

# Characterizing Quantum-Dot Cellular Automata

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# Chapter 1

## Approximations

### 1.1 Fixed charge model

Exact diagonalization scales exponentially with system size. For the full *grand canonical* QCA Hamiltonian, Eq. ??, only QCA devices of up to two cells are computationally feasible. Therefore, to access larger systems we need to introduce approximations. Approximating means to simplify. However, by carefully establishing successive approximations and their limits, we also reduce the problem to its essential ingredients and thus, hopefully, we gain a better understanding of the system. As a first step, we reduce the Hilbert space to a *fixed* number of particles per cell. We disallow any charge fluctuations, both for the system as a whole and for each individual cell. With that, we omit the chemical potential term in the Hamiltonian,  $\mu = 0$ , and prohibit inter-cell hopping. This is a major simplification. However, it is in line with the QCA idea: The approach requires a fixed number of charges per cell, typically two electrons, and cells are thought to interact only via Coulomb forces. In a sense we are shifting the starting point of our investigation. If the *fixed* charge approximation is not valid for a given system, then there is no hope of implementing QCA on it. On the other hand, for experimental systems like the atomic silicon quantum dots and for a given cell layout, it should always be possible, at least in principle, to tune the system parameters, especially the chemical potential, to get the system into the right particle number sector. The system has to be set up in a way that the two-electrons-per-cell sector is lowest in energy and other particle number sectors are sufficiently gapped out, that is, at an energy, compared to the ground state energy, much larger than temperature. Of course, in practice there are very clear limits as to how much the system parameters can be tuned and any QCA cell layout considered within the *fixed* charge approximation cannot necessarily be readily implemented on a given real-world material system.

For the *fixed* charge system, the state space scales as  $N_s = \binom{8}{2}^{N_c} = 28^{N_c}$  ( $N_c$  is the number of cells). Using symmetries, the largest block of the Hamiltonian matrix is the spin zero sector, of size  $N'_s = 16^{N_c}$ . On conventional computer hardware, systems of

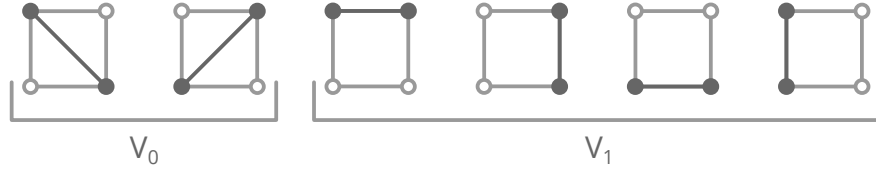


Figure 1.1: ...

up to four cells are possible, with memory requirements of 32GB. In practice, however, calculations for four-cell systems take too long and thus three cells is the practical limit for the *fixed*-number-of-particles-per-cell model.

## 1.2 Bond model

At its heart, QCA is a semi-classical idea. It relies solely on charge-charge interactions and ignores the particle spins. Therefore, as a next step in our quest to access larger system sizes, we neglect the spin degree of freedom in our model. The 28 states per cell of the *fixed* charge model can be reorganized into four doubly occupied dots and six bonds. The six bonds are illustrated in Fig. 1.1. Each bond corresponds to one spin singlet and three spin triplet states. The *bond* approximation only keeps one state for each bond and discards the doubly occupied states as well. With the *bond* model we thus assume that singlet and triplet states are qualitatively equivalent and energetically degenerate, and that doubly occupied dots are sufficiently gapped out, that is,  $U \gg T$ . As QCA ignores the spin, singlets and triplets should be qualitatively equivalent, but they are not quite degenerate. We expect that virtual double-occupancy lowers the energies of the singlet states and therefore introduces a small singlet-triplet splitting. Still, degeneracy is presumably not a bad assumption to start with and we will look at the singlet-triplet splitting in detail in due course. For the *bond* model the QCA Hamiltonian reduces to

$$H = - \sum_{\langle ij \rangle} t c_i^\dagger c_j + \sum_{i < j} V_{ij} (n_i - q) (n_j - q) . \quad (1.1)$$

With six bond states per cell, the Hilbert space of the *bond* model is  $N_s = 6^{N_c}$  ( $N_c$  the number of cells). Five and six cells are doable, with memory requirements of 460MB and 16GB, respectively, but for practical calculations five cells really is the limit. For the *bond* model there are no symmetries that can be exploited.

