

Following S. Datta, Lessons from Nanoelectronics B

Sec. 19.1.1 of Datta, Lessons...B

- see Lessons from Nanoelectronics: B, Sec. 17.4.1

```
In [ ]: import numpy as np
import matplotlib.pyplot as plt
from random import randint

from fuNEGF.models import LinearChain
```

```
In [ ]: N = 100 # keep even ; the number of atoms in the channel; the (first, last) atom i
eps_0 = 0 # eV; the on-site energy
t = 1.0 # eV; the hopping parameter
a = 1.0 # Angstrom; the lattice constant

N = 2 * (N // 2) # make sure N is even
```

Model definition

1. Hamiltonian

1D chain with nearest-neighbor hopping:

$$\hat{H}_{ij} = \begin{cases} \epsilon_0, & \text{if } i = j \\ t, & \text{if } i \neq j \end{cases}$$

or in matrix form

$$\hat{H} = \begin{bmatrix} \epsilon_0 & t & 0 & 0 & \cdots & 0 & 0 & 0 \\ t & \epsilon_0 & t & 0 & \cdots & 0 & 0 & 0 \\ 0 & t & \epsilon_0 & t & \cdots & 0 & 0 & 0 \\ 0 & 0 & t & \epsilon_0 & \cdots & 0 & 0 & 0 \\ \vdots & \vdots & \vdots & \vdots & \ddots & \vdots & \vdots & \vdots \\ 0 & 0 & 0 & 0 & \cdots & \epsilon_0 & t & 0 \\ 0 & 0 & 0 & 0 & \cdots & t & \epsilon_0 & t \\ 0 & 0 & 0 & 0 & \cdots & 0 & t & \epsilon_0 \end{bmatrix}$$

giving a dispersion relation (within an approximation -- periodic boundary conditions are actually *not* applied in the Hamiltonian above)

$$E(k) = \epsilon + 2t \cos(ka), \quad k \in \{0, 1, \dots, N-1\} \cdot 2\pi/(Na)$$

2. Self-energy

Self-energies for contact 1 and 2

$$\Sigma_1 = \begin{bmatrix} te^{ika} & 0 & 0 & \dots & 0 \\ 0 & 0 & 0 & \dots & 0 \\ 0 & 0 & 0 & \dots & 0 \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ 0 & 0 & 0 & \dots & 0 \end{bmatrix}, \quad \Sigma_2 = \begin{bmatrix} 0 & \dots & 0 & 0 & 0 \\ \vdots & \ddots & \vdots & \vdots & \vdots \\ 0 & \dots & 0 & 0 & 0 \\ 0 & \dots & 0 & 0 & 0 \\ 0 & \dots & 0 & 0 & te^{ika} \end{bmatrix}$$

with the broadening functions $\Gamma \equiv i [\Sigma - \Sigma^\dagger]$

$$\Gamma_1 = \frac{\hbar v}{a} \begin{bmatrix} 1 & 0 & 0 & \dots & 0 \\ 0 & 0 & 0 & \dots & 0 \\ 0 & 0 & 0 & \dots & 0 \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ 0 & 0 & 0 & \dots & 0 \end{bmatrix}, \quad \Gamma_2 = \frac{\hbar v}{a} \begin{bmatrix} 0 & \dots & 0 & 0 & 0 \\ \vdots & \ddots & \vdots & \vdots & \vdots \\ 0 & \dots & 0 & 0 & 0 \\ 0 & \dots & 0 & 0 & 0 \\ 0 & \dots & 0 & 0 & 1 \end{bmatrix}$$

where $v = dE/(\hbar dk) = -2at/\hbar \sin(ka)$ so that $\frac{\hbar v}{a} = -2t/\sin(ka)$

3. In-scattering

$$\Sigma_{1,2}^{\text{in}} = \Gamma_{1,2} \cdot f_{1,2}$$

where f_i is the Fermi-Dirac distribution function for contact i .

