approximate_bayesian_computation

```
cm_name: abc_30
dataframe in: data missing 30
description: Approximate Bayesian Computation for Time Series
diff_func_name: manhattan_metrics
diff_func_parameters: {}
model_method: approximate_bayesian_computation
name: approximate_bayesian_computation
parameters:
  algorithm: pydream
  decision_variables:
    keys:
    - max_keys
  decision_variables_names:
  - graph_structure
  epsilons:
  - 1
  ground_truth_topology:
    keys:
     - max_keys
  initial_points: 100
  n_chains: 3
  n draws: 15000
  n iterations: 100
  nfe: 15000
  num_pool: 1
  population_size: 100
  seed: 26
report_parameters: {}
running_time: 184095.6848502159
type: calibrationmodel
version: 1.0.0
```

Results

```
Summary CalibrationModel with solutions:
    graph structure Distance
0
     3.159753e+04 48.153855
1
     3.156725e+04 45.008375
2
     3.153696e+04 33.114240
3
     1.022303e+04 17.615843
4
     1.022303e+04 18.066367
24795 1.603224e-10 18.154785
24796
       1.605184e-10 17.961186
24797
       1.607144e-10 18.183463
24798
       1.609104e-10 19.795292
24799
       1.611064e-10 18.615566
```

with the most optimal solution:
graph_structure Distance round
0 0.0 16.669574 0.0
with an acceptance percentage of 31.216507682387213%

approximate_bayesian_computation

```
cm_name: abc_40
dataframe in: data missing 40
description: Approximate Bayesian Computation for Time Series
diff_func_name: manhattan_metrics
diff_func_parameters: {}
model_method: approximate_bayesian_computation
name: approximate_bayesian_computation
parameters:
  algorithm: pydream
  decision_variables:
    keys:
    - max_keys
  decision_variables_names:
  - graph_structure
  epsilons:
  - 1
  ground_truth_topology:
    keys:
     - max_keys
  initial_points: 100
  n_chains: 3
  n draws: 15000
  n iterations: 100
  nfe: 15000
  num_pool: 1
  population_size: 100
  seed: 26
report_parameters: {}
running_time: 185621.85700821877
type: calibrationmodel
version: 1.0.0
```

Results

```
Summary CalibrationModel with solutions:
    graph_structure Distance
      31597.533350 46.183660
0
1
      31567.248932 43.055642
2
      31536.964515 32.143851
3
      10223.034214 18.708822
4
        0.000000 18.689580
21576
          0.195521 19.527954
21577
          0.195668 19.655112
21578
          0.195815 19.051053
21579
          0.195962 18.745749
21580
          0.196110 18.314104
```

with the most optimal solution:
graph_structure Distance round
0 0.0 17.294976 0.0
with an acceptance percentage of 27.06290440931225%

Summary

Model Name	Model Method	Score	Difference Function	Dataframe	Duration
abc_40	approximate_bayesian_computation	0.96	manhattan_metrics	data_missing_40	185621.857 sec
abc_30	approximate_bayesian_computation	0.96	manhattan_metrics	data_missing_30	184095.685 sec