

ex07_solution

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Computational Quantum Physics - PHYS 463

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0.1 Solution 07 - Variational Monte Carlo (VMC)

```
[1]: import numpy as np
import matplotlib.pyplot as plt
from functools import partial

try:
    from tqdm.auto import tqdm # progressbar
except ImportError:
    tqdm = lambda x: x
```

```
[2]: # analytical solution for the ground state energy of the TFI in 1D with pbc

def ising1d_energy(L, Gamma):
    # Gamma is in units of the interaction strength J
    def Epsilon(k,h):
        eps = 1 + h**2 + 2 * h * np.cos(k)
        return 2 * np.sqrt(eps)

    i = np.arange(L)
    k = np.pi * ( 2 * i + 1)/L
    energy = Epsilon(k,Gamma).sum()
    return -0.5*energy
```

```
[3]: def err_rel(x, y):
    """relative error"""
    return np.abs((x-y)/y)
```

```
[4]: N = 16          # number of spins
Ns = 512           # number of samples
N_discard = 128    # how many samples to discard
```

```
[5]: # throughout this exercise we work with basis states in the z basis
# represented by arrays containing the quantum numbers -1 and +1

# the following function generates random basis states (uniformly sampled)

def random_states(N, size=1):
    return np.random.choice(np.array([-1,1]), size=(size, N))

random_states(N, 3)
```

```
[5]: array([[ 1, -1,  1,  1,  1,  1,  1,  1,  1, -1,  1,  1,  1,  1,  1,  1],
          [ 1, -1,  1, -1,  1,  1,  1, -1, -1,  1, -1,  1,  1,  1, -1,  1],
          [-1,  1,  1, -1, -1,  1, -1, -1, -1,  1,  1,  1, -1, -1,  1,  1]])
```

0.2 Exercise 7.1 : Mean-field Ansatz

0.2.1 a) Ansatz

```
[6]: def logpsi_mf(params, s):
    """Mean-field Ansatz"""
    Ns, N = s.shape
    d = params[:N]
    u = params[N:]
    return ((s == -1) * d + (s == +1) * u).sum(axis=-1)

def random_params_mf(N, stddev=0.1):

    return np.random.normal(0, stddev, size=(2*N))
```

```
[7]: # test it
x = random_states(N, 5)
params = random_params_mf(N)
logpsi_mf(params, x)
```

```
[7]: array([ 0.17967098,  0.20652151,  0.17270972,  0.08493403, -0.0767841 ])
```

```
[8]: # check the shape
assert logpsi_mf(params, x).shape == (len(x),)
```

0.2.2 b) Gradient

```
[9]: def grad_logpsi_mf(params, s):
    Ns, N = s.shape
    g = np.zeros((Ns,) + params.shape)
    g[:, :N] = (s == -1)
    g[:, N:] = (s == +1)
    return g
```

```
[10]: grad_logpsi_mf(params, x)
```

```
[10]: array([[1., 1., 0., 1., 0., 0., 1., 0., 1., 1., 1., 1., 1., 0., 0., 1.,
          0., 0., 1., 0., 1., 1., 0., 1., 0., 0., 0., 0., 1., 1., 0.],
          [1., 0., 1., 1., 1., 1., 0., 0., 1., 1., 0., 1., 1., 1., 1., 0.,
          0., 1., 0., 0., 0., 0., 1., 1., 0., 0., 1., 0., 0., 0., 0., 1.],
          [0., 1., 1., 1., 0., 0., 1., 0., 0., 0., 0., 0., 1., 0., 1., 0., 1.,
          1., 0., 0., 0., 1., 1., 0., 1., 1., 1., 1., 0., 1., 0., 1.],
          [0., 0., 1., 0., 1., 1., 0., 1., 0., 0., 1., 1., 1., 1., 0., 0.,
          1., 1., 0., 1., 0., 0., 1., 0., 1., 1., 0., 0., 0., 0., 1., 1.],
          [1., 0., 0., 0., 0., 0., 1., 1., 1., 0., 0., 0., 1., 0., 0., 0.,
          0., 1., 1., 1., 1., 1., 0., 0., 0., 1., 0., 0., 0., 0.,
          0., 1., 1., 1., 1., 1., 0., 0., 0., 1., 1., 1., 0., 1., 1., 1.]])
```

```
[11]: assert grad_logpsi_mf(params, x).shape == (len(x),)+params.shape
```

0.2.3 c) Direct sampling

```
[12]: def sample_direct_mf(logpsi, params, N, N_samples):
        assert logpsi == logpsi_mf
        p_down = np.exp(2 * params[:N])
        p_up = np.exp(2 * params[N:])
        pdf = p_up / (p_up + p_down)
        = np.random.uniform(size=(N_samples, N))
        samples = (pdf >) * (+1) + (1 - (pdf >)) * (-1)
        return samples
```

```
[13]: samples_direct = sample_direct_mf(logpsi_mf, params, N, Ns)
        samples_direct
```

```
[13]: array([[ 1,  1, -1, ..., -1, -1, -1],
          [-1,  1,  1, ..., -1,  1, -1],
          [ 1,  1, -1, ..., -1,  1,  1],
          ...,
          [-1, -1, -1, ..., -1, -1, -1],
          [ 1, -1,  1, ...,  1, -1, -1],
          [-1,  1,  1, ..., -1,  1,  1]])
```

```
[14]: assert samples_direct.shape == (Ns, N)
```

0.2.4 d) Monte Carlo Sampling

```
[15]: # we need a function to do the update move to obtain new configurations
        # we will use this as propose_fn below

        def single_spin_flip(x):
            assert x.ndim == 1
            N, = x.shape
```

```

x = x.copy()
i = np.random.randint(0, N)
x[i] *= -1
return x

```

```

[16]: s = np.array([-1,1,1,-1])

print(s)
print(single_spin_flip(s))

```

```

[-1  1  1 -1]
[-1  1  1  1]

```

```

[17]: def sample_step(logpsi, params, x, logpsi_x, propose_fn):
    """
    One sampling step of Monte Carlo Markov Chain.
    logpsi: function giving the log wave-function of a batch of sample given a
    ↪ set of parametes
           (params,x) -> log( x))
    params: parameters of the ansatz
    x: sample on which to do a step (shape (N,))
    logpsi_x: log-value of the wave-function for x
    propose_fn: update move
    """

    if logpsi_x is None:
        # for the sampling we work with a single sample
        # but our ansatz only supports batches
        # so we add a dummy batch dimension here
        logpsi_x = logpsi(params, np.expand_dims(x, 0))[0]

    # propose a new state
    x_proposed = propose_fn(x) # this function samples from T(x -> x')
    logpsi_x_proposed = logpsi(params, np.expand_dims(x_proposed, 0))[0]

    R = np.exp(2 * (logpsi_x_proposed - logpsi_x))
    u = np.random.uniform()
    accept = R > u

    if accept:
        return x_proposed, logpsi_x_proposed
    else:
        return x, logpsi_x

```

```

[18]: def sample_mcmc(logpsi, params, N, N_samples, N_discard, x0=None,
    ↪ propose_fn=single_spin_flip):
    """

```

Monte Carlo Markov Chains sampling. Given an ansatz, it samples randomly $\hookrightarrow N_{\text{samples}}$.

`logpsi`: function giving the log wave-funtion of a batch of sample given a $\hookrightarrow$ set of parametes

```
(params,x) -> log( (x))
params: parameters of the ansatz
N: number of sites
N_samples: number of samples to generate
N_discard: number of initial samples to discard (thermalization)
x0: initial configuration
      (if None, a random sample is drawn from the Hilbert space)
propose_fn: update move
"""

# Initialization
if x0 is None:
    x0 = random_states(N, 1)[0]

x = x0
logpsi_x = None

# Thermalization : we don't keep the samples
for i in range(N_discard):
    x, logpsi_x = sample_step(logpsi, params, x, logpsi_x, propose_fn)

# MCMC
samples = []
for i in range(N_samples):
    x, logpsi_x = sample_step(logpsi, params, x, logpsi_x, propose_fn)
    samples.append(x)

return np.vstack(samples)
```

```
[19]: samples_mcmc = sample_mcmc(logpsi_mf, params, N, Ns, N_discard)
      samples_mcmc
```

```
[19]: array([[ -1,  -1,   1, ...,  -1,   1,  -1],
             [ -1,   1,   1, ...,  -1,   1,  -1],
             [ -1,   1,   1, ...,  -1,   1,  -1],
             ...,
             [ -1,   1,  -1, ...,  -1,   1,  -1],
             [ -1,   1,  -1, ...,  -1,   1,  -1],
             [ -1,   1,  -1, ...,  -1,   1,  -1]])
```

```
[20]: assert samples_mcmc.shape == (Ns, N)
```

