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# Life Science Analytic Framework Overview

**Matt Becker**  
Advisory Life Science Industry Consultant  
SAS

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## Executive summary

The SAS Life Science Analytics Framework (LSAF) is a comprehensive platform for data governance and analysis in the life science industry. It provides end-to-end data management, from standards and study metadata management to reproducibility and traceability. LSAF is an open and flexible framework that can connect to other platforms using APIs, support open-source analytics like R and Python, and ingest various data sources. LSAF provides a comprehensive programming and version control environment with a customizable central repository, allowing customers to set up their own folder structure. It also offers collaboration, workload management, high availability, and a high-performance environment. The workspace status feature allows users to compare the status of files in their workspace with those in the repository. Audit trails are a crucial component of program management, providing a record of all modifications and actions taken within a program, allowing users to track changes and identify potential issues. Workflows are an essential part of LSAF, especially when it comes to managing the development, testing, and production of programs. These workflows involve multiple stages, including program development, quality control, and final sign-off or promotion.

# Features of LSAF



## Why LSAF?

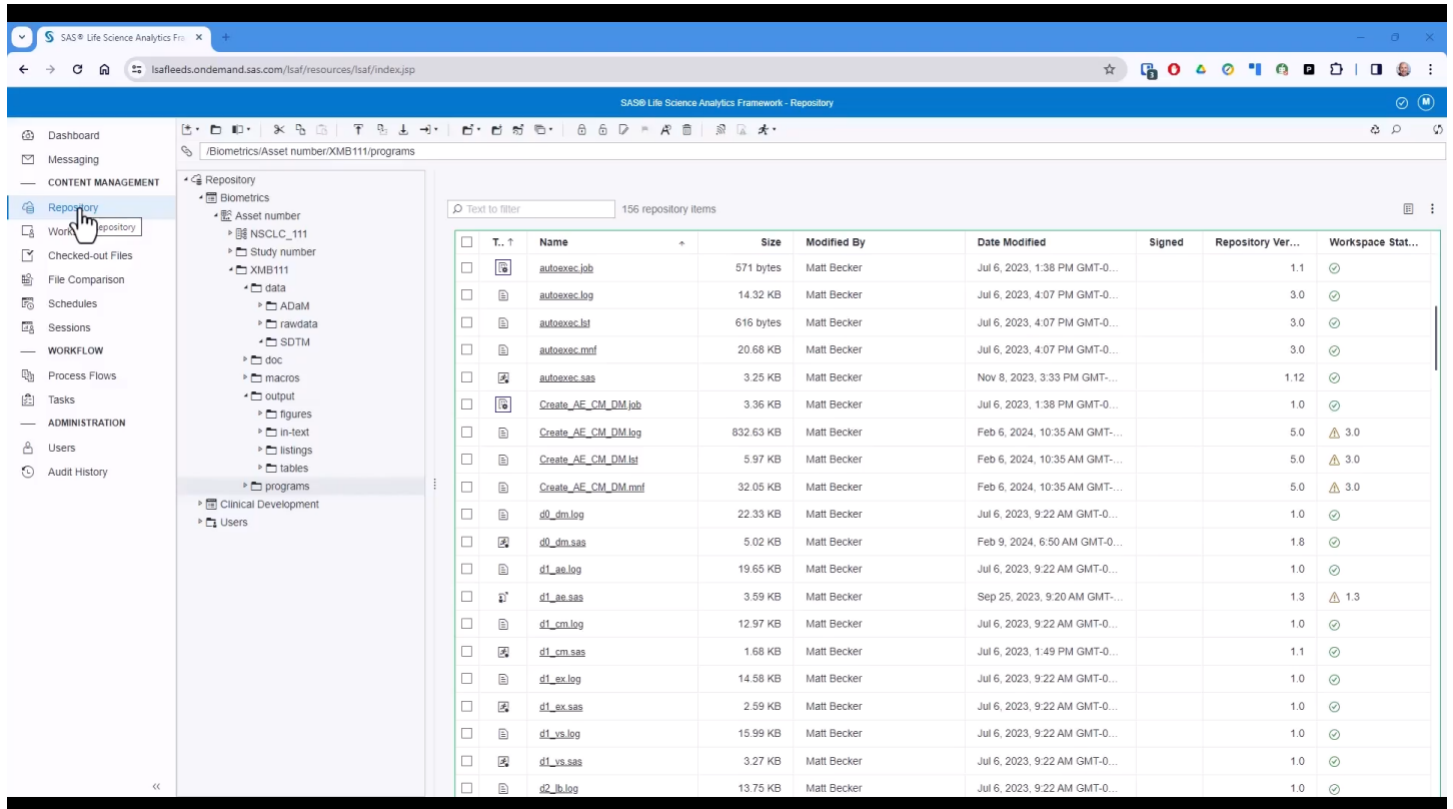
- **Single environment versus silos**  
Integrated platform for **data processing, data review, standards management, study metadata management, data analysis, programming and submission.**
- **Enabled regulatory compliance**  
Controlled **validated environment**, with complete **audit history** and **robust version controls** for managing the change history on all files.
- **Internal & External Collaboration**  
**Workflow capabilities, the content repository and permission models** allows customer to collaborate internally, but also to **share data** and analysis with CROs. It supports business processes and drives **automation.**
- **Standards & Study metadata management**  
Organize and integrate **Study metadata** and **Global Standards** from external **MDR systems.**
- **Reproducibility & Traceability**  
**Job concepts** and **manifest files** provides **full visibility** and allows for an analysis to be reproduced easily in the future. Job concept allows also for non-SAS programmers to run SAS programs via user friendly manner. **DefineXML** files can be made by a click of a button. Comparison, Adherence and Impact Assessment Reporting 'out of the box'.
- **Open & flexible framework**  
Even though an interactive programming editor is part of the solution, users can program in the environment of their liking **SAS, R, Python, etc.** The system is also open to integrate with Pinnacle 21, Jreview, Document Management systems and other solutions using WebDAV and very rich **JAVA** and **SAS Macro APIs.**

