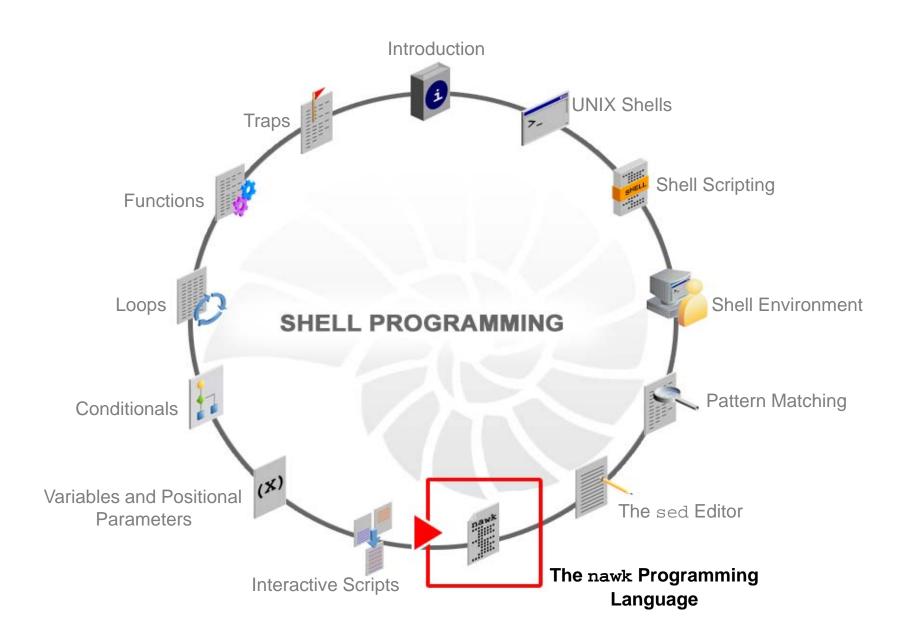
# The nawk Programming Language



# **Objectives**

After completing this lesson, you should be able to:

- Describe the capabilities of the nawk programming language
- Display output by using the print statement
- Perform pattern matching by using regular expressions
- Use the nawk built-in and user-defined variables

# **Agenda**

- Describing the capabilities of the nawk programming language
- Displaying output by using the print statement
- Performing pattern matching by using regular expressions
- Using the nawk built-in and user-defined variables

# awk Programming Language

- awk is named for its authors Aho, Weinberger, and Kernighan of AT&T Bell Labs.
- The awk programming language is:
  - A record-oriented language
  - A text processing, data extraction, and reporting tool
  - A language executed by the awk interpreter
- The language has three variations:
  - awk is the original and oldest version.
  - nawk (New awk) is the improved version adapted by most UNIX vendors.
  - gawk is the free GNU version.

# nawk Programming Language

#### The nawk programming language:

- Looks at data by records and fields
- Uses regular expressions
- Uses numeric and text variables and functions
- Uses command-line arguments

#### nawk Capabilities

Applications written in the nawk programming language provide the following capabilities:

- Filtering
- Numerical processing on rows and columns of data
- Text processing to perform repetitive editing tasks
- Report generation

#### nawk Command Format

Commands have the form:

```
nawk 'statement' input.file
```

Scripts are executed with:

```
nawk -f scriptfile input.file
```

# **Agenda**

- Describing the capabilities of the nawk programming language
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# **Printing Selected Fields**

- The print statement outputs data from the file.
- Command conventions:
  - Enclose the command in single quotation marks.
  - Enclose the command in braces { }.
  - Specify individual fields with \$1, \$2, \$3, and so on.
  - Specify individual records with \$0.

# **Printing Selected Fields**

```
$ cat data.file
northwest
                Joel Craiq 3.0.98
         NW
                Sharon Kelly 5.3.97
                                                  23
western WE
southwest SW
                Chris Foster 2.7.8
                                                  18
                May Chin 5.1 .95
                                                  15
southern SO
                Derek Johnson 5.0 .70
                                           4
southeast SE
                                                  17
eastern EA
                Susan Beal 4.4.8
                                                  20
northeast NE
                TJ Nichols 5.1 .94
                                                  13
                Val Shultz 4.5 .89
north NO
central CT
                Sheri Watson 5.7.94
                                                  13
$ nawk '{ print $3, $4, $2 }' data.file
Joel Craiq NW
Sharon Kelly WE
Chris Foster SW
May Chin SO
Derek Johnson SE
Susan Beal EA
TJ Nichols NE
Val Shultz NO
Sheri Watson CT
```