NEGF Equations

The *retarded Green's function*

$$\mathbf{G}^{\text{R}} = [\mathbf{E}\mathbf{I} - \mathbf{H} - \mathbf{\Sigma}]^{-1}$$

along with the *advanced Green's function*

$$\mathbf{G}^{\text{A}} = [\mathbf{G}^{\text{R}}]^\dagger$$

provide the *spectral function*

$$\mathbf{A} = i [\mathbf{G}^{\text{R}} - \mathbf{G}^{\text{A}}]$$

and are used to solve for the "electron occupation" Green's function ($\mathbf{G}^{\text{n}} \equiv -i\mathbf{G}^<$)

$$\mathbf{G}^{\text{n}} = \mathbf{G}^{\text{R}} \Sigma^{\text{in}} \mathbf{G}^{\text{A}}$$

which gives the *density matrix*

$$\hat{\rho} = \mathbf{G}^n / 2\pi.$$

Both, the self-energy Σ and the in-scattering term Σ^{in} are sums of the left contact and right contact, while the self-energy also contains an intrinsic term

$$\Sigma^{\text{in}} = \Sigma_1^{\text{in}} + \Sigma_2^{\text{in}}, \quad (1)$$

$$\Sigma = \Sigma_1 + \Sigma_2 + \Sigma_0 \quad (2)$$

NOTE: We use the (physically expressive) notation of S. Datta, where the self-energies and Green's functions in relation to the standard notation (on the right) are defined as

$$\Sigma \equiv \Sigma^{\text{R}}, \quad (3)$$

$$G^n \equiv -iG^<, \quad (4)$$

$$\Sigma^{\text{in}} \equiv -i\Sigma^<. \quad (5)$$

Class definition

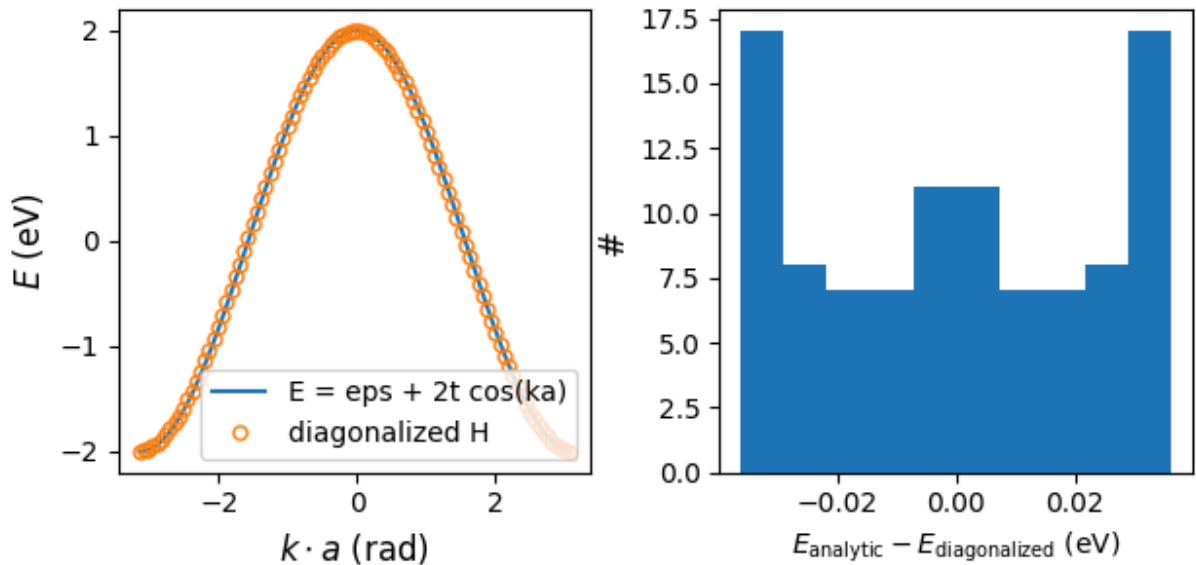
- packaged into the "fuNEGF" package; imported in the beginning