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SAS Life Science Analytics Framework (LSAF) is a single platform that provides end-to-end data governance and analytics solutions for the life science industry. With LSAF, users can easily govern, analyze, and submit data to regulatory authorities. The platform offers a validated environment that enables regulatory compliance, internal and external collaboration, and standards and study metadata management. LSAF also provides reproducibility and traceability, a central repository, and support for open-source analytics like R and Python. With LSAF, users can connect to various platforms using APIs, making it an open and flexible framework for data management and analytics.

# LSAF workspace and version control



The LSAF environment can be accessed via a browser and the repository view is where the final data, programs, and submission-type projects reside. The tree structure is customizable, and customers can set it up according to their preferences. Access control is also possible, allowing specific departments to see specific information. For instance, medical writers can be granted read-only access to the output folder for a particular study.

# Programming in LSAF

The screenshot displays the SAS Life Science Analytics Framework (LSAF) File Comparison tool. The interface is split into two panes for comparison. The left pane shows the workspace version of the file 'd0\_dm.sas' located at 'Workspace/Biometrics/Asset number/XMB111/programs'. The right pane shows the repository version of the same file. A table of modifications is shown between the two versions, with a red highlight on the entry for '27 \*\*\* 2/9/24 Made additional changes per QC findings'. The code editor below shows the SAS program code, with differences between the two versions highlighted in red. The code includes metadata, a data step, and a macro definition.

```
1 *****;
2 *** Program:      \biometrics\XMB111\programs\d0_dm.sas
3 *** Programmer:   Matt Becker
4 *** Date Created: 28Jan2010
5 ***
6 *** Input :       SDTH DM, DS, VS
7 ***
8 *** Output:       Derived DM dataset
9 ***
10 *** Purpose:      To create the derived DM dataset
11 ***
12 *** Comments:
13 ***
14 *** Software:     SAS 9 (Windows)
15 ***
16 *** Modifications:
17 ***
18 *** Date          Programmer      Description
19 *** -----
20 *** 04/27/22
21 *** 04/28/22
22 *** 8/15/23
23 *** 8/25/23
24 *** 10/6/23
25 *** 01/22/24 Added new dataset
26 *** 02/6/24 Comment
27 *** 2/9/24 Made edit per QC findings
28 *** 2/9/24 Made additional changes per QC findings
29 *****;
30 data dm;
31 set &randata._dm(drop=race);
32 length age attr _x_race dmtcn 8 _x_racecat $70 domain $2 usubjid $25 subjid $14 ageu brthotc dmtc $10 race $100 arm
33 country $3 sexn 4;
34 format brthotn dmtcn date9.;
35 domain='DM';
36 arm='Nicardopine';
37 armcd='A';
38 country='USA';
39 studyid='XMB111';
40 usubjide=trim(left(studyid)||'-'||trim(left(site)))||'-'||trim(left(randomno)));
41 subjid=trim(left(site)||'-'||trim(left(randomno)));
42 brthotc=put(brthotn,yyymmdd10.);
43 dmtcnday;
44 dmtc=put(dmtcn,yyymmdd10.);
45 ageround=(intck('month',brthotn,dmtcn) - (day(dmtcn) < day(brthotn))) / 12, 1);
46 ageu=Years;
47 attr=(0<age<2.9)*1 + (3<age<6.9)*2 + (7<age<11.9)*3 + (12<age<17.9)*4;
48 if attr not in(1, 2, 3, 4) then put "Age does not fit in any age group" usubjid;
```

LSAF provides a comprehensive environment for programming and version control. The workspace status feature allows users to check the status of files in their workspace, make changes, and compare their version to the repository version. The editor in the submission engine has additional tabs for submission inputs and outputs, which show the files, data, and macros used in the program and the output created. LSAF supports both SAS and R programming languages, and users can run programs in the same environment. Finally, users can check in their changes to the repository for the final version.

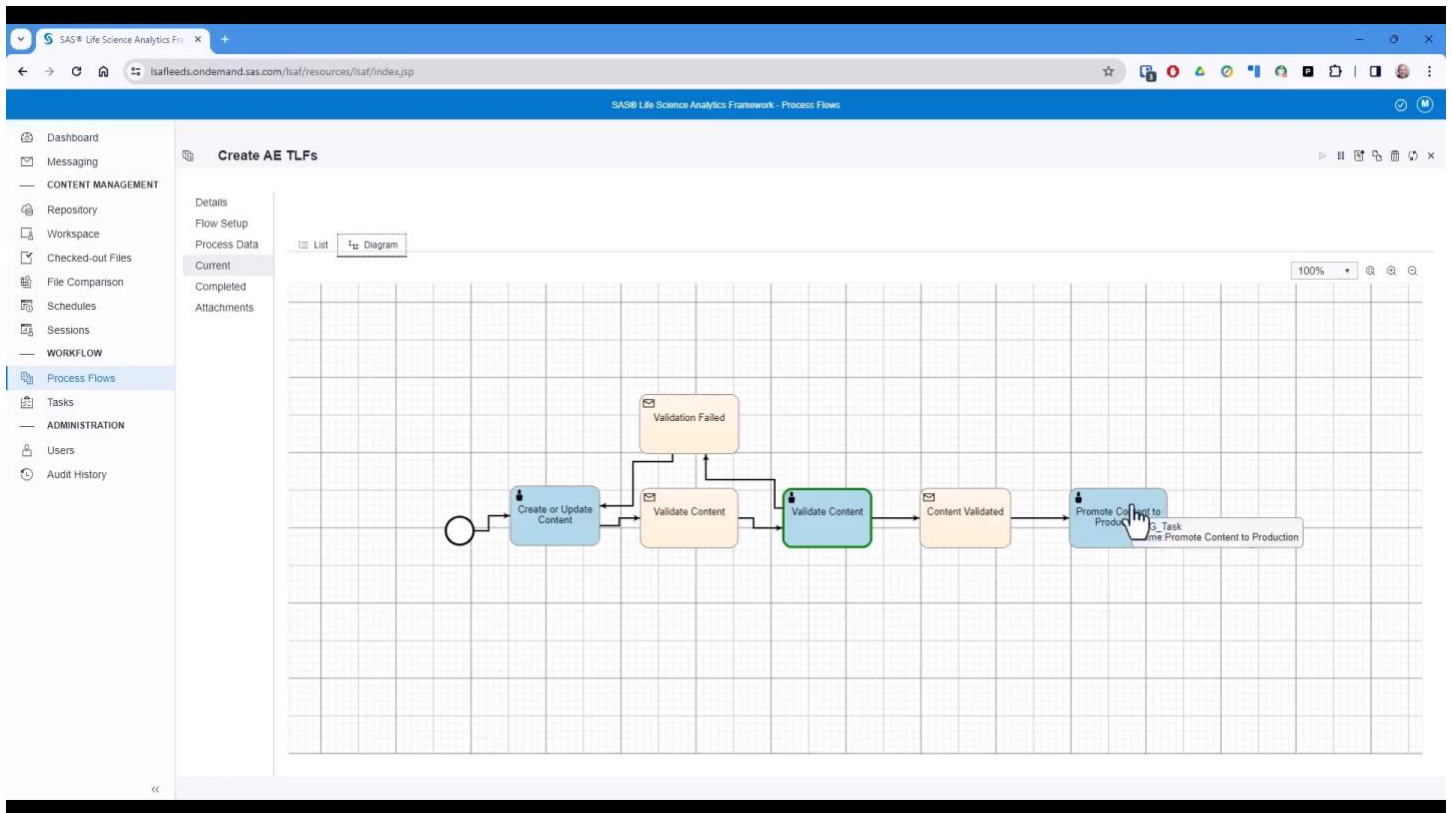
# Jobs in LSAF

The screenshot displays the SAS Life Science Analytics Framework (LSAF) interface. The browser address bar shows the URL: `lsafeeds.ondemand.sas.com/lsaf/resources/lsaf/index.jsp`. The page title is "SAS® Life Science Analytics Framework - Repository". The left sidebar contains navigation options: Dashboard, Messaging, CONTENT MANAGEMENT, Repository (selected), Workspace, Checked-out Files, File Comparison, Schedules, Sessions, WORKFLOW, Process Flows, Tasks, ADMINISTRATION, Users, and Audit History. The main content area shows the "Jobs" section for a job named "Create\_AE\_CM\_DM.job(v1.0)". A search bar indicates "5 submissions". Below the search bar is a table with the following columns: Run Status, Date Submitted, Version, Run-as User, Submitted By, Date Completed, Session Status, and Run Information. The table contains five rows of data, all with a status of "Job completed with program...".

Run Status	Date Submitted	Version	Run-as User	Submitted By	Date Completed	Session Status	Run Information
<input type="checkbox"/>	Feb 6, 2024, 10:35 AM GMT...	1.0	Matt Becker	Matt Becker	Feb 6, 2024, 10:35 AM GMT...		Job completed with program...