0.2.5 e) Connected Elements

```
[21]: def ising_hamiltonian(x,  $\Gamma$ =1, J=1):
        """
        TFI hamiltonian in 1D with pbc. Given a configuration x compute the all
        ↪connected x' and matrix elements Hxx' s.t. Hxx' != 0
        """

        # x is again a batch of samples:
        n_samples, n_sites = x.shape

        # there are n_sites + 1 connected states
        n_conn = n_sites + 1

        # initialize arrays
        x_prime = np.zeros((n_samples, n_conn, n_sites), dtype=x.dtype)
        mels = np.zeros((n_samples, n_conn))

        # states

        # diagonal: we take the first row to be the diagonal where x==x'
        # (this is arbitrary, we could have picked any other order)
        x_prime[:, 0] = x

        # off-diagonal:
        # the remaining rows are the off-diagonal terms
        # We have to flip one spin each, we do this in order
        flip_mask = np.eye(n_sites, dtype=int) # where to flip
        x_prime[:, 1:] = (1-flip_mask) * np.expand_dims(x, 1) + flip_mask * (- np.
        ↪expand_dims(x, 1))

        # matrix elements

        # diagonal
        mels[:, 0] = J * (x * np.roll(x, 1, axis=-1)).sum(axis=-1)

        # off-diagonal
        mels[:, 1:] = - $\Gamma$ 

        return x_prime, mels
```

```
[22]: xp, mels = ising_hamiltonian(x, 1, 1)
```

```
[23]: xp
```

```
[23]: array([[[-1, -1, 1, ..., 1, 1, -1],
              [ 1, -1, 1, ..., 1, 1, -1],
```

```

[-1, 1, 1, ..., 1, 1, -1],
...,
[-1, -1, 1, ..., -1, 1, -1],
[-1, -1, 1, ..., 1, -1, -1],
[-1, -1, 1, ..., 1, 1, 1]],

[[-1, 1, -1, ..., -1, -1, 1],
 [ 1, 1, -1, ..., -1, -1, 1],
 [-1, -1, -1, ..., -1, -1, 1],
 ...,
 [-1, 1, -1, ..., 1, -1, 1],
 [-1, 1, -1, ..., -1, 1, 1],
 [-1, 1, -1, ..., -1, -1, -1]],

[[ 1, -1, -1, ..., -1, 1, -1],
 [-1, -1, -1, ..., -1, 1, -1],
 [ 1, 1, -1, ..., -1, 1, -1],
 ...,
 [ 1, -1, -1, ..., 1, 1, -1],
 [ 1, -1, -1, ..., -1, -1, -1],
 [ 1, -1, -1, ..., -1, 1, 1]],

[[ 1, 1, -1, ..., -1, 1, 1],
 [-1, 1, -1, ..., -1, 1, 1],
 [ 1, -1, -1, ..., -1, 1, 1],
 ...,
 [ 1, 1, -1, ..., 1, 1, 1],
 [ 1, 1, -1, ..., -1, -1, 1],
 [ 1, 1, -1, ..., -1, 1, -1]],

[[-1, 1, 1, ..., 1, 1, 1],
 [ 1, 1, 1, ..., 1, 1, 1],
 [-1, -1, 1, ..., 1, 1, 1],
 ...,
 [-1, 1, 1, ..., -1, 1, 1],
 [-1, 1, 1, ..., 1, -1, 1],
 [-1, 1, 1, ..., 1, 1, -1]]])

```

```
[24]: mels
```

```

[24]: array([[ 0., -1., -1., -1., -1., -1., -1., -1., -1., -1., -1., -1., -1.,
        -1., -1., -1., -1.],
       [ 0., -1., -1., -1., -1., -1., -1., -1., -1., -1., -1., -1., -1.,
        -1., -1., -1., -1.],
       [-4., -1., -1., -1., -1., -1., -1., -1., -1., -1., -1., -1., -1.,
        -1., -1., -1., -1.],
       [ 0., -1., -1., -1., -1., -1., -1., -1., -1., -1., -1., -1., -1.,
        -1., -1., -1., -1.]])

```

```

-1., -1., -1., -1.],
[ 4., -1., -1., -1., -1., -1., -1., -1., -1., -1., -1., -1.,
-1., -1., -1., -1.]]

```

```

[25]: assert xp.shape == (len(x), N+1, N)
      assert mels.shape == (len(x), N+1)

```

0.2.6 f) Local operator

```

[26]: def compute_eloc(operator, logpsi, params, x):
      Ns, N = x.shape
      xp, mels = operator(x)
      logpsi_x = logpsi(params, x)
      # logpsi can only take batches of samples (with one single batch dimension)
      # so we have to flatten the input and unflatten the output
      logpsi_xp = logpsi(params, xp.reshape(-1, N)).reshape(xp.shape[:-1])
      eloc = (mels * np.exp(logpsi_xp - np.expand_dims(logpsi_x, -1))).sum(axis=-1)
      return eloc

```

```

[27]: eloc = compute_eloc(ising_hamiltonian, logpsi_mf, params, x)

```

```

[28]: assert eloc.shape == (len(x),)

```

0.2.7 g+h) Stochastic estimates

```

[29]: def expect(operator, logpsi, params, x):
      "compute the expectation value of an operator given a batch of samples"
      eloc = compute_eloc(operator, logpsi, params, x)
      E = eloc.mean()
      return E

```

```

[30]: expect(ising_hamiltonian, logpsi_mf, params, samples_direct)

```

```

[30]: -15.965010091574

```

```

[31]: expect(ising_hamiltonian, logpsi_mf, params, samples_mcmc)