Case studies

Construct (a pure) Hamiltonian

```
In [ ]: chain = LinearChain(N, eps_0, t, a, H_impurity=None, plot_dispersion=True)
```

$t = 1.0, \text{eps}_0 = 0$



- when periodic Hamiltonian enforced, the analytical dispersion relation indeed gives a precise dependence (energy error $\sim 10^{-15}$ eV)

- the order of the diagonalized eigenvectors needs to be imposed: $\{0, 2, 4, \dots, N, N-1, \dots, 1\}$

Impurity - Quantum Resistors in Series

- a constant on-site potential on a single (or multiple) sites

$$\hat{H} = \begin{bmatrix} \ddots & \vdots & \vdots & \vdots & \ddots \\ \cdots & \varepsilon & t & 0 & \cdots \\ \cdots & t & \varepsilon + U & t & \cdots \\ \cdots & 0 & t & \varepsilon & \cdots \\ \ddots & \vdots & \vdots & \vdots & \ddots \end{bmatrix}$$

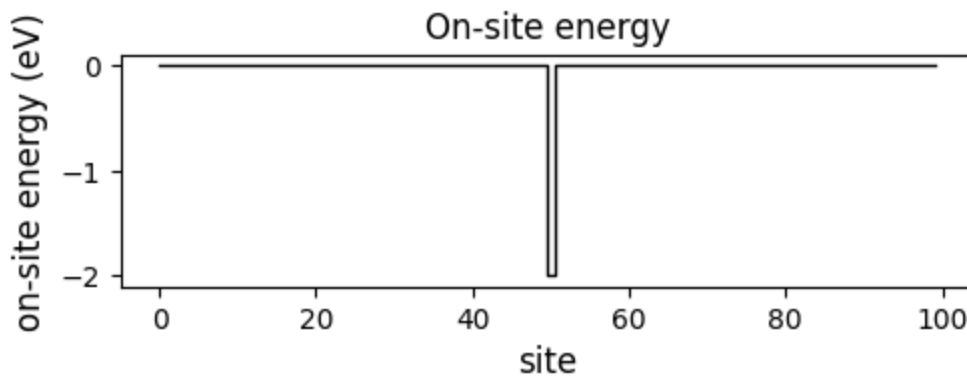
```
In [ ]: H_impurity = np.zeros((N, N), dtype=complex)

# (1) add a single impurity in the middle
H_impurity[N // 2, N // 2] += -2 * t

# (2) add two impurities symmetrically at <position_scatterer> from each end of the
# position_scatterer = 18
# H_impurity[position_scatterer, position_scatterer] += -t
# H_impurity[-position_scatterer, -position_scatterer] += -t

chain.add_H_impurity(H_impurity, plot_dispersion=False)

chain.plot_onsite_energy()
```



Plot the *Transmission function*

$$\bar{T}(E) = \text{Trace} \left[\mathbf{\Gamma}_1 \mathbf{G}^R \mathbf{\Gamma}_2 \mathbf{G}^A \right]$$

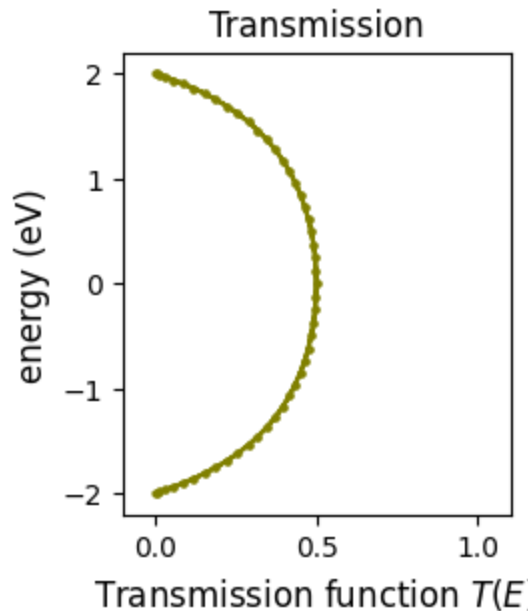
Cases:

- without impurities it is constant $T(E) = 1$
- with a single impurity of $U = -2t$, it reaches maximum of $T(E) = 0.5$

3. with two impurities of $U = -t$ each, the function looks almost the same, but with strong oscillations

```
In [ ]: chain.plot_transmission()
```

c:\Python310\lib\site-packages\matplotlib\cbook__init__.py:1335: ComplexWarning: Casting complex values to real discards the imaginary part
return np.asarray(x, float)



Phase / momentum relaxation

```
In [ ]: def plot_onsite_and_occupation(E_to_plot, D0_phase, D0_phase_momentum, N_sc):
    fig, axes = plt.subplots(2, 1, figsize=(4.5, 6), sharex=True, height_ratios=[1,
    plt.suptitle(
        r"occupation at $E = "
        + f"{E_to_plot:.2f}"
        + r"$ eV"
        + "\n"
        + r"phase relaxed with $D_0^{\mathrm{phase}} = "
        + f"{D0_phase:.2f}"
        + r"$ eV$^2$ "
        + "\n"
        + r"phase+momentum relaxed with $D_0^{\mathrm{ph,mom}} = "
        + f"{D0_phase_momentum:.2f}"
        + r"$ eV$^2$ ",
        fontsize=12,
    )

    chain.plot_onsite_energy(ax=axes[0])
    chain.plot_occupation(
        D0_phase, D0_phase_momentum, E_to_plot=E_to_plot, N_sc=N_sc, ax=axes[1]
    )

    # make x-axes common
```

```
axes[0].set_xlabel("")
plt.tight_layout()
```

```
In [ ]: E_to_plot = eps_0 + 0.5 * t
        N_sc = 70
```

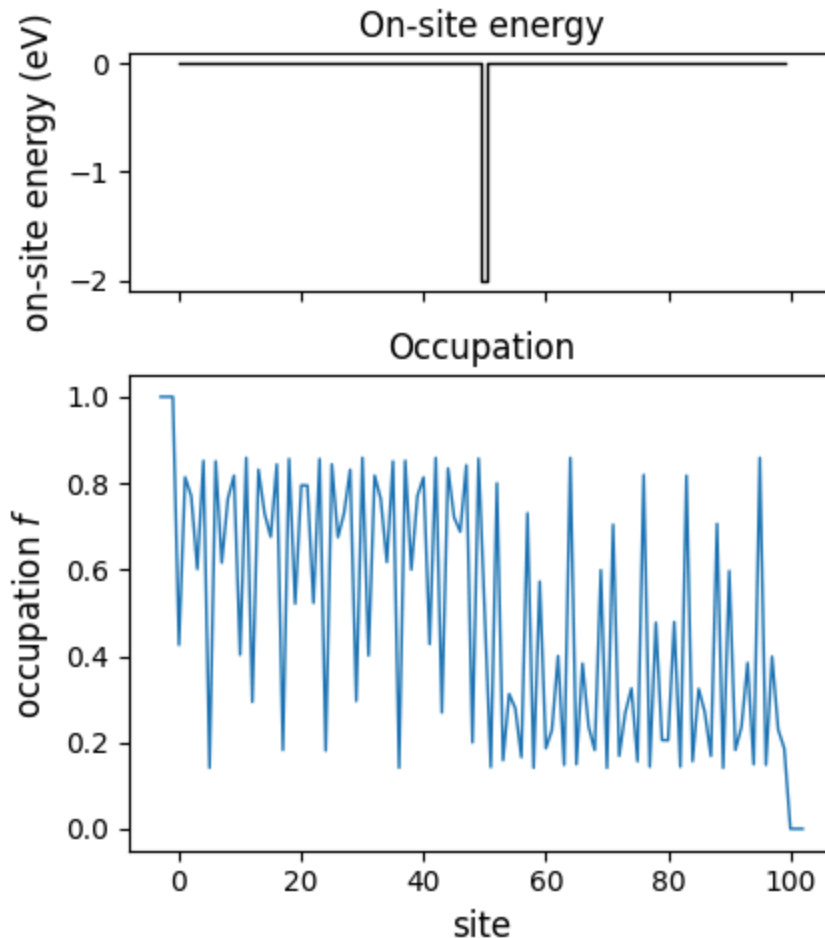
1) No relaxation

```
In [ ]: # no relaxation
        D0_phase = 0.00 * t**2
        D0_phase_momentum = 0.00 * t**2
        plot_onsite_and_occupation(E_to_plot, D0_phase, D0_phase_momentum, N_sc)
```

