently with the density matrix renormalization group (DMRG) technique [35–38]. This section is dedicated to transforming an *d*-dimensional noninteracting lattice Hamiltonian, onto an *equivalent*, solvable onedimensional chain [27].

A general Hamiltonian for impurity problems will have the form

$$H = H_l + H_{imp} + V_c. \tag{3}$$

Here, H_l is a single-particle tight-binding Hamiltonian of a lattice. H_{imp} and V_c describe the impurity and the coupling between impurity and lattice, respectively. We point out that this method is applicable regardless of the geometry or dimensionality of the lattice.

First, let one consider a single impurity problem and one orbital per site. More general cases of multiple orbitals and impurities will be discussed in the next section. We start by first considering the non-interacting band Hamiltonian without the impurity. In the presence of translational symmetry, this matrix can be readily diagonalized by transforming to a plane-wave basis. However, the impurity breaks this invariance but preserves other point group symmetries of the lattice, such as rotations and reflections. We would like to generate a single particle basis compatible with these. We get our inspiration from NRG: Wilson considered a free electron gas in the continuum and used a basis of partial waves centered at the position of the impurity. A lanczos transformation allowed him to map the problem onto an equivalent one dimensional chain that he solved by block decimation [30, 44] (NRG also introduces the idea of logarithmic discretization of energy scales and renormalization, that will not be used here). In our case, we carry out a similar procedure, but in the presence of the lattice. The intuition is as follows: consider a single electron state en real space representation corresponding to a particle at the lattice site connected to the impurity r_0 .

By applying the non-interacting term of the Hamiltonian H_l we obtain a new state that is a superposition of single-particle states at the neighboring sites; this is the first "Lanczos orbital." By continuing this procedure the wave-functions will expand radially outward, away from r_0 . However, the action of the hopping also makes these orbitals "contract" inward. In order to obtain an orthogonal basis, one needs to orthogonalize each new orbital with those obtained in prior iterations. This is precisely what the Lanczos recursion does for us. Formally, the procedure starts by choosing an appropriate single particle "seed" state as:

$$|\Psi_0\rangle = c_{r_0}^{\dagger} |0\rangle, \qquad (4)$$

where $c_{r_0}^{\dagger}$ creates an electron at site r_0 and $|0\rangle$ is the vacuum state. This site can be chosen according to two situations. First, for a Kondo impurity, it is the site the impurity will couple to. Alternatively, for a substitutional impurity, typically described with an Anderson-type model, the seed *is* the impurity site (see **Figure 1**). As it will be made clear, the mapping does not effect the "seed" state and, therefore, does not modify H_{imp} . In both cases, the mapping is performed in the same way.

Once the seed is chosen, the rest of the states are constructed with the following iterative procedure:

$$|\Psi_1\rangle = H_l |\Psi_0\rangle - a_0 |\Psi_0\rangle \tag{5}$$

$$|\Psi_n + 1\rangle = H_l |\Psi_n\rangle - a_n |\Psi_n\rangle - b_n^2 |\Psi_{n-1}\rangle \tag{6}$$

$$a_n = \frac{\langle \Psi_n | H_l | \Psi_n \rangle}{\langle \Psi_n | \Psi_n \rangle} \tag{7}$$

$$b_n^2 = \frac{\langle \Psi_n | \Psi_n \rangle}{\langle \Psi_{n-1} | \Psi_{n-1} \rangle}.$$
(8)

The equations for a_n and b_n are obtained by requiring the states to be orthogonal. Note, however, that at this stage the states are not normalized.

After this transformation, H_l is tri-diagonal and has the form:

$$H_{l} = \begin{pmatrix} a_{0} & b_{1} & 0 & 0 \\ b_{1} & a_{1} & b_{2} & 0 \\ 0 & b_{2} & a_{2} & b_{3} \\ 0 & 0 & b_{3} & \ddots \end{pmatrix} .$$
(9)

Equivalently, in second quantization it reads

$$H_l = \sum_{i=0}^{L} a_i \tilde{n}_i + \sum_{i=0}^{N-1} b_{i+1} (\tilde{c}_i^{\dagger} \tilde{c}_{i+1} + h.c.),$$
(10)

where \tilde{c}_i^{\dagger} , \tilde{c}_i are normalized creation and annihilation operators, respectively, $\tilde{n}_i = \tilde{c}_i^{\dagger} \tilde{c}_i$ is the particle number operator and *L* is the total length of the chain. The geometry of this new Hamiltonian is a chain as in **Figure 1**. The diagonal a_n terms are on-site potentials, while the b_n 's are the new hoppings along the chain.

