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: 16
report_parameters: {}
running_time: 294854.8771636486
type: calibrationmodel
version: 1.0.0
```

Results

```
Summary CalibrationModel with solutions:
   graph structure Distance
0
     22546.691528 33.126559
1
     22546.691528 31.594954
2
     22546.691528 27.285313
3
     22546.691528 27.474114
4
     22546.691528 27.312240
9047 34589.479087 17.496591
9048
     34589.479087 17.382516
9049
      34589.479087 17.495341
9050
      34589.479087 16.789311
9051
      34589.479087 16.938975
```

with the most optimal solution:
graph_structure Distance round
34589.479087 15.81305 34589.0
with an acceptance percentage of 0.008894225424143375%

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: 16
report_parameters: {}
running_time: 351222.3374669552
type: calibrationmodel
version: 1.0.0
```

Results

```
Summary CalibrationModel with solutions:
    graph_structure Distance
0
     22546.691528 31.985115
1
     22546.691528 32.103357
2
     22546.691528 31.146270
3
     22546.691528 29.244925
4
     22546.691528 29.744518
12433 34589.479087 17.601385
12434 34589.479087 17.287015
12435 34589.479087 17.357551
12436
       34589.479087 17.667397
12437
       34589.479087 17.680063
```

with the most optimal solution:
graph_structure Distance round
34589.479087 16.299042 34589.0
with an acceptance percentage of 0.008894225424143375%

Summary

Model Name	Model Method	Score	Difference Function	Dataframe	Duration
abc_40	approximate_bayesian_computation	0.98	manhattan_metrics	data_missing_40	351222.337 sec
abc_30	approximate_bayesian_computation	0.98	manhattan_metrics	data_missing_30	294854.877 sec