SIMBAD Users Manual. Version v.1.36

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## 1 Introduction

SIMBAD is a Particle-in-Cell (PIC) tracking code [1] designed to track particles in 3-D in an accelerator. Many of its algorithms are designed and optimized with an hadron synchrotron or storage ring in mind. In SIMBAD a "herd" of representative macro particles (in short: macros) is injected and propagated through the structure of the accelerator represented by a sequence of transfer maps. In the present version, transfer maps are produced by the code MAD, that stands for Methodical Accelerator Design. The effect of space charge in the presence of accelerator walls and the effects of impedance on the beam is calculated.

SIMBAD is the BNL version of ORBIT [2]. It has been designed and optimized to run on a parallel computer, however for limited tasks, it can also be run on a serial machine. The language used is C++, the default compiler is the gnu compiler. The code has also been installed on machines using different compilers, like the IBM AIX. Default graphics is provided by GNUPLOT [3].

The code is simply run by reading parameters from a configuration file, as follows:

| $>$ SIMBAD filename.conf | serial |
| :--- | :--- |
| $>$ mpirun -np N SIMBAD filename.conf | parallel linux |
| $>$ lload filename.cmd | IBM AIX |

The present SIMBAD User Manual is essentially a description of the configuration file and of the meaning of the various items in it. Before running SIMBAD one must edit the .conf file. A specific action is accomplished by uncomment the relative module.

## 2 Basic concepts

### 2.1 Nodes

Events happen sequentially during tracking. The propagation of the Herd through a machine element is one of such events, the calculation of Space Charge force is an event, dumping of the phase space coordinates of each macro is an event, etc. Events happen at locations in the lattice called Nodes. Each node has a name, an integer order number and a longitudinal coordinate. To start with, machine elements read from the MAD [4] output are given integer ordering number spaced by some input interval, say 1000, other nodes are later inserted between machine elements, just by assigning to them a number between these two numbers.

In SIMBAD the longitudinal distance variable is in a sense the independent variable. Sometimes the variable time is important for calculations, e.g. for space charge forces. In this case, space and time are transformed from one to the other

SIMBAD works using the Split Operator technique: some propagation in the "Bare Lattice", a calculation of some effect, a new propagation, an so on.

## 3 Preliminary Setting

The .conf file contains some preliminary settings. Some, marked with $\left(^{*}\right)$, here and in the following, should always been kept uncommented (activated)

| command | example | comment |
| :--- | :---: | :--- |
| string version $=$ | $" 1.36 "$ | version of SIMBAD |
| int runID $=$ | 1821 | run identification number |
| module Ring | $\left({ }^{*}\right)$ | this creates the ring ring basic structure |
| module SyncPart | $\left(*^{*}\right)$ | create Synchronous particle |
| module MacroHerd | $\left(*^{*}\right)$ | create MacroHerd |
| string herdName $=$ | "MainHerd" | name of the herd $[1]$ <br> module Parallel |
| initialize parallel computation <br> module Output <br> double mSync $=$ | 1.0 | activate output module <br> mass of Sychronous particle/mass of the proton] <br> double charge $=$ |

NOTES: [1] There may be more than one herd

## 4 Run commands

Logical variables, here and in the folowing, may have a value $[0 / 1=$ false/true $]$.

| command | example | comment |
| :--- | :---: | :--- |
| int nTurns $=$ | 1000 | number of turns to be completed |
| double tSync $=$ | 15. | start Kinetic Energy of Synchronous particle $[\mathrm{GeV}]$ |
| double matrixEnergy $=$ | 10. | kinetic Energy that was used to generate matrices $[1]$ |
| double harmonicNumber $=$ | 1.0 | harmonic number |
| int longTrackOnly $=$ | $[0 / 1]$ | longitudinal tracking only $[2]$ |
| int usePendulum $=$ | $[0 / 1]$ | use pendulum equation for phi $[3]$ |
| double rTime $=$ | 0.0 | initialize ring time |
| int injTintegererval $=$ | 1 | turn interval for injecting new particles |
| int nMcrsPTurn $=$ | 100000 | number of macroparticles to inject per turn |
| int maxMacros $=$ | 100000 | maximum number of macros to be injected $[4]$ |
| double nRealMacros $=$ | 5.0 e 11 | number of "real" particles |
| int singleNodeStep $=$ | $[0 / 1]$ | Interactive single step $[5]$ |
| int nnodes $=$ | 100 | number of nodes to traverse if nTurns $==0[6]$ |

NOTES: [1] This may be different than tSync. Defaults to tSync if not activated. It is there to initialize matrix elements scaling with energy. [2] Default value is false. [3] In this example all macros are injected at once. [4] use pendulum equation to track phi instead of matrices. default value is true. [5] Variable to specify interactive single stepping through nodes. [6] If nnodes $==0$ then all nodes will be traversed.

## 5 Injection

### 5.1 External Distribution

A distribution of particles for tracking is created by a preprocessor (see Sect. 14, or generated internally. The internal generation is inherited from $O R B I T$. We tend to favor an external generation.

| command | example | comment |
| :--- | :---: | :--- |
| module Injector | $(*)$ | activate Injector module |
| string injectorName $=$ | "Injector" | injector name |
| int injectorOrder $=$ | 2 |  |
| int injType $=$ | 1 | type of population at injection $[1]$ |
| string injFilename $=$ | "HESR-240-de-4.dis" | name of file containing particle coordinates |

NOTES: $[0=$ built in (Joho), $1=$ read from file $]$

### 5.2 Built-in Distribution

This second setting if built-in distribution (Joho) is chosen. This is borrowed from Accsim [5].

| command | example | comment |
| :---: | :---: | :---: |
| int injType $=$ | 0 | type of population at injection |
| string injXInitializer $=$ | "JohoXDist" | name of injection function for x |
| string injYInitializer $=$ | "JohoYDist" | name of injection function for y |
| string injLongInitializer $=$ | "JohoLDist" | name of injection function for longitudinal |
| double betaXInj $=$ | 13.588 | X distribution parameters for matching [1] |
| double alphaXInj $=$ | 1.838 |  |
| double epsXLimInj $=$ | 100.0 |  |
| double MXJoho = | 1.0 |  |
| double $\times$ TailFraction $=$ | 0.0 |  |
| double $\times$ TailFactor $=$ | 1.0 |  |
| double x0Inj $=$ | 0.0 |  |
| double xP0Inj $=$ | 0.0 |  |
| double MLJoho = | 1.0 | M value of Joho Logitudinal distribution |
| double lTailFraction $=$ | 0.0 |  |
| double lTailFactor $=$ | 1.0 |  |
| double phiLimInj $=$ | 180.0 | injection bucket [-phiLimInj, phiLimInj] in deg |
| double dELimInj $=$ | 0.01 | limiting energy spread in GeV |
| double deltaPhiBunch $=$ | 0.0 | longitudinal center of the bunch [deg] |
| int nLongInjBunch = | 1 |  |
| double phiMaxInj = | 180. |  |
| double phiMinInj $=$ | -180 |  |
| double EInjMean = | 0.0114 |  |
| double EInjSigma = | 0.01 |  |
| double EInjMin = | 0.0114 |  |
| double EInjMax = | 0.0114 |  |

NOTES: [1] for Joho epsXLimInj $=$ epsXRMSInj $* 2$ * (MXJoho+1) . Same for Y.

## 6 Units. MAD input

SIMBAD uses MKSA units for all calculations. Input and output of phase space coordinates are in [mm] and [mrad] for transverse coordinate and angle, [rad] for longitudinal coordinate $\Phi$ and $[\mathrm{GeV}]$ for energy coordinate $\Delta E$. Phase and $\Delta E$ are evaluated with respect to the phase and energy $E$ of the synchronous particle.
$M A D$ coordinates are in $[\mathrm{m}]$ and [rad] in the transverse phase space, in [m] for the phase -in $M A D$ it is $\Phi:-c \Delta t-$, and [ 0$]$ for the energy coordinate -in $M A D$ it is $\Delta E: \Delta E / p c-$ Some unit transformation is needed here and there. It is also important to note that the longitudinal coordinate, "position", of each accelerator element in $M A D$ is evaluated at the end of element [6].
$S I M B A D$ uses the output from $M A D-8$ at the present time. Work is in progress to make $M A D-X$ the standard.

Two MAD output files are used by SIMBAD: the ".twiss" file and an ".echo" file. The former contains twiss functions for each element, plus the value of some basic quantities for the lattice, like
the betatron tunes and the transition energy. This file is created with the command "twiss" in the input to $M A D$. The latter contains the transfer matrices for each element in the accelerator and the second order transfer maps. First order matrices are generated with the MAD commands "setopts, echo", then "select, flag=first". First and second order maps are generated with 'setopts, echo", then "select, flag=second".

SIMBAD reads and compares the two file to create a third file "Twiss Plus Matrices", or TPM, containing all the info necessary for bare lattice tracking. Commands and parameters are

| command | example | comment |
| :--- | :---: | :--- |
| module TransMats |  | activate Transfer matrix module |
| string madFileTWISS $=$ | "input/HESR1.twiss" | MAD Twiss file |
| string madFileTM $=$ | "input/HESR1.echo" | MAD echo file |
| string madreadOutput $=$ | "TPM.dat" | created Twiss+Maps |
| int second_order $=$ | 1 | $0=1 . s t$ order track, $1=2 . n d$ order $[1]$ |

NOTES: [1] must be matched by the appropriate mad files. By default 0 is assumed.

## 7 Acceleration

Acceleration is simulated in $S I M B A D$ by mimicking what happens in a real synchrotron control room. That is, a file containing a table of values of the main magnet field $B$ vs. time, plus RF voltage for different harmonics and RF phase at each time. The code interpolates through the table, calculates the approriate energy for each value of the field and applies it to the synchronous particle. The RF voltage is applied to each macro at the nodes corresponding to RF cavities. The .conf file elements are as follows

| command | example | comment |
| :--- | :--- | :--- |
| module RampBAccel <br> string rampBASpecFile $=$ | "input/Ramp_HESR1.in" | activate Ramp-B-Acceleration <br> name of rampBA input file |

The Ramp file contains informations for every ramp node with a line for each node. Line structure is (items separated by a blank space)

| node name | node index | table | type (RAMPBV or RAMPV) | subroutine | bend radius [m] |
| :---: | :---: | :---: | :---: | :---: | :---: |

Example of Ramp file

| RAMPB1 | 15 | RAMPB1.input | RAMPBV | INTERPOLATEBV | 7.0 |
| :--- | :--- | :--- | :--- | :--- | :--- |
| RAMPB2 | 30 | RAMPB2.input | RAMPBV | INTERPOLATEBV | 7.0 |
| $\ldots .$. |  |  |  |  |  |

There must be a table file for each node. A table type RAMPBV is structured as

| number or RF harmonics: integer |  |  |  |  |
| :---: | :---: | :---: | :---: | :---: |
| time [msec] $\mathrm{B}[\mathrm{T}]$ | Volt(1) [kV] | Phase(1) [rad] | Volt(2) | Phase(2) ... |
| .. |  |  |  |  |
| .. |  |  |  |  |

A table type RAMPV is structured as
number or RF harmonics: integer
time [msec] Volt(1) [kV] Phase(1) [rad] Volt(2) Phase(2) ...
..

## 8 Space Charge. Impedances

There are many methods to calculate transverse space charge self forces on a high intensity beam, all based on the solution of the Laplace-Poisson-Ampere equations. A force on a given particle P is due to the field directly generated at the particle' by every other particle Q in the beam, plus the
forces due to image charge and current on the walls of the accelerator vacum chamber. Radial forces transalte into radial angle kicks, longitudinal forces in longitudinal energy kicks.

In the present version of $S I M B A D$ only the transverse Poisson problem is solved, i.e. only wall charge images are considered, and image currents are not. In addition, only perfectly conducting walls are considered. Poisson equation can be written either in integral form or in differential form. In the first case, the solution is found in SIMBAD using two methods: "Brute Force" or direct numerical integration, and FFT, where the Poisson equation is reduced to a convolution. In the second case, the differential Poisson equation is solved by writing the Laplacian on a transverse mesh and inverting it using standard linear equation solvers, like LU decomposition, or iterative methods.

After a solution is found of the transverse problem, the longitudinal kick is calculated by the potential difference at any transverse $x, y$ location between adiacent slices.

Finite conductivity of the walls and variation of the vall geometry along the beam path can be represented by longitudinal and transverse impedance tables, with real an imaginary part of the impedance listed for each harmonic mode (frequency). Longitudinal and transverse kicks are calculated in SIMBAD, following a method developed for the one-dimensional FermiLab code ESME [7] by first performing an FFT analysis of the beam current, in order to find harmonic components of the current. Then, each component of the impedance, in [ohm], is combined with a component of the current, in $[\mathrm{A}]$, to generate a voltage kick, with the appropriate amplitude, in [volt], and phase. These kicks produce a distorsion in the phase space distribution that can lead to instabilities and beam losses. It must be pointed out that impedance effects develop very slowly, and then require that the herd would be followed for many thousand revolutions in the simulation. Also, FFT of the beam can be made up to a large number of frequency with the needed accuracy only if the number of macros used in the simulation is large. For impedance studies parallel computing is a must. At the present time impedance effects are only calculated once per turn, using impedance budget tables for the full accelerator structure.