While performing this recursion in practice, numerical errors are introduced for large *L* due to finite precision. There are two ways to reduce these errors: (1) Since $\langle \Psi_n | \Psi_n \rangle$ grows with *n*, normalize Ψ_{n-1} at each iteration. (2) To prevent loss



of orthogonalization, introduce re-orthogonalization. This can easily done with the following procedure:

$$|\Psi_n\rangle' = |\Psi_n\rangle - \sum_{j=0}^{n-1} \langle \Psi_j | \Psi_n \rangle | \Psi_j \rangle.$$
(11)

As an example, consider a 2D square lattice of linear dimension L. Instead of L^2 sites, one now has to keep only $\sim O(L)$ sites. This is indeed an exact canonical transformation. The remaining missing orbitals correspond to different symmetry sectors of the Hamiltonian and consequently are completely decoupled from the impurity. The impurity is only coupled to the "s-wave" channel. The seed site acts as center of symmetry for the group D_{4h} . The "s-wave" channel corresponds to the totally symmetric representation A_{1g} and is formed by the basis function $\{x^2 + y^2\}$. Different symmetry channels corresponding to different representations such as those shown in Figure 2 will form their own independent chains. An example of this is the B_{1g} representation formed by the $\{x^2 - y^2\}$ basis function which can be thought of as a "d-wave" channel. The key point of this mapping is that it leaves H_{imp} and V_c in Equation (3) unchanged after the transformation. The first site of the chain corresponds to a real lattice site in *real* space. Therefore, as mentioned, the impurity will only couple to this single site of the chain.

If the lattice is infinite, the recursion can be performed indefinitely. In practice, the iterations are stopped and the chain is cut after some sufficient number of iterations. This is equivalent to assuming the lattice has the same "shape and size" as the last orbital in the Lanczos iteration. Alternatively, one can consider a finite lattice of arbitrary shape. Once the boundary is reached by the orbitals, they will "bounce" or reflect back, retracing the



path to the seed site. At some point, we will find that $b_{n+1} = 0$, meaning that all orbitals within that symmetry sector have been exhausted and we have a chain of finite length. If the finite lattice has no symmetry elements, the chain length will be equal to the total number of lattice sites. For instance, for the case of a square lattice, Lanczos orbitals for a single impurity have a diamond-like shape (see **Figure 1**). If the lattice has a boundary with a different symmetry, the different symmetry sectors will mix.

The full many body problem is recovered by connecting the impurity back to the chain. The recursion has not affected the terms H_{imp} and V_c in Equation (3), so the impurity will now be connected to the first site of the chain (the seed orbital) by V_c without any changes. These Hamiltonian can now be solved with DMRG, or any method of choice.

2.1.2. Entanglement Reduction

As previously described, the Lanczos orbitals are defined by their symmetry properties, or channel, and their radial distance from the center of symmetry, which can be associated to the linear distance along the equivalent one-dimensional chain. These considerations provide for an intuitive understanding of the scaling of the entanglement. In the equivalent problem after the Lanczos transformation, the region enclosed by an area of "radius" L in the d-dimensional lattice maps onto $\sim L^{d-1}$ onedimensional chains, each corresponding to a different symmetry sector. The entanglement per channel between this region and the rest of the lattice is exactly the same as the entanglement between the first L sites of each chain and their complement. In cases where the system under consideration is gapless, the chain is a critical one-dimensional system (for square and cubic lattices, for instance) and the von Neumann entanglement entropy is proportional to log(L) [45, 46]. All channels contribute to the entropy with similar factors. This yields a final result proportional to $L^{d-1}\log(L)$. These simple arguments easily explain why free fermions in higher-dimensions have logarithmic corrections to the area law [47-50]. Remarkably, one only needs to solve the problem in the channel that is directly coupled to the impurity, reducing the entanglement by a factor of L^{d-1} !

2.1.3. Continued Fractions and Non-interacting Green's Functions

The Lanczos transformation described above can be used to obtain non-interacting Green's Functions (GF). As a matter of fact, it was originally introduced in the literature with this purpose and it is known as the continued fraction expansion, or method of the moments [31–34]. Consider the Green's Function,

$$G(\omega) = \sum_{n} \frac{|\Psi_n\rangle \langle \Psi_n|}{\omega - E_n}.$$
 (12)

Define a projection operator as $P_0 = |\Phi_0\rangle \langle \Phi_0|$, where $|\Phi_0\rangle$ is our seed state of the mapping. The projected GF is then defined as

$$\langle \Phi_0 | G(\omega) | \Phi_0 \rangle = G_0(\omega) = \sum_n \frac{|\langle \Psi_n | \Phi_0 \rangle|^2}{\omega - E_n}$$

= $\langle \phi_0 | \frac{1}{\omega - \hat{H}} | \phi_0 \rangle .$ (13)

Now, it is clear that the function $G_0(\omega)$ is just the 00 element of the matrix $(\omega \hat{I} - \hat{H})^{-1}$, where \hat{H} is the tri-diagonal Hamiltonian after the Lanczos transformation as in Equation (9). To compute this inverse, it is best to partition the matrix into blocks, then utilize properties of partitioned matrices. For example, an $m \times m$ matrix of the form:

$$\mathbf{M} = \begin{pmatrix} a & \mathbf{b}^T \\ \mathbf{b} & \mathbf{C} \end{pmatrix}, \tag{14}$$

where *a* is a scalar, **b** is an $(m - 1) \times 1$ vector and **C** is an $(m - 1) \times (m - 1)$ matrix. Then the 00 entry is given by,

$$\mathbf{M}_{00}^{-1} = \frac{1}{a - \mathbf{b}^T \mathbf{C}^{-1} \mathbf{b}}.$$
 (15)

In the current case, \mathbf{b}^T is just $(b_1, 0, 0, 0, ...)$, so the above equation reduces to, switching back to the a_n and b_n notation of Equation (9):

$$G_{00}^{-1} = \frac{1}{a_0 - b_1 G_{11}^{-1} b_1}.$$
(16)

The process is then repeated for \mathbf{G}_{11}^{-1} . This method can be used to find any matrix element of $G(\omega)$. After iterating, the result is a continued fraction:

$$G_{00}(z = \omega + i\delta) = \frac{1}{z - a_0 - \frac{b_1^2}{z - a_1 - \frac{b_2^2}{z - a_2 - \frac{b_n^2}{z - a_n}}}.$$
 (17)

Recall that $|\phi_0\rangle$ represents a site in real space of the lattice. The density of states (DOS) and local density of states (LDOS) for a system are defined, respectively as,

$$\rho(\omega) = \sum_{n} \langle \Psi_n | \Psi_n \rangle \,\delta(\omega - E_n) \tag{18}$$

$$\rho_0(\omega) = \sum_n |\langle \Psi_n | \phi_0 \rangle|^2 \delta(\omega - E_n) = -\frac{1}{\pi} \lim_{\eta \to 0} \operatorname{Im} G_0(\omega + i\eta).$$
(19)

The LDOS of any geometry can then be computed numerically. Notice the first site of the chain has the exact DOS of the lattice. One way to terminate the expansion is just to cut the chain at some sufficient chain length n, as in Equation (17). However, this results in poor resolution of the LDOS, even for quite large n. Instead, consider a continued fraction with all a_n and b_n constant, i.e., $a_n = a$ and $b_n = b$ for all n. The continued fraction, after an equivalence transform can be written as

$$Y(\eta) = \frac{\eta}{1 + \frac{\eta}{1 + \frac{\eta}{1 + \frac{\eta}{\dots}}}},$$
 (20)

where $\eta \equiv \frac{-b^2}{(z-a)^2}$ and the continued fraction is carried out indefinitely. This expression converges to a finite value for all complex *z* with finite imaginary part. This value is given by

$$Y(z) = \frac{1}{2}(1 + \sqrt{4\eta + 1}) - 1.$$
(21)

Using the square lattice again for an example, the hoppings b_n along the Lanczos chain converge to $b_{\infty} = 2$ as seen in **Figure 3A**. After a sufficient number of iterations, the values of a_n and b_n are approaching their asymptotic limits. This permits the insertion of the expression Y(z) into the continued fraction yielding the desired result,



Doing so, results in the ability to take the limit $\delta \rightarrow 0$, resulting in a well resolved LDOS as seen in **Figure 3B**.

This method can be generalized to the case when b_n is alternating between two or more values. This can happen if there is a gap in the band structure, such as when there are different on-site energies for opposite sublattices of a bipartite lattice. The continued fraction is then considered periodic and a similar but slightly more complicated expression can be found for Y(z).



FIGURE 3 | (A) Hopping terms along the equivalent chain for a square lattice. Values oscillate around and converge to 2, while $a_n = 0$ for all *n*. **(B)** Resulting LDOS. The length of the chain is L = 400 before insertion of the expression Y(z) as described in the text.

2.2.1. Block Lanczos

In this section, an extension of the previous method based on the block Lanczos transformation [51, 52] for multiple orbital or multi-impurity problems will be presented, following the original work in Shirakawa and Yunoki [29] and Allerdt et al. [28]. These considerations apply to different orbitals of the same atom, or impurities sitting at different lattice sites. As before, the first step is to choose the seed states. For illustrative purposes, let us consider a square lattice with two seeds at positions \mathbf{r}_1 and \mathbf{r}_2 , defined as:

$$egin{aligned} &|lpha_0
angle \,=\, c^{\dagger}_{\mathbf{r}_1} \,|0
angle \ &|eta_0
angle \,=\, c^{\dagger}_{\mathbf{r}_2} \,|0
angle \,, \end{aligned}$$

with the additional requirement that $\langle \alpha_0 | \beta_0 \rangle = 0$. Once the seeds are chosen, a new set of states can be obtained through the block Lanczos method [53]. The idea is the same: apply the Hamiltonian to each state and require orthogonality. The new set of states can be found through the iterative procedure as

$$\begin{aligned} |\alpha_{n+1}\rangle &= H |\alpha_n\rangle - a_n^{\alpha\alpha} |\alpha_n\rangle - a_n^{\alpha\beta} |\beta_n\rangle - b_n^{\alpha\alpha} |\alpha_{n-1}\rangle \\ &- b_n^{\alpha\beta} |\beta_{n-1}\rangle \end{aligned} \tag{23} \\ |\beta_{n+1}\rangle &= H |\beta_n\rangle - a_n^{\beta\beta} |\beta_n\rangle - a_n^{\beta\alpha} |\alpha_n\rangle - b_n^{\beta\beta} |\beta_{n-1}\rangle \\ &- b_n^{\beta\alpha} |\alpha_{n-1}\rangle \,. \end{aligned}$$

By requiring the states $|\alpha_n\rangle$, $|\beta_n\rangle$ to be orthogonal to all $|\alpha_{k< n}\rangle$, $|\beta_{k< n}\rangle$, the *a* and *b* coefficients can be found by solving the following matrix equations:

$$\begin{pmatrix} \langle \alpha_{n} | H | \alpha_{n} \rangle & \langle \beta_{n} | H | \alpha_{n} \rangle \\ \langle \alpha_{n} | H | \beta_{n} \rangle & \langle \beta_{n} | H | \beta_{n} \rangle \end{pmatrix} \begin{pmatrix} a_{n}^{\alpha \alpha} & a_{n}^{\alpha \beta} \\ a_{n}^{\beta \alpha} & a_{n}^{\beta \beta} \\ a_{n}^{\beta \alpha} & a_{n}^{\beta \beta} \end{pmatrix} \begin{pmatrix} \langle \alpha_{n} | \alpha_{n} \rangle & \langle \beta_{n} | \alpha_{n} \rangle \\ \langle \alpha_{n} | \beta_{n} \rangle & \langle \beta_{n} | \beta_{n} \rangle \end{pmatrix}$$
(25)

$$\begin{pmatrix} \langle \alpha_{n-1} | H | \alpha_n \rangle & \langle \beta_{n-1} | H | \alpha_n \rangle \\ \langle \alpha_{n-1} | H | \beta_n \rangle & \langle \beta_{n-1} | H | \beta_n \rangle \end{pmatrix} = \\ \begin{pmatrix} b_n^{\alpha \alpha} & b_n^{\alpha \beta} \\ b_n^{\beta \alpha} & b_n^{\beta \beta} \end{pmatrix} \begin{pmatrix} \langle \alpha_{n-1} | \alpha_{n-1} \rangle & \langle \beta_{n-1} | \alpha_{n-1} \rangle \\ \langle \alpha_{n-1} | \beta_{n-1} \rangle & \langle \beta_{n-1} | \beta_{n-1} \rangle \end{pmatrix}.$$
(26)

A solution to these is ensured by the fact that $\langle \alpha_n | \alpha_n \rangle \neq \langle \alpha_n | \beta_n \rangle$. The block Lanczos recursion will result in the geometry of a ladder as shown in **Figure 4**. Note that the states are not normalized at this stage. An important aspect of this method is that states within the same block are not generally orthogonal, i.e., $\langle \alpha_n | \beta_n \rangle \neq 0$ (resulting in the vertical rung coupling in the ladder). Since we are interested in an orthonormal basis, we can use a Gram Schmidt procedure—although there are many ways—, to orthogonalize them. This is done by choosing

$$|x_n\rangle = |\alpha_n\rangle; \qquad (27)$$

$$|y_n\rangle = |\beta_n\rangle - \langle \alpha_n |\beta_n\rangle |\alpha_n\rangle,$$
 (28)

which will also simplify the geometry causing some hoppings in the ladder to vanish. If the seed orbitals are spatially separated such that the Lanczos orbitals have not yet met or overlapped, the hoppings connecting the legs of the ladder will remain zero and the hoppings along the chain will be equal to the single seed case.

Our Hamiltonian can now be written in the desired tridiagonal form:

$$H_{\text{band}} = \begin{pmatrix} A_0 & B_1 & 0 & 0 & \cdots \\ B_1 & A_1 & B_2 & 0 & \\ 0 & B_2 & A_2 & B_3 & \\ 0 & 0 & B_3 & A_3 & \\ \vdots & & \ddots & \ddots \end{pmatrix},$$
(29)

where A_n and B_n are 2×2 matrices. These ideas can readily be generalized to an arbitrary number of impurities. For kimpurities, each A and B matrix will be $k \times k$. As long as the initial states are orthogonal, a new set of states can be found as

$$|\lambda_{n+1}\rangle = H |\lambda_n\rangle - \sum_{\lambda'} \left(a_n^{\lambda\lambda'} |\lambda'_n\rangle + b_n^{\lambda\lambda'} |\lambda'_{n-1}\rangle \right).$$
(30)

This matrix represents a new non-interacting tight-binding Hamiltonian where each block becomes a "unit cell" (there is no-translational invariance in this case, though).

