Detailed instructions for submitting Metabolon data to the Metabolomics Workbench (NMDR)

Please read the main NMDR tutorial first!!

https://www.metabolomicsworkbench.org/data/ds_tutorial.pdf

Page 7 of NMDR tutorial

(b): Begin the online submission of metadata and results

Start a new study from scratch (most common option, especially for new users) or

use the Metabolon template if the new submission is composed of Metabolon analyses (If your samples were analyzed by Metabolon, you MUST use this option) or

use an existing study as a template for a new submission



(b): Begin the online submission of metadata and results Entering Metabolon data

The Metabolon template on the Metabolomics Workbench has 4 different combined LC/MS methods: Low pH polar (LC/MS Pos early) Low pH Lipophilic (LC/MS Pos late) High pH (LC/MS Neg) HILIC (LC/MS Polar Neg) which correspond to 4 different sections (respectively) of the Metabolon results spreadsheet file: Pos Early Pos Late Neg Polar

Split your Metabolon results (1st column is metabolite names, subsequent columns are sample data) and metabolite metadata (1st column is metabolite names, subsequent columns are Pubchem id, KEGG, SMILES, etc) into these 4 sections based on the "PLATFORM" heading in the spreadsheet. Enter these data in the 4 "Data(Results)" sections of the submission form. Specify units of measurement for the data that you're providing (Unnormalized data, normalized-Imputed Data, log-transformed data, etc.)

PLATFORM in Metabolon results spreadsheet	Workbench template results section	LC/MS method	MS Polarity
Pos Early	1	Low pH polar (LC/MS Pos early)	POS
Pos Late	2	Low pH Lipophilic (LC/MS Pos late)	POS
Neg	3	High pH (LC/MS Neg)	NEG
Polar	4	HILIC (LC/MS Polar)	NEG

