





Biochemical assay Analyzer and CAlculator

BiAnaCA: A freely accessible tool for data analysis of end-point biochemical assay

BiAnaCA ~ 0.1 Tutorial



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and click "Perform"
Step 10: Click on Print Report to generate it
Contact Us

Introduction

A biochemical assay is an investigative (analytic) procedure in laboratory medicine, pharmacology, environmental biology, continuous delivery, and molecular biology for qualitatively assessing or quantitatively measuring the presence or amount or the functional activity of a target entity (the analyte) in cuvettes or 96- well Microplates using Colorimeter, Spectrophotometer or Microplate reader.



Applications

• BiAnaCA can be used not just for biochemical assays but any assay which involves spectrophotometric absorbance readings and you want to calculate the unknown concentration from the standard point or calibration curve method.

- Can be used well for 96 well plate but it is not restricted to 96 well it can be used for any number of large data set if you provide in the proper .csv file format.
- Can be used separately for linear regression curve analysis and statistical analysis also for any other applications
- Can generate .pdf reports by a software which can be help as regulatory requirement and documentary evidence
- It is totally free in nature and Fast, user friendly and efficient enough for academic or industrial research
- It is applicable not in just pharmacy or medical field but any clinical, biotech, analytical chemistry, agriculture and science field

Download and Installation

- Download the BiAnaCA for your system from <u>http://www.niper.ac.in/pi_dev_tools/BiAnaCA/index.php</u>
- To run the BiAnaCA you just need java run time environment (JRE) properly installed and updated as per your system configuration.
- Download and update latest java JRE.
- Download the zip folder and extract all files to a folder where you want to install it. You can install it anywhere but make sure that it has included executable BiAnaCA_version_0.1.jar file and a lib folder with demo folder inside lib folder. Do not move or Rename files within it.
- Click on BiAnaCA_version_0.1.jar to start the application.
- You can find the tutorial and sample files in the folder lib/demo/.

Step-wise workflow for BiAnaCA

Layout of workflow



Home screen



Step 1: Fill the experiment details in the given boxes

- Name of your assay
- Name of person, date and all other notes which you want to print in report fill them. You can copy paste with Ctrl+C/Ctrl+V functions.

N BiAnaCA	
<u>BiAnaCA</u> BioChemical Assay Ana	Nyzer and Calculator
HOME NEW PROJECT	
Project Detail Result Data Sheet Analysis Reports Graphical Plots Calibraton curve plots	
Project Detai	Data Upload box
Name of the assay:	Provide the absorbance data from?
Experiment performed by:	txt/.csv file upload Manual entry in data sheet
Experiment performed on:	Browse File Path Selected: (96-well plate data file)
Experimental notes:	Limit for outliers in replicates (in % C.V.):
Group Patterns	Calculation Panel
Fill in the details of samples for your assay experiment	Calculation method selection:
Number of groups: 1 🛛 🖓 + Standard (1 Smp)	Single Standard Multiple Standards
Number of sample replicates: 3 👘 🥑 + Blank (1 Smp) OK, Start	Enter the Standard concentration:
Group no. Group code No. of samples then do calculate Click on home	Mean Standard absorbance reading:
then select manual entry button	Dilution factor for sample readings:
Read me → rt will open table with data change samople code if u wish or else do submit there	Standard/sample concentration units:
and now click calculate do analysis.	Number of points in Std. calibration curve:
Reset Submit	Calculate Plot Equation Check Outliers
	•
Designed and developed by team BiAnaCA,	Dept. of Pharmacoinformatics, NIPER, S.A.S.Nagar, India.

Step 2: Fill the groups and sample replicate details

- Give the number of groups like control, treatment A, B, C etc. or groups like formulation A, B, C or group of compounds etc. excluding standard and blank.
- Tick the whether standard and/or blank were used or not.
- Enter the number of groups and click enter.
- Enter the number of replicates performed for each sample. E.g. 1 for single, 2 for duplicate and 3 for triplicate like wise. Click on **OK**, **Start**.
- Now enter the code of group and number of samples in each group. You can use tab and arrow keys for the navigation purpose.
- Click **Submit** for further process to upload the reading data or you can reset for new details.

BiAnaCA BioChemical Assay Analyzer and Calculator

HOME NEW PROJECT	
Project Detail	Data Upload box
Name of the assay: Demo	Provide the absorbance data from?
Experiment performed by: Your name	 ⊚ .txt/.csv file upload Manual entry in data sheet Standard at Start Standard at End
Experiment performed on: 8/2/16	Browse File Path Selected: (96-well plate data file)
Experimental notes:	Limit for outliers in replicates (in % C.V.):
Group Patterns Fill in the details of samples for your assay experiment Number of groups: Number of sample replicates: Group no. Multiple and the samples Multiple and the samples Multiple and the samples Group no. Group no. Group no. Group no. Group no. Group no. Multiple and the samples Multiple and the samples Multiple and the samples Multiple and the samples Multiple and the samples <td>Calculation Panel Calculation method selection:</td>	Calculation Panel Calculation method selection:
Reset Submit "	Calculate Plot Equation Check Outliers
Designed and developed by team BiA	naCA, Dept. of Pharmacoinformatics, NIPER, S.A.S.Nagar, India.



*To view the sample files go to Lib//demo// TestData.csv, or TestData.txt, or Test_96WellData.csv

BiAnaCA		
<u>BiAnaCA</u> BioChe	emical Assay An	alyzer and Calculator (Chout Contemporation Advance) Demo
HOME NEW PROJECT		
Project Detail Result Data Sheet Analysis Reports Graphical Pl	Calibraton curve plots	
Project Detail		Data Upload box
Name of the assay:	Demo	Provide the absorbance data from?
Experiment performed by:	User	● .txt/.csv file upload
Experiment performed on:		Browse :_browse_option.csv (96-well plate data file)
Experimental notes:	Temp: 25	Limit for outliers in replicates (in % C.V.):
Group Patterns		Calculation Panel
Fill in the details of samples for your assay e	xperiment	Calculation method selection:
Number of groups: 4	📝 + Standard (1 Smp)	Single Standard Multiple Standards
Number of sample replicates: 3	V + Blank (1 Smp) OK, Start	Enter the Standard concentration:
Group no. Group code No. of samples	then do calculate	Mean Standard absorbance reading: 0.01
1 Control 7 2 Treatment A 7	Click on home then select manual entry buttor	Dilution factor for sample readings:
3 Treatment B 8 4 Treatment A+B	change samople code if u wish	Standard/sample concentration units: sd
	and now click calculate do analysis.	Number of points in Std. calibration curve:
Reset	(III)	Calculate Plot Equation Check Outliers
	III	,
Designed a	nd developed by team BiAnaCA	A, Dept. of Pharmacoinformatics, NIPER, S.A.S.Nagar, India.