### 8.1 1-, 2-, and 3-dimension tracking

One dimension tracking is longitudinal tracking where we don't care about the transverse phase space that is only transformed along the lattice using the bare tune $M A D$ maps. Longitudinal transformations are done by solving the coupled discretized "pendulum equations" for $\Phi$ and $\Delta p / p$.

Two dimension tracking, that is sometimes called $2 \frac{1}{2}$, is performed by slicing the beam in many longitudinal partitions, and solving the space charge problem (see sec. /refsec:SpaceCharge) in each section. At each Transverse Space Charge node this operation is performed on each longitudinal slice, possibly with parallel computing. In this case, the longitudinal propagation is done by using the full $6 \times 61$.st order matrices and the $6 \times 6 \times 6$ second order transfer maps, or tensors.

Three dimension tracking is done by still slicing the beam, but solving the space charge problem with all macros in the herd reduced at the same time. We call this "freezing" the beam. Only in this case the space charge forces in all three dimensions are correctly calculated. The criterion for slicing the beam is that a slice should be a fraction of a $\beta$-function wave.

### 8.2 Longitudinal Space Charge. Longitudinal Impedances

Activate Longitudinal Space charge module based on FFT (don't use with freeze). This module is both calculating $L$ space charge with a theoretical formula and Long impedances [8], if the impedance file is present. If this module is on and Transverse Space Charge is also on, it must be int nLongSC $=0$ in TSC, otherwise SIMBAD would try to calculate the LSC twice with different methods

| command | example | comment |
| :--- | :---: | :--- |
| module FFTLongSC | "LSC1" | activate Longit. Space charge module |
| string fftLSCName $=$ | 770 | name of node |
| int fftLSCOrder $=$ | 128 | node index |
| int nLongBins $=$ | 5.0 | number of longitudinal bins |
| double b_a $=$ | $[0 / 1]$ | ratio between beam (round) and chamber $b / a$ |
| int useAvg $=$ | 10 | use averages |
| int nMacroLSCMin $=$ | minimum number of Macros $[1]$ |  |
| string fftLongSCSpecFile $=$ | input/ztab128.tab" | file of long. impedance data |

NOTES: [1] for LSC and TSC calcs to be done.
The file containing the longitudinal impedance data has format

| n0 | $\operatorname{real}(Z)$ | $\operatorname{imag}(Z)$ |
| :--- | :--- | :--- |
| n1 | $\operatorname{real}(Z)$ | $\operatorname{imag}(Z)$ |
| .. |  |  |

At the present only one table of L Imped is present, as a budget for the entire ring.

### 8.3 Transverse Space Charge

| command | example | comment |
| :--- | :---: | :--- |
| module TSpaceCharge |  | activate Transverse Space charge module |
| int nXBins $=$ | 32 | number of mesh points in x $[1]$ |
| int nYBins $=$ | 32 | number of mesh points in y |
| double hDimension $=$ | 160 | wall dimensions in mm $[2]$ |
| double vDimension $=$ | 160 |  |
| string wallsFile $=$ | 'input/walls.in" | input table for variable walls option [3] |
| int actPartLosses $=$ | $[0 / 1]$ | activate particle losses for collisions with walls |
| int tscCalcType $=$ | $[1 / 2 / 3 / 4]$ | method of calculating Transverse Space Charge [4] |
| double pOpt $=$ | 0.999922 | Jacobi spectral radius for SOR: optimal for 512x512 |
| double pOpt $=$ | 0.999912 | - optimal for $256 \times 256$ |
| double pOpt $=$ | 0.9997 | - optimal for $128 \times 128$ |
| double pOpt $=$ | 0.9986 | - optimal for $64 \times 64$ |
| double pOpt $=$ | 0.99469 | - optimal for $32 \times 32$ |
| double eps $=$ | 0.001 | smoothing parameter |
| double gridFactor $=$ | 0.001 | grid factor for extra spacing around distribution |
| int useMPISOR $=$ | $[0 / 1]$ | variable to enable parallel SOR solve [5] |
| double nMacsMinPercent $=$ | 0.01 | \% of macros for which TSC calcs will not be done $[6]$ |

NOTES: [1] the number of mesh points must be even. [2] At the present, need to be same value for H and V. If different, the largest is the default. These number can be overwritten by the variable wall option. Walls are perfectly conducting. [3] In a wall file each entry denotes a new section of wall [ mm ] vs. location around the machine [m]. The format is as follows: horiz[mm] vert[mm] sPos[m] [4] " 1 " is Sparse LU method, " 2 " is FFT, " 3 " is for SOR iterative method, " 4 " is BF. The grid must be square for all methods and the dimension of the chamber must be square for methods 1 and 3 . [5] This is only good with large grids. Make sure you test before committing to a production run. [6] For example, if the value is 1.0 then if the number of macros in a given element is less than one percent of the global number of macros the calcs will be skipped.

### 8.4 3-D

| command | example | comment |
| :--- | :---: | :--- |
| int nLongSC $=$ | 0 | no of times long. space charge is calculated per turn [1] |
| int longBins $=$ | 128 | number of bins to subdivide the beam for LSC calcs |
| int bmEnvLayout $=$ | $[1 / 0]$ | layout spacecharge nodes according to beam envelope <br> int betaFuncFact $=$ |
|  | 4 | variable to divide the space charge elements <br> - according to the length of the beta function |
| int freezeBeam $=$ | $[1 / 0]$ | variable to enable beam freezing [2] <br> int trimPhi $=$ |

NOTES: [1] the number of mesh points must be even. [2] At present only applies to iterative solver and only to serial runs.