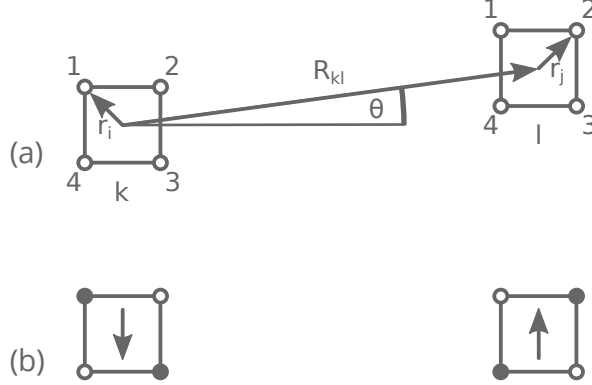


Figure 1.2: a) QCA cells  $k$  and  $l$ . b) The two-states-per-cell approximation identifies each cell with a spin  $\uparrow$  or  $\downarrow$ .

### 1.3 Ising model

A linear array of QCA cells where each cell has a state of logic 0 or 1 is reminiscent of a 1D spin  $\frac{1}{2}$  chain. Indeed, if we reduce the basis to only two states per cell, down from six states in the *bond* picture, we can map the QCA system to a transverse-field Ising model with long-ranging interactions. This is an attractive proposition: The smaller Hilbert space allows for larger system sizes with our exact diagonalization method; more importantly, the transverse-field Ising model is amenable to sign-problem-free Stochastic series expansion (SSE) quantum Monte Carlo schemes [1]. These methods do not scale exponentially<sup>1</sup> and consequently allow access to much larger systems. Last, but not least, such a mapping connects the QCA approach to the established and well studied Ising model. The prospect hinges on the assumption that the two-states-per-cell basis actually is a good approximation for QCA systems. And while bistable two-state cells are certainly the picture we have in mind when we talk about QCA, it is not *a priori* clear whether this is a correct physical picture.

We use the *bond* Hamiltonian (1.1) as the starting point. We had already discussed in the last chapter that such a Hamiltonian can be decomposed into single-cell terms and cell-cell interaction terms,

$$H = \sum_k H_k^c + \sum_{k < l} H_{kl}^{cc}. \quad (1.2)$$

In comparison, the transverse field Ising model is described by

$$\tilde{H} = - \sum_k \gamma S_k^x + \sum_{k < l} J_{kl} S_k^z S_l^z. \quad (1.3)$$

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<sup>1</sup>SSE quantum Monte Carlo methods roughly scale as  $N \ln N$  where  $N$  is the system size.

Thus, we would like to map the single cell term  $H_k^c$  to the transverse field term  $-\gamma S_k^x$  and the Coulombic cell-cell interaction  $H_{kl}^{cc}$  to the Ising term  $J_{kl} S_k^z S_l^z$ . Each cell  $k$  is identified with a pseudo spin  $S_k^z$ , specifically the logic 0 with a spin down state and the logic 1 with a spin up state, as illustrated in Fig. 1.2(b). We will first look at how the QCA cell can be represented by only two states and derive an approximate expression for the transverse field  $\gamma$ . Then we will use a multipole expansion to derive  $J_{kl}$  from the cell-cell Coulomb interaction.