Select the option, browse and upload the file as per given format or manually enter the data

Option 1: for uploading a .csv/.txt file with the absorbance readings sorted sample-wise

Format of Data.csv file (Save as .CSV coma separated value format from excel)

Format of Data.txt file (give tab -> to separate values and enter for next sample)

X	🛃 🤊 • (?) • 🛅 =					📔 H:\d	ata.txt - Notepad+	+ [Administ	rator]			
	File Home Inser	t Page Layo	ut Form	ulas Da	ata Review	File E	dit Search View	w Encoding	g Languag	e Settings	Macro	Run Plug
1	🚆 👗 Cut	Calibri	* 11	A A	= = =) 🖌 🖉 🕞		66 2	C 8	₽ <mark>₽</mark> ₽ 🔇 😵	ء 🗗 🔊
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	 Format Painter 	B 1 <u>0</u>				Cata			32.7			
_	Clipboard 🖓	F	ont	Fai		1	SAMPLECODE	OD1 OD2	OD3			
	A1 -	(* <i>f</i> x	SAMPLE	CODE		2	CTRL-C-1-H	0.071	0.07	0.072		
1	A	В	С	D	E	3	CTRL-C-1-M	0.053	0.056	0.059		
1	SAMPLECODE	OD1 O	D2 0	D3		4	CTRL-C-1-T	0.056	0.061	0.06		
2	CTRL-C-1-H	0.071	0.07	0.072	First row Titles	5	CTRL-C-1-HM	0.067	0.071	0.083		
3	CTRL-C-1-M	0.053	0.056	0.059	Thistrow Thes	6	CTRL-C-1-HT	0.055	0.057	0.055		
4	CTRL-C-1-T	0.056	0.061	0.06		7	CTRL-C-2-H	0.062	0.072	0.063		
5	CTRL-C-1-HM	0.067	0.071	0.083		8	CTRL-C-2-M	0.069	0.082	0.072		
6	CTRL-C-1-HT	0.055	0.057	0.055	Eirst column just	9	HSD-S-3-H	0.081	0.083	0.077		
7	CTRL-C-2-H	0.062	0.072	0.063	First column just	10	HSD-S-3-M	0.094	0.099	0.099		
8	CTRL-C-2-M	0.069	0.082	0.072	comple code	11	HSD-S-3-T	0.087	0.093	0.097		
9	HSD-S-3-H	0.081	0.083	0.077	sample coue	12	HSD-S-4-HM	0.09	0.096	0.095		
10	HSD-S-3-M	0.094	0.099	0.099		13	HSD-S-4-HT	0.125	0.131	0.123		
1:	HSD-S-3-T	0.087	0.093	0.097		14	HSD-S-4-H	0.14	0.152	0.149		
1:	HSD-S-4-HM	0.09	0.096	0.095	Complex in the	15	HSD-S-5-M	0 082	0.087	0 092		
13	HSD-S-4-HT	0.125	0.131	0.123	samples in the	16	UNV-U-6-M	0.073	0.07	0.072		
14	HSD-S-4-H	0.14	0.152	0.149	Commence	17	UNIX U C T	0.075	0.07	0.072		
15	HSD-S-5-M	0.082	0.087	0.092	Sequence of	11	UNX-0-6-1	0.062	0.07	0.062		
10	UNX-U-6-M	0.073	0.07	0.072		18	UNX-U-6-HM	0.08	0.091	0.091		
17	UNX-U-6-T	0.062	0.07	0.062	given group	19	UNX-U-6-HT	0.094	0.105	0.097		
18	UNX-U-6-HM	0.08	0.091	0.091	D 1 1	20	UNX-U-6-MT	0.074	0.074	0.069		
19	UNX-U-6-HT	0.094	0.105	0.097	Details	21	UNX-U-7-H	0.074	0.075	0.073		
20	UNX-U-6-MT	0.074	0.074	0.069		22	UNX-U-7-M	0.075	0.089	0.083		
2:	UNX-U-7-H	0.074	0.075	0.073		23	UNX-U-T-T	0.081	0.088	0.089		
22	UNX-U-7-M	0.075	0.089	0.083	C	24	UNX+HSD-US-	8-H 0.13	38 0.14	1 0.1	41	
23	UNX-U-T-T	0.081	0.088	0.089	Save as .csv file	25	UNX+HSD-US-	8-M 0.10	0.10	0.10	04	
24	UNX+HSD-US-8-H	0.138	0.141	0.141		26	UNX+HSD-US-	8-T 0.15	0.15	58 0.1	48	
25	UNX+HSD-US-8-M	0.105	0.105	0.104		27	UNX+HSD-US-	9-HM 0.14	16 0.15	55 0.1	57	
20	UNX+HSD-US-8-T	0.15	0.158	0.148		28	UNX+HSD-US-	9-HT 0.15	57 0.1	73 0.1	68	
27	UNX+HSD-US-9-HM	0.146	0.155	0.157	Template file	29	UNX+HSD-US-	9-MT 0.11	0.11	15 0.1	15	
28	UNX+HSD-US-9-HT	0.157	0.173	0.168		30	UNX+HSD-US-	10-H 0.10	0.10	0.10	02	
29	UNX+HSD-US-9-MT	0.111	0.115	0.115	Available in folder	31	UNX+HSD-US-	10-T 0.10	0.11	14 0.10	08	
30	UNX+HSD-US-10-H	0.104	0.102	0.102		32	STD 0.079	0.081	0.082			
3:	UNX+HSD-US-10-T	0.108	0.114	0.108	Lib\demo\data.csv	33	BLANK O	0 0	01002			
32	STD	0.079	0.081	0.082			DIRINI U	0 0				
33	BLANK	0	0	0								
34	1											

Option 2: for manual entry

Format for manual entry in data sheet. After entering data, you can also save it as a .csv file

ARE NEW PROJECT					
ata Entry Form	and the state of t	-			
/ Check the following	table. Then dick Ok Submit. Export CS	V To Save as file	Ok Sul	mit Decet Evport t	in CSV
/ check the following	grable. Then cloc ok submit, export ca	v to save as the.	UK 30	unic Reset Export o	
.no	Sample code	Obs 1	Obs 2	Obs 3	absorbance Data from 2
	1 Group Control: Sample 1				absorbance bata nom :
	2 Group Control: Sample 2				
	3 Group Control: Sample 3				Opened Enter in Data Chaot
	4 Group Control: Sample 4				
	5 Group Control: Sample 5				standard at End
	6 Group Control: Sample 6				(96 well plate data file)
	7 Group Control: Sample 7				
	8 Group Treatment A: Sa				i (in % C.V.) : 10 ≑
	9 Group Treatment A: Sa				
	10 Group Treatment A: Sa				
	11 Group Treatment A: Sa				valid CSV / TXT file.
	12 Group Treatment A: Sa				
market and	13 Group Treatment A: Sa				
Editab	C14 Group Treatment A: Sa				
	16 Group Treatment B: Sam				
Compati	17 Group Treatment B: Sam				election:
Sampi	19 Group Treatment B: Sam.				
	19 Group Treatment B: Sam				own Concentration 🔘 Std. Calibration Curve Regression
codes	20 Group Treatment B: Sam				Community Abanhara
coucs	21 Group Treatment B: Sam				Lard Concentration:
	22 Group Treatment B: Sam				
	23 Group Treatment A+B: S.				Reading mean ·
	24 Group Treatment A+B: S.				
	25 Group Treatment A+B: S.				
	26 Group Treatment A+B: S.				uple readings:
	27 Group Treatment A+B: S.				
	28 Group Treatment A+B: S.				contration Units
	29 Group Treatment A+B: S.				icentration onnes:
	30 Group Treatment A+B: S.				
	31 Standard				d Calibration curve
	32 Blank				
					ation Check Outliers

Option 3: for uploading unsorted absorbance readings in 96-or more well format

Format of 96 well plate readings in unsorted form (Save as .CSV coma separated value format from excel)

X	. 9 - C	- 🗄 🖙					-			Test_9	5WellData	
Fil	le Hor	ne Inse	rt Pagel	Layout F	ormulas	Data F	leview V	iew Dev	eloper	Acrobat		
Pad	Cut		Calibri	× 11	· A A	= =	 &	Wra	p Text	Genera		
- as	🗧 🍼 Form	at Painter	ns ⊥ ⊓			* = =		- Mer	ge & Cente	r * *	% ,	
	Clipboard	Es.		Font		G	Aligi	nment		5 I	lumber	
	A1		0	<i>f</i> _x 0.071	L							
	A	В	С	D	E	F	G	Н	1	J	К	
1	0.071	0.07	0.072	0.053	0.056	0.059	0.056	0.061				
2	0.06	0.067	0.071	0.083	0.055	0.057	0.055	0.062		кеаа	the inst	tructions
3	0.072	0.063	0.069	0.082	0.072	0.081	0.083	0.077		in tl	ne mess	sage box
4	0.094	0.099	0.099	0.087	0.093	0.097	0.09	0.096		Select	the opt	tion by tick
5	0.095	0.125	0.131	0.123	0.14	0.152	0.149	0.082		Standar	d and l	black at las
6	0.087	0.092	0.073	0.07	0.072	0.062	0.07	0.062		Stanuar		
7	0.08	0.091	0.091	0.094	0.105	0.097	0.074	0.074		Or	in the	starting
8	0.069	0.074	0.075	0.073	0.075	0.089	0.083	0.081				
9	0.088	0.089	0.138	0.141	0.141	0.105	0.105	0.104		He	ere it is	at last
10	0.15	0.158	0.148	0.146	0.155	0.157	0.157	0.173				
11	0.168	0.111	0.115	0.115	0.104	0.102	0.102	0.108				
12	0.114	0.108	0.079	0.081	0.082	0	0	0] Blanl	k readi	ngs	
13			stand	ard rea	dings						_	
14												

12 X 8 = 96 well plate readings

- If you select third option and uploaded file with just 96 readings and missing the sample codes then here is a trick for you to save and organize your file with code.
- First click this well icon to select 96 well plate file and then click upload file. Select same one.
- Now perform the calculation and on the next tab you may see concentrations but no sample codes
- So now come back to project detail tab and select the manual entry radio button and you
 will find the table filled with both sample code and readings. Now click ok submit and
 save the file if you wish.
- Again just click on calculate button and your tables with sample code will be updated.
- Here one more safety trick is used to prevent the user enter the characters or string in the some boxes where the number or integers are must so in that case box will be hanged red until you enter in the proper format.

- And anytime you need hint or help just have a look at message box it may give you some suggestions for the next step
- You can reset or restart the project in case if any strange mishap occurs.

For 384 or any m*n matrix well plates, user has to provide the csv or txt file with the readings sorted sample wise or in manual mode i.e., using first or second options only.

Step 4: Set the Outlier limit

- You can check the outliers if any, by setting the limits in % co-efficient of variation (%CV) between the replicate readings of same sample.
- By default set value is 10 % C.V.
- It will just notify the outlier detection analysis in message box.
- If you want to edit, you can edit the data in file and reupload it or you can continue with calculation.
- Now select the calculation method and proceed further to calculate the unknown concentration.

Step 5: Calculation of Concentration from Standard(s)

- Give the Dilution factor of sample solutions if any dilution performed or else default value of dilution factor is 1 (No extra dilution)
- Give the units of standard and sample concentration to be measured (sample concentrations will be calculated in same unit as of given standard concentrations)

Option 1: from a Single standard by simple cross multiplication

BiAnaCA BioChemical Assay Analyzer and Calculator

HOME NEW DOOTECT			
Project Detail Result Data Sheet Analysis Reports Graphical Plots	Calibraton curve plots		
		Patricia de co	*
Project Detail		Data Upioad box	
Name of the assay: Den	mo	Provide the absorbance data from?	
Experiment performed by:	ur name		a sheet
Experiment performed on:	2/16	Browse valid CSV / TXT file.	(96-well plate data file)
Experimental notes:	ecies name: Albino mice * emp:25 degABC Instrume *	Limit for outliers in replicates	s (in % C.V.): 10 *
Group Patterns		Calculation Panel	
Fill in the details of samples for your assay expe	eriment	Calculation method selection:	=
Number of groups: 4	🔽 + Standard (1 Smo)	Single Standard 💿 Multiple Standards	
Number of sample replicates:	Image: Plank (1 Smp) Image: OK, Start	Enter the Standard concentration:	100 Concentration Absorbance
Group no. Group code No. of samples	Welcome Your name	Mean Standard absorbance reading:	0.087
1 Control 7	You have total 4 Groups and Stand		
2 Treatment A 7	with each of Samples in replicates	Dilution factor for sample readings:	<u>*</u>
4 Treatment A+B 8	Give the data for group code and number of samples in each group:	Standard/sample concentration units:	mg/dl
	K	Number of points in Std. calibration curve	
Reset Submit		Calculate Plot Equation Check Outli	iers 🗸
<u> </u>	II		•
Designed and	developed by team BiAnaCA. I	Dept. of Pharmacoinformatics. NIPER. S.A.S.	Nagar, India.

Option 2: from multiple standard by Standard curve calibration and regression equation

- Enter the number of sample or points in the calibration curve, click the enter and in the side box that many cells will appear for concentration and corresponding readings
- Fill the data and click the Plot Equation, find the result in message box and graphical plot is generated in the last tab of calibration curve plot, view it.

BiAnaCA BioChemical Assay Analyzer and Calculator About O Help Demo HOME NEW PROJECT Project Detai Result Data Sheet Analysis Reports Graphical Plots Calibraton curve plots Project Detai Data Upload box

Name of the assay: Demo	Provide the absorbance data from?
Experiment performed by: Your name	⊙ .txt/.csv file upload ○ Manual entry in data sheet ○ Standard at Start ○ Standard at End ○ Standard at En
Experiment performed on: 8/2/16	Browse valid CSV / TXT file. (96-well plate data file)
Experimental notes:	Limit for outliers in replicates (in % C.V.): 10^{+}
Group Patterns	Calculation Panel
Fill in the details of samples for your assay experiment	Calculation method selection:
Number of groups: 4 ✓ + Standard (1 Smp) Number of sample replicates: 3 ☆ ✓ + Blank (1 Smp) Group no. Group code No. of samples 1 Control 7 2 Treatment A 7 3 Treatment B 8 4 Treatment A+B 8 Reset Submit	 Single Standard () Multiple Standards Enter the Standard concentration: Mean Standard absorbance reading: Dilution factor for sample readings: Standard/sample concentration units: Number of points in Std. calibration curve: Calculate Plot Equation Check Outliers
Designed and developed by team BiAnaCA I	Dept. of Pharmacoinformatics, NIPER, S.A.S.Nagar, India.

Standard Calibration curve plot



- You can view, edit, print, copy or save the plot in different formats from its properties option by right click on the plot.
- Linear regression equation along with the coefficient of determination (R square) is generated

```
Linear regression equation: y = mx + c
```

where,

y=absorbance,

m=slope,

x=concentration,

c=intercept

• Come back to Project detail screen and Click on activated calculate button and find the results on the screen in the next tabbed pane.

BiAnaCA BioChemical Assay Analyzer and Calculator

HOME NEW PROJECT	N. 6. H	
Project Detail Result Data Sheet Analysis Reports Graphical Project Detail	lots Calibraton curve plots	Data Upload box
Name of the assay: Experiment performed by:	Demo Your name	Original Control of Control
Experiment performed on:	8/2/16	Browse valid CSV / TXT file. (96-well plate data file)
Experimental notes:	Species name: Albino mice * Temp:25 degABC Instrume * *	Limit for outliers in replicates (in % C.V.):
Group Patterns Fill in the details of samples for your assay of	experiment	Calculation Panel Calculation method selection: Single Standard Multiple Standards
Number of sample replicates: 3		Enter the Standard concentration: Concentration Absorbance
Group no. Group code No. of samples	Standard Caliration Curve is as: Equation Y = mX+c. Where, Slope m = 0.001 and Intercept (Mean Standard absorbance reading: 200 0.165 300 0.234 Weight of the second line of the seco
2 Treatment A 3 Treatment B 4 Treatment A+B	06 8 Coefficient R-sqare value is: 0.9 8 Now click Calculate for calculati samples concentraion from this	Standard/sample concentration units: mg/d
Read me	> Jon. •	Number of points in Std. calibration curve: 5 Calculate Plot Equation Check Outliers Plot Equation
(Designed	m and developed by team BiAnaCA	Dent. of Pharmacoinformatics. NIPER, S.A.S.Nagar, India

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DIAL		siocnen		say Ana	liyzer a	na caic	ulator	Abou	t 🕜 Help	U Demo
ME NEW PR	ROJECT									
oject Detail	Result Data Sheet Analysis	Reports Graphical Plo	ts Calibraton curve pl	lots						
. no	Sample Code	Reading 1	Reading 2	Reading 3	Mean Abs.	Std. Error	std. dev	% C.V Co	ncentration n	otes
	CTRL-C-1-H	0.071	0.07	0.072	0.071	0.001	0.001	1,409	92.289	
	CTRL-C-1-M	0.053	0.056	0.059	0.056	0.002	0.003	5.357	74.496	
	CTRL-C-1-T	0.056	0.061	0.06	0.059	0.002	0.003	1.101	70.055	
	CTRL-C-1-HM	0.067	0.071	0.083	0.074	0.005	0.008	11.303	95.453 *	*Outliers
	CTRL-C-1-HT	0.055	0.057	0.055	0.056	0.001	0.001	2.0/4	/7.1	
	CTRL-C-2-H	0.062	0.072	0.063	0.066	0.003	0.005	8.387	85.963	
	CTRL-C-2-M	0.069	0.082	0.072	0.074	0.004	0.007	9.157	96.244	
	HSD-S-3-H	0.081	0.083	0.077	0.08	0.002	0.003	3.803	103.361	
	HSD-S-3-M	0.094	0.099	0.099	0.097	0.002	0.003	2.966	123.527	
	HSD-S-3-T	0.087	0.093	0.097	0.092	0.003	0.005	5.451	117.596	
	HSD-S-4-HM	0.09	0.096	0.095	0.094	0.002	0.003	3.432	119.177	
	HSD-S-4-HT	0.125	0.131	0.123	0.126	0.002	0.004	3.296	157.928	
	HSD-S-4-H	0.14	0.152	0.149	0.147	0.004	0.006	4.248	182.444	
	HSD-S-5-M	0.082	0.087	0.092	0.087	0.003	0.005	5.747	111.269	
	UNX-U-6-M	0.073	0.07	0.072	0.072	0.001	0.002	2.131	93.08	
	UNX-U-6-T	0.062	0.07	0.062	0.065	0.003	0.005	7.143	84.777	
,	UNX-U-6-HM	0.08	0.091	0.091	0.087	0.004	0.006	7.272	111.665	
	UNX-U-6-HT	0.094	0.105	0.097	0.099	0.003	0.006	5.763	125.109	
	UNX-U-6-MT	0.074	0.074	0.069	0.072	0.002	0.003	3.991	93.871	
	UNX-U-7-H	0.074	0.075	0.073	0.074	0.001	0.001	1.351	95.848	
	UNX-U-7-M	0.075	0.089	0.083	0.082	0.004	0.007	8.531	105.734	
	UNX-U-T-T	0.081	0.088	0.089	0.086	0.003	0.004	5.069	110.083	
	UNX +HSD-US-8-H	0.138	0.141	0.141	0.14	0.001	0.002	1.237	174.14	
	UNX+HSD-US-8-M	0.105	0.105	0.104	0.105	0	0.001	0.552	132.226	
	UNX+HSD-US-8-T	0.15	0.158	0.148	0.152	0.003	0.005	3.481	188.375	
	UNX+HSD-US-9-HM	0.146	0.155	0.157	0.153	0.003	0.006	3.838	189.166	
,	UNX+HSD-US-9-HT	0.157	0.173	0.168	0.166	0.005	0.008	4.931	204.982	
	UNX +HSD-US-9-MT	0.111	0.115	0.115	0.114	0.001	0.002	2.032	142.902	
)	UNX+HSD-US-10-H	0.104	0.102	0.102	0.103	0.001	0.001	1.125	129.854	
	UNX+HSD-US-10-T	0.108	0.114	0.108	0.11	0.002	0.004	3.149	138.553	
1	STD	0.079	0.081	0.082	0.081	0.001	0.002	1.894	103.756	
Home	This is the result data for	sample wise analysis.	Now, To Perform	the Next group wis	e data Analysis Click	on Data Analysis, o	r to change click on	Home to go back . D	ata Analysis :	a ^{next}
		Designed and	d developed by	v team BiAnaC	A, Dept. of Pha	rmacoinforma	itics, NIPER, S.	A.S.Nagar, India.	cuck	3

Step 6: Click "Calculate" button for Result data sheet containing sample wise concentration table

Step 7: Upon clicking "next", "Analysis Reports" tab will be opened

Analysis reports include Group-wise concentration data and Column statistics.

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NEW PROJECT							
	to Charak Analysis Peng	rte Compliant Plate Caliba					
up Conc Data	na sheet maryas repa		atori cui ve piots		Statistical Test Parameters		
rp1: Control	Grp2: Treatme	nt A Grp3: Trea	tment B	Grp4: Treatment A+B	Select your test		•
0.0	803.3333	/16.666/		1400.0			
0.0	9/3.3333	040.6667		1040.000/	Concontration	First coloct the group	as and then
6.667	923.3333	0/3.3333		1520.0	Concentration	perform test.	os anu crien
6 6667	1263 3322	700.000/		1520.0007	Select all		
6 6667	1470.0	720.000		1136 6667	Group wise data		
13 3333	870.0	823 3333		1026 6667			
5.5555	070.0	860.0		1100.0	Group 1		
<u> </u>				110010	Group 2	Alpha value (α):	
					Group 3		
					Group 4	0.05 (95% C.I.)	T
					Group 5	P value: 💿 One-tail	Two-tail
Group wise statistics	(Column Statistics)	·		Plot gr	aph Analysis report hox	Alpha	Perform
arameter	Grp 1: Control	Gro 2: Treatment A	Grn 3: Treatme	nt B Gro 4: Treatment	A+B starts new project	t-test results	
ample size	7	7	8	8			
an	650.476	1034.286	796.25	1302.083	Parameter Value	Parameter	Value
d.Error	30.967	91.051	38.773	89.134	GrootdusuCal	Group 2	
d.deviation	81.931	240.899	109.667	252.108	Sample size	Sample size	
ar.Ratio	6712.698	58032.275	12026.786	63558.532	Parameters		
c.v.	12.596	23.291	13.773	19.362	Mean ± S.E.M	Mean ± S.E.M	
inimum	556.667	803.333	646.667	1026.667	Group wise da	ta	-
quartile	560.0	870.0	718.333	1060.0	One (Thus to P	Al-1	
guartile	656.667	936.667	781.667	1268.333	Une-/Two- tail	Alpha value	
edian	736.667	1263.333	870.0	1525.0	P value	P value summary	
aximum	743.333	1470.0	986.667	1660.0	t-statistic	Null Ho rejection?	
						1	



Step 8: Click "Plot graph" for graphs of Group-wise and Sample-wise data



Right click on the graph and explore the properties, edit, view or save graphs

Step 9: Perform Statistical tests

- Select your test first:
 - Unpaired Student's t-test (for equal variance)
 - Paired Student's t-test
 - Unpaired t-test (unequal variance) (Welch correction)
 - One-sample Student's t-test
 - One-way ANOVA
 - Z-factor
- · Select alpha value from List
 - 0.05 (95% Confidence interval)
 - 0.01 (99% Confidence interval)
 - 0.001 (99.9% Confidence interval)
 - 0.1 (90% Confidence interval)
- Select P value
 - One-tail
 - Two-tail

