

# bayes\_models\_diagnostics

January 18, 2023

## 1 Overview

This is a python notebook to present all the model diagnostics of bayseian models used in Examining the Longitudinal Trajectory of Biopsychosocial Difficulties in Anorexia Nervosa Within a Bayesian Framework paper.

For more information on how the data was pre-processed and how the models where built please see [https://github.com/WMDA/BB\\_data/](https://github.com/WMDA/BB_data/)

### 1.1 Content

Section 1: Overview of notebook

Section 2: Libraries used in this note book and reading in the data/models .

Section 3: A Heatmap Plot

Section 4: KDE and trace plots for the mixed effects models

Section 5: Diagnostic Summaries of the mixed effects models

Section 6: Autocorrelation plots

## 2 Libraries used in this note book and reading in the data/models

Skip this section if just interested in model diagnostics

```
[12]: from functions.data_functions import save_pickle, load_pickle, load_data
import seaborn as sns
sns.set_style('darkgrid')
import matplotlib.pyplot as plt
import numpy as np
import arviz as az
```

```
[2]: models_summary = load_pickle('simpler_saved_models')
fitted_models = load_pickle('simpler_fitted_models')
```

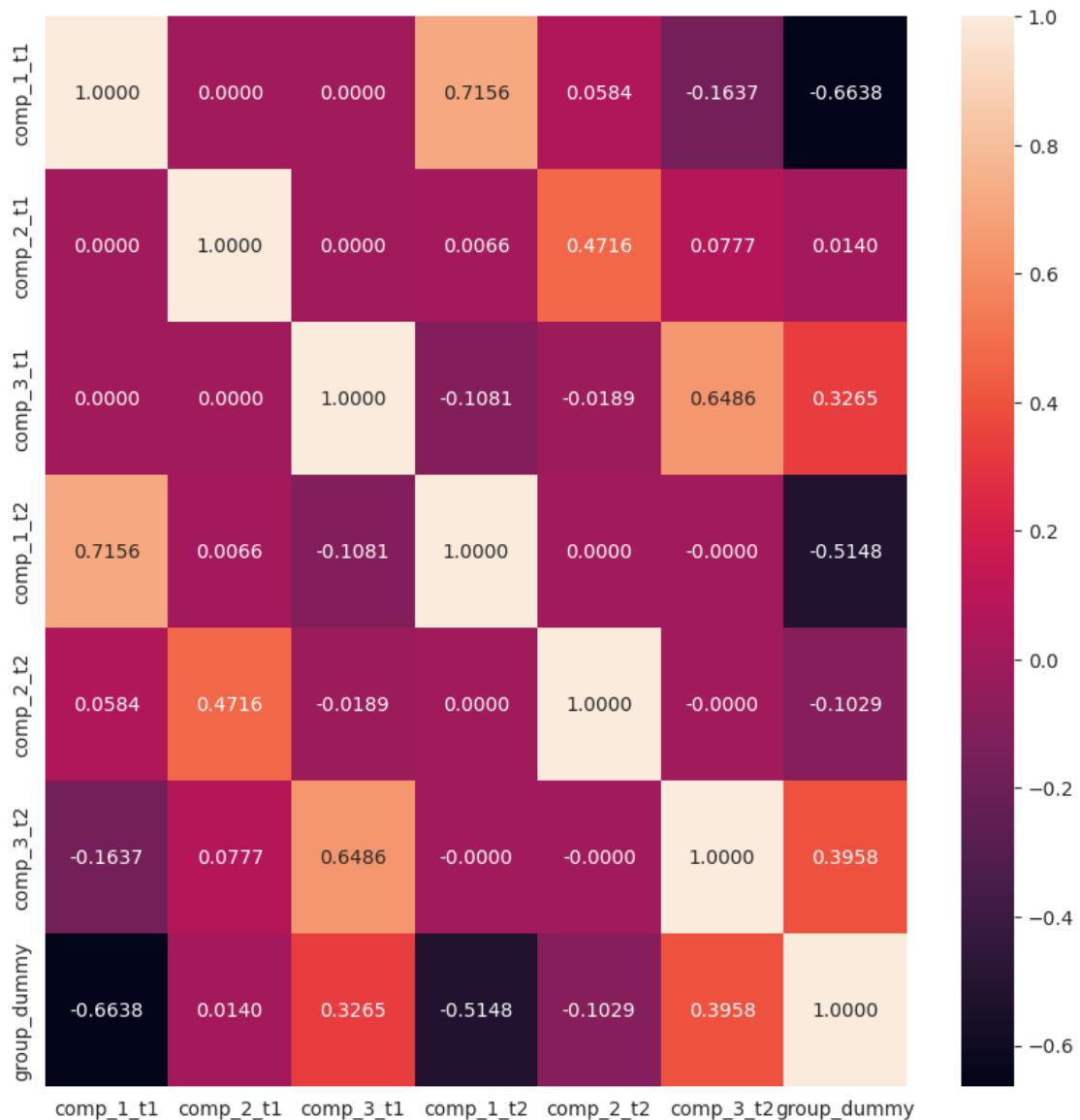
```
[3]: pca_df = load_data('BEACON', 'pca_df')
#Define number of components to loop over
comp = ['comp_1', 'comp_2','comp_3']
```

### 3 Heatmap

This is a heatmap of all the variables in the used in the models to check for multicollinearity

```
[4]: pca_df['group_dummy'] = pca_df['group'].apply(lambda group: 0 if group == 'AN' else 1)
plt.figure(figsize=(10, 10))
sns.heatmap(pca_df[['comp_1_t1', 'comp_2_t1', 'comp_3_t1', 'comp_1_t2', 'comp_2_t2', 'comp_3_t2', 'group_dummy']].corr(), annot=True, fmt=".4f")
```

