Scilab Bag Of Tricks

The Scilab-2.5 IAQ (Infrequently Asked Questions)

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sci-BOT – the Scilab Bag Of Tricks – is a collection of Scilab experience that come from every day use. We warn of common pitfalls, discuss stylistic issues, shed light on unknown spots, and show many different ways of increasing the performance of Scilab functions.

The document is not meant to be comprehensive or even suitable to a particular level of knowledge. Some sections are at the beginners level, some even surprise long-time users.

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Preface

Often we encounter technical problems that we have to solve, to overcome somehow, or just to work around. After having mastered the difficulty, we gladly add it to the knowledge-base in our mind, but from a certain level of difficulty we make notes in one form or the other. These notes then serve for later reference. A collection of related notes can be exploited to gain further insight in the class of problems it describes. Last but not least one can get ambitious to fill the holes of knowledge that an existing set of notes leaves open.

Richard B. Johnson

An expert in a particular computer language is really an expert in the work-arounds necessary to use this language to perform useful work. An ideal computer language would do exactly what it was told simply from reading a specification. In the absence of a specification, it would ask enough questions to produce such a specification, then it would generate the code necessary to perform the specified functions.

... 

Even C has its shortcomings which have to be handled with assembly language extensions. A Master Carpenter has many tools and is expert with most of them. If you only know how to use a hammer, every problem begins to look like a nail. Stay away from that trap. It bytes (sic).

This is the story of sci-BOT paraphrased. It started with bits of experience gathered in our heads and scattered e-mail correspondence. After more and more e-mails piled up, telling the same old stories one of the authors (lvd) decided to compile the problems and their solutions into a convenient format. Perl’s plain old documentation, POD, was chosen for its simplicity paired with a multitude of output formats. However, after 2000 plus lines it became clear that POD was missing a feature that would be needed more and more as sci-BOT would grow bigger: cross references. A more powerful documentation format and the associated tools had to be found. A two week web research resulted in one winner: DocBook (http://www.oasis-open.org/docbook/). The downside of the necessary switch of formats was that the previous work done with POD had to be converted into DocBook. Daytime work plus adding new material to sci-BOT plus converting the old work into the new format is too much for a single volunteer. So, a second idiotM-DEL author was searched and found (cls). His ten years of experience with the TeX/LaTeX typesetting system, his accuracy, and his intensity with which he attacks any obstacle made him the ideal choice for this madnessM-DEL project.

1. Outline

We open up talking about some of the most common syntactic pitfalls when using Scilab in Chapter 2. Finding that some of these syntax problems can be avoided with a clear programming style, the next chapter, Chapter 3, deals with coding issues. We then focus on the parts of Scilab that are not well
documented and therefore widely remain Chapter 4. A very brief discussion, Chapter 5, about the graphical capabilities follows. For many users not only enjoy Scilab’s nice user interface but demand high performance from the interpreter the massive chapter, Chapter 6, about performance issues covers these needs. It begins by introducing techniques suitable at a high level like vectorization which do not require low level programming and the dives down into the extension of Scilab by compiled routines. This is a vast field by itself. Therefore we devoted a full chapter, Chapter 7, to Scilab’s low level API. sci-BOT closes with Chapter 8 containing remarks on compiling and debugging as well as comments on the supplied documentation and available web pages.

2. Other Formats of sci-BOT

sci-BOT, the Scilab Bag-of-Tricks is available as SGML, as HTML, or several “printer-ready” versions. Each variant is available in different packing-/compression formats.

Alternate sci-BOT formats

SGML source distribution

The “Real Thing” (tm)! These are our SGML-sources. Building sci-BOT from source requires SGML DocBook version 4.0.

- SGML, tar (sci-bot-sgml.tar), MD5 (sci-bot-sgml.tar.md5)
- SGML, tar.gz (sci-bot-sgml.tar.gz), MD5 (sci-bot-sgml.tar.gz.md5)
- SGML, tar.bz2 (sci-bot-sgml.tar.bz2), MD5 (sci-bot-sgml.tar.bz2.md5)
- SGML, tar.Z (sci-bot-sgml.tar.Z), MD5 (sci-bot-sgml.tar.Z.md5)
- SGML, zip (sci-bot-sgml.zip), MD5 (sci-bot-sgml.zip.md5)

Web collection

This is sci-BOT in HTML; conveniently bundled for your offline reading pleasure.

- HTML, tar (sci-bot-html.tar), MD5 (sci-bot-html.tar.md5)
- HTML, tar.gz (sci-bot-html.tar.gz), MD5 (sci-bot-html.tar.gz.md5)
- HTML, tar.bz2 (sci-bot-html.tar.bz2), MD5 (sci-bot-html.tar.bz2.md5)
- HTML, zip (sci-bot-html.zip), MD5 (sci-bot-html.zip.md5)
Print versions

The printable versions are single files. By the way, you don’t have to print it; it looks great with Ghostview, too.

- \texttt{PostScript, ps} (sci-bot.ps), \texttt{MD5} (sci-bot.ps.md5)
- \texttt{PostScript, ps.gz} (sci-bot.ps.gz), \texttt{MD5} (sci-bot.ps.gz.md5)
- \texttt{PostScript, ps.bz2} (sci-bot.ps.bz2), \texttt{MD5} (sci-bot.ps.bz2.md5)
- \texttt{PostScript, ps.Z} (sci-bot.ps.Z), \texttt{MD5} (sci-bot.ps.Z.md5)
- \texttt{PostScript, zip} (sci-bot.ps.zip), \texttt{MD5} (sci-bot.ps.zip.md5)
- \texttt{Portable Document Format, pdf} (sci-bot.pdf), \texttt{MD5} (sci-bot.pdf.md5)
- \texttt{Portable Document Format, pdf.gz} (sci-bot.pdf.gz), \texttt{MD5} (sci-bot.pdf.gz.md5)
- \texttt{Portable Document Format, pdf.bz2} (sci-bot.pdf.bz2), \texttt{MD5} (sci-bot.pdf.bz2.md5)
- \texttt{Portable Document Format, zip} (sci-bot.pdf.zip), \texttt{MD5} (sci-bot.pdf.zip.md5)

3. Typographic Conventions Used in This Book

This section covers the conventions used in this book. Depending on how the version you are currently reading some fonts may look the same.

\textbf{Typographic Conventions}

\texttt{filename}

This font designates the name of a file. A filename optionally includes a path.

\texttt{user input}

This font is used for the user’s input. This refers only to things that can be typed in at the console.

\texttt{meta-variable}

This typeface is reserved for placeholders, i.e. stuff that always is replaced with the real input.

\texttt{literal piece of code}

We use this font to display literal pieces of code, variables, constant as well as operators.
variable
    Scilab variables of all kinds are marked up this way.

function
    Functions or procedures of all kinds are marked up this way.

command
    We use this font for shell commands, but also for Scilab commands.

environment-variable
    To distinguish environment variables from program variables a separate font is used.

4. Acknowledgments

Lydia van Dijk: To the CCMR system administrators Daniel Blakely and Berry Robinson for providing a rock solid multi-platform environment.

Christoph “Solo” Spiel: First of all thanks go to Lydia. Before working with her I did not know whether I am insane. This project has removed all doubt.

To F. Max “Tiger” Pitschi. You have shown me the difference between making software and hacking. Kick me again...
Chapter 1. Introduction

“I have read your posting as of ... to the Scilab newsgroup. It was very clear. Can you make a FAQ out of it?” Yes, we can, and here it is!

The hints, tricks, and information put together in sci-BOT come from our own experience (read: daily struggle), problems we have solved for our colleagues, and of course questions answered on the newsgroup. Therefore, this document is a rather loose collection of facts, and should not be read cover to cover.

What this document is not:

- An introduction to Scilab

- A replacement for reading the documentation
  IONSIO (“In Our Not So Humble Opinion”) folks who do not read the documentation get what they deserve. Scilab’s documentation is truly great, so why not using it? To get a command’s man-page type help at the command line. The same is achieved in the graphical environment with the Help button. If the exact command name is unknown, the powerful cousin of help, apropos jumps in. It can by used from the command line as well as from the Scilab Help Panel.

- Another FAQ list
  We do not follow the simple Question-and-Answer style. Instead we try to explore Scilab in all directions.

In the spirit of the OpenSource any helpful suggestion or correction concerning this collection will be acknowledged with the author’s name and email address. If you want to tell us of a mistake, or want an item added, please drop an email at <lydia_van_dijk@my-deja.com>.
Chapter 1. Introduction
Chapter 2. Common Pitfalls

The nice thing about Scilab? It is almost usable!

There are several peculiarities in Scilab’s way of interpreting an expression that will trip the unwary. Some of them are a result of “compatibility” to a certain commercial product of similar sounding name, others are home grown quirks.

2.1. The Infamous Dot

In Scilab a digit in front or after the decimal point is not enforced. This is similar to e.g. Fortran and C, but contrary to Ada. Therefore, the following three numbers are well formed

\[
\begin{align*}
87.492211 \\
.32493 \\
6857.
\end{align*}
\]

As an aside: \textit{digit+.0, digit+.}, and \textit{digit+ e.g. 123.0, 123.}, and 123 are considered identical. The last of the three examples, a decimal point at the end of the numeral, baffles users who want to invert a vector or matrix component-wise.

\[
\begin{align*}
\text{->} & \quad 1./ [1 \ 2 \ 3] \\
\text{ans} & = \\
! & 1. \quad 0.5 \quad 0.3333333 
\end{align*}
\]

Hey, but this is correct! Then, let us squeeze out the spaces in front of the ./ operator.

\[
\begin{align*}
\text{->} & \quad 1./ [1 \ 2 \ 3] \\
\text{ans} & = \\
! & 0.0714286 \\
! & 0.1428571 \\
! & 0.2142857
\end{align*}
\]

Oops! What happened? The last expression is not interpreted as

\[
(1) ./ ([1 \ 2 \ 3])
\]

but as
Chapter 2. Pitfalls

(1.) / ([1 2 3])

where the parentheses have been introduced for clarity. This behavior is described in SCI/README, and in the Scilab FAQ (http://www-rocq.inria.fr/scilab/faq/index.html).

We suggest to avoid whitespace that influences the calculation by not letting the decimal point stick out on either side. That way expressions with numerals will always be interpreted correctly. For our example this means

\[ \rightarrow 1.0 ./ [1 2 3] \]

\[ \text{ans} = \]

\[ 1. \quad 0.5 \quad 0.3333333 \]

which gives what we had in mind.

2.2. Vector Construction

The square bracket operator \([\])\) is a convenient tool to construct vectors. There even exists an idiom to build a matrix with brackets, which is shown in Example 2-1.

Example 2-1. Building a matrix column-by-column and row-by-row

```plaintext
mat = []
for i = 1:n
    row = []
    for j = 1:m
        ...
        expr = ...
        row = [row expr]
    end
    mat = [mat; row]
end
```

Rows are separated by semi-colons, which actually is straight forward. Columns are separated by commas, or spaces—and here comes trouble.

First, comma and space serve the same purpose and are interchangeable. Thus, the following expressions have the same result.

\[ [1 \ 2 \ 3 \ 4] \]
\[ [1,2,3,4] \]
\[ [1 \ 2 \ 3,4] \]
\[ [1 \ 2 \ 3,4] \]
Chapter 2. Pitfalls

[ 1, 2 3 , 4 ]

Second, a space is sometimes considered a column-separating space, sometimes a intra-expression space. This can lead to some confusion as the following three matrix definitions demonstrate. Who gets all three right without peeking at the answers?

\[-m1 = \begin{bmatrix} 1+%i & -1+%i \\ -1+%i & 1-%i \end{bmatrix}\]

\[
m1 = \begin{bmatrix} 1. + i & -1. + i \\ -1. + i & 1. - i \end{bmatrix}
\]

\[-m2 = \begin{bmatrix} 1 + %i & -1 + %i \\ -1 + %i & 1 - %i \end{bmatrix}\]

\[
m2 = \begin{bmatrix} 1. - 1. + 2.i \\ -1. + i & 1. - i \end{bmatrix}
\]

\[-m3 = \begin{bmatrix} 1 + %i & -1 + %i \\ -1 + %i & 1 - %i \end{bmatrix}\]

\[
m3 = \begin{bmatrix} 1. i & -1. + i \\ -1. + i & 1. - i \end{bmatrix}
\]

Confusion makes the programmer susceptible to writing code she did not intend. To make the matrix expression clear to you and to Scilab there are at least two possibilities.

1. Using no spaces in the construction of the elements of a matrix. This is e.g. demonstrated in \(m1\) above, or

2. Putting every compound expression in parentheses, like

\[-[(1 + %i) (-1 + %i); (-1 + %i) (1 - %i)]]\]

\[
ans = \begin{bmatrix} 1. + i & -1. + i \\ -1. + i & 1. - i \end{bmatrix}
\]

Both ways avoid the ambiguity.

Actually, matrices as simple as the ones shown in the examples can be arranged in a neat way. It is discussed in Section 3.1.2.
2.3. Last Newline

The last line in a Scilab script is ignored if it is not terminated by a newline (^J on UNI* systems, but most of the time written in C-style \n). This is emphasized at several places in the official Scilab documentation, but it is so common to forget it especially when using emacs that we repeat it here. However, emacs can be told *always* to add a final newline by adding (setq require-final-newline t) to the startup-file, .emacs. See “Learning GNU Emacs” [cameron:1996], Table C-8.

Another weapon against this kind of syntax flaw, and a few other pesky things, is e.g. the Perl-script shown in Example 2-2, which fixes part of the format of a Scilab script.

Example 2-2. Canonicalization of Scilab files

```perl
use Text::Tabs;
while (<>) {
    chomp; # remove newline if there is one
    tr/\200-\377/ /; # map 8-bit chars to spaces
    s/[\s]+$]/]; # kill whitespace at end of line
    $$_ = expand $$_; # convert tabs to spaces
    print “$_\n”; # print adding a newline
}
```

2.4. Variable Lifetime And Scoping

2.4.1. Local Variable Scoping

Scilab’s visibility rule for locally defined variables follow those of block structured languages: Variables local to a block shadow all variables of the same name not local this this block.

Variable `v` “shadows” variable `v’ means that `v’ is not accessible neither for reading nor for writing. What is available for manipulation is variable `v`.

Example 2-3. Shadowing of local variables

```
deff(’y = foo(x)’, ’a = 2*x, y = a + 1’)
->a = 1.0 // top level
a =
1.
->foo(3.5)
ans =
```


8.

```plaintext
-> a
a = 1.
-> foo(a)
ans = 3.
-> a
a = 1.
```

Example 2-3 demonstrates that the variable `a` which is local to function `foo` has no influence on the variable `a` in the surrounding environment. Even calling `foo` with a variable named `a` does not break this rule.

As usual in block structured languages variables from all enclosing scopes can be accessed, unless they are shadowed. Example 2-4 shows usage of variable `a` from an enclosing scope.

**Example 2-4. Accessing variables from the enclosing scope**

```plaintext
-> deff('y = bar(x)', 'y = a + 1')
-> a = 1 // top level
a = 1.
-> bar(3.5)
ans = 2.
-> bar(-1)
ans = 2.
-> a = 2
a = 2.
-> bar(-1)
ans = 3.
```

Now what is the “enclosing scope”? It is the call stack; Scilab scopes dynamically!

**Example 2-5. Dynamic scoping**

```plaintext
// scoping in Scilab
```
Chapter 2. Pitfalls

Example 2-5 deserves a close look. Dynamic scoping can be confusing for people used to e.g. C’s lexically scoped `auto` variables.

```c
/* lexical scoping in C */

void first_local(void);
void first(void);
void second(void);

int x = 1;

int
main(void)
{
    first_local(); /* prints 1 */
    first(); /* prints 1 */
    return 0;
}

void first_local(void)
{
    int x = 123; /* warning: unused variable 'x' */
    second();
}

void first(void)
{
    second();
}

void second(void)
{
    printf("%d\n", x);
}
```

```matlab
def('first_local()', 'x = '"foo"', second()');
def('first()', 'second()');
def('second()', 'disp(x)');

x = 1;
first_local() // prints 'foo'
first() // prints 1
```
But compare to Perl:\footnote{1}

\begin{verbatim}
# dynamical scoping with Perl's local variables

sub first_local {
    local $x = 'foo';
    second();
}

sub first {
    second();
}

sub second {
    print "$x\n";
}

$x = 1;
first_local();  # prints 'foo'
first();  # prints 1
\end{verbatim}

Dynamic scoping is an inherently dangerous feature for it might not be obvious where a variable gets its value.