```

```

[31]: -15.920906915118627

```

0.2.8 i) Gradient descent

```

[32]: def expect_and_grad(operator, logpsi, grad_logpsi, params, x):

      eloc = compute_eloc(operator, logpsi, params, x)
      E = eloc.mean()
      Dk = grad_logpsi(params, x)
      delta_eloc = eloc - eloc.mean(axis=0, keepdims=True)

```

```
Gk = 2 * np.expand_dims(delta_eloc, 1) * Dk
grad = Gk.mean(axis=0)
return E, grad
```

```
[33]: _, g = expect_and_grad(ising_hamiltonian, logpsi_mf, grad_logpsi_mf, params, x)
assert g.shape == params.shape
```

```
[34]: # we will pass this as opt_fn to the vmc function below
def sgd(params, grad, ):
    return params - * grad
```

```
[35]: def vmc(operator, sample_fn, opt_fn, logpsi, grad_logpsi, params, N, nsteps):

    energies = []

    for i in tqdm(range(nsteps)):

        # sample
        x = sample_fn(logpsi, params, N)
        # estimate energy and gradients
        E, grad = expect_and_grad(operator, logpsi, grad_logpsi, params, x)
        energies.append(E)
        # compute updated parameters
        params = opt_fn(params, grad)

    E = expect(operator, logpsi, params, x)
    energies.append(E)

    return params, energies
```

Mean-field Ansatz with direct sampling

```
[36]: # setup the direct sampler fixing the number of samples
sa = partial(sample_direct_mf, N_samples=Ns)

# set up the optimizer by fixing the learning rate
opt = partial(sgd, =0.01)

#set up the operator
ha = partial(ising_hamiltonian, Γ=1, J=1)

lpsi = logpsi_mf
glpsi = grad_logpsi_mf
# set up the initial guess for the parameters
par = random_params_mf(N, 0.1)

_, energies = vmc(ha, sa, opt, lpsi, glpsi, par, N, 1000)
plt.figure()
```

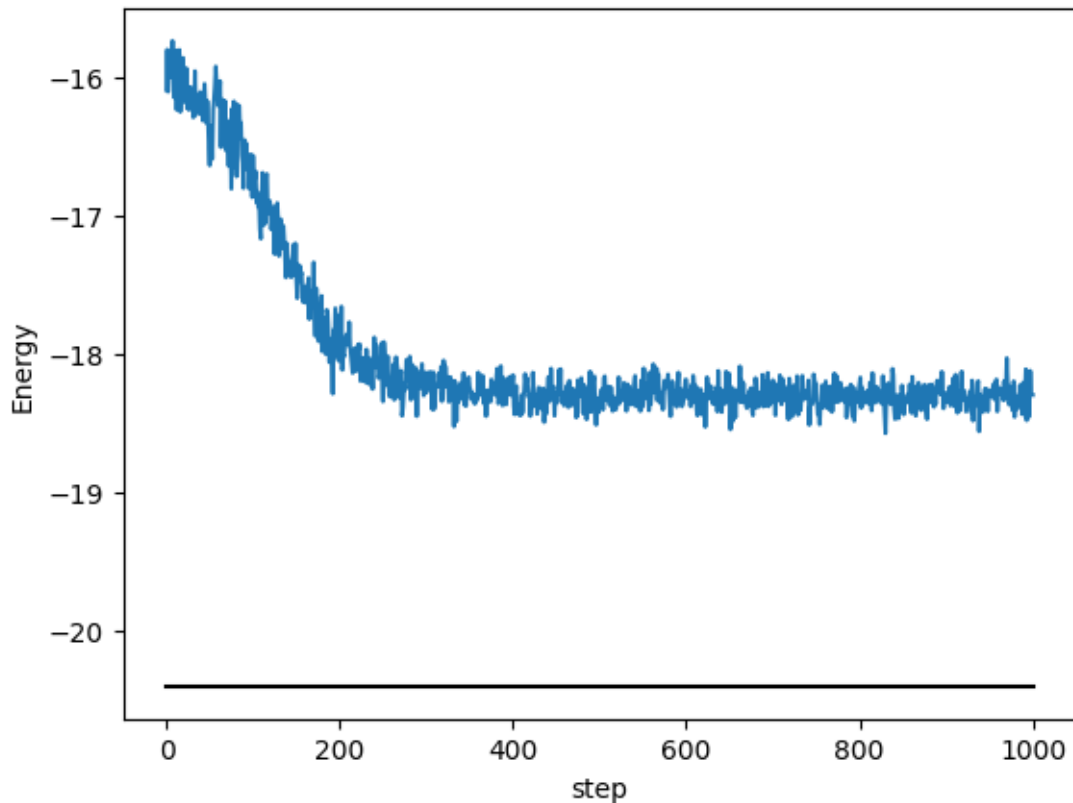
```

plt.plot(energies)
E0_analytical = ising1d_energy(N, 1)
plt.plot([0, len(energies)], [E0_analytical,]*2, color='k')
plt.xlabel('step')
plt.ylabel('Energy')
print('relative error: {:.2e}'.format(err_rel(E0_analytical, energies[-1])))
plt.show()