To arrive at a single-cell-basis with only two states we can, in principle, follow a similar prescription as for the fixed charge and bond approximations: We neglect high energy states which are assumed to be gapped out. In this case these are the four edge states with Coulomb energy  $V_1$ ,  $|\psi_Q\rangle = \{|3\rangle, |4\rangle, |5\rangle, |6\rangle\}$  in Fig. 1.1, where we have introduced  $|\psi_Q\rangle$  to denote the high-energy subspace of the single-cell Hilbert space. We only keep the low-energy, diagonal states  $|\psi_P\rangle = \{|1\rangle, |2\rangle\}$  with Coulomb energy  $V_0$ . Of course, these two are exactly our logic 0 and logic 1 state, or  $|1\rangle \doteq |\downarrow\rangle$  and  $|2\rangle \doteq |\uparrow\rangle$ , respectively. Here,  $|\psi_P\rangle$  denotes the low-energy subspace. For the high-energy states to be sufficiently gapped out we require  $\Delta V = V_1 - V_0 \gg T$ . In contrast to the fixed charge and bond models, merely truncating the Hilbert space is not sufficient for the Ising model. For our previous two approximations the Hamiltonian had remained essentially unchanged, apart from dropping no longer relevant terms, such as the chemical potential term or the Hubbard  $U$  term. The retained states were exactly the same states as in the full, untruncated model. But with only two states per cell the existing Hamiltonian (1.1) does not “work”: There is no process that takes the system from state  $|1\rangle$  to  $|2\rangle$ . Therefore, for the Ising approximation we need to derive an effective, low-energy Hamiltonian from the bond model. In the bond picture, for the system to transition from state  $|1\rangle$  to  $|2\rangle$  it can take different paths, for example  $|1\rangle \rightarrow |3\rangle \rightarrow 2$ , consisting of two hopping processes with an interim high-energy edge state. We will treat those processes perturbatively, as *virtual* excitations, and derive an effective hopping term between the two states  $|1\rangle$  and  $|2\rangle$ . This effective hopping term is precisely the transverse field  $\gamma$  which flips the spin in the Ising picture,  $-\gamma S_k^x = -\gamma \frac{1}{2} (S_k^+ + S_k^-)$ .

Using the basis depicted in Fig. 1.1 the single-cell bond Hamiltonian is very simple and can be written down explicitly. ...

To arrive at only two states per cell, we follow a similar prescription as for the *fixed* charge and *bond* approximations: We neglect high energy states which are assumed to be gapped out, in this case, the four edge states with Coulomb energy  $V_1 = V_0 + \Delta V$ ,  $|3\rangle, |4\rangle, |5\rangle$ , and  $|6\rangle$  in Fig. 1.1. We only keep the low energy states  $|1\rangle$  and  $|2\rangle$  with Coulomb energy  $V_0$ . Therefore, we require  $\Delta V \gg T$ . For the *fixed* charge and *bond* approximation we had only truncated the Hilbert space, but the Hamiltonian had remained unchanged, apart from removing no longer relevant terms, such as the chemical potential term or the Hubbard  $U$  term. The retained states were exactly the same states as in the full, untruncated model. The situation is more complex in the case of the Ising approximation. Simply discarding high energy states while keeping the same Hamiltonian is not sufficient,



for the two states  $|1\rangle$  and  $|2\rangle$  would not interact in any way. We need to derive an effective hopping term between the two states. The Ising approximation has to be derived as an effective low-energy model from the *bond* Hamiltonian.

For each cell  $k$  the two retained low energy states can be identified with a pseudo spin  $\sigma_l^z$ . More precisely, logic 0 corresponds to  $\downarrow$  and logic 1 to  $\uparrow$ , see Fig. 1.2(b). We had already seen in the last chapter that in the absence of inter-cell hopping the QCA Hamiltonian and thus also the simplified bond Hamiltonian, Eq. 1.1, can be decomposed into single-cell and cell-cell terms,

$$H = - \sum_k H_k^c + \sum_{k<l} H_{kl}^{cc}. \quad (1.4)$$

If the transverse field Ising model is a valid description of QCA systems, then we expect the single cell term  $H_k^c$  to map to a transverse field term  $-\gamma\sigma_k^x$  and the cell-cell term  $H_{kl}^{cc}$  to an Ising term  $J_{kl}\sigma_k^z\sigma_l^z$ . We will now first look at the single-cell term and then at the cell-cell interaction term.

To derive an effective Hamiltonian for the two-states-per-cell QCA basis we consider the *bond* Hamiltonian, Eq. 1.1 for a single cell. Using the basis depicted in Fig. 1.1 the single cell Hamiltonian matrix is very simple and can be written down explicitly.