ME NEW PROJECT								
oject Detail Result Dat oun Conc Data	a Sheet Analysis Reports	Graphical Plots Calibraton c	irve plots		Statistical Test Paramete	re.		
					bia abatan reser aramete			
Grp 1: Control	Gro2: Treatmen	t A Gro3: Treat	ment B (Grn4: Treatment A+B	Select your test			•
710.0	803, 3333	716.6667	1	400.0	Select your test			
60.0	973.3333	646,6667	1	046.6667	Unnaired Stude	nt's t-test (equal v	ariance)	
90.0	923.3333	873.3333	1	520.0	Paired Student's	st-test		
/36.6667	936.6667	986.6667	1	526.6667	Unpaired Stude	nt's t-test (unequa	l variance) (Welch correct	tion)
56.6667	1263.3333	723.3333	1/	660.0	One-sample Stu	dent's t-test		,
56.6667	1470.0	740.0	1	136.6667	One-way ANOV	A		
43.3333	870.0	823.3333	1	026.6667	Z-factor			
		860.0	1	100.0	Group 2		Alpha value (α):	
					Group 3			
					Group 4		0.05 (95% C.1.)	T
					Group 5		P value: 💿 One-tail	Two-tail
				-			Alaba	Desferre
			· · ·				мрта	Periorini
Group wise statistics (Column Statistics)			Plot graph	Analysis report box			
						t-te:	t results	
Parameter	Grp 1: Control	Grp 2: Treatment A	Grp 3: Treatment B	Grp 4: Treatment A+B	Parameter	Value	Parameter	Value
Sample size	7	7	8	8	Crearen 1		Common 2	1
/lean	650.476	1034.286	796.25	1302.083	Group 1		Group 2	
Std.Error	30.967	91.051	38.773	89.134	Sample size		Sample size	
std.deviation	81.931	240.899	109.667	252.108	Mean + S E M		Mean + S E M	-
Var.Ratio	6712.698	58032.275	12026.786	63558.532	Hear 1 0.1.14		Frear 1 0.2.11	
\$C.V.	12.596	23.291	13.773	19.362				
iinimum	556.667	803.333	646.667	1026.667	One-/Two-tail		Alpha value	
21 quartile	560.0	8/0.0	/18.333	1060.0	She / I we tall		Inpila value	
23 quartile	056.667	936.667	/81.667	1268.333	P value		P value summary	
Megian	/36.667	1263.333	8/0.0	1525.0	t-statistic		Null Ho rejection?	
	/43.333	14/0.0	986.667	1000.0	· statistic		itali no rejectioni	
Maximum								



(i) To perform Student's t-test or other t-tests, select two groups, alpha value, one or two tail and click "Perform"

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E NEW PROJECT									
ject Detail Result Da	ata Sheet Analysis Repor	rts Graphical Plots Calib	oraton curve plots						
oup Conc Data					Statistical Test Paramete	rs			
Srp 1: Control	Grp2: Treatmer	nt A Grp3: Tre	atment B	Grp4: Treatment A+B					
10.0	803.3333	716.6667		1400.0	Paired Student	's t-test		•	
50.0	973.3333	646.6667		1046.6667					
¥0.0	923.3333	873.3333		1520.0	Select the groups	for test:	Paired t-test:		
36.6667	936.6667	986.6667		1526.6667	Select all		Select two groups of they should have say	only me cample	
56.6667	1263.3333	723.3333		1660.0	_ Select all		size	me sample	
56.6667	1470.0	740.0		1136.6667	First group is co	ntrol group: 1 🚔	Select Alpha and P v	<i>r</i> alue	
43.3333	870.0	823.3333		1026.6667					
		860.0		1100.0	Group 1: Contro	ol			
					Group 2: Treatr	nent A	Alpha value (α):		
					Group 3: Treatr	nent B	0.05 (95% C.I.)	_	
					Group 4: Treatr	nent A+B	P 1 0 0 1 1		
				•			P value: 🔘 One-tai	I wo-tail	
Group wise statistics	s (Column Statistics)			Plot graph			Alpha	Perform	
					Applusis report how		13		
arameter	Grp 1: Control	Grp 2: Treatment A	Grp 3: Treatmen	t B Grp 4: Treatment A+B	Analysis report box	t-test res	ults		
ample size	7	7	8	8					
lean	650.476	1034.286	796.25	1302.083	Parameter	value	Parameter	value	
td.Error	30.967	91.051	38.773	89.134	Group 1	Control	Group 2	Treatment A	Ê
cd.deviation	81.931	240.899	109.667	252.108	Sample size	7	Sample size	7	1
ar.Ratio	6712.698	58032.275	12026.786	63558.532	Marris	(50.47(2)) 20.0(7	Marris	1024 2057 + 04 05	
	12.596	23.291	13.773	19.362	Mean ± S.E.M	650.4762 ± 30.967	Mean ± S.E.M	1034.2857 ± 91.05.	
c.v.	556.667	803.333	646.667	1026.667					-
C.V. inimum	560.0	870.0	718.333	1060.0	One-/Two-toil	Two Tail	Alpha value	0.05	
C.V. inimum 1 quartile		936.667	781.667	1268.333	one-/ i wo- tall	1.00 1 411	mpna value	0.03	-11
C.V. inimum 1 quartile 3 quartile	656.667	1763 333	870.0	1525.0	P value	0.00179	P value summary	**	
C.V. inimum 1 quartile 3 quartile edian	656.667 736.667	1203.333		12 6 6 7 7					-
C.V. nimum quartile guartile dian ximum	656.667 736.667 743.333	1470.0	986.667	1000.0	t-statistic	-3.99082	Null Ho rejection?	can be rejected	-

In Results table, you can find sample size, mean ± SEM, P value, T Statistic, Significance, Null Hypothesis rejection, deg. of freedom etc.

oject Detail Result Di	ata Sheet Analysis Repo	rts Graphical Plots Calib	raton curve plots						
roup Conc Data					Statistical Test Paramete	ers			
Grp 1: Control	Grp2: Treatme	nt A Grp3: Trea	atment B Gr	rp4: Treatment A+B					
710.0	803.3333	716.6667	14	00.0	Paired Student	's t-test		-	
560.0	973.3333	646.6667	10-	46.6667					
590.0	923.3333	873.3333	15	20.0	Select the groups	s for test:	Paired t-test:		
736.6667	936.6667	986.6667	15	26.6667	Calast all		Select two groups of	only	
556.6667	1263.3333	723.3333	16	60.0	Select all		size	me sample	
656.6667	1470.0	740.0	11	36.6667	First group is co	ntrol group: 🔢 🚔	Select Alpha and P	value	
743.3333	870.0	823.3333	10	26.6667					
		860.0	11	00.0	Group 1: Contro	pl			
					Group 2: Treatr	nent A	Alpha value (α):		
					Group 3: Treatr	nent B	0.05 (95% C.I.)	-	
					Group 4: Treatr	nent A+B	Develop of a set		
							P value: 🔘 One-tai	 I wo-tail 	
							10 C		
Group wise statistics	(Column Statistics)			Plot graph			Alpha	Perform	
Group wise statistics	(Column Statistics)			Plot graph	Analysis report hox		Alpha	Perform	
Group wise statistics	(Column Statistics)	Gro 7: Treatment A	Grp 3: Treatment B	Plot graph	Analysis report box	t_test res	Alpha	Perform	
Group wise statistics Parameter Sample size	Grp 1: Control	Grp 2: Treatment A	Grp 3: Treatment B	Plot graph Grp 4: Treatment A+B	Analysis report box	t-test resi	Alpha	Perform	
Group wise statistics Parameter Sample size Mean	Grp 1: Control 7 650.476	Grp 2: Treatment A 7 1034.286	Grp 3: Treatment E 8 796.25	B Grp 4: Treatment A+B 8 1302.083	Analysis report box Parameter	t-test rest	Alpha plts Parameter	Value	Re
Group wise statistics Parameter Sample size Mean Std.Error	(Column Statistics) Grp 1: Control 7 650.476 30.967	Grp 2: Treatment A 7 1034.286 91.051	Grp 3: Treatment E 8 796.25 38.773	Plot graph 3 Grp 4: Treatment A+B 8 1302.083 89.134	Analysis report box Parameter Group 1	Value Control	Appa Parameter Group 2	Value Treatment A	e Re
Group wise statistics Parameter Sample size Mean Std.Error Std.deviation	Grp 1: Control 7 650.476 30.967 81.931	Grp 2: Treatment A 7 1034.286 91.051 240.899	Grp 3: Treatment E 8 796.25 38.773 109.667	Plot graph 8 Grp 4: Treatment A+B 8 1302.083 89.134 252.108	Analysis report box Parameter Group 1 Sample size	t-test rest Value Control 7	Appa Parameter Group 2 Sample size	Value Treatment A 7	
Parameter Sample size Mean Std.Error Std.deviation Var.Ratio	Grp 1: Control 7 650.476 30.967 81.931 6712.698	Grp 2: Treatment A 7 1034.286 91.051 240.899 58032.275	Grp 3: Treatment E 8 796.25 38.773 109.667 12026.786	Plot graph 3 Grp 4: Treatment A+B 8 1302.083 89.134 252.108 63558.532	Analysis report box Parameter Group 1 Sample size	Value Control 7	Appa Parameter Group 2 Sample size	Value Treatment A 7	e <u>R</u> e
Group wise statistics Parameter Sample size Mean Std.Error Std.deviation Var.Ratio %C.V.	(Column Statistics) Grp 1: Control 7 650.476 81.931 6712.698 12.596	Grp 2: Treatment A 7 1034.286 91.051 240.899 58032.275 23.291	Grp 3: Treatment E 8 796.25 38.773 109.667 12026.786 13.773	Plot graph 8 Grp 4: Treatment A+B 8 1302.083 89.134 252.108 63558.532 19.362	Analysis report box Parameter Group 1 Sample size Mean ± S.E.M	Value Control 7 650.4762 ± 30.967	Parameter Group 2 Sample size Mean ± S.E.M	Value Treatment A 7 1034.2857 ± 91.05	« Re
Group wise statistics Parameter Sample size Mean Std.Error Std.deviation Var.Ratio %C.V. Minimum	(Column Statistics) Grp 1: Control 7 650.476 30.967 81.931 6712.698 12.596 556.667	Grp 2: Treatment A 7 1034.286 91.051 240.899 58032.275 23.291 803.333	Grp 3: Treatment E 8 796.25 38.773 109.667 12026.786 13.773 646.667	Plot graph 8 8 1302.083 89.134 252.108 65556.532 19.362 19.362 1026.667 1026.667	Analysis report box Parameter Group 1 Sample size Mean ± S.E.M	Value Control 7 650.4762 ± 30.967	Apna Parameter Group 2 Sample size Mean ± S.E.M	Value Treatment A 7 1034.2857 ± 91.05	« R e
Group wise statistics Parameter Sample size Mean Std.Error Std.deviation Var.Ratio %C.V. Minimum Ol quartile	(Column Statistics) Grp 1: Control 7 650.476 30.967 81.931 6712.698 12.596 556.667 560.0	Grp 2: Treatment A 7 1034.286 91.051 240.899 58032.275 23.291 803.333 870.0	Grp 3: Treatment E 8 796.25 38.773 109.667 12026.786 13.773 646.667 718.333	Plot graph B Grp 4: Treatment A+8 8 1302.083 85.134 252.108 63558.532 1026.667 1026.67 1060.0	Analysis report box Parameter Group 1 Sample size Mean ± S.E.M	Value Control 7 650.4762 ± 30.967	Parameter Group 2 Sample size Mean ± S.E.M	Value Treatment A 7 1034.2857 ± 91.05	« <u>R</u> e
Group wise statistics Parameter Sample size Mean Std.Error Std.deviation Var.Ratio &C.V. Minimum Q1 quartile Q3 quartile	(Column Statistics) Grp 1: Control 7 650.476 30.967 81.931 6712.698 12.596 556.667 560.0 656.667	Grp 2: Treatment A 7 1034.286 91.051 240.899 58032.275 23.291 803.333 870.0 936.667	Grp 3: Treatment E 8 796.25 38.773 109.667 12026.786 13.773 646.667 718.333 781.667	Plot graph 8 Grp 4: Treatment A+B 8 1302.083 89.134 252.108 63558.532 19.362 1026.667 1060.0 1268.333 1268.333	Analysis report box Parameter Group 1 Sample size Mean ± S.E.M One-/Two- tail	Value Control 7 650.4762 ± 30.967	Parameter Group 2 Sample size Mean ± S.E.M Alpha value	Value Treatment A 7 1034.2857 ± 91.05 0.05	* R ¢
Group wise statistics Parameter Sample size Mean Std.Error Std.deviation Var.Ratio &C.V. Minimum O1 quartile O3 quartile Median	(Column Statistics) Grp 1: Control 7 650.476 81.931 6712.698 12.596 556.667 560.0 656.667 736.667	Grp 2: Treatment A 7 1034.286 91.051 240.899 58032.275 23.291 803.333 870.0 936.667 1263.333	Grp 3: Treatment E 8 796.25 38.773 109.667 12026.786 13.773 646.667 718.333 781.667 870.0	Plot graph 8 902.083 91.134 252.108 63558.532 103.667 1026.667 1060.0 1265.333 1525.0	Analysis report box Parameter Group 1 Sample size Mean ± S.E.M One-/Two- tail P value	test rest Value Control 7 650.4762 ± 30.967	Parameter Group 2 Sample size Mean ± S.E.M Alpha value P value summary	Value Treatment A 7 1034.2857 ± 91.05 	* R ¢

(ii) To perform One-way ANOVA, select more than 2 groups, alpha value, one or two tail and click "Perform".

