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 *%ld\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

Jim Fulton, MIT X Consortium