Manual for slepton pair plus jet production in the POWHEG BOX V2

This document describes the settings and input parameters that are specific to the implementation of slepton pair production in association with a jet at the LHC within the POWHEG BOX V2 framework. The parameters that are common to all POWHEG BOX V2 implementations are given in the documents manual-BOX.pdf and Manyseeds.pdf in the POWHEG-BOX-V2/Docs directory. If you use our program, please quote Refs. [1-4].

Model parameters

Note that the current implementation has been tailored to left-handed slepton pair production in association with a jet. Right-handed sleptons are not yet supported. The masses of squarks that appear as virtual particles in the loop corrections are implicitly assumed to be degenerate. Mixing effects between squarks of different generations are entirely disregarded. Please contact the authors if you want to apply the code to scenarios that do not meet these limitations. Model parameters are read from the file param_card.dat in the subdirectory MODEL which is formated according to the SUSY Les Houches Accord (SLHA) [5,6] and can be taken from standard SUSY spectrum generators.

The treatment of the virtual corrections is specified by the flag qcdonly in the powheg.input file. If the switch is enabled, only Standard-Model type virtual corrections with a Drell-Yan-type structure are computed. Contributions with supersymmetric loop corrections and non-Drell-Yan type diagrams are neglected. While the numerical impact of the neglected terms is very small for current SUSY mass limits, the execution time of the code is halved when this switch is enabled.

Download and running of the code

Download the POWHEG BOX V2 following the instructions at the website

http://powhegbox.mib.infn.it/

and subsequently retrieve the subdirectory **dislepton-jet** into the directory **POWHEG-BOX-V2**. Before compiling the code make sure that:

- fastjet is installed and fastjet-config is in the path,
- lhapdf is installed and lhapdf-config is in the path,
- gfortran or ifort is in the path, and the appropriate libraries are in the environment variable LD_LIBRARY_PATH.

Go to the project directory

\$ cd POWHEG-BOX-V2/dislepton-jet

and compile the code with

\$ make

which may take some time. All three generated programs pwhg_main, lhef_analysis and main-PYTHIA-lhef are to be executed in a separate working directory. This directory has to contain a file powheg.input that specifies several technical parameters and, for parallel running, a file pwgseeds.dat with different seeds for each parallel computation. Samples of these files are provided in the subdirectory runs. For your own runs, create a subdirectory, e.g. named testrun, in dislepton-jet:

\$ mkdir testrun

enter the testrun directory

\$ cd testrun

and perform all your runs there. We recommend running the program in a parallel mode in several consecutive steps, with the following common settings in powheg.input:

```
foldcsi 2
foldy 2
foldphi 2
manyseeds 1
```

When executing

\$../pwhg_main

the program will ask you to

enter which seed

The program requires you to enter an index that specifies the line number in the pwgseeds.dat file where the seed of the random number generator to be used for the run is stored. All results generated by the run will be stored in files named *-[index].*. When running on parallel CPUs, make sure that each parallel run has a different index. The timings given in the following refer to the program compiled with gfortran and run on Intel Xeon E5-2670 CPUs with 2.6 GHz and 32 GB RAM each.

Step 1

In this first step, the importance sampling grids are generated. To speed up this step, set

fakevirt 1

in order to use fake virtuals that are proportional to the Born rather than the full virtual contributions for the preparation of the importance sampling grids. Make sure that the relevant technical parameters in powheg.input are set to the following values:

```
xgriditeration 1 parallelstage 1
```

For precision results, we recommend running about 100 parallel jobs with the number of calls set by

ncall1 200000

for each. In order to run the parallel jobs, use

\$../pwhg_main

When prompted

enter which seed

enter an index for each run (from 1 to 100). The pwgseeds.dat must contain at least 100 lines, each with a different seed.

The completion of the first iteration of the grid production takes roughly one hour of CPU per job. Once all jobs of the first iteration are completed, the grids are refined in further iterations. We recommend to perform at least a second iteration, setting

```
xgriditeration 2 parallelstage 1
```

and rerunning the program as in the first iteration. If more iterations are needed, the value of **xgriditeration** has to be adapted accordingly. Each iteration takes roughly the same amount of CPU time as the first one.

Step 2

In the directory where the previously generated grids are located, the generation of the NLO results can be performed, again in the parallel mode. Make sure to enable the full virtual amplitudes by deactivating the fakevirt switch. Set

```
ncal12 300000
xgriditeration 1
parallelstage 2
```

in powheg.input and run jobs in parallel, in the same way as explained for step 1 above. Upon the completion of step 2 (after about eight hours), for each parallel run with number [index] a file pwg-[index]-NLO.top is generated. These files contain the histograms defined in pwhg_analysis.F at NLO-QCD accuracy, if the variable bornonly is set to zero in powheg.input. Setting bornonly to 1 yields the respective LO results. In either case, the individual results of the parallel runs can be combined, for instance with the help of auxiliary files contained in the plot-aux directory of the POWHEG BOX V2 implementation VBF_Z_Z as described in the corresponding manual.

Step 3

Also this step can be run in parallel. With the settings

```
nubound 100000 parallelstage 3
```

each jobs takes roughly five minutes. The parallel execution of the program is performed as in the previous step.

Step 4

Set numerts to the number of events you want to generate per job, for example

numevts 100000

and run in parallel with

parallelstage 4

Generating the specified number of events takes about one day. At this point, files of the form pwgevents-[index].lhe are present in the run directory. The events can be merged into a single event file pwgevents.lhe by

```
$ cat pwgevents-*.lhe | grep -v "/LesHouchesEvents" > pwgevents.lhe
$ echo "</LesHouchesEvents>" >> pwgevents.lhe
```

Analyzing the events

It is straightforward to feed the combined event file pwgevents.lhe into a generic shower Monte Carlo program, within the analysis framework of each experiment. We also provide a sample analysis that computes several histograms and stores them in gnuplot output files. Doing

\$../lhef_analysis

analyzes the bare POWHEG BOX output and writes histograms to the file pwgLHEF_analysis.top. Parton showers, slepton decays, and further effects can be simulated when the events are handed over to PYTHIA [7] by

\$../main-PYTHIA-lhef

The corresponding histograms can be found in pwgPOWHEG+PYTHIA-output.top.

Various settings of PYTHIA can be modified by editing the file setup-PYTHIA-lhef.F prior to compilation. We modified PYTHIA in such a way that by setting the flag mstp(41)=1 slepton decays are activated. PYTHIA generates the decays according to the masses provided in param_card.dat. For the case of each slepton decaying into a lepton and the lightest neutralino, our sample analysis provides various histograms for the decay products.

References

- [1] B. Jäger, A. von Manteuffel, and S. Thier, Slepton pair production in association with a jet: NLO-QCD corrections and parton-shower effects, arXiv:1410.3802.
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- [7] T. Sjostrand, S. Mrenna, P. Z. Skands, PYTHIA 6.4 Physics and Manual, JHEP 0605 (2006) 026 [hep-ph/0603175].