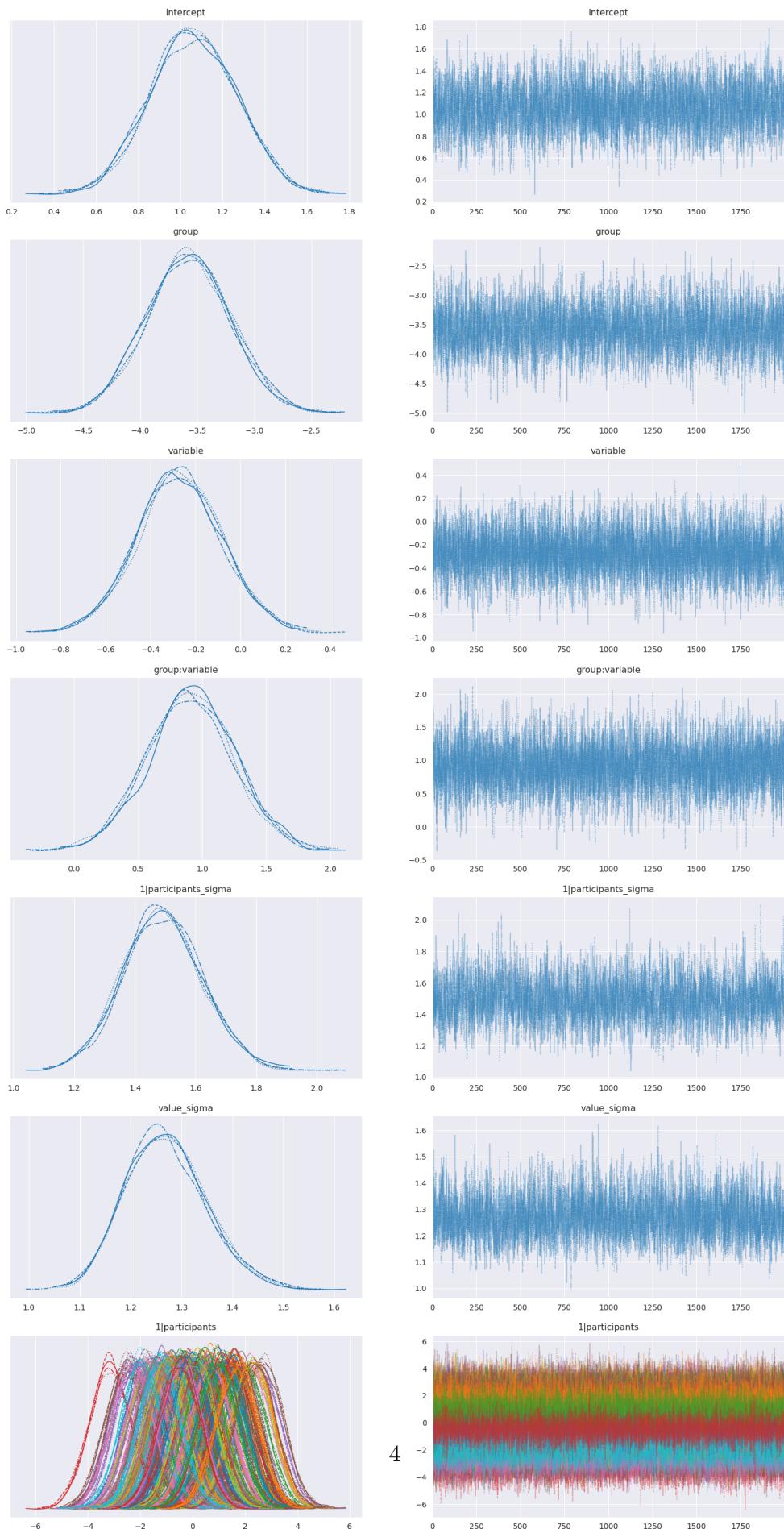
```
[4]: <AxesSubplot: >
```



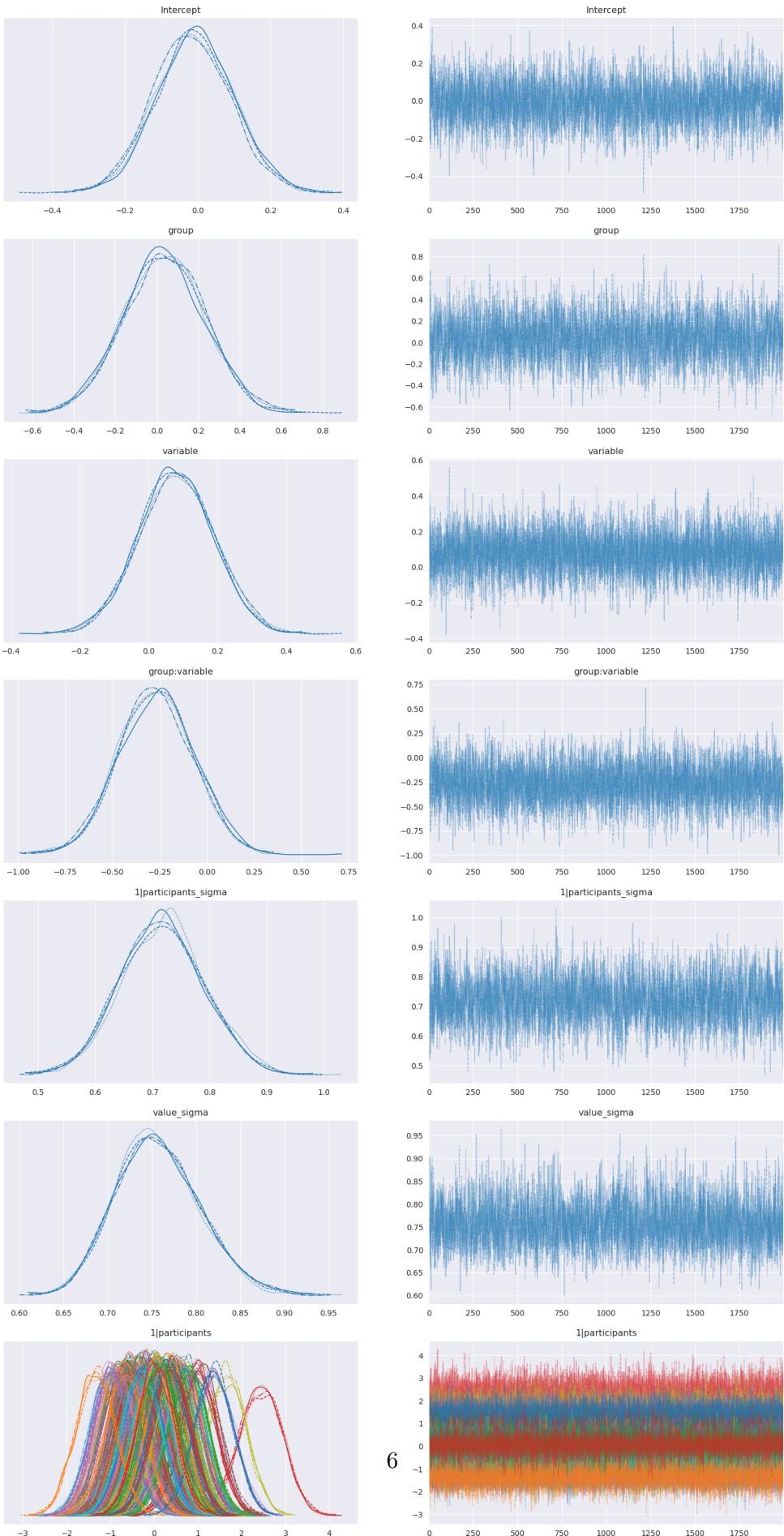
## 4 KDE and trace plots for the mixed effects models

### 4.1 