Intuitively, we can understand this geometry as follows: if the impurities are far apart, we can imagine performing



independent single-impurity mappings for each of them that yield two decoupled one-dimensional chains. At some point, when the length of the chains is about half the separation distance between impurities, the orbitals interfere and orthogonality is lost. Therefore, after re-orthogonalizing them, we end up introducing mixing that translates into a hopping term between the chains, leading to the resulting ladder.

For k impurities, it can be recognized as k coupled chains forming a $k \times L$ ladder. The new geometry is now quasi one-dimensional.

2.2.2. Symmetric and Anti-symmetric Channels

In the presence of inversion symmetry, the legs of the ladder can be decoupled into independent chains [28]. This method can be employed in two-impurity problems on simple lattices such as square, hexagonal, cubic, triangular, etc. An example of when it can not be used is for Bernal stacked graphene when the seeds are both on A-sites of opposite layers. Following Allerdt et al. [28], the first step is to choose the initial seed states. This symmetrization is similar to a folding transformation used in bosonization and NRG [54–58] and is carried out by choosing symmetric and antisymmetric (or bonding and anti-bonding) combinations of the original single particle states. For two seeds at \mathbf{r}_1 and \mathbf{r}_2 , they read:

$$|\psi_{0}^{+}\rangle = c_{+}^{\dagger} = \frac{1}{\sqrt{2}}(c_{\mathbf{r}_{1}}^{\dagger}|0\rangle + c_{\mathbf{r}_{2}}^{\dagger}|0\rangle);$$
 (31)

$$|\psi_{0}^{-}\rangle = c_{-}^{\dagger} = \frac{1}{\sqrt{2}}(c_{\mathbf{r}_{1}}^{\dagger}|0\rangle - c_{\mathbf{r}_{2}}^{\dagger}|0\rangle).$$
 (32)

Once the initial states are constructed, the Lanczos recursion is identical to the one seed case. The procedure is carried out for each initial state yielding two completely independent and orthogonal chains. Only after the impurities are introduced are these chains coupled. The resulting geometry is shown in **Figure 5** including the impurities. This transformation however, does not leave V_c unchanged. We illustrate it for the case of the two-impurity Kondo model introduced above in Equation (1). To write the many-body terms of the Hamiltonian in the new basis we use the inverse transformation,

$$c_{\mathbf{r}_{1,2}}^{\dagger} = \frac{1}{\sqrt{2}} \left(c_{+}^{\dagger} \pm c_{-}^{\dagger} \right),$$
 (33)



where $c^{\dagger}_{+}, c^{\dagger}_{-}$ are defined in Equations (31) and (32), respectively, and the fact that

$$S_i^z = \frac{1}{2} \left(n_{i\uparrow} - n_{i\downarrow} \right), \quad S_i^+ = c_{i\uparrow}^\dagger c_{i\downarrow}, \quad S_i^- = c_{i\downarrow}^\dagger c_{i\uparrow},$$
$$n_{i\sigma} = c_{i\sigma}^\dagger c_{i\sigma}, \qquad (34)$$

one arrives at the transformed Hamiltonian:

$$V_{c}' = \frac{J_{K}}{2} \left(\vec{S}_{1} + \vec{S}_{2} \right) \cdot \sum_{\mu,\nu,\gamma} c_{\gamma\mu}^{\dagger} \vec{\sigma}_{\mu\nu} c_{\gamma\nu} + \frac{J_{K}}{2} \left(\vec{S}_{1} - \vec{S}_{2} \right).$$
$$\sum_{\mu,\nu,\gamma} c_{\gamma\mu}^{\dagger} \vec{\sigma}_{\mu\nu} c_{-\gamma\nu}.$$
(35)

By applying this symmetry, not only the recursion is greatly simplified, but also that the equivalent problem reduces from a ladder to a chain. Since the real space entanglement on a chain is half of that on the ladder, this results into an exponential gain that dramatically improves the efficiency of the DMRG simulations.

2.3. Star Geometry

In certain cases, it can be beneficial to transform from a chain to a star geometry. It was recently shown that a star geometry results in lower ground state entanglement than the chain [59], making it ideal for DMRG calculations.

