2022/3/3 22:48 个人简历



Ryougi-yukiro

Python / Perl /R

▲ 基本信息

姓名 朱铭佳

学历 硕士

毕业院校 兰州大学

毕业年份 2022

┗ 联系方式

手机 18656474816

邮箱 z980907mj@gmail.com

个人主页 ryougi-yukiro.github.io

Github github.com/Ryougi-

vukiro

☆ 技能点

Perl

Markdown/htm

I

Python

R

▲ 技术栈

基因组 Nextdenovo BUSCO

转录组 clusterprofiler E2P2

enogg-mapper

其它 Population

ヘ 个人微信



■ 教育经历

• 兰州大学 - 生态学 (本科) 2015-2019

Learn biology and ecology, use free time learn R script and ggplot2 package during 2018-2019.

兰州大学 - 生物信息学(硕士) 2019-2022

work in current lab, and do a project about a QTP alpine herb.

🖨 工作经历

• Liu Lab - 研究生 (课题发表)

Learn Perl and Python and use the lastest software or tools to build new pipelines and then do project

• University - 课程内容(户外实习)

During 2016, profs led us to tianzhu and xiama, then led us to identify wild animals and plants for one month

🖺 个人项目

• 基于hexo搭建的个人博客

Use hexo to build personal blog, which is mainly used to record personal life and some laboratory routines; otherwise, some skills were also recorded

• 基于 clusterprofiler的GO富集流程

Use cluserprofiler 4.0 published in 2021 years, to build a pipeline for GO enrichment, which is better than other pipeline lab builded before

• 基于WGDI的多倍化分析流程

Use WGDI preprinted on bi'oArxiv, to build a pipelin for WGD events, which is betters than MCSCANX and MCSCAN. In addition, the pipeline could plot ks density together, which is easier than use R.

Others

gene family identified, transcriptome, SNP and Indel, SVs. And read article about single-cell RNA sequencing.

▼ 奖项与证书

 LZU of Life Sciences Second Prize and Field Internship Individual First Prize

During 2016 field internship, i got individual first prize (one of five people);During 2016-2018,i got LZU of Life Sciences Second Prize(6 people of 49 students in a class)

participating authors of two SCI article

During Liu Lab, i partipating two published articles (one is IF=4.0; one is IF=2.0). In addition, i partipated in an article of NC review, but no authors.

The first of SCI article under review(the second rounds)
Since 2020, i took a project about a QTP alpine herb with its omics sequencing, then writed an article and conrtibuted it to Molecular Ecology(under the second round of review)

❷ 自我评价

遑遑二十载, 书剑两无成。