

# Data Library Comparison Macro %COMPARE\_ALL

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# Data Library Comparison Macro %COMPARE\_ALL

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- Creates a report comparing all datasets within two libraries
  - Outputs to an Excel file (requires SAS 9.4+)
- 8 parameters (3 required, 5 optional)
- Creates four different types of worksheets in the report
- Contains navigation to easily move from one summary to another
- Contains error checking, documentation, and cleans up after itself



### Macro Parameters

#### Required

- BASE: designates the first library (libname not path) to be compared
- COMPARE: designates the second library for comparison
- OUTDOC: full file path and file name of output Excel document

### Optional

- ID: designate one or more variables as unique identifiers (example: Patient ID).
   Each dataset takes any ID variables that in this list that exist within it. They are sorted by the ID variables from left to right in the list.
- SELECT: optionally selects specific datasets from libraries for comparison
- CROSSTAB\_THRESHOLD: cut-point for displaying all unique values for a variable in comparison table
- IDSUMTABLE: determines how many ID variables are used for change summary table
- DEBUG: turns on options for debugging macro issues



### Macro Call Example

```
Libname frz18 'file-path 2018 version';
Libname frz19 'file-path 2019 version';
```

```
%COMPARE_ALL(
    BASE=frz18,
    COMPARE=frz19,
    OUTDOC=compare_all_example.xlsx,
    ID=patient_id cycle);
```



Example: ID=patient\_id cycle eval\_dt

Dataset 1: Includes variables patient\_id, age, sex, performance\_score

Dataset 2: Includes variables patient\_id, cycle, agent, dose

Dataset 3: Includes variables patient\_id, eval\_dt, toxicity, grade



Example: ID=patient\_id cycle eval\_dt

Dataset 1: Includes variables patient\_id, age, sex, performance\_score

Uses only patient\_id as ID variable

Dataset 2: Includes variables patient\_id, cycle, agent, dose Uses patient\_id and cycle as ID variables in that order

Dataset 3: Includes variables patient\_id, eval\_dt, toxicity, grade Uses patient\_id and eval\_dt as ID variables in that order



### **Macro Error Checking**

- The following items are checked by the macro:
  - If the BASE and COMPARE libraries exist and have been assigned
  - If OUTDOC is missing
  - If the CROSSTAB\_THRESHOLD and IDSUMTABLE parameters are not a number greater than 0



### Macro Error Checking

```
69 %include '~m080449/ccsic/mymacs/compare_all.sas';
1037 %compare_all(base=ifrz,compare=iinter2,id=protnum dcntr_id new_id,outdoc=~/ibm/data_changes.xlsx,
1038 select=,idsumtable=1);
ERROR: (Global: COMPARE) Library does not exist
ERROR: 1 pre-run errors listed
ERROR: Macro COMPARE_ALL will cease
COMPARE_ALL has finished processing, runtime: 0:00
1039
```

```
%compare_all(base=ifrz,compare=iinter,id=protnum dcntr_id new_id,outdoc=~/ibm/data_changes.xlsx,
1038 select=,idsumtable=-5);
ERROR: (Global: IDSUMTABLE) Must be greater than or equal to 0. -5 is not valid.
ERROR: 1 pre-run errors listed
ERROR: Macro COMPARE_ALL will cease
COMPARE_ALL has finished processing, runtime: 0:00
1039
```



### **Top Summary Worksheet**

- Overview of all datasets in either library
  - If SELECT parameter is used then only the datasets in the SELECT list will be shown
- Meta data is shown for both BASE and COMPARE versions
  - Date last updated, number of observations, number of variables
- High-level summary of differences
  - How many variables have had an attribute such as type changed
  - How many observations (based on matching ID variables) are in BASE but not COMPARE
  - How many observations (based on matching ID variables) are in COMPARE but not BASE
  - How many total variable values changed (based on matching ID variables)
- ID Variables Used for each dataset's comparison
- Libraries being compared are listed at top of sheet with file path
- Clicking a dataset name will navigate to that dataset's summary page
- BASE, COMPARE, Differences, and ID variables are colored consistently throughout report
- Differences are highlighted with color