Let us look at functions which try to change variables from an enclosing scope.

$\begin{verbatim}
->deff('y = baz(x)', 'a = 2*a, y = a + 1')
->a = 3 // top level
    a = 3.
->baz(1)
    ans = 7.
->baz(1)
    ans = 7.
->a
    a = 3.
\end{verbatim}$

Obviously, \texttt{a} is unchanged by the calls to \texttt{baz}. What happens is the following:
1. A local variable named `a` is created, and the contents of variable `a` from the enclosing scope is copied into it. Within `baz` the local `a` is changed.

2. When the thread of control leaves `baz` the previous value of `a` is restored.

In other words: A local variable *cannot* influence a variable of the same name in any enclosing scope. The only ways to “export” a – possibly modified – value is either via the list of return values (the preferred way), or with a `global` variable.

As strange as this may sound to programmers accustomed to languages that require an explicit declaration of all variables, this is a necessary feature in Scilab as variables are created when they are first written to (e.g. as in Python). If a local variable in a function would change a global variable or local variable of the same name in another function, adding a new function to an existing system or library became a maintenance nightmare.

### 2.4.2. Global Variables

The `global` attribute of a variable `var` is often misunderstood. It does not place `var` in an all-encompassing name space so that it could be accessed from everywhere without further ado. Instead, `global` places the variable `var` in a separate name space; separate from the interpreter’s name space and separate from all local functions’ name spaces. — and this is only the first half of the story.

```plaintext
->v = -1
v =
  -1.

->global('v')

->who('global')
ans =
v

->clear v

->who('global')
ans =
v

->deff('y = useglobal()', 'y = v')

->useglobal()
!-error 4
undefined variable : v
at line 2 of function useglobal called by :
```

24
useglobal()

As promised, this is only one half. After saying `global` the variable lives in its new name space, but it cannot be accessed. To work with it it must be imported explicitly, using the `global` again. Therefore, a slightly modified version of `useglobal` works.

```->deff('y = useglobal2()', 'global v, y = v')

->useglobal2()
ans =
 - 1.

->v = 1 + 2*%i
v =
 1. + 2.i

->useglobal2()
ans =
 - 1.
```

Now what if we want to access `v` from the interpreter level again? It must be imported just as it must be imported into any function.

```->global('v')

->v
v =
 - 1.

->v = 17 + 4
v =
 21.
```

```->clear v

->useglobal2()
ans =
 21.
```

One last hint: global variables even “survive” a restart. If this is not desired, `clearglobal` should be called in the user’s Scilab startup file, `~/.scilab`.

`clearglobal()`

will clear all global variables.
2.4.3. Clearing Variables

During everyday programming it is not necessary to explicitly remove variables from the work space. All local variables of a function die on exit from that function anyhow, and the variables in the global name space usually do not need special treatment.

However, there are conditions under which it is preferable to wipe out a variable completely. This happens if you need to avoid a pollution of the name space while working with the list of all variables, e.g. `who('local')`. The correct command to kill variable `v` is

```
clear v
```

Note that there are no parentheses. The assignment

```
v = []
```

sets `v` to the empty matrix. It does not remove the variable from the workspace. Global variables are cleared with `clearglobal`.

There is no need to worry if you do not understand how and why to kill a variable. This feature is only needed in very rare occasions.

Notes

1. The behavior of the C-example is reproduced by replacing `local` with `my`.
Chapter 3. Programming Style

The one and only general guideline to good programming style is: “Make it clear!” And one might extend that to “Make it clear; first of all to you and then to the poor person that takes over your project.” Every possible style feature of the language should be used to express the meaning of the code clearly.

3.1. Spacing and Formatting

Although often underestimated, the format, i.e. the visual layout of the source code itself can greatly help in the understanding of the actions described therein.

3.1.1. Intra-Expression Spacing

We often run into code like this

\[ x = a \cdot (x-y)^2 \cdot b \]

This is not bad, especially when typed at the command line for one-time use. But the expression is not as helpful to its understanding as could be. E.g. it can be improved by making the the precedence levels of the operators stand out.

\[ x = a \cdot c + b \cdot (x-y)^2 \]

Now, the assignment is intuitively clear at first glance. We use word “intuitive” here to make the reader alert of the consequences of formatting an expression the wrong way. Then our intuition will mislead us, as in

\[ x = a * c + (x-y)^2 \cdot b \]

3.1.2. Line Breaking

Breaking a long expression into lines can improve the readability dramatically. It is particularly recommended for matrix definitions with the square bracket operator. See also Section 2.2.

\[
\begin{bmatrix}
1+\%i & -1+\%i \\
-1+\%i & 1-\%i
\end{bmatrix}
\]

is superior to
If an arithmetic expression is split into lines the operator at which the split occurs always goes onto the next line. Preferred break points occur right before operators of equal precedence.

\[ d_2 = \text{fact} \times \left( \frac{a}{a+d} \times (b \times (1-\delta) + d \times \delta) - d \right) \times \left( \frac{P}{K} \right)^\theta \]

becomes for example

\[
\begin{align*}
 d_2 &= \text{fact} \times \left( \frac{a}{a+d} \times (b \times (1-\delta) + d \times \delta) - d \right) \times \left( \frac{P}{K} \right)^\theta \\
\end{align*}
\]

or

\[
\begin{align*}
 d_2 &= \text{fact} \times \left( \frac{a}{a+d} \times (b \times (1-\delta) + d \times \delta) - d \right) \times \left( \frac{P}{K} \right)^\theta \\
\end{align*}
\]

or more dramatic

\[
\begin{align*}
 d_2 &= \text{fact} \times \left( \frac{a}{a+d} \times (b \times (1-\delta) + d \times \delta) - d \right) \times \left( \frac{P}{K} \right)^\theta \\
\end{align*}
\]

The last way of breaking the expression is very LISP-like.

### 3.1.3. Setting Brackets Apart

If spaces right inside the parentheses or brackets of an expressions make the subexpression stand out more clearly, they should be used. That way

\[ B(k) = a_1 \times \exp(-b_1 P(k)/K(k) + b_2 Q(k)/K(k)) \]

becomes

\[ B(k) = a_1 \times \exp(-b_1 P(k)/K(k) + b_2 Q(k)/K(k)) \]
3.2. Indentation

Heavy indentation does not hurt! No, in fact it is a great help in finding out the control flow quickly. Let us start with a good example this time, Example 3-1.

Example 3-1. Function `whocat`

```matlab
function s = whocat(cat)
    // return all local variables, functions,
    // etc. that are in category cat.
    s = []; nl = who('local');
    for i = 1:size(nl, 1)
        typ = type(nl(i)) + Relations.
        if typ == cat then
            s = [s; nl(i)];
        end
    end
```

The `for` loop and the `if` branch are immediately recognizable. There are blank lines between the logical blocks of the function. They too aid the reader’s comprehension of `whocat`’s inner workings.

In longer functions the indentation becomes essential for the orientation of the maintainer. Here is an excerpt of a longer function, that would be terribly hard to understand if not massively indented.

```matlab
i = 1; j = 1;
while i <= n1 & j <= n2
    while i <= n1 & j <= n2
        if ~equ(lst1(i), lst2(j)), break, end
        i = i + 1;
        j = j + 1;
    end
    if i >= n1 | j >= n2, break, end
    i = icurs + 1;
    if equ(lst1(icurs), lst2(j)) then
        icurs = icurs + 1;
    end
    if icurs <= n1 then
        if equ(lst1(icurs), lst2(j)) then
```

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// record element(s) missing from lst1
for p = i : icurs-1
    this_diff = [lst1(p), string(-p)];
    diff = [diff; this_diff];
end
// re-sync
i = icurs;
end

The complete listing of this function can be found in Chapter 9.
The last example also shows that we are switching between several style paradigms:

• Neither the “One statement per line” rule is followed consistently,
  
  if equ(lst1(icurs), lst2(j)), break, end

  could be

  if equ(lst1(icurs), lst2(j)) then
    break
  end

• Nor is the intra-line spacing always consistent with the guidelines presented here:
  
  for p = i : icurs-1

  could be

  for p = i:icurs-1

The Golden Rule is that there are no golden rules... This is best known under the term ‘freedom’.

3.3. Choice Of Control Structures

Though not recognized as that by all programmers the flow control structures themselves are first class indicators of the codes workings. We consider three important cases here.
1. while vs. for,
2. if vs. select, and
3. strict block structure vs. premature return.

### 3.3.1. while/for

Expressed in words a for loop tells us:

- We know exactly how many iterations we shall need before we start looping.
- Nothing in the loop body will change this.

Whereas the while loop says:

- We must check whether we should loop at all, and
- we have to re-check after each iteration whether we need another round-trip.

**Corollary:** The termination condition of a while must be influenced in the loop’s body.

Compare the next to code snippets, the first calculating the average of a vector of numbers, the second searching zeroes of a given function.

```python
values = [1.0 2.0 3.0 4.0 5.0];
average = 0;
n = size(values, 'c'); // line 3
for i = 1:n
    average = average + values(i)
end;
average = average / n
```

Form line 3 on, we know the number of iterations, n; from the problem we know that nothing will change that. Thus a for-loop is adequate.

```python
def('[y, dy] = fun(x)', ..
    'y = -0.5 + 1.0 / (1.0 + x^2), ..
    dy = -2.0 * x / (y + 0.5)^2');
x0 = 0.76;
[y, dy] = fun(x0);
while abs(y) > sqrt(%eps)
    x = y / dy - x0;
    x0 = x;
    [y, dy] = fun(x)
```

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Assuming that the function $f_{\text{un}}$ and the start guess $x_0$ is given by the user, we do not know how many loops it will take for Newton’s algorithm to converge, if it does converge at all. (In the example it does.) Here, the while-loop expresses this lack of a-priori knowledge.

### 3.3.2. if/select

The relationship between if and select bears similarity with while and for respectively. In a select clause the different cases are known – and spelled out explicitly – before the thread of control enters the construct. There is a one to one relationship between the states of the selecting expression and the case branch taken. The else branch in a select works exactly as the else in an if.

**Example 3-2. Function fibonacci**

```plaintext
function f = fibonacci(n)
// return n-th Fibonacci number

select n
  case 0 then
    f = 1
  case 1 then
    f = 1
  else
    f = fibonacci(n - 1) + fibonacci(n - 2)
end
```

The selecting expression is not restricted to scalars. For example, vectors work too:

**Example 3-3. Function shape4**

```plaintext
function s = shape4(m)
// classify a 2x2 matrix according to its shape

select abs(m) <= %eps
  case [%t %t; ..
    %t %t] then
    s = "empty"
  case [%t %f; ..
    %f %t] then
    s = "diagonal"
end
```
case [%f %f; ..
    %t %f] then
    s = "upper triangular"
end

case [%t %t; ..
    %f %t] then
    s = "lower triangular"
end

case [%f %f; ..
    %f %f] then
    s = "full"
else
    s = "general"
end

An if clause is more flexible than a select clause, but at the price of being less expressive. Whenever a whole range of values has to be covered the if clause is the only way to go.

Example 3-4. Function mysign

function y = mysign(x)
    // re-write of the sign-function,
    // taking floating-point precision
    // into account
    if abs(x) < %eps
        y = 0.0
    elseif x >= %eps
        y = 1.0
    else
        y = -1.0
    end

3.3.3. Strict Block Structure/Premature Return

The paradigm of structured programming is: “Every block has one and only one entry point.” That’s it! Nothing is said about the number of exit points. The purists often misinterpret the paradigm, demanding a single exit point, too. We prefer our freedom and choose whatever we find adequate to the problem.

Here are two different implementations of an algorithm calculating the factorial of a given integral number.

function y = fact_block(x)

// faculty of x
// block-structured version

select x
  case 0 then
    y = 1
  case 1 then
    y = 1
  else
    y = x * fact(x - 1)
end

The two special cases 0, and 1 are tested separately and the general case is handled in the else branch.

function y = fact_early_ret(x)
  // faculty of x
  // early-return version
  if x >= 0 && x <= 1 then
    y = 1
    return
  end
  y = x * fact(x - 1)
end

This version immediately returns after having treated the special cases, leaving the general case to the “rest” of the function. In this very short function the advantages of the early return are not striking, however they are if there are many special cases to be handled. The “rest” of the function can then concentrate on the core of the problem without being obscured by deeply nested conditionals.

3.4. Size of a Function

There is a rule of thumb for the length of a C-function:

L. Torvalds

Functions should be short and sweet, and do just one thing. They should fit on one or two screenfuls of text (the ISO/ANSI screen size is 80x24, as we all know), and do one thing and do that well.

It is also true for Scilab functions with the exception that high level functions or functions that could eventually be called from the command line directly should be harnessed. See also: Section 4.2.2.
Therefore, they are usually much longer than just two screenful. But the structure decomposes quite naturally in two parts: the argument checking and the computation part. What remains true is that a Scilab function too should do only one thing and do that well.

For more information about programming style consult “The practice of programming” [kernighan:1999] which is centered around C-like languages but offers extremely valuable advice throughout. The Camel, [wall:1996] has a section that is called “Efficiency” in chapter 8. It is as insightful as it is fun to read for the authors discuss the various optimization directions. They do not hesitate to put up contradictory suggestions in the different optimizations paths.

Conclusion of this section: Whatever makes the code’s workings more obvious to the reader is good. In other words: “If it makes ya high, or saves you taxes, then – by any means – do it!”
Chapter 4. Unknown Spots

In this chapter we shed some light onto widely unknown features. Parts like the operator precedence unconsciously are exploited in every-day programming by all of us. Others like the use of function variables are truly unknown, at least to the average Scilab user. So, read on and become a Yedi^H^H^HScilab master.

4.1. Operator Precedence And Associativity

Strange but true, there is no listing of the precedence and associativity of neither class of Scilab’s operators anywhere in the documentation. So, we discuss the operator precedence and associativity in detail.

4.1.1. Numeric Operators

Table 4-1 displays a list of all numeric operators up to digraphs \(^1\), sorted in descending order of their precedence. An equal precedence value (column 1) means the operators are evaluated following the given associativity (column 3).

The table is generated with a Scilab script, i.e. we had the interpreter determine its own precedence rules, which is neat. These scripts are listed in Chapter 9.

Table 4-1. Arithmetic Operators

<table>
<thead>
<tr>
<th>precedence</th>
<th>operator</th>
<th>associativity</th>
<th>comment</th>
</tr>
</thead>
<tbody>
<tr>
<td>21</td>
<td>+</td>
<td>right</td>
<td>unary</td>
</tr>
<tr>
<td>20</td>
<td>^</td>
<td>right</td>
<td></td>
</tr>
<tr>
<td>20</td>
<td>.*</td>
<td>right</td>
<td></td>
</tr>
<tr>
<td>19</td>
<td>-</td>
<td>right</td>
<td>unary</td>
</tr>
<tr>
<td>8</td>
<td>*</td>
<td>non</td>
<td></td>
</tr>
<tr>
<td>8</td>
<td>/</td>
<td>left</td>
<td></td>
</tr>
<tr>
<td>8</td>
<td>.*</td>
<td>non</td>
<td></td>
</tr>
<tr>
<td>8</td>
<td>./</td>
<td>left</td>
<td></td>
</tr>
<tr>
<td>4</td>
<td>\</td>
<td>left</td>
<td></td>
</tr>
<tr>
<td>4</td>
<td>.\</td>
<td>left</td>
<td></td>
</tr>
</tbody>
</table>
4.1.2. Relational Operators

Scilab implements the usual gang of relational operators with some syntactic sugar of having two "unequality"-operators $<>$, and $\neq$. The relational operators' precedences rank in between the numeric and the logical operators like they do in many other modern programming languages. This allows for a minimal use of parentheses in larger expressions like

\[
\text{if } 2.0 * n > 1 + 1.0 \mid n / 3.0 \leq k \text{ then }
\]

\[
\text{...}
\]

\[
\text{end}
\]

which evaluates exactly the same way as

\[
\text{if } ((2.0 * n) > (1 + 1.0)) \mid ((n / 3.0) \leq k) \text{ then }
\]

\[
\text{...}
\]

\[
\text{end}
\]

just with much less line-noise.

