

## Running CYMA2DV with gfortran compiler on Linux system

- Download the package source file CYMA2DV-2025-10-31.tgz from [MEDOC](#) website :
- (**gfortran** compiler is required)
- Unpack the package by typing the following linux command :  
**tar -xvzf CYMA2DV-2025-10-31.tgz**
- Go to the folder CYMA2DV :  
**cd CYMA2DV**
- The folder contains the following files: intinc.dat, tembri.dat, paramod.dat, makefile, cyma2dv.f90, visu.f90
- The file to modify is ``paramod.dat"
- Run the code by typing :  
**make**  
**./cyma2dv**
- The output files are: cc2dhyv.log (iteration message), resu2dv.dat (emergent intensities for plotting line profiles), popc2dv.dat (hydrogen level populations), fort.8 (CPU time)
- The folder **results** contains the output files corresponding to a test case to be able to check if your results are good
- To visualize results, use the following program visu.f90 by typing :  
**gfortran -o visu visu.f90**  
**./visu**  
The output file is **cosbovi.ps**

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