

Reetika Sarkar – JMP Clinical

About Me

- PhD candidate (Year 4) in Computational Mathematics (Statistics) at UNC Greensboro
 - Started at SAS in May 2024 as a Graduate Intern in JMP Clinical Group reporting to Mann Geoffrey
 - Goal: Full-time Statistician/Developer role working in the Technology/Pharmaceutical Industry



Projects

- Validation of JMP Clinical safety evaluation reports to enhance reliability of the software

- Reports validated: Adverse Events (AE), Medical History (MH), Concomitant Medications (CM), Acute Kidney Injury (AKI), and Drug Induced Liver injury (DILI)
 - Dataset used: Nicardipine (a drug used to treat hypertension)

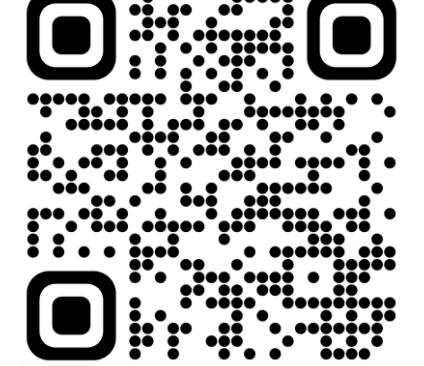
Analysis of Vaccine Adverse Events Reporting Systems (VAERS) safety data using JMP Clinical 19

- **SAS/JMP Courses and Trainings:**
 - SAS® Base Programming Fundamentals

- SAS® Programming 1: Essentials
 - SAS® Programming 2: Data Manipulation Techniques
 - SAS® Programming for R Users
 - Statistical Thinking for Industrial Problem Solving (STIPS)
 - Introduction to the JMP Scripting Language (JSL)

- **Hands-on Training with Team**

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Contact Information

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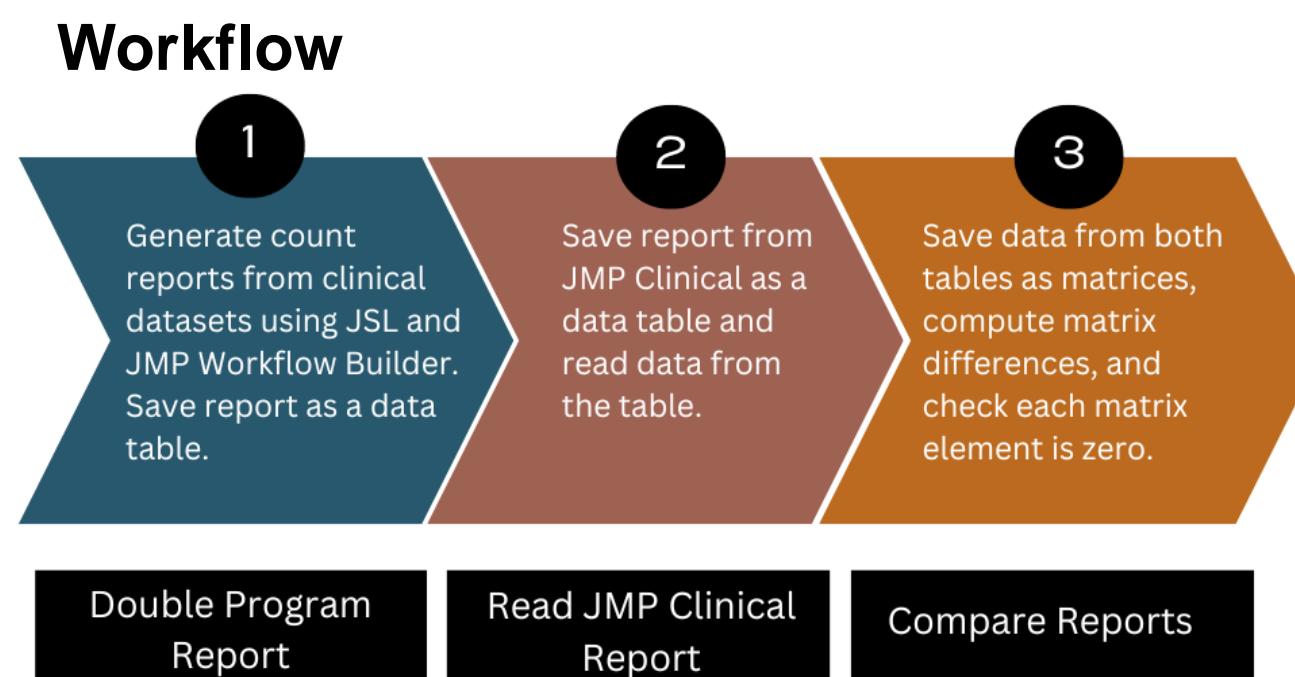
Double Programming and Unit Testing for Validation of JMP Clinical Safety Analysis Reports pertaining to Events, Findings, and Interventions

Challenge

Drug safety assessment in a clinical trial includes meticulous evaluation of a drug's potential side effects with the goal to ensure that the benefits outweigh the potential risks. It includes analysis of events, interventions, and findings data across treatment arms to report adverse findings. The goal of this project was to double program the reports mentioned above using JMP Scripting Language (JSL) and compare the findings to the reports generated in JMP Clinical.

Implications / Applications

The reports double programmed in this project would help in establishing a framework that validates the clinical safety reports generated using JMP Clinical. These reports would be made available to the users of JMP Clinical.



JMP Functionalities and JSL Code Overview

- Data Table: Open, Format, Tabulate, Subset, Summary, Transpose, Update, Get As Matrix, Select Matching Cells, Select Where, New Column, Set Selected Column, Delete Column, Exclude
 - Statistical: Max
 - Matrix: Loc
 - Programming: Length, N Items, Show
 - Conditional: If, For Each Row
 - JMP Workflow Builder

Example Code for AE Report

JMP Workflow Builder

The screenshot shows the JMP Workflow Builder interface. On the left, a code editor displays a JSL script for 'Workflow 353'. The script performs several steps: reading a report from a saved JMP Clinical file, deleting percentage columns, sorting by an aggregate column, generating tables using Tabulate(), opening data tables (ae.jmp and adsl.jmp), updating a data table, launching the Tabulate platform, and finally making the tabulate report into a data table. It also includes a comparison step to check if two matrices are the same.

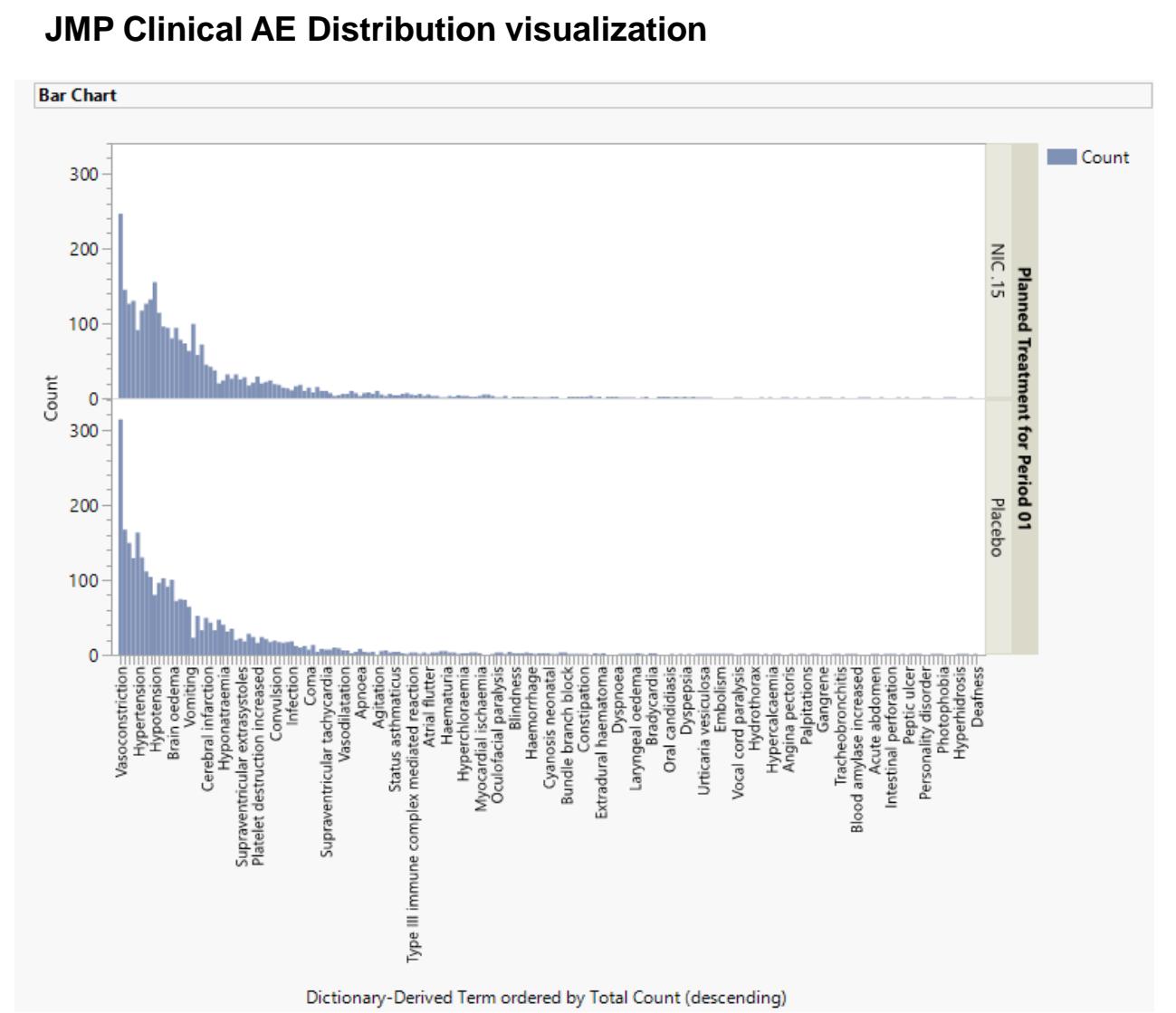
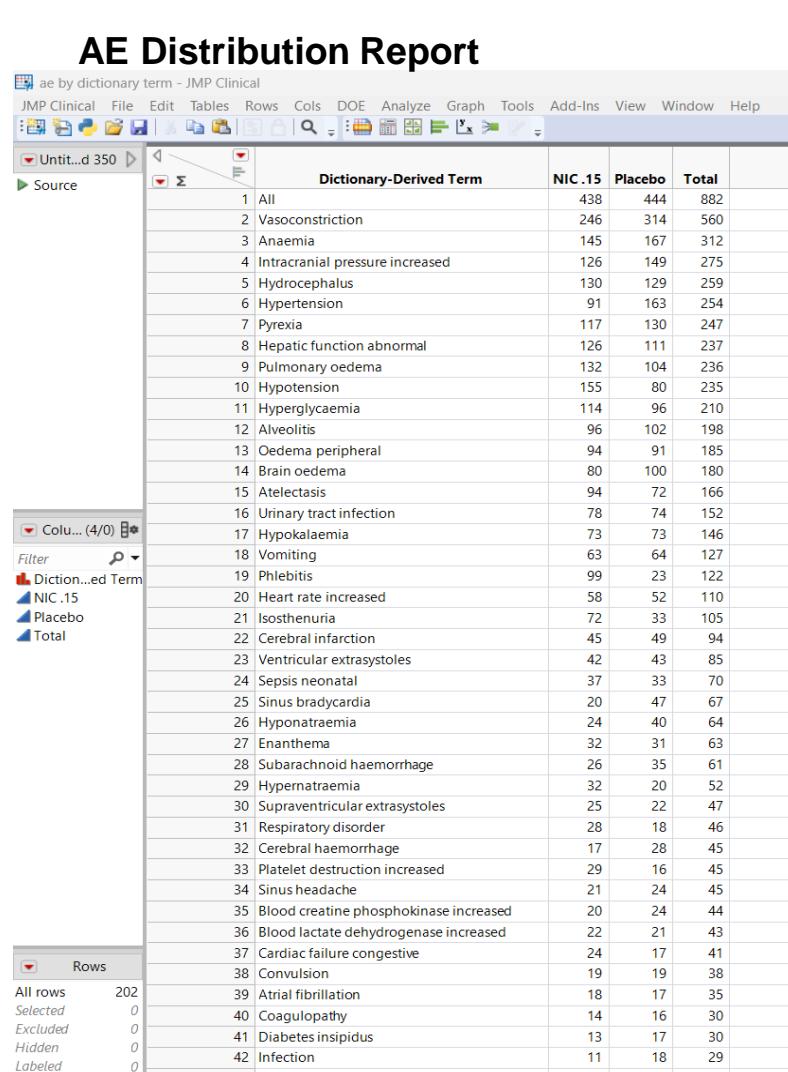
The central part of the interface is the 'Workflow Builder' panel, which lists the four steps defined in the script: 'Open Data Table: ae.jmp', 'Open Data Table: adsl.jmp', 'Update data table', and 'Launch platform: Tabulate'. Each step has a small icon and a 'Step Settings' button.