occupation at $E = 0.50$ eV

phase relaxed with $D_0^{\text{phase}} = 0.00$ eV²

phase+momentum relaxed with $D_0^{\text{ph, mom}} = 0.00$ eV²



2. Phase relaxation

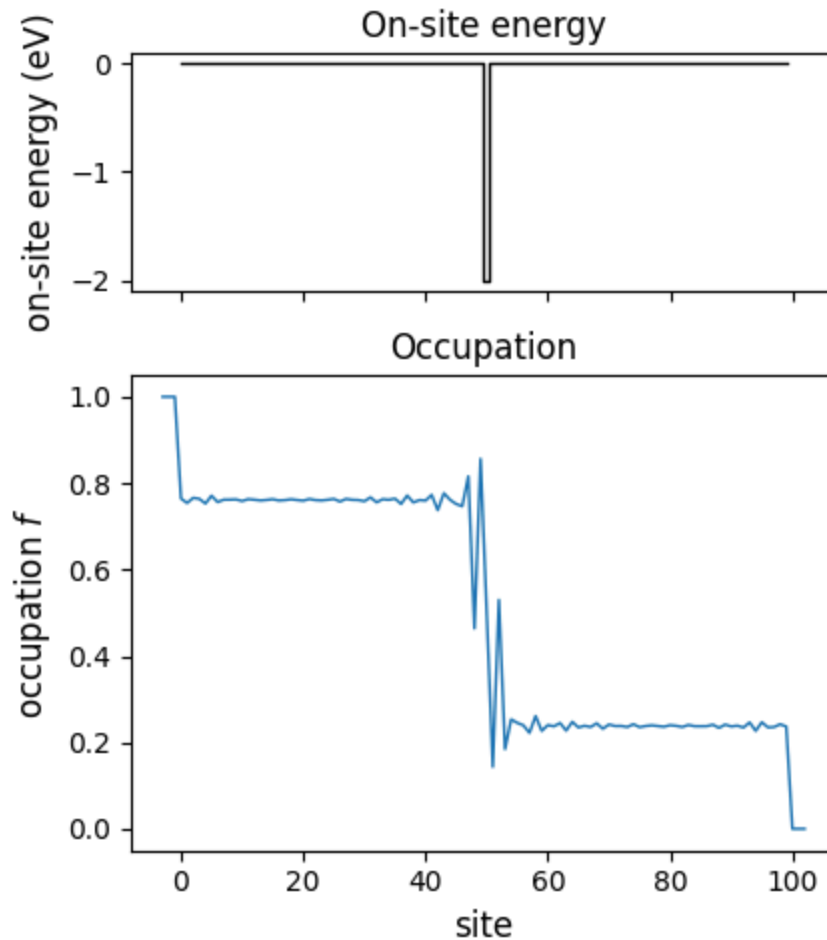
```
In [ ]: # only phase
        D0_phase = 0.09 * t**2
```

```
D0_phase_momentum = 0.00 * t**2
plot_onsite_and_occupation(E_to_plot, D0_phase, D0_phase_momentum, N_sc)
```