4.1.3. Logical Operators

There are three logical operators: $\&$, $|$, and $\sim$, meaning “and”, “or”, and “not”. The twiddle, $\sim$ has the unique syntactic property that any number of consecutive twiddles are allowed and evaluated. But unless you want to enter the obfuscated Scilab contest, sticking with one probably is best as e.g. 15 $\sim$ are as good as none, and therefore
returns `F`.

Table 4-2 shows the complete list of Scilab’s logical (also known as boolean) operators sorted according to decreasing precedence.

### Table 4-2. Boolean Operators

<table>
<thead>
<tr>
<th>operator</th>
<th>associativity</th>
<th>comment</th>
</tr>
</thead>
<tbody>
<tr>
<td>~</td>
<td>right</td>
<td>unary</td>
</tr>
<tr>
<td>&amp;</td>
<td>non</td>
<td></td>
</tr>
<tr>
<td></td>
<td></td>
<td>non</td>
</tr>
</tbody>
</table>

For the logical operators have boolean expressions as their arguments, it is time now to discuss the implicit promotion of numeric types to boolean type, something very familiar to C, Perl, and Python programmers. You have guessed right, the rule is: “Zero is false, everything else is true.” Here are some examples of that rule at work:

```plaintext
-> %t & 0
ans = F

-> %t & 0.1
ans = T

-> 6.34 | %f
ans = T

-> 6.34 | -0.3
ans = T
```

Scilab always evaluates boolean expressions completely. No operator is defined with short-circuit evaluation semantics.

```plaintext
-> deff('b = ret_false()', 'b = %f, disp("ret_false")');

-> ret_false() & ret_false()
ret_false
```
4.2. Functions

Functions are Scilab’s the main abstraction feature, thus they deserve a closer look.

4.2.1. Functions Without Arguments or Return Value

The “Introduction to Scilab”, SCI/doc/Intro.ps, solely explains functions that have one or more arguments, returning one or more values. If only one value is returned the square brackets in the function definition are optional. Therefore the function head

```matlab
function [y] = foo(x)
```

can be abbreviated to

```matlab
function y = foo(x)
```

However, this is 100% pure syntactic sugar. What is much more important – and a valuable feature – is the possibility of defining a function that returns nothing as

```matlab
function ext_print(x)
    printf("%f, %g", x, x)
endfunction
```

does. In Fortran parlance `ext_print` would be called a SUBROUTINE, whereas Ada programmers would term it a PROCEDURE.

Of similar importance is the definition of parameterless functions.

```matlab
function t = hires_timer()
    cps = 166e6
    t = rdtsc() / cps
endfunction
```

The parentheses after the function name are optional when defining the function, but not when calling it. For further information about the omission of parenthesis when calling a function, see Section 4.3.3.
4.2.2. Bulletproof Functions

If we want to write bulletproof Scilab functions, we have to take care that our functions get the right number of arguments which are furthermore of the correct type, and correct dimension. This is due to Scilab’s dynamic nature allowing us to pass arguments of different types, dimension etc. to a function.

We discuss the issues of writing robust function using Example 4-1 as an illustration. The complete function definition is given in Chapter 9.

Example 4-1. Function cat

```scilab
function [res] = cat(macname)
// Print definition of function 'macname'
// if it has been loaded via a library.

[nl, nr] = argn(0);
if nr ~= 1 then
    error("Call with: cat(macro_name)");
end
if type(macname) ~= 10 then
    error("Expecting a string, got a " ..
        + typeof(macname));
end
if size(macname, "**") ~= 1 then
    sz = size(macname);
    error("Expecting a scalar, got a " ..
        + sz(1) + "x" + sz(2) + " matrix")
end

[res, err] = evstr(macname);
if err ~= 0 then
    select err
        case 4 then
            disp(macname + " is undefined.");
            return;
        case 25 then
            disp(macname + " is a builtin function");
            return;
        else
            error("unexpected error", err);
    end // select err
end // err ~= 0
...
```
First, we check how many actual parameters function \texttt{cat} got. The built-in function \texttt{argn} returns the number of left-hand side (or output) variables \texttt{n1}, and then number of right-hand side (or input) values \texttt{nr}.

Ensuring the correct number of \texttt{input} arguments always is the first step. Otherwise we cannot assume whether even accessing a parameter is valid. The number of output values is not as critical, for calling a function with less output variables than specified in the function's signature causes the extra output values to be silently discarded.

After learning the number of actual rhs-parameters, we immediately check whether it is in the right range. In our example simply terminates with an error if the number of arguments is incorrect.

The next thing to address are the types of the arguments. Again we let the function fail with an error if it does not get what it wants, but this is not the only way possible.

It is conceivable that we convert from one type to another, say from numeric to string. Furthermore, it is possible that the type of the arguments determines the algorithm chosen, a feature normally advertised under the name "function overloading".

Finally, we examine the arguments' structure. A function can e.g. allow scalars only, or accept scalars and matrices. Here, we enforce a scalar. In other functions certain dimensional relations of several input parameters must be enforced. E.g. the matrix multiplication \(A \times B\) is only defined for \(\text{size}(A, 'c') == \text{size}(B, 'r')\).

Now we can start with the real work.

At first glance all this checking gizmos might seem exaggerated. To do it justice we should keep in mind that it is only necessary if a function must work reliably in different environments. All functions that a library exports belong to that class, because the library writer does not know how the functions will be used. Quick-and-dirty functions are a different thing, so are functions that are never called interactively.

### 4.2.3. Function Variables

Functions are a data type on their own right; therefore they themselves can be arguments to other functions, and can be elements in lists.

\[
\begin{align*}
\text{defn ('y = fun(x)'), 'if x > 0, y = sin(x); else, y = 1; end'}
\end{align*}
\]

\[
\begin{align*}
\text{fun(\%pi/2)}
\end{align*}
\]

\[
\begin{align*}
\text{ans} &=
\end{align*}
\]

\[
\begin{align*}
\text{1.}
\end{align*}
\]

\[
\begin{align*}
\text{fun(-3)}
\end{align*}
\]
As the example shows Scilab employs its usual copy-by-value semantics when assigning function-variables, consistent with the assignment of variables of any other data type.

### 4.2.4. Nested Function Definitions

Function definitions can be nested. The usual scoping rules apply. Online nested function definitions are some kind of awkward because of the massive number of quotes, but deffs in functions are easy to the eye.

**Example 4-2. Function tauc**

```plaintext
function [t, rmin, r0] = tauc(E0, M, s, D)

deff('U = Umorse(r, steepness, depth)', ..
    'e = exp(-r * steepness); ..
    U = depth*(e^2 - 2*e)');

// point of vanishing potential
deff('y = equ0(x)', 'y = Umorse(x, s, D)');

// reflection point
```

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deff('y = equ1(x)', 'y = Umorse(x, s, D) - E0');

deff('tau = integrand(x)', ...
   'tau = sqrt( M / (2*(E0 - Umorse(x, s, D))) )');

// rationalized units...
units = 10.0e-10 / sqrt(1.380662e-23 / 1.6605655e-27);

// calculate endpoints of definite integral
r0 = fsolve(-10.0, equ0);
rmin = fsolve(-10.0, equ1);

// evaluate definite integral
[t_unscaled, err] = intg(rmin, r0, integrand);
t = 2 * units * t_unscaled;

4.2.5. Functions as Parameters in Function Calls

As mentioned above, user-defined functions can be passed as parameters to (usually different) functions. Builtin functions have to be “wrapped” in user-defined functions before they can be used as parameters.

The following example defines a functional that implements a property of Dirac’s delta distribution.

\[
\text{deff('y = delta(a, foo)', 'y = foo(a)')}
\]

\[
\text{delta(cos)}
\]

!-error 25
bad call to primitive :cos

\[
\text{deff('y = mycos(x)', 'y = cos(x)')}
\]

\[
\text{delta(0, mycos)}
\]

ans =
1.

The next example is a bit more convoluted, but also closer to the real world. We define a new optimizer function, called minimize, which is based on Scilab’s optim function. minimize expects two vectors of data points xdata and ydata, a vector of initial parameters p_ini, the function to be minimized func, and an objective functional obj.

The advantage of defining separate model and objective functions is an increased flexibility as both can be replaced at will without changing the core minimization function minimize.
Example 4-3. Function minimize

```matlab
function [f, p_opt, g_opt] = minimize(xdata, ydata, ..
    p_ini, func, obj)

    // on-the-fly definition of the objective function
    deff('[f, g, ind] = _cost(p_vec, ind)', ..
        '[f_val, f_grad] = func(xdata, p_vec); ..
         [f, g] = obj(f_val - ydata, f_grad)');

    [f, p_opt, g_opt] = optim(_cost, p_ini);
```

The function `minimize` needs a model function `func` that returns the value and the gradient at all points \( x \) for a given vector of parameters \( p_{\text{vec}} \). Moreover we need the objective functional `obj` that gives the "cost" and the direction of steepest descent in parameter space.

In this example we choose a quadratic polynomial for the model, `my_model` and least squares for the objective `lsq`.

```matlab
function [f, g] = my_model(x, p)
    g = [ones(x), x, x.*x];
    f = p(1) + x.*(p(2) + x*p(3));

function [f, g] = lsq(diff, grad)
    f = 0.5 * norm(diff)^2;
    g = grad' * diff;
```

Given these definitions, we can call `minimize`:

```matlab
dx = [0.0 1.0 2.0 2.5 3.0]';
dy = [0.0 0.9 4.1 6.1 9.5]';
p_ini = [0.1 -0.2 0.9]';
[f_fin, p_fin, p_fingrad] = ..
    minimize(dx, dy, p_ini, my_model, lsq)
xbasc();
plot2d(dx, dy, -1);  // plot data points ...
xv = linspace(dx(1), dx($), 50)';
yv = my_model(xv, p_fin);
plot2d(xv, yv, l, "000"); // ... and optimized model function
```

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4.2.6. Functions in tlists

Currently the only complex data structure that allows for storage of functions is the typed list tlist.

FIXME: write it

4.2.7. macrovar

FIXME: write it

4.3. Miscellaneous Unknown Spots

4.3.1. Starting scilex

For debugging purposes it is sometimes desirable to start the main Scilab binary, scilex directly. Scilab is usually launched via the scilab script. Both, the script and the binary live in the SCI/bin directory. The script takes care of setting all environment variables and finally fires up scilex. On the other hand, if you want to run a debugger, say gdb, or ddd, or a profiler on Scilab then a manual invocation is the order of the day. Starting scilex is easy as long as you are not hooked by all the command-line editing goodies, and there is no need for any graphics. Actually, for minimum functionality only the environment variable SCI must be set, then we are all set to call scilex. A bash sequence to start Scilab “manually” could look as shown in Example 4-4.

Example 4-4. Manually launching scilex

lydia@orion:~$ cd /site/X11R6/src/scilab
lydia@orion:/site/X11R6/src/scilab$ SCI=`pwd`
lydia@orion:/site/X11R6/src/scilab$ export SCI
lydia@orion:/site/X11R6/src/scilab$ cd bin
lydia@orion:/site/X11R6/src/scilab/bin$ ./scilex -nw

============
Scilab-2.5
Copyright (C) 1989-99 INRIA
Startup execution:
loading initial environment
->

or shorter

lydia@orion:~$ export SCI=/site/X11R6/src/scilab
lydia@orion:~$ $SCI/bin/scilex -nw
       ===========
       Scilab
       ===========

       Scilab-2.5
       Copyright (C) 1989–99 INRIA

Startup execution:
loading initial environment
->

where we are assuming that Scilab is installed in /site/X11R6/src/scilab.

4.3.2. Tuple Assignment

The most commonly used form of assignment is single-assignment. Nonetheless, assigning multiple values in one statement is possible (and no surprise for Perl or Python programmers).

-> [x1 x2 x3] = (1, 2, 3)
  x3 = 3.
  x2 = 2.
  x1 = 1.

See: parents
4.3.3. Omitting Parentheses on Function Call

The parentheses of any one-parameter function can be omitted, if the function accepts a string argument. Moreover, the quotes for a literal string argument can be left out, too.

The is especially useful, when working interactively and loading functions or scripts. There is no need to type until your fingers bleed by saying

\[ \texttt{->getf("foo.sci")} \]

as the next two examples work just as well.

\[ \texttt{->getf "foo.sci"} \]

and even

\[ \texttt{->getf foo.sci} \]

is OK. Note that this is not only true for built-in, but also for user-defined functions.

Function \texttt{exec} is an exception to the rule that a semicolon suppresses any output of the preceding clause. \texttt{exec} does echo all commands when used without parenthesis despite a trailing semicolon, i.e.

\[ \texttt{->exec script.sci;} \]

with semicolon gives same results as

\[ \texttt{->exec('script.sci')} \]

without semicolon, whereas

\[ \texttt{->exec('script.sci');} \]

does not echo the commands in the script file.

Notes

1. The trigraph operators \texttt{.*.,./., and .\.} are left out.
2. Thanks to Glen Fulford for reporting this.
Chapter 5. Graphics

In this chapter we take care of the Achilles heel of Scilab, the graphics functions. Their user interface carries a lot of Fortran artifacts making it hard to remember the plot functions’ exact syntax or leaving the user in the dark about the possibilities.

We take off with the most commonly used graphics feature, plotting of functions and data. (This distinction is a bit artificial in Scilab as everything is discrete.) Then we descend in the hierarchy, looking at the graphics primitives Scilab offers. Finally, we wrap the chapter up with a discussion of the various GUI-functions.

5.1. Function- And Data-Plotting

FIXME: topics: - Plotting lines and points simultaneously - Plotting a different number of points in one graph

FIXME: Enrico or Stéphane this could become your section.

5.2. Graphics Primitives

FIXME: Enrico or Stéphane this could become your section.

5.3. User Interfaces

FIXME: Enrico or Stéphane this could become your section.
Chapter 6. Performance

In this chapter we discuss how expressions can be written to execute more quickly while doing the same thing. Scilab is powerful and flexible, therefore there are plenty of things one can do to speed up function execution. On the downside there are a lot of things that can be done the wrong way, slowing down the execution to a crawl.

In the first part we focus on high-level operations that are executed fast. The main class to name here are vectorized operations. Another class are all functions that are constructing or manipulating vectors or matrices as a whole. The second part of this chapter deals with the extension of Scilab through compiled functions for the sake of increased execution speed.

6.1. High-Level Operations

Not using vectorized operations in Scilab is the main source for suffering from a slow code. Here we present performance comparisons between different Scilab constructs that are semantically equivalent.

6.1.1. Vectorized Operations

The key to achieve a high speed with Scilab is to avoid the interpreter and instead make use of the built-in vectorized operations. Let us explain that with a simple example.

Say we want to calculate the standard scalar product $s$ of two vectors $a$ and $b$ which have the same length $n$. Naive as we are, we start with

```plaintext
s = 0 // line 1
i = 1 // line 2
while i <= n // line 3
    s = s + a(i) * b(i) // line 4
    i = i + 1 // line 5
end // line 6
```

Here Scilab re-interprets lines 3 to 5 in every round-trip, which in total is $n$ times. This results in slow execution. The example utilizes no vectorization at all. On the other hand it uses only very little memory as no vectors have to be stored.

The first step to get some vectorization is to replace the `while` with a `for` loop.
Line 2 is only interpreted once; the vector \( i = 1:n \) is set up and the loop body, line 3 is threaded over it. So, only line 3 is re-evaluated in each round trip.

OK, it is time for a really fast vector operation. In the previous examples the expression in the loop body has not been modified, but we can replace it with the element wise multiplication operator \( .* \) and replace the loop with the built-in \texttt{sum} function. (See also Section 6.1.3.3.)

\[
s = \text{sum}(a \; .* \; b)
\]

One obvious advantage is that we have a one-liner now. Is that as good as it can get? No, the standard scalar product is not only a built-in function it is also an operator:

\[
s = a \; \ast \; b'
\]

We summarize the timing results of a PII/330 Linux-system in Table 6-1.

\begin{table}[h]
\centering
\begin{tabular}{|c|c|}
\hline
\textbf{construct} & \textbf{MFLOPS} \\
\hline
while & 0.005 \\
for & 0.008 \\
\(.* \; \text{and} \; \text{sum} \) & 1.7 \\
\(\ast \) & 2.8 \\
\hline
\end{tabular}
\caption{Comparison of various vectorization levels}
\end{table}

In other words the speed ratio is 1:1.6:330:550. Of course the numbers vary from system to system, but the general trend is clear.

### 6.1.2. Avoiding Indexing

Accessing a vector- or matrix-element via indexing is slow. Sometimes the index cannot be avoided, but there are cases where it can. Compare

\[
\text{for } i = 1:n \\
\quad v(i) = i \\
\text{end}
\]

and
v = []
for i = 1:n
    v = [v, i]
end

The second snippet is not only faster, but in some circumstances may be clearer. Again there is a built-in operator that does the same job at lightning speed, the colon :, which is described in detail in Section 6.1.3.1.

v = 1:n

The speed ratio is approximately 1:1.5:5000.

In the next example, Example 6-1, the functions actually try to do something useful: they mirror a matrix along its columns or rows. We show different implementations of mirrorN that all do the same job, but utilize more and more of Scilab’s vector power with increasing function index N.

Example 6-1. Variants of a matrix mirror function

function b = mirror1(a, dir)
    // mirror matrix a along its
    // rows, dir = ’r’ (horizontal)
    // or along its columns, dir = ’c’ (vertical)

    [rows, cols] = size(a)
    select dir
    case ’r’ then
        for j = 1 : cols
            for i = 1 : rows
                b(i, j) = a(rows - i + 1, j)
            end
        end
    case ’c’ then
        for j = 1 : cols
            for i = 1 : rows
                b(i, j) = a(i, cols - j + 1)
            end
        end
    else
        error(“dir must be ’r’ or ’c’”)
    end

function b = mirror2(a, dir)
    // same as mirror 1
[rows, cols] = size(a)
b = []
select dir
case 'r' then
    for i = rows : -1 : 1
        b = [b; a(i, :)]
    end
case 'c' then
    for i = cols : -1 : 1
        b = [b, a(:, i)]
    end
else
    error("dir must be "r" or "c"")
end

function b = mirror3(a, dir)
// same as mirror 1
[rows, cols] = size(a)
select dir
case 'r' then
    i = rows : -1 : 1
    b = a(i, :)
case 'c' then
    i = cols : -1 : 1
    b = a(:, i)
else
    error("dir must be "r" or "c"")
end

function b = mirror4(a, dir)
// same as mirror 1
select dir
case 'r' then
    b = a($:-1:1, :)
case 'c' then
    b = a(:, $:-1:1)
else
    error("dir must be "r" or "c"");
end
Besides the performance issue discussed here the functions in Example 6-1 demonstrate how much expressiveness Scilab has got. The solutions look quite different, though they yield the same results. The benchmark results of all functions are plotted in Figure 6-1, and the discussion is found in Section 6.2.1. In brief the functions get faster from top to bottom, function \texttt{mirror1} is the slowest, \texttt{mirror4} the fastest.

### 6.1.2.1. $\$-$Constant

The last of the examples, \texttt{mirror4}, introduces a new symbol, the “highest index”, $\$ along a given direction. The dollar sign is \textit{only} defined in the index expression of a matrix. As 1 always is the lowest index, $\$ always is the highest. Please note that the dollar represents a constant, but that this constant varies across the expression! More precisely it varies with each matrix dimension. Let us make things clear by stating an example.

```plaintext
->m = [ 11 12 13; 21 22 23 ];
->m(2, $)
ans =
 23.

->m($, $)
ans =
 23.

->m(:, $/2 + 1)
ans =
  12.
  22.
```

### 6.1.2.2. Flattened Matrix Representation

The $\$ sign leads us to the flattened or vector-like representation of a matrix, if we rewrite the third line of the above example to

```plaintext
->m(1:$)^T
ans =
  11.
  21.
  12.
  22.
  13.
  23.
```

In general a \( r \times m \) matrix \( mat \) can be accessed in three ways:

- as a unit by saying \( mat \),
- by referencing its elements according to their row and column with \( mat(i, j) \), or
- via indexing into the flattened form \( mat(i) \).

The following equivalence holds: \( mat(i, j) = mat(i + (j - 1) \times n) \). Scilab follows Fortran in its way to store matrices in column-major form. See also the discussion of the function \( matrix \) in Section 6.1.3.3.

### 6.1.3. Built-In Vector-/Matrix-Functions

Functions discussed in this section:

- Colon Operator “:”
- \( \text{linspace} \)
- \( \text{logspace} \)
- \( \text{zeros} \)
- \( \text{ones} \)
- \( \text{eye} \)
- \( \text{diag} \)
- \( \text{rand} \)
- \( \text{find} \)
- \( \text{max} \)
- \( \text{min} \)
- \( \text{and} \)
- \( \text{or} \)
- \( \text{sum} \)
- \( \text{prod} \)
- \( \text{sort} \)
- \( \text{size} \)
- \( \text{matrix} \)
There are many built-in functions that work on vectors or matrices. Knowing what functions are available is important to avoid coding the same functionality with slow iterative expressions.

For further information about contemporary techniques of processing matrices with computers, the classical work “Matrix Computations” [golub:1996] is recommended.

### 6.1.3.1. Vector Generation

There are two built-in functions and one operator to generate a row-vector of numbers.

#### Vector Generation Functions

**Colon Operator “:”**

This syntax of the colon operator is

\[
\text{initial} [ : \text{increment}] : \text{final}
\]

with a default \text{increment} of +1. To produce the equivalent piece of Scilab code, we write

\[
\begin{align*}
x &= \text{initial} \\
v &= [x] \\
\text{while } x &\leq \text{final} - \text{increment} \\
  x &= x + \text{increment} \\
  v &= [v, x] \\
\text{end}
\end{align*}
\]

where \( v \) is the result. Note that the last element of the result always will be smaller or equal to the value \( \text{final} \).

**linspace**

The syntax of \text{linspace} is

\[
\text{linspace}(\text{initial}, \text{final} [, \text{length}])
\]

using a default of 100 for \text{length}. \text{linspace} returns a row-vector with \text{length} entries, which divide the interval \((\text{initial}, \text{final})\) in equal-length sub-intervals. Both endpoints, i.e. \text{initial} and \text{final} are always included.
logspace

logspace works much like linspace, and the following relation holds

\[
\text{logspace}(\text{initial}, \text{final}) = 10^{\text{linspace}(\text{initial}, \text{final})}
\]

After having discussed the most important vector generation functions, we now turn to functions that build a whole matrix at once.

### 6.1.3.2. Whole Matrix Construction

All of the functions shown in this section are capable to produce arbitrary matrices including the boundary cases of row- and column-vectors.

#### Matrix Generation Functions

**zeros**

As the name suggests this function produces a matrix filled with zeros. The two possible instantiations are with two scalar arguments

\[
\begin{align*}
    n &= 2 \\
    m &= 5 \\
    \text{mat} &= \text{zeros}(n, m)
\end{align*}
\]

or with one matrix argument

\[
\begin{align*}
    \text{mat1} &= \begin{bmatrix}
        4 & 2 \\
        4 & 5 \\
        3 & 5
    \end{bmatrix} \\
    \text{mat2} &= \text{zeros}(\text{mat1})
\end{align*}
\]

The first form produces the \( n \times m \) matrix \( \text{mat} \) made up from zeros, whereas the second builds the matrix \( \text{mat2} \) which has the same shape as \( \text{mat1} \), and is also consisting of zeros.

**Single scalar argument to zeros**

In the case of a single scalar argument \text{zeros} returns a \( 1 \times 1 \) matrix, the sole element being a zero.

Furthermore, note that
zeros()

is not allowed.

ones

The command is functionally equivalent to zeros. Instead of returning a matrix filled with 0.0 as zeros does, ones returns a matrix filled with 1.0. The only difference from the caller’s point of view is a third form which is permitted for ones, and that is calling the function without any arguments:

```matlab
->ones()
an =
  1.
```

eye

The eye function produces a generalized identity matrix, i.e. a matrix with all elements a(i, j) == 0.0 for i != j, and 1.0 for i == j. This command is functionally equivalent to zeros. The only extension is the usage without any argument, where the result automatically takes over the dimensions of the matrix in the subexpression it is used.

```matlab
->a=[2 3 4 3; 4 2 6 7; 8 2 7 4]
a =
  ! 2. 3. 4. 3. !
  ! 4. 2. 6. 7. !
  ! 8. 2. 7. 4. !
->a - 2*eye()
an =
  ! 0. 3. 4. 3. !
  ! 4. 0. 6. 7. !
  ! 8. 2. 5. 4. !
```

diag

Function diag constructs a diagonal matrix mat from the vector v, with v being mat’s main diagonal.

```matlab
->diag(2:2:8)
```
The second form of the `diag` function

\[ \text{diag}(v, k) \]

constructs a matrix that has its diagonal \( k \) positions away from the main diagonal, the diagonal being made up from \( v \) again. Therefore, \( \text{diag}(v) \) is the special case of \( \text{diag}(v, 0) \). A positive \( k \) denotes diagonals above, a negative \( k \) diagonals below the main diagonal.

\[-\text{diag}([1 1 1 1]) + \text{diag}([2 2 2], 1) + \text{diag}([-2 -2 -2], -1)\]

\[ \text{ans} = \]

\[ ! 1. \ 2. \ 0. \ 0. ! \]
\[ ! -2. \ 1. \ 2. \ 0. ! \]
\[ ! 0. \ -2. \ 1. \ 2. ! \]
\[ ! 0. \ 0. \ -2. \ 1. ! \]

`rand`

The `rand` function generates pseudo-random scalars and matrices. Again the function shares its two fundamental forms with `zeros`. Moreover, the distribution of the numbers can be chosen from `uniform` which is the default, and `normal`. The generator’s seed is set and queried with

\[ \text{rand}(’seed’, \text{new}_\text{seed}) \]

and

\[ \text{current}_\text{seed} = \text{rand}(’seed’) \]
6.1.3.3. Functions Operating on a Matrix as a Whole

find

In our opinion one of the most useful functions in the group of whole matrix functions is \texttt{find}. It takes a boolean expression of matrices (i.e. an expression which evaluates to a boolean matrix) as argument, and in form

\[
\text{index} = \text{find}(\text{expr})
\]

returns the indices of the array elements that evaluate to true, i.e. \%t in a vector. See also Section 6.1.2.2.

In the form

\[
[\text{rowidx, colidx}] = \text{find}(\text{expr})
\]

it returns the row- and column-index vectors separately. Here is a complete example.

\[
\begin{align*}
\rightarrow \text{a} &= \begin{bmatrix} 1 & -4 & 3; 6 & 2 & 10 \end{bmatrix} \\
\text{a} &= \\
&! 1. - 4. 3. ! \\
&! 6. 2. 10. ! \\
\rightarrow \text{index} &= \text{find}(\text{a} < 5) \\
\text{index} &= \\
&! 1. 3. 4. 5. ! \\
\rightarrow \text{a}(\text{index}) \\
\text{a} &= \\
&! 1. ! \\
&! - 4. ! \\
&! 2. ! \\
&! 3. ! \\
\rightarrow [\text{rowidx, colidx}] &= \text{find}(\text{a} < 5) \\
\text{colidx} &= \\
&! 1. 2. 2. 3. ! \\
\text{rowidx} &= \\
&! 1. 1. 2. 1. ! \\
\end{align*}
\]

The expressions \texttt{expr} can be arbitrarily complex, and they are not limited to a single matrix.

\[
\begin{align*}
\rightarrow \text{b} &= \begin{bmatrix} 1 & 2 & 3; 4 & 5 & 6 \end{bmatrix} \\
\text{b} &= \\
&! 1. 2. 3. ! \\
\end{align*}
\]
Chapter 6. Performance

! 4. 5. 6. !

-> a < 5
ans =
! T T T !
! F T F !

-> abs(b) >= 4
ans =
! F F F !
! T T T !

-> a < 5 & abs(b) >= 4
ans =
! F F F !
! F T F !

-> find( a < 5 & abs(b) >= 4 )
an =
4.

Last but not least find is perfectly OK on the left-hand side of an assignment. So, replacing all odd elements in a with 0 simply is

-> a(find(modulo(a, 2) == 1) ) = 0

a =
! 0. - 4. 0. !
! 6. 2. 10. !

To get the number of elements that match a criterion, just apply size(idxvec, '*') to the index vector idxvec of the find operation.

max, min

Searching the smallest or the largest entry in a matrix are so common that Scilab has separate functions for these tasks. We discuss max only as min behaves similarly.

To get the largest value saying

max_val = max(a)

is enough. The alternate form

-> [max_val, index] = max(a)
returns the position of the maximum element, too. The form of the index vector is the same as for size, i.e. [row-index, column-index]. Speaking of size, max has the forms max(mat, ‘r’), and max(mat, ‘c’), too.

```matlab
-> [max_val, rowidx] = max(b, ‘r’)
rowidx =
    ! 2.  2.  2. !
max_val =
    ! 4.  5.  6. !

-> [max_val, colidx] = max(b, ‘c’)
colidx =
    ! 3. !
    ! 3. !
max_val =
    ! 3. !
    ! 6. !
```

These forms return the maximum values of each row or column along with the respective indices of the elements’ rows or columns.

The third way of using max is with more than one matrix or scalar as arguments. All the matrices must be compatible, scalars are expanded to the full matrix size, like scalmat = scal * ones(mat). The return matrix holds the largest elements from all argument matrices.

```matlab
-> max(a, b, 3)
ans =
    ! 3.  3.  3. !
    ! 6.  5. 10. !
```

and, or

FIXME: write it!

sum, prod

FIXME: write it!
sort

FIXME: write it!

size

The size function handles all shape inquiries. It comes in four different guises. Assuming that mat is a scalar or matrix, size can be used as all-info-at-once function as in

\[ \text{[rows, cols]} = \text{size(mat)} \]

as row-only, or column-only function

\[ \text{rows} = \text{size(mat, 'r')} \]
\[ \text{cols} = \text{size(mat, 'c')} \]

and finally as totaling function

\[ \text{elements} = \text{size(mat, '*')} \]

matrix

A (hyper-)matrix can be reshaped with the matrix command. To keep things simple we demonstrate matrix with a 6x2-matrix.

\[ \text{a} = \begin{bmatrix} 1:6; 7:12 \end{bmatrix} \]
\[ \text{a} = \begin{bmatrix} 1. & 2. & 3. & 4. & 5. & 6. \\
                 7. & 8. & 9. & 10. & 11. & 12. \end{bmatrix} \]

\[ \text{matrix(a, 3, 4)} \]
\[ \text{ans} = \begin{bmatrix} 1. & 8. & 4. & 11. \\
                 7. & 3. & 10. & 6. \\
                 2. & 9. & 5. & 12. \end{bmatrix} \]

\[ \text{matrix(a, 4, 3)} \]
\[ \text{ans} = \begin{bmatrix} 1. & 3. & 5. \\
                 7. & 9. & 11. \\
                 2. & 4. & 6. \\
                 8. & 10. & 12. \end{bmatrix} \]
In contrary to the Fortran-9x function `RESHAPE`, matrix neither allows padding, nor truncation of the reshaped matrix. Put another way, for a \( m \times n \) matrix \( a \) the reshaped dimensions \( p \) and \( q \) must obey \( m \times n = p \times q \).

Matrix works by columnwise “filling” the contents of the original matrix \( a \) into an empty template of a \( p \) times \( q \) matrix. (See also Section 6.1.2.2.) If this a too hard to imagine, the second way to think of it is imagining \( a \) as a column vector of dimensions \( m \times n \) times 1 that is broken down column by column into a \( p \) times \( q \) matrix. In fact this is not pure imagination as in many situations there is the identity \( a(i,j) == a(i + n*(j - 1)) \) holds.

\[
-> a(2,4) \\
ans = \\
10.
\]

\[
-> a(8) \\
ans = \\
10.
\]

Moreover, the usual vector subscripting can be used to a matrix.

\[
-> a(:) \\
ans = \\
! 1. ! \\
! 7. ! \\
! 2. ! \\
! 8. ! \\
! 3. ! \\
! 9. ! \\
! 4. ! \\
! 10. ! \\
! 5. ! \\
! 11. ! \\
! 6. ! \\
! 12. !
\]

### 6.1.4. Evaluation Of Polynomials

Once upon a time there was a little Scilab newbie who coded an interface to the `optim` routine to make polynomial approximations easier. On the way an evaluate function for polynomials had to be written.
The author was very proud of herself because she knew the Right Thing (tm) to do in this case namely the Horner algorithm. Actually she has come up with two implementations!

Example 6-2. Naive functions to evaluate a polynomial

```matlab
function yv = peval1(cv, xv)
    // Evaluate polynomial given by the vector its coefficients cv
    // in ascending order, i.e. cv = [p q r] -> p + q*x + r*x^2
    // at all points listed in vector xv and return the
    // resulting vector.
    yv = cv(1) * ones(xv)
    px = xv
    for c = cv(2 : $)
        yv = yv + c * px
        px = px .* xv
    end
end
```

```matlab
function yv = peval2(cv, xv)
    // same as peval1
    yv = cv($);
    for i = length(cv)-1 : -1 : 1
        yv = yv .* xv + cv(i)
    end
end
```

So what is wrong with that? This code looks OK and it does the job. But from the performance viewpoint it is not optimal! The fact that Scilab offers a separate type for polynomials has been ignored. Even if we are forced to supply an interface with the coefficients stored in vectors the built-in function `freq` is preferable.

Example 6-3. Less naive functions to evaluate a polynomial

```matlab
function yv = peval3(cv, xv)
    // same as peval1, using horner()
    p = poly(cv, 't', 'coeff')
    yv = horner(p, xv)
end
```

```matlab
function yv = peval4(cv, xv)
    // same as peval1, using freq()
end
```

66
// The return value yv _always_ is a row-vector.

```plaintext
p = poly(cv, 't', 'coeff')
unity = poly(1, 't', 'coeff')
yv = freq(p, unity, xv)
```

Table 6-2 shows the speed ratios (each line is normalized separately) for a polynomial of degree 4 that we got on a P5/166 Linux system.

Table 6-2. Performance comparison of different polynomial evaluation routines

<table>
<thead>
<tr>
<th>evaluations</th>
<th>peval1</th>
<th>peval2</th>
<th>peval3</th>
<th>peval4</th>
</tr>
</thead>
<tbody>
<tr>
<td>5</td>
<td>3.5</td>
<td>4.2</td>
<td>1</td>
<td>7.0</td>
</tr>
<tr>
<td>1000</td>
<td>1.4</td>
<td>2.5</td>
<td>1</td>
<td>3.5</td>
</tr>
</tbody>
</table>

If we now decide to change our interface to take Scilab’s built-in polynomial type the evaluation with `freq` can again be accelerated by a factor of more than 3.

### 6.2. Extending Scilab

The brute force way of getting a better performance is rewriting an existing Scilab script in a low-level language as C, Fortran, or even assembler. This option should be chosen with care, because the rapid prototyping facilities of Scilab are lost. On the other hand if the interface of the function has settled, its performance is known to be crucial and it is of use in future projects then the translation into compiled code could be be worth the time and the grief.

In the first part of this section we compare different ways of integrating an external function into Scilab. We focus on the ease of integration versus the runtime overhead introduced. The second part deals with writing the low-level functions themselves, especially their interfaces.

#### 6.2.1. Comparison of the Link Overhead

We revive our matrix mirroring example from Section 6.1.2.

Our Fortran-77 version looks like this:

```fortran
subroutine mir(n, m, a, dir, b)
    *
    *   Mirror n*m-matrix a along direction prescribed by dir.
```
* If dir == 'r' then mirror along the rows, i.e. horizontally.
* Any other value for dir mirrors along the columns, i.e.
* vertically. The mirrored matrix is returned in b.
* implicit none

* ARGUMENTS
  integer n, m
  double precision a(n, m)
  character dir(*)
  double precision b(n, m)

* LOCAL VARIABLES
  integer i

* TEXT
  if (dir(1:1) .eq. 'c') then
    do 100, i = 1, m
      call dcopy(n, a(1, m+1-i), 1, b(1, i), 1)
    100  continue
  else
    do 200, i = 1, n
      call dcopy(m, a(n+1-i, 1), n, b(i, 1), n)
    200  continue
  end if
end

function b = mirf(a, dir)
// interface function for 'mir.f'
// Behavior is the same as mirror()

[n, m] = size(a)
b = zeros(n, m)

if dir == 'r' | dir == 'c' then
  b = fort('mir', ..
    n, 1, 'i', m, 2, 'i', a, 3, 'd', dir, 4, 'c', ..
    'out', ..
    [n, m], 5, 'd')
else
    error("dir must be "'r" or "c"')
end

OK, let’s lock-and-load. We are ready to rock!

    link('mir.o', 'mir')
    getf('mirf.sci')

The fast alternative to using **fort**, which dynamically creates an interface to a C or Fortran function is using **intersci**, which which creates an interface suitable for static loading.

**intersci** can create the Fortran glue code for a C or Fortran function to make it callable form the Scilab interpreter. The glue code is compiled (with a Fortran compiler) and linked to Scilab. **intersci** is described very well in the SCI/doc/Intro.ps. Anyhow, here is the description file for our current example. Finally it will supply us with a Scilab function called **mirai(a, dir)**.

```
iris  a  dir
   a     matrix  n  m
   dir    string  l
   b     matrix  n  m

mir  n  m    a  dir    b
   n  integer
   m  integer
   a  double
   dir  char
   b  double

out  sequence  b
```

We do not want to go into detail here, but a desc-file has three parts separated by blank lines: The description of the Scilab-level function’s signature (here: **mirai**), the same for the low-level function (here: **mir**), and finally the results’ structure. The signatures resemble Fortran or K&R-style C function definitions with the parenthesis missing. The process of passing a desc-file through **intersci**, compiling the low-level routine and the glue code can be automated. Example 6-4, a snippet of our Makefile.intersci shows the relevant rules.

**Example 6-4. Makefile for static Scilab interfaces via intersci**

```
ifdef SCI
    SCIDIR := $(SCI)
else
```
Chapter 6. Performance

```
SCIDIR := /site/X11R6/src/scilab-2.5
endif

%.f.pre: %.desc
    $(SCIDIR)/bin/intersci $*
    mv $*.f $*.f.pre

%.f: %.f.pre
    perl -pe 's#SCIDIR#$(SCIDIR)#' $< > $@

%.o: %.f
    $(FC) $(FFLAGS) -c $<
```

Running the automatically generated Fortran code through a filter (here: `perl`) is necessary to fix the lines `include 'SCIDIR/routines/stack.h'`. After everything is compiled a single Scilab command makes the new routine available to the user.

```
addinter([
    'mirai.o', 'mir.o',
    'mirai', 'mirai'
])
```

The first argument which almost always is a vector of strings tells Scilab the names of the object files to load. One of them is the interface code made by `intersci`. The rest are the user routines. The second argument specifies name of entry point into the interface routine. The third parameter is the name the new Scilab function will carry.

---

**Entry point of interface function**

`addinter`s second argument must be the name of the interface routine, i.e. the one generated by `intersci`. Using the low-level function’s entry point here causes Scilab to barf.

---

Why do we go through that tedious process? After all we are in the performance section, so what we want is speed, high speed, or even better the ultimate speed. Now we can compare all the variants as is done in Figure 6-1.
Performance comparison of \texttt{mirror[1-4]} and \texttt{mirai} on a P5/166 Linux box.
Performance comparison of mirror[1-4], and mirai on a 2-way PIII/550 Linux box.

If we compare the performance of our three Scilab mirror routines mirror1, mirror2, and mirror3 together with the two incarnations of the hard-coded routine mirf, and mirai, we reach the following conclusions.

- Scilab code that makes heavy use of indexing, like mirror1, is extremely slow no matter what problem size. Thumbs down on that one.

- Well written i.e. index-free Scilab code, like mirror4, performs very well. This is especially true for large vectors or matrices.

- The overhead of the fort-call in mirf is high; it is hard to amortize for that. fort is only justified in situations where a significant amount of time is spent in the low-level user-routine. Usually this will be the case for large problem sizes. Of course the cross-over point has to be determined separately in each case.

- Nothing can beat a compiled function that is integrated with addinter. mirai surpasses all other implementations. For small problem sizes the little overhead in comparison to all the other functions gives this function a factor 10 advantage, though, as the problems size increases mirai's lead is challenged by mirror4.
Conclusion: Never underestimate the power of the Emperor^H^H^H^H^H^H vectorized Scilab code.

6.2.2. Preparing And Compiling External Subroutines

In this section we will discuss the interfacing of C, C++, Fortran-77, Fortran-9x, or Ada routines with Scilab via `link` command. We restrict ourselves to the simple case of functions that expect exactly one double precision floating point parameter and return a double precision floating point result. Functions with that signature are required e.g. for the integration routine `intg`, or the root finder `fsolve`.

Before we dive into the language specific descriptions, let us point out the main features of Fortran we have to pay attention to when writing an interface in (another) language.

Function name mangling

A function named `FOO` (foo, or whatever capitalization is chosen) in the Fortran source can become a different symbol on code generation. This is compiler dependent. Most often an underscore “_” is prepended or appended. Sometimes the name is downcased, sometimes it is upcased.

**Tip:** The `nm()` command provides easy access to the symbols in an object file.

Call-by-reference

Fortran never passes the value of a parameter, but always a pointer to the parameter.

Arrays in column-major order

Arrays are stored so that their leftmost index varies fastest.

6.2.2.1. Fortran-77

Fortran-77 or, how do you want to ruin your day?

`lvd`

Extending Scilab with Fortran-77 is most straightforward. A Fortran-77 source for function `FALS` could look like this:

```fortran
double precision function fals(x)
```
double precision x
fals = sin(10.0d0 * x)
end

After compilation (e.g. `f77 -c fals.f`) the compiled code can be linked to Scilab and called with the integration routine.

```plaintext
link('fals.o', 'fals');
[res, aerr, neval, info] = ..
intals(0.0, 1.0, -0.5, -0.5, 'alg', 'fals')
```

### 6.2.2.2. Fortran-9x

*Fortran-90? Don’t worry, it can’t get much worse.*

A bloated, but portable Fortran-90 source for a function could look like this:

```plaintext
function fsm(x)
implicit none
integer, parameter :: idp = kind(1.0d0)
!
! arguments/return value
real(kind = idp), intent(in) :: x
real(kind = idp) :: fsm
!
! text
fsm = exp(x) / (1.0d0 + x*x)
end function fsm
```

After compilation (e.g. `f90 -c fsm.f90`) the compiled code can be linked to Scilab and called with an integration routine.

```plaintext
link('fsm.o', 'fsm');
[ires, ierr, neval] = intsm(0.0, 1.0, 'fsm')
```

### 6.2.2.3. (ANSI-) C

A simple C function meting out signature requirements has e.g. this shape:
#include <math.h>

double fgen(const double *x)
{
    if (*x > 0.0)
        return 1.0 / sqrt(*x);
    else
        return 0.0;
}

After compilation (e.g. `cc -c fgen.c`) the compiled code can be linked to Scilab and called with the integration routine.

```
link('fgen.o', 'fgen', 'c');
[ires, ierr, neval, info] = intgen(0.0, 1.0, 'fgen')
```

There are several ways to get the naming convention differences between Fortran and C right. We show three possible solutions for the case where C uses no decoration at all and Fortran appends one underscore.

```
/* (1) GNU C compiler */
double foo(const double *x) __attribute__((weak, alias ("foo_")));

/* (2) good preprocessor */
#define C2F(name) name_

/* (3) old preprocessor ;-) */
#define ANOTHERC2F(name) name/**/_
```

### 6.2.2.4. C++

A C++ source for a function could look like this:

```
#include <math.h>

extern "C" {
    double fgk(const double *x);
}

double fgk(const double *x)
{
    return 2.0 / (2.0 + sin(10.0 * M_PI * (*x)));
After compilation (e.g. `c++ -c fgen.c`) the compiled code can be linked to Scilab and called with the integration routine.

```
link('fgk.o', 'fgk', 'c');
[ires, ierr, neval, info] = ..
    intgk(0.0, 1.0, 'fgk', 0, %eps, '15-31')
```

### 6.2.2.5. Ada

For GNAT/Ada the package’s interface part pulls in the Fortran interface definitions. Is the simplest case the mathematical functions are only instantiated with the type `Double_Precision`. Ada requires to export every function’s interface separately, as is clear from the following example.

```ada
with Interfaces.Fortran;
use Interfaces.Fortran;
with Ada.Numerics.Generic_Elementary_Functions;

package TestFun is
    package Fortran_Elementary_Functions is new
        Ada.Numerics.Generic_Elementary_Functions (Double_Precision);
    use Fortran_Elementary_Functions;

    function foo(x : Double_Precision) return Double_Precision;
    pragma Export(Fortran, foo);
    pragma Export_Function(Internal => foo,
                            External => "foo_",
                            Mechanism => Reference,
                            Result_Mechanism => Value);
end TestFun;
```

According to the interface specification the package body looks like this:

```ada
package body TestFun is
    function foo(x : Double_Precision) return Double_Precision is
        begin
            return exp(x) / (1.0 + x*x);
        end foo;
    end TestFun;
```

The package is compiled as usual `gnatmake -O2 testfun.adb`. 


Make sure that there is a GNAT runtime library libgnat-3.12p.so. Your version number may be different. The ending so is critical, as libgnat-3.12p.so.1.7 will not make dlopen(3) happy. From now on everything is downhill and the function can be linked almost as usual.

Example 6-5. Linking (GNAT) Ada-code

```plaintext
link("testfun.o -L/site/gnat-3.12p/lib -lgnat-3.12p", "foo")
```

Again, the path to your gnat-library and the version numbers can differ.

In the case of several functions in the package it is preferable to rely on the extended dlopen(3) mechanism, and link the package/library combo with remembering the id of the shared library.

```plaintext
adacode = link("testfun.o -L/site/gnat-3.12p/lib -lgnat-3.12p", "foo")
```

Linking further functions from the library happens by referencing the number of the library.

```plaintext
link(adacode, "bar")
```

This saves space (Scilab’s TRS) and time (to execute the link). Speaking about saving... Users with a loader e.g. GNU ld, capable of incremental linking (e.g. -i, -r, -relocatable) can of course link testfun.o with the library before linking everything to Scilab. To complete the example, here comes the command-line:

```plaintext
ld -i -o testfun-lib.o testfun.o -L/site/gnat-3.12p/lib -lgnat-3.12p
```

In Scilab the arguments to link reduce to

```plaintext
link("testfun-lib.o", "foo")
```

### 6.2.3. Pushing It Further

What? What are you doing in this section? Still not satisfied with your functions’ performance?—Sorry, but there are no conventional ways to get more out of Scilab. Tinkering with the interface routines is not worth the effort. Some completely new approach is necessary.

If a problem is too tough, Scilab still can serve as a rapid prototyping environment. One sister program of Scilab, namely Tela (http://www.geo.fmi.fi/prog/tela.html) has been written for exactly this purpose. Prototyping with an interpreted language is currently going through a big revival with C (and C++) developers discovering Python.
As whenever optimization is the final goal, an extensive test suite is the base for success. So one way to proceed could be to develop test routines and reference implementation completely in Scilab. The next step is rewriting the routines *still* in Scilab to match the signatures of for example BLAS/LAPACK routines as closely as possible. The test suite can remain untouched in this step. The final step is to migrate the Scilab code to Fortran, C, or whatever, while making extensive use of BLAS/LAPACK. Ideally the test suite remains under Scilab and can be used to exercise the new standalone code.

### 6.3. Building an Optimized Scilab

One relatively easy way to increase Scilab’s performance is recompiling it with a good compiler and an optimized BLAS library. See also Section 8.3 to find out what optimized BLAS kernels are available and where to get them.

Our experience only suffices to explain the compilation on an ia32 Linux system. Here, gcc-2.95 (http://gcc.gnu.org) or pgcc-2.95 (http://www.goof.com/pgc/) are the compilers of choice.

The following options are a good starting point for further exploration. They apply to compiling Fortran and C code.

`-march=arch`

This option instructs gcc to generate code specifically for architecture *arch*. Among other things it sets `-mcpu=arch`. Furthermore, it forces `-malign-loops`, `-malign-jumps`, `-malign-functions`, and `-mpreferred-stack-boundary` to their optimum values for the selected architecture *without braking* the ABI. Therefore, it can be considered an optimization switch.

`-malign-double`

For systems with an original Intel P5 or above processor this option is an absolute *must*. Though it breaks the ABI, the gain in speed due to avoiding the misalignment penalty for 64bit-floats is tremendous, even on PPro (and derivative) systems with all write back caches enabled.

`-O2`

The workhorse optimization switch, `-O2`, activates a lot of optimizations. See node “Optimize Options” in gcc’s info file, e.g. `info -f/usr/info/gcc.info.gz -n "Optimize Options"`

The optimizations toggled on by `-O2` are well tested and do not produce excessively long text.
-funroll-all-loops

This switch increases the text size, speeding up some loops. YMMV.

Notes

1. Remember that the colon operator returns a row-vector.
Chapter 7. Scilab Core


Good morning Mister Tyler! Going down?

We are going down all the way right to the core, the core of Scilab. Though this is the most technical and most complex chapter, it is by no means true that writing a native Scilab function is unmanageable by for ordinary mortals. A strict programming discipline, patience, persistence, and a thorough knowledge of what makes up the stack-structures involved, let us overcome the difficulties.

7.1. Introduction To pAda

Instead of simply repeating the Fortran-77 statements that make up the Scilab stack, the API, etc., we introduce a new language that is better suited for this job: a pseudo Ada¹ (pAda), which is much more expressive. The syntax follows Ada, and the pAda types are mapped onto Fortran-77 types as listed in Table 7-1. What might look like an artificial complication, the introduction of new types, actually is a major simplification. First, the name of the type now makes clear exactly what it is used for. Second, distinct types designate distinct things, stuff that never should be mixed up. Third, the valid ranges of the sub-types are explicity mentioned.

Table 7-1. pAda to Fortran-77 type mappings

<table>
<thead>
<tr>
<th>pAda</th>
<th>Fortran-77</th>
</tr>
</thead>
<tbody>
<tr>
<td>Integer</td>
<td>INTEGER</td>
</tr>
<tr>
<td>Float</td>
<td>DOUBLE PRECISION, REAL*8</td>
</tr>
<tr>
<td>Boolean</td>
<td>LOGICAL</td>
</tr>
<tr>
<td>Character</td>
<td>CHARACTER</td>
</tr>
<tr>
<td>type String is array (1..N) of Character</td>
<td>CHARACTER*N</td>
</tr>
<tr>
<td>type ComplexFlag is (RealVariable, ComplexVariable)</td>
<td>INTEGER = 0, 1</td>
</tr>
<tr>
<td>subtype Natural is Integer range 0..Integer’Last</td>
<td>INTEGER</td>
</tr>
<tr>
<td>type ParameterStackAddress is new Integer range 1..Integer’Last</td>
<td>INTEGER</td>
</tr>
</tbody>
</table>
7.2. Internal Data Structure

FIXME: explain the parameter stack, data stack, etc.

7.2.1. Parameter Stack And Data Stack

FIXME: follow the docu in Internals

7.2.2. Storage of Complex Variables

FIXME: explain separate storage of two DOUBLE PRECISION parts instead of one DOUBLE COMPLEX

7.3. Writing Native Scilab Functions

In the following two sections we shall treat the “anatomy” of native, i.e. low-level Scilab functions. This will confront us with all the gory details of the stack, the low-level APL and the calling conventions. Having the “Guide for Developers”.Internals.ps (see also Section 8.2) ready is a good idea. Where the developer guide is at the end of its wits, a study of the source code is appropriate, especially the file SCI/routines/interf/stack1.f

We start out discussing simple functions. Simple in the sense that they are self-contained and only take non-function parameters as their arguments. In the second part we shall consider functions that take other functions (either Scilab functions or externals) as arguments.

7.3.1. Simple Functions

A typical native Scilab function proceeds as follows:

1. Check the number of input and output parameters.
2. Get the “pointers” to all actual input parameters; supply default values for optional parameters; issue warnings or errors as appropriate for too many or too few parameters.

3. Allocate space for all temporary variables, “workspaces”, etc.

4. It might be necessary to translate the input variables which are in Scilab format into the appropriate format for the worker routine. This is necessary for example if the worker routine uses \texttt{DOUBLE} \texttt{COMPLEX} (or \texttt{COMPLEX*16}) variables.

5. Perform the calculations or transformations that \textit{really} make up the procedure.

6. As in Step 4, it might be necessary to transform the results, now form the worker routine’s format back into Scilab format.

7. If necessary, allocate space on the Scilab stack and copy results to this space.

Now that the outline is clear we are ready to dissect a simple function: \texttt{ortho}. The function takes exactly one argument \(a\), that is a real or complex \(m \times n\) matrix. The single output parameter is a matrix of the same shape and type is the input matrix. The duty of \texttt{ortho} is to bring the columns of the input matrix into orthonormal form; to achieve this we employ the following LAPACK functions:

```plaintext
type Complex is record
   Re, Im : Float’Base;
end record;

type FloatVector is array (Positive range <>) of Float;
type ComplexVector is array (Positive range <>) of Complex;
type FloatMatrix is array (1..Lda, Positive range <>) of Float;
type ComplexMatrix is array (1..Lda, Positive range <>) of Complex;

procedure dgeqrf
   (M : in Natural;  
    N : in Natural;  
    A : in out FloatMatrix;  
    Lda : in Natural;  
    Tau : out FloatVector;  
    Work : out FloatVector;  
    Lwork : in Integer;  
    Info : out Integer);

procedure dorgqr
   (M : in Natural;  
    N : in Natural;  
    K : in Natural;  
    A : in out FloatMatrix;  
    Lda : in Natural;  
    Tau : out FloatVector;
```

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The two $[d|z]$geqrf-functions compute a QR-factorization of a real or complex $m$-by-$n$ matrix $a$, while the $[dor|zun]$gqr-functions generate an $m$-by-$n$ real or complex matrix $q$ with orthonormal columns. 
dcopy copies $N$ elements of the vector $X$ in increments of $IncX$ into the vector $Y$ using increments of $IncY$ on the output side. For a detailed description please consult the LAPACK User Guide or the appropriate manual pages.

Example 7-1 is one of the longest examples in the running text, but don’t be scared as we will explain line-by-line and variable-by-variable what is where and why.
Example 7-1. Simple native Scilab function

    subroutine ortho

Native functions are parameterless

    implicit none

Switch into weeny mode :-) 

* CONSTANTS
    integer realtype
    parameter (realtype = 0) – See Table 7-1 for type association

* LOCAL VARIABLES
    character*6 fname
    The name of the routine as string

        logical checklhs, checkrhs, creat, getmat
    Scilab API functions

    integer topk
    integer n, m, mattyp
    integer tausz, worksz, info
    integer areadr, aimadr, badr, tauadr
    integer wrkadr, rreadr, rimadr, dumadr

* EXTERNAL FUNCTIONS/SUBROUTINES
    external checklhs, checkrhs, creat, getmat
    Scilab API functions

    external error

    external dcopy, dgeqrf, dorgqr, zgeqrf, zungqr
    LAPACK/BLAS worker subroutines

* HEADER
    include '/site/X11R6/src/scilab/routines/stack.h' – Scilab API header

* TEXT
    fname = ‘ortho’
    Function name (for error messages)
    topk = top
    top is defined in stack.h
    rhs = max(0, rhs)

        if (.not. checkrhs(fname, l, l)) return ❶
if (.not. checklhs(fname, 1, 1)) return

* fetch input parameters
  if (.not. getmat(fname, topk, top - rhs + 1, $
      mattyp, m, n, areadr, aimadr)) return

    if (n * m .eq. 0) return

    Quick return on empty matrix
    tausz = min(m, n)

    Prescribed by man-page
    worksz = max(1, n)

    if (mattyp .eq. realtype) then
      real case

    * allocate temporary variables; all are real
      if (.not. cremat(fname, top + 1, realtype, tausz, 1, $
      tauadr, dumadr)) return
      if (.not. cremat(fname, top + 2, realtype, worksz, 1, $
      wrkadr, dumadr)) return
      if (.not. cremat(fname, top + 3, realtype, m, n, $
      badr, dumadr)) return

    * prepare worker routines’ input parameters
      call dcopy(n * m, stk(areadr), 1, stk(badr), 1)

    * call worker routines
      call dgeqrf(m, n, stk(badr), m, stk(tauadr), $
      stk(wrkadr), worksz, info)
      if (info .ne. 0) then
        Any error is considered fatal
        buf = fname // ’dgeqrf failed’
        call error(999)
        return
      endif

      call dorgqr(m, n, tausz, stk(badr), m, stk(tauadr), $
      stk(wrkadr), worksz, info)
      if (info .ne. 0) then
        Any error is considered fatal
        buf = fname // ’dorgqr failed’
        call error(999)
        return
endif

else
  complex case; mattyp != realtype

  allocate temporary variables,
  * use two REAL*8 for one COMPLEX*16 ➊
    if (.not. cremat(fname, top + 1, realtype, 2 * tausz, 1, $
      tauadr, dumadr)) return
    if (.not. cremat(fname, top + 2, realtype, 2 * worksz, 1, $
      wrkadr, dumadr)) return
    if (.not. cremat(fname, top + 3, realtype, 2 * m, 2 * n, $
      badr, dumadr)) return

  prepare worker routines’ input parameters, joining
  * two REAL*8 arrays into one COMPLEX*16 array ➋
    call dcopy(n * m, stk(areadr), 1, stk(badr), 2)
    call dcopy(n * m, stk(aimadr), 1, stk(badr + 1), 2)

  call worker routines ➌
    call zgeqrf(m, n, stk(badr), m, stk(tauadr), $
      stk(wrkadr), worksz, info)
    if (info .ne. 0) then
      Any error is considered fatal
        buf = fname // ’zgeqrf failed’
        call error(999)
        return
    endif

    call zungqr(m, n, tausz, stk(badr), m, stk(tauadr), $
      stk(wrkadr), worksz, info)
    if (info .ne. 0) then
      Any error is considered fatal
        buf = fname // ’zungqr failed’
        call error(999)
        return
    endif

endif

* get ready to exit
if (lhs .ge. 1) then ➏
  if (.not. cremat(fname, top, mattyp, m, n, $
    rreadr, rimadr)) return
if (mattyp .eq. realtype) then
   (10)
   call dcopy(m * n, stk(badr), 1, stk(rreadr), 1)
else
   call dcopy(m * n, stk(badr), 2, stk(rreadr), 1)
   call dcopy(m * n, stk(badr + 1), 2, stk(rimadr), 1)
endif
endif
end

1. Check the number of input and output parameters. Here the task is easy as we need one and write one. This line and the next correspond to Step 1.

2. Get the addresses as mentioned in Step 2 of the real and imaginary part of the matrix passed as only parameter to ortho. Note that getmat will return False if the parameter at the given parameter stack position is not a matrix of numbers.

   getmat is called with the second parameter, topk holding the value of the parameter stack pointer when the control flow entered ortho. This as well as the function name passed in fname is necessary for the cleanup and messaging in case of an error.

   The only parameter we use is on top of the parameter stack as top - rhs + 1 equals top in our case.

   On successful return getmat not only sets the data stack addresses areadr, and aimadr, but also tells us via mattyp whether the matrix is real complex, and via m, and n how large the matrix is.

   The following lines directly depend on the sizes passed back, calculating the necessary space for two scratch arrays.

3. Allocating space for the temporary variables tau, work, and b on the data stack is Step 3. tau and work are necessary because of the LAPACK routines used; b is a copy of a as the LAPACK routine works with the matrix in place, i.e. would mangle the input variable a. The temporaries are accessed the same way parameters are accessed: through indices into the data stack. These indices are tauadr, wrkadr, and badr. Their position on the parameter stack is top + 1, top + 2, and top + 3, respectively.

   We request a purely real storage for each of the three temporary variables with the third parameter being realtype = 0. Therefore the index for the imaginary part is a dummy index, dumadr.

   The sizes of the vectors or matrices have been computed before.

4. There is no “translation” to do in the real case. So Step 4 is easy. The input variable – of which we definitely know that it is real – is simply copied into the scratch space we have allocated on the data stack.
Chapter 7. Scilab Core

Note how the address of the matrices is passed. The idiom is \texttt{stk(index)}, where \texttt{index} has been obtained through a \texttt{get*-} or \texttt{cre*-} function. The mnemonic “stk” means data stack.

Everything is set up correctly and initialized. We have reached Step 5. The worker routines can take over now.

In the complex case the allocation of the temporaries variables requires a bit more thought, although it is again just Step 3. We know that the LAPACK routines need the complex vectors/matrices in packed form. Thus, we allocate one real (DOUBLE PRECISION) vector/matrix of twice the size each time thereby accommodating the the storage requirement of complex (DOUBLE COMPLEX) variables. Otherwise this step proceeds as in the real case.

Due to the different handling of complex variables in Scilab and in LAPACK, Step 4 requires two separate calls to the copy function.

\begin{verbatim}
call dcopy(n * m, stk(areadr), 1, stk(badr), 2)
call dcopy(n * m, stk(aimadr), 1, stk(badr + 1), 2)
\end{verbatim}

The first line says: “Copy \texttt{m} times \texttt{n} elements from the first position in the DOUBLE PRECISION variable \texttt{stk(areadr)} taking each entry (3rd parameter, read stride: 1) into the DOUBLE COMPLEX output variable \texttt{stk(badr)} filling every other entry (5th parameter, write stride: 2).” The second line does almost the same, but starts off writing at the second element \texttt{stk(badr + 1)}, therefore filling the imaginary parts into \texttt{stk(badr)}. This corresponds to Step 4.

Again we have reached Step 5; everything is set up correctly and initialized, and the worker routines can take over.

If there is an output variable, we copy the results into it. Otherwise, we skip the expensive copy operation.

(10) At this point a purely real result, \texttt{stk(rreadr)}, can simply be copied onto the output parameter, \texttt{stk(badr)}.

The situation is a bit more complicated for a complex result, as we have to split the DOUBLE COMPLEX result from LAPACK into two DOUBLE PRECISION matrices. Here are the crucial lines again:

\begin{verbatim}
call dcopy(m * n, stk(badr), 2, stk(rreadr), 1)
call dcopy(m * n, stk(badr + 1), 2, stk(rimadr), 1)
\end{verbatim}

The first line says: “Copy \texttt{m} times \texttt{n} elements from the first position in the DOUBLE COMPLEX result \texttt{stk(badr)} taking every other entry (3rd parameter, read stride: 2) into the DOUBLE PRECISION output variable \texttt{stk(rreadr)} filling each entry (5th parameter, write stride: 1).” The second line does almost the same, but starts off at the second element \texttt{stk(badr + 1)}, therefore copying the imaginary parts into \texttt{stk(rimadr)}. This way we are merging Step 6, and Step 7 into one.
7.3.2. Functionals

FIXME: write it!

7.4. Error Handling

We briefly discuss how to produce the three possible classes of errors: fatal, warning, and message in Scilab.

7.4.1. Fatal Errors

To signal a fatal error condition in an interface procedure, call `error` with the appropriate code. The codes can be looked up in SCI/routines/system/error.f.

Here is a code snippet that does this.

```fortran
if (ifail .eq. 2) then
   call error(1232)
   return
endif
```

If there is no suitable error message, place your own message (length <= 80 chars) in the global variable `buf`, and call `error` afterwards.

**Warning**
The string placed in `buf` must not be longer than 80 characters.

```fortran
if (ier .eq. 6) then
   buf = 'invalid limits'
   call error(32253)
   return
endif
```

Sideffect of calling `error`: The Scilab stack is cleaned up, it put back in the state it was before the interface routine has been entered.

On the Scilab interpreter level an error terminates the evaluation of whatever is currently evaluated (expression, file, or string), unless the trapping of errors has been modified by `errcatch`. See also: `errclear`, and `iserror`.

7.4.2. Warnings

To signal non-fatal error conditions (also known as soft-errors, or warnings), place a negative integer in \texttt{err2} and call \texttt{out} to display your warning message. Depending on the situation a \texttt{return} may be issued after that. The Scilab stack is \textit{not} clean up, which means all return values from the interface routine are passed back normally. This is the solution of choice if the user can decide how to proceed based on the return values.

Again, here is a small piece of code for demonstration.