$$H = \left( \begin{array}{cc|cccc} V_0 & 0 & -t & -t & -t & -t \\ 0 & V_0 & -t & -t & -t & -t \\ \hline -t & -t & V_1 & 0 & 0 & 0 \\ -t & -t & 0 & V_1 & 0 & 0 \\ -t & -t & 0 & 0 & V_1 & 0 \\ -t & -t & 0 & 0 & 0 & V_1 \end{array} \right) \quad (1.5)$$

$$= \begin{pmatrix} H_{PP} & H_{PQ} \\ H_{QP} & H_{QQ} \end{pmatrix}$$

Our aim is to truncate the basis to  $|\psi_P\rangle = \{|1\rangle, |2\rangle\}$  and derive an effective Hamiltonian  $\tilde{H}$  (with  $\tilde{H}|\psi_P\rangle = E|\psi_P\rangle$ ) which incorporates virtual excitations to the high energy states  $|\psi_Q\rangle = \{|3\rangle, |4\rangle, |5\rangle, |6\rangle\}$ . The time-independent Schrödinger equation  $H|\psi\rangle = E\psi$  is written

$$\begin{pmatrix} H_{PP} & H_{PQ} \\ H_{QP} & H_{QQ} \end{pmatrix} \begin{pmatrix} \psi_P \\ \psi_Q \end{pmatrix} = E \begin{pmatrix} \psi_P \\ \psi_Q \end{pmatrix} \quad (1.6)$$

Writing out the two matrix equations explicitly and eliminating  $|\psi_Q\rangle$  yields

$$H_{PP}|\psi_P\rangle + H_{PQ} \frac{1}{E - H_{QQ}} H_{QP} |\psi_P\rangle = E|\psi_P\rangle \quad (1.7)$$

and therefore

$$\tilde{H} = H_{PP} + H_{PQ} \frac{1}{E - H_{QQ}} H_{QP}. \quad (1.8)$$

Assuming that the system is predominantly in the subspace spanned by  $|\psi_P\rangle$  and additionally that the hopping is very small,  $t \ll V_1$ , we can write  $E \approx V_0$ . The effective Hamiltonian becomes

$$\begin{aligned}\tilde{H}_{ij} &= (V_0)_{ii} \delta_{ij} + (-t)_{ik} (V_0 - V_1)_{kk}^{-1} (-t)_{kj} \\ &= (V_0)_{ii} \delta_{ij} - \left( \frac{4t^2}{\Delta V} \right)_{ij},\end{aligned}\tag{1.9}$$

where we have used  $\Delta V = V_1 - V_0$ ,  $H_{PP} = (V_0)_{ii} \delta_{ij}$ ,  $H_{PQ} = (-t)_{ij}$ , and so on. We can subtract the constant diagonal term  $\tilde{H}_{ii} = V_0 - \frac{4t^2}{\Delta V}$  and arrive at

$$\tilde{H} = \begin{pmatrix} 0 & -\frac{4t^2}{\Delta V} \\ -\frac{4t^2}{\Delta V} & 0 \end{pmatrix}.\tag{1.10}$$

Comparing this with the transverse field term of the Ising model,

$$H_k = -\gamma \sigma_k^x = -\gamma \frac{1}{2} (\sigma_k^+ + \sigma_k^-) = \begin{pmatrix} 0 & -\frac{1}{2}\gamma \\ -\frac{1}{2}\gamma & 0 \end{pmatrix}\tag{1.11}$$

where we have used the basis  $|1\rangle \doteq |\downarrow\rangle$  and  $|2\rangle \doteq |\uparrow\rangle$ , we identify

$$\gamma = \frac{8t^2}{\Delta V}.\tag{1.12}$$

We now turn our attention to the cell-cell interaction term  $H_{kl}^{cc}$  which is

$$\begin{aligned}H_{kl}^{cc} &= \sum_{\substack{i \in k \\ j \in l}} V_{ij} (n_i - q) (n_j - q) \\ H_{kl}^{cc} &= \sum_{\substack{i \in k \\ j \in l}} (n_i - q) (n_j - q) |\mathbf{R}_{kl} + \mathbf{r}_j - \mathbf{r}_i|,\end{aligned}\tag{1.13}$$

where  $i$  and  $j$  sum over the four dots  $1 \dots 4$  of cell  $k$  and  $l$ , respectively;  $\mathbf{R}_{kl}$  denotes the vector between the centres of the cells, see Fig. 1.2(a). The cell-cell interaction term can be rewritten as

$$\begin{aligned}V &= \frac{e^2}{4\pi\epsilon} \sum_{ij} \frac{(n_i - q)(n_j - q)}{|\mathbf{R} + \mathbf{r}_j - \mathbf{r}_i|} \\ &= \frac{e^2}{4\pi\epsilon} \sum_{ij} \frac{n_i n_j - q(n_i + n_j) + q^2}{|\mathbf{R} + \mathbf{r}_{ij}|} \\ &= \frac{e^2}{4\pi\epsilon} \sum_{ij} \frac{n_i n_j - q(n_i + n_j)}{R} \left( 1 + 2 \frac{\mathbf{r}_{ij} \cdot \hat{\mathbf{R}}}{R} + \frac{r_{ij}^2}{R^2} \right)^{-1/2},\end{aligned}\tag{1.14}$$

where we dropped the constant  $q^2$  terms. Introducing  $x = 2\frac{r_{ij}\hat{\mathbf{R}}}{R}$  and  $y = \frac{r_{ij}^2}{R^2}$ , we expand  $(1 \pm x + y)^{-\frac{1}{2}}$  and keep only terms up to  $\mathcal{O}\left(\frac{a^4}{R^5}\right)$ . There are only four possibilities for two interacting cells:  $\uparrow\uparrow$ ,  $\downarrow\downarrow$ ,  $\uparrow\downarrow$ , and  $\downarrow\uparrow$ . Using this multipole expansion the four corresponding energies  $V_{kl}^{\uparrow\uparrow}$ ,  $V_{kl}^{\downarrow\downarrow}$ ,  $V_{kl}^{\uparrow\downarrow}$ , and  $V_{kl}^{\downarrow\uparrow}$  can be calculated separately. Due to the inherent geometrical symmetries of the problem and due to the vanishing dipole moments of the two cell states  $\uparrow$  and  $\downarrow$ , the obtained expressions are not too unwieldy. It turns out that  $V_{kl}^{\uparrow\downarrow} = V_{kl}^{\downarrow\uparrow}$ , but  $V_{kl}^{\uparrow\uparrow} \neq V_{kl}^{\downarrow\downarrow} \neq V_{kl}^{\uparrow\downarrow}$ . Hence we have a three-level system which we cannot hope to represent by a solely two-level Ising term  $J_{kl}S_k^zS_l^z$ . Instead we try to map to a cell-cell Hamiltonian of the form

$$H_{kl}^{cc} = J_{kl}S_k^zS_l^z + J'_{kl}(S_k^z + S_l^z). \quad (1.15)$$

For this Hamiltonian we have the energies (omitting the indices  $k$  and  $l$  for brevity)

$$E^{\uparrow\uparrow} - E^{\uparrow\downarrow} = 2J + 2J' \quad (1.16)$$

$$E^{\downarrow\downarrow} - E^{\uparrow\downarrow} = 2J - 2J' \quad (1.17)$$

which yields

$$J = \frac{1}{4} \left( E^{\uparrow\uparrow} + E^{\downarrow\downarrow} - 2E^{\uparrow\downarrow} \right) \quad (1.18)$$

$$J' = \frac{1}{4} \left( E^{\uparrow\uparrow} - E^{\downarrow\downarrow} \right). \quad (1.19)$$

By identifying  $E^{\uparrow\uparrow} = V_{kl}^{\uparrow\uparrow}$ ,  $E^{\downarrow\downarrow} = V_{kl}^{\downarrow\downarrow}$ , and so on, we obtain the final expressions

$$J = \frac{e^2}{4\pi\epsilon} \frac{1}{32} (9 - 105 \cos 4\theta) \frac{a^4}{R^5} \quad (1.20)$$

$$J' = \frac{e^2}{4\pi\epsilon} (1 - 2q) \left( \frac{3}{2} \sin 2\theta \frac{a^2}{R^3} + \frac{5}{4} \sin 2\theta \frac{a^4}{R^5} \right). \quad (1.21)$$

These results are remarkable in two ways. First,  $J$  does not depend on the compensation charge  $q$ . In turn, the Heiseberg-like interaction  $J_{kl}\sigma_k^z\sigma_l^z$  is independent of the cell net charge and represents only the quadrupole-quadrupole interaction, falling off as  $J \sim R^{-5}$ . Second, the  $J'$  term does depend on the compensation charge but vanishes for  $q = \frac{1}{2}$ , that is, charge neutral cells. For charge neutral cells QCA therefore maps directly to a normal (unmodified) Ising model. In all of our calculations we did not consider screening. Conceivably, in a real system with non-neutral cells, (static) screening might work like an effective compensating charge. Thus, a pure Ising model ( $J' = 0$ ) could very well be a reasonable description of a realistic QCA system.

We can also look at the angle dependence of the interactions. The  $J$  term is symmetric under rotations by  $90^\circ$ —which is what we expect intuitively—whereas the  $J'$  is symmetric

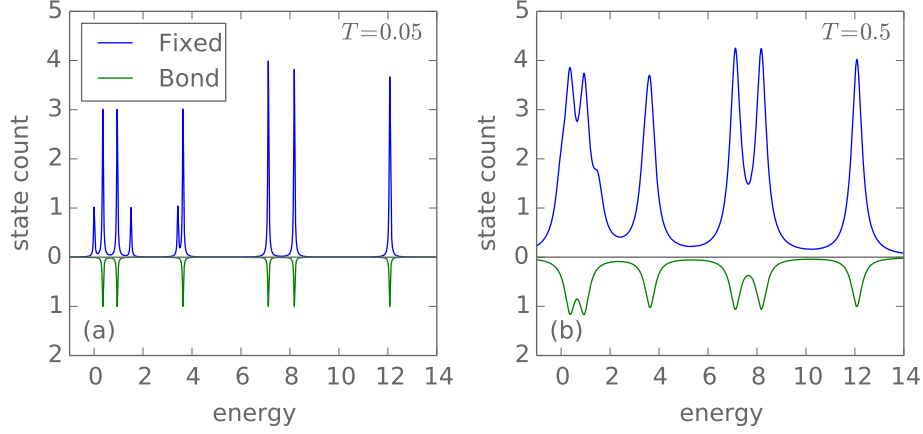


Figure 1.3: Low-energy density of states of a one cell QCA system for both the *fixed charge* and the *bond* model. The *bond* approximation does not reproduce the singlet-triplet splitting.

under rotations by  $180^\circ$ . Thus, by slowly turning on  $J'$  (by, for example, going from charge neutral to effectively negatively charged cells  $q = \frac{1}{2} \rightarrow q = 0$ ), we break the  $90^\circ$  rotational invariance. It is important to note that the  $J'$  term also breaks the symmetry between spin up and spin down cells. For  $J' \neq 0$ ,  $\uparrow\uparrow$  and  $\downarrow\downarrow$  have different energies and therefore also different polarization responses. For a practical implementation this is definitely not what we want! At the angles  $0^\circ$ ,  $90^\circ$ , and  $270^\circ$   $J'$  vanishes, so here we have a pure Heisenberg interaction even for  $q \neq 0$ . Conversely,  $J$  also vanishes at select angles, rendering the system bare any interaction for charge neutral cells, at least within the limits of our multipole expansion.