Alternative hypothesis models

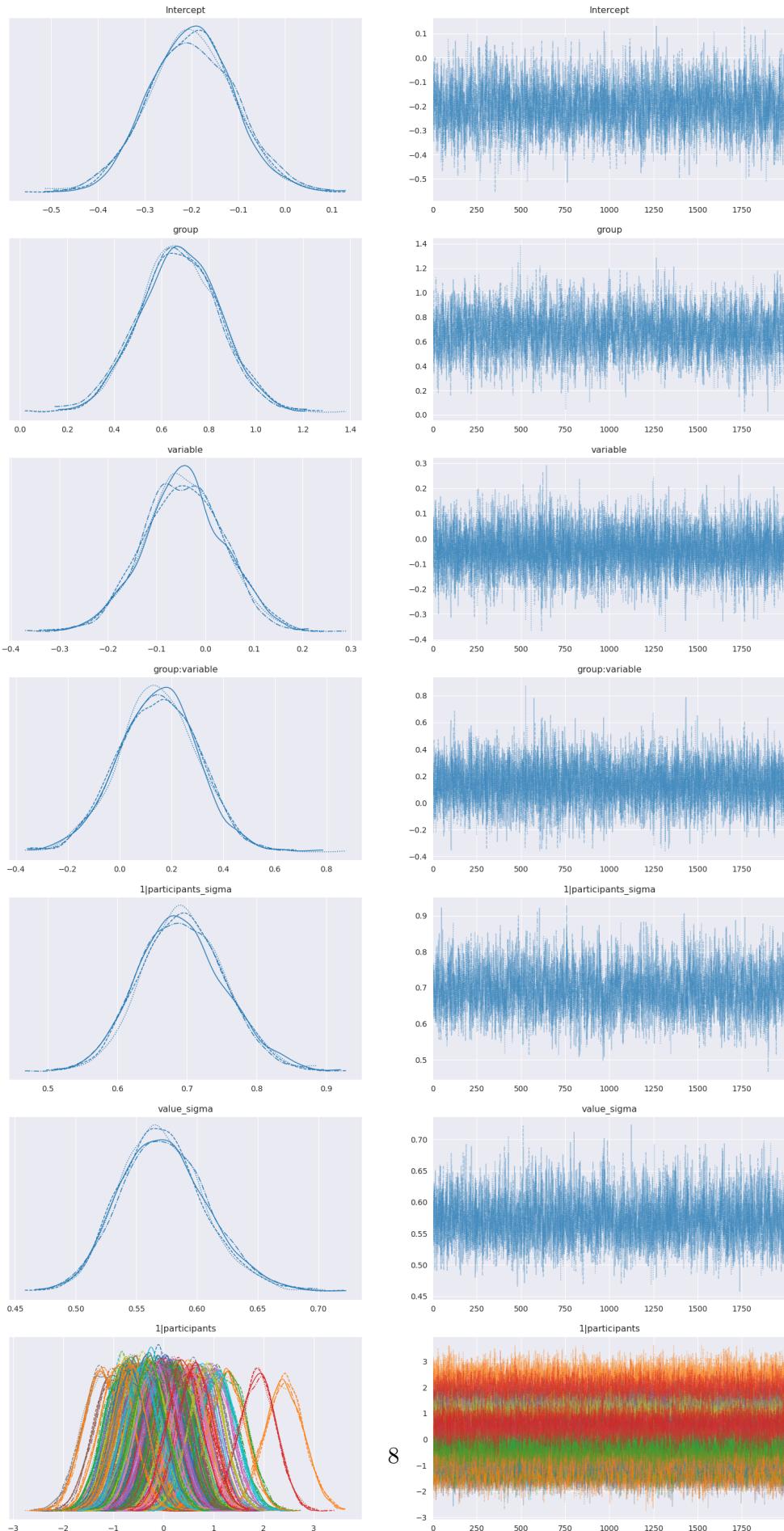
```
[5]: for component in comp:  
    az.plot_trace(fitted_models['alt'][component], figsize=(18,35))
```