"Chemical annotation" tab in Metabolon results file

A b CHEM [D] UB 49 50 62 90 93 98 1111 112 1132 2 136 1417 5 6101 1112 112 1132 2 136 143 171 5 179 180 192 192 192 192 207 2009 211 212 215 216	B C 400 400 400 209 305 305 305 209 305 209 209 209 209 209 209 209 209 209 209 209 209 209 209 209 209 209 209 209 209 209 209 209 400 209 400 209 400 400	OMP_ID}_ 1408 485 27665 38395 37483 528 1417 1549 531 1414 22842 38309 3127 32252	U E ETTER PATH 155357 Amino At 155357 Amino At 155829 Cofactors 143675 Lipid 146548 Energy 146398 Energy 124553 Amino Ac 146400 Lipid 124549 Carbohyc 124689 Lipid 124549 Lipid	VB_PATHWAY VB_PATHWAY VB_PATHWAY VALUE VA	C_SORT TYPE 537 NAMED 545 NAMED 4317 NAMED 2031 NAMED 1459 NAMED 1446 NAMED 272 NAMED 428 NAMED	INCHIKEY SMILES KIDHWZJLNCCCCN putress ATH6HQP NCCCCNC'spermi LDHMAVIFC[N+]1=CC1-met CQ5LTKIX*CCCCCC[C 12,13-1 YNOXCRMCC[C(CC(C 2-meth KPGXRSRH0=C[C(0)-alpha-1 HC72WEIE C01=C(0)-alpha-1 HC72WEIE C01=C(0	CHEMICAL_NAME cine idine ynicotinamide DiHOME ylcitrate ketoglutarate	PLC_NAME putrescine spermidine 1-methylnicotinamide 12,13-DiHOME 2-methylcitrate	CAS 110-60-1 124-20-9 1005-24-9 263399-3 6061-96-	N HEMSPIDE 13837702 1071 9 8305504 5 8412123 7 4573953	HMDB000 HMDB000 HMDB000 HMDB000 HMDB000	KEGG C00134 C00315 C02918 C14829 C02225	PUBCHEM 1045 1102 457 10236635 12898022	PLATFORM Pos Early Pos Early Pos Early Neg Polar
40 40 40 50 55 62 90 93 98 111 1 112 132 133 143 171 171 172 179 180 192 194 197 207 209 211 212 215 216	400 400 209 305 305 209 209 209 209 209 209 209 209 209 209	1408 485 27665 38395 37483 528 1417 1549 531 1414 228429 3829 3127 32352	155357 Amino Av 155305 Amino Av 155829 Cofactors 143675 Lipid 146548 Energy 146398 Energy 124553 Amino Ac 147711 Amino Ac 147400 Lipid 124549 Carbohyc 124689 Lipid 143797 Lipid	2 - Polyamine 2 Polyamine 3 Nicotinate Fatty Acid, TCA Cycle TCA Cycle 2 TCA Cycle 3 Tryptopha 3 Leucine, Is Mevalona	537 NAMED 545 NAMED 4317 NAMED 2031 NAMED 1459 NAMED 1446 NAMED 272 NAMED 428 NAMED	KIDHWZIL NCCCN putres ATHGHQP NCCCCNC(spermi LDHMAVIFC[N+]1=C(1-meth CQSLTKIX/CCCCCC(212,13-C YNOXCRMCC[C](CC](2-meth KPGXRSRHO=C(C(0)=a]pha-I VC2WEE OC1-C(C)	cine cine idine iylnicotinamide DiHOME iylcitrate ketoglutarate	spermidine 1-methylnicotinamide 12,13-DiHOME 2-methylcitrate	110-60-1 124-20-9 1005-24-9 263399-3 6061-96-1	13837702 1071 98305504 58412123 74573953	HMDB000 HMDB000 HMDB000 HMDB000 HMDB000	C00134 C00315 C02918 C14829 C02225	1045 1102 457 10236635 12898022	Pos Early Pos Early Pos Early Neg
50 55 62 90 93 98 111 12 132 2136 143 143 177 5 179 180 189 192 194 197 207 4 208 209 211 212 215 216	400 400 209 305 305 209 209 209 209 209 209 209 209 209 209	485 27665 38395 37483 528 1417 1549 531 1414 22842 38309 3127 32352	155305 Amino Av 155829 Cofactor: 143675 Lipid 146548 Energy 146398 Energy 124553 Amino Ao 147711 Amino Ao 147711 Amino Ao 146400 Lipid 124549 Carbohyo 124689 Lipid	ci Polyamine 5 Nicotinate Fatty Acid, TCA Cycle TCA Cycle 1 Tryptopha 1 Leucine, Is Mevalona	545 NAMED 4317 NAMED 2031 NAMED 1459 NAMED 1446 NAMED 272 NAMED 428 NAMED	ATHGHQP NCCCCNCispermi LDHMAVIFC[N+]1=C(1-meth CQSLTKIX/CCCCCC(212,13- TYNOXCRMCC[C(CC(2-meth KPGXRSRHO=C(C(0)=alpha-) HC7UHEFC1=C(1-c(-)-honore	idine iylnicotinamide DiHOME Nylcitrate ketoglutarate	spermidine 1-methylnicotinamide 12,13-DiHOME 2-methylcitrate	124-20-9 1005-24-9 263399-3 6061-96-	1071 9 8305504 5 8412123 7 4573953	HMDB000 HMDB000 HMDB000 HMDB000	C00315 C02918 C14829 C02225	1102 457 10236635 12898022	Pos Early Pos Early Neg
55 62 90 93 98 111 112 132 133 136 143 171 136 143 177 180 189 192 194 197 207 208 209 211 212 215 216	400 209 305 209 305 209 209 209 209 209 209 209 400 209 209	27665 38395 37483 528 1417 1549 531 1414 22842 38309 3127 32352	155829 Cofactor: 143675 Lipid 146548 Energy 146398 Energy 124553 Amino Ac 147711 Amino Ac 146400 Lipid 124549 Carbohyc 124689 Lipid 143797 Lipid	s Nicotinate Fatty Acid, TCA Cycle TCA Cycle :i Tryptopha :i Leucine, Is Mevalona	4317 NAMED 2031 NAMED 1459 NAMED 1446 NAMED 272 NAMED 428 NAMED	LDHMAVIf C[N+]1=CC1-meth CQSLTKIX/ CCCCCC(C 12,13-C YNOXCRM CC(CC(C 2-meth KPGXRSRHO=C(C(O)=alpha-I HC7HHEF OC1-C(C)	nylnicotinamide DiHOME nylcitrate ketoglutarate	1-methylnicotinamide 12,13-DiHOME 2-methylcitrate	1005-24- 263399-3 6061-96-	9 8305504 5 8412123 7 4573953	HMDB000 HMDB000 HMDB000	C02918 C14829 C02225	457 10236635 12898022	Pos Early Neg
62 90 93 98 111 112 132 136 143 171 172 179 180 189 192 194 194 197 207 208 209 211 222 215 216	209 305 305 209 305 209 209 209 209 209 209 400 209 209	38395 37483 528 1417 1549 531 1414 22842 38309 3127 32352	143675 Lipid 146548 Energy 146398 Energy 124553 Amino Ac 147711 Amino Ac 146400 Lipid 124549 Carbohyc 124689 Lipid 143797 Lipid	Fatty Acid, TCA Cycle TCA Cycle :i Tryptopha :i Leucine, Is Mevalona	2031 NAMED 1459 NAMED 1446 NAMED 272 NAMED 428 NAMED	CQSLTKIX/CCCCCC(C 12,13- YNOXCRMCC(C(CCCC2-meth KPGXRSRHO=C(C(O)=alpha- HC7HLELEOC1-C(C)=honor	DiHOME hylcitrate ketoglutarate	12,13-DiHOME 2-methylcitrate	263399-3 6061-96-	58412123 74573953	HMDB000 HMDB000	C14829 C02225	10236635 12898022	Neg