```plaintext
if (fail .eq. 1) then
  err2 = -6343
  call out(‘reached table limit’)
  return
endif
```

On interpreter level it is now mandatory to call \texttt{iserror} after a call to a routine that issues warnings like this. In the user-level error handler the error code \textit{must} be reset by \texttt{errclear} to allow for further warnings to be signaled.

A typical way of coping with these soft-errors in the interpreter level is shown in Example 7-2.

\textbf{Example 7-2. Handling of warnings in Scilab}

```plaintext
[z, n, info] = abraxas(a, b, foo, limit)
if iserror(-19) then
  errclear(-19)
  limit = limit / 2 // make it easier
  [z, n, info] = abraxas(a, b, foo, limit)
if iserror(-19) then
  errclear(-19)
  error(failed even with easy limit’);
end
end
```

7.4.3. Messages

Messages are the least severe class of errors. Sometimes they are not really errors, but just an additional information that something unexpected is going on. No news is good news.

We have already seen the appropriate subroutine in action. It is \texttt{out}.

```plaintext
if (iter .gt. 1000) then
```
7.5. Interface to Scilab’s Core

The interface to Scilab’s core is widely undocumented. To save the reader frequent lookups in the defining file, SCI/routines/interf/stack1.f, we have compiled the most important ones in the following sections: query, access and creation of objects.

7.5.1. Query

The functions in this group allow for retrieval of information about the parameters a function has been called with, and about the properties of objects on the stack.

7.5.1.1. checkrhs

Synopsis

function CheckRhs
  (FunctionName : in String;
   MinNumParameter : in Integer;
   MaxNumParameter : in Integer)
return Boolean;

Description

Check the number of actual parameters on the right-hand side to be in the range MinNumParameter : MaxNumParameter. Return true if it is in the range, otherwise raise error 77 associated with FunctionName.

Note that a function that is called without any parameters, i.e. Foo(), gets an Rhs of -1.

The semantics of CheckRhs are slightly goofy. If the number of actual input parameters is in the specified range, CheckRhs returns True, but it never returns False as it raises an error in this case.

Example

Ensure that at least 2, but not more than 5 parameters are passed to the function:

if (.not. checkrighs(fname, 2, 5)) return

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We have assumed that \texttt{fname} is set to the function’s name.

See also

CheckLhs, Rhs, Lhs

### 7.5.1.2. checklhs

**Synopsis**

```plaintext
function CheckLhs
    (FunctionName : in String;
     MinNumParameter : in Integer;
     MaxNumParameter : in Integer)
    return Boolean;
```

**Description**

Check the number of output variables, i.e. arguments on the right-hand side to be in the range \texttt{MinNumParameter} : \texttt{MaxNumParameter}. Return true if it is in the range, otherwise raise error 78 associated with \texttt{FunctionName}.

Note that it is no error to supply less output parameters than the function actually yields. The extra values are silently discarded. This is true for the case of zero output values, too; then \texttt{ans} gets the first output value. So, a function called without any output parameters gets an \texttt{Lhs} of 1.

The semantics of \texttt{CheckLhs} are slightly goofy. If the number of actual output parameters is in the specified range, \texttt{CheckLhs} returns \texttt{True}, but it never returns \texttt{False} as it raises an error in this case.

**Example**

Ensure that there are not more than 2 output parameters, when the function is called:

```plaintext
if (.not. checklhs(fname, 1, 2)) return
```

We have assumed that \texttt{fname} is set to the function’s name.

See also

CheckRhs, Rhs, Lhs
7.5.1.3. lhs

Synopsis

lhs : Integer

Description

The number of actual output parameters (i.e. those on the left-hand side of the assignment operator) is stored in the global variable lhs.

See also

CheckLhs, CheckRhs, Rhs

7.5.1.4. rhs

Synopsis

rhs : Integer

Description

The number of actual input parameters (i.e. those on the right-hand side of the assignment operator) is stored in the global variable rhs.

See also

CheckLhs, CheckRhs, Lhs

7.5.2. Access Object

The functions in this section grant the programmer access to parameters that are stored on the Scilab stack. The general working is always the same: An index to the current (i.e. as on entry of the function) top of the parameter stack, “BasePointer”, and an index to the desired argument, “StackPointer”, are passed to the API. On return the user gets all necessary information about the argument like sub-type and dimension and as important indices, “FooIndex”, into the Scilab heap which act like pointers to the actual contents. This way only meta-data is passed, saving time-consuming copy operations.
7.5.2.1. getmat

Synopsis

function GetMat
(FunctionName : in String;
BasePointer : in ParameterStackAddress;
StackPointer : in ParameterStackAddress;
IsComplex : out ComplexFlag;
Rows : out Natural;
Columns : out Natural;
RealIndex : out DataStackIndex;
ImaginaryIndex : out DataStackIndex)
return Boolean;

Description

Retrieve the address(es) and dimensions of a real or complex matrix from the parameter stack. The BasePointer must be set to the parameter stack pointer’s value on entry of the calling function. StackPointer points to the desired parameter on the parameter stack. If successful, GetMat returns True, and IsComplex, Rows, Columns, and RealIndex have valid values. If IsComplex = ComplexVariable then ImaginaryIndex is valid, too. If the parameter indexed by StackPointer is not a matrix then GetMat returns False.

The output parameter IsComplex indicates whether the matrix on the data stack is purely real or complex. In the first case RealIndex points to the matrix, in the second case RealIndex points to the real part of matrix, and ImaginaryIndex points to the imaginary part. In any case Rows and Columns are the number of rows and columns in the matrix.

Example

Fetch the addresses of a possible complex m times n matrix from position top of the parameter stack.

if (.not. getmat(fname, topk, top, iscmpx, m, n, are, aim)) return

It is assumed that fname has been set to the function’s name, and topk carries the position of the stack on entry to the calling function.

See also

GetRMat, GetRVect, GetVect
### 7.5.2.2. getrmat

**Synopsis**

```haskell
function GetRMat
  (FunctionName : in String;
   BasePointer  : in ParameterStackAddress;
   StackPointer : in ParameterStackAddress;
   Rows         : out Natural;
   Columns      : out Natural;
   RealIndex    : out DataStackIndex)
return Boolean;
```

**Description**

### 7.5.2.3. getrvect

**Synopsis**

```haskell
function GetRVect
  (FunctionName : in String;
   BasePointer  : in ParameterStackAddress;
   StackPointer : in ParameterStackAddress;
   Rows         : out Natural;
   Columns      : out Natural;
   RealIndex    : out DataStackIndex)
return Boolean;
```

**Description**

### 7.5.2.4. getvect

**Synopsis**

```haskell
function GetVect
  (FunctionName : in String;
   BasePointer  : in ParameterStackAddress;
   StackPointer : in ParameterStackAddress;
   IsComplex    : out ComplexFlag;
```

```
7.5.2.5. getscalar

Synopsis

function GetScalar
  (FunctionName : in String;
   BasePointer  : in ParameterStackAddress;
   StackPointer : in ParameterStackAddress;
   Index        : out DataStackIndex)
return Boolean;

Description

7.5.3. Create Object

The object creation functions are mainly used to setup temporary variables for the current procedure or the procedures to be called; they bear a lot of resemblance with the object access functions (see also Section 7.5.2). The difference is that a new object is created and therefore stack space is used.

7.5.3.1. Cremat

Synopsis

function Cremat
  (FunctionName : in String;
   StackPointer : in ParameterStackAddress;
   WantComplex  : in ComplexFlag;
   Rows         : in Natural;
   Columns      : in Natural;
Notes

1. We apologize to all Ada programmers for the abuse of the language, but Ada’s expressiveness and clarity are unmatched.
Chapter 8. Further Information

8.1. Coping With Scilab

Scilab is a large package no doubt about that. The source for version 2.5 comprises of more than 48 MB, and builds to over 88 MB on a i386-Linux system.

8.1.1. Distribution Size

We use several tools to cope with Scilab’s size and complexity. The most important ones are introduced in the following section.

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8.1.1.2. locate

The locate(1) command is the fast brother of the find(1) command. More precisely, locate accesses a precomputed database of filenames (usually /var/lib/locatedb; for its structure see locatedb(5)).

The database is generated by updatedb(1) with a find / -print and then mangled for faster access.

We have found a local filename database very useful for the work with large projects. Therefore, we setup two aliases that create and access a project-specific list of filenames.

```bash
alias upd='updatedb -output=./.locatedb -localpaths=.'
alias loc='locate -database=./.locatedb'
```

The upd sequence is typically run after a CVS checkout, add, or remove in the directory SCI.

We run loc whenever we are looking for a file in the Scilab distribution. This is much faster than running find every time, especially when working with a slow file server. The only inconvenience remaining is that loc must be executed in the directory where the database resides, here: SCI. However this is more
than compensated by the fact that `locate` does a substring search, i.e. given the filename `fpat` it returns all file- and directory names matching `*fpat*`.

If we want to scan the complete database and postprocess the output with our tools-of-choice, issuing a `loc` . and piping the output through the desired filters does the job.

### 8.1.1.3. Glimpse

What the `updatedb/locate` pair is for filenames the `glimpseindex/glimpse` pair is for file contents. `glimpseindex` generates a database that is accessed by the user via `glimpse`. So,

```
glimpse pattern
```

corresponds to a non-database backed command namely a recursive `grep` over a set of directories like

```
find . -print | xargs grep pattern
```

assuming that the database has been generated for `"."`. Again the fast version is so helpful that we have defined two aliases.

```
alias glidx='if test -f .glimpse_index; then
glimpseindex -H . -o -f -B .;
else
glimpseindex -H . -o -B .;
fi'
alias gl='glimpse -H .'
```

The first alias, `glidx`, is one line. It has been broken into several lines only to make its workings clear; namely if an index file already exists it is updated (`-f` option), otherwise it is generated from scratch.

Like our `locate` aliases everything is happening in the current directory which means that `glidx` should be called from SCI.

Glimpse is not part of the standard Linux distribution (at least not SuSE-6.2 and RedHat-6.1, the ones we checked). The University of Arizona currently hosts the [Glimpse home page](http://webglimpse.org/), and Glimpse can also be downloaded from [SC0's software archive](ftp://ftp.sco.com/skunkware/src/fileutil/), which is mirrored by [Sunsite UK](ftp://ftp.sunsite.org.uk/Mirrors/ftp.sco.com/skunkware/src/fileutil/).
8.1.2. Bug Hunting

In preparation of this document (lvd), and in our daily work (cls) we have found it very useful to have more than one Scilab. What? More than one running process? No, more than one binary of scilex. In fact three different versions all come in handy depending on the task.

**scilex binaries**

**Optimized Code**

The common name is “production quality code”, but Scilab is so far away from production quality that we refrain from using that term.

This scilex is built with all compiler optimizations enabled. Furthermore all compiler switches and options are specifically tuned for the machine the code will run on in the future (see also Section 6.3). Maximum performance is the only goal and no attempt is made to retain any debugging information.

**Debugging Code**

This scilex is not optimized, instead it carries complete debug information. Thus, it is ideally suited for interactive debugging sessions, and single-line tracing.

**Profiling Code**

The third variant is a profiling version of scilex that is not optimized for speed either.

Profiling is the first step of any optimization. But during our work with and on Scilab we have found it very helpful to be able to answer the notorious question: “Where is it burin’ the cycles?” Profiling – done right – is much faster than timing individual “suspects”, although analyzing the profiler output requires some skill.

8.2. Local Documents

Following documents come with every source distribution of Scilab. They live in the directory SCI/doc.
8.3. Hyperlinks

Here are a few links that are useful in connection with Scilab.

**Links**

INRIA’s official Scilab pages

- *Scilab Home page* (http://www-rocq.inria.fr/scilab/)
- *Scilab FAQ* (http://www-rocq.inria.fr/scilab/faq/index.html)
• Scilab FTP Site (ftp://ftp.inria.fr/INRIA/Projects/Meta2/Scilab/distributions/)

Pages Of Scilab Enthusiasts

• Enrico Segre’s page (http://www.polito.it/~segre/)
• Stéphane Mottelet’s page (http://wwwdma.utc.fr/~mottelet/scilab/)

Free Numerical Libraries

Netlib (http://www.netlib.org/index.html)
Netlib gathers a lot of free numerical libraries. Due to the nature of the business most of them are written in Fortran-77.

GAMS (http://math.nist.gov/)
The Guide to Available Mathematical Software. This is the lazy man’s entry point to Netlib. If you know your problem then just follow the decision tree until you reach the module that deals with it. Can it get any better than that?

Lapack (http://www.netlib.org/lapack/index.html)
The all time classic of the numerical libraries features linear equation, linear least squares, singular value, and (even generalized) eigenvalue solving in an orthogonal design: 4 precisions (real, double precision, complex, double complex), several matrix storage schemes (rectangular dense, symmetric/hermitian positive definite, banded, tridiagonal, ...).

Optimized BLAS (and sometimes more) Libraries

ATLAS (http://www.netlib.org/atlas/)
ATLAS is an acronym for Automatic Tuning of Linear Algebra Software. A novel approach to optimize the BLAS library for an arbitrary computers with deep memory hierarchies and pipelined functional units. In other words for any modern machine.
Chapter 8. Further Information

ASCI RED (http://www.cs.utk.edu/~ghenry/distrib/)

This is an acronym for Advanced Strategic Computing Initiative. In the course of their research they develop highly optimized BLAS libraries for PPro and later Intel processors.

D1G1TAL’s Extended Math Library (http://www.digital.com/info/hpc/software/dxml.html)

FIXME: Say something about it!

Intel’s Math Kernel Library (http://developer.intel.com/vtune/perflibst/mkl/) (MKL)

This is the one for the poor souls being trapped on the Dark Side. The MKL runs with the software-from-hell (aka Redmond/WA).


FIXME: Say something about it!

5UN’s Fortran High-Performance Library (http://www.sun.com/workshop/fortran/) (PerfLib)

FIXME: Say something about it!

5GI’s Scientific Library (http://www.sgi.com/software/scsl.html) (SCSL)

FIXME: Say something about it!

Other Free Mathematical Software

Pari (http://www.parigp-home.de/)

An extremely fast arbitrary precision calculator with a library that can be linked with user programs.

MuPAD (http://math-www.uni-paderborn.de/MuPAD/index.html)

Symbolic algebra program which is Maple alike. It features one of the most comprehensive integration libraries currently available.

Octave (http://www.che.wisc.edu/octave/)

Matlab3 compatible matrix package, whose core is written in C and C++; relies on classical Fortran libraries like LAPACK. The Octave libraries can be integrate easily in new user programs.
Chapter 8. Further Information

_Tela_ (http://www.geo.fmi.fi/prog/tela.html)

_Tela_ is the short form for tensor language. FIXME: more description about Tela needed.

Free Plotting Software

_GNUPlot_ (http://WWW.cs.dartmouth.edu/gnuplot_info.html)

Although the prefix is a pure coincidence with the _GNU project_ (http://www.gnu.org) it is one of the best 2d-plotting programs that are available with source code.

_PlotMTV_ (ftp://ftp.x.org/contrib/applications/)

An older, but still excellent program for 2d- and 3d-plots. The “philosophy” is different from GnuPlot as plotting instructions and data share the same file.
Chapter 9. Longer Examples

Welcome to our attic! Following the style of the BOT, the examples gathered here are an unsorted collection of hacks that has piled up over the years. A few functions are used or discussed in the earlier section, but were truncated to emphasized the important parts. Here you only find complete versions.

These example programs are free software; you can redistribute it and/or modify it under the terms of the GNU General Public License as published by the Free Software Foundation; either version 2 of the License, or (at your option) any later version.

This program is distributed in the hope that it will be useful, but WITHOUT ANY WARRANTY; without even the implied warranty of MERCHANTABILITY or FITNESS FOR A PARTICULAR PURPOSE. See the GNU General Public License at the end of this document for more details.

You should have received a copy of the GNU General Public License along with this program; if not, write to the Free Software Foundation, Inc., 59 Temple Place, Suite 330, Boston, MA 02111-1307 USA.

