User Guide for falcON

version of July 28, 2003

Summary. falcon is the "Force Algorithm with Complexity $\mathcal{O}(N)$)" which is described in Dehnen (2002). With this packages, you can use falcon in subroutine form as Poisson solver for particle based simulations. The package also has a full N-body code, based on falcon, called <code>gyrfalcon</code> ("GalaxY simulatoR using falcon"), which employs the N-body tool box NEMO. This code features individual adaptive time steps employing a block-step scheme, but can also be used in single-time-step mode (in which case momentum is exactly conserved).

1 Guarantee

This package comes with absolutely no guarantee whatsoever! The unpacking, installation, and usage of the code is entirely at the risk of the user alone.

2 Credit

Any scientific publication or presentation which has benefited from using any part of this package should quote the paper

Dehnen, W., 2002, JCP, 179, 27. (please find a pdf file of this paper in the subdirectory falcON/doc.)

3 Unpacking

After downloading the file falcon.tgz, unpack it typing

tar zxf falcON.tgz,

which should create the directory falcon with sub-directories src, inc, and doc, as well as several other files.

4 Installation

You need to make the library libfalcon. a and possibly the executables you want to use, see §§ below. The code is written entirely in C++ and it is strongly recommended to use a compiler that understands standard C++, e.g. GNU's gcc version 3.2 (or higher). If you want to use any other compiler than gcc, edit the file make.defs and change the entry for C++COMP. However, I cannot recommend using the Intel compiler (it produces slower code, I tried versions 6.0 and 7.0). The Makefile is intended for use with GNU make.

In order to allow the code to understand NEMO data format and parameter I/O, you must invoke NEMO **before** compilation.

To generate the library as well as a test program TestGrav and the N-body code gyrfalcON, type make

The making takes a little while but should not produce any warning or error messages. Otherwise something might be wrong. The executables live in a subdirectory

falcON/\$(MACHTYPE)_\$(OSTYPE),

where MACHTYPE and OSTYPE are environment variables unique to the machine type and operating system. In this way, you may have versions of the executables and the library (which is in subdirectory falcon/\$(MACHTYPE)_\$(OSTYPE)/lib) for several hosts on the same file system.

5 Testing falcON

Please run TestGrav in order to get some rough check on the validity of your library. Issuing the command

```
TestGrav 2 1 1000000 901 0.01 1
```

shall generate a Hernquist sphere with $N=10^6$ particles, build the tree (twice: once from scratch and once again) and compute the forces using a softening length of $\epsilon=0.01$ scale radii with the P_1 kernel (see §6). The output of this command may look like

```
time needed for set up of X_i:
time needed for falcON::grow():
                                                              2.64
time needed for falcON::grow():
                                                              1.6
time needed for falcON::approximate(): 9.03
state:
                                  tree built
root center:
root radius:
                                  0 0 0
                                 1024
bodies loaded:
                                1000000
total mass:
N_crit:
cells used:
                               353419
21
maximum depth:
current theta:
                                 0.6
softening length: 0.01 softening kernel:
softening kernel: P1
Taylor coeffs used: 84569
interaction statitics:
type approx direct (pairs) total

# body-body: - 0 (0) 0 = 0%

# cell-body: 2115758 477698 (1924073) 2593456 = 18.342%

# cell-cell: 11237586 254997 (3849070) 11492583 = 81.279%

# cell-self: - 53678 (13128506) 53678 = 0.38%

# total: 13353344 786373 (18901649) 14139717 = 100.000%
ASE(F)/<F^2> = 0.001598375617

max (dF)^2 = 0.8182717562

Sum m_i acc_i = -2.707049029e-09 1.63810987e-09 2.313722963e-10
```

Note that the second tree-build is somewhat faster then the original one. Note also the total-momentum change (last line) vanishes within floating point accuracy – that's a generic feature of falcon.

6 Choice of the Softening Kernel and Length

The code allows for various forms of the softening kernel, i.e. the function by which Newton's 1/r is replaced in order to avoid diverging near-neighbour forces. The following kernel functions are available ($x := r/\epsilon$)

name	density (is proportional to)	a_0	a_2	f
P_0	$(1+x^2)^{-5/2}$	∞	∞	1
P_1	$(1+x^2)^{-7/2}$	π	∞	1.43892
P_2	$7(1+x^2)^{-9/2} - 2(1+x^2)^{-7/2}$	0	∞	2.07244
P_3	$9(1+x^2)^{-11/2} - 4(1+x^2)^{-9/2}$	0	$-\pi/40$	2.56197

Note, that P_0 is the standard Plummer softening, however, **recommended** is the use of P_1 or P_2 . There are several important issues one needs to know about these various kernels.