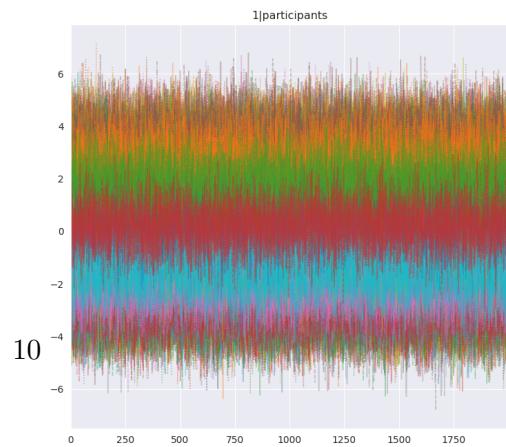
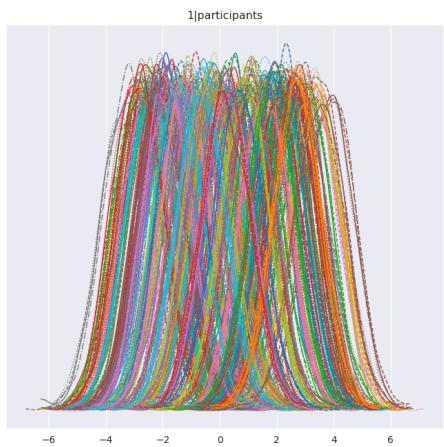
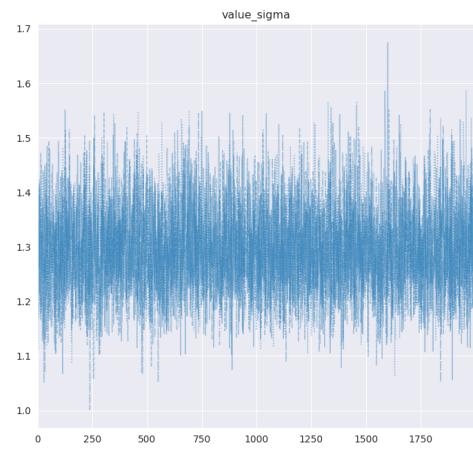
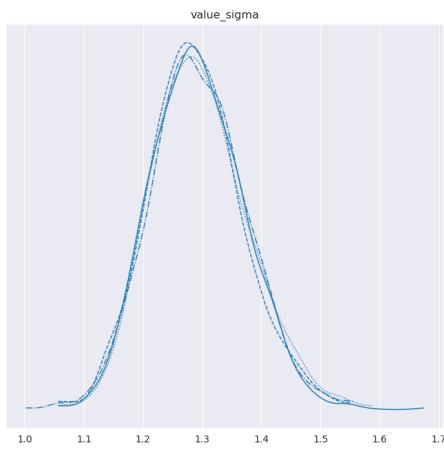
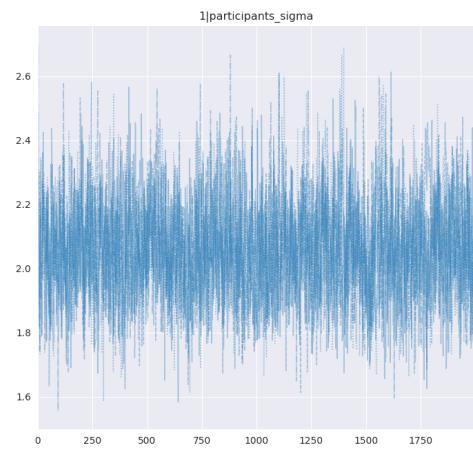
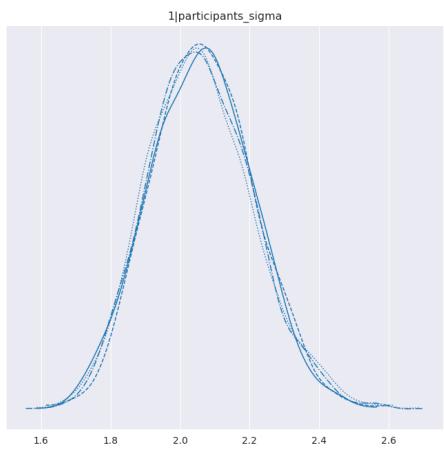
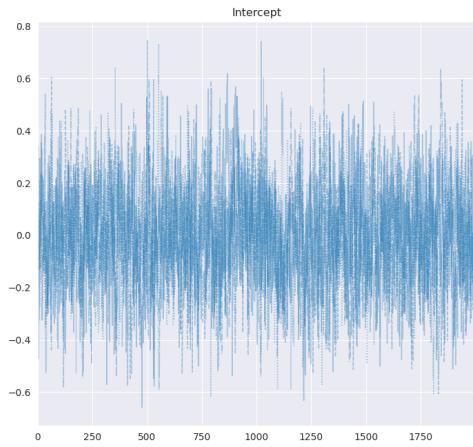
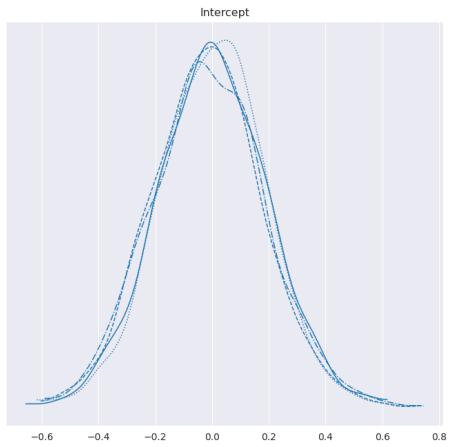