9.1. benchmark.sci

This example shows a benchmark function that tries hard to do better than others. In the first step the timer resolution is determined. Next the function under test is executed in a loop and the time taken is estimated. This time in turn is used for the final test. The number of loop iterations is chosen according to the preliminary test. Finally, the median of the timings is returned.

```matlab
function res = calibrate(max_len, n_avg, log_inc)
// determine the resolution of Scilab’s built-in timer
// Return vector with measured timer resolution(s)

[nl, nr] = argn()
if nr <= 2, log_inc = 1.1, end
if nr <= 1, n_avg = 31, end
if nr == 0, max_len = 100000, end

r = []
n = 1
while n <= max_len
    //disp(n)
    tv = []
    iter = 0:n
    for k = 1:n_avg
        timer()
        tv = [tv, timer()]
    end
    % Calculate median using the cumulative distribution
    % and sum the samples.
    r = [r, tv]
    n = n + 1
end
```

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```octave
function t = benchmark()
    verbose = %t
    min_test = 10 % minimum multiple of the timer
    std_test = 200 % as min_test but for real test
    n_avg = 31 % number of samples to calculate median
    log_inc = 2.0 % logarithmic increment in coarse test

    disp("+++ calibrating timer")
    resol = calibrate()
    if size(resol, 2) <= 2 then
        error("calibration failed")
    end
    if resol(1) ~= resol(2) then
        warning("calibration botched; proceeding anyway...")
    end
    t_resol = resol(1);
    if verbose
        disp("timer resolution is " + string(t_resol) + "s")
    end

    disp("+++ calibrating test")
    np = 1
    timer()
    mytest()
    t = timer()
    while t < min_test * t_resol
```

// disp(np, t, min_test * t_resol)
np = log_inc * np
.timer()
for i = 1:np
  mytest()
end
t = timer()
end
if verbose then
  disp("coarse, " + string(np) + " round trips in " + string(t) + "s")
end
if np == 1 then
  warning("slow procedure under test - time may be excessive")
end

// run real test
disp("+++ running test")
tc = t / np
ne = ceil(std_test * t_resol / tc)
if verbose then
disp("fine, test will take about " + string(ceil(tc * ne * n_avg)) + "s")
end

r = []
for k = 1:n_avg
  timer()
  for i = 1:ne
    mytest()
  end
  t = timer()
  r = [r; t]
end
if verbose then
disp("fine, " + string(ne) + " round trips in " + string(t) + "s")
end

// get median and return
r = sort(r)
// disp(r)
tpl = r($/2 + 1) / ne

function mytest()
exact = -2.5432596188;
z = abs( exact - intg(0, 2 * %pi, f) )

function y = f(x)
// y = x * sin(30 * x) / sqrt( 1 - ((x / (2 * %pi))^2) )
y = x

9.2. listdiff.sci

The function listdiff returns the differences of two vectors in the style of the \texttt{diff(1)} command. It is a funny example of doing something completely non-numerical with Scilab.

function diff = listdiff(lst1, lst2, equ)
// implement a \texttt{diff(1)} like function for vectors.
// The caller can supply a bool \texttt{equ(x, y)} function
// that will be used in all comparisons, otherwise
// operator \texttt{"=="} is used.
//
// RETURN VALUE
// \texttt{2-column vector describing the differences.}
// Column 1 contains the element and column 2
// the element's index. A \texttt{"+"} in front of the
// index means: "Extra element in \texttt{lst2}", a \texttt{"-"}
// means missing element in \texttt{lst1}.

[nl, nr] = argn(0);
select nr
  case 0 then
    error("Too few arguments. Got 0, require 2 or 3."),
  case 1 then
    error("Too few arguments. Got 1, require 2 or 3."),
  case 2 then
    deff('b = equ(s1, s2)', 'b = s1 == s2');
  case 3 then
    // caller supplied \texttt{equ()}
    if type(equ) == 13 then
      error("Function expected, got a " + typeof(equ) + "."),
    end
  else
    error("Too many arguments. Got " + string(nr) + " require 2 or 3."),
  end
if type(lst1) ~= 1 & type(lst2) ~= 1 then
  // none of the lists is empty
  if type(lst1) ~= type(lst2) then
    error("Both lists must be of the same type, or be empty.");
  end
end

fuzz = 10;

diff = [];
n1 = size(lst1, 1);
n2 = size(lst2, 1);

// special cases
if n1 == 0 & n2 == 0, return, end

if n1 == 0 then
  p = 1 : n2;
  diff = [lst2, "+" + string(p')];
  return;
end

if n2 == 0 then
  p = 1 : n1;
  diff = [lst1, string(-p')];
  return;
end

// general case (neither list is empty)
i = 1;
j = 1;
while i <= n1 & j <= n2
  while i <= n1 & j <= n2
    if ~equ(lst1(i), lst2(j)), break, end
    i = i + 1;
    j = j + 1;
  end
  if i >= n1 | j >= n2, break, end
  icurs = i;
  while icurs <= min(n1, i+fuzz)
if equ(lst1(icurs), lst2(j)), break, end
icurs = icurs + 1;
end
if icurs <= n1 then
    if equ(lst1(icurs), lst2(j)) then
        // record element(s) missing from lst1
        for p = i : icurs-1
            this_diff = [lst1(p), string(-p)];
            diff = [diff; this_diff];
        end
        // re-sync
        i = icurs;
    end
end

jcurs = j;
while jcurs <= min(n2, j+fuzz)
    if equ(lst1(i), lst2(jcurs)), break, end
    jcurs = jcurs + 1;
end
if jcurs <= n2 then
    if equ(lst1(i), lst2(jcurs)) then
        // record extra element(s) in lst2
        for p = j : jcurs-1
            this_diff = [lst2(p), "+" + string(p)];
            diff = [diff; this_diff];
        end
        // re-sync
        j = jcurs;
    end
end
end

9.3. whatis.sci

lvd: Enrico, please say something about your function here.

function rv = whatis(name_arr)
// NAME
// whatis - listing of variables in extended format
//
// CALLING SEQUENCE
// whatis()
// whatis(name_arr)
//
// PARAMETER
// name_arr : array of variables names
//
// DESCRIPTION
// whatis returns a column-vector with the names,
// types, and sizes of all local variables
// (first form), or only of the variables whose
// names (as strings!) are given in the matrix
// name_arr (second form).
//
// EXAMPLES
// whatis()
// whatis('my_mat')
// whatis(['foo' 'bar' 'baz'; 'foobar 'morefoo' 'foobaz'])
//
// SEE ALSO
// who, whos
//
// AUTHORS
// Enrico Segre, Lydia van Dijk
//
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// of the GNU General Public License, version 2.
//!

// LAST REVISION
// lvd, Fri Dec 3 01:01:45 UTC 1999

// TO DO/TO FIX
//
// - Accepting a regexp as an argument would be nice. This in turn
// leads to complete boolean expressions doing the variable selection
// resembling what the UNI* find utility does. Example:
// All vars ending in a 'v' that are complex and larger than
// 1000 words.
// - The behavior with undefined variables is unsatisfactory.

[nl, nr] = argn(0);
clear nl;
if nr == 0 then
// no arguments -> take all variables like whos() does
clear nr;
names_all = sort(who("get"));
end
clear nr;

if type(names_all) ~= 10 then
    error("Expecting a string or an array of strings, got a 
    
    + typeof(names_all) + ");
    return;
end

[nam, mem] = who("get"); // get memory usage of all local vars

// define isreal() for hypermatrices
def('b = %hm_isreal(hm)', ..
    'if size(hm, "***") == 0 then b = %t; ..
    else ..
    b = isreal(hm(1)); ..
    end');

def('b = %hm_isbool(hm)', ..
    'if size(hm, "***") == 0 then ..
    b = %t; ..
    else ..
    b = type(hm(1)) == 4; ..
    end');

def('b = %hm_isstring(hm)', ..
    'if size(hm, "***") == 0 then ..
    b = %t; ..
    else ..
    b = type(hm(1)) == 10; ..
    end');

def('b = %hm_isint(hm)', ..
    'if size(hm, "***") == 0 then ..
    b = %t; ..
    else ..
    b = type(hm(1)) == 8; ..
    end');

r = [];
for name = matrix(names_all, 1, size(names_all, "**")) do
    if isdef(name) then
        ..
end

114
clear var;
var = evstr(name);  // convert var’s name back into var

// type classification
//
ty = type(var);  // type number
select ty  // type 16 and 17 are not recognized
    case 16 then  // by the function typeof()
        tgenp = %f;  // we know the tlist’s type for these
        lab = var(1);  // vector of labels
        select lab(1)  // 1st label defines the type
            case "ar" then
                tnam = "ARMAX process";
            case "des" then
                tnam = "descriptor system";
            case "linpro" then
                tnam = "linear programming data";
            case "lss" then
                tnam = "linear system";
            case "r" then
                tnam = "rational";
            case "scs_tree" then
                tnam = "SCICOS navigator data";
            case "xxx" then
                tnam = "SCICOS menu data";
            else
                tnam = "generic tlist " + lab(1);
                tgenp = %t;
            end // select lab(1)
    case 17 then
        tnam = "hypermatrix";
    else
        tnam = typeof(var);  // type name, a string
    end // select ty
if ty==1 | ty==2 | ty==5 | ty==17 then
    // boolean, string, integral, real, or complex,
    // possibly sparse matrix or hypermatrix (yuck!)
    if hm_isbool(var) then
        tnam = "boolean " + tnam;
    elseif hm_isstring(var) then
        tnam = "string " + tnam;
    elseif hm_isint(var) then
        tnam = "int " + tnam;
    else
        if isreal(var) then

9.4. Auto-Determination of Precedence and Associativity

assoc.sci, prec.sci, and parser.sci are the scripts that determine the precedence and the associativity of the arithmetic Scilab operators. The results are used in Section 4.1.
### 9.4.1. assoc.sci

function a = assoc(oper, typ)
    // Return the associativity a of
    // operator oper which accepts type typ.
    // oper can be a matrix of operators.
    //
    // typ can be ‘n’ for numeric, or ‘b’ for boolean.
    // If typ is omitted, numeric is assumed.
    [nl, nr] = argn()
    if nr == 1 then
        typ = ’n’
    end
    select typ
        case ’n’ then
            args = string([1.1 1.2 1.5])
            deff(’b = equal(x, y)’, ’b = abs(x – y) < 1.2*%eps’) 
        case ’b’ then
            args = string([’%f’ ’%t’ ’%f’])
            deff(’b = equal(x, y)’, ’b = x == y’)
        else
            error(’unknown type ’ + typ)
    end
    a = []
    for op = oper
        expr = ’[’ + args(1) + op + op + args(3) + ’,’ + op + op + args(3) + ’,’ + op + op + args(3) + ’]’
        r = evstr(expr)
        //disp(r)
        if equal(r(2), r(3)) then
            a = [a ’non’]
        elseif equal(r(1), r(2)) then
            a = [a ’left’]
        elseif equal(r(1), r(3)) then
            a = [a ’right’]

        else
            // Additional code could be added here.
        end
    end

function p = prec(op1, op2)
    // determine the relative precedence of operator op1 vs op2
    // If operator op1 has a higher precedence than op2 then p = -1.
    // In the opposite case p = 1. If both have the same precedence
    // level p = 0

    args = string([1.1 1.2 1.5])
    deff('b = equal(x, y)', 'b = abs(x - y) < 1.2*%eps')

    expr = ['[' + args(1) + op1 + args(2) + op2 + args(3) + ']',
            '(' + args(1) + op1 + args(2) + ')
            op2 + args(3) + ']

    //disp(expr)
    r = evstr(expr)
    //disp(r)

    if equal(r(2), r(3)) then
        p = 0
    elseif equal(r(1), r(2)) then
        p = -1
    elseif equal(r(1), r(3)) then
        p = 1
    else
        error('could not determine precedence level')
    end

function p = prec1(uop, op)
    // determine what relative precedence the unary operator uop has
    // with respect to operator op. The return values are like those
    // of prec()

    args = string([1.1 1.2])
    deff('b = equal(x, y)', 'b = abs(x - y) < 1.2*%eps')

    expr = ['[' + uop + args(1) + op + args(2) + ']

    //disp(expr)
    r = evstr(expr)
    //disp(r)
    if equal(r(2), r(3)) then
        p = 0
    elseif equal(r(1), r(2)) then
        p = -1
    elseif equal(r(1), r(3)) then
        p = 1
    else
        error('could not determine precedence level')
    end
+ '(' + uop + args(1) + '+')' + op + args(2) + ']''

//disp(expr)
r = evstr(expr)
//disp(r)
if equal(r(1), r(2)) then
  p = -1
else
  p = 1
end

function p = lprec(op1, op2)
// determine relative precedence of the
// logical operators op1 and op2
v = ['%f' '%t']
for i = 1:2
  for j = 1:2
    for k = 1:2
      args = string([v(i) v(j) v(k)])
      expr = '[' + args(1) + op1 + args(2) + op2 + args(3) + ',' ..
      + '(' + args(1) + op1 + args(2) + ')' + op2 + args(3) + ',' ..
      + args(1) + op1 + '(' + args(2) + op2 + args(3) + ')]'
      //disp(expr)
r = evstr(expr)
      //disp(r)
      if r(2) == r(3) then
        p = 0
      elseif r(1) == r(2) then
        p = -1
        return
      elseif r(1) == r(3) then
        p = 1
        return
      else
        error('could not determine precedence level')
      end
    end
  end
end
end
9.4.3. parser.sci

// determine properties of Scilab’s parser:
// associativity and precedence level of operators

getf('assoc.sci');
getf('prec.sci');

numop1 = ['+' '−'];
numop2 = ['+' '−' '*' '/' '\' '>' '<' '.' '.*' './' '.\' '.^'];
logop1 = ['~'];
logop2 = ['&' '|'];

// inquire associativity
an = assoc(numop2, 'n');
ab = assoc(logop2, 'b');

// figure out the relative precedence of binary numeric operators
pm2 = [];
for i = numop2
    row = [
        for j = numop2
            row = [row prec(i, j)];
        end
    pm2 = [pm2; row];
end
[lev, idx] = sort( sum(pm2, 'r') );
lev = lev - min(lev) + 1; // minimum := 1

nop2 = numop2;
for op = numop1 // mark binary oparators that have a unary twin
    patch = find(op == nop2);
    nop2(patch) = op + '/2';
end

repl2 = [string(lev); nop2(idx); an(idx)]';

repl1 = [];
for i = numop1
    row = [
        for j = numop2
            row = [row, prec1(i, j)];
        end
    hop = numop2(find(row > 0.5)); // operators with higher precedence
Chapter 9. Longer Examples

```
minhop = 0;
for op = hop
    minhop = max( minhop, find(relp2(:, 2) == op) );
end
// now minhop is the index of the lowest precedence binary operator
// that has a higher precedence than the unary operator i, or 0 if
// there is none
if minhop == 0
    uop = evstr(relp2(1, 1)) + 1;
else
    uop = evstr(relp2(minhop, 1)) - 1;
end
relp1 = [relp1; [string(uop), i+'/1', 'right']];
end
//relp1
// Merge unary operators into matrix of binary operators
relp = [relp1; relp2];
[dummy, idx] = sort(evstr(relp(:, 1)));
relp(idx, :)
```

9.5. cat.sci

cat.sci defines the function cat which prints the source of a macro (function) if it is available. The argument-, type-, and size-checking part is used in Example 4-1.

```
function [res] = cat(macname)
// Print definition of function 'macname'
// if it has been loaded via a library.

// AUTHOR
// Lydia van Dijk
//
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// of the GNU General Public License, version 2.

[nl, nr] = argn(0);
if nr ~= 1 then
    error("Call with: cat(macro_name)");
end
```
if type(macname) ~= 10 then
    error("Expecting a string, got a " + typeof(macname));
end
if size(macname, "*") ~= 1 then
    sz = size(macname);
    error("Expecting a scalar, got a " ..
        + sz(1) + "x" + sz(2) + " matrix")
end

[res, err] = evstr(macname);
if err ~= 0 then
    select err
    case 4 then
        disp(macname + " is undefined.");
        return;
    case 25 then
        disp(macname + " is a builtin function");
        return;
    else
        error("unexpected error", err);
    end
    // select err
end

name = whereis(macname);
//disp("name = <" + name + ">");
if name == [] then
    disp(macname + " is defined, but its definition is unaccessible");
    clear ans;
    return;
end

cont = string(evstr(name)); // path (1) and contents (2..$) of library
fpath = cont(1);
if part(fpath, 1:4) == "SCI/" then
    fpath = SCI + "/" + part(fpath, 5:length(fpath));
end
fname = fpath + macname + ".sci";

[fh, err] = file("open", fname, "old");
if err ~= 0 then
    error("Could not open file " + fname, err);
end
text = read(fh, -1, 1, ")(a)";
file("close", fh);
write(%io(2), text);
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