<input type="checkbox"/>	Feb 6, 2024, 10:25 AM GMT...	1.0	Matt Becker	Matt Becker	Feb 6, 2024, 10:25 AM GMT...		Job completed with program...
<input type="checkbox"/>	Jan 22, 2024, 10:47 AM GMT...	1.0	Matt Becker	Matt Becker	Jan 22, 2024, 10:47 AM GMT...		Job completed with program...
<input type="checkbox"/>	Jul 6, 2023, 4:07 PM GMT-0...	1.0	Matt Becker	Matt Becker	Jul 6, 2023, 4:07 PM GMT-0...		Job completed with program...
<input type="checkbox"/>	Jul 6, 2023, 1:39 PM GMT-0...	1.0	Matt Becker	Matt Becker	Jul 6, 2023, 1:39 PM GMT-0...		Job completed with program...

In LSAF, jobs are used to create drive data sets, tables, listings, and figures. A job is a set of programs that run in a specific order, which can be SAS or R programs. One of the new features of LSAF is the impact analysis, which shows if any component of the job has changed since it was last run. The manifest file is a key component of the job, which shows the details of the job, the programs used, the inputs, and the outputs. LSAF also has search functionality and an audit trail, which stores all the information about modifications and comments made to the job.

# Workflows in LSAF



Workflows are an essential part of LSAF, and they are commonly used for dev-test prod. The process involves developing a program, QC-ing it, and then promoting it. In LSAF, workflows are assigned to individuals, and tasks are completed as they progress through the workflow. The progress of the workflow can be tracked, and tasks can be assigned to individuals. Once a task is completed, it can be approved or rejected. In this way, workflows ensure that the program is thoroughly tested and validated before it is promoted. Overall, workflows are an essential part of LSAF, and they help to ensure that the program is of the highest quality.





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