When all else fails: do-it-yourself programming

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Biomolecular simulations

Standard tasks:

- Run an MD simulation.
- Calculate atomic fluctuations.
- Visualize trajectory.
- ...
- ⇒ standard programs

Nonstandard tasks:

- Automatize a multistep operation.
- Extract data from a file.
- File format conversion.
- Implement new algorithms.
- Nonstandard visualization.
- ...
- ⇒ do-it-yourself programming

Programming options

Shell scripts:

- quick to write
- work on any (Unix) system
- very limited possibilities
- difficult to read
- very slow execution

Suitable for: automatization of simple operations

Low-level languages (C, C++, Fortran, ...):

- fast execution
- good development tools
- require significant experience
- long development times
- can be difficult to read

Suitable for: time-critical algorithms

Programming options

High-level languages (Perl, Python, Ruby, ...):

- complete programming languages
- easy to learn
- fast development
- easy to read
- interface to low-level languages
- slow execution

Suitable for: everything that is not time-critical

Empirical rule: 90% of any program is *not* time critical

Therefore:

- Everyone should use a high-level language.
- Developers of numerical methods must also know a low-level language.

High-level languages

Features:

- Interpreted (no compilation)
- High-level data types (lists, dictionaries, ...)
- No variable type declarations
- Automatic memory management
- Large library of existing code

Applications:

- Complete programs (file format conversion, ...)
- "Glue" between other programs
- "Scripting" of library code
- Mixed high/low-level programs

Python

Features:

- Clean syntax
- Object-oriented
 (→ problem-oriented data structures)
- Good low-level language interface (C, C++, Fortran)
- Free and portable implementation

Examples:

Sum up numbers in the third column of a text file

```
from Scientific.IO.TextFile import TextFile
from string import atof, split
sum = 0.
for line in TextFile('my_data'):
    sum = sum + atof(string.split(line)[2])
print sum
```

Python

Examples:

Calculate the radius of gyration of a protein

```
from MMTK.Proteins import Protein
import Numeric

protein = Protein('protein.pdb')
center = protein.centerOfMass()
r_sq = 0.
for atom in protein.atomList():
    mass = atom.mass()
    distance = atom.position()-center
    r_sq = r_sq + mass*distance*distance
r_sq = r_sq/protein.mass()
print Numeric.sqrt(r_sq)
```

Modules

Python code is structured into *modules*. Objects in another module must be *imported* before they can be used. One can import the module, individual objects from a module, or everything from a module.

Examples:

```
import Numeric
print Numeric.sin(Numeric.pi/3)
from Numeric import pi, sin
print sin(pi/3)
from Numeric import *
print sin(pi/3)
```

Modules can contain other modules:

```
from Scientific.Geometry import Vector
print Vector(1., -2., 0.5).length()

from Scientific import Geometry
print Geometry.Vector(1., -2., 0.5).length()

import Scientific
print Scientific.Geometry.Vector(1., -2.,
```

Basic math

Numbers

Integer 0, 1, 2, 3, -1, -2, -3

Real 0., 3.1415926, -2.05e30, 1e-4

Imaginary/Complex 1j, -2.5j, 3+4j

Special numbers: Numeric.pi, Numeric.e

Arithmetic

Addition 3+4, 42.+3, 1+0j

Subtraction 2-5, 3.-1, 3j-7.5

Multiplication 4*3, 2*3.14, 1j*3j

Division 1/3, 1./3., 5/3j

Power 1.5**3, 2j**2, 2**-0.5

Functions

Numeric.sqrt, Numeric.log, Numeric.log10, Numeric.exp, Numeric.sin, Numeric.cos, Numeric.tan, Numeric.arcsin, Numeric.arccos, Numeric.arctan, Numeric.sinh, Numeric.cosh

Text strings

Two forms: 'abc' or "abc"

Multiline strings:

''', 'This string extends over two lines.'''