## 1.4 Validity of the approximations

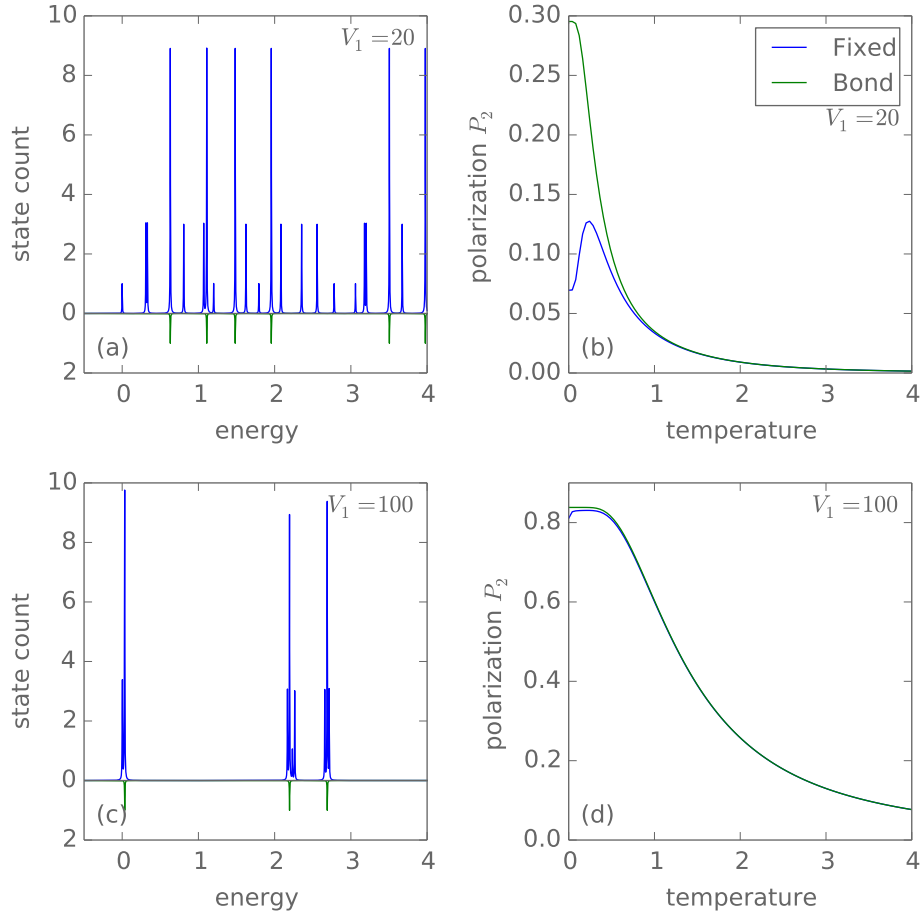


Figure 1.4: The two cell *fixed charge* and *bond* systems at  $V_1 = 20$  and  $V_1 = 100$ . (a)(c) Low-energy density of states. (b)(d) Output polarization  $P_2$  over temperature. For a small Coulomb repulsion the density of states curves look qualitatively very different (a) and the *bond* approximation does not work very well (b). At a larger Coulomb repulsion the density of states curves look much more alike (c) and the *bond* approximation works much better (d).

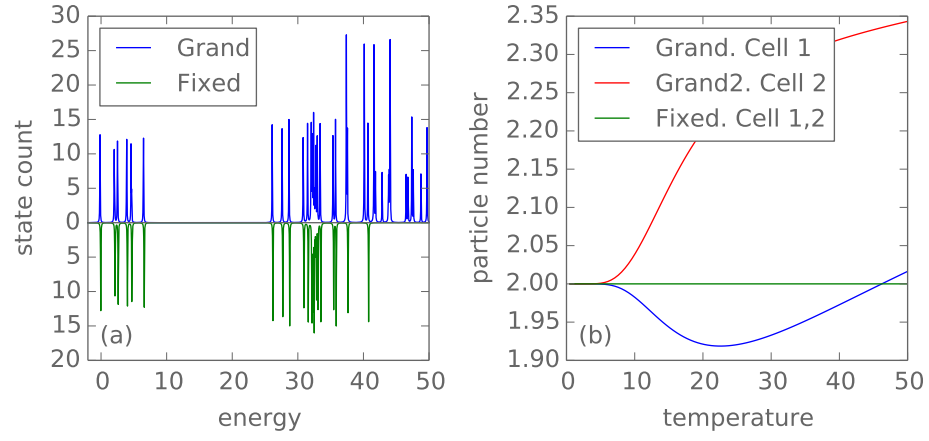


Figure 1.5: (a) Low-energy density of states of the exact *grand canonical* and the approximative *fixed charge* two cell QCA system. For small energies the curves agree perfectly (up to  $E \lesssim 35$ ). (b) Particle number per cell over temperature for the same two cell system. The curves diverge for  $T \gtrsim 5$ .