occupation at $E = 0.50$ eV

phase relaxed with $D_0^{\text{phase}} = 0.09$ eV²

phase+momentum relaxed with $D_0^{\text{ph, mom}} = 0.00$ eV²



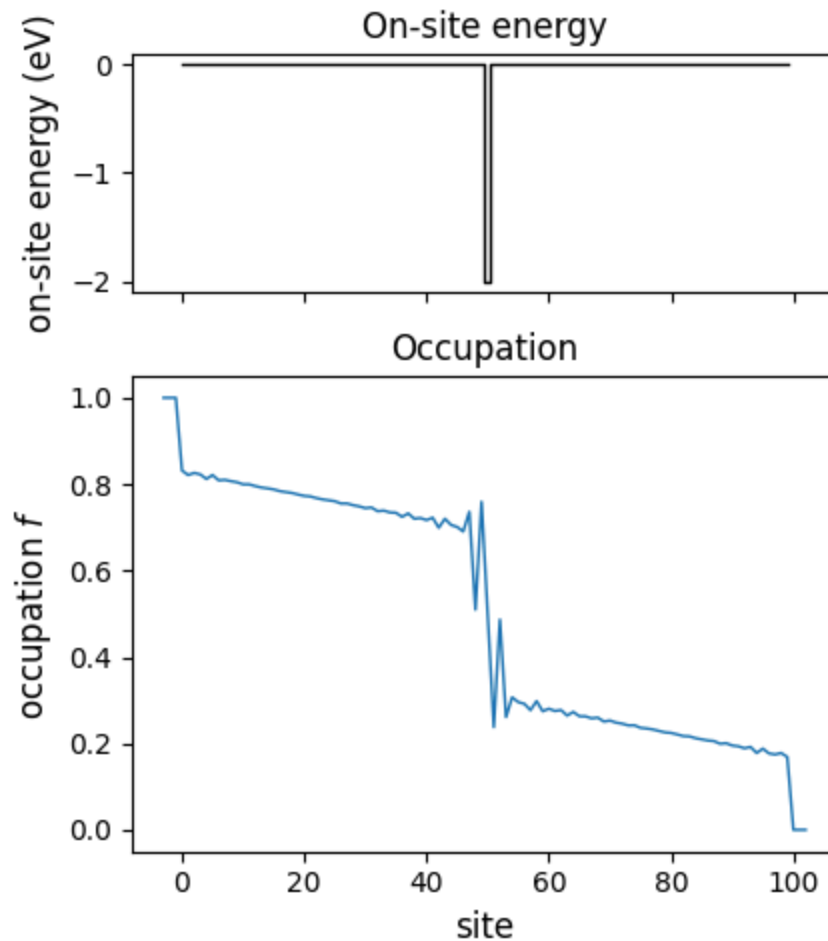
3. Phase and momentum relaxation

```
In [ ]: # phase and momentum
D0_phase = 0.09 * t**2
D0_phase_momentum = 0.03 * t**2
plot_onsite_and_occupation(E_to_plot, D0_phase, D0_phase_momentum, N_sc)
```

occupation at $E = 0.50$ eV

phase relaxed with $D_0^{\text{phase}} = 0.09$ eV²

phase+momentum relaxed with $D_0^{\text{ph, mom}} = 0.03$ eV²



- after each barrier, the potential drops (when we put $f=1$ on the left and $f=0$ on the right)
- wild oscillations in occupation can be conveniently damped by phase relaxation
- additional momentum relaxation causes a steady decrease in the potential

