Quartet Data Portal User Guide for

RNA-Seq Quality Assessment Report

1. Preparation

| QDP Address | http://chinese-quartet.org/ |
|-------------|--|
| OSSUtil | https://docs.chinese-quartet.org/tools/ossutil/ |
| Input files | The file name suffix should be: |
| | The data format that can be analysed online are pair-end |
| | FASTQ files, with suffixes |
| | of _R1.fastq.gz and _R2.fastq.gz or _R1.fq.gz and _R2.fq.gz. |
| | Your dataset must contain four groups D5, D6, F7 and M8, |
| | with at least 2 technical replicates in each group. |

2. Log on to QDP

2.1 QDP account

1)If you have a request to receive reference materials, please go to <u>http://chinese-quartet.org/#/materials</u> to request the reference materials, and the Quartet team will contact you to send the reference materials and send you a link to register your account.



2) If you do not have a request for reference materials, please send an email to **quartet@fudan.edu.cn** to request an account.

2.2 Log on to QDP

Address: http://chinese-quartet.org/



3. Upload RNA sequencing data

3.1 Select Transcriptomics> Upload Your Data> Run

| omics Transcriptomics Proteomics N | Aetabolomics | 0 |
|---|--|---|
| Upload Your Data Latest Upload your data to the data repo, and then you can use it in the pipeline. | RNA-Seq QC for Quartet vo.2.1 RNA Sequencing Quality Control Pipeline for Quartet. | O QC Report for Quartet RNA-Seq v0.22 Generate the QC Report for Quartet RNA- Seq data. |
| | Run Help Alignment | |
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3.2 Click the symbol called "Register & Upload Your Data"

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|---|---|---|------------------------------|-----------------------------|
| Home Quality Assessment Image: Comparison of the sessment Image: Comparison of the se | All Your Data | Help © View Your Data | 土 Metadata Template | Register & Upload Your Data |
| Total Started Finishe | Failed Please enter the name of dataset Q | | | 3 |
| genomics | Uploading Path © oss://quartet-data-portal/data/guoxr-22/practice/genomics/ | Started 2022-09-07 16:50 | Finished | New Token V Check |
| speed_test1 genomics | Uploading Path oss://quartet-data-portal/data/guoxr-22/speed_test1/genomics/ | Started 2022-09-02 10:19 | Finished 2022-09-06 15:02 | New Token Check |
| MGL_Quartet_test | Uploading Path © oss://quartet_data_portal/data/guoxr-22/MGi_Quartet_test/genomics/ | Started 2022-08-11 22:51 | Finished 2022-09-02 10:20 | New Token V Check O |
| | | | | < 1 > 30 / page ~ |

3.3 Fill in Dataset Name, DataType select Transcriptomics, and click Submit

| Home / Quality Assessment / Al | | Notices The uploading path will au access the same path as b | tomatically generate after you submit the form. If you want to before, please enter the same information. |
|--------------------------------|--|--|--|
| Total Started Finished | Failed Please enter the name of dataset Q. | Your Data Will Be Uploaded To | |
| practice | Uploading Path ◎ | 达 oss://quartet-data-portal/data/g | uoxr-22/MGI_Quartet_test/transcriptomics/ |
| genomics | ess://quartet-data-portal/data/guoxr-22/practice/genomics/ | * Dataset Name | * Contact Email |
| speed_test1 | Uploading Path © | MGI_Quartet_test | 🖾 guoxr22@m.fudan.edu.cn |
| genomics | oss://quartet-data-portal/data/guoxr-22/speed_test1/genomics/ | * Manager | * DataType |
| MGI_Quartet_test | Uploading Path © | 유 guoxr-22 | Transcriptomics |
| genomics | oss://quartet-data-portal/data/guoxr-22/MGI_Quartet_test/genomics/ | Description | |
| | | Please enter the description | |
| | | | Cancel |

3.4 Click the symbol called "New Token". Copy the autorizedCode, which is for subsequent ossbrowser login

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| Total Started Finished Failed Please entor Image: Started Finished Failed Please entor Image: Started Uploading Path © Started Started Finished Finished Image: Started Uploading Path © Started Started Started Finished Finished Image: Started Uploading Path © Started/Starte-tata-portal/starte Started/Starte-tata-portal/starte Finished Finished Image: Started Uploading Path © Started/Starte-tata-portal/starte Started/Starte-tata-portal/starte Finished Finished Image: Started Uploading Path © Started/Starte-tata-portal/starte Started/Starte-tata-portal/starte Finished New Token Image: Started Uploading Path © Started/Starte-tata-portal/starte Started/Starte-tata-portal/starte Finished New Token Image: Started Uploading Path © Started/Starte-tata-portal/starte Started/Starte-tata-portal/starte Finished New Token Image: Started Uploading Path © Started/Starte-tata-portal/starte Started/Starte-tata-portal/starte Finished New Token Image: Started </td <td>pload Your Data</td> | pload Your Data |
| MGL_Quartet_test Iranscriptomics Uploading Path © css:/guartet-data-ports/data/ genomics "eyjp2C1611NUU/SOVG1jRHk0RVVXDWS0YM12UXp2Q3pDVkMT1wic2 "cyjp2C1611NUU/SOVG1jRHk0RVVXDWS0YM12UXp2Q3pDVkMT1wic2 "cyjp2C1611NUU/SOVG1jRHk0RVVXDWS0YM12UXp2Q3pDVkMT1wic2 "cyjp2C1611NUU/SOVG1jRHk0RVVXDWS0YM12UXp2Q3pDVkMT1wic2 "cyjp2C1611NUU/SOVG1jRHk0RVVXDWS0YM12UXp2Q3pDVkMT1wic2 "cyp2C1611NUU/SOVG1jRHk0RVVXDWS0YM12UXp2Q3pDVkMT1wic2 "cyp2C1611NUU/SOVG1jRHk0RVVXDWS0YM12UXp2Q3pDVkMT1wic2 "cyp2C1611NUU/SOVG1jRHk0RVVXDWS0YM12UXp2Q3pDVkMT1wic2 "cyp2C1611NUU/SOVG1jRHk0RVVXDWS0YM12UXp2Q3pDVkMT1wic2 SX12TLUDdOMI poMRLKRP0Q27jazhf2jFdyS1Rvm0RkMTVKdWS0ULDEp (WSZ0Q60RHid)T1dFXYT7WK0SUB3CEX2KVp2Z8Ykp2ZXWT TdQ%mQVCMRdg1z1dFXYT7WK0SUB3CEX2KVp2Z8Ykp2ZXWT TdQ%mQVCMRdg1z1dVXTSFEXM010NRDEpkomRNHTVKdWFYK0X SNNQ2WzcUg4VWp2S1Z4XF1SX1yWRS0k0LRMFVKXdy ErReit1N2Z1MXCVM1SSkg3MXU/SHFKSm13MmcodC1XRF6VYtBN1/y1 erst/guartet-data-ports/data WSZ7060FSIN1Biccno2j271rE1VMF32Kb3MrVHDNVdXV Finished New Token | 0 |
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| speed_test1 Uploading Path Uploading Path Uploading Path Vert Vert Vert Vert Vert Vert Vert Vert | ✓ Check |
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| MGL_Quartet_test Uploading Path © Ca2x1v2xf1VFkQ1L0MipS2daYTNFR1kgKGR02FhttMpUdmF3dRVL055 Finished New Token genomics oss://guartet-data-portal/data mR3MGVIS iox2INRR1hkRTUSREx1NF2KaD2h0XNreVolbHBnc18U01p0 2022-09-02 10:20 New Token | Check |

3.5 Please follow the ossutil document to upload your data

ossutil is a user friendly command line tool to upload data to the Quartet Data Portal. It supports following operating systems: Windows, Linux, and macOS. You can download and install the ossutil version that best suits your requirements.