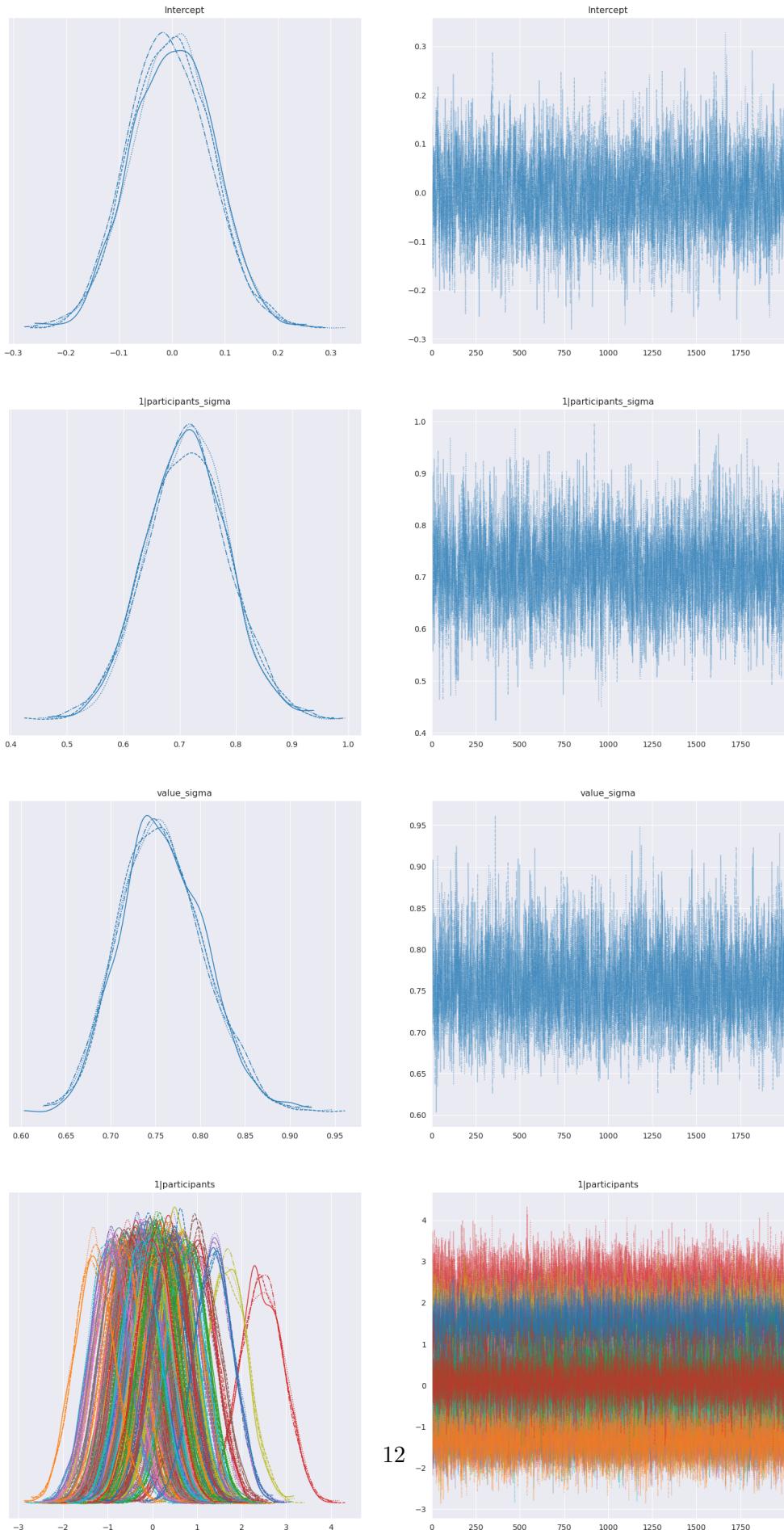


## 4.2 For the null hypothesis models

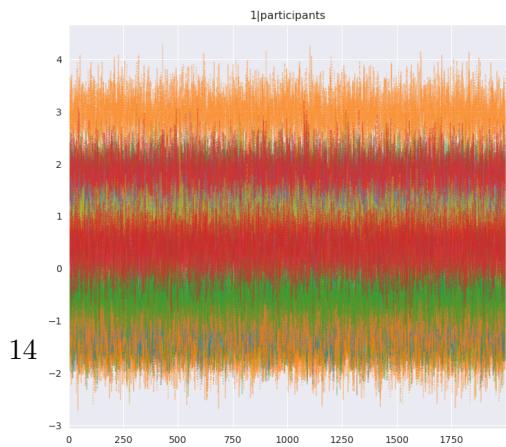
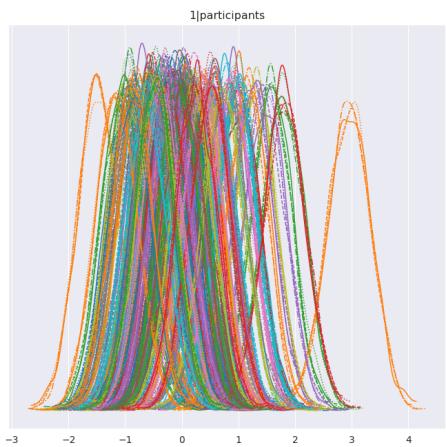
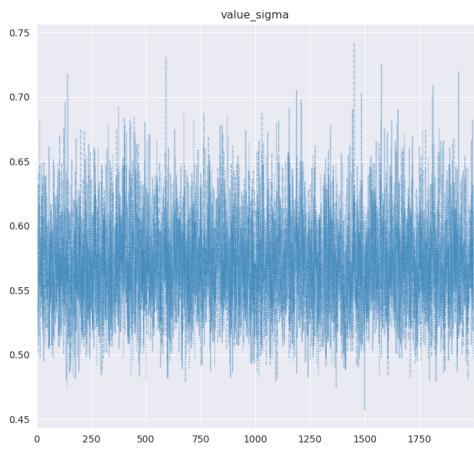
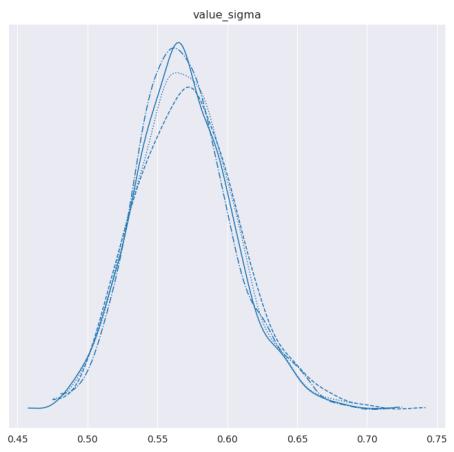
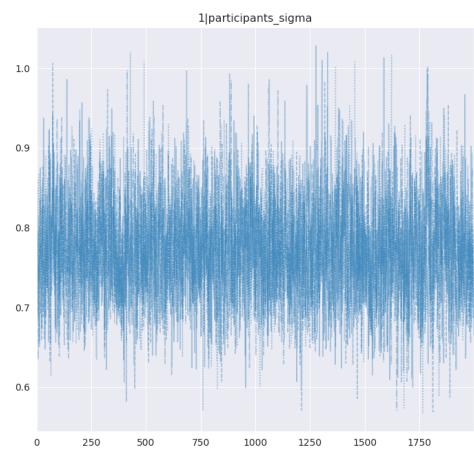
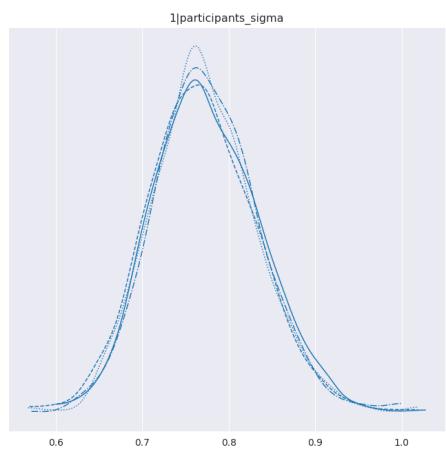
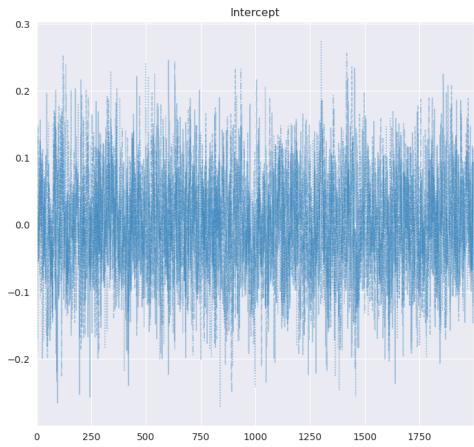
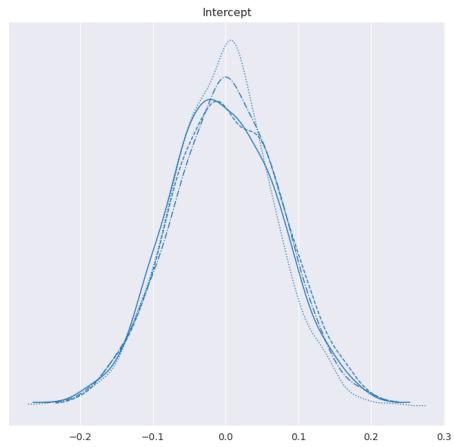
```
[6]: for component in comp:  
    az.plot_trace(fitted_models['null'][component], figsize=(18,35))
```











## 5 Diagnostic Summaries of the mixed effects models

### 5.1 Alternative Hypothesis

```
[7]: alt_variables = ['Intercept', 'group', 'group:variable', 'value_sigma',  
    ↴'1|participants_sigma']  
null_variables = ['value_sigma', '1|participants_sigma']
```

```
[8]: az.summary(fitted_models['alt']['comp_1'], kind='diagnostics',  
    ↴var_names=alt_variables )
```

```
[8]:
```

	mcse_mean	mcse_sd	ess_bulk	ess_tail	r_hat
Intercept	0.004	0.003	3194.0	4645.0	1.0
