NAME

    uniq – report repeated lines in a file

SYNOPSIS

    uniq [**−udc** [+n] [−n]] [input [output]]

DESCRIPTION

    Uniq reads the input file comparing adjacent lines. In the normal case, the second and succeeding copies of repeated lines are removed; the remainder is written on the output file. Note that repeated lines must be adjacent in order to be found; see *sort*(I). If the **−u** flag is used, just the lines that are not repeated in the original file are output. The **−d** option specifies that one copy of just the repeated lines is to be written. The normal mode output is the union of the **−u** and **−d** mode outputs.

    The **−c** option supersedes **−u** and **−d** and generates an output report in default style but with each line preceded by a count of the number of times it occurred.

    The *n* arguments specify skipping an initial portion of each line in the comparison:

        −*n*   The first *n* fields together with any blanks before each are ignored. A field is defined as a string of non-space, non-tab characters separated by tabs and spaces from its neighbors.

        +*n*   The first *n* characters are ignored. Fields are skipped before characters.

SEE ALSO

    sort(I), comm(I)