90 93 98 111 112 135 143 171 172 179 180 189 192 194 197 208 209 211 212 215 216	305 305 209 305 305 209 209 209 209 209 209 209 209 209	37483 528 1417 1549 531 1414 22842 38309 3127 32352	146548 Energy 146398 Energy 124553 Amino Ad 147711 Amino Ad 146400 Lipid 124549 Carbohyd 124689 Lipid 143797 Lipid	TCA Cycle TCA Cycle ti Tryptopha ti Leucine, Is Mevalona	1459 NAMED 1446 NAMED 272 NAMED 428 NAMED	YNOXCRM CC(C(CC(C 2-meth KPGXRSRH O=C(C(O)=alpha-l	nylcitrate ketoglutarate	2-methylcitrate	6061-96-	7 4573953	HMDB000	C02225	12898022	Polar
93 98 111 112 132 136 133 171 172 179 180 189 192 194 197 207 208 209 211 212 215 215	305 209 305 305 209 209 209 209 209 209 209 209 209	528 1417 1549 531 1414 22842 38309 3127 32352	146398 Energy 124553 Amino Ao 147711 Amino Ao 146400 Lipid 124549 Carbohyo 124689 Lipid 143797 Lipid	TCA Cycle i Tryptopha i Leucine, Is Mevalona	1446 NAMED 272 NAMED 428 NAMED	KPGXRSRHO=C(C(O)=alpha-l	ketoglutarate							FUIAI
98 111 112 132 136 143 143 143 171 172 179 180 189 192 194 197 207 208 209 211 220 209 211 222 215 226	209 305 305 209 209 209 209 400 209 209 400 209	1417 1549 531 1414 22842 38309 3127 32352	124553 Amino Au 147711 Amino Au 146400 Lipid 124549 Carbohyo 124689 Lipid 143797 Lipid	ti Tryptopha ti Leucine, Is Mevalona	272 NAMED 428 NAMED			alpha-ketoglutarate	328-50-7	250	HMDB000	C00026	51	Polar
111 112 132 136 143 171 172 179 180 189 192 194 197 207 208 209 211 222 215 216	305 305 209 209 209 209 400 209 209 400	1549 531 1414 22842 38309 3127 32352	147711 Amino Au 146400 Lipid 124549 Carbohyo 124689 Lipid 143797 Lipid	i Leucine, Is Mevalona	428 NAMED	I DEVENTE LE VELLE VOULE	enate	kynurenate	492-27-3	3712	HMDB000	C01717	3845	Neg
112 132 136 143 171 172 179 180 189 192 194 197 207 208 209 211 212 215 216	305 209 209 209 209 400 209 209 209 209	531 1414 22842 38309 3127 32352	146400 Lipid 124549 Carbohyc 124689 Lipid 143797 Lipid	Mevalona		DBXBTMS; CC(C(O)=C 3-bydr	oxvisobutvrate	3-hvdroxvisobutvrate	2068-83-	9 10140307	HMDB000	C01188 CC	11966314	Polar
132 136 143 171 172 179 180 189 192 194 197 207 208 209 211 212 215 216	209 209 209 209 209 400 209 209 209	1414 22842 38309 3127 32352	124549 Carbohyo 124689 Lipid 143797 Lipid	h Chueshusia	3176 NAMED	NPOAOTP CC(CC(0)= 3-bydr	oxy-3-methylglutarate	3-hydroxy-3-methylglutarate	503-49-1	4573695	HMDB000	C03761	1662 5459	Polar
136 143 171 172 179 180 189 192 194 197 207 208 209 211 222 215 216	209 209 209 400 209 209 209 400	22842 38309 3127 32352	124689 Lipid 143797 Lipid	UTUNCOINSIS	1237 NAMED	ACAHGYCOC(C(0)=(3-phos	sphoglycerate	3-nhosnhoglycerate	80731-10	10669764	HMDB000	C00597	724	Neg
143 171 172 179 180 189 192 194 197 207 208 209 211 212 212 215 216	209 209 400 209 209 209 209	38309 3127 32352	143797 Lipid	Primary Bi	3454 NAMED	BHOCOFE O[C@H]1[cholate		cholate	81-25-4	10005701	HMDB000	C00695	221493	Neg
171 172 179 189 192 194 197 207 208 209 211 212 215 216	209 209 400 209 209 400	3127 32352	143737 Lipiu	Fatty Acid	2054 NAMED	IVIEIOXAHCCCCCC///A-bydr	oxynonenal	4-bydroxypopenal	1280/6-6	51620		C21642	5282244	Nog
172 179 180 189 192 194 197 207 208 209 211 212 215 216	400 209 209 400	32352	124615 Nucleotic	Puripo Mo	4121 NAMED	EDGOSTZI O=C1C(N= hypox	anthine	hypoxanthine	68-04-0	768		C00262	700 12520	Neg
172 179 180 189 192 194 197 207 208 209 211 212 215 216	209 209 400	32332	155012 Nucleotic	Purine Me	4121 NAMED	LIVER UPD (NC(N1)=N) guarin		guanina	72 40 5	700		00202	12520962	Dec Farly
179 180 189 192 194 197 207 208 209 211 212 215 216	209 209 400	20200	142740 Linid	Forthe Arial	4184 NAIVIED	VERVSORC CCCCC (C= 0.10 D		guanine	73-40-5	/44	HIVIDBUUU	C14929	135398034	Pos Early
180 189 192 207 208 209 211 212 215 216	400	30399	135060 Lipid	Fatty Acid,	1569 NAMED	AEBROQUUOCICCCCC "	(19-2-6)		203399-3	40142232		C14020	5300040	Neg
189 192 194 197 207 208 209 211 212 215 216	400	1400	122309 Lipia	Long Chair	116 NAMED			Indicate (18:20b)	00-33-3	4444105	HIVIDB000	C01595	5280450	Neg Des Cashi
192 194 197 207 208 209 211 212 215 216	a () ()	1498	155380 Amino Ac	I Lysine Me	116 NAMED	WIXINKLEU: C[N+J(C)(CN6,N6,	ino-trimethyllysine	No,No,No-trimethyllysine	23284-33	- 140379	HIMDB000	<u>C03793</u>	440120	Pos Early
194 197 207 208 209 211 212 215 216	400	37496	150406 Amino Ad	Polyamine	538 NAMED	KLZGKIDSECC(NCCCCN-acet	yiputrescine	N-acetyiputrescine	18233-70	- 109095	HIVIDB000	<u>CU2/14</u>	122350	Pos Early
197 207 208 209 211 212 215 216	305	2829	14/946 Amino Ad	I Methionin	437 NAMED	PYUSHNKI CSCCC(C(CN-form	nyimethionine	N-formy(methionine	4289-98-	3388809	HMDB000	<u>CU3145</u>	439750	Polar
207 208 209 211 212 215 216	209	42382	124939 Amino Ad	.i Methionin	446 NAMED	OC1C(N2CS-aden	osyinomocysteine (SAH)	S-adenosylhomocysteine (SAH)	979-92-0	16788012	HMDB009	<u>C00021</u>	439155	Neg
208 209 211 212 215 216	209	2831	123233 Nucleotid	ePurine Me	4143 NAMED	IVOMOUV NC1=NC=1 adeno:	sine 3',5'-cyclic monophosphate (cAl	adenosine 3',5'-cyclic monophospha	60-92-4	268	HMDB000	<u>C00575</u>	6076	Neg
209 211 212 215 216	209	3108	126001 Nucleotid	ePurine Me	4139 NAMED	XTWYTFM NC1=NC=1 adenos	sine 5'-diphosphate (ADP)	ADP	20398-34	- 5800	HMDB000	<u>C00008</u>	6022	Neg
211 212 215 216	400	32342	155907 Nucleotid	I Purine Me	4140 NAMED	UDMBCSS NC1=NC=Fadenos	sine 5'-monophosphate (AMP)	AMP	149022-2	C10239183	HMDB000	<u>C00020</u>	6083	Pos Early
212 215 216	400	1553	155395 Nucleotid	IePurine Me	4170 NAMED	OLXZPDW NC1=C(N= 2'-deox	xyadenosine	2'-deoxyadenosine	16373-93	- 13135	HMDB000	<u>C00559</u>	13730	Pos Early
215 216	400	1419	155365 Amino Ac	:i Polyamine	556 NAMED	WUUGFSXNC1=NC=15-meth	nylthioadenosine (MTA)	5-methylthioadenosine (MTA)	2457-80-	9 388321	HMDB000	<u>C00170</u>	439176	Pos Early
216	209	558	125963 Cofactors	Nicotinate	4327 NAMED	SDMADEZ NC1=NC=Nadenos	sine 5'-diphosphoribose (ADP-ribose	adenosine 5'-diphosphoribose (ADP	68414-18	- 3674071	HMDB000	<u>C00301</u>	192	Neg
	400	38325	156507 Nucleotic	le Purine Me	4115 NAMED	4000 NC(C1=C(I AICA ri	ibonucleotide	AICA ribonucleotide	3031-94-	5 195	HMDB000	<u>C04677</u>	65110	Pos Early
219	305	1763	147132 Carbohyo	I Pentose Pl	1255 NAMED	YXJDFQJKEO[C@@H] ribose	1-phosphate	ribose 1-phosphate	50-99-7,5	8388373	HMDB000	<u>C00620</u>	439236	Polar
229	402	1110	157891 Lipid	Long Chair	1574 NAMED	YZXBAPSD CCCCC/C= arachic	donate (20:4n6)	arachidonate (20:4n6)	506-32-1	392692	HMDB000	<u>C00219</u>	444899	Pos Late
230	305	57752	213350 Lipid	Fatty Acid	1486 NAMED	JDEPVTUUCCCCC/C= arachic	donoyl CoA	arachidonoyl CoA	799812-9	121402992	HMDB000	<u>C02249</u>	25243941,	Polar
231	400	1638	155435 Amino Ar	:i Urea cycle	483 NAMED	ODKSFYD>N=C(N)NC arginin	ie	arginine	1119-34-3	2 227	HMDB000	C00062	6322	Pos Early
232	305	15497	147954 Amino Ar	i Urea cycle	484 NAMED	KDZOASG(NC(C(O)=Carginin	osuccinate	argininosuccinate	156637-5	8805	HMDB000	C03406	60150382	Polar
233	400	32354	155914 Cofactors	Ascorbate	4348 NAMED	CIWBSHSkOCC(O)C1 ascorb	ate (Vitamin C)	ascorbate (Vitamin C)	134-03-2	10189562	HMDB000	D00018,CC	54670067	Pos Early
234	400	443	155304 Amino Ar	i Alanine ar	36 NAMED	CKUMWT: NC(C(O)=Casparta	ate	aspartate	56-84-8	5745	HMDB000	C00049	5960	Pos Early
240	209	32197	123321 Amino Ar	i Tyrosine N	181 NAMED	JVGVDSSUOC(C(O)=(3-(4-hy	/droxyphenyl)lactate	3-(4-hydroxyphenyl)lactate (HPLA)	6482-98-	0 9010	HMDB000	C03672	9378	Neg
244	400	55	155295 Nucleotic	le Pyrimidine	4240 NAMED	UCMIRNVINCCC(O)= beta-a	lanine	beta-alanine	107-95-9	5234	HMDB000	C00099	239	Pos Early
249	400	1768	155446 Amino Ar	i Histidine N	84 NAMED	CQOVPNP.OC([C@H]carnos	ine	carnosine	305-84-0	388363	HMDB000	C00386	439224	Pos Early
250	402	2137	158009 Cofactor	Hemoglob	4412 NAMED	QBUVFDK CC1=C(/C=bilivero	din	biliverdin	55482-27	-10628548	HMDB000	C00500	5280353	Pos Late
252	305	1437	147106 Energy	TCA Cycle	1449 NAMED	KDYFGRW OC(CCC(O succina	ate	succinate	110-15-6	1078	HMDB000	C00042	1110	Polar
254	305	542	145710 Lipid	Ketone Bo	1939 NAMED	WHBMMVCC(O)CC(C3-hvdr	oxvbutvrate (BHBA)	3-hvdroxybutyrate (BHBA)	625-72-9	428	HMDB000	C03197 CC	94318,441	Polar
266	402	63	164177 Lipid	Sterol	3193 NAMED	HVYWMO CC1(C2C(Ccholest	terol	cholesterol	57-88-5	4937803	HMDB000	C00187	5997,6432	Pos Late
267	400	34396	156099 Lipid	Phospholi	2234 NAMED	YHHSONZIC[N+](C)(Ccholine	phosphate	phosphocholine	72556-74	119298	HMDB000	C00588	1014	Pos Early
269	209	62804	220430 Amino Ar	Glutathior	593 NAMED	JYKWMJBLO[C@H]1[CoA-al	utathione*	CoA-glutathione*	6477-52-	7 10128089		C00920	46873828	Neg
270	209	46322	145265 Cofactor	Pantothen	4343 NAMED	RGJOEKW CC/COP/O coepzy	/me A	CoA	85-61-0 1	8311	HMDB000	C00010	87642	Neg
270	200	5082	126005 Lipid	Corticoste	3298 NAMED	OMEXVET: CC/C/CC1) cortico	sterone	corticosterone	50-22-6	5550	HMDR000	C02140	5753	Neg
275	400	512	155307 Amino Av	Creatine N	529 NAMED	DDRIAAN(CN(C(N1)= creation	line	creatinine	60-27-5	568	HMDB000	C00791	588	Pos Farly
273	400	25627	156206 Amino A	i Glutathior	576 NAMED	ZUKDVRW/ NC(C(NCC system	widwine	cysteinyldycine	102/6-19	58762	HMDB000	C01410	120/08	Pos Early
210	400	2272	155/60 Nucleotic	Durimiding	1248 NAMED	IERUI VCD NC1-NC/N ortidia	e 5'-mononhosnhate (5'-CMP)	CMD	62-27-6	5414407	HMDR000	000055	6121	Pos Early
202	205	2372	146412 Cark-L.	di Aminosura			e 5 - monophosphate (5 -CIVIP)	dusesamine 6 phosphate	3616 42	0012767		000252	440007	Polor
290	305	21260	147100 Carbonyo	n Aminosug	103 NAMED		amme-o-phosphate	glucosamine-o-phosphate	102102 5	10052204		C00352	440997	Delee
291	305	31200	14/190 Carbohyo	in Giycolysis,	1224 NAMED	VERNOHX: OCIOCICC glucose	e o-priosphate	giucose o-pnospnate	103192-5	5 19952394	HIVIDB000	<u>C00092</u>	5958	Polar
294	305	1469	140438 Carbohyc	II Fructose, I	1354 NAMED	VFRKUHX: OP(OC[C@mannd	ose-o-pnosphate	mannose 6-phosphate	104872-9	419969422	HIVIDB000	<u>CUU275</u>	439198	Polar
296	0.05	1474	14/108 Carbohy	Il Pentose Pl	1247 NAMED	FNZLKVNUOCC([C@Fribulos	se 5-phosphate	ribulose 5-phosphate	18265-46	- 388327	HMDB000	C00199	439184	Polar
297	305	A 17 A 17	A DOA CALLS IN		0460 0400	Manual Disconsister of the		1.	400 70 -					P I I
< > D	305 402	17747	158164 Lipid	Sphingosir	3160 NAMED	WWUZIQCCCCCCC sphine	osine	sphingosine	123-78-4	4510275	HMDB000	C00319	5280335	Pos Late

"Chemical annotation" tab in Metabolon results file: Sort on "Platform" column to get 4 datasets (Pos Early, Pos Late, Neg and Polar)

Α	В	С	D	Е	F	G	Н	1	J		k	<	L		м	N	0	Р	Q	R
CHEM_ID	LIB_ID	COMP_ID	LIB_ENTR	ER_PATH		AY_SORT	ТҮРЕ	INCHIKE	Y SMILI	ES	CHEMICA	AL_NAME	PLOT_NAME		CAS	HEMSPIDE	HMDB	KEGG	PUBCHEM	PLATFORM
62	209	38395	143675	Lipid	Fatty Acid,	2031 NA	AMED	CQSLTKI	¢ ccccc	C(C 12,13	-DiHOME	-	12,13-DiHOME		263399-35	8412123	HMDB000	C14829	10236635	Neg
98	209	1417	124553	Amino Ac	i Tryptopha	272 NA	MED	HCZHHE	F OC1=C	(C= kynur	renate		kynurenate		492-27-3	3712	HMDB000	C01717	3845	Neg
132	209	1414	124549	Carbohyd	l Glycolysis,	1237 NA	MED	IACAHG		D)=(3-pho	osphoglycerate		3-phosphoglycerate	:	80731-10-	10669764	HMDB000	C00597	724	Neg
136	209	22842	124689	Lipid	Primary Bi	3454 NA	MED	BHQCQF	F'O[C@I	H]1[chola	te		cholate	:	81-25-4		HMDB000	C00695	221493	Neg
143	209	38309	143797	Lipid	Fatty Acid,	2054 NA	MED	JVJFIQYA	нссссс	C(/(4-hyd	droxynonenal		4-hydroxynonenal		128946-65	1630	HMDB000	C21642	5283344	Neg
171	209	3127	124615	Nucleotid	Purine Me	4121 NA	MED	FDGQST	ZJ O=C1C	(N=hypo	xanthine		hypoxanthine		68-94-0	768	HMDB000	C00262	790,13539	Neg
179	209	38399	143740	Lipid	Fatty Acid,	2032 NA	MED	XEBKSQS	GCCCCC	/C= 9,10-I	Dihome		9,10-DiHOME	:	263399-34	8142232	HMDB000	C14828	9966640	Neg
180	209	1105	125969	Lipid	Long Chair	1568 NA	MED	OYHQOL	uoc(cc	CCC linole	eate (18:2n6)		linoleate (18:2n6)		60-33-3	4444105	HMDB000	<u>C01595</u>	5280450	Neg
197	209	42382	124939	Amino Ac	i Methionin	446 NA	MED		OC1C(N2C S-ade	enosylhomocyste	eine (SAH)	S-adenosylhomocysteine	(SAH)	979-92-0	16788012	HMDB009	C00021	439155	Neg
207	209	2831	123233	Nucleotid	ePurine Me	4143 NA	MED	ΙνοΜοι	VNC1=N	IC=Naden	osine 3',5'-cyclic	monophosphate (cA	Aladenosine 3',5'-cyclic mor	nophospha	60-92-4	268	HMDB000	<u>C00575</u>	6076	Neg
2 208	209	3108	126001	Nucleotid	lePurine Me	4139 NA	MED	XTWYTF	M NC1=N	IC=Naden	osine 5'-diphosp	ohate (ADP)	ADP	:	20398-34-	5800	HMDB000	<u>C00008</u>	6022	Neg
215	209	558	125963	Cofactors	Nicotinate	4327 NA	MED	SDMADE	Z NC1=N	IC=Naden	osine 5'-diphosp	horibose (ADP-ribos	adenosine 5'-diphosphori	ibose (ADP	68414-18-	3674071	HMDB000	C00301	192	Neg
240	209	32197	123321	Amino Ac	i Tyrosine N	181 NA	MED	JVGVDSS	UOC(C(C	D)=(3-(4-ł	hydroxyphenyl)l	actate	3-(4-hydroxyphenyl)lactat	te (HPLA)	6482-98-0	9010	HMDB000	<u>C03672</u>	9378	Neg
269	209	62804	220430	Amino Ac	i Glutathior	593 NA	MED	JYKWMJ	BLO[C@I	H]1[CoA-	glutathione*		CoA-glutathione*		6477-52-7	10128089		C00920	46873828	Neg
270	209	46322	145265	Cofactors	Pantothen	4343 NA	MED	RGJOEK	v cc(co	P(O coenz	zyme A		CoA	:	85-61-0,18	311	HMDB000	<u>C00010</u>	87642	Neg
272	209	5983	126005	Lipid	Corticoste	3298 NA	MED	OMFXVF	Ticc(c(c	C1) cortic	osterone		corticosterone	!	50-22-6	5550	HMDB000	C02140	5753	Neg
301	209	34302	126135	Cofactors	Ascorbate	4352 NA	MED	SBJKKFF	12OCC(C	10C dehy	droascorbate		dehydroascorbate		490-83-5	812	HMDB000	<u>C05422</u>	440667	Neg
302	209	1114	125714	Lipid	Secondary	3480 NA	MED	KXGVEG	M CC12[0	@(deox)	ycholate		deoxycholate	:	83-44-3	389318	HMDB000	C04483	222528,44	Neg
327	209	2134	124607	Cofactors	Riboflavin	4334 NA	MED	vwwqx	NOC(C(D)C(flavin	adenine dinucl	eotide (FAD)	FAD		146-14-5,8	559059	HMDB000	C00016	643975	Neg
347	209	2849	123235	Nucleotid	lePurine Me	4178 NA	MED	RQFCJAS	X NC1=N	IC2=guan	osine 5'-monopl	hosphate (5'-GMP)	5'- GMP	!	5550-12-9	741	HMDB000	C00144	6804	Neg
348	209	1411	124547	Nucleotid	le Purine Me	4201 NA	MED	YKBGVTZ	Y OC1=C	(N=2'-de	oxyguanosine		2'-deoxyguanosine		961-07-9	163230	HMDB000	C00330	13539859	Neg
355	209	59	123153	Amino Ac	i Histidine N	70 NA	MED	HNDVDC	UNC(C)=Chistid	line		histidine		5934-29-2	752	HMDB000	C00135	6274	Neg
361	209	1123	123191	Nucleotid	ePurine Me	4120 NA	MED	UGQMR	VFO[C@F	H]1[inosir	ne		inosine	!	58-63-9	21241953	HMDB000	C00294	6021	Neg
409	209	1303	123199	Energy	TCA Cycle	1452 NA	MED	BJEPYKJF	Y OC(C(C	D)=(malat	te		malate		6915-15-7	510	HMDB003	C00149,C	C 525	Neg
424	209	1336	125720	Lipid	Long Chair	1511 NA	MED	IPCSVZSS	voc(cc	CCC palmi	itate (16:0)		palmitate (16:0)	!	57-10-3	960	HMDB000	C00249	985	Neg
439	209	1358	125724	Lipid	Long Chair	1515 NA	MED	QIQXTH		CCC stear	ate (18:0)		stearate (18:0)	1	57-11-4	5091	HMDB000	C01530	5281	Neg
452	209	33447	126083	Lipid	Long Chair	1530 NA	MED	SECPZKH	в ссссс	C\C palmi	itoleate (16:1n7))	palmitoleate (16:1n7)		373-49-9	393216	HMDB000	C08362	445638	Neg
461	209	42109	126986	Energy	Oxidative	1469 NA	MED	NBIIXXV	JZO=P(O)(O)phos	phate		phosphate		7664-38-2	1032	HMDB000	<u>C00009</u>	1061	Neg
463	209	597	124535	Carbohyd	di Glycolysis,	1238 NA	MED	DTBNBX	NO=P(O)(O(phos	phoenolpyruvat	e (PEP)	phosphoenolpyruvate (PE	EP)	10526-80-	980	HMDB000	<u>C00074</u>	1005	Neg
482	209	527	123163	Carbohyd	di Glycolysis,	1240 NA	MED	JVTAAEK	c:cc(o)	C(O) lactat	te		lactate		79-33-4	592	HMDB000	<u>C00186</u>	612	Neg
491	209	1651	123217	Cofactors	Vitamin Bt	4451 NA	MED	RADKZD	VI OC1=C	(C)I pyrid	oxal		pyridoxal		65-22-5	1021	HMDB000	<u>C00250</u>	1050	Neg
492	209	5331	152420	Cofactors	Vitamin Bé	4450 NA	MED	NGVDGC	NO=P(O)(O)pyrid	oxal phosphate		pyridoxal phosphate		41468-25-	1022	HMDB000	<u>C00018</u>	1051	Neg
500	209	1827	123225	Cofactors	Riboflavin	4333 NA	MED	AUNGAN	IROC(C(C	D)C(ribofl	lavin (Vitamin B2	2)	riboflavin (Vitamin B2)	;	83-88-5	431981	HMDB000	<u>C00255</u>	493570	Neg
519	209	1365	125728	Lipid	Long Chair	1508 NA	MED	TUNFSR	IVOC(CC	CCC myris	state (14:0)		myristate (14:0)	1	544-63-8	10539	HMDB000	<u>C06424</u>	11005	Neg
522	209	35670	128273	Cofactors	Thiamine	4423 NA	MED	YXVCLPJ	Q [°] CC1=N	C=C thiam	nin diphosphate		thiamin diphosphate		154-87-0	8715	HMDB000	<u>C00068</u>	1132	Neg
523	209	15798	164715	Cofactors	Thiamine	4422 NA	MED	HZSAJDV	VCC1=N	C=C thiam	nin monophospł	nate	thiamin monophosphate		532-40-1	2627905	HMDB000	<u>C01081</u>	1131,3382	Neg
535	209	606	123185	Nucleotid	le Pyrimidine	4220 NA	MED	DRTQHJ	OC1C(N(C uridir	ne		uridine	!	58-96-8	5807	HMDB000	<u>C00299</u>	6029	Neg
536	209	1412	123201	Nucleotid	le Pyrimidine	4236 NA	MED	MXHRCP	NO=C(N	C1=2'-de	oxyuridine		2'-deoxyuridine	1	951-78-0	13118	HMDB000	C00526	13712	Neg
821	209	33442	125870	Nucleotid	le Pyrimidine	4222 NA	MED	HZIOZCL	E>O[C@]	1([l pseud	douridine		pseudouridine		1445-07-4	21403010	HMDB000	<u>C02067</u>	15047	Neg
872	209	2183	124827	Nucleotid	le Pyrimidine	4275 NA	MED	IQFYYKK	MO=C(N	C(C thym	idine		thymidine	1	50-89-5	5585	HMDB000	<u>C00214</u>	5789	Neg
882	209	604	126810	Nucleotid	le Pyrimidine	4278 NA	MED	RWQNB	REO=C1C	(C)= thym	ine		thymine		65-71-4	1103	HMDB000	<u>C00178</u>	1135	Neg
888	209	1642	125736	Lipid	Medium C	1495 NA	MED	GHVNFZ	- ccccc	CCC capra	ate (10:0)		caprate (10:0)	:	334-48-5	2863	HMDB000	<u>C01571</u>	2969	Neg
891	209	1121	125718	Lipid	Long Chair	1513 NA	AMED	KEMQGT	RO=C(O)CC marg	arate (17:0)		margarate (17:0)	!	506-12-7	10033	HMDB000	2259	10465	Neg
892	209	1356	125722	Lipid	Long Chair	1517 NA	MED	ISYWECE	DE CCCCC	CCC nona	decanoate (19:0))	nonadecanoate (19:0)		646-30-0	12071	HMDB000	<u>C16535</u>	12591	Neg
893	209	1118	125716	Lipid	Long Chair	1518 NA	AMED	VKOBVW	xcccc	CCCarach	nidate (20:0)		arachidate (20:0)	!	506-30-9	10035	HMDB000	<u>C06425</u>	10467	Neg
913	209	15586	124633	Carbohyd	lı Glycogen I	1300 NA	MED	GUBGYT	AIO[C@(@H]malto	ose		maltose		6363-53-7	388469	HMDB000	<u>C00208</u>	10991489	Neg
926	209	32489	125850	Lipid	Medium C	1490 NA	AMED	FUZZWV	x(ccccc	C(O capro	oate (6:0)		caproate (6:0)		142-62-1	8552	HMDB000	<u>C01585</u>	8892	Neg
932	209	32492	125852	Lipid	Medium C	1492 NA	AMED	WWZKQ	нссссс	CCC capry	/late (8:0)		caprylate (8:0)		124-07-2	370	HMDB000	<u>C06423</u>	379	Neg
980	209	1361	125726	Lipid	Long Chair	1510 NA	MED	WQEPLU	UO=C(O)CC penta	adecanoate (15:	0)	pentadecanoate (15:0)		10002-84-	13249	HMDB000	<u>C16537</u>	13849	Neg
1021	209	1494	123209	Amino Ac	ci Glutathior	579 NA	AMED	ODHCTX	KIO=C1C	C[C 5-oxo	oproline		5-oxoproline	!	98-79-3	7127	HMDB000	C02237,C	c ⁴ 39685,74	Neg
1024	209	1508	124797	Cofactors	Pantothen	4337 NA	MED	GHOKW	G ⁻ O=C(N	CCC panto	othenate		pantothenate (Vitamin B5	5) :	137-08-6	6361	HMDB000	<u>C00864</u>	6613	Neg
1026	209	1600	124589	Lipid	Phospholi	2239 NA	MED	SUHOOT	K NCCOF	P(O) phos	phoethanolamir	ne	phosphoethanolamine (P	PE)	1071-23-4	990	HMDB000	<u>C00346</u>	1015	Neg
1080	209	48465	154416	Lipid	Eicosanoic	2150 NA	AMED	4000	ccccc	\C= 5-KET	TE		5-KETE		106154-18	1765	HMDB001	C14732	5353355	Neg
$\langle \rangle$	Data I	Key & Expla	anation	Chemica	al Annotatior	Sample	e Meta	Data	Peak Are	ea Data	Batch-norma	lized Data Batch	n-norm Imputed Data	Mass_extrac	ted-norm	Data	Log Transfe	ormed Da	ata	+

