

**Student Name: Guangyu Lin**

**Collaboration Statement:**

Total hours spent: a week

I discussed ideas with these individuals:

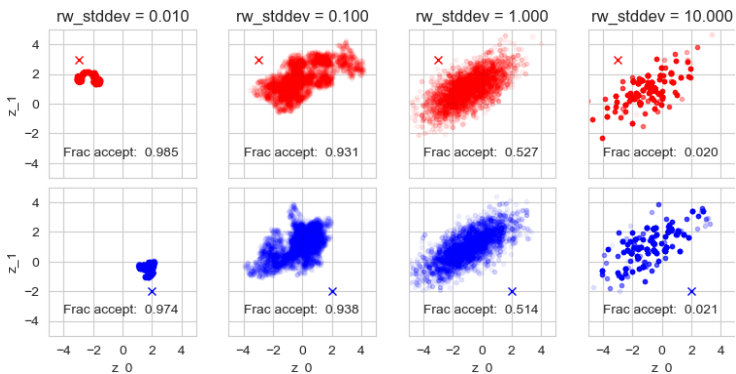
- Mingyang Wu
- TODO
- ...

I consulted the following resources:

- office hour with Professor and TA
- textbook
- ...

By submitting this assignment, I affirm this is my own original work that abides by the course collaboration policy.

# 1a

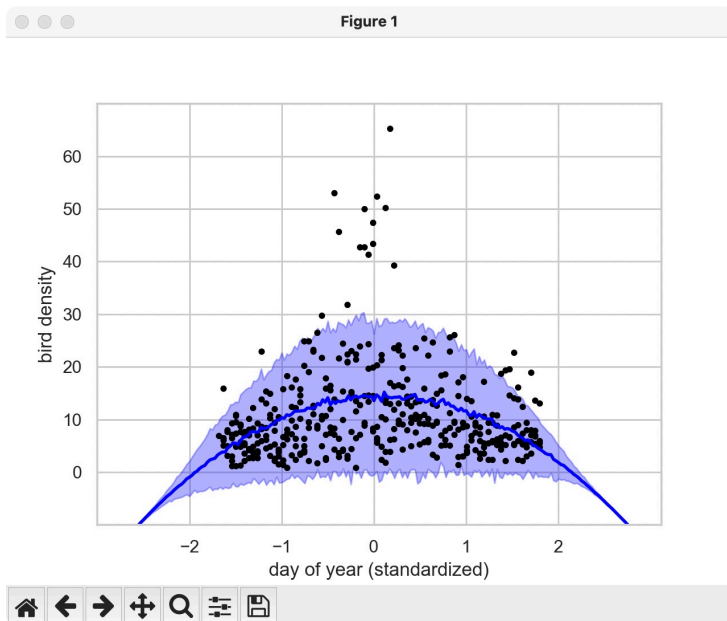


# 1b

No, we are not confident towards the MCMC chain has converged. There are two reasons. First, based on 1a, we know the graph is converged better when accept rate is 0.5 than 0.9. Therefore, we can't

guarantee when accept rate is 0.8, the MCMC chain has converged. Second, it only tried one trial and we can't guarantee it stated with a good position or not. If we can try more trials with different start points, we may have more confident to say it is converged.

## 2a



## 2b

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Order	test score
0	-4.513134
2	-4.218459

## 2c

---

```
1  def calc_score(list_of_z_D,  
2      phi_RM, t_R):  
3      ''' Calculate per-example  
4      score averaged over provided  
5      test set of size R  
6      Args  
7      ----  
8      list_of_z_D : list of  
9      ndarray
```

```

7         List of samples of
parameters, assumed to be from
target posterior
8         phi_RM : 2D array, shape (R,
M)
9         Feature vectors for each
of the examples in test set of
size R
10        t_R : 1D array, shape (R,)
11        Output values for each
of the examples in test set of
size R
12
13        Returns
14        -----
15        score : float
16        Per-example log pdf of
all t values in test set
17        using Monte-Carlo
approximation to marginal
likelihood
18        '''
19        S = len(list_of_z_D)

```

```

20     log_likelihoods = []
21     for ss in range(S):
22         z_ss_D = list_of_z_D[ss]
23         mean_R, stddev_R =
unpack_mean_N_and_stddev_N(z_ss_
D, phi_RM)
24         log_likelihoods.append(
25
        scipy.stats.norm.logpdf(t_R,
loc=mean_R, scale=stddev_R)
26     )
27     # Compute score formula
for ss-th sample (see
instructions)
28     # Hint: Use
unpack_mean_N_and_stddev_N
29     # TODO aggregate across all
S samples
30     # Hint: use
scipy.special.logsumexp to be
numerically stable
31     log_likelihoods =
np.asarray(log_likelihoods)

```

```
32     score =  
    np.mean(scipy.special.logsumexp(  
    log_likelihoods, axis = 0) -  
    np.log(S))  
33     return score # TODO FIXME
```