

*SMOG-SERVER.ORG*

**THE SMOG  
USER GUIDE**

*EXCERPT ON USING WHAM.jar*

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[HTTP://SMOG-SERVER.ORG/SB\\_ANALYSIS.HTML](http://smog-server.org/SB_ANALYSIS.HTML)

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# Contents

<b>1</b>	<b>Weighted Histogram Analysis - WHAM.jar</b>	<b>2</b>
1.1	Introduction . . . . .	2
1.2	Downloading WHAM.jar . . . . .	2
1.3	Citing WHAM.jar . . . . .	3
1.4	Running WHAM.jar . . . . .	3
1.5	Configuration file . . . . .	3
1.5.1	Basic configuration file: compute density of states ( $\Omega$ ), i.e. run WHAM	3
1.5.2	Full configuration parameter list . . . . .	4
1.6	Compute $F(Q), S(Q), E(Q)$ at $T$ from $\Omega(E, Q)$ or $\Omega(E, Q_1, Q_2)$ . . . . .	4
1.7	Compute $C_V(T), E(T), S(T), F(T)$ from $\Omega(E, \dots)$ . . . . .	5
1.8	Compute $\langle Q(T) \rangle$ from $\Omega(E, Q)$ . . . . .	5
1.9	Compute $\langle Q_2(Q_1) \rangle$ from $\Omega(E, Q_1, Q_2)$ . . . . .	6
1.10	Compute $\langle Q_3(Q_1, [Q_2^{\text{low}}, Q_2^{\text{high}}]) \rangle$ from $\Omega(E, Q_1, Q_2, Q_3)$ . . . . .	6
1.11	Reweighting of umbrella sampling . . . . .	7
1.11.1	Harmonic umbrella . . . . .	7

## 1 Weighted Histogram Analysis - WHAM.jar

### 1.1 Introduction

This section describes a Java application `WHAM.jar` which at its heart uses a well established algorithm described here [1]. `WHAM.jar` is tailored for the sort of analysis that is most often performed on SBMs, including free energy perturbation, thermal averaging of reaction coordinates and free energy as a function of (1 or 2) reaction coordinates. The basic operation involves providing `WHAM.jar` with a set of histograms from constant temperature simulation with various temperatures and umbrella parameters, and a configuration file. Based on the options set in the configuration file (or at the command line) `WHAM.jar` will perform the appropriate analysis. The WHAM algorithm itself [1] provides an optimal density of states ( $\Omega$ ). From the  $\Omega$  many thermodynamic quantity of interest can be calculated.

### 1.2 Downloading WHAM.jar

`WHAM.jar` is available at [smog-server.org](http://smog-server.org) ([http://smog-server.org/SB\\_analysis.html](http://smog-server.org/SB_analysis.html)). There are no plans to make the source code publicly available. Any feature requests or source code requests should be directed to [info@smog-server.org](mailto:info@smog-server.org).

## 1.3 Citing WHAM.jar

WHAM.jar is part of the SMOG group of tools, so please cite [2].

## 1.4 Running WHAM.jar

Like any java application, no compilation is necessary, but a virtual machine is required; WHAM.jar requires JRE 1.6.0\_29 or greater. At the command line, the basic syntax is

```
user$ java [-Xmx1000m] -jar WHAM.jar --config configurationFile [ argument... ]
```

-Xmx1000m assigns 1000 MB of RAM to the Java virtual machine heap. With large histograms (4 or 5 dimensions) the default heap allocation can run out which gives the following error:

```
java.lang.OutOfMemoryError: Java heap space
```

## 1.5 Configuration file

### 1.5.1 Basic configuration file: compute density of states ( $\Omega$ ), i.e. run WHAM

```
numDimensions 2 #number of dimensions in the dos histogram
                #energy and Q, energy is always in the first column, can have up to 3 reaction coordinates

kB 0.008314     #Boltzmann constant

run_wham       #creates optimal density of states, writes to dosFile, comment out to not run

dosFile dos     #density of states output filename for run_wham, input filename for analysis (i.e. run_cv)

threads 1       #on multicore systems you can use up to 8 threads to speed up the calculation

### energy binning ###
numBins 142
start -88
step 2

### reaction coordinate 1 binning ###
numBins 50
start 0
step 2

### reaction coordinate 2 binning, change numDimensions to 3 ###
## numBins 50
## start 0
## step 2

### list of histogram filenames and their temperatures ###
numFiles 4 # number of histogram files listed below
name data/hist144 temp 144
name data/hist145 temp 145
name data/hist146 temp 146
```

```
name data/hist147 temp 147
```

- The Boltzmann constant (`kb`) that relates the energy units used in the histogram files to the temperature units given with `temp`. The value shown here is appropriate for Gromacs and SMOG, where the temperature is the Gromacs temperature in Kelvin.
- The total considered range for the energy binning is [`start` , `start + numBins*step`]. Adjust `step` to balance detail (finer `step`) with noise reduction (larger `step`). Also, a larger `step` leaves a smaller memory footprint, and runs faster. So large `step` is useful for preliminary tests.
- The histogram files are designated with relative paths with respect to the directory where `WHAM.jar` is launched. Histogram files are space delimited columns with the following format:

```
energy reaction_coord_1 [ more_reaction_coords ] [ umbrella_coords ]
```

```
-23.779612 117
-37.528976 124
-52.246330 123
-43.539364 132
-41.074669 129
-34.133888 130
-25.081692 125
6.063239 117
51.440716 102
21.739162 114
4.184849 112
37.821884 117
-10.340809 121
```

This file (`hist144`) has an energy column and one reaction coordinate column. The configuration file designates it was run at a temperature of 144.

### 1.5.2 Full configuration parameter list

The following will give a full list of configuration options:

```
user$ java -jar WHAM.jar -help
```

## 1.6 Compute $F(Q)$ , $S(Q)$ , $E(Q)$ at $T$ from $\Omega(E, Q)$ or $\Omega(E, Q_1, Q_2)$

Add the following to compute  $F(Q)$  from  $\Omega(E, Q)$ . You can substitute  $Q$  for your favorite reaction coordinate, and it is always given in the second column

```

run_free          #comment out to not run, reads dosFile

run_free_out free #prefix for the free energy curves

startTF 135      #first temperature to compute F(Q)

deltaTF 0.1      #step in temperature

ntempsF 200     #total F(Q) to generate

```

- There will be `ntempsF` files with names like `$run_free_out$temp`. Output files have four columns in the following format:

```

reaction_coord_value free_energy enthalpy entropy

```

- If two reaction coordinates are given  $\Omega(E, Q_1, Q_2)$ , `run_free` assumes you want two dimensional analysis  $F(Q_1, Q_2)$ .

```

reaction_coord_1_value reaction_coord_2_value free_energy enthalpy entropy

```

## 1.7 Compute $C_V(T)$ , $E(T)$ , $S(T)$ , $F(T)$ from $\Omega(E, \dots)$

Add the following to compute the heat capacity  $C_V(T)$  (also the temperature dependence of energy, entropy, or free energy) from  $\Omega(E, \dots)$ .

```

run_cv           #comment out to not run, reads dosFile

run_cv_out cv    #filename for the temperature curves

startT 135      #starting temperature

deltaT 0.1      #step in temperature

ntemps 200     #total temps to generate

```

- There will be one output file with five columns in the following format:

```

temperature Cv enthalpy free_energy entropy

```

## 1.8 Compute $\langle Q(T) \rangle$ from $\Omega(E, Q)$

Add the following to compute the thermal expectation of a reaction coordinate  $Q_1$  with respect to reaction coordinate  $Q_2$ .

```

run_coord        #comment out to not run, reads dosFile

run_coord_out coord #filename for the coordinate curve

```

```

startTC 110      #temperature to start at
deltaTC 0.1     #step in temperature
ntempsC 200     #total temps to generate

```

- There will be one output file with two columns in the following format:

```
temperature <Q>
```

## 1.9 Compute $\langle Q_2(Q_1) \rangle$ from $\Omega(E, Q_1, Q_2)$

Add the following to compute the thermal expectation of a reaction coordinate  $Q_2$  with respect to reaction coordinate  $Q_1$ .

```

run_coord      #comment out to not run, reads dosFile
run_coord_out coord #filename prefix for the coordinate curve
startTC 110     #temperature to use

```

- There will be one output file `$run_coord_out$startTC` with two columns in the following format:

```
Q1 <Q2(Q1)>
```

## 1.10 Compute $\langle Q_3(Q_1, [Q_2^{\text{low}}, Q_2^{\text{high}}]) \rangle$ from $\Omega(E, Q_1, Q_2, Q_3)$

Add the following to compute the thermal expectation of a reaction coordinate  $Q_3$  with respect to reaction coordinate  $Q_1$  and averaged over a range of  $Q_2$ .

```

run_coord      #comment out to not run, reads dosFile
run_coord_out coord #filename prefix for the coordinate curve
startTC 110     #temperature to use
lowQ2 38       #start of range for averaging third reaction coord
               #note that internally it uses the binning, so bins (not values)
               #greater than lowQ2 are considered
highQ2 42      #end of range for averaging third reaction coord
               #note that internally it uses the binning, so bins (not values)
               #less than highQ2 are considered

```

- There will be one output file `$run_coord_out$startTC` with two columns in the following format:

```
temperature <Q3(Q1, [Q2])>
```

## 1.11 Reweighting of umbrella sampling

The umbrella coordinate is always given as the last columns in the input. `numDimensions` should be equal to the number of columns, which is `numUmbrella+1+number_of_reaction_coords`, where the +1 comes from the energy column. The energies (first column) are assumed to be without the umbrella included.

### 1.11.1 Harmonic umbrella

The potential for a harmonic umbrella is  $V = \frac{k}{2}(Q-Q_0)^2 \rightarrow \Delta E = \text{umbrella\_k} * (\text{column\_3} - \text{umbrella\_0})^2$

```
numDimensions 3      #energy reaction_coord umbrella_coord
numUmbrella 1       #number of umbrella_coord
umbrellaType harmonic #harmonic umbrellas
.
.
.
### list of histogram filenames, temperatures, umbrella stiffness and umbrella centers ###
numFiles 4 # number of histogram files listed below
name data/hist144.40 temp 144 umbrella_k 0.5 umbrella_0 40
name data/hist144.80 temp 144 umbrella_k 0.5 umbrella_0 80
name data/hist146.40 temp 146 umbrella_k 0.5 umbrella_0 40
name data/hist146.80 temp 146 umbrella_k 0.5 umbrella_0 80
```

- Important! The energy column is the energy without the umbrella perturbation included.

## References

- [1] Kumar, S, Rosenberg, J, Bouzida, D, & Swendsen. (1992) The weighted histogram analysis method for free-energy calculations on biomolecules. I. The method. *J. Comput. Chem.* **13**, 1011.
- [2] Noel, J. K, Levi, M, Raghunathan, M, Lammert, H, Hayes, R. L, Onuchic, J. N, & Whitford, P. C. (2016) SMOG 2: A Versatile Software Package for Generating Structure-Based Models. *PLoS computational biology* **12**, e1004794.