On the right, the 'Script Window - JMP Clinical' shows the full JSL script. The 'JMP Log History' panel at the bottom lists the actions taken during the workflow execution, including 'Report snapshot: ...SUBJID - Tabulate', 'Close Data Table: ...group by USUBJID', 'Close Data Table: Subset of lb', 'Close Data Table: lb', 'Open Data Table: ae.jmp', and 'Open Data Table: adsljmp'.

```

1 Names Default To Here( 1 );
2
3 // STEP 1: Read report from JMP Clinical saved as a data table
4 review_report_data = Open(
5   "C:\Users\resark\OneDrive - SAS\Documents\NicardipineAE181\Nicardipine_AE by Dictionary Term\NicardipineAETermReport181.jmp"
6 );
7
8 // Delete percentage columns from table
9 Column( review_report_data, 3 ) << Set Selected;
10 Column( review_report_data, 5 ) << Set Selected;
11 Column( review_report_data, 7 ) << Set Selected;
12 review_report_data << Delete Columns();
13
14 // Sort data table by aggregate column
15 review_report_data << Sort(
16   By( Column( review_report_data, 4 ), Column( review_report_data, 1 ) ),
17   Replace Table,
18   Order( Descending, Ascending )
19 );
20
21 // STEP 2: Generate tables using Tabulate()
22 // Open Data Table: adsl.jmp
23 Open( "$CLINICAL_HOME/Sample Data/Nicardipine/ADaM/adsl.jmp" );
24
25 // Open Data Table: ae.jmp
26 Open( "$CLINICAL_HOME/Sample Data/Nicardipine/SDTM/ae.jmp" );
27
28 // Update data table
29 Data Table("ae") << Update(
30   With( Data Table( "adsl" ) ),
31   Match Columns( :USUBJID = :USUBJID )
32 );
33
34 // Tabulate from updated table
35 tabulate_report_data = (Data Table("ae") << Tabulate(
36   ID( :USUBJID ),
37   Add Table(
38     Column Table(
39       Grouping Columns( :TRT01P ),
40       Add Aggregate Statistics( :TRT01P )
41     ),
42     Row Table(
43       Grouping Columns( :AEDECOD ),
44       Add Aggregate Statistics( :AEDECOD )
45     )
46   )
47 )) << Make Into Data Table;
48
49 // Sort data table by aggregate column
50 tabulate_report_data << Sort(
51   By( Column( tabulate_report_data, 4 ), Column( tabulate_report_data, 1 ) ),
52   Replace Table,
53   Order( Descending, Ascending )
54 );
55
56 // STEP 3: Compare tables and report if they are same
57 // Extract numerical columns from data table into a matrix
58 review_report_matrix = Round( review_report_data << Get As Matrix() );
59 tabulate_report_matrix = Round( tabulate_report_data << Get As Matrix() );
60
61 Diff = review_report_matrix - tabulate_report_matrix;
62
63 If( Length( Loc( Diff, 0 ) ) == N Items( tabulate_report_matrix ),
64   Print( "All values in the two tables are the same." ),
65   Print( "All values in the two tables are not the same." )
66 );

```



Example Code for AKI/DILI Report

JMP Clinical AKI Report

Parameter	Analysis Visit	n (%)	Mean	Mean Change from Baseline
Creatinine	Visit 1	16 (3.6)	6 (1.3)	2.3 (0.2, 4.3)
	Visit 2	9 (2.0)	11 (2.4)	-0.4 (-2.3, 1.5)
	Visit 3	11 (2.5)	5 (1.1)	1.4 (-0.4, 3.1)

Lab Name	NIC.15 (N = 447)	Placebo (N = 455)	Risk Difference for NIC.15 over Placebo	
Creatinine	Level 1 (>=1.5 x baseline)	16 (3.6)	6 (1.3)	2.3 (0.2, 4.3)
	Level 2 (>=2 x baseline)	9 (2.0)	11 (2.4)	-0.4 (-2.3, 1.5)
	Level 3 (>=3 x baseline)	11 (2.5)	5 (1.1)	1.4 (-0.4, 3.1)

Double programmed AKI Report tabulation and comparison

```

118 // STEP 3: Compare tables and report if they are same
119 // Extract numerical columns from both data tables into a matrix
120 review_report_matrix = Round( review_report_data << Get As Matrix() );
121 tabulate_report_matrix = Round( tabulate_report_data << Get As Matrix() );
122
123 Diff = review_report_matrix - tabulate_report_matrix;
124
125 If( Length( Loc( Diff, 0 ) ) == N Items( tabulate_report_matrix ),
126     Print( "All values in the two tables are the same." ),
127     Print( "All values in the two tables are not the same." )
128 );

```

JMP Clinical window showing the 'Untitled 356' data table. The table has columns: Category Level, NIC.15, and Placebo. The data shows the count of patients in each category for both groups.

Category Level	NIC.15	Placebo
1 >=1.5 x baseline	16	6
2 >=2 x baseline	9	11
3 >=3 x baseline	11	5

```

tabulate_report_data = DataTable("Untitled 356");
review_report_data = DataTable("AKITable_Clinical19");

```

