### Exact Diagonalization Notes

Tsung-Han Lee

May 6, 2015

#### Hamiltonians:

$$H = t \sum_{i=i}^{N-1} (c_{i\sigma}^{\dagger} c_{i+1\sigma} + c_{i+1\sigma}^{\dagger} c_{i\sigma}) + \sum_{i=i}^{N} \epsilon_{i} c_{i\sigma}^{\dagger} c_{i\sigma} + \sum_{i=i}^{N} U_{i} c_{i\uparrow}^{\dagger} c_{i\uparrow} c_{i\downarrow}^{\dagger} c_{i\downarrow}$$
a) Anderson model:  $\epsilon_{i} = \epsilon \delta_{i,1}$  and  $U_{i} = U \delta_{i,1}$ . Only the first site has

- Coulomb repulsion and atomic energy.
  - b) Hubbard model:  $\epsilon_i = \epsilon$  and  $U_i = U$ .

The basis state representation is chosen in the convention  $c_{1\uparrow}^{\dagger}...c_{N\uparrow}^{\dagger}c_{1\downarrow}^{\dagger}...c_{N\downarrow}^{\dagger}|0>$ , such that the hopping term for the nearest neighbor is always positive. In the code, the basis is represented as 0 (empty), -1 (spin down), 1 (spin up), and 2 (double occupied), so the basis looks like  $|0,1,-1,2\rangle$ . The basises is build based on two symmetries: the total charge (N) and the total spin  $S_Z$ . The quantum number is represented as Q and  $S_Z$ , each Q and  $S_Z$  has subspace r. Normally the highly symmetric state has the largest subspace, e.g. for N=4, Q = 0 and  $S_Z = 0$  has r = 1...400 400 basises. Noted that in the note, the real  $S_Z$  need to be devided by 2, because the code's  $S_Z$  is set to integer for the convinience of building basis. Similarly, the real total charge Q need to be added by a total size of the chain N, since it runs from  $-N \leq Q \leq N$  for the convinience of building basis.

The full exact diagonalization is performed in each subspace (block diagonal for each Q and  $S_Z$ ), and the energies is recorded in "energiesXXXXX.dat".

#### 2 Zero Temperature Spectral Function and Green's function:

Here we discuss the implementation of zero temperature Lehmann representation for spectral function. For simplicity, we calculate the spectral function at

the first site with spin up, 
$$A_{1\uparrow}(\omega)$$
. 
$$A_{1\uparrow}(\omega) = \sum_{Q,S_Z,r} |_{diag} < 0, 0, 0 | c_{1\uparrow}| Q, S_Z, r >_{diag} |^2 \delta(\omega - E_{0,0,0} + E_{Q,S_Z,r}) + \sum_{Q,S_Z,r} |_{diag} < 0, 0, 0 | c_{1\uparrow}^{\dagger}| Q, S_Z, r >_{diag} |^2 \delta(\omega - E_{Q,S_Z,r} + E_{0,0,0})$$

Since we know that only the state  $Q = 1, S_Z = 1$  or  $Q = -1, S_Z = -1$ , for  $c_{1\uparrow}$  and  $c_{1\uparrow}^{\dagger}$  respectly, can survive for the bracket, we only need to focus on their subspace r. Noted that the subscript diag means the  $|Q,S_Z,r>_{diag}$  is the eigenvector for the energy  $E_{Q,S_Z,r}$ . Here we focus on the  $_{diag}<0,0,0|c_{1\uparrow}|1,1,r>_{diag}$  term, the other term can be calculated similarily.

The  $_{diag} < 0, 0, 0 | c_{1\uparrow} | 1, 1, r >_{diag}$  is not easy to calculate, because each eigenstate  $|Q, S_Z, r >_{diag}$  is the mix state of the original basis  $|Q, S_Z, r >_{old}$ , where  $|Q, S_Z, r >_{old}$  is the primitive basis, e.g. |0, 1, -1, 2 > for  $Q = 0, S_Z = 0$ . However, the  $_{old} < 0, 0, 0 | c_{1\uparrow} | 1, 1, r >_{old}$  is easy to calculate. So we can do a basis transformation:

$$_{old} < 0, 0, 0 \\ |c_{1\uparrow}|1, 1, r>_{diag} = \sum_{r'} \sum_{r''} {}_{diag} < 0, 0, 0 \\ |0, 0, r'>_{old} {}_{old} < 0, 0, r' \\ |c_{1\uparrow}|1, 1, r''>_{old} {}_{old} < 1, 1, r'' \\ |1, 1, r>_{diag}$$

