

NAME

`xlsatoms` - list interned atoms defined on server

SYNOPSIS

xlsatoms [-options ...]

DESCRIPTION

Xlsatoms lists the interned atoms. By default, all atoms starting from 1 (the lowest atom value defined by the protocol) are listed until unknown atom is found. If an explicit range is given, *xlsatoms* will try all atoms in the range, regardless of whether or not any are undefined.

OPTIONS

-display *dpy*

This option specifies the X server to which to connect.

-format *string*

This option specifies a *printf*-style string used to list each atom *<value,name>* pair, printed in that order (*value* is an *unsigned long* and *name* is a *char **). *Xlsatoms* will supply a newline at the end of each line. The default is *%lu\t%s*.

-range [*low*]-[*high*]

This option specifies the range of atom values to check. If *low* is not given, a value of 1 assumed. If *high* is not given, *xlsatoms* will stop at the first undefined atom at or above *low*.

-name *string*

This option specifies the name of an atom to list. If the atom does not exist, a message will be printed on the standard error.

-version Print out the program version and exit.

SEE ALSO

X(7), Xserver(1), xprop(1)

ENVIRONMENT**DISPLAY**

to get the default host and display to use.

AUTHOR

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