PyVIN, Pyomo based optimization model

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This document describes the modeling structure of PyVIN, a new version of <u>CALVIN</u> modeled on <u>Pyomo</u> (Python Optimization modeling Objects). Pyomo supports the formulation and analysis of mathematical models for optimization applications using algebraic modeling language format (AMPL).

PyVIN separates model structure from the data. So, the core of the model does not depend on data. Similar to CALVIN, PyVIN is a linear programming model, and currently uses GLPK as solver. The objective is to minimize total cost on each arc, subject to physical and regulatory constraints.

Objective function:

$$minimize Z = \sum_{i} \sum_{j} \sum_{k} c_{ijk} X_{ijk}$$

Subject to

$$\sum_{i} \sum_{k} X_{jik} = \sum_{i} \sum_{k} a_{ijk} X_{ijk}$$
$$X_{ijk} \le u_{ijk}$$
$$X_{ijk} \ge l_{ijk}$$

where Z is total cost of flows throughout the networks, X_{ijk} flow on the kth arc leaving node i toward node j; c_{ijk} economic costs or loss of benefits; a_{ijk} gains/losses on flows in arc ijk; u_{ijk} upper bound on arc ijk; and l_{ijk} lower bound on arc ijk.

A simple Pyomo model consists of

Set

Set data used to define a model instance

Param

Parameter used to define a model instance

Var

Decision variable

Objective

Function that is minimized or maximized

Constraint

Expressions that impose restrictions on variable values

The Core Structure of the Model (pyvin.py)

First three lines are Python packages. First line ensures that division arguments are converted to floating number before the division. The second line makes symbols used by Pyomo known to Python.

```
from __future__ import division
from pyomo.environ import *
import itertools
```

The following line declares that the model will be an abstract model

```
model = AbstractModel()
```

Declare set of nodes defined in data file

```
# Nodes in the network
model.N = Set()
```

Declare set of links within possible combination of nodes (N*N) and number of piecewise links (k)

```
# Network arcs
model.k = Set()
model.A = Set(within=model.N*model.N*model.k)
```

Declare model parameters. For 'source' and 'sink' nodes, mass balance constraint (flow rule) is not calculated. Other parameters are upper bound, lower bound, amplitude, and cost.

```
# Source node
model.source = Param(within=model.N)
# Sink node
model.sink = Param(within=model.N)
# Flow capacity limits
model.u = Param(model.A)
# Flow lower bound
model.l = Param(model.A)
# Link amplitude (gain/loss)
model.a = Param(model.A)
# Link cost
model.c = Param(model.A)
```

Declare decision variables, flows on each arch. A decision variable should be a positive real number.

```
# The flow over each arc
model.X = Var(model.A, within=NonNegativeReals)
```

Objective function to be minimized. In this model, the objective is to minimize total cost, which is the sum of flow on each arc multiplied by unit cost.

```
# Minimize total cost
def total_rule(model):
    return sum(model.c[i,j,k]*model.X[i,j,k] for (i,j,k) in model.A)
model.total = Objective(rule=total rule, sense=minimize)
```

Define upper bound constraint, such as reservoir storage capacity, canal conveyance capacity or pumping capacity

```
# Enforce an upper bound limit on the flow across each arc
def limit_rule_upper(model, i, j, k):
    return model.X[i,j,k] <= model.u[i,j,k]
model.limit upper = Constraint(model.A, rule=limit rule upper)</pre>
```

Define lower bound constraint. Lower bound constraints usually represent minimum in-stream flow requirements.

```
# Enforce an upper bound limit on the flow across each arc
def limit_rule_upper(model, i, j, k):
    return model.X[i,j,k] <= model.u[i,j,k]
model.limit_upper = Constraint(model.A, rule=limit_rule_upper)</pre>
```

The following lines are used to speed up the mass balance constraint

```
arcs_in = {}
arcs_out = {}

def arc_list_hack(model, i,j,k):
    if j not in arcs_in:
        arcs_in[j] = []
    arcs_in[j].append((i,j,k))

    if i not in arcs_out:
        arcs_out[i] = []
    arcs_out[i].append((i,j,k))
    return [0]

model._ = Set(model.A, initialize=arc_list_hack)
```

Declare mass balance constraints for nodes. For each node, flow entering to node (inflow) must be equal to flow leaving that node (outflow).

```
# Enforce flow through each node (mass balance)
def flow_rule(model, node):
    if node in [value(model.source), value(model.sink)]:
        return Constraint.Skip
    outflow = sum(model.X[i,j,k]/model.a[i,j,k] for i,j,k in arcs_out[node])
    inflow = sum(model.X[i,j,k] for i,j,k in arcs_in[node])
    return inflow == outflow
model.flow = Constraint(model.N, rule=flow rule)
```

<u>Data Structure of the Model</u> (data.dat)

The data file contains a list of nodes, set of piecewise arc parameters (k), parameters for which mass balance constraint will not be calculated, and list of network links with their properties.

Declare list of nodes

```
set N :=
D94.2001-08-31
D94.2001-09-30
INBOUND.2001-08-31
INBOUND.2001-09-30
OUTBOUND.2001-09-30
SINK
SOURCE
SR_CLE.2001-08-31
SR CLE.2001-09-30;
```

Declare set of 'k' parameters. Currently maximum number of piecewise arcs is assumed 10

```
set k := 0 1 2 3 4 5 6 7 8 9 10;
```

Super source and super sink parameters, where all inflows come and drain, respectively

```
param source := SOURCE;
param sink := SINK;
```

Declare the matrix for network links. This matrix consists of origin node (i), terminal node (j), piecewise arc (k), cost (c), amplitude (a), lower bound (l), and upper bound (u).

Installing Pyomo:

In order to run Pyomo, Python must be installed in your computer. Pyomo can run both with Python 2 and Python 3. However, installing GLPK, a linear programming solver required to run a Pyomo model, is easy to install with Python 3 for Windows users. So, we will only consider installation process for Python 3

Steps:

Install Python 3 through Anaconda:

Anaconda is required to install Pyomo. There are other ways to install Pyomo, such as pip install pyomo. However, currently solvers cannot be installed with pip command. Also, Anaconda already included Python, so there is no need to install Python separately. Anaconda package can be installed from https://www.continuum.io/downloads. Please, make sure to select Python version 3 or later.

Install Pyomo:

Once you install Anaconda, open Command Prompt (for Windows users: go to Start and search for Command Prompt). The latest version of Pyomo and installation commands can be found at https://conda.anaconda.org/cachemeorg.

In Command Prompt, type conda install -c cachemeorg pyomo=4.2.10784

There are also extra Pyomo packages that need to be installed. In order to install extra packages, in Command Prompt, type conda install -c cachemeorg pyomo.extras=2.0

Please note that, currently the channel Cachemeorg supports on Windows-64 bit. If you are running Windows-32 bit, pyomo and pyomo.extras are available in other channels (such as conda-forge). The list of channels can be retrieved with this command: anaconda search -t conda pyomo

Install Solver:

Currently, GNU linear programming solver GLPK is employed. However, the model can work with different solvers.

In order to install GLPK solver, in Command Prompt, type conda install -c cachemeorg glpk=4.57

Running the Model:

The abstract model file (pyvin.py) and the data file (data.dat) must be in the same directory in order to run the model. Open Command Prompt and update current directory to point where your model files are. In Windows, Command Prompt can be easily opened in current directory with Alt+Shift+F and select 'open command window here'

In Command Prompt opened in the same directory as your model files (*.py and *.dat), type

```
pyomo solve --report-timing --solver=glpk --solver-suffix=dual pyvin.py
data.dat --json
```

This command will show how much time is spent in each process and will save results into 'results.json' file. This command will also report dual values for nodes and constraints. Please note that, GLPK solver returns only nonzero values.