# Formatting with the print Statement

- Anything in double quotation marks is a string constant that can be used:
  - In print statements
  - As values to be assigned to variables among other things
- You can also use some special formatting characters:

Character	Octal Value	Meaning
\t	\011	Tab
\n	\012	Newline
	\007	Bell
	\042	"
	\044	\$
	\045	%

# Formatting with the print Statement

```
$ nawk '{ print $3, $4 "\t" $2 }' data.file
Joel Craig
                      NW
Sharon Kelly
                      WE
Chris Foster
                      SW
May Chin
                      SO
Derek Johnson
                      SE
Susan Beal
                      EΑ
TJ Nichols
                      NE
Val Shultz
                      NO
Sheri Watson
                      CT
```

# **Agenda**

- Describing the capabilities of the nawk programming language
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# **Using Regular Expressions**

Regular expression metacharacters can be used in the pattern.

```
$ nawk '/east/' data.file
southeast SE Derek Johnson 5.0 .70
                                                 17
eastern EA Susan Beal 4.4.8 5
                                                 20
northeast NE TJ Nichols 5.1 .94
                                                 13
$ nawk '/east/ { print $1, $5, $4 }' data.file
southeast 5.0 Johnson
eastern 4.4 Beal
northeast 5.1 Nichols
$ nawk '/east/ { print $1, $5 "\t" $4 }' data.file
southeast 5.0 Johnson
eastern 4.4 Beal
northeast 5.1 Nichols
$ nawk '/^east/' data.file
         EA Susan Beal 4.4.8
                                                 20
eastern
```

# **Using Regular Expressions**

\$ nawk '/.	9/' da	ta.file				
northwest	NW	Joel Craig	3.0 .98	3	4	
western	WE	Sharon Kelly	5.3 .97	5	2	
southern	SO	May Chin	5.1 .95	4	1	
northeast	NE	TJ Nichols	5.1 .94	3	1	
north	NO	Val Shultz	4.5 .89	5	9	
central	CT	Sheri Watson	5.7 .94	5	1	
\$ nawk '/\.9/' data.file						
northwest	NW	Joel Craig	3.0 .98	3	4	
western	WE	Sharon Kelly	5.3 .97	5	2	
southern	SO	May Chin	5.1 .95	4	1	
northeast	NE	TJ Nichols	5.1 .94	3	1	
central	CT	Sheri Watson	5.7 .94	5	1	

# **The Special Patterns**

There are two special patterns that are not used to match text in a file.

- BEGIN: An action to take before reading any lines
- END: An action to take after all lines are read and processed

# **The Special Patterns**

```
$ nawk 'BEGIN { print "Eastern Regions\n" }; /east/ {
print $5, $4 }' data.file
Eastern Regions
5.0 Johnson
4.4 Beal
5.1 Nichol
$ nawk 'BEGIN {
> print "Eastern Regions\n"}; /east/ {print $5, $4}'
data.file
Eastern Regions
5.0 Johnson
4.4 Beal
5.1 Nichols
```

# **The Special Patterns**

```
$ nawk 'BEGIN { print "Eastern Regions\n"}; /east/ {print
$5, $4}
> END {print "Eastern Region Monthly Report"}' data.file
Eastern Regions

5.0 Johnson
4.4 Beal
5.1 Nichols
Eastern Region Monthly Report
```

# Using nawk Scripts

- A nawk script is a collection of nawk statements (patterns and actions) stored in a text file.
- To instruct nawk to read the script file, use the command:

```
nawk -f script_file data_file
```

# Using nawk Scripts

```
$ cat report.nawk
BEGIN {print "Eastern Regions\n"}
/east/ {print $5, $4}
END {print "Eastern Region Monthly Report"}

$ nawk -f report.nawk data.file
Eastern Regions

5.0 Johnson
4.4 Beal
5.1 Nichols
Eastern Region Monthly Report
```

# Using nawk Scripts

```
$ cat report2.nawk
BEGIN {print "** Acme Enterprises **"}
BEGIN {print "Eastern Regions\n"}
/east/ {print $5, $4}
END {print "Eastern Region Monthly Report"}
$ nawk -f report2.nawk data.file
** Acme Enterprises **
Eastern Regions
5.0 Johnson
4.4 Beal
5.1 Nichols
Eastern Region Monthly Report
```

# **Agenda**

- Describing the capabilities of the nawk programming language
- Displaying output by using the print statement
- Performing pattern matching by using regular expressions
- Using the nawk built-in and user-defined variables

#### **Built-in Variables**

- As nawk processes an input file, it uses several variables.
- You can provide a value to some of these variables, whereas other variables are set by nawk and cannot be changed.
- A variable value can be a number, a string, or a set of values in an array.

Name	<b>Default Value</b>	Description
FS	Space or tab	The input field separator
OFS	Space	The output field separator
NR		The number of records from the beginning of the first input file

# Input Field Separator (FS)

- The default input field separator (FS) is white space, which can be either a space or a tab.
- Frequently, other characters can separate the input, such as a colon or comma.
- You can set the input field separator variable with the -F
   option or set the value with an assignment.

```
nawk -F: 'statement' filename
nawk 'BEGIN { FS=":" } ; statement' filename
```

# Input Field Separator (FS): Example

```
$ nawk 'BEGIN { FS=":" }; { print $1, $3 }' /etc/group
root 0
other 1
bin 2
sys 3
adm 4
uucp 5
mail 6
tty 7
lp 8
nuucp 9
staff 10
daemon 12
sysadmin 14
nobody 60001
noaccess 60002
nogroup 65534
```

# Input Field Separator (FS): Example

```
$ cat report3.nawk
BEGIN { FS=":" }
{ print $1, $3 }
$ nawk -f report3.nawk /etc/group
root 0
other 1
bin 2
sys 3
adm 4
uucp 5
mail 6
tty 7
lp 8
nuucp 9
staff 10
daemon 12
sysadmin 14
nobody 60001
noaccess 60002
nogroup 65534
```

# Output Field Separator (OFS)

- The default OFS is a space.
- In the print statement, a comma specifies using the OFS.
- If you omit the comma, the fields run together.
- You can also specify a field separator directly in the print statement, as in the following three lines:

```
$ nawk '{ print $3 $4 $2 }' data.file
$ nawk '{ print $3, $4, $2 }' data.file
$ nawk '{ print $3, $4 "\t" $2 }' data.file
```

# Output Field Separator (OFS): Example

```
$ nawk 'BEGIN { OFS="\t" } ; { print $3, $4, $2 }'
data.file
Joel
         Craig
                        NW
Sharon
          Kelly
                        WE
Chris
          Foster
                        SW
         Chin
May
                        SO
Derek
          Johnson
                        SE
Susan Beal
                        EA
TJ
         Nichols
                        NE
Val
        Shultz
                        NO
Sheri
          Watson
                        CT
```

# Number of Records (NR)

- The number of records (NR) variable counts the number of input lines read from the beginning of the first input file.
- The variable's value is updated each time another input line is read.

# Number of Records (NR): Example

```
$ more report4.nawk
{ print $3, $4, $2 }
END { print "The number of employee records is " NR }
$ nawk -f report4.nawk data.file
Joel Craig NW
Sharon Kelly WE
Chris Foster SW
May Chin SO
Derek Johnson SE
Susan Beal EA
TJ Nichols NE
Val Shultz NO
Sheri Watson CT
The number of employee records is 9
```

#### **User-Defined Variables**

- nawk allows you to create your own variables.
- Variable names should not conflict with the nawk variables or function names.
- The value assigned to a variable can be:
  - A string (string of characters)
  - A numeric
  - A literal value ("hello")
  - The contents from a field from the input file

# **User-Defined Variables: Example**

```
$ cat numexample.nawk
{ counter = counter + 1 }
{ print $0 }
END { print "*** The number of records is " counter }
$ nawk -f numexample.nawk data.file
northwest
               Joel Craiq 3.0 .98
         NW
western WE Sharon Kelly 5.3.97
                                               23
southwest SW Chris Foster 2.7.8
                                               18
               May Chin 5.1 .95
southern SO
                                               15
               Derek Johnson 5.0 .70
southeast SE
                                               17
eastern EA Susan Beal 4.4.8
                                               20
northeast NE TJ Nichols 5.1 .94
                                               13
north NO Val Shultz 4.5 .89
central CT Sheri Watson 5.7.94
                                               13
*** The number of records is 9
```

# **User-Defined Variables: Example**

```
$ cat numexample2.nawk
{ total = total + $8 }
{ print "Field 8 = " $8 }
END { print "Total = " total }
$ nawk -f numexample2.nawk data.file
Field 8 = 4
Field 8 = 23
Field 8 = 18
Field 8 = 15
Field 8 = 17
Field 8 = 20
Field 8 = 13
Field 8 = 9
Field 8 = 13
Total = 132
```

# **User-Defined Variables: Example**

```
$ cat numexample3.nawk
{ total = total + $8 }
{ print $0 }
END { print "The total of field 8 is " total }
$ nawk -f numexample3.nawk data.file
                Joel Craiq 3.0 .98
northwest
         NW
                                                4
                Sharon Kelly 5.3.97
                                                23
western WE
               Chris Foster 2.7 .8
southwest SW
                                                18
               May Chin 5.1 .95
southern SO
                                                15
               Derek Johnson 5.0 .70
southeast SE
                                                17
eastern EA Susan Beal 4.4.8
                                                20
northeast
        NE
               TJ Nichols 5.1 .94
                                                13
north NO Val Shultz 4.5.89
central CT Sheri Watson 5.7.94
                                                13
The total of field 8 is 132
```

```
$ nawk '/N[EOW] { print NR, $0 }' data.file
                     Joel Craig 3.0 .98 3
1 northwest
         NW
7 northeast NE TJ Nichols 5.1 .94 3
           NO Val Shultz 4.5 .89 5
8 north
$ nawk 'BEGIN { count = 0 }
> /N/ { print NR, $0; count = count + 1 }
> END { print "count of North regions is", count }'
data.file
1 northwest
                     Joel Craig 3.0 .98 3
         NW
         NE TJ Nichols 5.1 .94 3
7 northeast
                                             13
8 north NO Val Shultz 4.5.89 5
count of North regions is 3
```

```
$ nawk '{ print "Record:", NR, $NF }' data.file
Record: 1 4
Record: 2 23
Record: 3 18
Record: 4 15
Record: 5 17
Record: 6 20
Record: 7 13
Record: 8 9
Record: 9 13
```

```
$ nawk '{ print "Record:", NR, "has", NF, "fields." }' \
raggeddata.file
Record: 1 has 8 fields.
Record: 2 has 6 fields.
Record: 3 has 8 fields.
Record: 4 has 7 fields.
Record: 5 has 5 fields.
Record: 6 has 6 fields.
Record: 7 has 7 fields.
Record: 8 has 5 fields.
Record: 9 has 6 fields.
```

```
$ nawk '{ print "Field 1 has", length($1), "letters." }'
raggeddata.file
Field 1 has 9 letters.
Field 1 has 2 letters.
Field 1 has 9 letters.
Field 1 has 8 letters.
Field 1 has 9 letters.
Field 1 has 7 letters.
Field 1 has 2 letters.
Field 1 has 3 letters.
Field 1 has 3 letters.
Field 1 has 7 letters.
```

# **Writing Output to Files**

In any statement that generates an output, it is possible to have that output redirected to a file name instead, as follows:

Use the redirection symbol, >, to send data to a file.

```
$ nawk '{ print $2, $1 > "textfile" }' data.file
$ cat textfile

NW northwest
WE western
SW southwest
SO southern
SE southeast
EA eastern
NE northeast
NO north
CT central
```

Use two redirection symbols, >>, to append to a file.

#### The printf() Statement

- The printf() statement allows you to print a character string.
- This string can contain place holders that represent other values.
- These place holders can represent integers, floating points, characters, and character strings.
- Syntax:

```
printf ("string_of_characters" [ , data_values ] )
```

**Note:** For every place holder in the character string of printf(), there must be a value following the character string. These values must be separated by a comma.

# The printf() Statement

```
$ nawk '{ printf "%10s %3d \n", $4, $7 }' data.file
Craig 3
Kelly 5
Foster 2
Chin 4
Johnson 4
Beal 5
Nichols 3
Shultz 5
Watson 5
```

**Note:** The printf statement does not use the OFS variable because of the printf statement's inherent ability to format the output line. So adjusting the fields to appear in column format is done by preceding the format specification with a digit, which refers to the number of spaces that the value should consume.

# **Summary**

In this lesson, you should have learned how to:

- Describe the capabilities of the nawk programming language
- Display output by using the print statement
- Perform pattern matching by using regular expressions
- Use the nawk built-in and user-defined variables

# Practice 7 Overview: The nawk Programming Language

This practice covers the following topics:

- Using nawk and Regular Expressions
- Using nawk to Create a Report