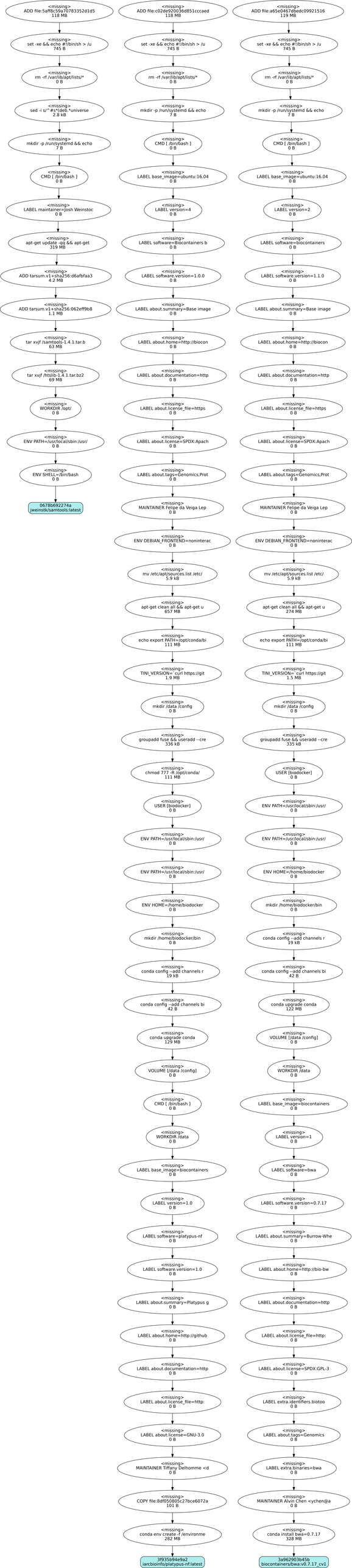
<missing>
ADD file:5aff8c59a70783352d1d5
118 MB
<missing>
set -xe && echo #!/bin/sh > /u
745 B
<missing>
rm -rf /var/lib/apt/lists/*
0 B
<missing>
sed -i s/^ *#s*tdep *universe
2.8 kB
<missing>
mkdir -p /run/systemd && echo
7 B
<missing>
CMD [/bin/bash]
0 B
<missing>
LABEL maintainer=Josh Weinstoc
0 B
<missing>
apt-get update -qq && apt-get
319 MB
<missing>
ADD tarsum.v1+sha256:06afbfaa3
4.2 MB
<missing>
ADD tarsum.v1+sha256:062eff9b8
1.1 MB
<missing>
tar xvjf /samtools-1.4.1.tar.b
63 MB
<missing>
tar xvjf /htslib-1.4.1.tar.bz2
69 MB
<missing>
WORKDIR /opt/
0 B
<missing>
ENV PATH=/usr/local/sbin:/usr/
0 B
<missing>
ENV SHELL=/bin/bash
0 B
0678b692274a
jweinstk/samtools:latest
<missing>
ADD file:c02de9200364b851cccaed
118 MB
<missing>
set -xe && echo #!/bin/sh > /u
745 B
<missing>
rm -rf /var/lib/apt/lists/*
0 B
<missing>
mkdir -p /run/systemd && echo
7 B
<missing>
CMD [/bin/bash]
0 B
<missing>
LABEL base_image=ubuntu:16.04
0 B
<missing>
LABEL version=4
0 B
<missing>
LABEL software=Biocontainers b
0 B
<missing>
LABEL software.version=1.0.0
0 B
<missing>
LABEL about.summary=Base image
0 B
<missing>
LABEL about.home=http://biocon
0 B
<missing>
LABEL about.documentation=http
0 B
<missing>
LABEL about.license_file=https
0 B
<missing>
LABEL about.license=SPDX:Apach
0 B
<missing>
LABEL about.tags=Genomics,Prot
0 B
<missing>
MAINTAINER Felipe da Veiga Lep
0 B
<missing>
ENV DEBIAN_FRONTEND=noninterac
0 B
<missing>
mv /etc/apt/sources.list /etc/
5.9 kB
<missing>
apt-get clean all && apt-get u
657 MB
<missing>
echo export PATH=/opt/conda/bi
111 MB
<missing>
TINI_VERSION= curl https://git
1.9 MB
<missing>
mkdir /data /config
0 B
<missing>
groupadd fuse && useradd --cre
336 kB
<missing>
chmod 777 -R /opt/conda/
111 MB
<missing>
USER [biodocker]
0 B
<missing>
ENV PATH=/usr/local/sbin:/usr/
0 B
<missing>
ENV PATH=/usr/local/sbin:/usr/
0 B
<missing>
ENV HOME=/home/biodocker
0 B
<missing>
mkdir /home/biodocker/bin
0 B
<missing>
conda config --add channels r
19 kB
<missing>
conda config --add channels bi
42 B
<missing>
conda upgrade conda
129 MB
<missing>
VOLUME [/data /config]
0 B
<missing>
CMD [/bin/bash]
0 B
<missing>
WORKDIR /data
0 B
<missing>
LABEL base_image=biocontainers
0 B
<missing>
LABEL version=1.0
0 B
<missing>
LABEL software=platypus-nf
0 B
<missing>
LABEL software.version=1.0
0 B
<missing>
LABEL about.summary=Platypus g
0 B
<missing>
LABEL about.home=http://github
0 B
<missing>
LABEL about.documentation=http
0 B
<missing>
LABEL about.license_file=http:
0 B
<missing>
LABEL about.license=GNU-3.0
0 B
<missing>
MAINTAINER Tiffany Delhomme <d
0 B
<missing>
COPY file:8df050805c27bce6072a
101 B
<missing>
conda env create -f /environme
282 MB
3f935b94e9a2
iarcbioinfo/platypus-nf:latest
<missing>
ADD file:e65e04674dbedc09921516
119 MB
<missing>
set -xe && echo #!/bin/sh > /u
745 B
<missing>
rm -rf /var/lib/apt/lists/*
0 B
<missing>
mkdir -p /run/systemd && echo
7 B
<missing>
CMD [/bin/bash]
0 B
<missing>
LABEL base_image=ubuntu:16.04
0 B
<missing>
LABEL version=2
0 B
<missing>
LABEL software=biocontainers
0 B
<missing>
LABEL software.version=1.1.0
0 B
<missing>
LABEL about.summary=Base image
0 B
<missing>
LABEL about.home=http://biocon
0 B
<missing>
LABEL about.documentation=http
0 B
<missing>
LABEL about.license_file=https
0 B
<missing>
LABEL about.license=SPDX:Apach
0 B
<missing>
LABEL about.tags=Genomics,Prot
0 B
<missing>
MAINTAINER Felipe da Veiga Lep
0 B
<missing>
ENV DEBIAN_FRONTEND=noninterac
0 B
<missing>
mv /etc/apt/sources.list /etc/
5.9 kB
<missing>
apt-get clean all && apt-get u
274 MB
<missing>
echo export PATH=/opt/conda/bi
111 MB
<missing>
TINI_VERSION= curl https://git
1.5 MB
<missing>
mkdir /data /config
0 B
<missing>
groupadd fuse && useradd --cre
335 kB
<missing>
USER [biodocker]
0 B
<missing>
ENV PATH=/usr/local/sbin:/usr/
0 B
<missing>
ENV PATH=/usr/local/sbin:/usr/
0 B
<missing>
ENV HOME=/home/biodocker
0 B
<missing>
mkdir /home/biodocker/bin
0 B
<missing>
conda config --add channels r
19 kB
<missing>
conda config --add channels bi
42 B
<missing>
conda upgrade conda
122 MB
<missing>
VOLUME [/data /config]
0 B
<missing>
WORKDIR /data
0 B
<missing>
LABEL base_image=biocontainers
0 B
<missing>
LABEL version=1
0 B
<missing>
LABEL software=bwa
0 B
<missing>
LABEL software.version=0.7.17
0 B
<missing>
LABEL about.summary=Burrow-Whe
0 B
<missing>
LABEL about.home=http://bio-bw
0 B
<missing>
LABEL about.documentation=http
0 B
<missing>
LABEL about.license_file=http:
0 B
<missing>
LABEL about.license=SPDX:GPL-3
0 B
<missing>
LABEL extra.identifiers.biotoo
0 B
<missing>
LABEL about.tags=Genomics
0 B
<missing>
LABEL extra.binaries=bwa
0 B
<missing>
MAINTAINER Alvin Chen <ychen@a
0 B
<missing>
conda install bwa=0.7.17
328 MB
3a9629033b45b
biocontainers/bwa:v0.7.17_cv1