https://docs.chinese-quartet.org/tools/ossutil/

4. Run RNA-Seq app

4.1 Select Transcriptomics> RNA-Seq QC for Quartet> Run

| | Metabolomics | () H |
|---|--|--|
| Upload Your Data Intent Upload your data to the data repo, and then you can use it in the pipeline. | RNA-Seq QC for Quartet vo.2.1 RNA Sequencing Quality Control Pipeline for Quartet. | QC Report for Quartet RNA-Seq v0.2.7 Generate the QC Report for Quartet RNA- Seq data. |
| Run O Help Share | Run Help description | Run Help Share |
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4.2 Click the symbol called "Start Assessment"

| Quality Assessment ② Reference Materials & Multiomics Data 🖻 Quality Assessment ③ Reference Datasets | 🕜 💬 💽 Guo 🗛 |
|---|---------------------------|
| Home / Quality Assessment / RNA-Seq Data / RNA-Seq Pipeline | |
| RNA-Seq Pipeline | Help Start Assessment |
| Total Running Finished Failed Iizhihui/quartet-rnaseq-qc-v0.2.1 V | 0 |
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| | 1 General Information | 2 Assessment Parameters | 3 Submit | | | |
| | * Assessment Name : | MGI_Quartet_test_new | | | | |
| | * Description : | MGL_Quartet_test_new | | | | |
| | | * | | | | |
| | * Started Time : | 2022-10-14 11:04:37 | | | | |
| | * Group : | Quartet Team | | | | 0 |
| | * Assessment App : | lizhihui/quartet-maseq-qc-v0.2.1 v | | | | |
| | | Reset | | | | ? |
| | | | | | | 0 |

4.3 Fill in Assessment Name and Description. Then click Next.

4.4 Fill in Sample information

Sample ID Fill in the numbers; Select the read files and click upload. The parameters other than read file can be selected as default files or you can upload your own files. Assessment Parameters can be downloaded and checked.

| General Information 2 Assessment Parameters 3 Submit | |
|---|---|
| * Sample ID ③ | |
| * Read1 Files | |
| + Select Files D5_S16_R1_001.fastq.gz × | |
| * Read2 Files | |
| + Select Files D5_S16_R2_001.fastq.gz × | |
| * FastqScreen Conf File | |
| oss://pgx-reference-data/fastq_screen_reference/fastq_screen.conf v | 0 |
| * References of Fastq Screen | |
| oss://pgc-reference-data/fastq_screen_reference/ | ? |
| | |

| | | G (V) G | eneral Information Assessment P | Parameters | 3 Submit |
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| Grou | up | | Quartet Team | | Show Jobs |
| Appl | old | | fd63809bc013bec3664b072d3465ba75 | | Show Assessments |
| Desc | jectName cription | | MGI_Quartet_test_new MGI_Quartet_test_new | | |
| Appl | Name | | lizhihui/quartet-rnaseq-qc-v0.2.1 | | Run Assessment |
| | | (m) | read1 | | read? |
| 48 | sample_id 89eb0af-bdca-499e-996c- | oss://qu | artet-data-portal/data/guoxr-22/MGI_Quartet_test/transcriptomic | cs/D5_S16_R1_001.fastg.gz | oss://guartet-data-portal/data/guoxr-22/MGI_Quartet_test/ |
| 48 c1 | sample_id 89eb0af-bdca-499e-996c- 1d6ea07feba | oss://qu | artet-data-portal/data/guoxr-22/MGI_Quartet_test/transcriptomic | cs/D5_S16_R1_001.fastq.gz | oss://quartet-data-portal/data/guoxr-22/MGI_Quartet_test/ |
| 48 c1 | sample_id 89eb0af-bdca-499e-996c- 1d6ea07feba | oss://qu | artet-data-portal/data/guoxr-22/MGLQuartet_test/transcriptomi | cs/D5_S16_R1_001.fastq.gz | oss://quartet-data-portal/data/guoxr-22/MGLQuartet_test/ |
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| 48 c1 | sample_id 89eb0af-bdca-499e-996c- 1d6ea07feba | oss://qu | artet-data-portal/data/guoxr-22/MGL_Quartet_test/transcriptomi | cs/D5_S16_R1_001.fastq.gz | oss://quartet-data-portal/data/guoxr-22/MGI_Quartet_test/ |

4.5 Click the symbol called "Run Assessment"

4.6 Check the status of Run Assessment

| Quartet ② Overview 四 Reference Materials 。 | å, Multiomics Data 🛛 🕅 Quality | Assessment 🛛 🕢 Reference Data | asets | 0 💬 💽 G | A ol |
|---|--------------------------------|-------------------------------|--------|------------------|--------|
| Home / Quality Assessment / RNA-Seq Data / RNA-Seq Pipeline | | | | | |
| RNA-Seq Pipeline | | | 0 | Help Start Asses | sment |
| Total Running Finished Failed lizhihul/quartet-maseq-q | c-v0.2,1 v | | | | 2 |
| aaaaaaaa | Started | Finished | Status | View | Delete |
| aaaaaaaa | 2022-10-14 18:40 | 0000-00-00 00:00 | | | |
| MGLQuartet_test_new | Started 2022-10-14 11:04 | Finished 2022-10-14 19:50 | Status | View | Delete |
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5. Results and Report

5.1 Select Transcriptomics> QC Report for Quartet RNA Seq-Run

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|---|--|---|
| Upload Your Data Latest Upload your data to the data repo, and then you can use it in the pipeline. | RNA-Seq QC for Quartet v3.2.1 RNA Sequencing Quality Control Pipeline for Quartet. | QC Report for Quartet RNA-Seq O222 Generate the QC Report for Quartet RNA-Seq data. |
| | | Run G Help Share |
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|--------------------------------------|--------------------------|-------------------|------------------|----------------|----------------|--------|---------------|
| ome / Quality Assessment / RNA-Seq I | Data / RNA-Seq QC Report | | | | | | |
| RNA-Seq QC Report | | | | | | | New QC Report |
| Report Name | Category | Report Tool | Version | Created At | Finished At | Status | Action |
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| .5 Follow the St | cps as pi | ompicu | | | | | | |
|---|---------------------|--|--|---|--|--|--|--------------|
| Quartet O Overview | A Reference Mat | New QC Report | | | | | | |
| Home / Quality Assessment / RNA-Seq Da | ata / RNA-Seq QC Re | Step1: Choose Report | Step2: Load | I Project & Submit | | | | |
| Report Name | Category | Notices Please select the est | xpected quality a | sessment tool from the | following list, after selecting th | ne tool, the correspo | nding guidance message v | will appear |
| | | Category N | lame | | Description | | Version | Githul |
| | | | C Report for Qua | rtet RNA-Seq | Generate the QC Report for Q | uartet RNA-Seq data | v0.2.2 | C |
| | | | | | | | | |
| Cuartet O Overview Home / Guality Assessment / RNA-Seq DC RNA-Seq OC Report | A Reference Mat | New QC Report Step1: Choose Report | Step2: Load | Project & Submit | | | | |
| Report Name | Category | Notices Please select the c results file generate whether the corres | ompleted project ad by the `RNA-S ponding analysis | from the dropdown mer eq QC for Quartet` app has been completed pro | u below and the QC report too and generate a QC report. If yo perly. | l will automatically e ou do not find the exp | xtract the required data fro pected project, please che | om the ck |
| | | MGI_Quartet_test_new | | | | | | |
| Quartet @ Overview मण्डहाइ Home / Quality Assessment / RNA-Seq Dr | A Reference Materia | als A Multionics Date | a 🖻 Quality | Assessment 🥝 R | leference Datasets | | Cancel ⑦ 으 C | Subm Guo |
| RNA-Seq QC Report | | | | | | | New QC | Report |
| Report Name | Category | Report Tool | Version | Created At | Finished At | Status | Action | |
| MGI_Quartet_test_new | ٢ | quartet-rseqc-report | v0.2.2 | 2022-10-14 22:00 | 2022-10-14 22:00 | \odot | © Result 止 Do | ownload |
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5.3 Follow the Steps as prompted