The term  $_{diag} < 0, 0, 0|0, 0, r'>_{old}$  and  $_{old} < 1, 1, r''|1, 1, r>_{diag}$  are the orthonormal transformation matrices, which is stored in the code variable "states". Thus, the braket can be easily evaluate using the simple matrix multiplication. We first evaluate the matrix  $_{old} < 0, 0, r'|c_{1\uparrow}|1, 1, r''>_{old}$  by simple index look up, then multiply it by transform matrix.

$$_{diag} < 0, 0, 0 | c_{1\uparrow} | 1, 1, r >_{diag(1 \times M)} = U_{0,0(1 \times L)}^{T} \cdot _{old} < 0, 0, r' | c_{1\uparrow} | 1, 1, r'' >_{old(L \times M)} \cdot U_{1,1(M \times M)}$$

The dimension for  $U_{0,0}^T$  is  $1\times L$ , where L is the size of subspace  $Q=0, S_Z=0$ . The dimension for  $U_{1,1}$  is  $M\times M$ , where M is the size of subspace  $Q=1, S_Z=1$ . After obtained  $_{diag}<0,0,0|c_{1\uparrow}|1,1,r>_{diag}$ , the square of it is called spectral weight, which is printed out in "spectralWeightsXXXXX.dat". Pluging in the energies  $E_{1,1,r}$  and the corresponding spectralweight  $|_{diag}<0,0,0|c_{1\uparrow}|1,1,r>_{diag}|^2$  into  $A_{1\uparrow}(\omega)$  we obtain the electron part of the spectral function. The hole part  $|_{diag}<0,0,0|c_{1\uparrow}|-1,-1,r>_{diag}|^2$  can be calculated similarily. We can check that:

$$\sum_{r} (|_{diag} < 0, 0, 0|c_{1\uparrow}| - 1, -1, r >_{diag} |^{2} + |_{diag} < 0, 0, 0|c_{1\uparrow}|1, 1, r >_{diag} |^{2}) = 1$$

the total spectralweight is 1. Finally, the spectralfunction is outputed as "spectralFunctionXXXXXX.dat". The Green's function can be evaluated by the Kramer-Kronig relation:

$$G_{1\uparrow}(\omega) = \int_{-\infty}^{\infty} d\omega' \frac{A(\omega')}{\omega - \omega'}$$
 and the output is "greenFunctionXXXXX.dat".

### 3 Examples:

# Hubbard at half filling

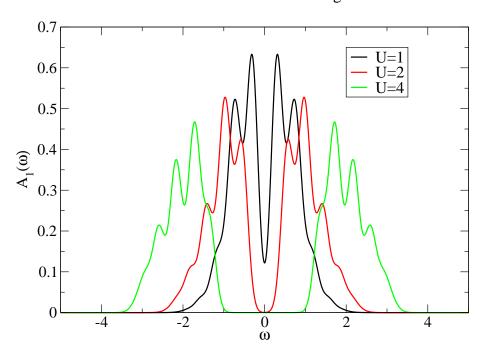


Figure 1:  $\mu=-U/2,\,t=-0.5$  , and broadening is 0.2.

# Anderson at half filling

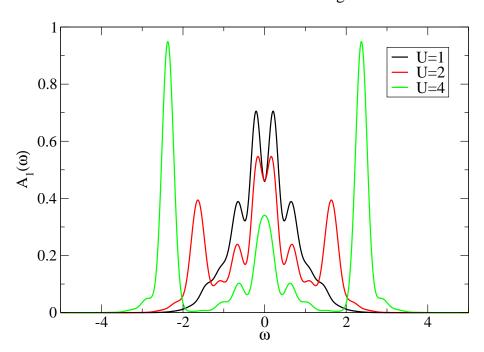


Figure 2:  $\mu=-U/2,\,t=-0.5$  , and broadening is 0.2.