

pyrpipe report

file name:pyrpipe_logs/2019-12-24-16_14_55_pyrpipe.log

Summary

Time start: 2019-12-24 16:14:55 Time end: 2019-12-24 16:31:03
Total time: 0:16:08

Num commands: 19

Num failed commands: 0

Num passed commands: 19

Total programs: 6

Programs: hisat2-build,samtools,hisat2,stringtie,fasterq-dump,prefetch

Details

command

```
prefetch -O /home/usingh/work/urmi/hoap/test/athalData/  
sraData/SRR976159 SRR976159
```

stdout

```
2019-12-24T22:14:57 prefetch.2.9.3: 1) Downloading  
'SRR976159'...  
2019-12-24T22:14:57 prefetch.2.9.3: Downloading via https...  
2019-12-24T22:15:08 prefetch.2.9.3: 1) 'SRR976159' was  
downloaded successfully
```

stderr

Program	Return code	Start	Runtime	Version
prefetch	0	19-12-24 16:14:57	0:00:11	prefetch : 2.9.3 (2.9.3-1)

command

```
prefetch -0 /home/usingh/work/urmi/hoap/test/athalData/  
sraData/SRR978411 SRR978411
```

stdout

```
2019-12-24T22:15:09 prefetch.2.9.3: 1) Downloading  
'SRR978411'...  
2019-12-24T22:15:09 prefetch.2.9.3: Downloading via https...  
2019-12-24T22:15:20 prefetch.2.9.3: 1) 'SRR978411' was  
downloaded successfully
```

stderr

Program	Return code	Start	Runtime	Version
prefetch	0	19-12-24 16:15:08	0:00:11	prefetch : 2.9.3 (2.9.3-1)

command

```
prefetch -0 /home/usingh/work/urmi/hoap/test/athalData/  
sraData/SRR971778 SRR971778
```

stdout

```
2019-12-24T22:15:20 prefetch.2.9.3: 1) Downloading  
'SRR971778'...  
2019-12-24T22:15:20 prefetch.2.9.3: Downloading via https...  
2019-12-24T22:15:32 prefetch.2.9.3: 1) 'SRR971778' was  
downloaded successfully
```

stderr

Program	Return code	Start	Runtime	Version
prefetch	0	19-12-24 16:15:20	0:00:12	prefetch : 2.9.3 (2.9.3-1)

command

```
fasterq-dump -e 8 -f -t /home/usingh/work/urmi/hoap/test/
athalData/sraData -O /home/usingh/work/urmi/hoap/test/
athalData/sraData/SRR976159 -o SRR976159.fastq /home/usingh/
work/urmi/hoap/test/athalData/sraData/SRR976159/SRR976159.sra
```

stdout

```
spots read      : 4,728,806
reads read      : 9,457,612
reads written    : 9,457,612
```

stderr

Program	Return code	Start	Runtime	Version
fasterq-dump	0	19-12-24 16:16:22	0:00:24	fasterq-dump : 2.9.6

command

```
fasterq-dump -e 8 -f -t /home/usingh/work/urmi/hoap/test/
athalData/sraData -O /home/usingh/work/urmi/hoap/test/
athalData/sraData/SRR978411 -o SRR978411.fastq /home/usingh/
work/urmi/hoap/test/athalData/sraData/SRR978411/SRR978411.sra
```

stdout

```
spots read      : 4,200,625
reads read      : 8,401,250
reads written    : 8,401,250
```

stderr

Program	Return code	Start	Runtime	Version
fasterq-dump	0	19-12-24 16:16:46	0:00:24	fasterq-dump : 2.9.6

command

```
fasterq-dump -e 8 -f -t /home/usingh/work/urmi/hoap/test/
athalData/sraData -O /home/usingh/work/urmi/hoap/test/
athalData/sraData/SRR971778 -o SRR971778.fastq /home/usingh/
work/urmi/hoap/test/athalData/sraData/SRR971778/SRR971778.sra
```

stdout

```
spots read      : 5,432,321
reads read      : 10,864,642
reads written   : 10,864,642
```

stderr

Program	Return code	Start	Runtime	Version
fasterq-dump	0	19-12-24 16:17:11	0:00:31	fasterq-dump : 2.9.6

command

```
hisat2-build -p 8 -a -q /home/usingh/work/urmi/hoap/test/
athalData/ref/Arabidopsis_thaliana.TAIR10.dna.toplevel.fa /
home/usingh/work/urmi/hoap/test/athalData/sraData/athalIndex/
athalInd
```

stdout

stderr

Program	Return code	Start	Runtime	Version
hisat2-build	0	19-12-24 16:17:45	0:00:39	hisat2-build version 2.1.0 64-bit Built on login-node03 Wed Jun 7 15:52:43 EDT 2017 Compiler: gcc version 4.8.2 (GCC) Options: -O3 -m64 -msse2 -funroll-loops - g3 -DPOPCNT_CAPABILITY Sizeof {int, long, long long, void*, size_t, off_t}: {4, 8, 8, 8, 8, 8}

command

```
hisat2 --dta-cufflinks -p 10 -x /home/usingh/work/urmi/hoap/
test/athalData/sraData/athalIndex/athalInd -1 /home/usingh/
work/urmi/hoap/test/athalData/sraData/SRR976159/
SRR976159_1.fastq -2 /home/usingh/work/urmi/hoap/test/
athalData/sraData/SRR976159/SRR976159_2.fastq -S /home/
usingh/work/urmi/hoap/test/athalData/sraData/SRR976159/
SRR976159_hisat2.sam
```

stdout

```
4728806 reads; of these:
  4728806 (100.00%) were paired; of these:
    214957 (4.55%) aligned concordantly 0 times
    4401400 (93.08%) aligned concordantly exactly 1 time
    112449 (2.38%) aligned concordantly >1 times
    ----
    214957 pairs aligned concordantly 0 times; of these:
      39783 (18.51%) aligned discordantly 1 time
    ----
    175174 pairs aligned 0 times concordantly or
discordantly; of these:
    350348 mates make up the pairs; of these:
      194409 (55.49%) aligned 0 times
      150911 (43.07%) aligned exactly 1 time
      5028 (1.44%) aligned >1 times
97.94% overall alignment rate
```

stderr

Program	Return code	Start	Runtime	Version
hisat2	0	19-12-24 16:18:50	0:01:12	/home/usingh/lib_urmi/software/hisat2-2.1.0/hisat2-align-s version 2.1.0 64-bit Built on login-node03 Wed Jun 7 15:53:42 EDT 2017 Compiler: gcc version 4.8.2 (GCC) Options: -O3 -m64 -msse2 -funroll-loops -g3 -DPOPCNT_CAPABILITY Sizeof {int, long, long long, void*, size_t, off_t}: {4, 8, 8, 8, 8, 8}

command

```
hisat2 --dta-cufflinks -p 10 -x /home/usingh/work/urmi/hoap/test/athalData/sraData/athalIndex/athalInd -1 /home/usingh/work/urmi/hoap/test/athalData/sraData/SRR978411/SRR978411_1.fastq -2 /home/usingh/work/urmi/hoap/test/athalData/sraData/SRR978411/SRR978411_2.fastq -S /home/usingh/work/urmi/hoap/test/athalData/sraData/SRR978411/SRR978411_hisat2.sam
```

stdout

```
4200625 reads; of these:
  4200625 (100.00%) were paired; of these:
```

```

202812 (4.83%) aligned concordantly 0 times
3890411 (92.62%) aligned concordantly exactly 1 time
107402 (2.56%) aligned concordantly >1 times
----
202812 pairs aligned concordantly 0 times; of these:
 39649 (19.55%) aligned discordantly 1 time
----
163163 pairs aligned 0 times concordantly or
discordantly; of these:
 326326 mates make up the pairs; of these:
 183385 (56.20%) aligned 0 times
 137749 (42.21%) aligned exactly 1 time
  5192 (1.59%) aligned >1 times
97.82% overall alignment rate

```

stderr

Program	Return code	Start	Runtime	Version
hisat2	0	19-12-24 16:20:02	0:01:05	/home/usingh/lib_urmi/software/ hisat2-2.1.0/hisat2-align-s version 2.1.0 64-bit Built on login-node03 Wed Jun 7 15:53:42 EDT 2017 Compiler: gcc version 4.8.2 (GCC) Options: -O3 -m64 -msse2 -funroll-loops -g3 -DPOPCNT_CAPABILITY Sizeof {int, long, long long, void*, size_t, off_t}: {4, 8, 8, 8, 8, 8}

command

```

hisat2 --dta-cufflinks -p 10 -x /home/usingh/work/urmi/hoap/
test/athalData/sraData/athalIndex/athalInd -1 /home/usingh/
work/urmi/hoap/test/athalData/sraData/SRR971778/
SRR971778_1.fastq -2 /home/usingh/work/urmi/hoap/test/
athalData/sraData/SRR971778/SRR971778_2.fastq -S /home/
usingh/work/urmi/hoap/test/athalData/sraData/SRR971778/
SRR971778_hisat2.sam

```

stdout

```

5432321 reads; of these:
 5432321 (100.00%) were paired; of these:
  650714 (11.98%) aligned concordantly 0 times
 4600911 (84.70%) aligned concordantly exactly 1 time
 180696 (3.33%) aligned concordantly >1 times
----

```

```

650714 pairs aligned concordantly 0 times; of these:
  16819 (2.58%) aligned discordantly 1 time
----
633895 pairs aligned 0 times concordantly or
discordantly; of these:
  1267790 mates make up the pairs; of these:
    1145159 (90.33%) aligned 0 times
    116666 (9.20%) aligned exactly 1 time
    5965 (0.47%) aligned >1 times
89.46% overall alignment rate

```

stderr

Program	Return code	Start	Runtime	Version
hisat2	0	19-12-24 16:21:08	0:01:24	/home/usingh/lib_urmi/software/ hisat2-2.1.0/hisat2-align-s version 2.1.0 64-bit Built on login-node03 Wed Jun 7 15:53:42 EDT 2017 Compiler: gcc version 4.8.2 (GCC) Options: -O3 -m64 -msse2 -funroll-loops -g3 -DPOPCNT_CAPABILITY Sizeof {int, long, long long, void*, size_t, off_t}: {4, 8, 8, 8, 8, 8}

command

```

samtools view -@ 8 -o /home/usingh/work/urmi/hoap/test/
athalData/sraData/SRR976159/SRR976159_hisat2.bam -b /home/
usingh/work/urmi/hoap/test/athalData/sraData/SRR976159/
SRR976159_hisat2.sam

```

stdout

stderr

Program	Return code	Start	Runtime	Version
samtools	0	19-12-24 16:24:52	0:00:33	samtools 1.9 Using htslib 1.9 Copyright (C) 2018 Genome Research Ltd.

command

```
samtools sort -@ 8 -o /home/usingh/work/urmi/hoap/test/athalData/sraData/SRR976159/SRR976159_hisat2_sorted.bam /home/usingh/work/urmi/hoap/test/athalData/sraData/SRR976159/SRR976159_hisat2.bam
```

stdout

```
[bam_sort_core] merging from 0 files and 8 in-memory blocks...
```

stderr

Program	Return code	Start	Runtime	Version
samtools	0	19-12-24 16:25:25	0:00:17	samtools 1.9 Using htlib 1.9 Copyright (C) 2018 Genome Research Ltd.

command

```
samtools view -@ 8 -o /home/usingh/work/urmi/hoap/test/athalData/sraData/SRR978411/SRR978411_hisat2.bam -b /home/usingh/work/urmi/hoap/test/athalData/sraData/SRR978411/SRR978411_hisat2.sam
```

stdout

stderr

Program	Return code	Start	Runtime	Version
samtools	0	19-12-24 16:25:43	0:00:30	samtools 1.9 Using htlib 1.9 Copyright (C) 2018 Genome Research Ltd.