group[HC]	0.006	0.005	3311.0	5059.0	1.0
group:variable[HC, t2]	0.004	0.003	6509.0	5701.0	1.0
value_sigma	0.002	0.001	2697.0	4278.0	1.0
1 participants_sigma	0.003	0.002	2101.0	3993.0	1.0

```
[9]: az.summary(fitted_models['alt']['comp_2'], kind='diagnostics',  
    ↴var_names=alt_variables)
```

```
[9]:
```

	mcse_mean	mcse_sd	ess_bulk	ess_tail	r_hat
Intercept	0.002	0.001	2852.0	4230.0	1.0
group[HC]	0.004	0.003	2847.0	4499.0	1.0
group:variable[HC, t2]	0.003	0.002	5177.0	6231.0	1.0
value_sigma	0.001	0.001	2708.0	4788.0	1.0
1 participants_sigma	0.002	0.001	2341.0	4528.0	1.0

```
[10]: az.summary(fitted_models['alt']['comp_3'], kind='diagnostics',  
    ↴var_names=alt_variables)
```

```
[10]:
```

	mcse_mean	mcse_sd	ess_bulk	ess_tail	r_hat
Intercept	0.002	0.001	2229.0	3852.0	1.0
group[HC]	0.003	0.002	2557.0	3974.0	1.0
group:variable[HC, t2]	0.002	0.002	5459.0	5436.0	1.0
value_sigma	0.001	0.000	2866.0	5044.0	1.0
1 participants_sigma	0.001	0.001	2333.0	4097.0	1.0