### 8.5 Transverse Impedances

Transverse Impedances require an FFT of the beam transversely, to cople impedances with transverse beam modes

| verse beam modes | example | lomment |
| :--- | :---: | :--- |
| command |  | activate module |
| module FFT_TImpedance | "TIMPED1" | name of node |
| string ffTTImpedName $=$ | 780 | node index |
| int fftTImpedOrder $=$ | 32 | number of transverse bins |
| int nTImpedBins $=$ | 1.0 | ratio of beam height to width |
| double b_a_TImped $=$ | $[0 / 1]$ | use transverse averages |
| int useAvg_TImped $=$ | $[0 / 1]$ | minumum number of Macros for calcs |
| int nMacsMinTImped $=$ | input/FFTTImped.input" | T impedances table |
| string fftTimpedSpecFile $=$ | inple |  |

The FFT_TImpedance input file contains the information for the impedance values of the node. The file should contain (items separated by a blank)

| (nTImpedBins) lines with 9 columns per line: |
| :--- |
| nfrequency number |
| real(ZXImped_nplus) |
| imag(ZXImped_nplus) |
| real(ZXImped_minus) |
| imag(ZXImped_minus) |
| real(ZYImped_nplus) |
| imag(ZYImped_nplus) |
| real(ZYImped_minus) |
| imag(ZYImped_minus) |

## 9 Special Machine Elements

SIMBAD allows the insertion in the lattice of elements that have a special purpose, like foils, thin lenses or collimators

### 9.1 Foil

In some proton machines negative ions are injected and stripped to protons on a thin foil. Foils can also be placed in the lacttice for special purposes. The scattering and propagation through a foil is calculated. Hits to the foils are counted.

| command | example | comment |
| :--- | :---: | :--- |
| module Foil |  | activate Foil module |
| string foilName $=$ | "Foil" | name of the foil |
| int foilOrder $=$ | 2 | node order of the foil |
| int useFoilScat $=$ | $[0 / 1]$ | flag to use foil scattering |
| double xfmin $=$ | -100. | min. foil coordinate in x[mm] |
| double xfmax $=$ | -25. | maximum foil in x $[2]$ |
| double yfmin $=$ | -100.0 | miniumum foil in y |
| double yfmax $=$ | 100.0 | maximum foil in y |
| double foilThickness $=$ | 300. | thickness of the foil $\left[\mathrm{mgram} / \mathrm{mm}^{2}\right]$ |
| double foilFac $=$ | 0.665 | scale edge so that $1 \%$ miss foil[1] |

NOTES: [1] Foil factor $=1.33 /(($ Injection::MXJoho +1$) / 2) .[2]$ inj $->x 0$ Inj + sqrt (fabs(foilFac * inj-¿betaXInj * inj->epsXLimInj))

### 9.2 Bump

Programmable orbit bump. Three types of bump. Most likely: injection bumps are simulated in this module. If a bump table is used

| command | example | comment |
| :--- | :---: | :--- |
| module IdealBump1 |  | activate Ideal Bump type 1 |
| string bump1Name $=$ | "UP-BUMP" | Bump 1 name |
| int bump1Order $=$ | 1 | Bump 1 node index |
| module IdealBump2 |  | activate Ideal Bump type 2 |
| string bump2Name $=$ | "DOWN-BUMP" | Bump 2 name |
| int bump2Order $=$ | 4 | Bump 2 node indx |
| int bump1UD $=$ | $[0 / 1]$ | Up/Down bump1, 0=down, $1=$ up |
| string bumpFileName $=$ | "input/bump.tab" | file from which to read bump.ramp |
| double tBmp0 $=$ | 0.0 | initial Bump time |
| double tBmpF $=$ | 1. | final Bump time |
| double eFTX $=$ | 8 | e-fold time ratio (after this, the bump is gone) |
| double eFTY $=$ | 0.00 | e-fold time |

If a bump table is NOT used, then

| command | example | comment |
| :--- | :---: | :--- |
| string bumpFunct $=$ | EFoldBump" | bump function to be used |
| double $x B m p 0=$ | -30. | initial value of the x-bump |
| double $\mathrm{xBmpF}=$ | 0. | final value of the x-bump |
| double $\mathrm{xPBmp} 0=$ | -5. | initial value of the xP-bump |
| double $\mathrm{yBmp}=$ | 0.0 | initial value of the y-bump |
| double $\mathrm{yBmpF}=$ | 0.0 | final value of the y-bump |
| double yPBmp $0=$ | 0.0 | initial value of the yP-bump |

### 9.3 Integrable Lens 2D

| command | example | comment |
| :--- | ---: | :--- |
| module IntegLens2D <br> string intLens2DSpecFile $=$ | "input/IntLens2D.in" | activate Integrable 2D Lens <br> name of integrable 2D Lens input file |

The IntegLens2D input file contains the information for each node. The format of the file must be as follows (a blank between items)

| name of the node <br> non linear coefficient | name of output file <br> linear coefficient | node order number | first line <br> :second line |
| :--- | :--- | :--- | :--- |

skip a line and repeat for the next IntegLens node

Example of IntegLens file
IL1 15
$5.0 \quad 2.0$
IL2 25
$7.0 \quad 1.0$

### 9.4 Rectangular Aperture

| command | example | comment |
| :--- | :--- | :--- |
| module RectAperture <br> string rectApFileName $=$ | "input/RectAp.in" | activate the module <br> name of input file |

The RectAperture input file contains the information for each node. The format of the file must be as follows (a blank between items)

| node name <br> xmin | node index |  |  |  | first line |
| :--- | :--- | :--- | :--- | :--- | :--- |
| xmax | ymin | ymax | transparent[0/1] | :second line |  |

skip a line and repeat for the next RectAperture node
Example of Rectangular Aperture file:

| RectAp1 | 5 |  |  |  |
| :--- | :--- | :--- | :--- | :--- |
| 10.0 | 25.0 | 5.0 | 50.0 | 0 |

RectAp2 ....

### 9.5 Momentum Aperture

| command | example | comment |
| :--- | :--- | :--- |
| module MomentumAperture <br> string momApFileName $=$ | "input/MomentAp.in" | activate module <br> name of input file |

The MomentumAperture input file contains the information for each node. The format of the file must be as follows (a blank between items)

| name of the node <br> $\max (\mathrm{dp} / \mathrm{p})$ | node order number <br> calcFreq | :first line <br> :second line |
| :--- | :--- | :--- |
| skip a line and repeat for the next MomentumAperture node |  |  |

Example of Momentum Aperture file:
MomentAp1 5
10.03

### 9.6 Thin Multipole

| command | example | comment |
| :--- | :--- | :--- |
| module ThinMPole <br> string thinMPoleSpecFile $=$ | "input/ThinMPole.in" | activate module <br> name of input file |

The ThinMPole input file contains the information for each node. The format of the file must be as follows (a blank between items)

| name of the node <br> order of multipole | node order number <br> integrated strength[1] | skew variable | :first line |
| :--- | :--- | :--- | :--- |
| :second line |  |  |  |

skip a line and repeat for next ThinMPole node
NOTES: [1] integrated strength of the field expansion (kl).

Example of Thin Multipole file:

| TMP1 | 15 |  |
| :--- | :--- | :--- |
| 1 | 2.0 | 0 |
|  |  |  |
| TMP2 | 25 |  |
| 2 | 3.0 | 1 |

### 9.7 Lattice Kicks

| command | example | comment |
| :--- | :--- | :--- |
| module LatKicks <br> string tKickInput $=$$\quad$ "input/TKicks.in" | activate module <br> name of input file |  |

### 9.8 RF Cavity

To be used if you have a RF cavity at constant voltage and fixed beam energy (No Ramp)

| command | example | comment |
| :--- | :--- | :---: |
| module RFCavity <br> string rfSpecFile $=$ | "input/RFCav.in" | activate module |
| name of input file |  |  |

The RFCavity input file contains the information for each RFCavity. The format of the file must be as follows (a blank between items)

| node name | no of harmonics | node index | function | :first line |
| :--- | :--- | :--- | :--- | :--- |
| $\operatorname{volt}(1)[\mathrm{kv}]$ | harmonic number | phase(1) |  | :second line |
| $\operatorname{volt}(2)[\mathrm{kv}]$ | harmonic number | phase(2) |  | :third line |
| $\ldots$ |  |  |  |  |

skip a line and repeat for the next RFCavity node
Example of RF Cavity file

| RF1 | 2 | 75 | CONSTVOLTS |
| :--- | :--- | :--- | :--- |
| 40.0 | 1.0 | 0.0 |  |
| -20.0 | 2.0 | 0.0 |  |
|  |  |  |  |
| RF2 | 2 | 100 | CONSTVOLTS |
| 41.0 | 3.0 | 0.0 |  |
| -21.0 | 2.0 | 0.0 |  |

## 10 Output

| command | example | comment |
| :---: | :---: | :---: |
| string of1 | "Ring.dat" | name of file containing the ring structure |
| int showSingleNode $=$ | [0/1] | display node information [no/yes] [1] |
| int outputScreenToFile $=$ | [0/1] | output screen data to file [no/yes] |
| string outputFileName $=$ | "screen.dat" | name of output file [2] |
| int dumpLostParts $=$ | [0/1] | dump lost particles to a file [no/yes] |
| string lostPartsFileName | "LostParts.dat" | name of file to contain lost particles |
| int showSingleNode $=$ | [0/1] | display some info to screen after every node |
| NOTES: [1] switch to display some node information to screen after every node completes. [2] Output file contains: |  |  |
|  |  |  |
| Turn Time n-Macros | n-Reals hits/p |  |

## 11 Plot

Graphic output is done with GNUPLOT. One can run GNUPLOT interactively, i.e. getting continuous grapic output on the screen while $S I M B A D$ is running, or dump data to graphic postscript files. Interactive plot is not possible with parallel runs. Command and setting for GNUPLOT are

| command | example | comment |
| :--- | :---: | :--- |
| module GPlot |  | activate GPlot module |
| string gpNodename $=$ | "gplot" | GNUPLOT node name |
| int gpPSOut $=$ | $[0 / 1]$ | output plots to postscript [no/yes] <br> string gpFilenameBase $=$ <br> int gpScrOut $=$ <br> inESR1.gp" <br> int gpNodes $=$ |
| base name of GPlot postscript files |  |  |
| int gpInitialOIndex $=$ | $[0 / 1]$ | output plots to screen [no/yes] <br> int gpOrderSpacing $=$ |
| int gpActTurnInt $=$ | 15 | number of GPlot nodes |
| int startPlots $=$ | 40 | GPlot beginning node index |
| int stopPlots $=$ | 20 | GPlot node order spacing |
| int prune $=$ | 600 | turn activation interval (..every n turns..) |
| int x_yPlot $=$ | 10 | turn number in which to start plotting |
| int x_xpPlot $=$ | $[0 / 1]$ | prunes the number of particles plotted $[1]$ |
| plot y vs x [no/yes] |  |  |
| int y_ypPlot $=$ | $[0 / 1]$ | plot xp vs x [no/yes] |
| int phi_dEPlot $=$ | $[0 / 1]$ | plot yp vs y [no/yes] |
| int w_dots $=$ | $[0 / 1]$ | plot dE vs phi [no/yes] |
| string gpScrSize $=$ | $[0 / 1]$ | allows feature "with dots" [no/yes] |
| int setRanges $=$ | $800 \times 800 "$ | screen size |

NOTES: [1] i.e. if prune $=10$ then only 1 in 10 particles will be plotted.
[2] $0=$ set autoscale. If 1 , set ranges.
If setRanges $=1$, no autoscale, plot ranges are prescribedd as

| int setRanges $=$ | 1 |  |
| :--- | :---: | :--- |
| double xy_xMin $=$ | -100.0 | x-y range |
| double xy_xMax $=$ | 100.0 |  |
| double xy_yMin $=$ | -100.0 |  |
| double xy_yMax $=$ | 100.0 |  |
| double x_xp_xMin $=$ | -100.0 | x-xp range |
| double x_xp_xMax $=$ | 100.0 |  |
| double x_xp_yMin $=$ | -100.0 |  |
| double x_xp_yMax $=$ | 100.0 |  |
| double y_yp_xMin $=$ | -100.0 | y-yp range |
| double y_yp_xMax $=$ | 100.0 |  |
| double y_yp_yMin $=$ | -100.0 |  |
| double y_yp_yMax $=$ | 100.0 |  |
| double phi_dE_xMin $=$ | -3.14 | phi-dE range |
| double phi_dE_xMax $=$ | 3.14 |  |
| double phi_dE_yMin $=$ | -0.01 |  |
| double phi_dE_yMax $=$ | 0.01 |  |

## 12 Diagnostics

$S I M B A D$ has many diagnostics modules, inherited from $O R B I T$. Grdually, we are moving out of them, relying instead on post processing the fundamental output files, especially the Phase Dump. To generate a Phase Dump file we need some commands.

### 12.1 Phase Space Diagnostics

| command | example | comment |
| :--- | ---: | :--- |
| module PhaseSpace <br> string phaseSpaceFName $=$ | "input/PhaseSpace.in" | activate module <br> name of input file |

The PhaseSpace input file contains the information necessary for each PhaseSpace node. The format of the file MUST be as follows. The first line contains these records (separated by blanks)

```
name of the PS Diagnostics node
name of the file to which dump information
node index
turn interval for activation of the node[1]
Freeze beam [1/0=yes/no]
modulo for output of particles [2]
skip a line and repeat for the next PhaseSpace Diagnostic node
```

NOTES: [1] f the turn interval activation number is 1, then diagnostic information will be dumped every turn, else every n turns. [2] $1=$ all particles and must be 0 .

Example of Phase Spase Diagnostics file

| PSnode1 | "PSDump1.dat | 1425 | 0 | 10 | 10 |
| :--- | :--- | :--- | :--- | :--- | :--- |
| PSnode2 | "PSDump2.dat | 1725 | 0 | 10 | 10 |
| .. |  |  |  |  |  |

### 12.2 Tune Diagnostics

Betatron and Synchrotron tune are also calculated in SIMBAD with various methods.
For the betatron tunes, a simple method is to count the number of transverse oscillations by every macro per turn, or, better to compare the average wavelength of a transverse oscillation with the length of one turn, for many turns. This method, still embedded int SIMBAD proper, requires a large number of turns for a good accuracy, and for the very same reason cannot be used to follow the evolution of the tune of a specific macro or a number of macros turn by turn. A second method for the betatron tune, also in the body of $S I M B A D$, is based on the FFT of orbits. This method can give interesting results -e.g. higher order tunes- but requires again many turns.

An alternative method, much more powerful, is based on the analysis of the eigen values of the one turn matrix. Betatron and synchrotron tunes can be calculated at every turn, using the phase space for the six preceding turns, with great accuracy. This method is impemented in a Post Processor, or Utility subroutine, described in Sec. 15.

The built-in routines for tune calculations are driven by the following

| command | example | comment |
| :--- | :---: | :--- |
| module Tunes |  | activate module |
| string tunesFileName $=$ | "input/tunes.in" | name of input file |
| int startTune $=$ | 400 | turn to start the tune calculation $[1]$ |
| int stopTune $=$ | 499 | turn to stop the tune calculation |
| int calcCoherentTune $=$ | $[0 / 1]$ | flag to calculate the coherent tune |
| string coherentTuneFilename $=$ | "CoherentTune.out" | coherent tune output file |
| int calcIncoherentTune $=$ | $[0 / 1]$ | flag to calculate the incoherent tune |

NOTES: [1] Should have stopTune - startTune i 10 to get a good statistics. - 1 means never.
The Tunes input file contains the information necessary for each Tunes node. The format of the file must be as follows (a blank between items)

```
name of the node
The name of file to which dump information
node order number
    Example of Tune Diagnostics file
| TunesNode1 "tunes1.dat" 13
```


### 12.3 Momentum Diagnostics

| command | example | comment |
| :--- | :---: | :--- |
| module Moment |  | activate module |
| string momentFileName $=$ | "input/Moment.in" | name of input file |
| int momentOrder $=$ | 2 | order to which moments are calculated |

The format of the file must be as follows. The first line contains these records (items separated by blanks)

| name of the node <br> The name of file to which dump information <br> node order number <br> turn interval |
| :--- |
| skip a line and repeat for the next RectAperture node |
| MomNode1 "MomDump1.dat 1512 0 <br> 100    <br> MomNode2 "MomDump2.dat 1712 0 <br> 100    <br> ..    |

### 12.4 TSC Kicks Diagnostics

| command | example | comment |
| :--- | :---: | :--- |
| module TSCKicks <br> string tscKicksSpecFile $=$ | "input/tscKicks2D.in" | activate module <br> configuration file |

The TSCKicks input file contains the information required for each TSCKicks diagnostic node. The format of the file must be as follows (items separated by blanks)

| name of the node |
| :--- |
| name of file to which dump information |
| NTSC element name [1] |
| turn interval |
| start turn |
| stop turn |

NOTES: [1] NTSC is the transverse space charge element number that should immediately precede this diagnostic.

### 12.5 Accelerate Diagnostics

| command | example | comment |
| :--- | :---: | :--- |
| module Accelerate |  |  |
| string accelFileName $=$ | "input/Accel.in" | configuration file |
| int calculateBucket $=$ | $[0 / 1]$ | flag to enable calculation of bucket |

The Accelerate input file contains the information required for each Accelerate diagnostic node. The format of the file MUST be as follows

```
    name of the node
    name of output file
    node order number
    integer to specify the type of RF associated [1]
    ordinal number of the specific RF node associated [2]
    turn interval
    skip a line and repeat for the next RectAperture node
NOTES: [1] 0 = an RFCavity node, and 1 = a RampBAccel node. [2] starting with zero
```

Output contains

| Turn | Time | n-Macros | T-Sync | B_f | beta | phase | Volt | w_synch | E_bck | E_bnc | A_bck | A_bnc | dp/p |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | $[\mathrm{msec}]$ |  | $[\mathrm{GeV}]$ |  |  | $[\mathrm{deg}]$ | $[\mathrm{kV}]$ | $[\mathrm{Hz}]$ | $[\mathrm{MeV}]$ | $[\mathrm{MeV}]$ | $[\mathrm{eV}-\mathrm{s}]$ | $[\mathrm{eV}-\mathrm{s}]$ | $\%$ |

For example, if you have a single RFCavity node in the ring at position 20 you might use the following...
| AccNode1 "AccDump1.dat" $21 \quad 0 \quad 1 \quad 100$
If you had 3 RampBAccel nodes in the ring at positions 51, 101, and 201 you might use the following...

| AccNODE2 | "AccDump2.dat" | 52 | 1 | 0 | 100 |
| :--- | :--- | :--- | :--- | :--- | :--- |
| AccNODE3 | "AccDump3.dat" | 102 | 1 | 1 | 75 |
| AccNODE4 | "AccDump4.dat" | 202 | 1 | 2 | 50 |

### 12.6 Canonical Coordinate Diagnostics

| command | example | comment |
| :--- | ---: | :--- |
| module CanonicalCoords <br> string ccFileName $=$ | "input/CanonCoords.in" | activate module <br> name of configuration file |

The Canonical Coordinate conf file contains the information required for each CC diagnostic node. The format of the file must be as follows (items separated by blanks)

| name of the node | name of output file | node index |
| :--- | :--- | :--- |
| turn interval |  |  |
| skip a line and repeat for the next CCDiag node |  |  |

Example
| CCNode1 CC1.dat 81

### 12.7 Transverse Emittance Diagnostics

| command | example | comment |
| :--- | :---: | :--- |
| module Emittance <br> string emitFileName $=$ | "input/TEmit.in" | activate module <br> name of config file |

The Transverse Emittance conf file contains the information required for each TEM diagnostic node. The format of the file must be as follows (items separated by blanks)


### 12.8 Longitudinal Emittance Diagnostics

| command | example | comment |
| :--- | :---: | :--- |
| module LongEmittance <br> string lEmitFileName $=$ | "input/LEmit.in" | name of conf file |

The Longintudinal Emittance input file contains the information necessary for each LongEmit node. The format of the file must be as follows (items separated by blanks)

| name of the node | name of output file | node index |
| :--- | :--- | :--- |
| skip a line and repeat for the next LEM node |  |  |

Example
| LEMNode1 "LEM1.dat" 850

### 12.9 Action Diagnostics

| command | example | comment |
| :--- | :--- | :--- |
| module Actions <br> string actionsFileName $=$ | "input/Actions.in" | activate Action Diagnostics module <br> name of conf file |

The Actions input file contains the information necessary for each Actions node. The format of the file must be as follows (items separated by blanks)

| name of the node | name of output file | node index | turn interval |
| :--- | :--- | ---: | :--- |
| skip a line and repeat for the next RectAperture node |  |  |  |

Example
| ActNode1 "Act1.dat" $9 \quad 15$

### 12.10 Stat Lat Diagnostics

| command | example | comment |
| :--- | :---: | :--- |
| module StatLat <br> string statlatFileName $=$ | "input/StatLat.in" |  | | activate StatLat Diagnostic module |
| :--- |
| name of conf file |

The StatLat input file contains the information necessary for each StatLat node. The format of the file must be as follows (items separated by blanks)

| name of the node | name of output file | node index | turn interval |
| :--- | :--- | ---: | ---: |
| skip a line and repeat for the next RectAperture node |  |  |  |

Example
| SLNode1 "SL1.dat" 925

### 12.11 Check Points

Enable if program checkpointing is desired

| command | example | comment |
| :--- | :---: | :--- |
| module CheckPoint |  | activate module |
| int chkPntTurnInt $=$ | 25 | turn interval for writing checkpoint files |
| int cleanChkPnt $=$ | $[0 / 1]$ | clean checkpoint files on exit |

## 13 Modularity, Pre- and Post-Processors

Our effort is to make the structure of $S I M B A D$ as modular as possible, with the use of an expanding number of Pre- and Post- processors. The purpose is twofold: (i) we find it desirable to decrease the sheer size of the code, devolving many tasks to auxiliary codes than are used only if needed; (ii) we want SIMBAD to migrate and work within the "Unified Accelerator Library", or $U A L$, as a wrapper, and for this modularity is a prerequisite.

The old $O R B I T$ was full of preparatory routines and diagnostic modules. A preparatory routine is the one described in section 6 above, that reads the output of $M A D$. This is still part of the present version of SIMBAD, but is gradually being replaced by a pre-processor that will be able to read the optics not only from other versions of $M A D$, but also from other codes, like TEEPOT or others. Another pre-processor creates the initial particle distribution in the herd. Since any diagnostics is based on some analysis of the output of the phase space ( 6 coordinates), we considered it more profitable to do these analyses with post-mortem processors instead than burden the basic tracking code with specialized routines.

## 14 Pre Processors

A collection of pre-processors are available for $S I M B A D$. One, of course is $M A D$, if we dare to call it a pre processor. $M A D$ creates the input optics files that contain the twiss functions and the transfer maps, as described in Sect. 6. Only $M A D-8$ is compatible with $S I M B A D$ so far.

### 14.1 MAD

A typical MAD-8 input (e.g. HESR1.mad) must contain the following commands

| example of MAD command lines | comment |
| :--- | :--- |
| $\mathrm{PC}:=14.5$ | particle momentum |
| beam, particle=proton, momentum $=\mathrm{PC}$ | set up particle [1] |
| call filename='HESR1.lat' | lattice file |
| setopts, echo | insures that maps are output to .echo file |
| select, flag=SECOND, range $=$ full | writes transfer maps |
| twiss, tape='HESR1.twiss',deltap $=00$ | produces and writes twiss.file [2] |

NOTES: [1] This is important for the longitudinal components of the transport maps that depend on energy. [2] The created .echo file will carry the same name as the .mad file. In our example HESR1.echo

### 14.2 MAKEPOP: Particle Distribution Creator

The input particle distribution for tracking can either be creates inside $S I M B A D$ or read from an external file, as described in Sect. 5.

An external distribution is created with the Fortran-77 code MAKEPOP2, based on formalism presented by G.Franchetti [9]. The operation of the code is as follows
$>$ makepop $2<$ input_file
An example of input file containing the parameters for the distribution is

| variable | value | comment |
| :--- | :--- | :--- |
| \&par |  | namelist group begin mark |
| npart $=$ | 10000 | number of particles in the distribution |
| Tdist $=$ | 'G1' | transverse distribution=random Gaussian [1] |
| iseed $=$ | 1812 | seed for random number generation |
| alfax $=$ | -0.33569 | Twiss- $\alpha_{x}$ |
| betax_m $=$ | 68.8921 | Twiss- $\beta_{x}$ |
| ex_m_rad $=$ | $1 . d-6$ | X-emittance [m-rad] |
| alfay $=$ | -4.4297 | Twiss- $\alpha_{y}$ |
| betay_m $=$ | 148.107 | Twiss- $\beta_{y}$ |
| ey_m_rad $=$ | $1 . d-6$ | Y-emittance [m-rad] |
| dx_m $=$ | $0 . d 0$ | horizontal displacement of beam |
| dy_m $=$ | $0 . d 0$ | vertizontal displacement of beam |
| LDist $=$ | 'LG' | longitudinal distribution Linear-Gaussian $[2]$ |
| Dphi_deg $=$ | 240. | r.m.s. $\Phi$ width [deg] |
| dEo $=$ | 0.014 | r.m.s. $\Delta E$ half-width [GeV] |
| outfile $=$ | 'HESR1-e5-240-em-6-de-3.dis' | generated distribution file |
| $/$ |  | namelist group end mark |

NOTES: [1] Transverse options are 'G1' (Gaussian), 'L' (Linear), 'KV' (Kapchinskij-Vladimirskij), 'WB' (water bag). [2] Longitudinal distributions are: 'LG' (Linear-Gaussian), 'PG' (ParabolicGaussian), 'GG' (Gaussian-Gaussian)

## 15 Post Processors

A collection of post-processing routines are in the folder "utils" of the SIMBAD distribution. They are mostly operating on the Phase Space output file. The operation of each PostProc is described below; a description of their operation appears on the screen by simply invoking the routine.

First, let us recall that the Phase Space output consists of a file containing the macro number and the six phase space coordinates of the macro. The file is arranged in sectors, on for each turn dumped, separated by two blank lines. This arrangement permits to plot phase space diagrams for each turn separately using the "i" (index) option of GNUPLOT. The "prune" capability of the Phase Space Diagnostics module of SubSec. /refsubs:PSDiag allows to dump to the file only a subset of the macros used in the simulation, in order to limit the size of the file itself.

## 15.1 ttunes

This C++ routine operates on the Phase Space output file. Its purpose is to calculate the betatron and synchrotron tune from the eigenvalues of the One Turn Matrix (OTM) built from six turns of each macroparticle [10]. The output of ttunes can be either a file containing the tunes for all turn is dumped or a specific turn. Another alternative dump contains the OTM and the eigenvalues for a specific particle.

