A data cleaner's cookbook - Windows carriage returns

On Linux and Mac machines, a newline is built with just one character, the UNIX linefeed '\n' ('LF'). On Windows computers, a newline is created using two characters, one after the other: '\r\n' ('CRLF'), where '\r' is called a 'carriage return' ('CR'). Carriage returns aren't necessary in a data table and can cause problems in data cleaning. (For examples of problems, see this BASHing data post.)

There are several ways to find CR characters. You can use **sed -n 'l'** to visualise any '\r' in a table, and **grep** to select out the lines with a CR and print their line numbers. Alternatively, a CR character will be shown as '^M' if you use **cat -v**, where the '-v' option shows non-printing characters other than tabs and linefeeds. In the example below, the file **winCR** has an invisible Windows carriage return at the end of the first line:

```
$ cat winCR
aaa bbb
ccc ddd
eee fff

$ sed -n 'l' winCR
aaa\tbbb\r$
ccc\tddd$
eee\tfff$

$ sed -n 'l' winCR | grep -n "\\r"
1:aaa\tbbb\r$

$ cat -v winCR | grep -n "\^M"
1:aaa bbb^M$
```

It's wise to run these commands with **grep**'s '-c' option first rather than '-n'. The '-c' option returns only the *number* of lines with a CR, and if that number is big, you avoid having large number of lines printed at high speed in your terminal. If your **grep** supports Perl-type regular expressions, you can count '\r' characters directly.

```
$ sed -n 'l' winCR | grep -c "\\r"

$ cat -v winCR | grep -c "\^M"

$ grep -cP "\r" winCR
```

Another command to find carriage returns is **file**, which will report on line endings if they're different from a single linefeed, but won't count them:

```
$ file example.csv
example.csv: ASCII text, with CRLF line terminators
```

The easiest way to remove all Windows carriage returns from **table** is with **tr**:

```
$ tr -d '\r'  table_without_CR
```

Deleting all the carriage returns could be a mistake, however, if any of them are within data items.

1 of 2

The screenshot below shows a real-world example. In the file **afd1**, I used **sed** to replace each of the 2 carriage returns in line 67893 with a single whitespace. Note that this was an 'in-place' edit with **sed**'s '-i' option.

```
grep -n "\^M"
 cat -v afd1
                safrina Spilopyra safrina Reid & Beatson, 2010C
HRYSOMELIDAE
                Spilopyra
                                         safrina
                                                         Synonym
                                         2010
synonym Species
                        Reid & Beatson
                                                         2d32135
3-4268-467d-8335-f5802ece5a50
                                 20110419T01:26:11.999+0000
99f0a0b-3651-4174-a87f-ca286902ea0a
                                         20120323T01:38:19.026+0
                4f91518a-ae5e-4912-9505-3cd5f08557ce
                                                         Reid, C
000
.A.M. & Beatson, M.
                        2010
                                Revision of the Australo-Papuan
 genus <i>Spilopyra</i> Baly^M(Coleoptera: Chrysomelidae: Spilo
pyrinae)
                                Zootaxa
!--MARK-->Reid, C.A.M. & Beatson, M. 2010.
                                           Revision of the Aust
ralo-Papuan genus <i>Spilopyra</i> Baly^M(C)leoptera: Chrysomel
idae: Spilopyrinae). <!--MARK--><!--MARK--><em>Zootaxa</em> <st
rong>2692</strong>: 1-32<!--MARK-->
                                                 Article in Jour
        999deae1-59ee-41db-9159-3d43c22206af
                                                 20110419T01:03:
41.080+0000
                96466227-221c-4922-ac97-1ac2dd946d0a
 sed -i '67893s/\r/ /g' afd1
 sed -n '67893p' afd1
                safrina Spilopyra safrina Reid & Beatson, 2010C
HRYSOMELIDAE
                                         safrina
                Spilopyra
                                                         Synonym
                        Reid & Beatson
synonym Species
                                         2010
                                                         2d32135
3-4268-467d-8335-f5802ece5a50
                                 20110419T01:26:11.999+0000
99f0a0b-3651-4174-a87f-ca286902ea0a
                                         20120323T01:38:19.026+0
000
                4f91518a-ae5e-4912-9505-3cd5f08557ce
                                                         Reid. C
.A.M. & Beatson, M.
                        2010
                                Revision of the Australo-Papuan
 genus <i>Spilopyra</i> Baly (Coleoptera: Chrysomelidae: Spilop
yrinae) 1-32
                        Zootaxa
K-->Reid, C.A.M. & Beatson, M. 2010.
                                     Revision of the Australo-P
apuan genus <i>Spilopyra</i> Baly (Coleoptera: Chrysomelidae: S
pilopyrinae). <!--MARK--><!--MA
                                RK--><em>Zootaxa</em> <strong>26
92</strong>: 1-32<!--MARK-->
                                         Article in Journal
99deae1-59ee-41db-9159-3d43c22206af
                                         20110419T01:03:41.080+0
000
        96466227-221c-4922-ac97-1ac2dd946d0a
```

2 of 2 3/19/2019, 12:47 PM