USEFUL BASH ONE-LINERS

... and useful Bash features

FILES...

echo "Hello World" > file
echo "Hello Mars" >>file
>file

Write text to file

Append text to file

Truncate file (DANGER!)

while read -r x y z _; do
 echo \$x \$y \$z
done <file</pre>

Loop over first three fields in file

HEREDOC

cat <<EOF >file
This HEREDOC will have
\$variables substituted
EOF

Cat <<"EOF" >file
This HEREDOC will be used
\$verbatim
EOF

PROCESS MANAGEMENT

```
program &
jobs

fg 1
[Ctrl+Z]

bg 1
```

SORTING

sort file | uniq -c

export LC_ALL=C

Locale configuration changes sorting *sigh*

sort -k 1,1 -k2,2n file

E.g., sort BED file (better use bedtools sort with genome/FAI file)

sort -u file

Remove duplicate lines from file

Print lines without

duplicates, with count

WORKING WITH (TAB) DELIMITED FILES

Select fields from TSV file

Select fields from CSV file

Delete all jobs with the given job name from SGE queue (see also xargs on later slide)

COMMAND SUBSTITUTION

Read from/write into bash commands

```
cat <(grep pattern in) >out
```

```
cat FILE | tee >(grep pattern >out2) >out1
```

• Use result of bash result in variable

```
for x in $(find /src -name foo.txt); do echo $x; done
```

VARIABLE SUBSTITUTION

• Trim endings

```
var=file.vcf.gz
echo ${var%.vcf.gz} # => "file"
```

• Perform string replacement

```
var=foo.bar
echo ${var/i/o} # => "fio.bar"
echo ${var//i/o} # => "fii.bar"
```

• Use default values

```
threads=${threads:4}
```

RENAME STRINGS IN MULTIPLE FILES

- Rename string_old in string_new in all files in /path
 find /path -type f -exec sed -i 's/string_old/string_new/g'
 \;
 - Example:

```
#Create the files
mkdir tmp_dir
echo 'acgt' > tmp_dir/file1
echo 'acgt' > tmp_dir/file2
#Edit the files
find tmp_dir -type f -exec sed -i 's/acgt/ACGT/g' {} \;
cat tmp_dir/*
```

XARGS, I LIKE

Read command line parts from stdin
 find . -name 'pattern.' | xargs grep PATTERN

KILL ALL CHROME PROCESSES

• Read command line parts from stdin

```
for p in $(ps -A | grep chrom | awk '{print $1}');
do kill -9 $p; done
```

IN-PLACE SEARCH AND REPLACE

```
If you like Perl:
perl -p -i -e 's/needle/pin/g' haystack.txt
If you prefer sed:
sed -i -e 's/needle/pin/g' haystack.txt
```

THE UNOFFICIAL BASH STRICT MODE

```
#!/bin/bash
set -euo pipefail
IFS=$'\n\t'
```

http://redsymbol.net/articles/unofficial-bash-strict-mode/

LESS

```
Large files can be easily viewed with less
$ less <filename>
The options -N and -S are useful to display files with many
columns: -S suppresses linebreaks, and -N adds rownumbers:
$ less -NS <file_with_lots_of_rows_and_cols>
```

