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Genomic Sequencing and Analysis Facility Online Submission Portal

WE HAVE A NEW LIMS SO BE SURE TO USE IT FOR SUBMITTING NEW REQUESTS

To submit a new project or get detailed pricing information, please use the online job submission portals.

1. Sign up for an account

- Anyone at the University of Texas at Austin with a "utexas.edu" or a "austin.utexas.edu" email address will need to email the lab to request and account
 - Please email info@utgsaf.org to start the process
- All others can use the self registration portal by [following this link](#)
 - This includes outside academic labs, commercial labs and University of Texas affiliates not at Austin
- **IMPORTANT: Both external and internal users will receive TWO emails from either Lockbox or Salesforce, they might go to spam. You MUST set up your security questions before attempting to login**

2. Once you have an account use the [GSAF Requester Portal](#)

- Important notes on sample submissions
 - [See this web page for sample input guidelines.](#)
 - Metagenomics and TagSeq
 - [DNA/RNA concentrations should be normalized before submission](#)

3. [Job Info TOOL](#)

- Use this link to check job details without logging into portal
- You will not be able to access files from this link
- This resource is great for researchers who just need to check the parameters of previous projects

4. [Sequencing Information](#)

- Link provides access to FastQC and MulitQC reports
- Tag-Seq jobs-TagSeq Deduplication Analysis

5. [Online Job Estimator](#), quick estimates without putting in a project request. This is only for estimates, for a formal quote you need to submit a project. Please note that not all services will be in the estimator.

Browse the wiki: