# Submission Infrastructure

Submission UI URL: https://submissions-ui.mlcommons.org/

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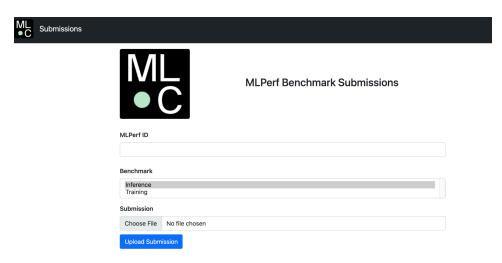
## **Objectives**

- Make a submission UI that allows submitters to upload tarballs containing their results. Allow submitters to run verification tests and preview their results as well
- Deploy infrastructure and make the UI available to submitters

# **Usage**

I'll briefly describe the process to submit your results.

- Before the submission process begins you will receive an email containing an ID (Submitter ID). You will need this in order to upload your results. (This information will be sent to the email you provided in the registration survey). A link to access the web UI will also be provided.
- 2. When you access the web UI, you should see a page like this:



- 3. Using this interface, enter the ID that was provided to you, select the appropriate benchmark, and upload from your computer a tarball containing the results of running the inference benchmark. Take into account the <u>structure</u> of the submission folder. The tarball should be computed in the following way (or in a way that is equivalent to):
  - o tar -cvzf <tarfile\_name>.gz <submission\_folder>

Example submission structure:

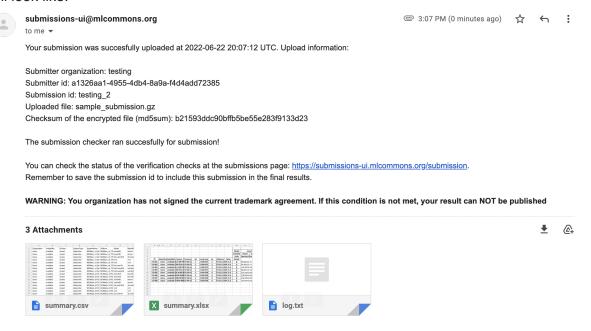
- root\_folder (or any name you prefer)
  - Closed
    - <submitter\_name>
      - code
      - . ...
      - results
      - systems
  - Open
    - <submitter\_name>
      - code
      - ...
      - results
      - systems

For that case, you should tar the **root\_folder** 

After you upload the tarball, a submission\_id will be generated and sent to you by email and will be displayed in the UI. The submission checker will be run as well and the results will be sent to you along with the submission\_id. By default, you can upload as many tarballs as you want, but only the last one that passes the submission\_checker will be included in the final results.

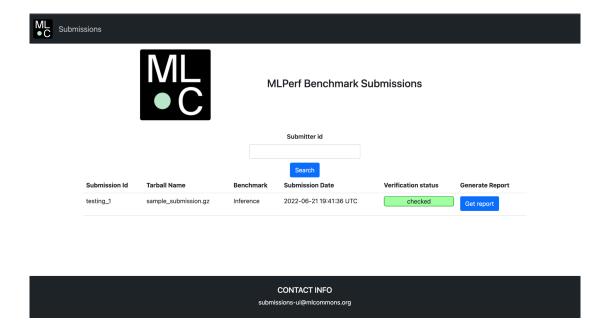
- If additional tarballs need to be chosen, the submitter needs to contact Pablo with the details (with tarball ids)
- The tarballs you choose should have a different <submitter\_name> in the previous folder structure. E.g: Google, Google\_1 or Qualcomm\_edge, Qualcomm\_datacenter.
- If you want to request multiple submitter\_ids, the submitter needs to contact Pablo. The tarballs you choose should have a different <submitter\_name> in the previous folder structure.
- 4. You will receive an email that contains the results of the submission checker. This will tell you if the submission passed or not. Here is what an email for a successful submission

#### will look like:



Here is a description of the files you'll receive for a successful submission:

- **summary.csv**: Here are the results for all your systems summarized in a csv file. You'll only get this if your submission was successful.
- summary.xlsx: Here are the results for all your systems summarized in a spreadsheet. Similar to what the final report should look like. You'll only get this if your submission was successful.
- o **log.txt**: The log output of the submission checker. This checks if your submission is valid.
- 5. In the upper left corner of the interface, you will find a link to the submissions page. (Also notice that the image of the MLCommons logo is a link that takes you back to the main page). There you will be able to see the submission that is currently included in the final results. This page should look like this:



## **Frequent Issues and Questions**

 The loading bar gets stuck at 100% when uploading a submission. In this case, the file is uploaded correctly, but some frontend error prevents the submission checker from running.



If you get this issue you can try one of the following to solve the issue:

- 1. Wait for a couple of seconds (<1 min). It can take some time to complete the request
- 2. This issue doesn't occur very often, therefore you can try to upload your submission again.
- 3. If the error persists, you can:
  - a. Go to the submissions page
  - b. Use your submitter id to look for your submissions.
  - c. You should see the following button that you can use to run the submission checker:

uploading

Check submission