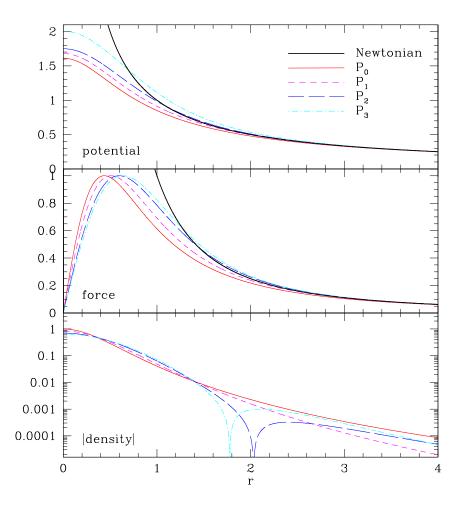


Figure 1: Potential, force, and density for the softening kernels of the table, including the standard Plummer softening (P_0) . The softening lengths ϵ are scaled such that the maximum force equals unity. The kernels $P_{>0}$ approach Newtonian forces more quickly at larger r than does P_0 . The kernels P_2 and P_3 have slightly super-Newtonian forces (and negative densities) in their outer parts, which compensate for the sub-Newtonian forces at small r.

First, the softening length ϵ is just a parameter and using the same numerical value for it but different kernels corresponds in effect to different amounts of softening. Actually, this softening is strongest for the Plummer sphere: at fixed ϵ , the maximal force is smallest. In order to obtain comparable amounts of softening, larger ϵ are needed with all the other kernels. An idea of the factor by which ϵ has to be enlarged can be obtained by setting ϵ such that the maximum possible force between any two bodies are equal for various kernels. The last column in the previous table gives these factors. Note, that using a larger ϵ with other than the P_0 kernel does **not** mean that your resolution goes down, it in fact increases, see Dehnen (2001), but the Poisson noise is more suppressed with larger ϵ . It is recommended not to use Plummer softening, unless (i) you want $\epsilon \equiv 0$, (ii) in 2D simulations, as here ϵ is the average scale-height of the disk, and, perhaps, of (iii) in simulations made to compare with others that use Plummer softening (for historical reasons).

Second, as shown in Dehnen (2001), Plummer softening results in a strong force bias, due to its slow convergence to the Newtonian force at $r \gg \epsilon$. This is quantified by the measure a_0 , which for P_0 is infinite. In Dehnen (2001), I considered therefore other kernels (not mentioned above), which have finite support, ie. the density is exactly zero for $r \ge \epsilon$. This discontinuity makes them less useful for the tree code (which is based on a Taylor expansion of the kernel). In order to overcome this difficulty, the kernels P_1 to P_3 , which are continuous in all derivatives, have been designed as extensions to the Plummer softening, but with finite a_0 (P_1), zero a_0 but infinite a_2 (P_2), or even zero a_0 and finite a_2 (P_3).

7 Choice of the Tolerance Parameter

The code falcON approximates an interaction between two nodes, if their critical spheres don't overlap. The critical spheres are centered on the nodes' centers of mass and have radii

$$r_{\rm crit} = r_{\rm max}/\theta$$
 (1)

where $r_{\rm max}$ is the radius of a sphere that is guaranteed to contain all bodies of the node (bodies have $r_{\rm max}=0$), while θ is the tolerance parameter. The default is to use a mass-dependent $\theta=\theta(M)$ with $\theta_0\equiv\theta(M_{\rm tot})$ being the parameter, see Dehnen (2002). For near-spherical systems or groups of such systems, θ_0 of 0.6 gives relative forces error of the order of 0.001, which is generally believed to be acceptable. However, the force error might often be dominated by discreteness noise, in which case a larger value does no harm. For disk systems, however, a smaller tolerance parameter, e.g. $\theta_0=0.5$, might be a better choice.

The recommendation is to either stick to θ_0 no larger than about 0.6, or perform some experiments with varying θ_0 (values larger than 0.8, however, make no sense, as there is hardly any speed-up).

8 Use of falcON as Poisson Solver

8.1 With C++

In order to make use of the code, you need to insert the C macro

#include <falcON.h>

somewhere at the beginning of your C++ source code. Make sure that the compiler finds the file falcON.h by including -I falcON/inc among your compiler options. The usage of the code in your application is explained in gory detail in the file falcON.h (don't forget that class falcON lives in namespace nbdy). In order to make an executable, add the linker options -LfalcON/\$(MACHTYPE)_\$(OSTYPE)/lib -lfalcON -lm (expand the macros in your makefile) so that the library is loaded.

