Appendix D

The PYTHTB Package

Numerous examples and exercises in this book make use of the PYTHTB software package. This package, which is written in the PYTHON programming language, is designed to allow the user to construct and solve tight-binding (TB) models of the electronic structure of finite clusters and of systems that display periodicity in one or more dimensions, such as polymer chains, ribbons, slabs, and 3D crystals. It is also designed to provide convenient features for computing geometric or topological properties such as Berry phases, Berry curvatures, and Chern numbers. The code was developed beginning in 2010 and is maintained principally by Sinisa Coh and David Vanderbilt, although significant contributions have been provided by others (see http://www.physics.rutgers.edu/pythtb/about.html#History).

The examples and exercises used in this book assume the use of version 1.7.2 of PYTHTB, which is the current version at the time of this writing. PYTHTB version 1.7.2 is designed for compatibility with both PYTHON2.7 and PYTHON3.X (PYTHON2.6 and below are not recommended). If you already have one of these PYTHON versions and PIP installed on a Linux system, you should be able to install the latest version of PYTHTB just by issuing the command

```
pip install pythtb --upgrade
```

at a command prompt if you have root permission, or

```
pip install pythtb --upgrade --user
```

to install it in your home folder if you do not. If you wish to install the package without PIP or otherwise need further help with installation, see the package web pages at http://www.physics.rutgers.edu/pythtb. You will also need access to the standard NUMPY (numerical) and MATPLOTLIB (plotting) PYTHON packages; install these too if they are not already present.

The example programs, together with their printed and plotted outputs, are available at http://www.cambridge.org/9781107157651#resources. This is referred to as "the website" in the remainder of this appendix. All efforts will be made to keep future upgrades of PYTHTB backwardcompatible with the example programs provided here, but in case of doubt you can obtain version 1.7.2 specifically by issuing the command

```
pip install pythtb==1.7.2
```

or download and install PYTHTB 1.7.2 from the "Installation" link on the website. Each program starts with the header lines

```
#!/usr/bin/env python
from __future__ import print_function # python3 style print
```

The first identifies the script as a PYTHON program. The second is ignored when running under PYTHON3, but is needed for compatibility with PYTHON2 so that print is interpreted as a function, as is standard in PYTHON3. These two lines are typically followed by some comment lines (starting with '#') that identify the program.

The next few lines import the PYTHTB module and any other modules needed for execution of the script. Typical entries include

```
from pythtb import *
import numpy as np
import matplotlib.pyplot as plt
```

The first line imports the PYTHTB module. All of the example programs also make use of the NUMPY mathematical subroutine library, which is imported on the second line in such a way that NUMPY 'func' will be called as np.func. This line is actually unnecessary because the import is done automatically when PYTHTB is imported, but it may be included to make the code more readable; this has been done in the first example in Appendix D.1, but not in subsequent ones. Finally, the import of matplotlib.pyplot on the third line provides a MATLIB-like plotting environment for those example programs that produce plotted output.¹

PYTHTB defines two *classes*: tb_model and wf_array.² A command such as my_model=tb_model (*arguments*) defines an *instance* of the tb_model class, an object that carries all the relevant information about the TB model. The (*arguments*) given initially define some general characteristics of the model (dimensionality, whether it is a spinor model, and so on), but then parameters such as on-site energies and hopping amplitudes are set subsequently by making

¹ The precise appearance of a plot may depend on the installed version of MATPLOTLIB (version 1.5.3 was used here) and on any matplotlibre configuration files that may be present.

It also defines a third, w90, which provides an interface to the WANNIER90 code package (Mostofi et al., 2008, 2014). We shall not make use of this class here.

use of the set_onsite and add_hop methods. For periodic models, the next step is usually to set up a mesh of k-points on which the model is to be solved, with the solution typically done by calling the solve_all method. Alternatively, an array object can be defined as an instance of the wf_array class and solved by the solve_on_grid method of this class; this is especially useful when computing Berry phases and curvatures, since the wf_array class provides methods specifically for such purposes. Finally, the results are printed or plotted for inspection.

All this is best illustrated by browsing the example programs and their outputs in the sections that follow.

D.1 Water Molecule

Program h2o.py computes the eigenstates of a TB model for the water molecule as described in Section 2.2.2. The first few lines define the molecular geometry (bond length and angle) and the TB parameters (s and p site energies and s-s and p-p hoppings). The lat= line specifies three basis vectors (here Cartesian), in terms of which the atom coordinates will be given, and the orb line specifies those coordinates. The next few lines define the TB model my_model, while my_model.display prints a summary of it and my_model.solve_all solves for the eigenvalues and eigenvectors. Since the model is described by a real Hamiltonian, the imaginary parts of the eigenvectors are zero to numerical accuracy and are discarded. The last few lines print the result; np.set_printoptions is a numpy function that sets the format for subsequent print statement, and the eigenvalues and eigenvectors are printed at the end.

h2o.py

```
#!/usr/bin/env python
from __future__ import print_function # python3 style print
# -----
# Tight-binding model for H2O molecule
# ------
# import the pythtb module
from pythtb import *
import numpy as np
# geometry: bond length and half bond-angle
b=1.0; angle=54.0*np.pi/180
# site energies [O(s), O(p), H(s)]
eos=-1.5; eop=-1.2; eh=-1.0
# hoppings [O(s)-H(s), O(p)-H(s)]
ts=-0.4; tp=-0.3
```

```
# define frame for defining vectors: 3D Cartesian
lat=[[1.0,0.0,0.0],[0.0,1.0,0.0],[0.0,0.0,1.0]]
\# define coordinates of orbitals: O(s,px,py,pz) ; H(s) ; H(s)
orb=[ [0.,0.,0.], [0.,0.,0.], [0.,0.,0.], [0.,0.,0.],
      [b*np.cos(angle), b*np.sin(angle),0.],
      [b*np.cos(angle),-b*np.sin(angle),0.] ]
# define model
my model=tbmodel(0,3,lat,orb)
my model.set onsite([eos,eop,eop,eop,eh,eh])
my model.set hop(ts,0,4)
my model.set hop(ts,0,5)
my model.set hop(tp*np.cos(angle),1,4)
my_model.set_hop(tp*np.cos(angle),1,5)
my_model.set_hop(tp*np.sin(angle),2,4)
my model.set hop(-tp*np.sin(angle),2,5)
# print model
my_model.display()
# solve model
(eval, evec) = my_model.solve_all(eig_vectors=True)
# the model is real, so OK to discard imaginary parts of eigenvectors
evec=evec.real
# optional: choose overall sign of evec according to some specified rule
# (here, we make the average oxygen p component positive)
for i in range(len(eval)):
 if sum(evec[i,1:4]) < 0:
    evec[i,:] = - evec[i,:]
# print results, setting numpy to format floats as xx.xxx
np.set printoptions(formatter={'float': '{: 6.3f}'.format})
# print eigenvalues and real parts of eigenvectors, one to a line
print(" n eigval
                     eiqvec")
for n in range(6):
    print(" %2i %7.3f " % (n,eval[n]), evec[n,:])
```

D.2 Benzene Molecule

Program benzene.py, discussed in Section 2.2.2, computes the eigenstates for a TB model of the p_{π} manifold (i.e., p orbitals oriented normal to the plane of the molecule) for benzene (CH₆). The structure is much like that of h2o.py in the preceding section. See also Ex. 2.5.

benzene.py

```
r=1.2
                                  # distance of atoms from center
orb=np.zeros((6,2),dtype=float)  # initialize array for orbital positions
for i in range(6):
                                  # define coordinates of orbitals
  angle=i*np.pi/3.0
  orb[i,:] = [r*np.cos(angle), r*np.sin(angle)]
# set site energy and hopping amplitude, respectively
ep = -0.4
t = -0.25
# define model
my model=tbmodel(0,2,lat,orb)
my model.set onsite([ep,ep,ep,ep,ep,ep])
my model.set hop(t,0,1)
my_model.set_hop(t,1,2)
my_model.set_hop(t,2,3)
my_model.set_hop(t,3,4)
my_model.set_hop(t,4,5)
my model.set hop(t,5,0)
# print model
my model.display()
# solve model and print results
(eval, evec) = my model.solve all(eig vectors=True)
\mbox{\#} print results, setting numpy to format floats as \mbox{xx.xxx}
np.set printoptions(formatter={'float': '{: 6.3f}'.format})
# print eigenvalues and real parts of eigenvectors, one to a line
print(" n eigval eigvec")
for n in range(6):
    print(" %2i %7.3f " % (n,eval[n]), evec[n,:].real)
```

D.3 bcc Li Crystal

Program li.py computes the band structure of bcc Li based on a TB model with one s orbital per atomic site, as introduced in Section 2.2.4. Now the vectors lat have to be a set of primitive real-space lattice vectors that define the periodicity of the crystal, and subsequent coordinates must be given in terms of these. In particular, the neighboring orbitals specified by the calls to set_hop are given in these lattice coordinates, with the translation to Cartesian coordinates given in comments at the end of each line. Orbital locations also need to be defined in lattice coordinates, but there is only one atom in the unit cell and we place it at the origin. (The value of the lattice constant has no effect on the results to be computed here, so it is set to unity for convenience.)

The next few lines after my_model.display() specify a list of three special k points, or "nodes," that define the path along which the band structure will be computed and plotted. The path and label variables are lists of the node coordinates and their labels, respectively, and the k_path method takes path as input and constructs a list of k points tracing a path along straight-line segments between these nodes. The returned variables k_vec and k_dist are arrays containing the coordinates of the k points and corresponding accumulated distance

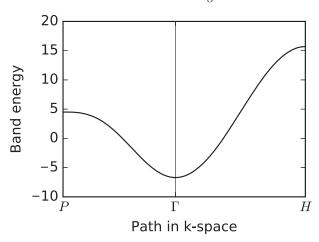


Figure D.1 Band structure of bcc Li as generated by the li.py program.

along the path for each point, while k_node gives the accumulated distance just for the nodes. These are used in the calls to the PYPLOT plotting routines to generate the band-structure plot which is output as li_band.pdf, shown in Fig. D.1. This reproduces the results shown in Fig. 2.4 in Section 2.2.3.

Note that the node positions in path must be specified in *reciprocal lattice coordinates*, as linear combinations of the primitive reciprocal lattice vectors that are dual to the real-space ones. (The same convention is used for the output array k_vec.) It is the responsibility of the user to specify path in this way, but the "k_path report" generated by the k_path function is useful for checking that this has been done correctly. For this program, for example, see the lines between k_path report begin and k_path report end in the printed output on the website.

li.py

```
#!/usr/bin/env pvthon
from __future__ import print_function # python3 style print
# 3D model of Li on bcc lattice, with s orbitals only
from pythtb import * # import TB model class
import matplotlib.pyplot as plt
# define lattice vectors
lat=[[-0.5, 0.5, 0.5],[ 0.5,-0.5, 0.5],[ 0.5, 0.5,-0.5]]
# define coordinates of orbitals
orb=[[0.0,0.0,0.0]]
# make 3D model
my model=tb model(3,3,lat,orb)
# set model parameters
\# lattice parameter implicitly set to a=1
Es= 4.5 # site energy
t = -1.4
         # hopping parameter
```

```
# set on-site energy
my model.set onsite([Es])
# set hoppings along four unique bonds
# note that neighboring cell must be specified in lattice coordinates
# (the corresponding Cartesian coords are given for reference)
my_model.set_hop(t, 0, 0, [1,0,0])  # [-0.5, 0.5, 0.5] cartesian
my_model.set_hop(t, 0, 0, [0,1,0])  # [ 0.5,-0.5, 0.5] cartesian
my_model.set_hop(t, 0, 0, [0,0,1])  # [ 0.5, 0.5,-0.5] cartesian
my_model.set_hop(t, 0, 0, [1,1,1])  # [ 0.5, 0.5, 0.5] cartesian
# print tight-binding model
my model.display()
# generate k-point path and labels
# again, specified in reciprocal lattice coordinates
k P = [0.25, 0.25, 0.25]
                                          # [ 0.5, 0.5, 0.5] cartesian
k_{Gamma} = [ 0.0, 0.0, 0.0]
                                           # [ 0.0, 0.0, 0.0] cartesian
k H = [-0.5, 0.5, 0.5]
                                          # [ 1.0, 0.0, 0.0] cartesian
path=[k P,k Gamma,k H]
label=(r'$P$',r'$\Gamma $',r'$H$')
(k_vec,k_dist,k_node) = my_model.k_path(path,101)
print('starting calculation')
print('----')
print('Calculating bands...')
# solve for eigenenergies of Hamiltonian on
# the set of k-points from above
evals=my model.solve all(k vec)
# plotting of band structure
print('Plotting band structure...')
# first make a figure object
fig, ax = plt.subplots(figsize=(4.,3.))
# specify horizontal axis details
ax.set_xlim([0,k_node[-1]])
ax.set_xticks(k_node)
ax.set_xticklabels(label)
for n in range(len(k node)):
  ax.axvline(x=k node[n], linewidth=0.5, color='k')
# plot bands
ax.plot(k dist,evals[0],color='k')
# put title
ax.set_xlabel("Path in k-space")
ax.set ylabel("Band energy")
# make a PDF figure of a plot
fig.tight_layout()
fig.savefig("li bsr.pdf")
print('Done.\n')
```

D.4 Alternating Site Model

Program chain_alt.py computes the band structure of the 1D alternating site chain model of Fig. 2.6 in Section 2.2.4. The lattice constant is set to unity for convenience. The orbitals at x = 0 and x = 1/2 are assigned site energies Δ and

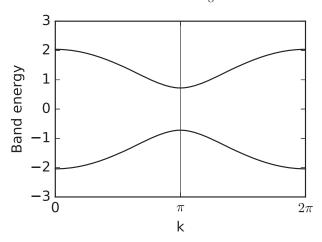


Figure D.2 Band structure resulting from the 1D alternating chain model of Fig. 2.6 as computed by chain_alt.py.

 $-\Delta$, respectively, and the alternating hopping strengths are $t + \delta t$ and $t - \delta t$. This time a subroutine set_model is defined to specify the model; this will be convenient later when looping over parameters of the model. The resulting plot appears in Fig. D.2.

chain_alt.py

```
#!/usr/bin/env python
from __future__ import print_function # python3 style print
# Chain with alternating site energies and hoppings
from pythtb import *
import matplotlib.pyplot as plt
# define function to set up model for a given paramter set
def set model(t,del t,Delta):
  # 1D model with two orbitals per cell
 lat=[[1.0]]
 orb=[[0.0],[0.5]]
 my_model=tbmodel(1,1,lat,orb)
  # alternating site energies (let average be zero)
 my_model.set_onsite([Delta,-Delta])
  # alternating hopping strengths
 my model.add hop(t+del t, 0, 1, [0])
 my model.add hop(t-del t, 1, 0, [1])
  return my model
# set reference hopping strength to unity to set energy scale
t = -1.0
# set alternation strengths
del t=-0.3
             # bond strength alternation
Delta= 0.4
             # site energy alternation
# set up the model
my model=set model(t,del t,Delta)
```

```
# construct the k-path
(k_vec,k_dist,k_node) = my_model.k_path('full',121)
k lab=(r'0',r'$\pi$',r'$2\pi$')
# solve for eigenvalues at each point on the path
evals=my model.solve all(k vec)
# set up the figure and specify details
fig, ax = plt.subplots(figsize=(4.,3.))
ax.set xlim([0,k node[-1]])
ax.set xticks(k node)
ax.set xticklabels(k lab)
ax.axvline(x=k node[1],linewidth=0.5, color='k')
ax.set_xlabel("k")
ax.set ylabel("Band energy")
# plot first and second bands
ax.plot(k_dist,evals[0],color='k')
ax.plot(k_dist,evals[1],color='k')
# save figure as a PDF
fig.tight layout()
fig.savefig("chain alt.pdf")
```

D.5 Graphene

Program graphene.py computes the band structure for the model of the π orbitals of graphene discussed in Section 2.2.4 and illustrated in Fig. 2.7(a). The conventional Wigner–Seitz Brillouin zone (BZ) is shown in Fig. 2.7(b) with high-symmetry points labeled; the my_model.k_path line of the program specifies a path from Γ to K to M and back to Γ (see the "k_path report" in the printed output available on the website). The resulting band structure plot appears as Fig. 2.8 in Section 2.2.4.

graphene.py

```
#!/usr/bin/env python
from __future__ import print_function # python3 style print
# Simple model of pi manifold of graphene

from pythtb import * # import TB model class
import matplotlib.pyplot as plt

# define lattice vectors
lat=[[1.0,0.0],[0.5,np.sqrt(3.0)/2.0]]
# define coordinates of orbitals
orb=[[1./3.,1./3.],[2./3.,2./3.]]

# make 2D tight-binding graphene model
my_model=tb_model(2,2,lat,orb)

# set model parameters
delta=0.0
t=-1.0
```

```
my model.set onsite([-delta,delta])
my_model.set_hop(t, 0, 1, [ 0, 0])
my_model.set_hop(t, 1, 0, [ 1, 0])
my model.set hop(t, 1, 0, [0, 1])
# print out model details
my model.display()
# list of k-point nodes and their labels defining the path for the
  band structure plot
path=[[0.,0.],[2./3.,1./3.],[.5,.5],[0.,0.]]
label=(r'$\Gamma $',r'$K$', r'$M$', r'$\Gamma $')
# construct the k-path
nk=121
(k vec, k dist, k node) = my model.k path(path, nk)
# solve for eigenvalues at each point on the path
evals=my model.solve all(k vec)
# generate band structure plot
fig, ax = plt.subplots(figsize=(4.,3.))
# specify horizontal axis details
ax.set xlim([0,k node[-1]])
ax.set_ylim([-3.4,3.4])
ax.set_xticks(k_ node)
ax.set xticklabels(label)
# add vertical lines at node positions
for n in range(len(k node)):
 ax.axvline(x=k node[n],linewidth=0.5, color='k')
# put titles
ax.set xlabel("Path in k-space")
ax.set ylabel("Band energy")
# plot first and second bands
ax.plot(k dist,evals[0],color='k')
ax.plot(k_dist,evals[1],color='k')
# save figure as a PDF
fig.tight_layout()
fig.savefig("graphene.pdf")
```

D.6 Trimer Molecule

Program trimer.py computes the eigenfunctions of the trimer molecule discussed in Section 3.2.4. In addition, it produces plots of Berry phases and Berry curvatures of the ground state resulting from variation of the parameters.

A set_model subroutine has been defined so as to return an instance of the model for a given set of the parameters defined in Eq. (3.43), and another subroutine get_evecs calls set_model and returns the computed eigenvectors. Predefining these functions simplifies the process of looping over the values of φ and α as needed later. This time one of the hoppings is generally complex, so TR symmetry is broken and the eigenvectors are generally complex as well.

The plotted output of this program appears as Fig. 3.11 in the main text. For pedagogical purposes the Berry phase is computed in several ways. First, for a single given value of α , the program computes the discrete Berry phase explicitly according to Eq. (3.1) as φ runs from 0 to 2π on a dense mesh (increments of $\pi/30$) at a fixed value of α . The product of the inner products appearing in Eq. (3.1) is accumulated in the variable prod and the Berry phase is computed explicitly by taking the imaginary part of the log using the NUMPY function angle.

Second, on a coarse mesh of α values (in increments of $\pi/6$), the eigenvectors on a dense mesh of φ values are stored into a 1D PYTHTB wave-function array object evec_array that is initialized as an instance of the PYTHTB wf_array class. The Berry phase is computed by the berry_phase method associated with this class, and the results are plotted as the black dots in Fig. 3.11(a). The same procedure is repeated to compute the Berry phases in the α direction on a coarse mesh of φ values and plot them as the dots in Fig. 3.11(b).

Next, a 2D wf_array object (also named evec_array) is filled with eigenvectors on a fine mesh in both the φ and α directions. The berry_phase method is then used to compute the Berry phases first in the φ direction and then in the α direction, and these (together with values shifted by integer multiples of 2π) are plotted as the continuous lines in Fig. 3.11(a–b). By default, the berry_phase procedure enforces a smooth evolution of the Berry phase, so we do not see the same kind of discontinuities that appeared when using the previous method.