```

0%| | 0/1000 [00:00<?, ?it/s]

relative error: 1.15e-01



Mean-field with MCMC sampling

```

[37]: sa = partial(sample_mcmc, N_samples=Ns, N_discard=N_discard)
      opt = partial(sgd, =0.01)
      ha = partial(ising_hamiltonian, Γ=1, J=1)

      lpsi = logpsi_mf
      glpsi = grad_logpsi_mf
      par = random_params_mf(N, 0.1)

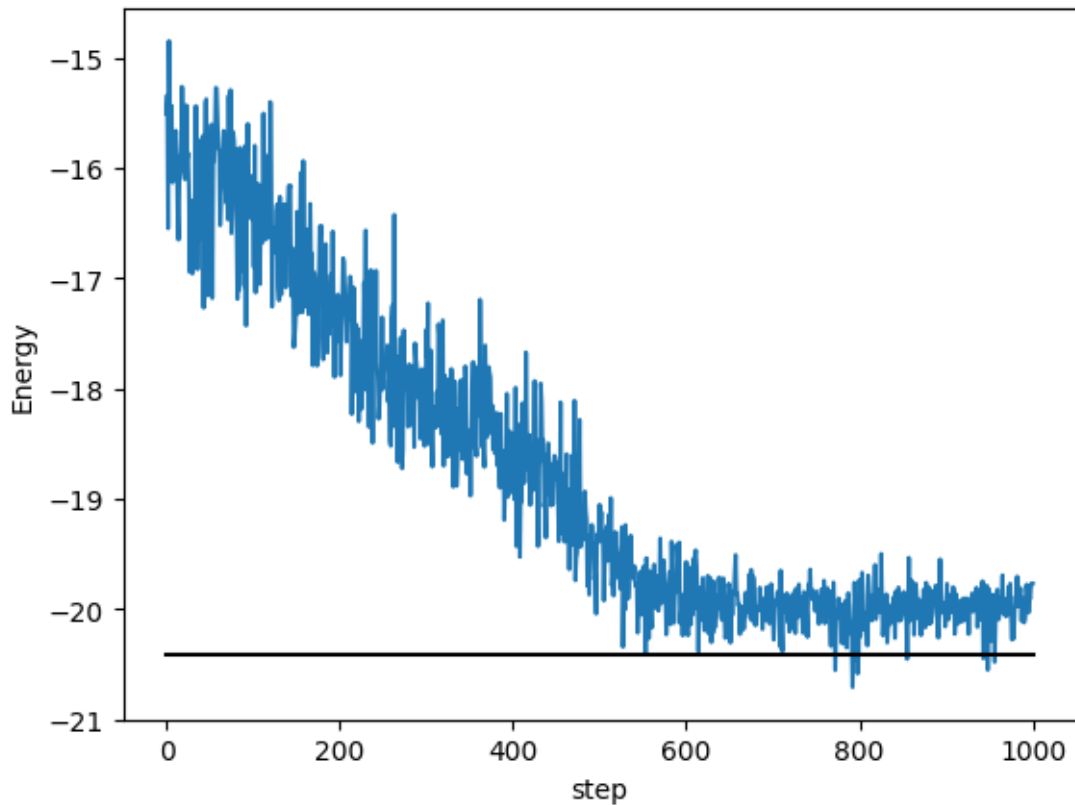
```

```
_, energies = vmc(ha, sa, opt, lpsi, glpsi, par, N, 1000)

plt.figure()
plt.plot(energies)
E0_analytical = ising1d_energy(N, 1)
plt.plot([0, len(energies)], [E0_analytical,]*2, color='k')
plt.xlabel('step')
plt.ylabel('Energy')
print('relative error: {:.2e}'.format(err_rel(E0_analytical, energies[-1])))
plt.show()
```