Extracting the number of divergences, tree depth, energy steps and acceptance rate.

```
[13]: for component in comp:  
    print(f'Number of divergences for {component} alt model:', int(np.  
        ↴sum(fitted_models['alt'][component].sample_stats.diverging)))
```

```

    print(f'Number of times tree depth hit 10 for {component} alt model:', int(np.sum(fitted_models['alt'][component].sample_stats.tree_depth == 10)))
    print(f'Number of energy steps > 0.3 for {component} alt model:', int(np.sum(fitted_models['alt'][component].sample_stats.energy < 0.3)))
    print(f'Number of steps where acceptance rate < 0.95 for {component} alt model:', int(np.sum(fitted_models['alt'][component].sample_stats.acceptance_rate < 0.95)))
    print('\n\n')

```

Number of divergences for comp\_1 alt model: 0  
Number of times tree depth hit 10 for comp\_1 alt model: 0  
Number of energy steps > 0.3 for comp\_1 alt model: 0  
Number of steps where acceptance rate < 0.95 for comp\_1 alt model: 3074

Number of divergences for comp\_2 alt model: 0  
Number of times tree depth hit 10 for comp\_2 alt model: 0  
Number of energy steps > 0.3 for comp\_2 alt model: 0  
Number of steps where acceptance rate < 0.95 for comp\_2 alt model: 3213

Number of divergences for comp\_3 alt model: 0  
Number of times tree depth hit 10 for comp\_3 alt model: 0  
Number of energy steps > 0.3 for comp\_3 alt model: 0  
Number of steps where acceptance rate < 0.95 for comp\_3 alt model: 3186

## 5.2 Null Hypothesis

[14]: az.summary(fitted\_models['null']['comp\_1'], kind='diagnostics', var\_names=null\_variables)

	mcse_mean	mcse_sd	ess_bulk	ess_tail	r_hat
value_sigma	0.001	0.001	3209.0	4753.0	1.0
1 participants_sigma	0.004	0.003	1427.0	2394.0	1.0

[15]: az.summary(fitted\_models['null']['comp\_2'], kind='diagnostics', var\_names=null\_variables)

	mcse_mean	mcse_sd	ess_bulk	ess_tail	r_hat
value_sigma	0.001	0.001	2821.0	4512.0	1.0
1 participants_sigma	0.002	0.001	2411.0	4181.0	1.0

```
[16]: az.summary(fitted_models['null']['comp_3'], kind='diagnostics',  
    ↪var_names=null_variables)
```

```
[16]:
```

	mcse_mean	mcse_sd	ess_bulk	ess_tail	r_hat
value_sigma	0.001	0.000	3160.0	4755.0	1.0
1 participants_sigma	0.001	0.001	2353.0	4323.0	1.0

Extracting the number of divergences, tree depth, energy steps and acceptance rate.

```
[17]: for component in comp:  
    print(f'Number of divergences for {component} alt model:', int(np.  
        ↪sum(fitted_models['null'][component].sample_stats.diverging)))  
    print(f'Number of times tree depth hit 10 for {component} alt model:',  
        ↪int(np.sum(fitted_models['null'][component].sample_stats.tree_depth == 10)))  
    print(f'Number of energy steps > 0.3 for {component} alt model:', int(np.  
        ↪sum(fitted_models['null'][component].sample_stats.energy < 0.3)))  
    print(f'Number of steps where acceptance rate < 0.95 for {component} alt  
        ↪model:', int(np.sum(fitted_models['null'][component].sample_stats.  
        ↪acceptance_rate < 0.95)))  
    print('\n\n')
```