For examples of code using falcON.h, see the files TestGrav.cc and TestPair.cc in subdirectory src/mains/, which may be compiled by typing make TestGrav and make TestPair and produce a short summary of their usage when run without arguments.

8.2 With C

In order to make use of the code, you need to insert the C macro

#include <falcON_C.h>

somewhere at the beginning of your C source code. Make sure that the compiler finds the file fal-cON_C.h by including -I falcON/inc among your compiler options. The usage of the code in your application is explained in gory detail in the file falcON_C.h. In order to make an executable, add the linker options -LfalcON/\$(MACHTYPE)_\$(OSTYPE)/lib -lfalcON -lstdc++ -lm (expand the macros in your makefile) so that the library is loaded.

For examples of code using falcON_C.h, see the files TestGravC.cc and TestPairC.cc in subdirectory src/mains/, which may be compiled by typing make TestGravC and make TestPairC and produce a short summary of their usage when run without arguments.

8.3 With FORTRAN

In order to make use of the code, you need to insert

INCLUDE 'falcON.f'

somewhere at the beginning of your FORTRAN program. Make sure that the compiler finds the file fal-con.f by including -I falcon/inc among your compiler options. The usage of the code in your application is explained in gory detail in the file falcon.f. In order to make an executable, add the linker options -Lfalcon/\$(MACHTYPE)_\$(OSTYPE)/lib -lfalcon -lstdc++ -lm (expand the macros in your makefile) so that the library is loaded.

For examples of code using falcON.f, see the files TestGravF.F and TestPairF.F in subdirectory src/mains/, which may be compiled by typing make TestGravF and make TestPairF. Just run these programs, they are self-explanatory and provide some statistics output. You may also use the input files given and run them as TestGravF < treeF.in and TestPairF < pairF.in.

9 The N-Body Code gyrfalcON

The package also contains a full N-body code, called "gyrfalcON" (GalaxY simulatoR using falcON)¹. If you want to use this code, you need first to install and invoke the N-body tool box NEMO, version 3.0.13 or higher², see http://www.astro.umd.edu/nemo. It is recommended to configure NEMO with configure --enable-single --enable-lfs.

```
Then type

make gyrfalcON

which should produce the executable gyrfalcON in the subdirectory (add it to your $PATH)

falcON/$(MACHTYPE)_$(OSTYPE).

gyrfalcON comes with the usual NEMO help utility: calling

gyrfalcON help=h

produces the following overview over the options.
```

```
in
                 : input file
                                                                       [???]
                 : file for primary output; required, unless resume=t []
out
                 : final integration time [default: never]
                                                                       [1
tstop
                : time between primary outputs; 0 -> every step
                                                                       [1]
step
logfile
                : file for log output
                                                                       [-]
stopfile
               : stop simulation as soon as file exists
                                                                       []
               : # blocksteps between log outputs
                                                                       [1]
                : file for secondary output stream
                                                                       []
               : time between secondary outputs; 0 -> every step
                                                                       [0]
theta
               : tolerance parameter at M=M_tot
                                                                       [0.60]
harow
               : grow fresh tree every 2^hgrow smallest steps
                                                                       [0]
Ncrit
               : max # bodies in un-split cells
                                                                       [6]
                : softening length OR maximum softening length
eps
                                                                       [0.05]
                : softening kernel of family P_n (P_0=Plummer)
                                                                       [1]
kernel
                : tau_min = (1/2)^hmin
                                                                       [6]
hmin
                : # time-step levels
Nlev
                                                                       [1]
fac
                : tau = fac / acc
                                             \ If more than one of
                                                                       []
                : tau = fph / pot
                                                these is non-zero,
fph
                                            [ ]
                : tau = fpa * sqrt(pot)/acc | we use the minimum
                                                                       []
fpa
                : tau = fea * sqrt(eps/acc) /
fea
                                                tau.
                                                                       []
resume
                : resume old simulation? that implies:
                   - read last snapshot from input file
                   - append primary output to input (unless out given) [f]
                 : list of output specifications. Recognizing:
give
                   m: mass
                                                         (default)
                   x: position
                                                         (default)
                    v: velocity
                                                         (default)
                    a: acceleration
                    p: potential
                    P: external Pot (added to pot before output)
                    l: time-step level (if they exist)
                                                                       [mxv]
                    f: body flag
                : list of specifications for secondary output
give2
                                                                       [mxv]
                : Newton's constant of gravity
                                                                       [1]
Grav
potname
                : name of external potential
                                                                       [ ]
potpars
                : parameters of external potential
                                                                       []
potfile
                : file required by external potential
                                                                       []
                : primary output for t=tstart?
startout
                                                                       [t]
                 : primary output for t=tstop?
lastout
                                                                       [t]
VERSION
                 : 20-mar-2003 Walter Dehnen
                   compiled Mar 20 2003, 13:02:47
                                                                       [1.5.7]
```

¹Called "YancNemo" in former versions of this package.