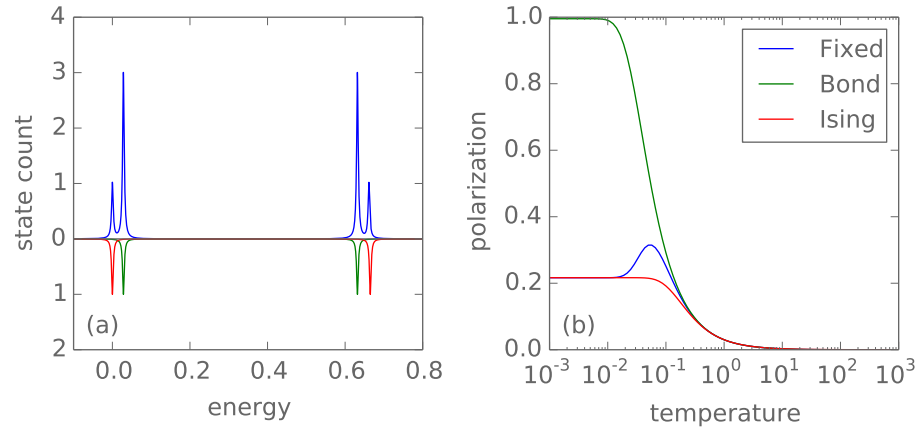


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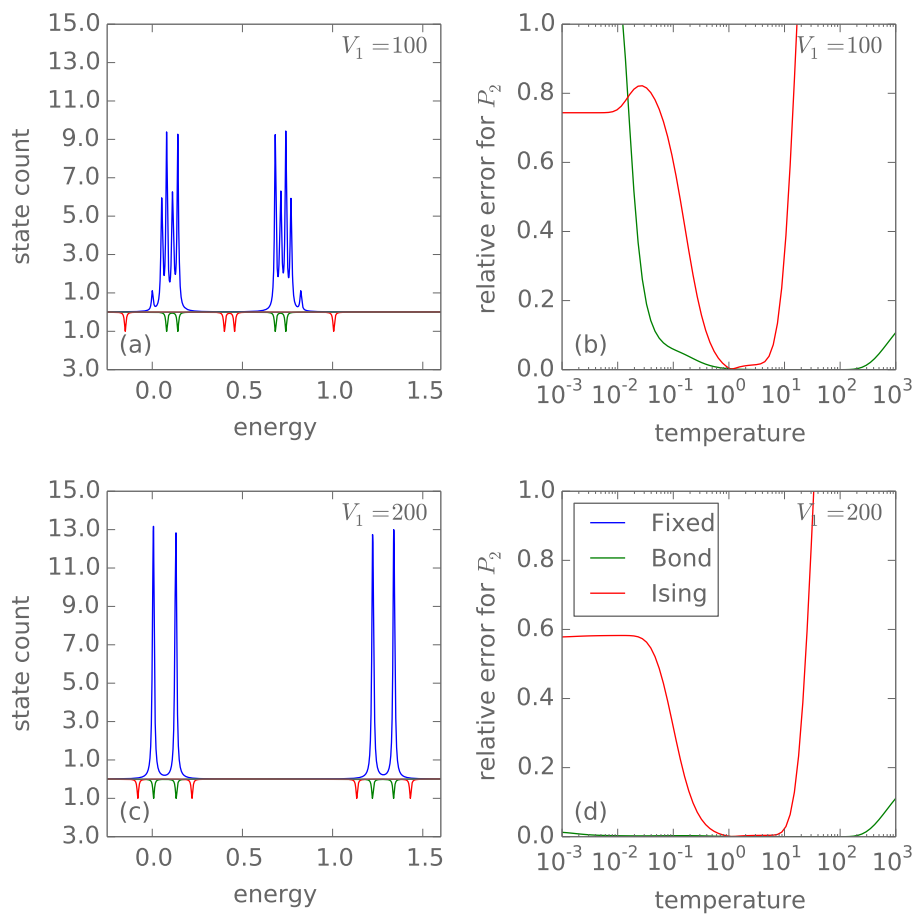


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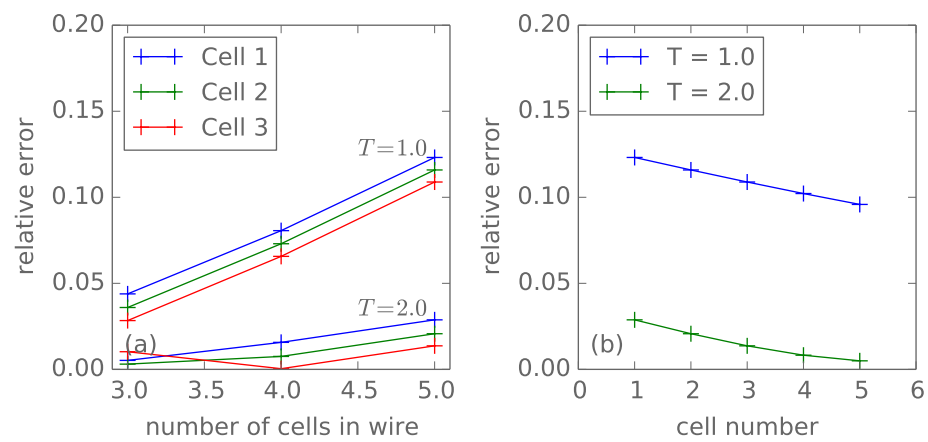


Figure 1.8: ...



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