"Peak area Data" tab in Metabolon results file

A	B	C	D	E	F	G	H		J	K	L	M	N	0	P	Q	R	S	T	
PARENT_SAMPLE_NAME	49	50	55	62	90	93	98	111	112	132	136	143	171	172	179	180	189	192	194	1
WASH-02116	452223	3895163	579521	468682	2970954	3749072			1863447	16833276	1038381	5527007	46310586	340131	392831	9.1E+09	19713662		911546	299
WASH-02117	506/1/	4583423	828044	161483	3110/1/	3884827	75602	219800	2378842	1/66132/	286/58	4265011	36109629	259006	224002	1.13E+10	26925663	54917	1259469	35
WASH-02118	605861	6158099	610499	537652	35/3398	6397640	78402	400000	2106469	10/42633	24/48/	4459762	42153612	294639	647369	1.28E+10	218/1686	41862	1059919	254
WASH-02119	441/51	4153682	303872	318184	4430110	3640053	68/65	109383	1908509	16220075	1898216	518338	38699707	202536	208805	6.51E+09	21386361	36088	1055856	31
WASH-02120	805913	6700700	1282324	126101	4083535	2662570	08044 E019E	107900	2423502	10239975	408449	1080034	39559541	208/11	282907	1.125.10	232/1494	46041	1070020	28
WASH-02121	1210522	10746472	1207400	117547	2025199	1696390	50105	122026	2010317	110006400	E16442	E 490E 22	27202751	293227	102622	0.155.00	23001314	40941 E110E	1070039	16
WASH-02122	832389	5788404	1615321	84901	2941955	3631780	85685	155050	2163237	12449156	283164	9421384	46654715	302389	105052	1 22E+10	27086612	68635	1235271	26
WASH-02125	929364	7668152	1418806	250094	2509682	3774100	72811	254419	22103237	13282071	1184432	1403858	39859283	312510	248208	5.09F+09	24331196	66361	1328296	20
WASH-02125	281687	2638282	531414	278891	1450477	5561934	72011	212428	1061857	6827723	140095	1256697	38294424	243503	209622	9.82E+09	13786631	26364	704702	190
WASH-02127	347989	2962047	776812	230655	2937948	6191605	41842	201933	1334843	6287164	593986	464709	41737056	255850	217231	2.63E+09	18826021	37906	914876	250
WASH-02128	322323	2502108	541188	200000	781021	3148661	12012	202000	1139584	8984219	163264	1449207	32051659	224466	104617	7.42E+09	11604740	19991	598135	20
WASH-02129	322225	3387646	1029117	141466	4079801	7050846	26471		1579708	9515811	11880905	2280852	47597637	188203	101673	1.01F+10	15902259	33533	880489	32
WASH-02130	541158	4761027	1032922	374300	4432035	7120633	81898	424133	2732850	5830215	293444	2252090	43984500	274982	246771	1.03E+10	20401007	55095	1165911	29
WASH-02131	1351471	11226342	2557445	138535	3960003	2600718	76429	373898	2429806	10186090	357261	2270758	43668981	300358	178907	6.74E+09	21104936	83497	1223775	20
WASH-02132	633176	4962499	1048678	238471	2513251	4068241		286872	1678963	7408478	149143	1299512	36857444	267244		9.87E+09	21054475	40134	1201646	18
WASH-02133	476658	3239360	977307	118147	946545	2797872	76823	297922	1362085	7770003	97722	3639983	28679425	276517	129554	4.83E+09	9853218	37413	726802	13
WASH-02134	1333529	12280021	1734416	226921	4299507	5015963	104724	430355	2781239	13707284	295223	4742948	40390420	293952	200701	6.81E+09	19733833	72700	1342306	14
WASH-02135	1012028	5694174	1812076	133784	2500685	3773585	80607	297467	1780622	5530944	206118	1839597	47164794	330849		8.48E+09	18965672	64138	1246164	20
WASH-02136	478579	9184206	1055198		5015913	6041946	90058	404987	2203541	9612669	565271	624946	40213318	244919		2.93E+09	20398081	43604	1299973	22
WASH-02137	767842	7094598	1495681	267037	3291609	4501810	116855	403515	2282611	8698422	1684768	2918166	52139203	271598	158700	8.49E+09	21673162	51912	1299340	280
WASH-02138	360871	6576576	985798	226303	113736	1180590	72911	232240	870818	2054978		345235	42207487	298546	339363	4.47E+09	3209201	49824	466205	12
WASH-02139	286139	4562681	833049	280938	1652145	2503899	68502	518281	1013587	3028341		2125700	65404478	392940	448285	8.99E+09	9156338		938358	16
WASH-02140	537298	8014111	1070191	368687	944889	1602599	52360	363098	1405278	2964360	122178	1066450	74314550	223748	425993	1.43E+10	8010512		823900	254
WASH-02141	380950	4036562	1097747	291444	726152	1806675	73149	350249	1135204	3366571	123030	2767905	75358930	394703	258210	6.47E+09	6521075	21745		200
WASH-02142	531362	4969552	1074331	197660	2754975	2620883	147219	779053	1698283	2470274	37690	1396714	95477943	558123	263579	9.41E+09	6724135	119504	1420421	229
WASH-02143	350808	4267671	1010103	162104	1522391	2366760	32446	521508	991234	2439319		2570266	63411455	478931	309859	4.7E+09	8094222	36037	1061621	20
WASH-02144	508999	5128872	1035422	640576	2581054	3249721	38858	344780	1100873	4220228	107887	1647480	80594236	313681	492380	1.56E+10	12845859		953729	299
WASH-02145	594570	7311398	1114854	242101	1053573	2560641	97967	390064	4791523	11201316	47333	2580081	72326128	230203	202057	8.75E+09	14186025	27221	1133087	24
WASH-02146	534526	5542335	1327826	119966	1612459	3330442	90879	456035	2292206	17647299		381029	70877806	224707	180199	9.62E+09	17446537	22125	1303324	25
WASH-02147	473407	5294052	928595	147430	1489568	1884485	108889	599693	1127689	4211967	33224	1427310	82158155	495653	293791	8E+09	8368288	81018	870504	16:

"Peak area Data" needs to be transposed into a new tab (Transposed peak area data) and chem_id needs to be mapped to chemical name and Platform (from Chemical annotation tab)

A	В	С	D E	F	G	Н	- E	J K	L	М	N	0	Р	Q	R			
CHEM_ID	CHEMICAL_NAME	PLATFORM	I PLATFORM	CHEMICAL_NAME	PARENT_SAMPLE_NAME	E WASH-02116 W	ASH-021WAS	SH-021WASH	I-021WASH-02	1WASH-021W	VASH-021V	WASH-021	WASH-021V	VASH-021	WASH-021WAS			
62 12	2,13-DiHOME	Neg	Pos Early	putrescine	49	452223	506717 60	05861 441	805913	594692	1319533	832389	929364	281687	347989 32			
98 ky	nurenate	Neg	Pos Early	spermidine	50	3895163 4	583423 615	58099 4153	6989570	6790709 10	0746472	5788404	7668152	2638282	2962047 250			
132 3-	phosphoglycerate	Neg	Pos Early	1-methylnicotinamide	e 55	579521	828044 61	10499 363	1282324	1086786	1387408	1615321	1418806	531414	776812 54			
136 ch	nolate	Neg	Neg	12,13-DiHOME	62	468682	161483 53	37652 318	3184 531691	. 126191	117547	84901	250094	278891	230655			
143 4-	hydroxynonenal	Neg	Polar	2-methylcitrate	90	2970954 3	110717 357	73398 4430	4083535	1568864	3035188	2941955	2509682	1450477	2937948 78			
171 hy	ypoxanthine	Neg	Polar	alpha-ketoglutarate	93	3749072 3	884827 639	97640 3640	0053 5727567	2662570	4686280	3631780	3774100	5561934	6191605 314			
179 9,3	10-DiHOME	Neg	Neg	kynurenate	98		75602 7	78402 68	68644	50185	52258	85685	72811		41842			
180 lin	noleate (18:2n6)	Neg	Polar	3-hydroxyisobutyrate	111		219800	109	383 187985	127327	133036		254419	212428	201933			
197 S-a	adenosylhomocysteine (SAH)	Neg	Polar	3-hydroxy-3-methylgl	ut 112	1863447 2	378842 210	06469 1908	3509 2423502	2016517	2175406	2163237	2210292	1061857	1334843 113			
207 ad	denosine 3',5'-cyclic monophosphate	e (cAMP) Neg	Neg	3-phosphoglycerate	132	16833276 17	661327 1074	42633 7755	6479 16239975	26838468 13	1988649 1	12449156	13282071	6827723	6287164 898			
208 ad	denosine 5'-diphosphate (ADP)	Neg	Neg	cholate	136	1038381	286758 24	47487 1898	468449	654614	516443	283164	1184432	140095	593986 16			
215 ad	denosine 5'-diphosphoribose (ADP-r	ribose) Neg	Neg	4-hydroxynonenal	143	5527007 4	265011 4											
240 3-	(4-hydroxyphenyl)lactate	Neg	Neg	hypoxanthine	171	46310586 36	109629 42			1 A								
269 Co	oA-glutathione*	Neg	Pos Early	guanine	172	340131	259006	F2		✓ J <mark>x</mark> =VL	_OOKUP(G	i2,A:B,2,FA	ALSE)					
270 со	penzyme A	Neg	Neg	9,10-DiHOME	179	392831	224002	A		В			C		DE	F	G	н
272 со	orticosterone	Neg	Neg	linoleate (18:2n6)	180	9098076797 1.	13E+10 1.	1 CHEM I	D	CHEMICA			ΡΙΑΤΕ	DRM	PLATEOR		CHEM ID	WASH-02116 WASH-021WAS
301 de	ehydroascorbate	Neg	Pos Early	N6,N6,N6-trimethyllys	sir 189	19713662 26	925663 21	2 6	- 2 12 13-DiHO	MF			Neg		Pos Farly	nutrescine	49	452223 506717 60
302 de	eoxycholate	Neg	Pos Early	N-acetylputrescine	192		54917	3 0	8 kynurenate				Neg		Pos Early	spermidine	50	3805163 4583422 615
327 fla	avin adenine dinucleotide (FAD)	Neg	Polar	N-formylmethionine	194	911546 1	259469 1	1 13	2 2 nhocmhor	theoreto			Neg		Pos Early	1 mothylniaetinomide	50	E70E21 828044 61
347 gu	uanosine 5'-monophosphate (5'-GM	1P) Neg	Neg	S-adenosylhomocyste	eir 197	2992368 3	525992 2	4 15 F 12	sz s-priospriog	giycerate			Neg		POSEATIY	12 12 Dillona	55	379321 828044 01
348 2'-	-deoxyguanosine	Neg	Neg	adenosine 3',5'-cyclic	m 207	420775	349546	5 13	o cholate				Neg		Neg	12,13-DIHOIVIE	62	408082 101483 53
355 his	istidine	Neg	Neg	adenosine 5'-diphosp	h: 208	17829829 19	567207 19	6 14	3 4-nydroxyn	onenal			Neg		Polar	2-methylcitrate	90	2970954 3110717 357
361 ind	osine	Neg	Pos Early	adenosine 5'-monoph	10 209	187872060 2.	14E+08 2.	/ 1/	1 hypoxanthi	ne			Neg		Polar	alpha-ketoglutarate	93	3749072 3884827 639
409 m	alate	Neg	Pos Early	2'-deoxyadenosine	211	111429	119838	8 17	9 9,10-DiHON	/IE			Neg		Neg	kynurenate	98	75602 7
424 pa	almitate (16:0)	Neg	Pos Early	5-methylthioadenosir	ne 212	7230096 7	490975 8	9 18	0 linoleate (1	8:2n6)			Neg		Polar	3-hydroxyisobutyrate	111	219800
439 ste	earate (18:0)	Neg	Neg	adenosine 5'-diphosp	h 215	5065001 5	037979 8 ¹	10 19	7 S-adenosylł	nomocysteine	e (SAH)		Neg		Polar	3-hydroxy-3-methylglut	112	1863447 2378842 210
452 pa	almitoleate (16:1n7)	Neg	Pos Early	AICA ribonucleotide	216	65272	135229 1	11 20	7 adenosine	3',5'-cyclic mo	onophosp	hate (cAN	AP) Neg		Neg	3-phosphoglycerate	132	16833276 17661327 1074
461 ph	hosphate	Neg	Polar	ribose 1-phosphate	219	10035389 9	081342 9 1	12 20	8 adenosine	5'-diphospha	ate (ADP)		Neg		Neg	cholate	136	1038381 286758 24
463 ph	hosphoenolpyruvate (PEP)	Neg	Pos Late	arachidonate (20:4n6) 229	946354 1	322323 1 1	13 21	5 adenosine 5	5'-diphospho	oribose (Al	DP-ribose) Neg		Neg	4-hydroxynonenal	143	5527007 4265011 445
482 lao	ctate	Neg	Polar	arachidonoyl CoA	230	85290	81362 1	14 24	0 3-(4-hydrox	yphenyl)lact	ate		Neg		Neg	hypoxanthine	171	46310586 36109629 4215
491 py	yridoxal	Neg	Pos Early	arginine	231	380650966 3.	39E+08 3. 1	15 26	9 CoA-glutath	nione*			Neg		Pos Early	guanine	172	340131 259006 29
492 py	, yridoxal phosphate	Neg	Polar	argininosuccinate	232	116510	127865 1	16 27	0 coenzyme A	4			Neg		Neg	9.10-DiHOME	179	392831 224002 64
500 rib	boflavin (Vitamin B2)	Neg	Pos Early	ascorbate (Vitamin C)	233	9907570 7	928587 12 1	17 27	2 corticostero	one			Neg		Neg	linoleate (18:2n6)	180	9098076797 1.13E+10 1.28
519 m	vristate (14:0)	Neg	Pos Early	aspartate	234	352308882 5.	21E+08 1. 1	18 30	1 debydroaso	orhate			Neg		Pos Farly	N6 N6 N6-trimethyllysir	189	19713662 26925663 2187
522 th	niamin diphosphate	Neg	Neg	3-(4-hydroxyphenyl)la	ac 240	3192734 7	253383 2	10 20	2 de averte de	*-			Nee		Dee Control	N to deside	103	13713002 20323003 2107
523 th	niamin monophosphate	Neg	Pos Early	beta-alanine	244	15556985 19	434152 1730	00907 18599	856 19894174	19443945 20	0062589 1	8117737	20276420 1	6475044	16727922 145			
535 ur	ridine	Neg	Pos Early	carnosine	249	33880836 29	607345 4138	80611 33054	394 19667518	22549756 13	3300957 2	20282700	12400255 2	5782149	35001395 2144			
536 2'-	-deoxyuridine	Neg	Pos Late	biliverdin	250	787256	931612 68	83782 876	356 837303	782139	908806	751253	866764	618602	760121 48			
821 ps	seudouridine	Neg	Polar	succinate	252	58324082 70	123806 5722	28034 1.07E	+08 47986814	60446788 79	9385158 6	5366972	62440908 4	4023117	65429215 3756			
872 th	ymidine	Neg	Polar	3-hydroxybutyrate (B	HI 254	309817	227750 45	56673 220	267 565219	261654	356287	250251	435055	1003685	951145 39			
882 th	, nymine	Neg	Pos Late	cholesterol	266	76831562 87	118536 8140	02571 94741	746 88796872	79139455 83	3496013 8	35110691	97077000 7	1130571	95826566 7359		- L	
888 ca	, aprate (10:0)	Neg	Pos Early	choline phosphate	267	133091811 1.	07E+08 1.3	3E+08 1.32E	+08 1.12E+08	96203030	1.3E+08	1.21E+08	1.27E+08 9	6371978	87655097 816	Use vio	окир ти	Inction
891 m	argarate (17:0)	Neg	Neg	CoA-glutathione*	269	10177646 7	299369 699	94737 7620	778 5361912	8924962	6956630	7482218	6810473	5251092	6810397 67			
892 no	onadecanoate (19:0)	Neg	Neg	coenzyme A	270	1089193 1	006735 226	55115 1164	649 1383771	1647865	1147963	509139	1174901	927291	1568802 91	to man	sham id	+0
893 ar	rachidate (20:0)	Neg	Neg	corticosterone	272	835316	568112 47	77274 518	3920 311460	606640	189446	690947	859007	246732	478892 38	to map of	linem iu	lo
913 m	altose	Neg	Pos Farly	creatinine	275	96237943 97	823434 9669	96778 94271	117 1 09F+08	1 08E+08 99	9343577 9	99584383	1 06E+08 7	8186706	92285583 6759	•	—	
926 ca	aproate (6:0)	Neg	Pos Early	cysteinylglycine	278	435015	409416 78	84755 481	686 517466	432128	366438	438528	348707	178786	247395 1	motahol	ito name	bnc
932 ca	aprvlate (8:0)	Neg	Pos Early	cytidine 5'-mononhos	ap 282	4067428 4	679323 476	59664 4749	556 4512069	5873288	5883627	5088953	4743794	2478710	3143149 290	metabol	ite name	
980 ne	entadecanoate (15:0)	Neg	Polar	glucosamine-6-phosp	ha 290	1733002 2	978093 540	92398 2277	192 6205981	4772909	4116051	3642374	2869232	657988	1268148 86			
1021 5-4	-oxoproline	Neg	Polar	glucose 6-nhosphate	291	30225858 48	045592 8955	57228 381/19	918 1 07F±08	80756267 7	6235750 6	54835766	45152902 1	2743833	28640559 173	nlattorm	1	
1021 5-	antothenate	Nor	Dolar	mannose-6-nhosphate	201	72/12220 11	066607 1793	31970 7890	813 215/2700	15270702 14	6250870 1	3768036	10273702	3523006	6603523 420	Plation	•	
1024 pa	hosphoethanolamine	Neg	Polar	ribulose 5-phosphate	296	19971	16207.8 00	468.9 00		61758	153091	128851	10043.2	88010	182462 3			
1020 pi	.KETE	Neg	Postate	sphingosine	290	7873434 0	085811 044	50731 10/00	694 9023074	8701405 10	0483862	0002609	11380069	6290470	7589568 710			
1000 3-1	Data Kay & Durkansting	micel Annotation	Cample Mate Dat	Deals Are- Data	ransnosed Peak area data	Datah	lized Date	Dat-h		ata 14-	0-10500Z	5552000	ta 1-	5250479	, 555506 / I			
dy Se Accessi	ibility lovestigate	Annotation	Sample Meta Data		tanoposed i cak area data		izeu Data	Datch-noi	ini imputed Da	atd Mass_	_extracted	-norm Da	LOG	nanstorm	ieu Data			

Sort the results table on the "Platform " column

Now you've got all 4 results datasets separated in the "Transposed peak area data" tab and 4 metabolite metadata sets separated in the "Chemical Annotation" tab