## **Top Summary Worksheet**

	A	В	С	D	E	F	G	Н			K	1
- 10		D	C	U	E	F	G	П		J	I.	L
1	Base Library (FREEZE): /frozen_data/	+										
2	Compare Library (LIVE): /live_data/	- ly										
3	Summary of Datasets Compared	PRODUCE CONTRACTOR OF THE PRODUCE CONTRACTOR							4-7-1000			
4		Base			Compare			Any Differe		I de la constantina della cons	I m	
		Last Updated	A CONTRACTOR OF THE PARTY OF TH	Number of	Last Updated	Number of		The second second	Lost	New	Data	ID Variables Used
5	Dataset Name		Variables	Observations		Variables	Observations	Attributes	Observations	Observations	Changes	
6	AE_MAX_GRADE	02/12/2019	20	16316	11/03/2019	21	12163	0	6369	2216	0	protnum, dcntr_id
7	BASELINE	02/12/2019	12	40928	11/03/2019	13	38368	0	2560	0	0	protnum, dcntr_id
8	BIOMARKERS	02/12/2019	5	22494	01/07/2020	7	21206	3	1726	438	9812	protnum, dcntr_id
9	DZCHAR_PRIORTRT	02/12/2019	19	40928	11/03/2019	20	38368	0	2560	0	11	protnum, dcntr_id
	DZ_ASSESS	02/12/2019	27	133619	11/03/2019	28	133471	1	148	0	0	protnum, dcntr_id,
10				the second	8.0000000000000000000000000000000000000		THE STATE OF THE S		3333	1,550	5-50	merged_day
11	LABS_BASELINE	02/12/2019	11	35758	11/03/2019	12	35164	0	594	0	0	protnum, dcntr_id
	LABS	02/12/2019	23	235656	11/03/2019	24	234760	0	896	0	0	protnum, dcntr_id, visit,
12												study_day
13	MET_SITES	02/12/2019	27	32029	11/03/2019	27	31266	0	763	0	0	protnum, dentr_id
14	OUTCOMES	02/12/2019	37	40081	11/03/2019	20	37766	2	2315	0	5827	protnum, dcntr_id
15	PRIMARY_SITE	02/12/2019	7	29336	11/03/2019	8	27657	0	1679	0	0	protnum, dentr id
	PRIOR CHEMO	02/12/2019	6	28841	11/03/2019	7	18379	0	10462	0	0	protnum, dcntr_id, start_dt,
16												end dt
17	PROT_TRT	02/12/2019	11	40928	11/03/2019	13	28395	1	12533	0	24141	protnum, dcntr_id
18	PROT_TRT_MAINT				11/03/2019	16	4660	0				
19	PROT_TRT_SEQUENCE				11/03/2019	22	9746	0		V)		
	SUBSEQUENT_CHEMO	02/12/2019	16	16357	11/03/2019	17	16327	0	30	0	0	protnum, dcntr_id, start_dt
	VITALS	02/12/2019	11	164559	11/03/2019	12	163301	0	1258	0	0	protnum, dentr id, visit



### **Dataset Summary Worksheet**

- Overview of variables in each dataset that exists in both libraries
- Displays basic meta data for each variable
  - Type, length, label and format
  - Highlights any meta data that has changed
- Marks each variable being used as an ID variable in green
- Summary of differences by variable
  - Whether variable exists in BASE but not COMPARE
  - Whether variable exists in COMPARE but not BASE
  - Whether any observations (based on matching ID variables) have changed values
- Any variables with differences are marked in red
  - Clicking a red variable will navigate to that variable's summary page
- Navigation to top summary worksheet is in the header



## **Dataset Summary Worksheet**

Dataset Name: PROT_TRT	В	7	70 To	100		G	11.00		7, 1		7	100
	+											
ID Variables: protnum dcntr id	10:											
Return to Top Summary	-	Dane			- 2	Compara				Ame Differ		
	1011 111 0	Base				Compare				Any Differences		
Variable Name	ID Variables?	Туре	Label	Format	Length	Type	Label	Format	Length	A TANK TO THE COMMON CO	New Variable	No. Data Changes
PROTNUM	Yes	Numeric	Study Name	PROT	8	Numeric	Study Name	PROT	8	No	No	20000000
DCNTR_ID	Yes	Character	Patient ID	S	25	Character	Patient ID	5	25	No	No	
NEW_ID	No	Character				Character	ARCAD ID		25	No	Yes	
LINE_TRIAL	No	Numeric	1st, 2nd, or >=3 Line Study	LINE_TRIAL	8	Numeric	1st, 2nd, or >=3 Line Study	LINE_TRIAL	8	No	No	0
STUDY_START_DT	No	Numeric	Study Start Date	MMDDYY	8	Numeric	Study Start Date	MMDDYY	8	No	No	0
ARM STRAT INDEX	No	Numeric	Arm Stratification	Medeument)	8	Numeric	Arm Stratification	NAME OF THE PARTY	8	No	No	123
ARM_STRAT_TEXT	No	Character	Arm Stratification (Decode)		40	Character	Arm Stratification (Decode)		100	No	No	2291
TARGET	No	Numeric	Regimen Includes Any Target Agents?	TARGET	8	Numeric	Regimen Includes Any Target Agents?	TARGET	8	No	No	0
CHEMO_BACKBONE	No	Character				Character	Chemotherapy Backbone		40	No	Yes	
ANG	No	Numeric	Regimen Includes Any Angiogenic Agents?	ANG	8	Numeric	Regimen Includes Any Angiogenic Agents?	ANG	8	No	No	0
EGFR	No	Numeric	Regimen Includes Any Anti-EGFR Agents?	EGFR	8	Numeric	Regimen Includes Any Anti-EGFR Agents?	EGFR	8	No	No	0
TRT_DAYS	No	Numeric	Duration (Days) of Protocol Treatment (Excluding Strategy Trials)		8	Numeric	Duration (Days) of Protocol Treatment (Excluding Strategy Trials)		8	No	No	0
STUDY END DT	No	Numeric	Study End Date	MMDDYY	8	Numeric	Study End Date	MMDDYY	8	No	No	0