command

```
samtools sort -@ 8 -o /home/usingh/work/urmi/hoap/test/athalData/sraData/SRR978411/SRR978411_hisat2_sorted.bam /home/usingh/work/urmi/hoap/test/athalData/sraData/SRR978411/SRR978411_hisat2.bam
```


stdout

```
[bam_sort_core] merging from 0 files and 8 in-memory  
blocks...
```

stderr

Program	Return code	Start	Runtime	Version
samtools	0	19-12-24 16:26:13	0:00:15	samtools 1.9 Using htslib 1.9 Copyright (C) 2018 Genome Research Ltd.

command

```
samtools view -@ 8 -o /home/usingh/work/urmi/hoap/test/  
athalData/sraData/SRR971778/SRR971778_hisat2.bam -b /home/  
usingh/work/urmi/hoap/test/athalData/sraData/SRR971778/  
SRR971778_hisat2.sam
```

stdout

stderr

Program	Return code	Start	Runtime	Version
samtools	0	19-12-24 16:26:28	0:00:41	samtools 1.9 Using htslib 1.9 Copyright (C) 2018 Genome Research Ltd.

command

```
samtools sort -@ 8 -o /home/usingh/work/urmi/hoap/test/  
athalData/sraData/SRR971778/SRR971778_hisat2_sorted.bam /  
home/usingh/work/urmi/hoap/test/athalData/sraData/SRR971778/  
SRR971778_hisat2.bam
```

stdout

```
[bam_sort_core] merging from 0 files and 8 in-memory  
blocks...
```

stderr

Program	Return code	Start	Runtime	Version
samtools	0	19-12-24 16:27:10	0:00:19	samtools 1.9 Using htslib 1.9 Copyright (C) 2018 Genome Research Ltd.

command

```
stringtie -G /home/usingh/work/urmi/hoap/test/athalData/ref/Arabidopsis_thaliana.TAIR10.45.gtf -o /home/usingh/work/urmi/hoap/test/athalData/sraData/SRR976159/SRR976159_hisat2_sorted_stringtie.gtf /home/usingh/work/urmi/hoap/test/athalData/sraData/SRR976159/SRR976159_hisat2_sorted.bam
```

stdout

stderr

Program	Return code	Start	Runtime	Version
stringtie	0	19-12-24 16:28:55	0:00:48	2.0.4

command

```
stringtie -G /home/usingh/work/urmi/hoap/test/athalData/ref/Arabidopsis_thaliana.TAIR10.45.gtf -o /home/usingh/work/urmi/hoap/test/athalData/sraData/SRR978411/SRR978411_hisat2_sorted_stringtie.gtf /home/usingh/work/urmi/hoap/test/athalData/sraData/SRR978411/SRR978411_hisat2_sorted.bam
```

stdout

stderr

Program	Return code	Start	Runtime	Version
stringtie	0	19-12-24 16:29:43	0:00:38	2.0.4

command

```
stringtie -G /home/usingh/work/urmi/hoap/test/athalData/ref/
Arabidopsis_thaliana.TAIR10.45.gtf -o /home/usingh/work/urmi/
hoap/test/athalData/sraData/SRR971778/
SRR971778_hisat2_sorted_stringtie.gtf /home/usingh/work/urmi/
hoap/test/athalData/sraData/SRR971778/
SRR971778_hisat2_sorted.bam
```

stdout

stderr

Program	Return code	Start	Runtime	Version
stringtie	0	19-12-24 16:30:22	0:00:41	2.0.4

Environment Information

Programs		
name	version	path
hisat2-build	hisat2-build version 2.1.0 64-bit Built on login-node03 Wed Jun 7 15:52:43 EDT 2017 Compiler: gcc version 4.8.2 (GCC) Options: -O3 -m64 -msse2 -funroll-loops -g3 -DPOPCNT_CAPABILITY Sizeof {int, long, long long, void*, size_t, off_t}: {4, 8, 8, 8, 8, 8}	/home/usingh/lib_urmi/softwares/hisat2-2.1.0/hisat2-build
samtools	samtools 1.9 Using htlib 1.9 Copyright (C) 2018 Genome Research Ltd.	/home/usingh/anaconda3/envs/pyrpipeTest/bin/samtools
hisat2	/home/usingh/lib_urmi/softwares/hisat2-2.1.0/hisat2-align-s version 2.1.0 64-bit Built on login-node03 Wed Jun 7 15:53:42 EDT 2017 Compiler: gcc version 4.8.2 (GCC) Options: -O3 -m64 -msse2 -funroll-loops -g3 -DPOPCNT_CAPABILITY Sizeof {int, long, long long, void*, size_t, off_t}: {4, 8, 8, 8, 8, 8}	/home/usingh/lib_urmi/softwares/hisat2-2.1.0/hisat2
stringtie	2.0.4	/home/usingh/lib_urmi/softwares/stringtie-2.0.4.Linux_x86_64/stringtie

fasterq-dump	fasterq-dump : 2.9.6	/home/usingh/lib_urmi/softwares/sratoolkit.2.9.6-1-centos_linux64/bin/fasterq-dump
prefetch	prefetch : 2.9.3 (2.9.3-1)	/home/usingh/lib_urmi/softwares/sratoolkit.2.9.6-1-centos_linux64/bin/prefetch

System Information	
Time at collection	19-12-24 16:14:55
Python	Python 3.7.5 (default, Oct 25 2019, 15:51:11) [GCC 7.3.0]
Operating system	Linux-3.16.0-4-amd64-x86_64-with-debian-8.9
CPU	8 logical CPU cores

sys.modules
'sys'
'builtins'
'_frozen_importlib'
'_imp'
'_thread'
'_warnings'
'_weakref'
'zipimport'
'_frozen_importlib_external'
'_io'
'marshal'
'posix'
'encodings'
'codecs'
'_codecs'
'encodings.aliases'
'encodings.utf_8'
'_signal'
'__main__'
'encodings.latin_1'
'io'
'abc'
'_abc'
'_bootlocale'
'_locale'
'site'

'os'
'stat'
'_stat'
'posixpath'
'genericpath'
'os.path'
'_collections_abc'
'_sitebuiltins'
'types'
'importlib'
'importlib._bootstrap'
'importlib._bootstrap_external'
'warnings'
'importlib.util'
'importlib.abc'
'importlib.machinery'
'contextlib'
'collections'
'operator'
'_operator'
'keyword'
'heapq'
'_heapq'
'itertools'
'reprlib'
'_collections'
'functools'
'_functools'
'mpl_toolkits'
'sphinxcontrib'
'runpy'
'pkgutil'
'weakref'
'_weakrefset'
'ipykernel'
'ipykernel._version'
'ipykernel.connect'
'__future__'
'json'
'json.decoder'
're'
'enum'
'sre_compile'
'_sre'

'sre_parse'
'sre_constants'
'copyreg'
'json.scanner'
'_json'
'json.encoder'
'subprocess'
'time'
'signal'
'errno'
'_posixsubprocess'
'select'
'selectors'
'collections.abc'
'math'
'threading'
'traceback'
'linecache'
'tokenize'
'token'
'IPython'
'IPython.core'
'IPython.core.getipython'
'IPython.core.release'
'IPython.core.application'
'atexit'
'copy'
'glob'
'fnmatch'
'logging'
'string'
'_string'
'shutil'
'zlib'
'bz2'
'_compression'
'_bz2'
'lzma'
'_lzma'
'pwd'
'grp'
'traitlets'
'traitlets.traitlets'
'inspect'

'dis'
'opcode'
'_opcode'
'six'
'struct'
'_struct'
'traitlets.utils'
'traitlets.utils.getargspec'
'traitlets.utils.importstring'
'ipython_genutils'
'ipython_genutils._version'
'ipython_genutils.py3compat'
'ipython_genutils.encoding'
'locale'
'platform'
'traitlets.utils.sentinel'
'traitlets.utils.bunch'
'traitlets._version'
'traitlets.config'
'traitlets.config.application'
'decorator'
'traitlets.config.configurable'
'traitlets.config.loader'
'argparse'
'gettext'
'ast'
'_ast'
'ipython_genutils.path'
'random'
'hashlib'
'_hashlib'
'_blake2'
'_sha3'
'bisect'
'_bisect'
'_random'
'ipython_genutils.text'
'textwrap'
'ipython_genutils.importstring'
'IPython.core.crashhandler'
'pprint'
'IPython.core.ultratb'
'pydoc'
'urllib'

'urllib.parse'
'IPython.core.debugger'
'bdb'
'IPython.utils'
'IPython.utils.PyColorize'
'IPython.utils.coloransi'
'IPython.utils.ipstruct'
'IPython.utils.colorable'
'pygments'
'pygments.util'
'IPython.utils.py3compat'
'IPython.utils.encoding'
'IPython.core.excolors'
'IPython.testing'
'IPython.testing.skipdoctest'
'pdb'
'cmd'
'code'
'codeop'
'IPython.core.display_trap'
'IPython.utils.path'
'IPython.utils.process'
'IPython.utils._process_posix'
'pexpect'
'pexpect.exceptions'
'pexpect.utils'
'pexpect.expect'
'pexpect.pty_spawn'
'pty'
'tty'
'termios'
'ptyprocess'
'ptyprocess.ptyprocess'
'fcntl'
'resource'
'ptyprocess.util'
'pexpect.spawnbase'
'pexpect.run'
'IPython.utils._process_common'
'shlex'
'IPython.utils.decorators'
'IPython.utils.data'
'IPython.utils.terminal'
'IPython.utils.sysinfo'

'IPython.utils._sysinfo'
'IPython.core.profiledir'
'IPython.paths'
'tempfile'
'IPython.utils.importstring'
'IPython.terminal'
'IPython.terminal.embed'
'IPython.core.compilerop'
'IPython.core.magic_arguments'
'IPython.core.error'
'IPython.utils.text'
'pathlib'
'ntpath'
'IPython.core.magic'
'getopt'
'IPython.core.oinspect'
'IPython.core.page'
'IPython.core.display'
'binascii'
'mimetypes'
'IPython.lib'
'IPython.lib.security'
'getpass'
'IPython.lib.pretty'
'datetime'
'_datetime'
'IPython.utils.openpy'
'IPython.utils.dir2'
'IPython.utils.wildcard'
'pygments.lexers'
'pygments.lexers._mapping'
'pygments.modeline'
'pygments.plugin'
'pygments.lexers.python'
'pygments.lexer'
'pygments.filter'
'pygments.filters'
'pygments.token'
'pygments.regexopt'
'pygments.unistring'
'pygments.formatters'
'pygments.formatters._mapping'
'pygments.formatters.html'
'pygments.formatter'

'pygments.styles'
'IPython.core.inputtransformer2'
'typing'
'typing.io'
'typing.re'
'IPython.core.interactiveshell'
'asyncio'
'asyncio.base_events'
'concurrent'
'concurrent.futures'
'concurrent.futures._base'
'socket'
'_socket'
'ssl'
'_ssl'
'base64'
'asyncio.constants'
'asyncio.coroutines'
'asyncio.base_futures'
'asyncio.format_helpers'
'asyncio.log'
'asyncio.events'
'contextvars'
'_contextvars'
'asyncio.base_tasks'
'_asyncio'
'asyncio.futures'
'asyncio.protocols'
'asyncio.sslproto'
'asyncio.transports'
'asyncio.tasks'
'asyncio.locks'
'asyncio.runners'
'asyncio.queues'
'asyncio.streams'
'asyncio.subprocess'
'asyncio.unix_events'
'asyncio.base_subprocess'
'asyncio.selector_events'
'pickleshare'
'pickle'
'_compat_pickle'
'_pickle'
'IPython.core.prefilter'

'IPython.core.autocall'
'IPython.core.macro'
'IPython.core.splitinput'
'IPython.core.alias'
'IPython.core.builtin_trap'
'IPython.core.events'
'backcall'
'backcall.backcall'
'IPython.core.displayhook'
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