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sql-ledger/oldstable 3.0.8-1 all
  Web based double-entry accounting program

ssake/oldstable 3.8.4-1 all
  genomics application for assembling millions of very short DNA sequences

staden/oldstable 2.0.0+b11-2 armhf
  DNA sequence assembly (Gap4/Gap5), editing and analysis tools

staden-common/oldstable 2.0.0+b11-2 all
  Architecture independent files for Staden

tandem-mass/oldstable 1:20151215-4 armhf
  mass spectrometry software for protein identification

tandem-mass-dbg/oldstable 1:20151215-4 armhf
  mass spectrometry software for protein identification - debug symbols

tar-split/oldstable 0.9.13-1 armhf
  tar archive assembly/disassembly

texify/oldstable 1.20-3 all
  Beautify source code for use with LaTeX

tiemu/oldstable 3.03-nogdb+dfsg-3 armhf
  Texas Instruments calculators emulator (without GDB)

transdecoder/oldstable 3.0.1+dfsg-1 all
  find coding regions within RNA transcript sequences

transdecoder-doc/oldstable 3.0.1+dfsg-1 all
  find coding regions within transcripts

trinityrnaseq-examples/oldstable 2.2.0+dfsg-2 all
  RNA-Seq De novo Assembly common example and testing files

ufc/oldstable 2.3.0-1 armhf
  unified code generation interface for form-compilers

ufc-2.0/oldstable 2.0.5-3+rp11 all
  unified code generation interface for form-compilers

ufc-doc-2.0/oldstable 2.0.5-3+rp11 all
  documentation for UFC

velvetoptimiser/oldstable 2.2.5-5 all
  automatically optimise Velvet do novo assembly parameters

x86dis/oldstable 0.23-6 armhf
  Frontend to libdisasm

z80asm/oldstable 1.8-1 armhf
  assembler for the Zilog Z80 microprocessor

z80dasm/oldstable 1.1.4-1 armhf
  disassembler for the Zilog Z80 microprocessor

pi@raspberrypi:~/in2060 $
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