Control characters: 'end of line:\n'

Concatenation: 'abc' + 'def'/ gives 'abcdef'

Repetition: 3*"ab" gives "ababab"

Indexing: 'abc'[0] is 'a', 'abc'[1:] is 'bc'

String operations in module string: strip, split, join, atoi, atof, find, index, replace and many others

Lists

Lists are sequences of arbitrary objects:

```
some_prime_numbers = [2, 3, 5, 7, 11]
names = ['Smith', 'Jones']
a_mixed_list = [3, [2, 'b']]
```

Elements and subsequences are accessed by indexing:

```
print names[0]
names[1] = 'Python'
print a_mixed_list[-1]
some_prime_numbers[3:] = [17, 19, 23, 29]
```

Concatenation: [1, 2]+['a'] gives [1, 2, 'a']

Repetition: 3*[0] is [0, 0, 0]

Some more list operations:

```
names.append('van Rossum')
some_prime_numbers.sort()
a_mixed_list.reverse()
print len(names)
print names.index('Smith')
```

Dictionaries

Dictionaries map (almost) arbitrary *keys* to arbitrary *values*.

Examples:

```
atomic_mass = {'H': 1., 'C': 12., 'S': 32.}
atomic_mass['0'] = 16.
print atomic_mass['0'] + 2*atomic_mass['H']

Lookup with default value:
atomic_mass.get('X', 0.)
```

Applications:

- Storing tables
- Counting (counted objects are keys!)
- Classifying (values are lists)
- ...

Flow control

Conditions:

```
if a > b:
    print 'a > b'
elif a == b:
    print 'a == b'
else:
    print 'a < b'</pre>
```

Loops over sequences (lists, strings, ...):

```
for x in [1.2, 2.5, -0.2]:
    ...

for n in range(10):
```

Conditional loops:

The block structure is defined by *indentation*.

Functions

```
def square(x):
   return x**2
print square(3)
Multiple return values:
def sum_and_difference(a, b):
   return a+b, a-b
x, y = sum_and_difference(1, 2)
Variables created in a function are local:
def square(x):
   sq = x**2
   return sq
sq = 5.
print square(1.)
print sq
prints 5., not 1.
```

Error handling

```
def inverse(x):
    return 1./x
print inverse(0)
```

Output:

```
Traceback (innermost last):
   File "demo.py", line 3, in ?
     print inverse(0)
   File "demo.py", line 2, in inverse
     return 1./x
ZeroDivisionError: float division
```

Intercepting an error:

```
try:
    print inverse(0)
except ZeroDivisionError:
    print 'Division by Zero'
```

Programming by creating *problem-specific* data types and *operations* for these data types.

Example: a simple representation for atoms
from Scientific.Geometry import Vector

class Atom:

def __init__(self, name, position):
 self.name = name
 self.position = position

def translateBy(self, vector):
 self.position = self.position+vector

atom = Atom('C-alpha', Vector(1., 0., 0.))
atom.translateBy(Vector(0.1, -0.2, 0.05))

Procedural approach:

from Scientific.Geometry import Vector
atom = ['C-alpha', Vector(1., 0., 0.)]
atom[1] = atom[1] + Vector(0.1, -0.2, 0.05)

Problem: code that works on atom data must

Advantages of OOP:

 Encapsulation: users of a class don't need to know its implementation. All code that knows how atom data are stored is located in the class Atom.

Advantage: the implementation can be changed without affecting client code.

 Data representations are more universal than procedures.

Advantage: classes can be reused for other projects.

Classes directly represent real-world objects.

Advantage: code becomes easier to understand.

Functions defined in classes are called *methods*. The class of an object determines which method is called (*polymorphism*).

Suppose we also have a Molecule class:

```
class Molecule:
```

Then x.translateBy(v)

- calls Atom.translateBy if x is an atom
- calls Molecule.translateBy if x is a molecule
- reports an error if x is something else

Advantage: higher-level code can be written such

Inheritance is the definition of a data type as a *specialization* of an existing one.

```
class Ion(Atom):
    def __init__(self, name, position, charge):
        Atom.__init__(name, position)
        self.charge = charge

ion = Ion('Fe, Vector(0., 1., 0.), 2.)
ion.translateBy(Vector(-0.1, -0.3, 0.05))
```

An Ion inherits all the features of an Atom, and adds its own ones (the charge).

Advantage: several specializations of general classes can be written without changing existing (tested!) code and without duplicating code.

Profiting from existing code

Python standard library:

- string and text processing
- object serialization: write any object to a file
- thread support
- database interfaces
- profiling and debugging
- most internet protocols
- complete Web server
- access to common data and file formats, including HTML and XML

Other free libraries and packages:

- Numerical Python, Scientific Python
- plotting
- signal processing
- interfaces to scientific data formats
- distributed objects (CORBA etc.)
- Zope (Web server management system)