## compare

November 12, 2017

## 1 Reproducibility between tools

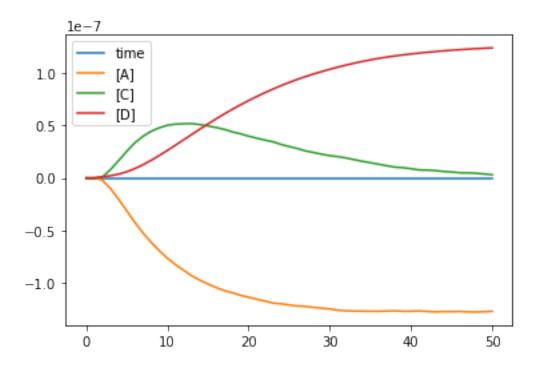
The following scripts checks that solutions are reproducibile between tools. Reproducibility of the model simulations was tested by comparing the numerical SOA results between sbmlutils and iBioSim for models with unique solutions. Results were assumed as numerical identical if the absolute difference for every time point  $t_k$  for all dynamical FBA species  $c_k$  in the model was smaller than the tolerance  $\epsilon = 1E-3$ , i.e.,

```
abs(c_i(t_k)_{shmlutils} - c_i(t_k)_{ibiosim}) \le \epsilon \ \forall c_i, t_k
In [31]: """
         Helper class for comparing simulation results.
         import pandas as pd
         from matplotlib import pyplot as plt
         from pprint import pprint
         import warnings
         class DataSetsComparison(object):
              """ Comparing two simulation results.
              Currently only supports comparison between two datasets.
              eps = 1E-3 # tolerance for comparison
              def __init__(self, files, dfs, columns=None):
                  self.files = files
                  self.columns = columns
                  for df in dfs:
                      # check that identical number of timepoints
                      if len(df) != len(dfs[0]):
                          raise ValueError("DataFrames have different length: \
                                             {} != {}".format(len(df), len(dfs[0])))
                  if columns:
                      assert len(self.files) == len(self.columns)
                      for column in self.columns:
                          assert len(column) == len(self.columns[0])
```

```
self.read_dfs(dfs)
    self.diff = self.df_diff()
def read_dfs(self, dfs):
    """ Read the dataframes using the files and given column ids."""
    self.dfs = []
    for k, df in enumerate(dfs):
        file = self.files[k]
        if self.columns:
            cols = self.columns[k]
            try:
                df1 = df[cols]
            except KeyError:
                pprint(df.columns)
                raise
            df1.columns = self.columns[0] # unify columns
            self.dfs.append(df1)
        else:
            # no columns specified, necessary to figure out the mapping
            print("-"*40)
            print(file)
            print("-"*40)
            pprint(df.columns)
    return self.dfs
def df diff(self):
    """ DataFrame of all differences between the files."""
    return self.dfs[0]-self.dfs[1]
def is_equal(self):
    """ Check if DataFrames are identical within numerical tolerance."""
    return abs(self.diff.abs().max().max()) <= DataSetsComparison.eps</pre>
def info(self):
    pprint(self.files)
    pprint(self.columns)
def print_diff(self):
    print("\n# Elements")
    print(self.diff.shape)
    print("\n# Maximum column difference")
    print(self.diff.abs().max())
```

```
print(self.diff.abs().max().max())
                print("\n# Datasets are equal (diff <= eps={})".format(self.eps))</pre>
                print(self.is_equal())
                if not (self.is_equal()):
                    warnings.warn("Datasets are not equal !")
            def plot_diff(self):
                for cid in self.diff.columns:
                    plt.plot(self.diff[cid], label=cid)
                plt.legend()
                plt.show()
            def report(self):
                print("*" * 80)
                self.info()
                self.print_diff()
                print("*" * 80)
                self.plot_diff()
                self.diff
1.1 toy_wholecell
In [32]: wholecell_version = 14
        files = [
                "./toy_wholecell/toy_wholecell_mk_v14-sbmlutils_dt1.0_tend50.0.csv",
                "./toy_wholecell/toy_wholecell_mk_v14-ibiosim_dt1.0_tend50.0.csv"
        wholecell_dsc = DataSetsComparison(
            files=files,
            dfs=[pd.read_csv(file) for file in files],
            columns = [
                 ["time", "[A]", "[C]", "[D]"],
                ["time", "A", "C", "D"],
        )
        wholecell_dsc.report()
********************************
['./toy_wholecell/toy_wholecell_mk_v14-sbmlutils_dt1.0_tend50.0.csv',
 './toy_wholecell/toy_wholecell_mk_v14-ibiosim_dt1.0_tend50.0.csv']
[['time', '[A]', '[C]', '[D]'], ['time', 'A', 'C', 'D']]
# Elements
(51, 4)
```

print("\n# Maximum element difference")



## 1.2 toy\_atp

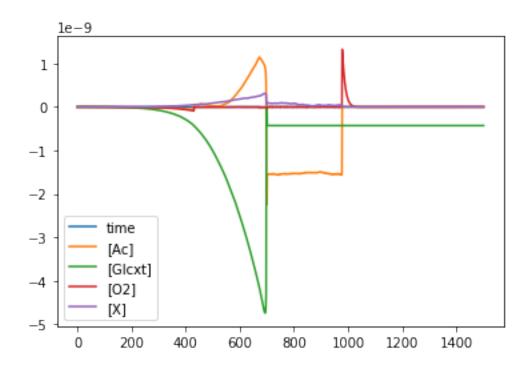
```
pd.read_csv(files[0], sep="\t"),
             pd.read_csv(files[1])
         ],
         columns = [
             ["time", "[adp]", "[atp]", "[pyr]", "[glc]"],
             ["time", "adp", "atp", "pyr", "glc"],
         ]
     )
     atp_dsc.report()
                                               Traceback (most recent call last)
    ValueError
    <ipython-input-33-e7957bbc5a3f> in <module>()
            columns = [
                ["time", "[adp]", "[atp]", "[pyr]", "[glc]"],
     12
---> 13
                ["time", "adp", "atp", "pyr", "glc"],
     14
            ]
     15 )
    <ipython-input-31-afecd26f9d49> in __init__(self, files, dfs, columns)
                    # check that identical number of timepoints
     19
     20
                    if len(df) != len(dfs[0]):
---> 21
                        raise ValueError("DataFrames have different length:
     22
     23
                if columns:
```

ValueError: DataFrames have different length:

## 1.3 diauxic

16 != 151

```
["time", "[Ac]", "[Glcxt]", "[02]", "[X]"],
               ["time", "Ac", "Glcxt", "O2", "X"],
           ]
        )
        diauxic_dsc.report()
*******************************
['./diauxic_growth/diauxic_growth_lw_v4-sbmlutils_dt0.01_tend15.0.csv',
'./diauxic_growth/diauxic_growth_lw_v4-ibiosim_dt0.01_tend15.0.csv']
[['time', '[Ac]', '[Glcxt]', '[O2]', '[X]'], ['time', 'Ac', 'Glcxt', 'O2', 'X']]
# Elements
(1501, 5)
# Maximum column difference
time
         3.552714e-15
[Ac]
         2.245718e-09
[Glcxt]
         4.743860e-09
Γ027
         1.326935e-09
[X]
         3.114851e-10
dtype: float64
# Maximum element difference
4.74386013805e-09
# Datasets are equal (diff <= eps=0.001)</pre>
True
*******************************
```



In []: