bash command

```
$ mpirun -np NUM_PROCESSORS MMPBSA.py.MPI -O -i INPUT_FILE.in -o
  ↪ gb_results.dat -sp SOLVATED_COMPLEX.parm7 -cp complex.parm7
  ↪ -rp receptor.parm7 -lp ligand.parm7 -y
  ↪ PRODUCTION_TRAJECTORY.nc
```