A pop-up window of ANOVA results appears

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	L CL L Applysis Dopo	rta la la la la la la			
ject Detail Result Da oun Conc Data	ita Sneet Analysis Repu	Graphical Plots Calibra	aton curve plots		Statistical Test Parameters
conc butu					
Grp 1: Control	Grp2: Treatme	nt A Grp3: Trea	tment B G	rp4: Treatment A+B	
710.0	803.3333	716.6667	14	00.0	One-way ANOVA
60.0	973.3333	646.6667	10	46.6667	
90.0	923.3333	873.3333	15	20.0	Select the groups for test: One-way ANOVA: Select more
36.6667	936.6667	986.6667	15	26.6667	than two groups and
56.6667	1263.3333	723.3333	16	60.0	
56.6667	1470.0	740.0	11	36.6667	V First group is control group:
/43.3333	870.0	823.3333	10	26.6667	
		860.0	11	00.0	Group 1: Control
					Group 2: Treatment A Alpha value (α):
					Group 3: Treatment B
					Group 4: Treatment A+B
				-	P value: O One-tail O Two-tail
Course when a table to be	(Column Challetter)			Distansh	Alpha Perform
Group wise statistics	(column statistics)			Plot graph	A3
		Cra 2: Treatment A	Gro 3: Treatment F	Gro 4: Treatment A+B	Analysis report box
Parameter	Grp 1: Control		orp of fredericite	orp in frequience and	
Parameter	Grp 1: Control	Grp 2: freatment A	8	8	
Parameter Sample size Mean	Grp 1: Control 7 650,476	7 1034-286	8	8	Parameter Value Parameter Value
Parameter Sample size Mean Std.Error	Grp 1: Control 7 650.476 30.967	7 1034.286 91.051	8 796.25 38.773	8 1302.083 89.134	Parameter Value Parameter Value Group 1 Message ₩200 marsie Rest
Parameter Sample size Mean Std.Error Std.deviation	Grp 1: Control 7 650.476 30.967 81.931	7 1034.286 91.051 240.899	8 796.25 38.773 109.667	8 1302.083 89.134 252.108	Parameter Volue Purameter Volue Group 1 Message Res (
Parameter Sample size Mean Std.Error Std.deviation Tar.Ratio	Grp 1: Control 7 650.476 30.967 81.931 6712.698	7 1034.286 91.051 240.899 58032.275	8 796.25 38.773 109.667 12026.786	8 1302.083 89.134 252.108 63558.532	Parameter Value Porsmeter Value Group 1 Message Rescue F-value for ANOVA: 17:55243131338694
Parameter Sample size Mean Std.Error Std.deviation Var.Ratio SC.V.	Grp 1: Control 7 650.476 30.967 81.931 6712.698 12.596	7 1034.286 91.051 240.899 58032.275 23.291	8 796.25 38.773 109.667 12026.786 13.773	8 1302.083 89.134 252.108 63558.532 19.362	Parameter Value Parameter Value Group 1 Message Image: Complexity of the second s
Parameter Sample size 4ean Std.Error Std.deviation /ar.Ratio SC.V. 4inimum	Grp 1: Control 7 650.476 30.967 81.931 6712.698 12.596 556.667	ap 2: Treatment A 7 1034.286 91.051 240.899 58032.275 23.291 803.333	8 796.25 38.773 109.667 12026.786 13.773 646.667	8 1302.083 89.134 252.108 63558.532 19.362 1026.667	Parameter Value Parameter Value Group 1 Message Image: Comparison of the state of the sta
Parameter Sample size dean Std.Error Std.deviation /ar.Ratio &C.V. dinimum D1 quartile	Grp 1: Control 7 650.476 30.967 81.931 6712.698 12.596 556.667 560.0	7 1034.286 91.051 240.899 58032.275 23.291 803.333 870.0	8 796.25 38.773 109.667 12026.786 13.773 646.667 718.333	8 1302.083 89.134 252.108 63558.532 19.362 1026.667 1060.0	Parameter Value Purameter Value Group 1 Message Image: Comparison of the state of the sta
Parameter Sample size dean Std.Error Std.deviation /ar.Ratio dc.V. dinimum 21 quartile 33 quartile	Grp 1: Control 7 650.476 30.967 81.931 6712.698 12.596 556.667 560.0 656.667	7 1034.286 91.051 240.899 58032.275 23.291 803.333 870.0 936.667	8 796.25 38.773 109.667 12026.786 13.773 646.667 718.333 781.667	8 1302.083 89.134 252.108 63558.532 19.362 1026.667 1060.0 1268.333	Parameter Value Parameter Value Group 1 Message Image: Comparison of the state of the sta
Parameter Sample size Mean Std.Error Std.deviation Var.Ratio &C.V. Minimum D1 quartile D3 quartile dedian	Grp 1: Control 7 650.476 30.967 81.931 6712.698 12.596 556.667 560.0 656.667 736.667	Op2 Treatment A 7 1034.286 91.051 240.899 58032.275 23.291 803.333 870.0 936.667 1263.333	8 796.25 38.773 109.667 12026.786 13.773 646.667 718.333 781.667 870.0	8 1302.083 89.134 252.108 63556.532 19.362 1026.667 1060.0 1268.333 1525.0	Parameter Value Parameter Value Group 1 Message mer≪a= Res/0 Sample size F-value for ANOVA: 17.55243131318694 P value for ANOVA: 17.55243131318694 Mean ± S.E P value for ANOVA: 17.55243131318694 P value for ANOVA: 17.55243131318694 One-/Two-tr At significance level 0.05. Null Ho can be rejected. E One-/Two-tr OK OK
Parameter sample size fean td.Error id.deviation 'ar.Ratio C.V. finimum 1 quartile i3 quartile iedian laximum	Grp 1: Control 7 650.476 30.967 81.931 6712.698 12.596 556.667 556.667 736.667 743.333	application application 7 1034.286 91.051 240.899 58032.275 23.291 803.333 870.0 936.667 1263.333 1470.0 1470.0	8 796.25 38.773 109.667 12026.786 13.773 646.667 718.333 781.667 870.0 986.667	8 1302.083 89.134 252.108 63558.532 19.362 1026.667 1060.0 1288.333 1525.0 1660.0	Parameter Value Parameter Value Group 1 Message Image: Comparison of the state of the sta

ANOVA results can also be viewed in the "Graphical Plots" tab.



Step 10: Click on Print Report to generate it

Report and Result files are generated in a folder dedicated to project name and it will open up directly on your screen





Contact Us

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