

**NAME**

mkopr - make an ocean project

**SYNOPSIS**

**mkopr** project

**PARAMETERS**

*project* Absolute or relative path of the project to be created.

**DESCRIPTION**

*Mkopr* is a symbolic link to the ICELLES(1ICD) script. The *ICELLES* script is a front-end to the real *mkopr* script. This script is to be sourced by the *ICELLES* script. It creates a NELSI project directory for the OCEANPROCESS (default "fishbone") and imports the libraries "primitives", "digilib", "analib" and "bonding" into that project. The libraries must first be installed with the script ICELLES(1ICD).

**ENVIRONMENT VARS****ICDPATH**

The full path to the place where the Ocean & Nelsis tree lives. This environment variable must be set!

**OCEANPROCESS**

The name of the ocean process and image to use, e.g. "fishbone", "gatearray" or "octagon". The default process is "fishbone".

**BINLIBS**

Is set to ICDPATH/share/lib/celllibs/OCEANPROCESS. This variable is used by the real script and is not an environment variable. It is the path to the directory containing the binary version of the libraries.

**SRCLIBS**

Is set to ICDPATH/share/lib/celllibs/OCEANPROCESS/src. This variable is used by the real script and is not an environment variable. It is the path to the directory containing the ascii version of the libraries, as well as the template directory.

**EXAMPLES**

```

% mkopr myproject
----- creating fishbone project myproject -----
----- importing primitives -----
----- importing digilib8_93 -----
----- importing analib8_00 -----
----- importing bonding11_93 -----
----- copying default config files -----

...done. Enjoy your new project "myproject" !!!
% cd myproject

```

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**FILES**

```

ICDPATH/share/lib/process/processlist
SRCLIBS/mkopr      # the ocean process real mkopr script
SRCLIBS/template/* # place of default config files
BINLIBS/*         # place of imported libraries

```

**SEE ALSO**

ICELLES(1ICD), mkopr(1ICD), mkepr(1ICD), mkropr(1ICD), mkvopr(1ICD).