To work correctly, one should calculate the tunes at a location in the machine where the $\beta$-Twiss function for X and Y are as different as possible from each other. This will facilitate the attribution of the eigenvalue to the appropriate coordinate.

Let us invoke the routine
$>$ ttunes
usage:
ttunes $-i<$ filename_in $>-x<$ betaX $>-y<$ betaY $>[-m<M-p a r t n o j-n<$ recno $>][$ a $<$ average num $>]$
The input, filename_in should contain the phasespace output with data dumped each turn.
betaX should be the betaX value associated with the phasespace output
betaY should be the betaY value associated with the phasespace output
If the calculated matrix is desired M-partno is the particle number
for which the matrix calculated after the initial records are processed
and recno is the record number in which to print the matrix will be stored to file "matrix.dat".
The processing of the file will end at that point.
average_num is the number of turns over which the tunes are averaged. This means that the tune output
will be average_num turns per record.
average_num must be an integer greater than 1
Data output, consisting of partno, tunex, tuney, tunez, alphaX, betaX, alphaY, betaY, alphaZ, betaZ (OTM), is on stdout, while status output, which consists of record (turn) number and particle count, is on stderr.

Example to write tunes (and macro number) for each macro and each turn to a file "tune.dat". In this example the values of the : $\beta$-functs at the chosen locasion were $69[\mathrm{~m}]$ and $148[\mathrm{~m}]$, respectively | ttunes -i PSDump.dat -x 69. -y 148. > tunes.dat

Example to write the OTM for macro no. 10, at turn 100, to a file"matrix-10.100.dat" (first record has index 0 )
| ttunes -i PSDump.dat -x 69. -y 148. -m 10 -n $99>$ matrix-10.100.dat

## 15.2 tunecontour

This $\mathrm{C}++$ routine was designed to operate on output files created by the Tune Diagnostic module of SubSec. 12.2. It creates $G N U P L O T$ compatible file for 2 D and 3 D contours of betatron data in the $Q_{x}-Q_{y}$ plane.

Let us invoke the routine

```
> tunecontour
        usage:
tunecontour -d <dimension> -i <filename_in> -o <filename_out>
contours may be plotted using gnuplot with the following commands:
set cntrparam levels auto <number>
set nosurface
set contour
set view 0,0
splot '<filename_out>' w l
```


## 15.3 picktunes

This C++ routine was designed to operate on output files created by the Tune Diagnostic module of SubSec. 12.2. It picks up the betatrn tune of specific particles

Let us invoke the routine
$>$ picktune
usage:
picktunes -i <filename_in> $[-\mathrm{t}<0,1>]$
The input file should contain the tunes output.
Enter the particle numbers on std input.
The last entry should be the word "end"
The output is printed on stdout
If - t has argument 0 , the format of the input is assumed to be the ORBIT tunes output file.
If -t has argument 1 , the format of the input is assumed to be the output from ttunes.
The default is 1 .

## 15.4 pickparts

This C++ routine was designed to operate on output files created by the Phase Space Diagnostic module of SubSec. 12.1. It picks up the phase space coordinates of specific particles for all turns dumped.

Let us invoke the routine
$>$ pickparts
usage:
pickparts -i <filename_in> -o <filename_out> [-ns]
the input file should contain the phasespace output
enter the particle numbers on std input
the last entry should be the word "end"
the output file will contain the phasespace for those particles
-ns specifies that no spaces should be added between records

## 15.5 partsemit

This $\mathrm{C}++$ routine was designed to operate on output files created by the Phase Space Diagnostic module of SubSec. 12.1. It calculates the transverse emittance of the Herd for all turns dumped with the formula

$$
\epsilon_{x}=\left[<\hat{x}^{2}><{\hat{p_{x}}}^{2}>-<\hat{x}{\hat{p_{x}}}^{2}\right]^{1 / 2}, \hat{x}=x-<x>, \hat{p_{x}}=p_{x}-<p_{x}>
$$

Let us invoke the routine
$>$ partsemit
usage:
partsemit -a <alpha> -b < beta> -e <emmit> -l <lowerbnd> -u <upperbnd> -i <file_in>
each particle is tested against the c-s invariant < - lowerbnd-([upperbnd-)*emmit
the input file should be a PhaseSpace diagnostic file
the output will be printed to stdout
that are within the emittance range specified

## 15.6 longemit

This $\mathrm{C}++$ routine was designed to operate on output files created by the Phase Space Diagnostic module of SubSec. 12.1. It calculates the longitudinal emittance of the Herd

Let us invoke the routine
$>$ longemit
usage:
longemit -i <filename_in> -s $<\mathrm{E}>[-\mathrm{r}<$ record $\sharp>][-\mathrm{e}<\mathrm{E} 0>]$
the input file should contain the phasespace output
$<\mathrm{E}>$ is the energy of the synchronous particle ( GeV )
$<$ record $\sharp>$ denotes the record number in the phasespace output
$<$ record $\sharp \mathrm{y}>$ should index starting from 0
$<\mathrm{E} 0>$ is the rest energy of the particle $(\mathrm{GeV})$, default is proton

## 15.7 extractrec

This $\mathrm{C}++$ routine was designed to operate on output files created by the Phase Space Diagnostic module of SubSec. 12.1. It selects a single record (turn) from the file

Let us invoke the routine
$>$ extractrec
usage:
extractrec -i <filename_in> [-r <record $\sharp>$ ]
the input file should contain the phasespace output
<record $\sharp>$ denotes the record number in the phasespace output
$<$ record $\sharp>$ should index starting from 0
default value is 0

## 15.8 binphsp

This C++ routine was designed to operate on output files created by the Phase Space Diagnostic module of SubSec. 12.1. It bins and creates 3D histograms of the phase space

Let us invoke the routine
> binphsp
usage:

```
binphsp -x <xdim> -y < ydim> [-xparm < 1-6>][-yparm < 1-6>]
[-x\operatorname{max}<x\operatorname{max}>][-x\operatorname{min}<x\operatorname{min}>][-y\operatorname{max}<y\operatorname{max}>][-ymin<ymin}>
[-t<type>][-z<plotZero >] -i<inputfile>
```

binphsp bins phasespace parameters yparm vs xparm using a grid (xmax-xmin) * (ymax-ymin) in size with xdim * ydim grid points the input file should be the PhaseSpace output file the type may be specified as 0 or 1 where $0=2 \mathrm{D}$ and $1=3 \mathrm{D}$ plotZero may take values of 0 or 1 where $0=$ no and $1=$ yes plotZero should be set to 1 if contour lines are desired this will result in greater output volume

## References

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