

# Manual for electroweak $W^+W^+jj$ production in the POWHEG-BOX-V2

The `VBF_Wp_Wp` program is an implementation of the electroweak  $W^+W^+jj$  production cross section within the POWHEG BOX framework. It complements the `Wp_Wp-j-j` program, which provides the QCD-induced contributions to  $W^+W^+jj$  production in hadronic collisions.

This document describes the input parameters that are specific to the implementation of the EW channel in the POWHEG-BOX-V2. Note that the recommended settings are very similar to those for the QCD production mode. The parameters that are common to all POWHEG-BOX-V2 implementations are described in the POWHEG-BOX-V2/Docs directory.

If you use this program, please quote Refs. [1–3].

## Running the program

Download the POWHEG-BOX-V2, following the instructions at the web site

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

In addition, retrieve the `VBF_Wp_Wp` directory as explained, and go to the new directory via

```
$ cd POWHEG-BOX-V2/User-Processes-V2/VBF_Wp_Wp
```

Running is most conveniently done in a separate directory, for instance do

```
$ mkdir testrun
```

The directory must contain the `powheg.input` file and, for parallel running, a `pwgseeds.dat` file (see `manual-BOX.pdf` and `Manyseeds.pdf`).

Before compiling make sure that:

- `fastjet` is installed and `fastjet-config` is in the path,
- `lhapdf` is installed and `lhapdf-config` is in the path,
- `gfortran`, `ifort` or `g77` is in the path, and the appropriate libraries are in the environment variable `LD_LIBRARY_PATH`.

If `LHAPDF` or `fastjet` are not installed, the code can still be run using a dummy analysis routine and built-in PDFs, see the `Makefile` in `VBF_Wp_Wp`.

After compiling, enter the `testrun` directory:

```
$ cd testrun
```

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 program can be run in several steps. Each new step requires the completion of the previous step.

## Step 1

At this point the user has to decide whether the weak bosons are to be generated on-shell (`zerowidth = 1`) or off-shell, distributed according to a Breit-Wigner distribution (`zerowidth = 0`). By default on-shell weak bosons are generated. The user can also select the decay modes of the weak bosons by assigning appropriate PDG code to the parameters `vdecaymodew1` and `vdecaymodew2` in `powheg.input`. Note that only leptonic decays are supported. The parameters `vdecaymodew1` and `vdecaymodew2` are to be set to -11, -13 or -15 for decays to positrons, anti-muons, or anti-taus, respectively. It is also possible to generate events with combinations of positron and anti-muon decays, in which case `vdecaymodew1` and `vdecaymodew2` at the same time should be set to -113. A combination of all leptons (positron, anti-muons, and anti-taus) can be obtained by setting `vdecaymodew1` and `vdecaymodew2` to -135. The template analysis file `pwhg_analysis.f` needed in subsequent steps of the analysis is designed for the  $e^+\nu_e\mu^+\nu_\mu$  mode. In order to obtain predictions for  $pp \rightarrow W^-W^-jj$ , one can do a calculation for  $\bar{p}\bar{p} \rightarrow W^+W^+jj$ , treat the decay leptons of the  $W^+$  bosons as if they were negatively charged and reverse the momentum directions for parity-odd distributions. In the POWHEG BOX, predictions for anti-protons can be obtained by setting the input parameters `ih1` and `ih2` to -1 in the `powheg.input` file.

We recommend to generate the grid with the option `fakevirt 1` in `powheg.input`. When using this option, the virtual contribution is replaced by a fake one proportional to the Born term. This speeds up the generation of the grid.

In the POWHEG-BOX-V2 grids can be generated in the parallel mode. To that end, set

```
manyseeds 1
xgriditeration 1
parallelstage 1
```

For a default setup one needs about 50–100 jobs with the number of calls set by

```
ncall1 10000
```

for each. Run the program via

```
$../pwhg_main
```

When prompted

```
enter which seed
```

enter 1 or any other valid seed number.

The program stops automatically after the compilation of this step. We recommend performing at least 3 iterations. After completion of the first iteration, change the value of `xgriditeration` according to the respective iteration, and re-do step 1.

## Step 2

To produce true NLO results comment out the `fakevirt` token from `powheg.input`. The runs must be performed in the directory where the previously generated grids are stored.

The integration and upper bound for the generation of `btilde` can be performed with 50-100 runs with 5000-10000 calls each. Set for instance

```
ncall2 5000
itmx2 1
```

in `powheg.input`.

Folding numbers that are appropriate for runs at LHC energy are:

```
foldcsi 5 ! number of folds on csi integration
foldy 5 ! number of folds on y integration
foldphi 10 ! number of folds on phi integration
```

In addition, adapt the values for steering the parallel mode of the program to

```
xgriditeration 1
parallelstage 2
```

Run jobs in parallel, in the same way as explained for step 1 above.

Upon the completion of this step, for each parallel run a file `pwg-*-NLO.top` is generated (where the `*` denotes the integer identifier of the run). These files contain the histograms defined in `pwg_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 with the help of the `mergedata.f` file contained in the `plot-aux` directory. To this end, just compile the file by typing, e.g.,

```
$ gfortran -o mergedata mergedata.f
```

and run the resulting executable. in your run directory `mergedata 1 *NLO.top`. The program expects a number between 1 and 5 and a list of files to merge. If no number or no list is specified, the user will be prompted by the program to enter them. Running `mergedata` without any arguments will also explain what the 5 different options are. `mergedata` will combine the histograms into a file called `fort.12`. We provide a file `plot-aux/genplots.sh` and `plot-aux/gnuplotsplit.gp` which can be used to plot the resulting histograms. First run `genplots.sh`

```
./genplots.sh file nameoutput
```

or if a comparison between two different runs is wanted

```
./genplots.sh file1 file2 nameoutput
```

This will result in a file called `genplots.gp`. In the later case the file `plot-aux/pastegnudata.f` has to be compiled and either put in the working directory or in the users path. After `genplots.sh` has been run, the user can produce a set of `.eps` files with `gnuplot` running

```
./gnuplotsplit.gp genplots.gp
```

### Step 3

Also this step can be run in parallel. We recommend to set

```
ncall1 500000  
itmx1 4  
ncall2 500000  
itmx2 4  
nubound 100000  
parallelstage 3
```

The program will stop after completion of this step. The parallel execution of the program is performed as in the previous steps.

### Step 4

Set `numevts` to the number of events you want to generate per process, for example

```
numevts 5000
```

and run in parallel.

At this point, files of the form `pwgevents-[index].lhe` are present in the run directory.

Count the events:

```
$ grep '/event' pwgevents-*.lhe | wc
```

The events can be merged into a single event file by

```
cat pwgevents-*.lhe | grep -v '/LesHouchesEvents' > pwgevents.lhe
```

## Analyzing the events

It is straightforward to feed the `*.lhe` events 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 `topdrawer` output files.

Doing (from the `VBF_Wp_Wp` directory)

```
$ make lhef_analysis
```

```
$ cd testrun
```

```
../lhef_analysis
```

analyses the bare POWHEG BOX output, creating the topdrawer file `LHEF_analysis.top`. The targets `main-HERWIG-lhef` and `main-PYTHIA-lhef` are instead used to perform the analysis on events fully showered using HERWIG or PYTHIA. Various setting of the Monte Carlo can be modified by editing the files `setup-PYTHIA-lhef.f` and `setup-HERWIG-lhef.f` respectively.

## References

- [1] B. Jäger, C. Oleari, D. Zeppenfeld, *Next-to-leading order QCD corrections to  $W^+W^+jj$  and  $W^-W^-jj$  production via weak-boson fusion*, Phys. Rev. **D80** (2009) 034022. [arXiv:0907.0580 [hep-ph]].
- [2] B. Jäger, G. Zanderighi, *NLO corrections to electroweak and QCD production of  $W^+W^+$  plus two jets in the POWHEG BOX*, JHEP **1111** (2011) 055. [arXiv:1108.0864 [hep-ph]].
- [3] S. Alioli, P. Nason, C. Oleari and E. Re, *A general framework for implementing NLO calculations in shower Monte Carlo programs: the POWHEG BOX*, JHEP **1006** (2010) 043. [arXiv:1002.2581 [hep-ph]].