Finally, the berry_flux method is applied to the same 2D evec_array array to compute the Berry flux through each of the 60×60 plaquettes, and the results are divided by parametric area to obtain the Berry curvature and used to generate the contour plot in Fig. 3.11(c). The PYPLOT subplots method is used to combine the three panels into a single PDF output file.

trimer.py

```
orb=np.zeros((3,2),dtype=float)
 orb[0,:]=[0.,1.] # orbital at top vertex
orb[1,:]=[-sqr32,-0.5] # orbital at lower left
orb[2,:]=[ sqr32,-0.5] # orbital at lower right
 # compute hoppings [t 01, t 12, t 20]
 # s is distortion amplitude; phi is "pseudorotation angle"
 tpio3=2.0*np.pi/3.0
 t=[t0+s*np.cos(phi), t0+s*np.cos(phi-tpio3), t0+s*np.cos(phi-2.0*tpio3)]
 # alpha is fraction of flux quantum passing through the triangle
 # magnetic flux correction, attached to third bond
 t[2] = t[2] *np.exp((1.j) *alpha)
 # set up model (leave site energies at zero)
 my_model=tbmodel(0,2,lat,orb)
 my_model.set_hop(t[0],0,1)
 my_model.set_hop(t[1],1,2)
my_model.set_hop(t[2],2,0)
 return(my_model)
# ------
# define function to return eigenvectors for given (t0,s,phi,alpha)
def get evecs(t0,s,phi,alpha):
 my model=set model(t0,s,phi,alpha)
 (eval, evec) = my_model.solve_all(eig_vectors=True)
 return(evec)
                # evec[bands,orbitals]
# begin regular execution
# -----
# for the purposes of this problem we keep t0 and s fixed
t \cdot 0 = -1 \cdot 0
s = -0.4
ref model=set model(t0,s,0.,1.) # reference with phi=alpha=0
ref model.display()
# define two pi
twopi=2.*np.pi
# -----
# compute Berry phase for phi loop explicitly at alpha=pi/3
# -----
alpha=np.pi/3.
n phi=60
psi=np.zeros((n_phi,3),dtype=complex)
                                    # initialize wavefunction array
for i in range (n phi):
 phi=float(i)*twopi/float(n phi)
                                     # 60 equal intervals
 psi[i] = get evecs(t0,s,phi,alpha)[0]
                                    # psi[i] is short for psi[i,:]
                                       # final [0] picks out band 0
prod=1.+0.j
for i in range(1, n phi):
 prod=prod*np.vdot(psi[i-1],psi[i]) # <psi 0|psi 1>...<psi 58|psi 59>
prod=prod*np.vdot(psi[-1],psi[0]) # include <psi_59|psi_0>
berry=-np.angle(prod)
                                  # compute Berry phase
print("Explicitly computed phi Berry phase at alpha=pi/3 is %6.3f"% berry)
# compute Berry phases for phi loops for several alpha values
# using pythtb wf array() method
```

```
alphas=np.linspace(0.,twopi,13)  # 0 to 2pi in 12 increments
berry_phi=np.zeros_like(alphas)  # same shape and type array (empty)
print("\nBerry phases for phi loops versus alpha")
for j,alpha in enumerate(alphas):
  # let phi range from 0 to 2pi in equally spaced steps
  n phi=61
 phit=np.linspace(0.,twopi,n phi)
  # set up empty wavefunction array object using pythtb wf array()
  # creates 1D array of length [n phi], with hidden [nbands, norbs]
  evec array=wf array(ref model,[n phi])
  # run over values of phi and fill the array
  for k,phi in enumerate(phit[0:-1]):  # skip last point of loop
    evec_array[k] =get_evecs(t0,s,phi,alpha)
  evec_array[-1] = evec_array[0] # copy first point to last point of loop
  # now compute and store the Berry phase
 berry_phi[j] = evec_array.berry_phase([0]) # [0] specifices lowest band
 print("%3d %7.3f %7.3f"% (j, alpha, berry phi[j]))
# compute Berry phases for alpha loops for several phi values
# using pythtb wf array() method
phis=np.linspace(0.,twopi,13) # 0 to 2pi in 12 increments
berry alpha=np.zeros like(phis)
print("\nBerry phases for alpha loops versus phi")
for j,phi in enumerate(phis):
 n alpha=61
 alphat=np.linspace(0.,twopi,n alpha)
  evec array=wf array(ref model, [n alpha])
  for k,alpha in enumerate(alphat[0:-1]):
   evec array[k] = get evecs(t0,s,phi,alpha)
 evec array[-1] = evec array[0]
 berry alpha[j] = evec array.berry phase([0])
 print("%3d %7.3f %7.3f"% (j, phi, berry_alpha[j]))
# now illustrate use of wf array() to set up 2D array
# recompute Berry phases and compute Berry curvature
# -----
n phi=61
n alp=61
n_cells=(n_phi-1) * (n_alp-1)
phi=np.linspace(0.,twopi,n_phi)
alp=np.linspace(0.,twopi,n alp)
evec_array=wf_array(ref_model,[n_phi,n_alp]) # empty 2d wavefunction array
for i in range(n phi):
 for j in range(n alp):
   evec array[i,j] = get evecs(t0,s,phi[i],alp[j])
evec_array.impose_loop(0)
                          # copy first to last points in each dimension
evec array.impose loop(1)
bp of alp=evec array.berry phase([0],0) # compute phi Berry phases vs. alpha
bp of phi=evec array.berry phase([0],1) # compute alpha Berry phases vs. phi
# compute 2D array of Berry fluxes for band 0
flux=evec array.berry flux([0])
print("\nFlux = %7.3f = 2pi * %7.3f"% (flux, flux/twopi))
```

```
curvature=evec array.berry flux([0],individual phases=True)*float(n cells)
# plots
fig,ax=plt.subplots(1,3,figsize=(10,4),gridspec kw={'width ratios':[1,1,2]})
(ax0,ax1,ax2)=ax
ax0.set xlim(0.,1.)
ax0.set ylim(-6.5, 6.5)
ax0.set xlabel(r"$\alpha/2\pi$")
ax0.set ylabel(r"Berry phase $\phi(\alpha)$ for $\varphi$ loops")
ax0.set title("Berry phase")
for shift in (-twopi, 0., twopi):
  ax0.plot(alp/twopi,bp_of_alp+shift,color='k')
ax0.scatter(alphas/twopi,berry_phi,color='k')
ax1.set_xlim(0.,1.)
ax1.set_ylim(-6.5,6.5)
ax1.set_xlabel(r"$\varphi/2\pi$")
ax1.set_ylabel(r"Berry phase $\phi(\varphi)$ for $\alpha$ loops")
ax1.set_title("Berry phase")
for shift in (-twopi, 0., twopi):
 ax1.plot(phi/twopi,bp of phi+shift,color='k')
ax1.scatter(phis/twopi,berry alpha,color='k')
X=alp[0:-1]/twopi + 0.5/float(n alp-1)
Y=phi[0:-1]/twopi + 0.5/float(n phi-1)
cs=ax2.contour(X,Y,curvature,colors='k')
ax2.clabel(cs, inline=1, fontsize=10)
ax2.set title("Berry curvature")
ax2.set xlabel(r"$\alpha/2\pi$")
ax2.set xlim(0.,1.)
ax2.set ylim(0.,1.)
ax2.set ylabel(r"$\varphi/2\pi$")
fig.tight layout()
fig.savefig("trimer.pdf")
```

D.7 Berry Phase of Alternating Site Chain

Program chain_alt_bp.py returns to the same alternating chain model presented in Appendix D.4 and computes the Berry phase of the lower band as the Bloch wavevector k is cycled around the BZ, as presented in Section 3.4. The Berry phase is again calculated first explicitly and then again using the wf_array method. In the explicit calculation, one has to be sure to account for the extra phase factor in Eq. (3.74) as discussed on p. 108. The wf_array method, by contrast, takes care of this automatically.

chain_alt_bp.py

```
#!/usr/bin/env python
from __future__ import print_function # python3 style print
# Chain with alternating site energies and hoppings
from pythtb import *
import matplotlib.pyplot as plt
```

```
# define function to set up model for a given parameter set
def set model(t,del t,Delta):
 lat=[[1.0]]
 orb=[[0.0],[0.5]]
 my model=tbmodel(1,1,lat,orb)
 my model.set onsite([Delta,-Delta])
 my_model.add_hop(t+del_t, 0, 1, [0])
 my_model.add_hop(t-del_t, 1, 0, [1])
 return my model
# set parameters of model
t=-1.0 # average hopping
del_t=-0.3  # bond strength alternation
Delta= 0.4  # site energy alternation
my_model=set_model(t,del_t,Delta)
my model.display()
# explicit calculation of Berry phase
# -----
# set up and solve the model on a discretized k mesh
              # 60 equal intervals around the unit circle
(k vec,k dist,k node) = my model.k path('full',nk,report=False)
(eval, evec) = my_model.solve_all(k_vec, eig_vectors=True)
evec=evec[0] # pick band=0 from evec[band,kpoint,orbital]
              # now just evec[kpoint,orbital]
# k-points 0 and 60 refer to the same point on the unit circle
# so we will work only with evec[0],...,evec[59]
# compute Berry phase of lowest band
prod=1.+0.j
for i in range(1,nk-1):
                                 # <evec 0 evec 1>...<evec 58 evec 59>
 prod*=np.vdot(evec[i-1],evec[i]) # a*=b means a=a*b
# now compute the phase factors needed for last inner product
orb=np.array([0.0,0.5]) # relative coordinates of orbitals
phase=np.exp((-2.j)*np.pi*orb)  # construct phase factors
evec_last=phase*evec[0]  # evec[60] constructed from evec[0]
prod*=np.vdot(evec[-2],evec_last) # include <evec_59|evec_last>
print("Berry phase is %7.3f"% (-np.angle(prod)))
# Berry phase via the wf_array method
# fill with eigensolutions
berry phase=evec array.berry phase([0]) # Berry phase of bottom band
print("Berry phase is %7.3f"% berry phase)
```

D.8 Infinite and Finite Three-Site Chain

Program chain_3_site.py solves a 1D TB model with three sites per unit cell and computes the Wannier function centers in the multiband context as discussed in Section 3.6.3. The nearest-neighbor hoppings are all equal, but the site energies are

modulated according to Eq. (3.132) in such a way that the evolution of parameter λ from 0 to 2π represents a sliding charge-density wave, as illustrated in Fig. 1.4 in Chapter 1.

The Wannier centers are first computed for the periodic bulk via calculations of multiband Berry phases, first for the lowest-energy band, then for the next lowest, then by computing the centers of the maximally localized Wannier functions (MLWFs) from the multiband Berry phases for the two lowest bands taken as a group (this requires specifying berry_evals=True in the call to berry_phase).

Next, the finite-system MLWF centers are computed for a finite chain of 10 unit cells cut from the infinite bulk model using the <code>cut_piece</code> method of the PYTHTB <code>tb_model</code> class. Again this is done for the lowest band alone (more precisely, the 10 lowest eigenstates), the second band (next 10 eigenstates), and the joint group including both (20 lowest states). The MLWF centers are obtained by diagonalizing the 10×10 or 20×20 X position matrix of Eq. (3.129) using the <code>position_hwf</code> method of the <code>PYTHTB</code> package.

The resulting printout is reproduced on p. 134. The format is designed to facilitate comparison between the two sets of results. In each case it is evident that while the Wannier centers deviate from the bulk positions near the ends of the chain, they converge rapidly to those bulk values as one goes deeper into bulk-like region of the chain.

chain_3_site.py

```
#!/usr/bin/env python
from future import print function # python3 style print
# Chain with three sites per cell
from pythtb import *
import matplotlib.pyplot as plt
# define function to construct model
def set model(t,delta,lmbd):
    lat=[[1.0]]
    orb=[[0.0],[1.0/3.0],[2.0/3.0]]
    model=tb_model(1,1,lat,orb)
    model.set hop(t, 0, 1, [0])
    model.set_hop(t, 1, 2, [0])
    model.set hop(t, 2, 0, [1])
    onsite 0=delta*(-1.0)*np.cos(2.0*np.pi*(lmbd-0.0/3.0))
    onsite 1=delta*(-1.0)*np.cos(2.0*np.pi*(lmbd-1.0/3.0))
    onsite 2=delta*(-1.0)*np.cos(2.0*np.pi*(lmbd-2.0/3.0))
    model.set onsite([onsite 0,onsite 1,onsite 2])
    return (model)
# construct the model
t = -1.3
delta=2.0
lmbd=0.3
my model=set model(t,delta,lmbd)
```

```
# compute the results on a uniform k-point grid
evec_array=wf_array(my_model,[21]) # set array dimension
evec array.solve on grid([0.])
                                         # fill with eigensolutions
# obtain Berry phases and convert to Wannier center positions
# constrained to the interval [0.,1.]
wfc0=evec array.berry phase([0])/(2.*np.pi)%1.
wfc1=evec_array.berry_phase([1])/(2.*np.pi)%1.
x=evec array.berry phase([0,1],berry evals=True)/(2.*np.pi)%1.
awfc0=x[0]
qwfc1=x[1]
print ("Wannier centers of bands 0 and 1:")
print((" Individual"+" Wannier centers: "+2*"%7.4f") % (wfc0,wfc1))
print((" Multiband "+" Wannier centers: "+2*"%7.4f") % (gwfc1,gwfc0))
print()
# construct and solve finite model by cutting 10 cells from infinite chain
finite model=my model.cut piece(10,0)
(feval, fevec) = finite_model.solve_all(eig_vectors=True)
print ("Finite-chain eigenenergies associated with")
print(("Band 0:"+10*"%6.2f")% tuple(feval[0:10]))
print(("Band 1:"+10*"%6.2f")% tuple(feval[10:20]))
# find maxloc Wannier centers in each band subspace
xbar0=finite model.position hwf(fevec[0:10,],0)
xbar1=finite model.position hwf(fevec[10:20,],0)
xbarb=finite model.position hwf(fevec[0:20,],0)
print ("\nFinite-chain Wannier centers associated with band 0:")
print((10*"%7.4f")% tuple(xbar0))
x=10*(wfc0,)
print(("Compare with bulk:\n"+10*"\%7.4f")\% x)
print ("\nFinite-chain Wannier centers associated with band 1:")
print((10*"%7.4f")% tuple(xbar1))
x=10*(wfc1,)
print(("Compare with bulk:\n"+10*"%7.4f")% x)
print ("\nFirst 10 finite-chain Wannier centers associated with bands"+
  "0 and 1:")
print((10*"%7.4f")% tuple(xbarb[0:10]))
x=5*(gwfc0,gwfc1)
print(("Compare with bulk:\n"+10*"%7.4f")% x)
```

D.9 Adiabatic Cycle for Infinite Three-Site Chain

Program chain_3_cycle.py treats the same model as in Appendix D.8, but this time treating λ as a cyclic parameter that runs from 0 to 2π and focusing on tracking the bulk Wannier center position of the lowest band through this cycle. The approach is almost the same as is used in the "one-dimensional cycle of 1D tight-binding model" example program in the standard PYTHTB distribution, except for the usage of the wf_array method in two-dimensional (k,λ) space there, which has the advantage of providing automatic enforcement of continuity with respect to λ . By contrast, continuity is enforced explicitly in the version here in the lines following the comment line #enforce smooth evolution of xbar. The plotted output of this program appears as Fig. 4.5 in Section 4.2.3.

chain_3_cycle.py

```
#!/usr/bin/env python
from __future__ import print_function # python3 style print
# Chain with three sites per cell - cyclic variation
from pythtb import *
import matplotlib.pyplot as plt
# define function to construct model
def set model(t,delta,lmbd):
   lat=[[1.0]]
   orb=[[0.0],[1.0/3.0],[2.0/3.0]]
   model=tb model(1,1,lat,orb)
   model.set hop(t, 0, 1, [0])
   model.set hop(t, 1, 2, [0])
   model.set hop(t, 2, 0, [1])
   onsite 0=delta*(-1.0)*np.cos(lmbd)
   onsite 1=delta*(-1.0)*np.cos(lmbd-2.0*np.pi/3.0)
   onsite 2=delta*(-1.0)*np.cos(lmbd-4.0*np.pi/3.0)
   model.set onsite([onsite 0,onsite 1,onsite 2])
   return (model)
def get xbar(band, model):
    evec array=wf array(model,[21])
                                            # set array dimension
    evec_array.solve_on_grid([0.])
                                            # fill with eigensolutions
   wfc=evec_array.berry_phase([band])/(2.*np.pi) # Wannier centers
   return(wfc)
# set fixed parameters
t = -1.3
delta=2.0
# obtain results for an array of lambda values
lmbd=np.linspace(0.,2.*np.pi,61)
xbar=np.zeros like(lmbd)
for j,lam in enumerate(lmbd):
    my model=set model(t,delta,lam)
    xbar[j] =get_xbar(0,my_model)
                                     # Wannier center of bottom band
# enforce smooth evolution of xbar
for j in range(1,61):
   delt=xbar[j]-xbar[j-1]
    delt=-0.5+(delt+0.5)%1. # add integer to enforce |delt| < 0.5
   xbar[j] = xbar[j-1] + delt
# set up the figure
fig, ax = plt.subplots(figsize=(5.,3.))
ax.set_xlim([0.,2.*np.pi])
ax.set_ylim([-0.6,1.1])
ax.set_xlabel(r"Parameter $\lambda$")
ax.set ylabel(r"Wannier center position")
xlab = [\overline{r}"0", r"\$ \pi'3\$", r"\$2 \pi'3\$", r"\$ \pi'\$4 \pi'3\$", r"\$5 \pi'3\$", r"\$2 \pi'3\$"]
ax.set_xticks(np.linspace(0.,2.*np.pi,num=7))
ax.set xticklabels(xlab)
ax.plot(lmbd,xbar,'k')
                           # plot Wannier center and some periodic images
ax.plot(lmbd,xbar-1.,'k')
ax.plot(lmbd,xbar+1.,'k')
ax.axhline(y=1.,color='k',linestyle='dashed') # horizontal reference lines
ax.axhline(y=0.,color='k',linestyle='dashed')
fig.tight layout()
fig.savefig("chain 3 cycle.pdf")
```

D.10 Surface Properties of Alternating Site Model

Program chain_alt_surf.py returns to the alternating site model of Appendices D.4 and D.7, but now focuses on computing the surface properties of a finite chain. Two types of variation are considered: modification of the site energy of the last orbital on the chain, and modification of the bulk Hamiltonian along an adiabatic cycle corresponding to a charge pump. The calculations are discussed in Section 4.5.3, where the resulting plot is presented as Fig. 4.16. For each kind of parameter variation, three plots are presented, showing the variation of the energy eigenvalues, the Wannier center positions (computed along the same lines as in Appendix D.8), and the surface charge computed using the ramp-function method of Eqs. (4.78) and (4.79).