0%| | 0/1000 [00:00<?, ?it/s]

relative error: 3.22e-02

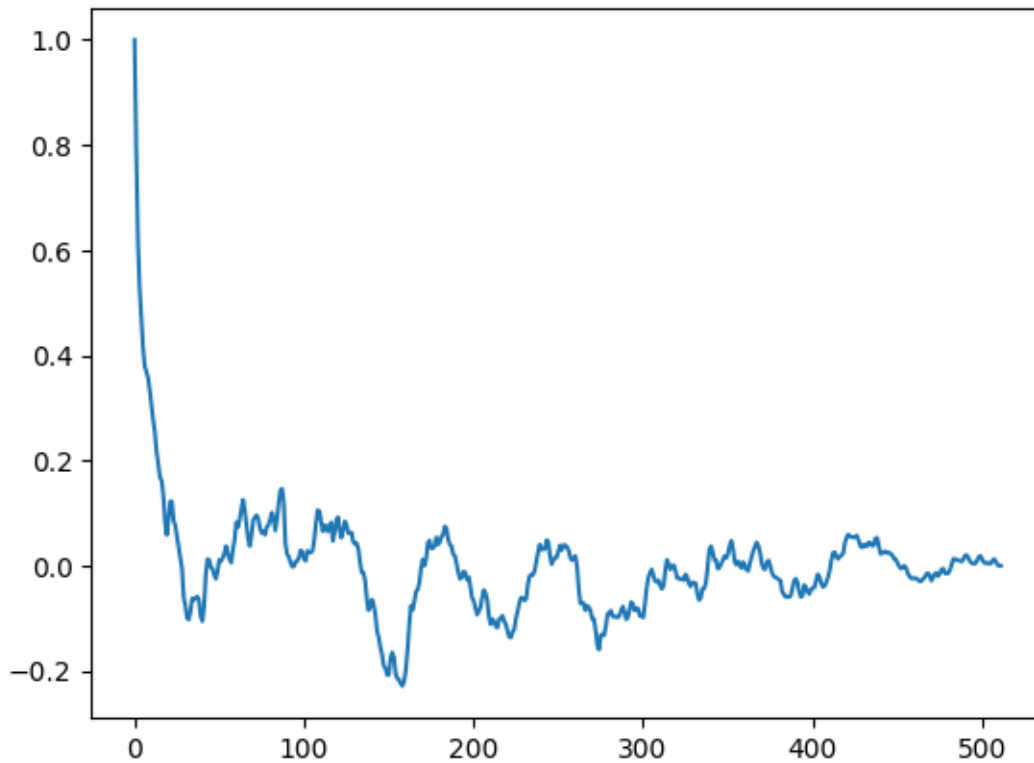


0.2.9 j) Auto-correlation time (bonus)

```
[38]: def corr_fn_fft(g):
      n = len(g)
      A = np.fft.rfft(g - g.mean(axis=0), n=2*n)
      B = A * A.conj() / len(g)
      return np.fft.irfft(B)[:n] / ((g**2).mean(axis=0) - g.mean(axis=0)**2)
```

```
[39]: par_mf = random_params_mf(N, 0.1)
      x = sample_mcmc(logpsi_mf, par_mf, N_samples=Ns, N=N, N_discard=N_discard)
      eloc = compute_eloc(ising_hamiltonian, logpsi_mf, par_mf, x)
```

```
[40]: = corr_fn_fft(eloc)
      plt.figure()
      plt.plot()
      plt.show()
```



```
[41]: jcut = np.argmin( > 0)
      = 0.5 + ([:jcut]).sum()
```

```
[41]: 8.19596157065558
```

0.3 Exercise 7.2

0.3.1 a) Nearest-neighbour Jastrow

```
[42]: def logpsi_jastrow_nearest(params, s):
    Ns, N = s.shape
    J1 = params
    return J1 * (s * np.roll(s, -1, axis=-1)).sum(axis=-1) # nearest neighbors
    ↪(pbc)

def grad_logpsi_jastrow_nearest(params, s):
    Ns, N = s.shape
    return (s * np.roll(s, -1, axis=-1)).sum(axis=-1, keepdims=True) # nearest
    ↪neighbors (pbc)

def random_params_jastrow_nearest(N, stddev=0.1):
    return np.random.normal(0, stddev, size=1)
```

```
[43]: # check
params_jastrow = random_params_jastrow_nearest(N, 0.1)
assert logpsi_jastrow_nearest(params_jastrow, x).shape == (len(x),)
assert grad_logpsi_jastrow_nearest(params_jastrow, x).shape ==
    ↪(len(x), len(params_jastrow))
```

