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 $\epsilon = 1E-5$, i.e.,

```
abs(c_i(t_k)_{sbmlutils} - c_i(t_k)_{ibiosim}) \le \epsilon \ \forall c_i, t_k
```

```
In [1]: """
        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.
            HHHH
            eps = 1E-5 # 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])
```

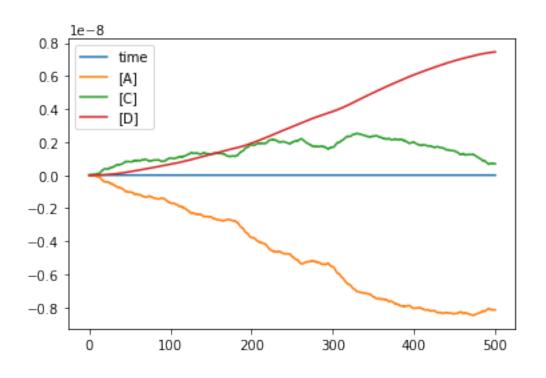
```
self.read_dfs(dfs)
    self.diff = self.df_diff()
def read_dfs(self, dfs):
    """ Read the dataframes using the files and given column ids."""
    self.dfs = []
    for k, df in enumerate(dfs):
        file = self.files[k]
        if self.columns:
            cols = self.columns[k]
            try:
                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("\n# Maximum element difference")
    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
```

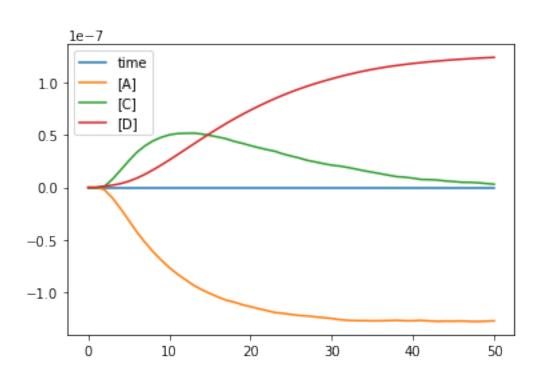
1.1 toy_wholecell

```
In [2]: wholecell_version = 14
        filesets = [
                "./toy_wholecell/toy_wholecell_mk_v14-sbmlutils_dt0.1_tend50.0.csv",
                "./toy_wholecell_toy_wholecell_mk_v14-ibiosim_dt0.1_tend50.0.csv"
            ],
            Γ
                "./toy_wholecell/toy_wholecell_mk_v14-sbmlutils_dt1.0_tend50.0.csv",
                "./toy_wholecell/toy_wholecell_mk_v14-ibiosim_dt1.0_tend50.0.csv"
            ],
                "./toy_wholecell/toy_wholecell_mk_v14-sbmlutils_dt5.0_tend50.0.csv",
                "./toy_wholecell_toy_wholecell_mk_v14-ibiosim_dt5.0_tend50.0.csv"
            ],
        1
        for files in filesets:
            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_mk_v14-sbmlutils_dt0.1_tend50.0.csv',
'./toy_wholecell_mk_v14-ibiosim_dt0.1_tend50.0.csv']
[['time', '[A]', '[C]', '[D]'], ['time', 'A', 'C', 'D']]
# Elements
(501, 4)
# Maximum column difference
      1.421085e-14
\lceil A \rceil
      8.488962e-09
[C]
      2.533890e-09
ΓD٦
      7.456576e-09
dtype: float64
# Maximum element difference
8.48896153371e-09
# Datasets are equal (diff <= eps=1e-05)</pre>
******************************
```



```
*******************************
['./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)
# Maximum column difference
       0.000000e+00
time
\lceil A \rceil
       1.280189e-07
[C]
       5.191503e-08
[D]
       1.242827e-07
dtype: float64
# Maximum element difference
1.28018868395e-07
# Datasets are equal (diff <= eps=1e-05)</pre>
True
```



```
************************
```

```
['./toy_wholecell/toy_wholecell_mk_v14-sbmlutils_dt5.0_tend50.0.csv',
   './toy_wholecell/toy_wholecell_mk_v14-ibiosim_dt5.0_tend50.0.csv']
[['time', '[A]', '[C]', '[D]'], ['time', 'A', 'C', 'D']]
```

Elements

(11, 4)

Maximum column difference

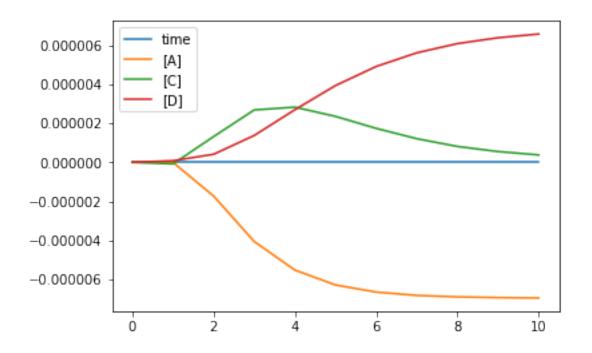
time 0.000000 [A] 0.000007 [C] 0.000003 [D] 0.000007 dtype: float64

Maximum element difference

6.95619353408e-06

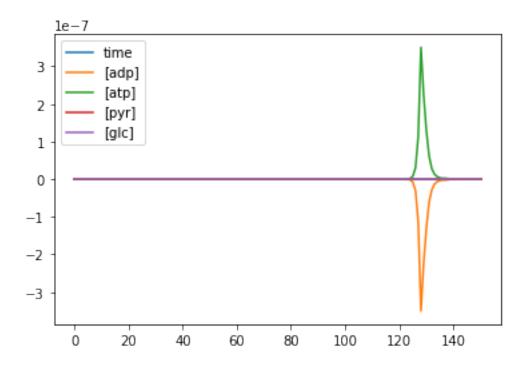
Datasets are equal (diff <= eps=1e-05)</pre>

True



1.2 toy_atp

```
In [3]: files = [
           "./toy_atp/toy_atp_mk_v12-sbmlutils_dt0.1_tend15.csv",
           "./toy_atp/toy_atp_mk_v12-ibiosim_dt0.1_tend15.csv"
       atp_dsc = DataSetsComparison(
          files=files,
          dfs = [
              pd.read_csv(files[0], sep="\t"),
              pd.read_csv(files[1])
          ],
           columns = [
              ["time", "[adp]", "[atp]", "[pyr]", "[glc]"],
              ["time", "adp", "atp", "pyr", "glc"],
          1
       )
       atp_dsc.report()
********************************
['./toy_atp/toy_atp_mk_v12-sbmlutils_dt0.1_tend15.csv',
'./toy_atp/toy_atp_mk_v12-ibiosim_dt0.1_tend15.csv']
[['time', '[adp]', '[atp]', '[pyr]', '[glc]'],
['time', 'adp', 'atp', 'pyr', 'glc']]
# Elements
(151, 5)
# Maximum column difference
time
       1.776357e-15
       3.505205e-07
[adp]
[atp]
      3.505205e-07
[pyr]
       4.263256e-14
[glc]
      1.781908e-14
dtype: float64
# Maximum element difference
3.50520538227e-07
# Datasets are equal (diff <= eps=1e-05)</pre>
**********************************
```



1.3 diauxic

```
In [4]: files = [
           "./diauxic_growth/diauxic_growth_lw_v4-sbmlutils_dt0.01_tend15.0.csv",
           "./diauxic_growth/diauxic_growth_lw_v4-ibiosim_dt0.01_tend15.0.csv",
       diauxic_dsc = DataSetsComparison(
           files=files,
           dfs = [
               pd.read_csv(files[0]),
               pd.read_csv(files[1])
           ],
           columns = [
               ["time", "[Ac]", "[Glcxt]", "[O2]", "[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]', '[02]', '[X]'], ['time', 'Ac', 'Glcxt', '02', 'X']]
```

```
# Elements (1501, 5)
```

Maximum column difference

time 3.552714e-15
[Ac] 2.245718e-09
[Glcxt] 4.743860e-09
[02] 1.326935e-09
[X] 3.114851e-10

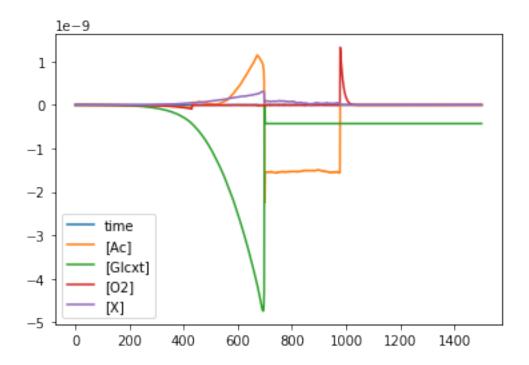
dtype: float64

Maximum element difference

4.74386013805e-09

Datasets are equal (diff <= eps=1e-05)</pre>

True



In []: