
PhreeqPy Documentation

Release 0.4.0

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PHREEQPY - PYTHON TOOLS FOR PHREEQC

1.1 Introduction

PhreeqPy is Open Source software and provides Python tools to work with [PHREEQC](#).

Currently it provides access to the new [IPhreeqc](#) interface without the need to run a COM server and therefore also works on non-Windows systems. [IPhreeqc](#) is described in more detail in this [publication](#).

Please [let us know](#) what you do with PhreeqPy or if things do not work as expected. There is a [mailing list](#) for PhreeqPy. Please [subscribe](#) to get your questions about PhreeqPy answered.

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1.2 Installation

Pythons (tested): Python 2.6, 2.7, 3.3, 3.4, 3.5, 3.6, 3.8, 3.7, 3.9 PyPy

Pythons (untested but should work since using ctypes): IronPython, Jython 2.7

Platforms: Unix/Posix and Windows

PyPI package name: [phreeqpy](#)

Installation with *pip*:

```
pip install -U phreeqpy
```

You need an [IPhreeqc](#) shared library for your operating system. PhreeqPy comes with shared libraries for 32-bit Windows, 32-bit Linux and 64-bit Mac OS X. There is no 64-bit Windows distributed with PhreeqPy. Please install the **'Windows COM 64-bit'** module of [IPhreeqc](#) and use `phreeqpy.iphreeqc.phreeqc_com` as `phreeqc_mod` instead of `import phreeqpy.iphreeqc.phreeqc_dll as phreeqc_mod` as shown in the example. If you use Anaconda or Miniconda make sure to:

```
conda install pywin32
```

before you install PhreeqPy.

You may download an appropriate library from here: <ftp://brrftp.cr.usgs.gov/pub/charlton/iphreeqc/>

For example for Linux:

```
wget ftp://brrftp.cr.usgs.gov/pub/charlton/iphreeqc/iphreeqc-2.18.4-6386.tar.gz
tar -xzvf iphreeqc-2.18.4-6386.tar.gz
cd iphreeqc-2.18.4-6386
./configure
make
make check
sudo make install
```

Then either use the full path to the shared library when making an instance of `phreeqc_dll.IPhreeqc`

```
phreeqc = phreeqpy.iphreeqc.phreeqc_dll.IPhreeqc('/full/path/to/libiphreeqc.so')
```

or copy the shared object into `phreeqpy/iphreeqc` replacing the existing one. For example:

```
sudo cp /usr/local/lib/libiphreeqc.so /path/to/site-packages/PhreeqPy-0.2.0-py2.7.egg/
↪phreeqpy/iphreeqc/libiphreeqc.so.0.0.0
```

1.3 Benchmark Test Comparing PhreeqPy to External Processes, COM and C++

This publication demonstrates how PhreeqPy can be used for reactive transport modeling.

Müller M., Parkhurst D.L., Charlton S.R. (2011) [Programming PHREEQC Calculations with C++ and Python - A Comparative Study](#) , In: Maxwell R., Poeter E., Hill M., Zheng C. (2011) MODFLOW and More 2011 - Integrated Hydrological Modeling, Proceedings, pp. 632 - 636.

CLASS IPHREEQC

This is the main class to work with the IPhreeqc interface.

You can *add your own shared library for IPhreeqc*.

Access PHREEQC-DLL via ctypes.

This is exchangeable with the COM interface.

class phreeqpy.iphreeqc.phreeqc_dll.**IPhreeqc** (*dll_path=None*)
Wrapper for the IPhreeqc DLL.

Connect to DLL and create IPhreeqc.

The optional *dll_path* takes a path to the IPhreeqc shared library. If not provided it tries to select an appropriate library. Make sure you have the right library for your operating system. You may download one from here: <ftp://brrftp.cr.usgs.gov/pub/charlton/iphreeqc/>

See the PhreeqPy documentation for help on compiling a IPhreeqc shared library.

accumulate_line (*line*)
Put line in input buffer.

add_error (*phc_error_msg*)
Add an error message to Phreeqc.

add_warning (*phc_warn_msg*)
Add an warning message to Phreeqc.

clear_accumlated_lines ()
Clear the input buffer.

property column_count
Get number of columns in selected output.

property component_count
Return the number of components.

create_iphreeqc ()
Create a IPhreeqc object.

destroy_iphreeqc ()
Delete the current instance of IPhreeqc.

get_component (*index*)
Get one component.

get_component_list ()
Return all component names.

get_error_string()

Retrieves the error messages.

get_selected_output_array()

Get all values from selected output.

get_selected_output_column(*col*)

Get all values for one column from selected output.

get_selected_output_row(*row*)

Get all values for one from selected output.

get_selected_output_value(*row, col*)

Get one value from selected output at given row and column.

load_database(*database_name*)

Load a database with given file_name.

load_database_string(*input_string*)

Load a database from a string.

static raise_ipq_error(*error_code*)

There was an error, raise an exception.

raise_string_error(*errors*)

Raise an exception with message from IPHREEQC error.

property row_count

Get number of rows in selected output.

run_string(*cmd_string*)

Run PHREEQC input from string.

set_output_file_off()

Turn on writing to selected output file.

set_output_file_on()

Turn on writing to selected output file.

set_selected_output_file_off()

Turn on writing to selected output file.

set_selected_output_file_on()

Turn on writing to selected output file.

exception phreeqpy.iphreeqc.phreeqc_dll.PhreeqcException

Error in Phreeqc call.

class phreeqpy.iphreeqc.phreeqc_dll.VAR

Struct with data type and data values.

See Var.h in PHREEQC source.

class phreeqpy.iphreeqc.phreeqc_dll.VARUNION

Union with types.

See Var.h in PHREEQC source.

EXAMPLES FOR PHREEQPY

This is an example for the use PhreeqPy for one-dimensional advection. It is the example 11 from the PHREEQC users manual.

```
"""Advection with DLL or COM server.

Using MODFIY we update the concentration on every
time step. We shift by one cell per step.
"""

from __future__ import print_function

import sys

# Simple Python 3 compatibility adjustment.
if sys.version_info[0] == 2:
    range = xrange

import os
import timeit

MODE = 'dll' # 'dll' or 'com'

if MODE == 'com':
    import phreeqpy.iphreeqc.phreeqc_com as phreeqc_mod
elif MODE == 'dll':
    import phreeqpy.iphreeqc.phreeqc_dll as phreeqc_mod
else:
    raise Exception('Mode "%s" is not defined use "com" or "dll".' % MODE)

def make_initial_conditions():
    """
    Specify initial conditions data blocks.

    Uniform initial conditions are assumed.
    """
    initial_conditions = """
TITLE Example 11.--Transport and ion exchange.
SOLUTION 0 CaCl2
    units          mmol/kgw
    temp            25.0
    pH              7.0    charge
    pe              12.5    O2(g)  -0.68
    Ca              0.6
```

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```

    Cl          1.2
    SOLUTION 1  Initial solution for column
      units          mmol/kgw
      temp           25.0
      pH             7.0      charge
      pe             12.5     O2(g)   -0.68
      Na             1.0
      K              0.2
      N(5)           1.2
      END
    EXCHANGE 1
      equilibrate 1
      X              0.0011
    END
    """
    return initial_conditions

def make_selected_output(components):
    """
    Build SELECTED_OUTPUT data block
    """
    headings = "-headings   cb   H   O   "
    for i in range(len(components)):
        headings += components[i] + "\t"
    selected_output = """
    SELECTED_OUTPUT
      -reset false
    USER_PUNCH
    """
    selected_output += headings + "\n"
    #
    # charge balance, H, and O
    #
    code = '10 w = TOT("water")\n'
    code += '20 PUNCH CHARGE_BALANCE, TOTMOLE("H"), TOTMOLE("O")\n'
    #
    # All other elements
    #
    lino = 30
    for component in components:
        code += '%d PUNCH w*TOT("%s")\n' % (lino, component)
        lino += 10
    selected_output += code
    return selected_output

def initialize(cells, first=False):
    """
    Initialize IPhreeqc module
    """
    phreeqc = phreeqc_mod.IPhreeqc()
    phreeqc.load_database(r"phreeqc.dat")
    initial_conditions = make_initial_conditions()
    phreeqc.run_string(initial_conditions)
    components = phreeqc.get_component_list()
    selected_output = make_selected_output(components)

```

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```

phreeqc.run_string(selected_output)
phc_string = "RUN_CELLS; -cells 0-1\n"
phreeqc.run_string(phc_string)
conc = get_selected_output(phreeqc)
inflow = {}
initial = {}
for name in conc:
    if first:
        inflow[name] = conc[name][0]
    else:
        inflow[name] = conc[name][1]
        initial[name] = conc[name][1]
task = initial_conditions + "\n"
task += "COPY solution 1 %d-%d\n" % (cells[0], cells[1])
task += "COPY exchange 1 %d-%d\n" % (cells[0], cells[1])
task += "END\n"
task += "RUN_CELLS; -cells %d-%d\n" % (cells[0], cells[1])
task += selected_output
phreeqc.run_string(task)
conc = get_selected_output(phreeqc)
for name in conc:
    value = [initial[name]] * len(conc[name])
    conc[name] = value
return phreeqc, inflow, conc

def advect_step(phreeqc, inflow, conc, cells):
    """Advect by shifting concentrations from previous time step.
    """
    all_names = conc.keys()
    names = [name for name in all_names if name not in ('cb', 'H', 'O')]
    for name in conc:
        # shift one cell
        conc[name][1:] = conc[name][:-1]
        conc[name][0] = inflow[name]
    modify = []
    for index, cell in enumerate(range(cells[0], cells[1] + 1)):
        modify.append("SOLUTION_MODIFY %d" % cell)
        modify.append("\t-cb      %e" % conc['cb'][index])
        modify.append("\t-total_h %f" % conc['H'][index])
        modify.append("\t-total_o %f" % conc['O'][index])
        modify.append("\t-totals")
        for name in names:
            modify.append("\t\t%s\t%f" % (name, conc[name][index]))
    modify.append("RUN_CELLS; -cells %d-%d\n" % (cells[0], cells[1]))
    cmd = '\n'.join(modify)
    phreeqc.run_string(cmd)
    conc = get_selected_output(phreeqc)
    return conc

def get_selected_output(phreeqc):
    """Return calculation result as dict.

    Header entries are the keys and the columns
    are the values as lists of numbers.
    """

```

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```

output = phreeqc.get_selected_output_array()
header = output[0]
conc = {}
for head in header:
    conc[head] = []
for row in output[1:]:
    for col, head in enumerate(header):
        conc[head].append(row[col])
return conc

def run(ncells, shifts, specie_names):
    """Do one run in one process.
    """
    cells = (1, ncells)
    phreeqc, inflow, conc = initialize(cells, first=True)
    outflow = {}
    for name in specie_names:
        outflow[name] = []
    for _counter in range(shifts):
        # advect
        conc = advect_step(phreeqc, inflow, conc, cells)
        for name in specie_names:
            outflow[name].append(conc[name][-1])
    return outflow

def write_outflow(file_name, outflow):
    """Write the outflow values to a file.
    """
    fobj = open(file_name, 'w')
    header = outflow.keys()
    for head in header:
        fobj.write('%20s' % head)
    fobj.write('\n')
    for lineno in range(len(outflow[head])):
        for head in header:
            fobj.write('%20.17f' % outflow[head][lineno])
        fobj.write('\n')

def main(ncells, shifts):
    """Run different versions with and without multiprocessing
    """

    def measure_time(func, *args, **kwargs):
        """Convenience function to measure run times.
        """
        start = timeit.default_timer()
        result = func(*args, **kwargs)
        return result, timeit.default_timer() - start

    print('Dimensions')
    print('=====')
    print('number of cells: ', ncells)
    print('number of shifts ', shifts)
    specie_names = ('Ca', 'Cl', 'K', 'N', 'Na')

```

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```
outflow, run_time = measure_time(run, ncells, shifts, specie_names)
if not os.path.exists('data'):
    os.mkdir('data')
write_outflow('data/out.txt', outflow)
print('run time:', run_time)
print("Finished simulation\n")

if __name__ == '__main__':
    main(ncells=40, shifts=120)
```


CHANGELOG HISTORY

4.1 Changes between versions 0.2 and 0.3

- Added `set_output_file_off()` and `set_output_file_on()` that call IPhreeqc's `SetOutputFileOn`

4.2 Changes between versions 0.1 and 0.2

- Added more IPhreeqc functions.
- Added support for Mac OS X.
- Added error handling turning IPhreeqc errors into Python exceptions.
- Added Python 3 compatibility. Tested with Python 3.3.
- Added documentation in addition to example on website.

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Über diese Website erheben wir keine personenbezogenen Daten.

6.1 1. Datenerhebung und -protokollierung

Wir speichern von Ihrem Besuch Log-Dateien mit diesem Inhalt:

- pseudonymisierte IP-Adresse (ohne Personenbezug, siehe Erklärung unten)
- Datum und Uhrzeit des Zugriffs
- Art des Zugriffes (z.B. GET oder POST)
- besuchten Seiten innerhalb unseres Angebots
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- übertragene Datenmenge
- Meldung über erfolgreichen Abruf
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Bei uns werden nur pseudonymisierte IP-Adressen von Besuchern der Website gespeichert. Auf Webserver-Ebene erfolgt dies dadurch, dass im Logfile standardmäßig stat der tatsächlichen IP-Adresse des Besuchers z.B. 123.123.123.123 eine IP-Adresse 123.123.123.XXX gespeichert wird, wobei XXX ein Zufallswert zwischen 1 und 254 ist. Die Herstellung eines Personenbezuges ist nicht mehr möglich.

6.2 2. Verwendung von Cookies

Wir verwenden keine Cookies.

DATA PROTECTION STATEMENT

English Version for your convenience. The German version below is the legally binding one.

This page describes the collection, processing, and usage of your personal data while visiting this website. Your data is protected within the legal regulations. This page explains what data are collected while visiting this website and how they are used.

We do not collect personal via this website.

7.1 1. Data Collection and Data Logging

We store log files about your visit with this content:

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- protocol (e.g. HTTP 1.1)
- amount of transmitted data
- referring domain (e.g. google.com)
- type of webbrowsers

We do not save hostname or Ip address of your request. The collected data is not personal and don't allow to find out who is visiting our website.

We only store pseudonymised IP addresses of visitors to the website. At the web server level, this happens by default by storing an IP address 123.123.123.XXX in the log file instead of the visitor's actual IP address, for example, 123.123.123.123. The "XXX" is a random value between 1 and 254, so it is no longer possible to establish the true identity of the visitor.

7.2 2. Use of Cookies

We do not use cookies.

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