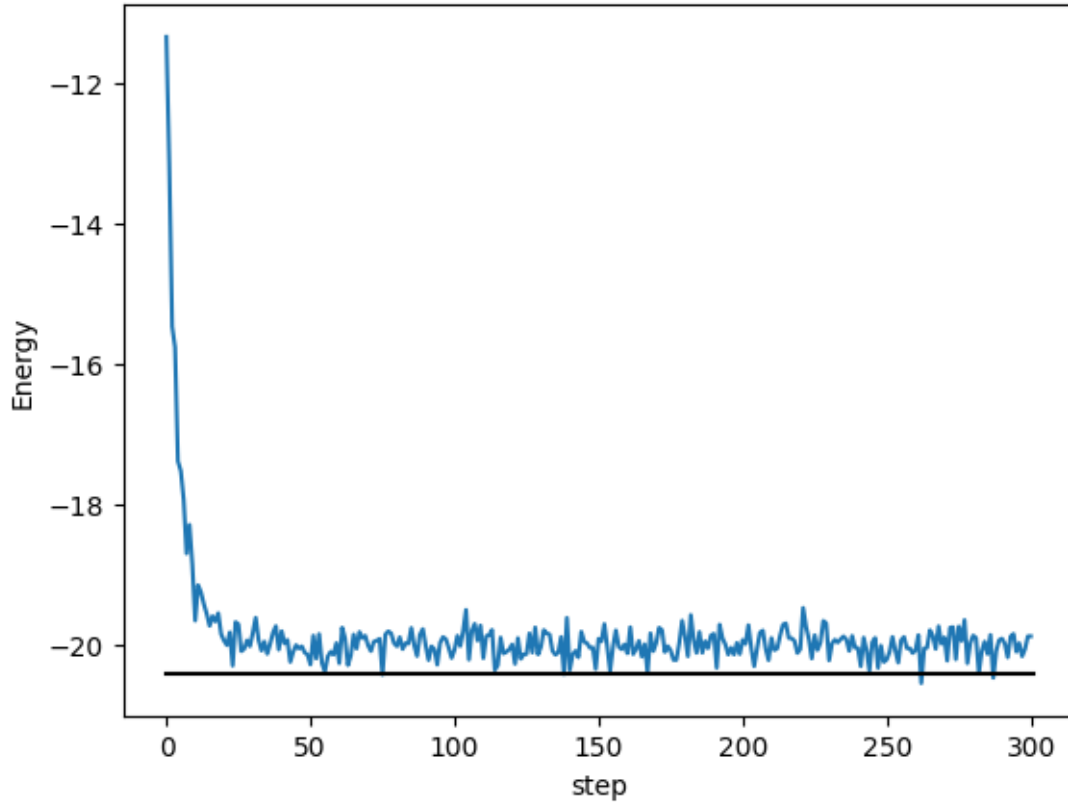
```
[44]: sa = partial(sample_mcmc, N_samples=Ns, N_discard=N_discard)
opt = partial(sgd, =0.001) # needs a fairly small learning rate
ha = partial(ising_hamiltonian, Γ=1, J=1)

lpsi = logpsi_jastrow_nearest
glpsi = grad_logpsi_jastrow_nearest
par = random_params_jastrow_nearest(N, 0.1)

_, energies = vmc(ha, sa, opt, lpsi, glpsi, par, N, 300)
plt.figure()
plt.plot(energies)
E0_analytical = ising1d_energy(N, 1)
plt.plot([0, len(energies)], [E0_analytical,]*2, color='k')
plt.xlabel('step')
plt.ylabel(('Energy'))
print('relative error: {:.2e}'.format(err_rel(E0_analytical, energies[-1])))
plt.show()
```

0%| | 0/300 [00:00<?, ?it/s]