chain_alt_surf.py

```
#!/usr/bin/env python
from __future__ import print_function # python3 style print
# Chain with alternating site energies and hoppings
# Study surface properties of finite chain
from pythtb import *
import matplotlib as mpl
import matplotlib.pyplot as plt
# to set up model for given surface energy shift and lambda
def set_model(n_cell,en_shift,lmbd):
  # set parameters of model
  t=-1.0 # average hopping
  Delta=-0.4*np.cos(lmbd) # site energy alternation
  del t=-0.3*np.sin(lmbd) # bond strength alternation
  # construct bulk model
  lat=[[1.0]]
  orb=[[0.0],[0.5]]
  bulk model=tbmodel(1,1,lat,orb)
  bulk model.set onsite([Delta, -Delta])
  bulk_model.add_hop(t+del_t, 0, 1, [0])
  bulk model.add hop(t-del t, 1, 0, [1])
  # cut chain of length n_cell and shift energy on last site
  finite_model=bulk_model.cut_piece(n_cell,0)
  finite_model.set_onsite(en_shift,ind_i=2*n_cell-1,mode='add')
  return finite model
# set Fermi energy and number of cells
Ef=0.18
n cell=20
n_orb=2*n_cell
# set number of parameter values to run over
n param=101
# initialize arrays
params=np.linspace(0.,1.,n_param)
eig sav=np.zeros((n orb,n param),dtype=float)
```

```
xbar sav=np.zeros((n orb,n param),dtype=float)
nocc_sav=np.zeros((n_param),dtype=int)
surf_sav=np.zeros((n_param),dtype=float)
count=np.zeros((n orb),dtype=float)
# initialize plots
mpl.rc('font', size=10) # set global font size
fig, ax=plt.subplots(3,2,figsize=(7.,6.),
    gridspec kw={'height ratios':[2,1,1]},sharex="col")
# loop over two cases: vary surface site energy, or vary lambda
for mycase in ['surface energy', 'lambda']:
  if mycase == 'surface energy':
    (ax0,ax1,ax2)=ax[:,0] # axes for plots in left panels
    ax0.text(-0.30,0.90,'(a)',size=22.,transform=ax0.transAxes)
    lmbd=0.15*np.pi*np.ones((n param),dtype=float)
    en shift=-3.0+6.0*params
    abscissa=en shift
  elif mycase == 'lambda':
    (ax0,ax1,ax2)=ax[:,1] # axes for plots in right panels
    ax0.text(-0.30,0.90,'(b)',size=22.,transform=ax0.transAxes)
    lmbd=params*2.*np.pi
    en_shift=0.2*np.ones((n_param),dtype=float)
    abscissa=params
  # loop over parameter values
  for j in range(n param):
    # set up and solve model; store eigenvalues
    my model=set model(n cell,en shift[j],lmbd[j])
    (eval, evec) = my model.solve all(eig vectors=True)
    # find occupied states
   nocc=(eval<Ef).sum()
    ovec=evec[0:nocc,:]
    # get Wannier centers
   xbar_sav[0:nocc,j] =my_model.position_hwf(ovec,0)
    # get electron count on each site
    # convert to charge (2 for spin; unit nuclear charge per site)
    # compute surface charge down to depth of 1/3 of chain
    for i in range(n orb):
      count[i] = np.real(np.vdot(evec[:nocc,i],evec[:nocc,i]))
    charge=-2.*count+1.
    n_cut=int(0.67*n_orb)
    surf sav[j]=0.5*charge[n cut-1]+charge[n cut:].sum()
    # save information for plots
    nocc sav[j]=nocc
    eig sav[:,j]=eval
  ax0.set_xlim(0.,1.)
  ax0.set ylim(-2.8, 2.8)
  ax0.set ylabel(r"Band energy")
  ax0.axhline(y=Ef,color='k',linewidth=0.5)
  for n in range(n orb):
    ax0.plot(abscissa,eig sav[n,:],color='k')
  ax1.set xlim(0.,1.)
```

```
ax1.set ylim(n cell-4.6,n cell+0.4)
  ax1.set yticks(np.linspace(n cell-4, n cell, 5))
  \#ax1.set_ylabel(r"\$\bar{x}$")
  ax1.set_ylabel(r"Wannier centers")
  for j in range(n param):
   nocc=nocc sav[j]
    ax1.scatter([abscissa[j]]*nocc,xbar sav[:nocc,j],color='k',
      s=3., marker='o', edgecolors='none')
 ax2.set ylim(-2.2,2.2)
 ax2.set yticks([-2.,-1.,0.,1.,2.])
 ax2.set ylabel(r"Surface charge")
 if mycase == 'surface energy':
    ax2.set xlabel(r"Surface site energy")
 elif mycase == 'lambda':
    ax2.set xlabel(r"$\lambda/2\pi$")
  ax2.set xlim(abscissa[0],abscissa[-1])
 ax2.scatter(abscissa, surf sav, color='k', s=3., marker='o', edgecolors='none')
  # vertical lines denote surface state at right end crossing the
  # Fermi energy
  for j in range(1,n_param):
    if nocc_sav[j] != nocc_sav[j-1]:
      n=min(nocc sav[j], nocc sav[j-1])
      frac=(Ef-eig sav[n,j-1])/(eig sav[n,j]-eig sav[n,j-1])
      a jump=(1-frac)*abscissa[j-1]+frac*abscissa[j]
      if mycase == 'surface energy' or nocc sav[j] < nocc sav[j-1]:</pre>
        ax0.axvline(x=a_jump,color='k',linewidth=0.5)
        ax1.axvline(x=a_jump,color='k',linewidth=0.5)
        ax2.axvline(x=a jump,color='k',linewidth=0.5)
fig.tight layout()
plt.subplots adjust(left=0.12,wspace=0.4)
fig.savefig("chain alt surf.pdf")
```

D.11 Band Structure of the Haldane Model

Program haldane_bsr.py computes the band structure of the Haldane model of Eq. (5.1) for four different parameter sets, with the plotted output appearing as Fig. 5.2 in Section 5.1.1. In addition to plotting the band structure, the program places filled or open circles on the bands at the high-symmetry K and K' points according to which site dominates the character of the lower-energy band, thereby highlighting the band inversion.

haldane_bsr.py

```
#!/usr/bin/env python
from __future__ import print_function # python3 style print
# Band structure of Haldane model
from pythtb import * # import TB model class
import matplotlib.pyplot as plt
# set model parameters
```

```
delta=0.7 # site energy shift
         # real first-neighbor hopping
t = -1.0
t2=0.15
          # imaginary second-neighbor hopping
def set model(delta,t,t2):
  lat=[[1.0,0.0],[0.5,np.sqrt(3.0)/2.0]]
  orb=[[1./3.,1./3.],[2./3.,2./3.]]
 model=tb model(2,2,lat,orb)
 model.set_onsite([-delta,delta])
  for lvec in ([ 0, 0], [-1, 0], [ 0, -1]):
   model.set_hop(t, 0, 1, lvec)
  for lvec in ([ 1, 0], [-1, 1], [ 0,-1]):
   model.set hop(t2*1.j, 0, 0, lvec)
  for lvec in ([-1, 0], [ 1,-1], [ 0, 1]):
    model.set_hop(t2*1.j, 1, 1, lvec)
  return model
# construct path in k-space and solve model
path=[[0.,0.],[2./3.,1./3.],[.5,.5],[1./3.,2./3.], [0.,0.]]
label=(r'$\Gamma $',r'$K$', r'$M$', r'$K^\prime$', r'$\Gamma $')
(k vec, k dist, k node) = set model(delta, t, t2).k path(path, 101)
# set up band structure plots
fig, ax = plt.subplots(2,2,figsize=(8.,6.),sharex=True,sharey=True)
ax=ax.flatten()
t2 values=[0.,-0.06,-0.1347,-0.24]
labs=['(a)','(b)','(c)','(d)']
for j in range(4):
  my model=set model(delta,t,t2 values[j])
  evals=my model.solve all(k vec)
 ax[j].set xlim([0,k node[-1]])
 ax[j].set xticks(k node)
  ax[j].set xticklabels(label)
  for n in range(len(k node)):
    ax[j].axvline(x=k_node[n],linewidth=0.5, color='k')
  ax[j].set_ylabel("Energy")
  ax[j].set_ylim(-3.8,3.8)
  for n in range(2):
    ax[j].plot(k_dist,evals[n],color='k')
  # filled or open dots at K and K' following band inversion
  for m in [1,3]:
    kk=k node[m]
    (en, ev) =my_model.solve_one(path[m],eig_vectors=True)
    if np.abs(ev[0,0]) > np.abs(ev[0,1]):
                                           #ev[band,orb]
      en=[en[1],en[0]]
    ax[j].scatter(kk,en[0],s=40.,marker='o',edgecolors='k',
      facecolors='w',zorder=4)
    ax[j].scatter(kk,en[1],s=40.,marker='o',color='k',zorder=6)
  ax[j].text(0.20,3.1,labs[j],size=18.)
# save figure as a PDF
fig.tight layout()
fig.savefig("haldane bsr.pdf")
```

D.12 Berry Curvature of the Haldane Model

Program haldane_bcurv.py generates contour plots of the Berry curvature of the Haldane model for three parameter sets. The model is solved on a 2D mesh of k-points using the solve_on_grid method of wf_array. Then the berry_flux method is used in two different ways: once with individual_phases=True to obtain the flux through each plaquette of the mesh, and once with the default False value to obtain the total flux through the entire 2D BZ. When normalized, the first gives the Berry curvature, which is plotted using the contour method of PYPLOT, and the second gives the Chern number, which is printed out. The plot appears as Fig. 5.3 in Section 5.1.1.

haldane_bcurv.py

```
#!/usr/bin/env python
from future import print function # python3 style print
# Berry curvature of Haldane model
from pythtb import * # import TB model class
import matplotlib.pyplot as plt
# define setup of Haldane model
def set model(delta,t,t2):
  lat = [[1.0, 0.0], [0.5, np.sqrt(3.0)/2.0]]
  orb=[[1./3.,1./3.],[2./3.,2./3.]]
  model=tb model(2,2,lat,orb)
  model.set onsite([-delta,delta])
  for lvec in ([ 0, 0], [-1, 0], [ 0,-1]):
    model.set_hop(t, 0, 1, lvec)
  for lvec in ([ 1, 0], [-1, 1], [ 0,-1]):
    model.set_hop(t2*1.j, 0, 0, lvec)
  for lvec in ([-1, 0], [ 1,-1], [ 0, 1]):
   model.set_hop(t2*1.j, 1, 1, lvec)
  return model
# miscellaneous setup
delta=0.7 # site energy shift
t = -1.0
           # real first-neighbor hopping
nk=61
dk=2.*np.pi/(nk-1)
k0 = (np.arange(nk-1)+0.5)/(nk-1)
kx=np.zeros((nk-1,nk-1),dtype=float)
ky=np.zeros((nk-1,nk-1),dtype=float)
sq3o2=np.sqrt(3.)/2.
for i in range(nk-1):
  for j in range(nk-1):
    kx[i,j] = sq302*k0[i]
    ky[i,j] = -0.5*k0[i]+k0[j]
fig, ax=plt.subplots(1,3,figsize=(11,4))
labs=['(a)','(b)','(c)']
```

```
# compute Berry curvature and Chern number for three values of t2
for j,t2 in enumerate([0.,-0.06,-0.24]):
 my_model=set_model(delta,t,t2)
 my_array=wf_array(my_model,[nk,nk])
 my array.solve on grid([0.,0.])
  bcurv=my_array.berry_flux([0],individual_phases=True)/(dk*dk)
  chern=my_array.berry_flux([0])/(2.*np.pi)
 print('Chern number =', "%8.5f"%chern)
  # make contour plot of Berry curvature
 pos lvls= 0.02*np.power(2.,np.linspace(0,8,9))
 neq lvls=-0.02*np.power(2.,np.linspace(8,0,9))
  ax[j].contour(kx,ky,bcurv,levels=pos lvls,colors='k')
 ax[j].contour(kx,ky,bcurv,levels=neg_lvls,colors='k',linewidths=1.4)
  # remove rectangular box and draw parallelogram, etc.
  ax[j].xaxis.set_visible(False)
  ax[j].yaxis.set_visible(False)
  for loc in ["top", "bottom", "left", "right"]:
    ax[j].spines[loc].set_visible(False)
  ax[j].set(aspect=1.)
 ax[j].plot([0,sq3o2,sq3o2,0,0],[0,-0.5,0.5,1,0],color='k',linewidth=1.4)
  ax[j].set_xlim(-0.05,sq3o2+0.05)
  ax[i].text(-.35,0.88,labs[i],size=24.)
fig.savefig("haldane bcurv.pdf")
```

D.13 Hybrid Wannier Centers and Edge States of the Haldane Model

Program haldane_topo.py produces plots of the hybrid Wannier center flow in the bulk, and of the edge band structure of a finite-width ribbon, for the Haldane model with two different parameter sets covering both the trivial and topological cases. The plotted output appears as Fig. 5.4 in Section 5.1.1. The methods used to obtain the bulk Wannier centers should be familiar by now. The cut_piece method of the tb_model class is applied to the 2D bulk model my_model to obtain the new 1D ribbon_model, extending indefinitely in the horizontal direction but of finite width vertically, here chosen to be 20 cells high. The band structure of the ribbon is plotted using the PYPLOT scatter method, instead of as a continuous line, so that the identity of the states localized at the top and bottom edges of the ribbon can be distinguished. This is done by assigning each plotted point a weight that is normally 1, but that is reduced for states at the bottom edge. These are identified by using the position_expectation method of PYTHTB to compute the component of the position_expectation value in the vertical direction for each Bloch eigenvector.

haldane_topo.py

```
#!/usr/bin/env python
from __future__ import print_function # python3 style print
# Band structure of Haldane model
```

```
from pythtb import * # import TB model class
import matplotlib.pyplot as plt
# define setup of Haldane model
def set model(delta,t,t2):
  lat=[[1.0,0.0],[0.5,np.sqrt(3.0)/2.0]]
 orb=[[1./3.,1./3.],[2./3.,2./3.]]
 model=tb model(2,2,lat,orb)
 model.set_onsite([-delta,delta])
  for lvec in ([ 0, 0], [-1, 0], [ 0,-1]):
   model.set_hop(t, 0, 1, lvec)
  for lvec in ([ 1, 0], [-1, 1], [ 0,-1]):
   model.set hop(t2*1.j, 0, 0, lvec)
  for lvec in ([-1, 0], [ 1,-1], [ 0, 1]):
    model.set_hop(t2*1.j, 1, 1, lvec)
 return model
# set model parameters and construct bulk model
delta=0.7 # site energy shift
t = -1.0
            # real first-neighbor hopping
nk=51
# For the purposes of plot labels:
   Real space is (r1,r2) in reduced coordinates
  Reciprocal space is (k1,k2) in reduced coordinates
# Below, following Python, these are (r0,r1) and (k0,k1)
# set up figures
fig, ax=plt.subplots(2,2,figsize=(7,6))
# run over two choices of t2
for j2,t2 in enumerate([-0.06,-0.24]):
  # solve bulk model on grid and get hybrid Wannier centers along r1
  # as a function of k0
 my model=set model(delta,t,t2)
 my array=wf array(my model,[nk,nk])
 my_array.solve_on_grid([0.,0.])
 rbar_1 = my_array.berry_phase([0],1,contin=True)/(2.*np.pi)
  # set up and solve ribbon model that is finite along direction 1
 width=20
 nkr=81
  ribbon model=my model.cut piece(width,fin dir=1,glue edgs=False)
  (k_vec,k_dist,k_node)=ribbon_model.k_path('full',nkr,report=False)
  (rib_eval,rib_evec)=ribbon_model.solve_all(k_vec,eig_vectors=True)
 nbands=rib eval.shape[0]
  (ax0, ax1) = ax[j2,:]
  # hybrid Wannier center flow
 k0=np.linspace(0.,1.,nk)
  ax0.set xlim(0.,1.)
  ax0.set_ylim(-1.3,1.3)
 ax0.set xlabel(r"$\kappa 1/2\pi$")
  ax0.set_ylabel(r"HWF centers")
  for shift in (-2.,-1.,0.,1.):
    ax0.plot(k0,rbar 1+shift,color='k')
  # edge band structure
 k0=np.linspace(0.,1.,nkr)
```

```
ax1.set xlim(0.,1.)
 ax1.set_ylim(-2.5, 2.5)
  ax1.set_xlabel(r"$\kappa_1/2\pi$")
  ax1.set ylabel(r"Edge band structure")
  for (i,kv) in enumerate(k0):
    # find expectation value <rl> at i'th k-point along direction k0
    pos exp=ribbon model.position expectation(rib evec[:,i],dir=1)
    # assign weight in [0,1] to be 1 except for edge states near bottom
    weight=3.0*pos exp/width
    for j in range (nbands):
      weight[j] = min(weight[j],1.)
    # scatterplot with symbol size proportional to assigned weight
    s=ax1.scatter([k_vec[i]]*nbands, rib_eval[:,i],
         s=0.6+2.5*weight, c='k', marker='o', edgecolors='none')
# save figure as a PDF
aa=ax.flatten()
for i, lab in enumerate(['(a)','(b)','(c)','(d)']):
  aa[i].text(-0.45,0.92,lab,size=18.,transform=aa[i].transAxes)
fig.tight_layout()
plt.subplots adjust(left=0.15, wspace=0.6)
fig.savefig("haldane topo.pdf")
```

D.14 Entanglement Spectrum of the Haldane Model

Program haldane_entang.py produces plots of the entanglement spectrum for the Haldane model for two parameter sets, one in the trivial phase and one in the topological phase. The entanglement spectrum, mentioned briefly at the end of Section 5.1.1, is defined as follows. For an infinite 1D chain, one arbitrarily chooses a dividing surface that partitions the chain into a lower region A and an upper region B. The entanglement spectrum is then given by computing the eigenvalues of the reduced density matrix $\rho_A = \mathcal{P}_A \rho \mathcal{P}_A$, where $\rho_{ij} = \sum_{n}^{\text{occ}} \psi_{nk}^*(i) \psi_{nk}(j)$ is the one-particle density matrix in the TB basis and \mathcal{P}_A is the projector onto subregion A. For a 2D system, one plots these eigenvalues for the effective 1D system extending along lattice direction \mathbf{a}_2 as a function of κ_1 , as shown in Fig. D.3. In practice, haldane_entang.py obtains the entanglement spectrum from the same finite-width ribbon constructed in haldane_topo.py, partitioned in the middle; edge states on the physical boundary of the ribbon have little influence since they are far from the dividing surface.

A comparison of Fig. D.3(b) and Fig. 5.4(c) shows that an upward flow of the hybrid Wannier function (HWF) centers with increasing κ_1 corresponds to a downward flow of the ρ_A eigenvalues. This is only natural, since an upward migration of Wannier functions corresponds to a depletion of occupation in the lower region A.

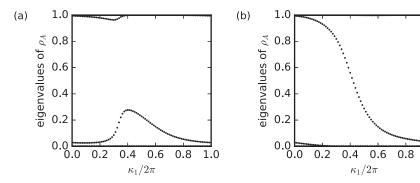


Figure D.3 Entanglement spectrum of the Haldane model with $\Delta=0.7$ and $t_1=-1$. A ribbon geometry of width 20 unit cells along the κ_2 direction was subdivided into bottom and top halves A and B, respectively; eigenvalues of the reduced density matrix ρ_A (see text) are plotted vertically. (a) Trivial phase $(t_2=0.10)$ with C=0. (b) Topological phase $(t_2=-0.24)$ with C=1.

The origin of the name "entanglement spectrum" is associated with the definition of an *entanglement entropy* given by $\text{Tr}\left[-\rho_A \ln \rho_A - (1-\rho_A) \ln (1-\rho_A)\right]$. This object is dominated by eigenvalues of ρ_A lying in the middle of the interval [0,1], so roughly speaking it counts the number of states that cannot be localized entirely in A or in B, but instead are "entangled" across the dividing surface.

The flow of the single-particle entanglement spectrum, as shown in Fig. D.3, can be used as a tool to extract the Chern index of a 2D magnetic insulator, much like the edge band structures and HWF flow patterns in Fig. 5.4. The three methods are, in a sense, close cousins. However, the HWF flow approach has the advantage of requiring only a simple bulk calculation, with no introduction of artificial bounding surfaces or edge terminations. When generalized to the many-body context, however, the entanglement entropy becomes a useful tool for the study of strongly correlated topological states with long-range entanglement, such as those mentioned in Section 5.5.3.

haldane_entang.py

```
#!/usr/bin/env python
from __future__ import print_function # python3 style print
# Entanglement spectrum of Haldane model

from pythtb import * # import TB model class
import matplotlib.pyplot as plt

# define setup of Haldane model
def set_model(delta,t,t2):
    lat=[[1.0,0.0],[0.5,np.sqrt(3.0)/2.0]]
    orb=[[1./3.,1./3.],[2./3.,2./3.]]
    model=tb_model(2,2,lat,orb)
    model.set_onsite([-delta,delta])
    for lvec in ([ 0, 0], [-1, 0], [ 0,-1]):
```

```
model.set hop(t, 0, 1, lvec)
  for lvec in ([1, 0], [-1, 1], [0,-1]): model.set_hop(t2*1.j, 0, 0, lvec)
  for lvec in ([-1, 0], [ 1,-1], [ 0, 1]): model.set_hop(t2*1.j, 1, 1, lvec)
  return model
# set model parameters and construct bulk model
delta=0.7 # site energy shift
           # real first-neighbor hopping
t = -1.0
# set up figures
fig, ax=plt.subplots(1,2,figsize=(7,3))
# run over two choices of t2
for j2,t2 in enumerate([-0.10,-0.24]):
  my model=set model(delta,t,t2)
  # set up and solve ribbon model that is finite along direction 1
  width=20
  nkr=81
  ribbon_model=my_model.cut_piece(width,fin_dir=1,glue_edgs=False)
  (k_vec,k_dist,k_node)=ribbon_model.k_path('full',nkr,report=False)
  (rib eval, rib evec) = ribbon model.solve all(k vec, eig vectors=True)
  nbands=rib eval.shape[0]
  ax1=ax[j2]
  # entanglement spectrum
  k0=np.linspace(0.,1.,nkr)
  ax1.set xlim(0.,1.)
  ax1.set ylim(0.,1)
  ax1.set xlabel(r"$\kappa 1/2\pi$")
  ax1.set ylabel(r"eigenvalues of $\rho A$")
  (nband, nk, norb) = rib evec.shape
  ncut=norb/2
  nocc=nband/2
  for (i,kv) in enumerate(k0):
    # construct reduced density matrix for half of the chain
    dens mat=np.zeros((ncut,ncut),dtype=complex)
    for nb in range (nocc):
      for j1 in range(ncut):
        for j2 in range(ncut):
          dens_mat[j1,j2] += np.conj(rib_evec[nb,i,j1])*rib_evec[nb,i,j2]
    # diagonalize
    spect=np.real(np.linalg.eigvals(dens mat))
    # scatterplot
    s=ax1.scatter([k_vec[i]]*nocc, spect,
         s=4, c='k', marker='o', edgecolors='none')
# save figure as a PDF
aa=ax.flatten()
for i,lab in enumerate(['(a)','(b)']):
  aa[i].text(-0.45,0.92,lab,size=18.,transform=aa[i].transAxes)
fig.tight layout()
plt.subplots adjust(left=0.15,wspace=0.6)
fig.savefig("haldane_entang.pdf")
```

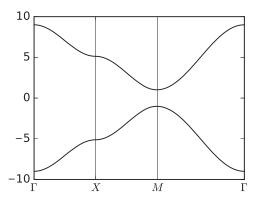


Figure D.4 Band structure of the 2D checkerboard model.

D.15 Band Structure of the Checkerboard Model

Program checkerboard.py computes the band structure for the 2D checkerboard model introduced in Ex. 5.3. The output appears here as Fig. D.4.

checkerboard.py

```
#!/usr/bin/env python
from __future__ import print_function # python3 style print
from pythtb import * # import TB model class
import matplotlib.pyplot as plt
# set geometry
lat=[[1.0,0.0],[0.0,1.0]]
orb=[[0.0,0.0],[0.5,0.5]]
my model=tbmodel(2,2,lat,orb)
# set model
Delta = 5.0
t 0
        = 1.0
tprime = 0.4
my_model.set_sites([-Delta,Delta])
my_model.add_hop(-t_0, 0, 0, [ 1, 0])
my_model.add_hop(-t_0, 0, 0, [ 0, 1])
my_model.add_hop( t_0, 1, 1, [ 1, 0])
my_model.add_hop( t_0, 1, 1, [ 0, 1])
my_model.add_hop( tprime*1j, 1, 0, [ 0, 1])
                             , 1, 0, [ 0, 0])
my model.add hop(-tprime
my model.add hop(-tprime*1j, 1, 0, [ 1, 0])
my_model.display()
# generate k-point path and labels and solve Hamiltonian
path=[[0.0,0.0],[0.0,0.5],[0.5,0.5],[0.0,0.0]]
k_lab=(r'\$\backslash Gamma \$',r'\$X\$', r'\$M\$', r'\$\backslash Gamma \$')
\begin{tabular}{ll} $(\bar{k}\_vec,k\_dist,k\_node)=my\_model.k\_path(path,121) \\ evals=my\_model.solve\_all(k\_vec) \end{tabular}
# plot band structure
fig, ax = plt.subplots(figsize=(4.,3.))
```

```
ax.set_xlim([0,k_node[-1]])
ax.set_xticks(k_node)
ax.set_xticklabels(k_lab)
for n in range(len(k_node)):
    ax.axvline(x=k_node[n], linewidth=0.5, color='k')
ax.plot(k_dist,evals[0],color='k')
ax.plot(k_dist,evals[1],color='k')
fig.savefig("checkerboard bsr.pdf")
```

D.16 Band Structure of the Kane-Mele Model

Program kanemele_bsr.py computes the band structure of the Kane–Mele model of Eq. (5.17) in Section 5.2.2 for two parameter sets exemplifying the normal and topological regimes. This is a spinor model, so we set nspin=2 when creating the model. To simplify the specification of the spin-dependent hoppings in the model, we have predefined Pauli matrices σ_x , σ_y , and σ_z in Cartesian directions, and then σ_a , σ_b , and σ_c in three relevant in-plane directions related by 120° rotations. Note that mode="add" is needed when we specify a second component of the first-neighbor hoppings.

The results are plotted as Fig. 5.12. Filled and open circles are added to denote the orbital character for the highest occupied and lowest unoccupied states at K and K' to highlight the band inversion. Here we have cheated, with the filled and open circles simply exchanged in the second plot; a proper implementation should follow the example of haldane_bsr.py in Appendix D.11.

kanemele_bsr.py

```
#!/usr/bin/env python
from __future__ import print_function # python3 style print
# Tight-binding 2D Kane-Mele model
# C.L. Kane and E.J. Mele, PRL 95, 146802 (2005)
from pythtb import * # import TB model class
import matplotlib.pyplot as plt
# set model parameters
delta=0.7 # site energy
t=-1.0  # spin-independent first-neighbor hop
rashba=0.05  # spin-flip first-neighbor hop
soc list=[-0.06,-0.24] # spin-dependent second-neighbor hop
def set model(t,soc,rashba,delta):
  # set up Kane-Mele model
  lat=[[1.0,0.0],[0.5,np.sqrt(3.0)/2.0]]
  orb=[[1./3.,1./3.],[2./3.,2./3.]]
  model=tb model(2,2,lat,orb,nspin=2)
  model.set onsite([delta,-delta])
  # definitions of Pauli matrices
  sigma_x=np.array([0.,1.,0.,0])
  sigma_y=np.array([0.,0.,1.,0])
```

```
sigma z=np.array([0.,0.,0.,1])
  r3h = np.sqrt(3.0)/2.0
  sigma_a= 0.5*sigma_x-r3h*sigma_y
  sigma_b= 0.5*sigma_x+r3h*sigma_y
  sigma c=-1.0*sigma x
  # spin-independent first-neighbor hops
  for lvec in ([ 0, 0], [-1, 0], [ 0, -1]):
   model.set hop(t, 0, 1, lvec)
  # spin-dependent second-neighbor hops
  for lvec in ([ 1, 0], [-1, 1], [ 0,-1]):
    model.set hop(soc*1.j*sigma z, 0, 0, lvec)
  for lvec in ([-1, 0], [ 1,-1], [ 0, 1]):
   model.set hop(soc*1.j*sigma z, 1, 1, lvec)
  # spin-flip first-neighbor hops
  model.set_hop(1.j*rashba*sigma_a, 0, 1, [0, 0], mode="add")
  model.set_hop(1.j*rashba*sigma_b, 0, 1, [-1, 0], mode="add")
  model.set_hop(1.j*rashba*sigma_c, 0, 1, [ 0,-1], mode="add")
 return model
# construct path in k-space and solve model
path=[[0.,0.],[2./3.,1./3.],[.5,.5],[1./3.,2./3.], [0.,0.]]
label=(r'$\Gamma $',r'$K$', r'$M$', r'$K^\prime$', r'$\Gamma $')
(k_vec,k_dist,k_node) = set_model(t,0.,rashba,delta).k_path(path,101)
# set up band structure plots
fig, ax = plt.subplots(1,2,figsize=(8.,3.))
labs=['(a)','(b)']
for j in range(2):
  my model=set model(t,soc list[j],rashba,delta)
  evals=my model.solve all(k vec)
 ax[j].set xlim([0,k node[-1]])
 ax[j].set xticks(k node)
  ax[j].set_xticklabels(label)
  for n in range(len(k node)):
    ax[j].axvline(x=k_node[n],linewidth=0.5, color='k')
  ax[j].set_ylabel("Energy")
  ax[j].set_ylim(-3.8,3.8)
  for n in range(4):
    ax[j].plot(k dist,evals[n],color='k')
  for m in [1,3]:
    kk=k node[m]
    en=my model.solve one(path[m])
    en=en[1:3] # pick out second and third bands
    if j==1:
                # exchange them in second plot
      en=[en[1],en[0]]
    ax[j].scatter(kk,en[0],s=40.,marker='o',color='k',zorder=6)
    ax[j].scatter(kk,en[1],s=40.,marker='o',edgecolors='k',
      facecolors='w',zorder=4)
  ax[j].text(-0.45,3.5,labs[j],size=18.)
# save figure as a PDF
fig.tight layout()
plt.subplots adjust(wspace=0.35)
fig.savefig("kanemele bsr.pdf")
```

D.17 Hybrid Wannier Centers and Edge States of the Kane-Mele Model

Program kanemele_topo.py produces plots of the hybrid Wannier center flow in the bulk, and of the edge band structure of a finite-width ribbon, for the Kane—Mele model with two different parameter sets covering both the trivial and topological cases, following the example of haldane_topo.py in Appendix D.13. The plotted output appears as Fig. 5.13 in Section 5.2.2.

kanemele_topo.py

```
#!/usr/bin/env python
from __future__ import print_function # python3 style print
# Tight-binding 2D Kane-Mele model
# C.L. Kane and E.J. Mele, PRL 95, 146802 (2005)
from pythtb import * # import TB model class
import matplotlib.pyplot as plt
# set model parameters
delta=0.7  # site energy
t=-1.0  # spin-independent first-neighbor nop
soc=0.06  # spin-dependent second-neighbor hop
rashba=0.05  # spin-flip first-neighbor hop
              # spin-independent first-neighbor hop
soc_list=[-0.06,-0.24] # spin-dependent second-neighbor hop
def set model(t,soc,rashba,delta):
  # set up Kane-Mele model
  lat=[[1.0,0.0],[0.5,np.sqrt(3.0)/2.0]]
  orb=[[1./3.,1./3.],[2./3.,2./3.]]
  model=tb_model(2,2,lat,orb,nspin=2)
  model.set_onsite([delta,-delta])
  # definitions of Pauli matrices
  sigma_x=np.array([0.,1.,0.,0])
  sigma_y=np.array([0.,0.,1.,0])
  sigma_z=np.array([0.,0.,0.,1])
  r3h = np.sqrt(3.0)/2.0
  sigma a= 0.5*sigma x-r3h*sigma y
  sigma b= 0.5*sigma x+r3h*sigma y
  sigma c=-1.0*sigma x
  # spin-independent first-neighbor hops
  for lvec in ([ 0, 0], [-1, 0], [ 0,-1]):
    model.set hop(t, 0, 1, lvec)
  # spin-dependent second-neighbor hops
  for lvec in ([ 1, 0], [-1, 1], [ 0,-1]):
    model.set_hop(soc*1.j*sigma_z, 0, 0, lvec)
  for lvec in ([-1, 0], [ 1,-1], [ 0, 1]):
    model.set_hop(soc*1.j*sigma_z, 1, 1, lvec)
  # spin-flip first-neighbor hops
  model.set_hop(1.j*rashba*sigma_a, 0, 1, [ 0, 0], mode="add")
model.set_hop(1.j*rashba*sigma_b, 0, 1, [-1, 0], mode="add")
  model.set_hop(1.j*rashba*sigma_c, 0, 1, [ 0,-1], mode="add")
  return model
# For the purposes of plot labels:
   Real space is (r1,r2) in reduced coordinates
```

```
Reciprocal space is (k1,k2) in reduced coordinates
# Below, following Python, these are (r0,r1) and (k0,k1)
# set up figures
fig, ax=plt.subplots(2,2,figsize=(7,6))
nk=51
# run over two choices of t2
for je, soc in enumerate (soc list):
  # solve bulk model on grid and get hybrid Wannier centers along r1
  # as a function of k0
 my model=set model(t,soc,rashba,delta)
 my array=wf array(my model,[nk,nk])
 my_array.solve_on_grid([0.,0.])
 rbar = my_array.berry_phase([0,1],1,berry_evals=True,contin=True)/
    (2.*np.pi)
  # set up and solve ribbon model that is finite along direction 1
 width=20
 nkr=81
 ribbon_model=my_model.cut_piece(width,fin_dir=1,glue_edgs=False)
  (k_vec,k_dist,k_node) = ribbon_model.k_path('full',nkr,report=False)
  (rib eval, rib evec) = ribbon model.solve all(k vec, eig vectors=True)
 nbands=rib eval.shape[0]
  (ax0,ax1)=ax[je,:]
  # hybrid Wannier center flow
 k0=np.linspace(0.,1.,nk)
 ax0.set xlim(0.,1.)
  ax0.set ylim(-1.3,1.3)
  ax0.set_xlabel(r"$\kappa_1/2\pi$")
 ax0.set_ylabel(r"HWF centers")
 ax0.axvline(x=0.5,linewidth=0.5, color='k')
  for shift in (-1.,0.,1.,2.):
    ax0.plot(k0,rbar[:,0]+shift,color='k')
    ax0.plot(k0,rbar[:,1]+shift,color='k')
  # edge band structure
 k0=np.linspace(0.,1.,nkr)
  ax1.set_xlim(0.,1.)
  ax1.set_ylim(-2.5, 2.5)
  ax1.set xlabel(r"$\kappa 1/2\pi$")
  ax1.set_ylabel(r"Edge band structure")
  for (i, kv) in enumerate(k0):
    # find expectation value <rl> at i'th k-point along direction k0
    pos exp=ribbon model.position expectation(rib evec[:,i],dir=1)
    # assign weight in [0,1] to be 1 except for edge states near bottom
    weight=3.0*pos exp/width
    for j in range (nbands):
      weight[j] = min(weight[j],1.)
    # scatterplot with symbol size proportional to assigned weight
    s=ax1.scatter([k vec[i]]*nbands, rib eval[:,i],
         s=0.6+2.5*weight, c='k', marker='o', edgecolors='none')
        ax0.text(-0.45,0.92,'(a)',size=18.,transform=ax0.transAxes)
        ax1.text(-0.45,0.92,'(b)',size=18.,transform=ax1.transAxes)
```

```
# save figure as a PDF
aa=ax.flatten()
for i,lab in enumerate(['(a)','(b)','(c)','(d)']):
    aa[i].text(-0.45,0.92,lab,size=18.,transform=aa[i].transAxes)
fig.tight_layout()
plt.subplots_adjust(left=0.15,wspace=0.6)
fig.savefig("kanemele topo.pdf")
```

D.18 Fu-Kane-Mele Model

Program fkm. py solves the Fu–Kane–Mele model of Eq. (5.28) in Section 5.3.4. This is again a spinor model, and the spin-dependent hoppings are now specified using the convention that spin is a NUMPY array containing the coefficients of $\mathbb{1}$, σ_x , σ_y , and σ_z respectively, with the list of directions dir_list specifying the last three of those coefficients (the first is set to zero).

The program generates two plot files, fkm_bsr.pdf and fkm_topo.pdf. The first is the band-structure plot that appears as Fig. D.5. The second is a plot of the hybrid Wannier center flow in two different κ_1 - κ_2 planes, at $\kappa_3 = 0$ and at $\kappa_3 = \pi$, which appears as Fig. 5.19 in Section 5.3.4.

fkm.py

```
#!/usr/bin/env python
from __future__ import print_function # python3 style print
# Three-dimensional Fu-Kane-Mele model
# Fu, Kane and Mele, PRL 98, 106803 (2007)
from pythtb import * # import TB model class
import matplotlib.pyplot as plt
# set model parameters
t=1.0 # spin-independent first-neighbor hop
```

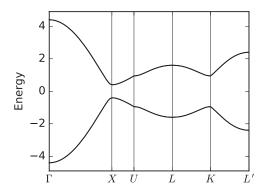


Figure D.5 Band structure of the Fu–Kane-Mele model of Eq. (5.28). Parameter values are $t_0 = 1$, $\Delta t = 0.4$, and $\lambda_{SO} = 0.125$.

```
# modification to t for (111) bond
dt = 0.4
soc=0.125 # spin-dependent second-neighbor hop
def set model(t,dt,soc):
  # set up Fu-Kane-Mele model
  lat=[[.0,.5,.5],[.5,.0,.5],[.5,.5,.0]]
  orb=[[0.,0.,0.],[.25,.25,.25]]
  model=tb model(3,3,lat,orb,nspin=2)
  # spin-independent first-neighbor hops
  for lvec in ([0,0,0],[-1,0,0],[0,-1,0],[0,0,-1]):
    model.set hop(t,0,1,lvec)
 model.set hop(dt,0,1,[0,0,0],mode="add")
  # spin-dependent second-neighbor hops
  lvec\_list = ( [1,0,0] \,, [0,1,0] \,, [0,0,1] \,, [-1,1,0] \,, [0,-1,1] \,, [1,0,-1] \,)
 dir_list=([0,1,-1],[-1,0,1],[1,-1,0],[1,1,0],[0,1,1],[1,0,1])
  for j in range(6):
    spin=np.array([0.]+dir_list[j])
    model.set_hop( 1.j*soc*spin,0,0,lvec_list[j])
    model.set_hop(-1.j*soc*spin,1,1,lvec_list[j])
 return model
my model=set model(t,dt,soc)
my_model.display()
# first plot: compute band structure
# construct path in k-space and solve model
path=[[0.,0.,0.],[0.,.5,.5],[0.25,.625,.625],
         [.5,.5,.5],[.75,.375,.375],[.5,0.,0.]]
label=(r'$\Gamma$',r'$X$',r'$U$',r'$L$',r'$K$',r'$L^\prime$')
(k vec,k dist,k node) = my model.k path(path,101)
evals=my_model.solve_all(k_vec)
# band structure plot
fig, ax = plt.subplots(1,1,figsize=(4.,3.))
ax.set_xlim([0,k_node[-1]])
ax.set_xticks(k_node)
ax.set_xticklabels(label)
for n in range(len(k_node)):
  ax.axvline(x=k node[n],linewidth=0.5, color='k')
ax.set_ylabel("Energy")
ax.set_ylim(-4.9,4.9)
for n in range(4):
 ax.plot(k dist,evals[n],color='k')
fig.tight layout()
fig.savefig("fkm bsr.pdf")
# second plot: compute Wannier flow
# initialize plot
fig, ax = plt.subplots(1,2,figsize=(5.4,2.6),sharey=True)
# Obtain eigenvectors on 2D grid on slices at fixed kappa 3
# Note physical (kappa 1, kappa 2, kappa 3) have python indices (0,1,2)
```

```
kappa2 values=[0.,0.5]
labs=[r'$\kappa_3$=0',r'$\kappa_3$=$\pi$']
nk=41
dk=1./float(nk-1)
wf=wf_array(my_model,[nk,nk])
#loop over slices
for j in range(2):
 for k0 in range(nk):
    for k1 in range(nk):
      kvec=[k0*dk,k1*dk,kappa2 values[j]]
      (eval, evec) = my model.solve one(kvec, eig vectors=True)
      wf[k0,k1] = evec
 wf.impose pbc(mesh dir=0,k dir=0)
 wf.impose_pbc(mesh_dir=1,k_dir=1)
 hwfc=wf.berry_phase([0,1],dir=1,contin=True,berry_evals=True)/
    (2.*np.pi)
 ax[j].set_xlim([0.,1.])
 ax[j].set_xticks([0.,0.5,1.])
 ax[j].set_xlabel(r"$\kappa_1/2\pi$")
 ax[j].set_ylim(-0.5,1.5)
 for n in range(2):
    for shift in [-1.,0.,1.]:
      ax[j].plot(np.linspace(0.,1.,nk),hwfc[:,n]+shift,color='k')
  ax[j].text(0.08,1.20,labs[j],size=12.,bbox=dict(facecolor='w',
    edgecolor='k'))
ax[0].set_ylabel(r"HWF center $\bar{s}_2$")
fig.tight_layout()
plt.subplots adjust(left=0.15, wspace=0.2)
fig.savefig("fkm topo.pdf")
```