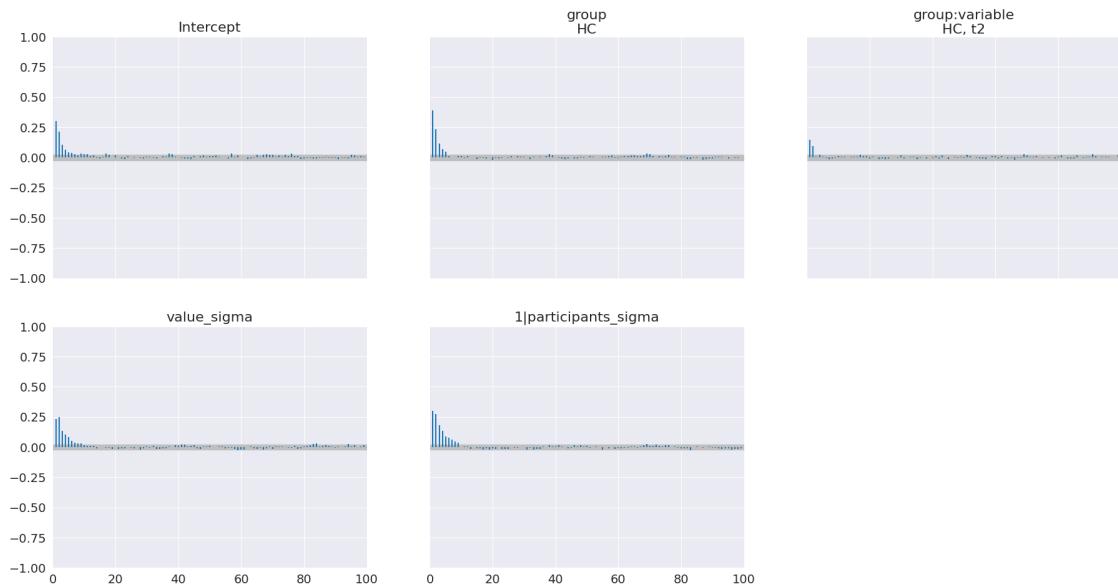
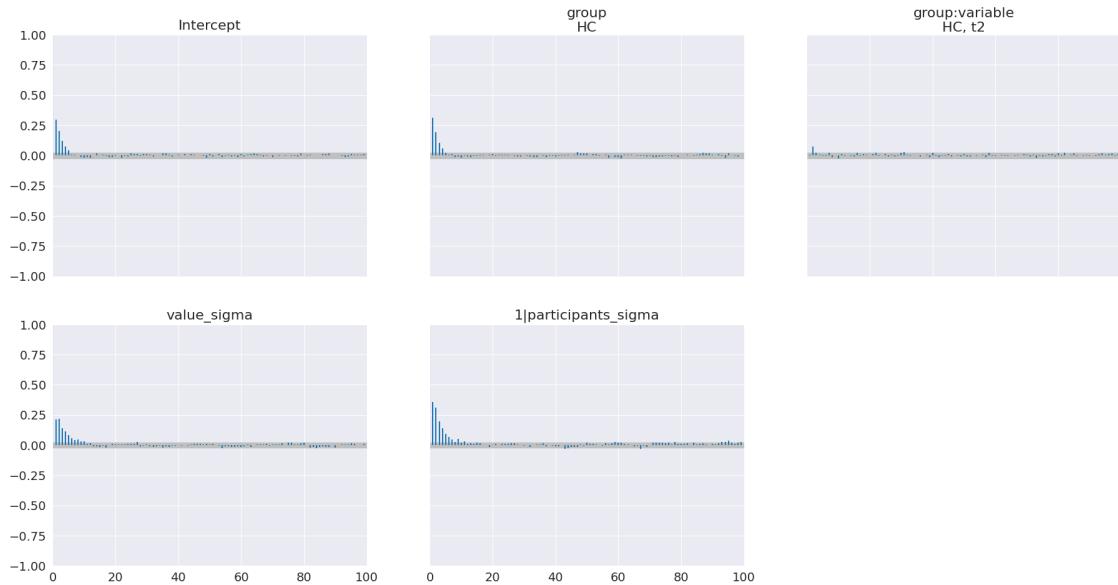
```
Number of divergences for comp_1 alt model: 0  
Number of times tree depth hit 10 for comp_1 alt model: 0  
Number of energy steps > 0.3 for comp_1 alt model: 0  
Number of steps where acceptance rate < 0.95 for comp_1 alt model: 3319
```

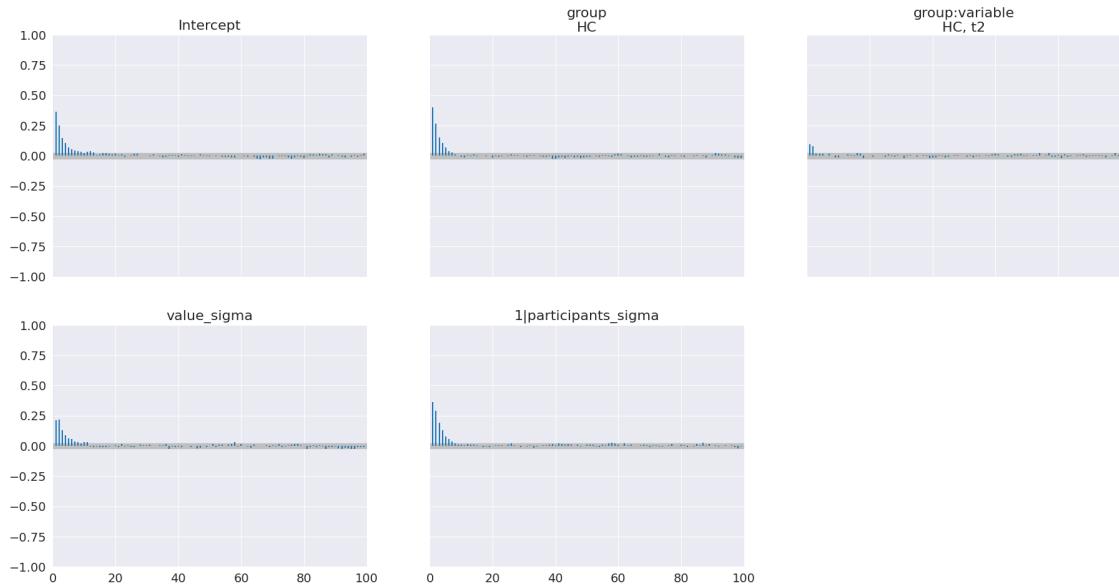
```
Number of divergences for comp_2 alt model: 0  
Number of times tree depth hit 10 for comp_2 alt model: 0  
Number of energy steps > 0.3 for comp_2 alt model: 0  
Number of steps where acceptance rate < 0.95 for comp_2 alt model: 3123
```

```
Number of divergences for comp_3 alt model: 0  
Number of times tree depth hit 10 for comp_3 alt model: 0  
Number of energy steps > 0.3 for comp_3 alt model: 0  
Number of steps where acceptance rate < 0.95 for comp_3 alt model: 3150
```

## 6 Autocorrelation plots

```
[ ]: for component in comp:  
    az.plot_autocorr(fitted_models['alt'][component], combined=True,  
                     var_names=alt_variables)
```





```
[ ]: for component in comp:
    az.plot_autocorr(fitted_models['null'][component], combined=True, □
    ↵var_names=null_variables)
```