relative error: 2.63e-02



0.3.2 b) Nearest+next-nearest-neighbour Jastrow

```
[45]: def logpsi_jastrow_next_nearest(params, s):
    Ns, N = s.shape
    J1 = params[0]
    J2 = params[1]
    res1 = logpsi_jastrow_nearest(J1, s)
    res2 = J2 * (s * np.roll(s, -2, axis=-1)).sum(axis=-1) # next-nearest
    ↪ neighbors (pbc)
    return res1+res2

def grad_logpsi_jastrow_next_nearest(params, s):
    Ns, N = s.shape
    J1 = params[0]
    # J2 = params[1]
    g1 = grad_logpsi_jastrow_nearest(J1, s)
    g2 = (s * np.roll(s, -2, axis=-1)).sum(axis=-1, keepdims=True) #
    ↪ next-nearest neighbors (pbc)
    return np.concatenate([g1,g2], axis=1)

def random_params_jastrow_next_nearest(N, stddev=0.1):
```

```
return np.random.normal(0, stddev, size=2)
```

```
[46]: sa = partial(sample_mcmc, N_samples=Ns, N_discard=N_discard)
      opt = partial(sgd, =0.001)
      ha = partial(ising_hamiltonian,  $\Gamma$ =1, J=1)

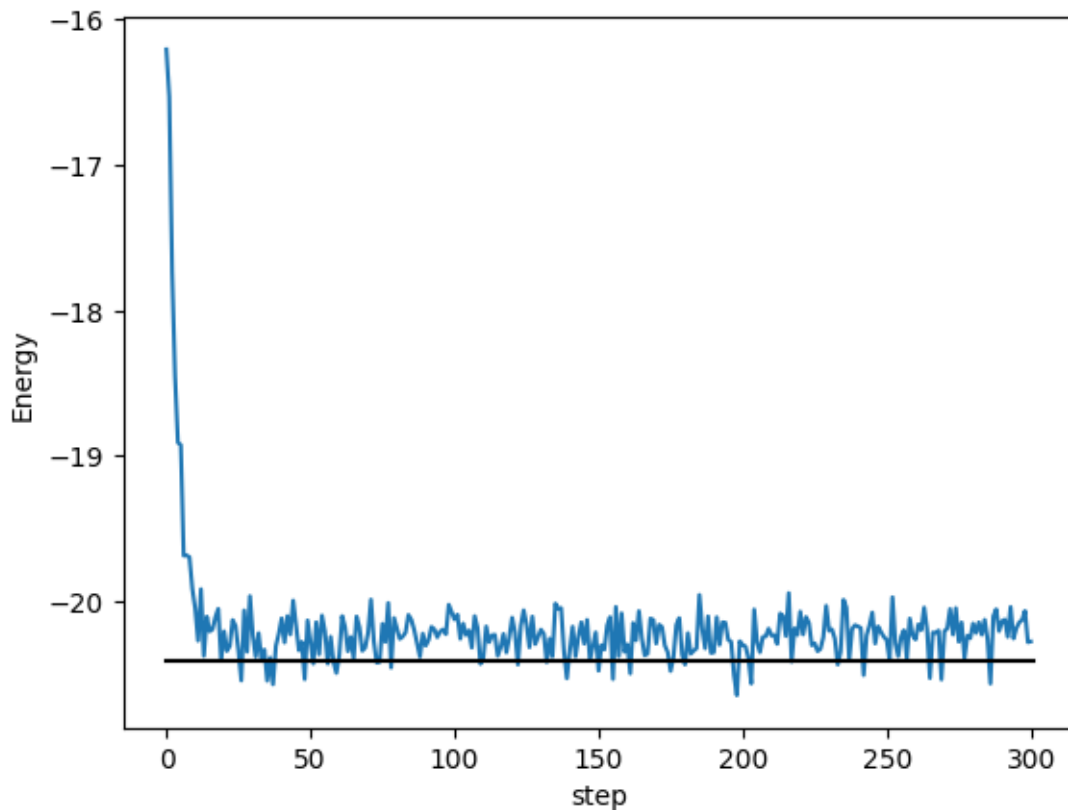
      lpsi = logpsi_jastrow_next_nearest
      glpsi = grad_logpsi_jastrow_next_nearest
      par = random_params_jastrow_next_nearest(N, 0.1)

      _, energies = vmc(ha, sa, opt, lpsi, glpsi, par, N, 300)

      plt.figure()
      plt.plot(energies)
      EO_analytical = ising1d_energy(N, 1)
      plt.plot([0, len(energies)], [EO_analytical,]*2, color='k')
      plt.xlabel('step')
      plt.ylabel(('Energy'))
      print('relative error: {:.0.2e}'.format(err_rel(EO_analytical, energies[-1])))
      plt.show()
```

0%| | 0/300 [00:00<?, ?it/s]

relative error: 6.54e-03



0.3.3 c) Arbitrary n-neighbours Jastrow

```
[47]: def logpsi_jastrow_n_neighbours(params, s):
    Ns, N = s.shape
    n = params.shape[0]

    res = 0
    for i in range(n):
        res += params[i]*(s * np.roll(s, -(i+1), axis=-1)).sum(axis=-1)

    return res

def grad_logpsi_jastrow_n_neighbours(params, s):
    Ns, N = s.shape
    n = params.shape[0]

    g = []
    for i in range(n):
        g.append( (s * np.roll(s, -(i+1), axis=-1)).sum(axis=-1, keepdims=True)
        ↪ )

    return np.concatenate(g, axis=1)

def random_params_jastrow_n_neighbours(N, n=1, stddev=0.1):
    assert n>=1 and n<N
    return np.random.normal(0, stddev, size=n)

[48]: sa = partial(sample_mcmc, N_samples=Ns, N_discard=N_discard)
    opt = partial(sgd, =0.001)
    ha = partial(ising_hamiltonian, Γ=1, J=1)

    E0_analytical = ising1d_energy(N, 1)

    E_converged = []
    delta_E = []
    for n in range(1, N//2+1): #<- we only go up to N//2 since we have pbc's
        lpsi = logpsi_jastrow_n_neighbours
        glpsi = grad_logpsi_jastrow_n_neighbours
        par = random_params_jastrow_n_neighbours(N, n, 0.1)

        _, energies = vmc(ha, sa, opt, lpsi, glpsi, par, N, 300)

        E_converged.append( energies[-1] )
        delta_E.append( err_rel(E0_analytical, energies[-1]) )
```

```
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```

```
[49]: plt.figure()
plt.plot(np.arange(1,N//2+1), E_converged)
plt.plot([0,N//2+1], [E0_analytical,E0_analytical])
plt.xlabel('Number of neighbours')
plt.ylabel('Final Energy')
plt.show()
```

