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# Rocky Enterprise Linux 9.2 Manual Pages on command 'xlsatoms.1'

## \$ man xlsatoms.1

XLSATOMS(1)

**General Commands Manual** 

XLSATOMS(1)

NAME

xlsatoms - list interned atoms defined on server

### **SYNOPSIS**

xlsatoms [-options ...]

### **DESCRIPTION**

XIsatoms lists the interned atoms. By default, all atoms starting from 1 (the lowest atom value defined by the protocol) are listed until un? known atom is found. If an explicit range is given, xIsatoms 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 un? signed long and name is a char \*). XIsatoms will supply a new? line at the end of each line. The default is %Id\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 er? ror.

#### -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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X Version 11 xlsatoms 1.1.2 XLSATOMS(1)