approximate_bayesian_computation

```
cm_name: abc_70
dataframe in: data missing 70
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: 21
report_parameters: {}
running_time: 186576.00573682785
type: calibrationmodel
version: 1.0.0
```

Results

```
Summary CalibrationModel with solutions:
    graph structure Distance
0
      2782.768614 20.698156
1
      2782.768614 20.001156
2
        0.000000 19.994814
3
        0.000000 18.900103
4
        0.000000 19.946007
10399
          0.000000 18.973322
10400
          0.000000 19.267233
10401
          0.000000 18.522132
10402
          0.000000 18.579473
10403
          0.000000 19.317698
```

with the most optimal solution:
graph_structure Distance round
0 0.0 17.157248 0.0
with an acceptance percentage of 0.011117781780179222%

approximate_bayesian_computation

```
cm_name: abc_80
dataframe in: data missing 80
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: 21
report_parameters: {}
running_time: 186609.7592880726
type: calibrationmodel
version: 1.0.0
```

Results

```
Summary CalibrationModel with solutions:
   graph structure Distance
0
      2782.768614 21.192830
1
      2782.768614 20.578201
2
       0.000000 20.439641
3
       0.000000 19.378192
4
       0.000000 19.185049
8984
         0.000000 19.672666
8985
         0.000000 19.038400
8986
         0.000000 19.646849
8987
         0.000000 20.847256
8988
         0.000000 22.601109
```

with the most optimal solution:
graph_structure Distance round
0 0.0 18.177476 0.0
with an acceptance percentage of 0.011117781780179222%

Summary

Model Name	Model Method	Score	Difference Function	Dataframe	Duration
abc_80	approximate_bayesian_computation	0.96	manhattan_metrics	data_missing_80	186609.759 sec
abc_70	approximate_bayesian_computation	0.96	manhattan_metrics	data_missing_70	186576.006 sec