The single impurity case is presented here and same considerations can easily be extended to multiple impurities. This mapping corresponds to a second unitary transformation on top of the original lattice-to-chain conversion. In order to do this mathematically and numerically, the matrix H_l is split into two blocks (similar to the partition when calculating the LDOS with the continued fractions). One contains just the seed state, which remains unaltered and the other block contains the rest. Explicitly, Equation (9) is split into

$$H_{l} = \begin{pmatrix} \frac{a_{0} \mid b_{1} \mid 0}{b_{1} \mid H'} \\ 0 \mid \end{pmatrix}.$$
 (36)

The block corresponding to the non-interacting chain H' is then diagonalized. The couplings to the seed site are then $V_n = b_1 a'_n$, where a'_n are the values of the n^{th} eigenvector of H' on the site



 H'_{00} (or the second site of the chain). Equation (36) then becomes

$$H_{l} = \begin{pmatrix} a_{0} & V_{1} & V_{2} & V_{3} & \cdots \\ V_{1} & \epsilon_{1} & 0 & 0 & \cdots \\ V_{2} & 0 & \epsilon_{2} & 0 & \cdots \\ V_{3} & 0 & 0 & \epsilon_{3} & \cdots \\ \vdots & \vdots & \vdots & \vdots & \ddots \end{pmatrix} ,$$
(37)

where a_0 corresponds seed of the Lanczos transformation and the center sites in **Figures 6A,B**. The new bath energies ϵ_n are just the eigenvalues of H'.

To extend this idea to two impurities is straight forward. The Hamiltonian after the Lanczos transformation is again split into blocks, where now the first block contains both seed states. Obtaining the new hoppings and bath energies is identical to the single impurity case. The new form of Equation (29) will then be

$$H_{l} = \begin{pmatrix} a_{0}^{\alpha\alpha} & a_{0}^{\alpha\beta} & V_{1}^{\alpha} & V_{2}^{\alpha} & \cdots \\ a_{0}^{\beta\alpha} & a_{0}^{\beta\beta} & V_{1}^{\beta} & V_{2}^{\beta} & \cdots \\ V_{1}^{\alpha} & V_{1}^{\beta} & \epsilon_{1} & 0 & \cdots \\ V_{2}^{\alpha} & V_{2}^{\beta} & 0 & \epsilon_{2} & \cdots \\ \vdots & \vdots & \vdots & \vdots & \ddots \end{pmatrix}.$$
 (38)

The resulting geometry is shown in **Figure 7**. Depending on the symmetry of the lattice and impurities, certain V_n 's will vanish, as in the case where the symmetric/antisymmetric seeds are used. Here, the most general form is shown.

In some circumstances, the impurities entangle strongly to some states in the metallic substrate within a window of width $\sim J_K$ around the Fermi energy, while a large fraction remains disentangled. When this occurs, it is convenient to use the star geometry, which allows one to reach extremely large systems, practically in the thermodynamic limit. The advantage here is that after the transformation to the star, some bath sites (in the interacting problem) will be double occupied and some will be empty. Observing that these states play no role in the physics, they can be safely discarded. For an application we refer the reader to Allerdt et al. [60].



3. DMRG SIMULATIONS

In the block Lanczos approach, the equivalent problem consists of a ladder with k legs, where k is the number of impurities. As discussed above, by means of a folding transformation one can reduce the entanglement by a factor of 2, with the consequent reduction in the number of DMRG states needed in the calculation. Typically, for S = 1/2 impurities we take a total system size to be L = 4n (including impurities), such that each impurity can form part of a collective RKKY state or its own Kondo cloud (it has been already observed that Kondo does not develop in chains of length L = 4n + 2 [61]). For instance, L = 204 corresponds approximately to a "sphere" around the two impurities of radius $\sim\,$ 100. As explained above and described in Büsser et al. [27], we assume that the lattice has the same shape/symmetry as the orbitals. When this is the case, orbitals never "bounce back" inward and the Lanczos recursion can be stopped at an arbitrary number of iterations without introducing any bias. Typically, we consider system sizes at least 4 times larger than the maximum inter-impurity distance. Notice that each impurity configuration will generate a new mapping and the shape of the lattice will be modified as a consequence, depending on the relative position of the impurities and the number of Lanczos iterations. This could potentially lead to artifacts and care must be taken to make sure that no finite size effects are introduced.

Since the energy difference between the ground-state and the first excited state can be very small ($\sim 10^{-6}$), we fix the truncation error at 10^{-9} in all simulations. This translates into a number of DMRG states of the order of 3,000 or more in most cases. Achieving this level of accuracy in a ladder geometry (without the bonding-antibonding symmetrization) would be practically impossible due to the larger entanglement. However, as previously pointed out, it is possible to work in the star geometry directly. This is particularly ideal in the strong coupling limit, since the impurity is mostly entangled to an electron at position r_0 . The rest of the free electrons remain disentangled and, in "momentum" (or energy) space, they practically form a product state. In this basis, the number of states required to achieve the same accuracy is reduced dramatically.

4. ENERGY SCALES

As suggested by Doniach [62] (see also [63]), one could define a binding energy (or "Kondo temperature") for forming a Kondo singlet $T_K \simeq e^{-1/J_K}$, or an RKKY state, $T_{RKKY} \sim J_K^2$ and a competition between these two energy scales will dictate which phase will win. Another consideration that plays an important role in finite systems is the energy level spacing Δ that introduces another competing energy scale which can become significant in, for example, quantum dots [64]. The Kondo effect in small or confined systems has been referred to as the "Kondo Box." It is expected that Δ will dominate the physics when $T_K \sim \Delta$. Large amounts of work have been dedicated to studies of the competition between T_K and Δ , which is a complicated problem, as well as scaling in the presence of finite bandwidth [65]. A fully developed resonance (fully screened impurity) requires a finite density of states at the Fermi level. In this regime, even/odd effects in chain length and particle number are also observed [63, 66].

In order to understand the relevance of these scales in numerical simulations we first analyze the single impurity case at zero temperature. According to renormalization group arguments, the system flows to a strong coupling fixed point characterized by a single energy scale—the Kondo temperature T_K . In this regime, the impurity and the conduction electrons form a scattering state, the "Kondo cloud," that has a characteristic extension R_K [22–24, 67–70] that depends on J_K (or T_K). At distances of the order of R_K electrons are more entangled to the impurity. In finite systems where the conduction electrons are confined to a "Kondo box" [64, 71–74] *all* the electrons may all be inside the Kondo cloud, without an "outside." This regime would correspond to the "crossover" between weak coupling and strong coupling in the RG flow.

The strong coupling limit can be explicitly realized by taking $J_K \gg W$, where W is the bandwidth, and physically results into a tightly bound singlet formed by the impurity and a localized electron at r_0 . This corresponds to $R_K = 0$. As J_K is reduced, the impurity will become correlated with electrons farther and farther from it and R_K will increase. If the system is finite, at some point we will find $R_K \sim L$ and the Kondo singlet will extend to the entire volume and the impurity will couple mostly to one electron at the Fermi level. According to the analysis presented in Yang and Feiguin [75], practically a single conduction electron is responsible for most of the screening: In the weak-coupling limit, this electron is precisely at the Fermi level [63], while in the strong-coupling limit, it corresponds to the localized orbital directly in contact with the impurity spin. In both cases, the remaining conduction electrons form a completely disentangled Fermi sea. As a consequence, it is important in finite systems to have a single occupied state at the Fermi energy in order to realize screening (otherwise, the impurity will behave as a local moment, decoupled from the Fermi sea). This situation, however, is far from the universal, strong coupling regime [65].

To make these observations more explicit, we study the behavior of gap Δ_0 to the first (triplet) excited state as a function

of system size. We focus our discussion at $J_K = 1$, where the effects are more obvious. In the thermodynamic limit the system is gapless, $\Delta_0 \rightarrow 0$ as $L \rightarrow \infty$: it costs no energy to flip a spin infinitely far from the impurity. In addition, we introduce the "correlation gap," defined as:

$$\Delta_K = \frac{\langle g.s.|S^-_{imp}HS^+_{imp}|g.s.\rangle}{\langle g.s.|S^-_{imp}S^+_{imp}|g.s.\rangle} - E_0.$$
(39)

The first term is a variational energy obtained by flipping the spin of the impurity with the S_{imp}^+ operator. This action will disentangle the impurity from the chain and destroy the Kondo singlet. Equivalently, one could apply the S_{imp}^z operator to transform the Kondo singlet into a "Kondo triplet." Both quantities are equivalent in the SU(2) symmetric case. This estimate will yield what we identify as the minimum energy cost for destroying the Kondo state, which can also be related to the Kondo temperature.

For small systems the energy spacing Δ is large and the gap Δ_0 is determined by the correlation gap Δ_K . As we increase the size of the Fermi sea, the level spacing is reduced and, at some point, Δ will become smaller than the correlation gap. When this happens, the system gap Δ_0 follows the behavior of the level spacing and shrinks as 1/L, as clearly observed in **Figure 8**: the system undergoes a crossover from the weak Kondo-box-like regime to the universal strong coupling regime and Δ_K becomes a system-independent characteristic energy scale.

One way to reach this regime in finite systems is by introducing "damped boundary conditions" [76]. In this particular case, we multiply the hoppings by a factor $\lambda^{-n/2}$, where λ is a number larger than one and *n* is the position along the chain, counting from L_b sites before the boundary (we pick $L_b = 15$ in this case). This is reminiscent of Wilson leads, which have been successfully used in a number of contexts (besides NRG calculations) and are known to alleviate finite-size effects [77, 78] by making the energy spacing smaller around the Fermi energy. To illustrate this behavior we also show Δ , Δ_0 and Δ_K



for different system sizes and $J_K = 1$. We find that, as long as the "bulk" level spacing is large (for $\lambda = 1$), the gap Δ does not depend on the chain length, but only on λ and L_b (which is kept constant). Therefore, now one needs to focus on the behavior of the different quantities with λ and not L. As we see, for all values of λ considered here Δ is smaller than Δ_K and is practically equal to Δ_0 . In addition, we see that while Δ gets smaller with increasing λ , the correlation gap Δ_K stays constant, indicating that this is basically a measure of the correlation energy in the thermodynamic limit.