# Data Changes Summary Worksheet

- Summary of changes within dataset, follows Dataset Summary worksheet
- Summarizes lost observations, new observations, and number of data changes across specified ID variables
  - Maximum number of ID variables used is determined by IDSUMTABLE
- Each variable with data changes is listed along with a summary based on variable type and CROSSTAB\_THRESHOLD
  - If number of unique changes <= CROSSTAB\_THRESHOLD then each unique change is listed along with a count
  - If number of unique changes > CROSSTAB\_THRESHOLD then values are summarized as "Non-missing" or "Missing" with counts
  - Numeric variables also have minimum and maximum changed value
- Navigation to top summary and dataset summary worksheet is in the header



# Data Changes Summary Worksheet

4	Α	В	С	D	E	F	G	Н	1	J
1	Summary of Pi	ROT_TRT_INTE	RMEDIATE Sui	mmary of Data (	Changes		1.	421		- 11
2	All Available ID	Variables: prof	num dentr_id							
3	Note: Only firs	t 1 ID variable(s	) is used in su	mmary						
1	Return to Top	Summary				MATE				
5	Study Name	Control of the Contro	New Observations	N Data Changes	Variable	Base	Compare	N	Mininum	Maximum
	Study 1	6	0	171	arm_strat_text	XELOX	CAPOX	171		
	Study 2	0	0	471	arm_strat_text	Cap	Capecitabine	156		
						Cap+Bev	Capecitabine + Bevacizumab	157		
						Cap+Bev+ Mitomycin	Capecitabine + Bevacizumab + Mitomycin	158		
)	Study 2	0	0	0						
	Study 3	825	0					1		
,	Study 4	0	0	0						
,	Study 5	0	0	463	arm_strat_text	BSC	Best Supportive Care	232		
4	- 82					BSC + panitumumab	Best Supportive Care + Panitumumab	231		
5	Study 6	0	0	923	arm_strat_text	5FULV+Bev	5FULV + Bevacizumab	110		
3	164					IFL+Bev	IFL + Bevacizumab	402		
7						IFL+Placebo	IFL + Placebo	411		



# Data Changes Summary Worksheet

A	А	В	С		E	F	G	Н	1
1	Summary of Bl	OMARKERS Su	ımmary of Data	Changes		20		120	12
2	All Available ID	Variables: prot	tnum dcntr_id						
3	Note: No ID va	riables used in	summary						
4	Return to Top	Summary						16.00	
5	Lost Observations	New Observations	N Data Changes	Variable	Base	Compare	N	Mininum	Maximum
6	1726	438	9812	braf		MT	183		
7						WT	1250		
В					MT		6		
9					WT		66		Ì
0				kras		MT	1256		
1					5	WT	2111	1 5	11
2				ras		MT	2381		
13						WT	2559		j



# Variable Changes Summary Worksheet

- Created for each variable that had any value changes
- Replicates the output from traditional COMPARE procedure
  - Displays observations with data changes
  - Lists each ID variable, base and compare values, and absolute/percent change if the variable is numeric
- Navigation to top summary and dataset summary worksheet is in the header

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# Variable Changes Summary Worksheet

Α	В	С	D	E	F	G
Dataset Name:	PROT_TRT_	INTERMEDIATE				
Variable Name	(label): arm_	strat_text (Arm	Stratification (Decode))			
Return to Top	Summary					
D Variables			-V	191	10.515	
Study Name	Patient ID	Observation	Base	Compare	Absolute Change	Percent Change
Study 1	1	1	FOLFOX4 - bevacizumab	FOLFOX4 + Bevacizumab		
Study 1	101	4	bevacizumab	Bevacizumab		
Study 1	102	5	FOLFOX4 - bevacizumab	FOLFOX4 + Bevacizumab		
Study 1	105	8	FOLFOX4 - bevacizumab	FOLFOX4 + Bevacizumab		
Study 1	106	9	FOLFOX4 - bevacizumab	FOLFOX4 + Bevacizumab		
Study 1	108	11	FOLFOX4 - bevacizumab	FOLFOX4 + Bevacizumab		
Study 1	109	12	FOLFOX4 - bevacizumab	FOLFOX4 + Bevacizumab		
Study 1	11	13	bevacizumab	Bevacizumab		
Study 1	110	14	bevacizumab	Bevacizumab		
Study 1	111	15	bevacizumab	Bevacizumab		

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- The COMPARE\_ALL macro is a powerful tool for comparing multiple versions of the same library
- Creates an easy to read Excel report with built in navigation to easily jump to the needed worksheet
- The macro is available for download on the <u>SAS Communities</u> <u>page</u>



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