²Older versions of this package contained a non-NEMO code, called "YANC". This code was never properly tested and has hence been deprecated.

The last column indicates the default value, with '[???]' indicating that the value for the keyword must be given. A filename '-' means that output is written to stdout and can be piped into another command or that input is expected from stdin, which can be a pipe. A filename '.' means that no output is made at all.

9.1 Log Output

Log output is written every logstep blocksteps to logfile. The default, logfile=-, i.e. log output is written to stdout, prevents the usage of the pipe to transfer N-body data. Thus, in order to use piping you must specify some logfile.

9.2 Data I/O

The initial conditions, including the simulation time, are read in from the file given with the parameter in and **must** be in NEMO snapshot format. Unless the keyword resume=t (see below) the first snapshot in this file is used.

The code allows for two distinct data outputs, one into out and another optional one into out2. Output is made every step and step2 time units, respectively. The option lastout=t, which is default, ensures that the last snapshot of the simulation is written to out, in any case, even if it is not an integer number of steps from the initial simulation time.

The type of N-body data written is controlled by the options give and give2, which are character strings containing the letters m, x, v, a, p, P, 1, f, indicating that the masses, positions, velocities, accelerations, N-body potentials, external potentials (if present, see below)³, time-step levels, and body flags shall be given. The default is give=mxv, i.e. masses, positions, and velocities.

Traditionally on linux systems, there is a limit of 2Gb on the size of files. This will cause trouble with NEMO snapshot files, since the snapshots of all output times are written to one file. To overcome this, you must (i) configure NEMO appropriately (use configure --enable-lfs when installing and (ii) ensure that your file systems supports large files - consult your system administrator.

9.3 Gravity Parameters

The parameters eps and kernel control the softening, see \$6. The parameter Ncrit sets the maximum number of bodies in a leaf cell of the tree. The default value of Ncrit is set to yield highest performance. The parameter 'theta' gives the tolerance parameter θ_0 , the opening angle for the root cell. It defaults to 0.6, which results typically in relative force errors of a few 10^{-3} . For simulations of disk-like systems, however, it might be better to use a smaller value, like $\theta_0 = 0.5$, see §7.

9.4 Time Step and Simulation Time

The (shortest) time step is controlled by the parameter hmin and is equal to

$$\tau_{\min} = 2^{-\text{hmin}}.$$

The simulation stops either if the simulation time has been advanced to the value of the parameter tstop (if given), or if a file whose name has been given with the parameter stopfile is found to exist. The latter may be used to stop the simulation based on the results of some analysis program (which gets the data piped in via out2 at every time step and creates an appropriate stopfile if some condition is satisfied). If tstop equals the initial simulation time, the initial forces are computed and, if so desired, output is made for this time only.

9.5 Resuming an Old Simulation

An old simulation may be resumed by setting the keyword resume=t. In this case, the last snapshot from the input file is taken as initial conditions and output is appended to the input file.

³ Unfortunately, the present version of NEMO does not support more than one potential output per body. This means that on output the sum of the desired potentials is written, while input of an external potential is impossible.

9.6 Adding an External Potential

On top of the N-body forces, an external potential may be added using the usual NEMO keywords potname, potpars, and potfile. When potname is given, a corresponding shared object file is loaded dynamically and the potential is initialized with the parameters given with potpars or the data file given with potfile, whatever applies.

9.7 Adaptive Time Stepping

You can use gyrfalcON without adaptive time steps, by setting 'Nlev=1' (default). In this case, also much of the (small) overhead that is necessary for adaptive time steps is avoided, a standard leap-frog time integrator is used, and the total momentum (center-of-mass motion) is conserved exactly (within floating-point accuracy).

When Nlev> 1, a block-step scheme with Nlev time-step levels is used, i.e. the longest step contains $2^{\text{Nlev}-1}$ shortest steps. The bodies' individual time-step levels are adapted in an (almost) time symmetric fashion to be on average

$$\tau = \min \left\{ \frac{\text{fac}}{|a|}, \frac{\text{fph}}{|\Phi|}, \text{fpa} \frac{\sqrt{|\Phi|}}{|a|}, \text{fea} \sqrt{\frac{\epsilon}{|a|}} \right\}, \tag{2}$$

where Φ , a, and ϵ are the gravitational potential and acceleration and the softening length. The parameters fac, fph, fpa, and fea determine the stepping. If either of them is zero, it is ignored.

