Starling-May18 Projects/Katarina Stuart/KStuart.Starling-Aug18/Sv3_Genome/Annotation/2020-04-06.BUSCOlong

PDF Version generated by

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or

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Table of Contents

2020-04-06.BUSCOlong 2



Katarina Stuart (z5188231@ad.unsw.edu.au) - Sep 30, 2020, 11:39 PM NZDT

BUSCO-long

#!/bin/bash

#PBS -N 2020-05-05.BUSCOLong.pbs

#PBS -I nodes=1:ppn=40

#PBS -I mem=56gb

#PBS -I walltime=200:00:00

#PBS -j oe

#PBS -M katarina.stuart@student.unsw.edu.au

#PBS -m ae

GENOMEDIR=/srv/scratch/z5188231/KStuart.Starling-Aug18/Sv3_Genome/Sv3.4_GenomeAnnotation/data_2020/genome_assembly GENBASE=Sturnus vulgaris 2.3.simp

GENOME=\${GENOMEDIR}/\${GENBASE}.fasta

OUTDIR=\${GENBASE}.busco

MODEL=/srv/scratch/z5188231/KStuart.Starling-Aug18/Sv3_Genome/Sv3.4_GenomeAnnotation/data/BUSCO.2018-08-21/aves_odb9

module purge

module load python/3.7.3 blast+/2.2.31 hmmer/3.2.1 augustus/3.3.2 emboss/6.6.0 busco/3.0.2b

export AUGUSTUS_CONFIG_PATH=/srv/scratch/z5188231/programs/augustus

export BUSCO_CONFIG_FILE=/home/z5188231/busco/3.0.2b/config/config.ini

cd /srv/scratch/z5188231/KStuart.Starling-Aug18/Sv3 Genome/Sv3.4 GenomeAnnotation/data 2020/buscolong

mkdir \$OUTDIR

cd \$OUTDIR

python3 /apps/busco/3.0.2b/scripts/run_BUSCO.py -o \$GENBASE -i \$GENOME -I \$MODEL --cpu 40 -m genome --tmp \$TMPDIR --long

Ran once (end 15/04/20), Timed out after 200 hrs. resubmitting to Katana with the --restart tag

3rd run restart - 24/04/2020

Resubmmitted with new genome version 5/5/20.

python3 /apps/busco/3.0.2b/scripts/run_BUSCO.py -i ./GCF_001522545.3_Parus_major1.1_genomic.fna -o Parus_major1.1_genomic.busco -m genome -l \${BUSCOSET}/aves_odb9/ -c 32 -f

$BUSCOSET = /srv/scratch/z5188231/KStuart. Starling-Aug18/Sv3_Genome/Sv3.4_GenomeAnnotation/data/BUSCO.2018-08-21$

submitted job to katana 22/8/18

[z5188231@katana scripts]\$ qsub 2018-08-22.1.BUSCOLong.pbs 6679258.katana.science.unsw.edu.au

timed out after 200 hrs. resubmitting to Katama with the --restart tag

[z5188231@katana scripts]\$ qsub 2018-08-22.1.BUSCOLong.pbs 6683522.katana.science.unsw.edu.au