Multiple resistors

```
In [ ]: # RELAXATION
D0_phase = 0.12 * t**2
D0_phase_momentum = 0.03 * t**2
N_sc = 90

# IMPURITY HAMILTONIANS
H_imp_clean = np.zeros((N, N))
H_imp_single = np.zeros((N, N))
H_imp_single[N // 2, N // 2] = -2 * t
H_imp_double = np.zeros((N, N))
H_imp_double[5 * N // 12, 5 * N // 12] = -t
```



```

H_imp_double[3 * N // 4, 3 * N // 4] = -t

# multiple random impurities
H_imp_multiple = np.zeros((N, N))
N_random_impurities = randint(3, 7)
potential_ratios = [randint(1, 5) for _ in range(N_random_impurities)]
potentials = [
    potential_ratios[i] / sum(potential_ratios) * (-2 * t)
    for i in range(N_random_impurities)
]
positions = [randint(1, N - 1) for _ in range(N_random_impurities)]
for i in range(N_random_impurities):
    H_imp_multiple[positions[i], positions[i]] = potentials[i]

H_impurities = [H_imp_clean, H_imp_single, H_imp_double, H_imp_multiple]

impurity_titles = [
    "clean",
    r"single impurity $U=-2t$",
    r"double imp.  $2 \times (U=-t)$",
    r"randomimps. (total $U=-2t$)",
]

# CALCULATE AND PLOT
fig, axes = plt.subplots(
    5,
    len(H_impurities),
    figsize=(2.5 * len(H_impurities) + 2, 8),
    height_ratios=[0.9, 1, 1, 1, 1.3],
    layout="constrained",
)
fig.set_constrained_layout_pads(
    w_pad=0.35
)
# , h_pad=4./72., hspace=0./72., wspace=0./72.)
for i, H_imp in enumerate(H_impurities):
    chain = LinearChain(N, eps_0, t, a, H_impurity=H_imp, plot_dispersion=False)
    chain.plot_onsite_energy(axes[0, i])
    chain.plot_occupation(
        D0_phase=0, D0_phase_momentum=0, E_to_plot=E_to_plot, N_sc=N_sc, ax=axes[1, i]
    )
    chain.plot_occupation(
        D0_phase=D0_phase,
        D0_phase_momentum=0,
        E_to_plot=E_to_plot,
        N_sc=N_sc,
        ax=axes[2, i],
    )
    chain.plot_occupation(
        D0_phase=D0_phase,
        D0_phase_momentum=D0_phase_momentum,
        E_to_plot=E_to_plot,
        N_sc=N_sc,
        ax=axes[3, i],
    )
    chain.plot_transmission(axes[4, i])

```

```

# labels, titles, limits, etc.
axes[0, 0].set_ylabel("On-site (eV)")
axes[1, 0].set_ylabel("Occupation")
axes[2, 0].set_ylabel("Occupation")
axes[3, 0].set_ylabel("Occupation")
axes[4, 0].set_ylabel("Energy (eV)")
for j in range(1, len(H_impurities)):
    for i in range(5):
        axes[i, j].set_yticklabels([])
        axes[i, j].set_ylabel("")
        axes[i, j].tick_params(axis="y", direction="in")
# set all titles empty string
for j in range(len(H_impurities)):
    axes[4, j].set_xlabel("Transmission")
    for i in range(5):
        axes[i, j].set_title("")
    axes[0, j].set_title(impurity_titles[j], fontsize=14)
    for i in range(3):
        axes[i, j].set_xticklabels([])
        axes[i, j].set_xlabel("")
        # ticks inside
        axes[i, j].tick_params(axis="x", direction="in")
text_left = N // 2
text_top = 0.8
axes[1, 0].text(
    text_left,
    text_top,
    "no relaxation" + r"$\rightarrow$",
    fontsize=12,
    rotation=0,
    ha="center",
    va="center",
)
axes[2, 0].text(
    text_left,
    text_top,
    "phase\nrelaxed" + r"$\rightarrow$",
    fontsize=12,
    rotation=0,
    ha="center",
    va="center",
)
axes[3, 0].text(
    text_left,
    text_top,
    "phase+momentum\nrelaxed" + r"$\rightarrow$",
    fontsize=12,
    rotation=0,
    ha="center",
    va="center",
)
# all on-site energy plots have the same y-axis
margin = 0.08
y_max = max([axes[0, i].get_ylim()[1] for i in range(len(H_impurities))])
y_min = min([axes[0, i].get_ylim()[0] for i in range(len(H_impurities))])
for j in range(len(H_impurities)):

```

```
axes[0, j].set_ylim(
    [y_min - margin * (y_max - y_min), y_max + margin * (y_max - y_min)]
)
```