In the multi-impurity problem the correlation gap cannot distinguish between Kondo or RKKY physics and the origin of the correlation effects. In the two-impurity case, one compares Δ_K against *twice* the value for a single impurity. The difference indicates the energy gained by forming a correlated RKKY state vs. to the one gained by two independent Kondo singlets.

5. APPLICATIONS

The single impurity mapping can be used to study Kondo physics on graphene [79] or multi-band problems such as phosphorene, for instance [80]. It was found that, in the case of graphene, edge states tend to hybridize with the impurity spin introducing a competition between bulk and edge-induced Kondo screening.

The multi-impurity version allows one to explore a broad range of interesting open problems. Although briefly suggested in Büsser et al. [27], the block Lanczos method was formally introduced in Shirakawa and Yunoki [29] to study spin-spin correlations between an impurity and conduction electrons in graphene. The folding transformation was later used to investigate the competition between Kondo and RKKY physics on different geometries with small [28] and large spin [81] and a number of surprising results were found. Notably, there are important non-perturbative effects that in some cases make Kondo more robust than an RKKY state, even at short distances [28] of the order of a few lattice sites. This is a consequence of lattice effects that make the RKKY interaction very weak when impurities are sitting at positions where single particle wave-functions interfere destructively. See for instance Figure 1, the single particle wave functions that expand away from the impurity site have very small amplitude on the opposite sublattice, hindering the possibility of mediating an RKKY exchange interaction between impurities. Moreover, due to the fast decay of the RKKY interaction, Kondo screening becomes the dominant effect at relatively short inter-impurity separation. Interestingly, in the spin-1 problem, it was found that impurities can be practically screened by the substrate, while simultaneously forming an RKKY state [81].

The method is particularly suited to study the structure of the correlations around a single impurity. In this case, one seed site is connected to the impurity and the second one is used to measure the two-point correlations between the conduction electron in the substrate and the impurity spin [29]. For instance, one could place the impurity at the edge or bulk of the material and calculate the correlations around substitutional impurities and adatoms at the edge of a topological insulator [82, 83]. Unlike previous calculations that consider the coupling of the impurity to one-dimensional effective modes [84, 85], in this formulation the helical liquid arises naturally as an edge effect of the twodimensional bulk. Similarly, one could place the impurity on metallic (111) surfaces and study the influence of bulk states and Shockley surfaces states on the RKKY interaction [60].

Remarkably, this approach can be extended to a onedimensional interacting "wire" connected to a substrate [86, 87]. The many-body physics in this case is contained in the wire (for instance, a Hubbard chain), while the substrate remains non-interacting. The mapping leads to Lanczos orbitals that describe cylindrical "shells" centered at the wire as their axis of symmetry. The generalization takes advantage of the translational invariance of the system along the wire direction that results into the wire being connected to an array of independent semiinfinite 1D chains, each characterized by a momentum k_x . In this representation the chain Hamiltonian becomes non-local due to the interaction terms. Other mappings are possible, mixing real-space and momentum representation. It is convenient to preserve the locality of the interaction terms in the chain by using the real-space representation of it. As a price, the tunneling terms between the chain and the substrate become non-local. In the case of an insulating substrate, the length of the chains can be truncated and the system can be solved with the DMRG.

6. CONCLUSIONS

We have discussed a powerful method to study multi-impurity problems in realistic lattice geometries and configurations. This approach can readily be generalized to study generic band

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structures, multi-orbital problems, and magnetic molecules. It can be used in conjunction with first-principles band structure calculations, where the information about the metallic substrate is calculated using density functional theory. Quantum chemistry methods such as CASPT2 or NEVPT2 can be used to precisely calculate the hybridization terms in the structure of the seed state, paving the way toward a first principles modeling and a better understanding of correlation effects in quantum impurity problems.

We point out that any quadratic Hamiltonian can be mapped onto an equivalent one-dimensional system following this prescription, including problems with spin-orbit interaction, disorder and superconductors (at the mean field/BdG level).

Although the DMRG method was used as a solver for the effective one-dimensional impurity Hamiltonian, other methods are equally applicable, such as the embedded cluster approximation [24] or quantum Monte Carlo. One could also use these techniques to study time-dependent [88, 89] and thermodynamic properties [90], or to calculate spectral functions [91, 92].

AUTHOR CONTRIBUTIONS

All authors listed have made a substantial, direct and intellectual contribution to the work, and approved it for publication.

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Conflict of Interest Statement: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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