In order to make a sensible choice for the parameters hmin, Nlev, fac, fph, fpa, and fea, use the following method. (i) Decide on the smallest time step: think what time step you would use in a single-time-step leap-frog scheme and then set τ_{\min} to about half of that. (ii) Decide on the largest time step, whereby ensuring that orbits in regions of very low density are accurately integrated when using the above criterion (2). (iii) Do some tests with varying fac, fph, fpa, and fea (set tstop=0), in order to check that the distribution of bodies over the time steps is reasonable, in particular there should be a few percent in the smallest time step.

When using adaptive time stepping, it may be worth your while to use a larger value for Ncrit than default. This reduces the time of tree-building and increases that for the force computation. However with adaptive time stepping the relative contribution of tree-building to the total CPU time budget is much larger than without.

Note that using this scheme is sensible only if you really have a very inhomogeneous stellar system, because otherwise, the simple single-time-step leap-frog is only slightly less efficient but somewhat more accurate. In particular, with the block-step scheme, the total momentum is not conserved, but with the single-time-step leap-frog it is.

9.8 An Example

In order to integrate a Plummer sphere with $N=10^5$ particles, you may issue the command

```
mkplummer - 100000 seed=1 scale=1 | gyrfalcON - plum.snp tstop=10 eps=0.1
```

which first creates initial conditions from a Plummer model, which are then piped into gyrfalcON. gyrfalcON creates an output file 'plum.snp' containing output every full time unit until time t=10. The log output looks like

```
#
  "gyrfalcON - plum.snp tstop=10 eps=0.1 VERSION=1.5.3"
         Thu Feb 27 09:19:58 2003
# run at
     by
          "dehnen"
     on "milkyway"
                                                       build
                                                                                 accum
                          0.50316 0.0010761 6e-09
                          0.50317 0.0010762 1.9e-09
0.015625
          -0.147379
                                                      0.14
                                                             0.98
                                                                        1.12
                                                                                  2.35
                          0.50316 0.0010762 5.6e-09
                                                             0.98
0.03125
           -0.1473791
                                                                                  3.52
```

0.046875	-0.147379	0.50317 0.0010762	5.1e-09	0.14	0.99	1.14	4.68
•							
•							
•							
9.9531	-0.1473785	0.49925 0.0010798	4.1e-09	0.14	0.98	1.13	745.378
9.9688	-0.1473783	0.49926 0.0010799	5.8e-09	0.14	1	1.14	746.548
9.9844	-0.1473778	0.49927 0.0010799	2.3e-09	0.14	0.98	1.14	747.718
10	-0.1473785	0.49927 0.0010799	5.8e-09	0.15	0.99	1.14	748.888

The column | v_cm| gives the center-of-mass motion, which stays constant (within floating point precision) due to the momentum-conserving nature of falcon. The last four columns contain the CPU time in seconds spent on the tree building, force computation, and full time step, as well as the accumulated time.

10 addgravity and getgravity

The public version of the falcON package contains two further NEMO executables. addgravity simply adds acceleration and potential to every body in the snapshots of a NEMO snapshot file. getgravity computes the gravity generated by the particles (sources) in the snapshots of a NEMO snapshot file at the positions of the particles (sinks) in another snapshot. This is useful for, for instance, computing the rotation curves of N-body galaxies.

11 Bugs and Features

11.1 Test-Particles

falcon does not support the notion of a test particle, i.e. a body with zero mass. Such bodies will never get any acceleration (that is because the code first computes the force, which is symmetric and hence better suited for mutual computations, and then divides by the mass to obtain the acceleration). To overcome this, you may use tiny masses, but note that the forces created by such light bodies will be computed, even if they are tiny.

Actually, this is exactly what we do in getgravity.

11.2 Bodies at Identical Positions

The code cannot cope with more than Ncrit bodies at an identical position (within floating point accuracy). Such a situation would result in an infinitely deep tree; the code aborts with an error message.

11.3 Unknown Bugs

A bug that lead falcon or gyrfalcon to occasionally crash with 'Segmentation fault' I have recently tracked down and debugged (as of 3rd April 2003). However, there seems still to be a similar bug around, which is not reproducible and hence hard to track down and weed out. Measures have been taken to solve this problem eventually. If you ever encounter a problem that you think might be a bug and which is not mentioned in this documentation, please report it to me (wdehnen@aip.de). Thanks.

12 References

Dehnen, W., 2001, MNRAS, 324, 273 Dehnen, W., 2002, JCP, 179, 27