

NAME

highlay - high-light the layout of conductors and/or devices

SYNOPSIS

highlay [-npeFvmidL -l mask -w width -c outcell] [-N n1,n2... -P p1,p2... -E e1,e2...] cell [namefile]

OPTIONS

- n** Select nets.
- p** Select ports.
- e** Select devices (fets and bipolar transistors).
- L** High-light with the conductor masks (default: all original masks that are present at that position).
- l mask** Use specified mask for high-lighting (default: all original masks that are present at that position).
- w width**
The width of an area with which a selected conductor area should be extended (can only be used in combination with the option -l, default is 0).
- c outcell**
Name of output cell (default: HIGH_OUT).
- o** Do not include the original cell as a sub-cell.
- F** Include the original cell at the same level.
- v** Verbose mode. Print the nets, ports and devices that are selected.

The following options are only appropriate when using match results (no namefile is specified).

- m** Select matching items.
- i** Select inconclusive items.
- d** Select deficient items.
- N n1,n2...**
Select the nets that are part of groups *n1,n2...*
- P p1,p2...**
Select the ports that are part of groups *p1,p2...*
- E e1,e2...**
Select the devices that are part of groups *e1,e2...*

DESCRIPTION

Highlay(IICD) is a program for high-lighting one or more (extracted) conductors and/or devices in a layout-editor/viewer. In order to use *highlay* for a particular cell, first run the layout-to-circuit extractor *space(IICD)* for this cell using the option **-x**.

The conductors (nets/ports) and devices that are high-lighted can be specified in two different ways: (1) they are specified in a namefile, or (2) they are selected from the output of the circuit comparison program *match(IICD)*.

When a namefile is specified, the names of the nets, ports and/or devices that are high-lighted are read from this file. The names must be separated by tabs, spaces and/or newline characters. For the devices, the database names must be used. These are the names that are obtained when *xsls(IICD)* or *xspice(ICD)* is used with the option **-d**, but without the prefix character. Nets, ports and/or devices are selected for high-lighting according to the use of the options **-n**, **-p** and/or **-e**.

When no namefile is specified, *highlay* will read output of the program *match* for the specified cell. In that case, it will select nets, ports and/or devices according to the use of the options **-m**, **-i**, **-d**, **-n**, **-N**, **-p**, **-P**, **-e** and/or **-E**. A net, port or device is high-lighted if and only if it is selected by one of the options **-m**, **-i** or **-d** and by one of the options **-n**, **-N**, **-p**, **-P**, **-e** or **-E**. In order to obtain correct input for *highlay* from *match*, the latter program should be run using the option **-edif**.

Highlay will generate a new cell that contains (default) the high-lighted parts of the specified cell, plus a copy of the specified cell as a sub-cell. The name of the new cell that is generated is (default) HIGH_OUT. This name can be changed by using the option **-c**. The nets, ports and/or devices that are selected for high-lighting are (default) displayed in their original mask colors. An alternative mask for high-lighting can be specified by using the option **-l**.

NOTE

Note that names that are printed in spice descriptions as in_1_ in_2_, might be known in the layout database - and must therefore be referred to in the namefile of *highlay* - as in[1] in[2].

EXAMPLE

An example of a namefile "latch.hil" for *highlay* is as follows:

```
vdd
a1
a2
_T2
```

This namefile may be used to obtain a cell HIGH_OUT in which the nets vdd, a1 and a2 and the device _T1 of cell latch are high-lighted, as follows.

```
$ highlay -n -e latch latch.hil
```

EXAMPLE2

Name wildcards can also be used in the namefile. For example to select all names:

```
*
```

For example to select names "grp1" to "grp12":

```
grp*
```

For example to select only names "grp11" and "grp12":

```
grp??
```

Note that only one '*' may be used in a name. Characters behind the '*' are not supported.

AUTHOR

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SEE ALSO

match(IICD), *space(IICD)*.