Analysis Summary

SUBMISSION DETAILS

Analyzed By	LCMSDATA-2	
Analysis Date	02/27/2023	
Script Version	Script_v5.0	
Acquired Date	2/23/2023	
Project Code	SolidBio MicroDyst	
Instrument	Q2	
Project Leader	Shubha	
Processed By	Marcy Matthews	
Notebook Code	MDM000428	

FILES

MIDA Database(s):

Compound Reports:

 $D: Compound \ Reports \ LJZ230218_Insulin_02-24-23Qv \ Test \ LJZ230218_Insulin_0025ug_IP_MS_rep1Q2-27Pxxx_Compound \ Report B06 rev01.xls$

 $D: Compound \ Reports \\ LJZ230218_Insulin_02-24-23Qv\\ Test1\\ \\ 0008H_LJZ230218_Insulin_D025ug_Std_MS_rep1Q2-27Pxxx_Compound \\ ReportB06rev01.xls$

 $D: Compound\ Reports \\ LJZ230218_Insulin_02-24-23Qv\\ Test1\\ 0002H_LJZ230218_Insulin_D05ug_IP_MS_rep1Q2-27Pxxx_Compound\\ ReportB06rev01.xls$

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 $D: Compound \ Reports \ LJZ230218_Insulin_02-24-23Qv \ Test \ 1 \ 0007H_LJZ230218_Insulin_D2ug_Std_MS_rep1Q2-27Pxxx_Compound \ ReportB06rev01.xls$

ANALYSIS DETAILS

Sample	Body Water	Min. MIDA for EMx
025ug_IP-0003H	1.0%	0.025
025ug_Std-0008H	1.0%	0.025
05ug_IP-0002H	1.0%	0.025
1ng_IP-0004H	1.0%	0.025
2ug_IP-0001H	1.0%	0.025
2ug_Std-0007H	1.0%	0.025

PARAMETERS

Parameter	Value
Rt Difference Filter	1.7
Total Abundance Filter	30000
RMS Error Filter	0.015
DB Score Filter	1.0
Base Peak Abundance Filter	30000
% Theo EM0 Upper Limit	1.4
% Theo EM0 Lower Limit	-0.4
Peptide Std Dev Filter	2.0
Isotopomer Std Dev Filter	0.2
Use All Isotopomers for f	No
Combine peptides	Yes
Min. MIDA for EMx	human25
SILAC Masses	4
Sample:SILAC Ratio Upper Limit	20.0
Sample:SILAC Ratio Upper Limit	0.02
SILAC Std Dev Filter	2.0
Fractionated Samples	Yes
Saturation Limit	10000000.0
Offset Correction Slope	[1.0]
Offset Correction Intercept	[0.0]