LESS

seqid source type start end score strand phase ID gene_id transcript_id gene_type gene_status gene_name transcript_type transcript_status transcr ipt_name level havana_gene Parent havana_transcript exon_number exon_id ont tag protein_id ccdsid length mean_length
chrī HAVANA gene 11869 14412 NA + NA gene1 ENSG00000223972.4 ENSG00000223972.4 pseudogene KNOWN DDX11L1 pseudogene KNOWN DDX11L1 2 OTTHUMG00000000961.2 character(0) NA NA NA Character(0) character(0) NA NA 2543 2543
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KNOWN DDX11L1-002 2 OTTHUMG00000000061.2 gene1 OTTHUMT00000362751.1 NA NA character(0) character(0) NA NA 2540 2318.75 chr1 HAVANA exon 11869 12227 NA + NA exon1 ENSG00000223972.4 ENST00000456328.2 pseudogene KNOWN DDX11L1 processed_transcript KNOWN DDX11L1
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chr1 NA intron 12228 12612 NA + NA intron1 ENSG00000223972.4 ENST00000456328.2 pseudogene KNOWN DDX11L1 processed_transcript KNOWN DDX11L1
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-002 2 OTTHUMG000000000961.2 transcript1 OTTHUMT00000362751.1 2 ENSE00003582793.1 character(0) character(0) NA NA 1654 1224
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chr1 NA intron 12228 12612 NA + NA intron3 ENSG00000223972.4 ENST00000515242.2 pseudogene KNOWN DDX11L1 transcribed_unprocessed_pseudogene KNOWN DDX11L1-201 3 OTTHUMG00000000961.2 transcript2 NA 1 ENSE00002234632.1 character(0) character(0) NA NA 886 983.83333333333
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pseudogene KNOWN DDX11L1-202 3 OTTHUMG0000000961.2 gene1 NA NA NA character(0) character(0) NA NA 2535 2318.75
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chr1 ENSEMBL exon 13661 14409 NA + NA exon10 ENSG00000223972.4 ENST00000518655.2 pseudogene KNOWN DDX11L1 transcribed_unprocessed_pseudogene KNOWN DDX11L1-202 3 OTTHUMG00000000961.2 transcript3 NA 4 ENSE00002303382.1 character(0) character(0) NA NA 1479 1224
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_pseudogene KNOWN DDX11L1-001 2 OTTHUMG00000000961.2 gene1 OTTHUMT00000002844.2 NA NA "c(""PGO:0000005"", ""PGO:0000019"")" character(0) NA NA 1660 2318.75
chr1 HAVANA exon 12010 12057 NA + NA exon11 ENSG00000223972.4 ENST00000450305.2 pseudogene KNOWN DDX11L1 transcribed_unprocessed_pseudogene
KNOWN DDX11L1-001 2 OTTHUMG00000000961.2 transcript4 OTTHUMT00000002844.2 1 ENSE00001948541.Ī "c(""PG0:0000005"", ""PG0:00000019"")" character(0) N
A NA 626 1224 chr1 NA intron 12058 12178 NA + NA intron8 ENSG00000223972.4 ENST00000450305.2 pseudogene KNOWN DDX11L1 transcribed unprocessed pseudogene
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A NA 626 1224
chr1 NA intron 12228 12612 NA + NA intron9 ENSG00000223972.4 ENST00000450305.2 pseudogene KNOWN DDX11L1 transcribed_unprocessed_pseudogene KNOWN DDX11L1-001 2 OTTHUMG00000000961.2 transcript4 OTTHUMT00000002844.2 2 ENSE00001671638.2 "c(""PG0:0000005"", ""PG0:0000019"")" character(0) N
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chrl HAVANA exon 12613 12697 NA + NA exon13 ENSG00000223972.4 ENST00000450305.2 pseudogene KNOWN DDX11L1 transcribed_unprocessed_pseudogene
gff3v19.tsv

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	1	segid	source	type	start	end	score	strand	phase	ID	gene_id transcri	pt_id	gene_type	gene_status	gene_name	transcr	ipt_type tran:	script_statu	us
		chr1	HAVANA	gene	11869	14412	NA	+	NA	gene1	ENSG00000223972.		ENSG00000223972.	4 pseudog			pseudogene		DDX11L1
		chr1	HAVANA	transcr		11869	14409	NA	+	NA			00223972.4	ENST00000456328			KNOWN DDX1		
		chr1 chr1	HAVANA NA	exon intron	11869 12228	12227 12612	NA NA	+	NA NA	exon1	ENSG00000223972.		ENST00000456328. ENST00000456328.				processed_tr		KNOWN KNOWN
		chr1	NA HAVANA	exon	12613	12721	NA	+	NA	exon2	ENSG00000223972.		ENST00000456328.				processed_tra		KNOWN
	7	chr1	NA	intron		13220	NA	+	NA		ENSG00000223972.		ENST00000456328.				processed_tr		KNOWN
	8	chr1	HAVANA	exon	13221	14409	NA	+	NA	exon3	ENSG00000223972.		ENST00000456328.				processed tr		KNOWN
	9	chr1	ENSEMBL	transcr	ript	11872	14412	NA		NA	transcript2	ENSG0000	00223972.4	ENST00000515242				IL1 transcr	ibed_unp
		chr1	ENSEMBL		11872	12227	NA		NA	exon4	ENSG00000223972.		ENST00000515242.				transcribed_		
		. chr1	NA	intron		12612	NA		NA		ENSG00000223972.		ENST00000515242.				transcribed_		
		chr1	ENSEMBL		12613	12721	NA		NA NA	exon5	ENSG00000223972.		ENST00000515242.				transcribed_		
		chr1 chr1	NA ENSEMBL	intron	12722 13225	13224 14412	NA NA	+	NA NA	exon6	ENSG00000223972.		ENST00000515242. ENST00000515242.				transcribed_ transcribed		
		chr1		transcr		11874	14409	NA	+	NA			00223972.4	ENST00000518655				IL1 transcr	
		chr1	ENSEMBL		11874	12227	NA	+	NA	exon7	ENSG00000223972.		ENST00000518655.				transcribed_		
		chr1	NA	intron	12228	12594	NA	+	NA		ENSG00000223972.		ENST00000518655.				transcribed_		
	18	chr1	ENSEMBL	exon	12595	12721	NA		NA	exon8	ENSG00000223972.		ENST00000518655.	2 pseudog			transcribed_		
		chr1	NA	intron		13402	NA		NA		ENSG00000223972.		ENST00000518655.				transcribed_		
		chr1	ENSEMBL		13403	13655	NA		NA	exon9	ENSG00000223972.		ENST00000518655.				transcribed_		
		chr1	NA	intron	13656 13661	13660 14409	NA NA	+	NA	intron7			ENST00000518655.				transcribed_		
		chr1 chr1	ENSEMBL HAVANA	transcr		12010	13670	NA	NA +	exon10 NA	ENSG00000223972.		ENST00000518655. 00223972.4	.2 pseudog ENST00000450305			transcribed_ KNOWN DDX1		
		chr1	HAVANA	exon	12010	12057	NA	+	NA	exon11	ENSG00000223972.		ENST00000450305.				transcribed_		
		chr1	NA	intron		12178	NA	+	NA		ENSG00000223972.		ENST00000450305.				transcribed_		
		chr1	HAVANA	exon	12179	12227	NA		NA	exon12	ENSG00000223972.		ENST00000450305.				transcribed_		
		chr1	NA	intron	12228	12612	NA		NA	intron9	ENSG00000223972.		ENST00000450305.				transcribed_		
		chr1	HAVANA	exon	12613	12697	NA	+	NA	exon13	ENSG00000223972.		ENST00000450305.				transcribed_		
		chr1	NA	intron	12698	12974	NA		NA	intron1				00450305.2	pseudogene	KNOWN	DDX11L1 tran		
		chr1	HAVANA NA	exon intron	12975	13052	NA NA	+	NA		ENSG00000223972.		ENST00000450305.			KNOWN	transcribed_		
		chr1	HAVANA	exon	13053 13221	13220 13374	NA	+	NA NA	intron1 exon15	1 ENSG0000 ENSG00000223972.		ENST0000450305.	00450305.2 .2 pseudoc	pseudogene gene KNOWN		DDX11L1 tran: transcribed_		
		chr1	NA	intron	13375	13452	NA	+	NA	intron1				00450305.2	pseudogene	KNOWN	DDX11L1 tran		
		chr1	HAVANA	exon	13453	13670	NA	+	NA	exon16	ENSG00000223972.		ENST00000450305.				transcribed_		
		chr1	HAVANA	gene	14363	29806	NA		NA	gene2	ENSG00000227232.		ENSG00000227232.			WASH7P	pseudogene	KNOWN	
		chr1	ENSEMBL			14363	24886	NA		NA			00227232.4	ENST00000541675				7P unproces	
		chr1	ENSEMBL		14363	14829	NA		NA	exon17	ENSG00000227232.		ENST00000541675.			WASH7P	unprocessed_		
		chr1	NA	intron	14830	14969	NA		NA	intron1				00541675.1	pseudogene	KNOWN	WASH7P unpr		
		chr1	ENSEMBL NA	exon	14970 15039	15038 16853	NA NA	_	NA NA	exon18 intron1	ENSG00000227232.		ENST00000541675.			WASH7P KNOWN	unprocessed_		
		chr1 chr1	ENSEMBL		16854	17055	NA	_	NA	exon19	4 ENSG00000 ENSG00000227232.4		ENST00000541675.	00541675.1 .1 pseudoc	pseudogene gene KNOWN	WASH7P	WASH7P unprocessed_		
		chr1	NA	intron	17056	17232	NA		NA	intron1				0541675.1	pseudogene	KNOWN	WASH7P unpr		
		chr1	ENSEMBL		17233	17364	NA		NA		ENSG00000227232.		ENST00000541675.			WASH7P	unprocessed_		
	44	chr1	NA	intron	17365	17497	NA		NA	intron1	6 ENSG0000	0227232.		00541675.1	pseudogene	KNOWN	WASH7P unpr		
		chr1	ENSEMBL		17498	17504	NA		NA	exon21	ENSG00000227232.		ENST00000541675.			WASH7P	unprocessed_		
		chr1	NA	intron		17605	NA		NA	intron1				00541675.1	pseudogene	KNOWN	WASH7P unpr		
		chr1	ENSEMBL		17606	17742	NA		NA	exon22	ENSG00000227232.		ENST00000541675.			WASH7P	unprocessed_		
		chr1	NA ENSEMBL	intron	17743 17915	17914 18061	NA NA		NA NA	intron1	8 ENSG0000 ENSG00000227232.4		ENST0000541675.	00541675.1	pseudogene gene KNOWN	KNOWN WASH7P	WASH7P unprunprocessed		
		chr1	NA	intron	18062	18267	NA	_	NA	intron1				.1 pseudog 90541675.1	pseudogene	KNOWN	WASH7P unpr		
		chr1	ENSEMBL		18268	18369	NA		NA		ENSG00000227232.		ENST00000541675.			WASH7P	unprocessed		
		chr1	NA	intron		24733	NA		NA	intron2				0541675.1	pseudogene	KNOWN	WASH7P unpr		
	53	chr1	ENSEMBL		24734	24886	NA		NA	exon25	ENSG00000227232.	4	ENST00000541675.	1 pseudog	gene KNOWN	WASH7P	unprocessed_	seudogene	KNOWN
		chr1		transcr		14363	29370	NA		NA			00227232.4	ENST00000438504			KNOWN WASH	7P unproces	ssed_pse
		chr1	ENSEMBL		14363	14829	NA		NA		ENSG00000227232.		ENST00000438504.			WASH7P	unprocessed_	oseudogene	KNOWN
		chr1	NA	intron	14830	14969	NA		NA	intron2				00438504.2	pseudogene	KNOWN	WASH7P unpr		
		chr1 chr1	ENSEMBL NA	intron	14970 15039	15038 15795	NA NA	_	NA NA	exon27 intron2	ENSG00000227232.4 2 ENSG0000		ENST00000438504.	.2 pseudog 00438504.2		WASH7P KNOWN	unprocessed_ WASH7P unpr	oseudogene	KNUWN
		chr1	ENSEMBL		15796	15901	NA	_	NA NA	exon28	ENSG00000227232.4		ENST0000438504.		pseudogene gene KNOWN	WASH7P	unprocessed_		
		chr1	NA	intron		15903	NA		NA	intron2				0438504.2	pseudogene	KNOWN	WASH7P unpr		
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