E1	\sim : $\times \checkmark f_x$ platform												
	АВ	С	D E	F	G	н	1	J	к	L	М	Ν	0
1 CH	EM ID CHEMICAL NAME	PLATFORM	PLATFORM	CHEMICAL NAME	CHEM ID	WASH-02116	WASH-021	WASH-021	WASH-021	WASH-021	WASH-021	WASH-021	WASH-0
2	62 12,13-DiHOME	Neg	Neg	12,13-DiHOME	62	468682	161483	537652	318184	531691	126191	117547	849
3	98 kynurenate	Neg	Neg	kynurenate	98		75602	78402	68765	68644	50185	52258	856
4	132 3-phosphoglycerate	Neg	Neg	3-phosphoglycerate	132	16833276	17661327	10742633	7755479	16239975	26838468	11988649	124491
5	136 cholate	Neg	Neg	cholate	136	1038381	286758	247487	1898216	468449	654614	516443	2831
6	143 4-hydroxynonenal	Neg	Neg	4-hydroxynonenal	143	5527007	4265011	4459762	518338	1680034	6867986	5480533	94213
7	171 hypoxanthine	Neg	Neg	hypoxanthine	171	46310586	36109629	42153612	38699707	39559541	40404808	37203751	466547
8	179 9,10-DiHOME	Neg	Neg	9,10-DiHOME	179	392831	224002	647369	208805	582967	280209	183632	
9	180 linoleate (18:2n6)	Neg	Neg	linoleate (18:2n6)	180	9098076797	1.13E+10	1.28E+10	6.61E+09	6.52E+09	1.12E+10	9.15E+09	1.22E+
10	197 S-adenosylhomocysteine (SAH)	Neg	Neg	S-adenosylhomocysteine (SAH)	197	2992368	3525992	2541187	3180488	2802713	2519401	1660326	26775
11	207 adenosine 3',5'-cyclic monophosphate (cAMP) Neg	Neg	adenosine 3',5'-cyclic monophosphate (cAMP)	207	420775	349546	436772	289541	353215	376982	421234	2964
12	208 adenosine 5'-diphosphate (ADP)	Neg	Neg	adenosine 5'-diphosphate (ADP)	208	17829829	19567207	19447017	16253400	19331734	19665441	16119902	193059
13	215 adenosine 5'-diphosphoribose (ADP-ribose)	Neg	Neg	adenosine 5'-diphosphoribose (ADP-ribose)	215	5065001	5037979	8302545	4243314	6123915	5955234	3956941	43991
14	240 3-(4-hydroxyphenyl)lactate	Neg	Neg	3-(4-hydroxyphenyl)lactate	240	3192734	7253383	2670469	2805752	4045479	4335856	2422332	32988
15	269 CoA-glutathione*	Neg	Neg	CoA-glutathione*	269	10177646	7299369	6994737	7620778	5361912	8924962	6956630	74822
16	270 coenzyme A	Neg	Neg	coenzyme A	270	1089193	1006735	2265115	1164649	1383771	1647865	1147963	5091
17	272 corticosterone	Neg	Neg	corticosterone	272	835316	568112	477274	518920	311460	606640	189446	69094
18	301 dehydroascorbate	Neg	Neg	dehydroascorbate	301	32454432	21083647	32216119	19388828	36235128	32484331	29583050	226733
19	302 deoxycholate	Neg	Neg	deoxycholate	302	626659	336491.4	282028	402366.3	235622.7	472615	310452	138040
20	327 flavin adenine dinucleotide (FAD)	Neg	Neg	flavin adenine dinucleotide (FAD)	327	8808331	7452926	7844323	6840745	6582727	9071945	6816842	69431
21	347 guanosine 5'-monophosphate (5'-GMP)	Neg	Neg	guanosine 5'-monophosphate (5'-GMP)	347	5485435	6606740	5424499	7769851	6706212	5392608	6976651	70254
22	348 2'-deoxyguanosine	Neg	Neg	2'-deoxyguanosine	348	99267	135890	97149	97107	102892	98502	70779	923
23	355 histidine	Neg	Neg	histidine	355	39813651	40862048	40067084	38140008	41099521	40907574	36531342	421208
24	361 inosine	Neg	Neg	inosine	361	273033982	2.29E+08	2.9E+08	2.3E+08	2.85E+08	2.74E+08	2.37E+08	2.62E+
25	409 malate	Neg	Neg	malate	409	451367710	5.03E+08	3.48E+08	5.29E+08	4.48E+08	3.77E+08	4E+08	5.5E+
26	424 palmitate (16:0)	Neg	Neg	palmitate (16:0)	424	6968920389	8.57E+09	9.49E+09	5.38E+09	4.9E+09	8.29E+09	7.04E+09	8.26E+
27	439 stearate (18:0)	Neg	Neg	stearate (18:0)	439	4750878530	6.45E+09	6.58E+09	3.52E+09	2.68E+09	5.35E+09	5.09E+09	5.95E+
28	452 palmitoleate (16:1n7)	Neg	Neg	palmitoleate (16:1n7)	452	474668336	1.12E+09	1.01E+09	3.21E+08	4.56E+08	1.26E+09	6.75E+08	1.11E+
29	461 phosphate	Neg	Neg	phosphate	461	466110671	4.17E+08	3.84E+08	4.16E+08	3.57E+08	3.8E+08	3.8E+08	4.26E+
30	463 phosphoenolpyruvate (PEP)	Neg	Neg	phosphoenolpyruvate (PEP)	463	1467211	1322880	747880	355454	1137711	2705133	583540	5023
31	482 lactate	Neg	Neg	lactate	482	1147394447	9.75E+08	1.34E+09	7.83E+08	1.13E+09	1.17E+09	9.45E+08	9.66E+
32	491 pyridoxal	Neg	Neg	pyridoxal	491	263607	365083	331220	295021	258874	320411	137846	2060
33	492 pyridoxal phosphate	Neg	Neg	pyridoxal phosphate	492	718694	599769	546396	509240	603413	606029	558852	5846
34	500 riboflavin (Vitamin B2)	Neg	Neg	riboflavin (Vitamin B2)	500	548423	637973	633848	585615	622532	625225	571184	6019
35	519 myristate (14:0)	Neg	Neg	myristate (14:0)	519	166341682	2.33E+08	2.54E+08	88022375	71566430	1.88E+08	1.56E+08	2.56E+
36	522 thiamin diphosphate	Neg	Neg	thiamin diphosphate	522	1659434	1598486	1619093	1247999	1637294	1759398	1480493	17917
37	523 thiamin monophosphate	Neg	Neg	thiamin monophosphate	523	1333949	999492	1032664	900805	1045031	1116207	1141045	11081
38	535 uridine	Neg	Neg	uridine	535	78488682	76840706	84666589	64956103	73458404	71737826	66085122	749766
39	536 2'-deoxyuridine	Neg	Neg	2'-deoxyuridine	536	1394212	1042296	1642898	1036776	1287672	1651282	1024170	14589
40	821 pseudouridine	Neg	Neg	pseudouridine	821	1341272	1435741	1096983	1327491	1633531	1596007	1359471	17692
41	872 thymidine	Neg	Neg	thymidine	872	4096292	3627678	4421489	3945145	4199339	3738773	3489902	37886
42	882 thymine	Neg	Neg	thymine	882	90737		61397		148828	108020	181888	1363
43	888 caprate (10:0)	Neg	Neg	caprate (10:0)	888	6644989	7723423	4727246	5184881	4538658	4252997	6246371	49523
A.A.	001 margarata (17:0)	Neg	Nor		901	122279640	1 705.00	2 105.00	64262010	E714E009	1 645.00	1 355.00	2.145.0

Create separate tabs for Pos Early, Pos Late, Neg and Polar datasets

Delete the chem_id and platform columns

These datasets are then ready to be copied/pasted into the results sections of the online submission form Order of analyses: 1:Pos Early, 2:Pos Late, 3:Neg 4:Polar

А	B C D E F G H I J K	K L M N O P
1 CHEMICAL_NAME	WASH-021WASH-0	SH-021WASH-021WASH-021WASH-021WASH-021WASH-0
2 12,13-DiHOME	468682 161483 537652 318184 531691 126191 117547 84901 250094 2788	78891 230655 141466 374300 1385
3 kynurenate	75602 78402 68765 68644 50185 52258 85685 72811	41842 26471 81898 764
4 3-phosphoglycerate	16833276 17661327 10742633 7755479 16239975 26838468 11988649 12449156 13282071 68277	27723 6287164 8984219 9515811 5830215 101860
5 cholate	1038381 286758 247487 1898216 468449 654614 516443 283164 1184432	
6 4-hydroxynonenal	5527007 4265011 4459762 518338 1680034 6867986 5480533 9421384 1403858	
7 hypoxanthine	46310586 36109629 42153612 38699707 39559541 40404808 37203751 46654715 39859283	Start/Edit Data Submission Examples of study design and data layouts Online Study Submission Tutorial(pdf)
8 9,10-DIHOME	392831 224002 647369 208805 582967 280209 183632 248208	
9 linoleate (18:2n6)	9.1E+09 1.13E+10 1.28E+10 6.61E+09 6.52E+09 1.12E+10 9.15E+09 1.22E+10 5.09E+09	mwTab Identifier: Javier Munoz Briones 24 20230930 161810 Return to start
10 S-adenosylhomocysteine (SAH)	2992368 3525992 2541187 3180488 2802713 2519401 1660326 2677573 2264655	Analysis: Payarsad hasa POSITIVE (1)
11 adenosine 3',5'-cyclic monophosphate (cAMP)	420775 349546 436772 289541 353215 376982 421234 296435 354779	Analysis. Reverseu phase Positive (1)
12 adenosine 5'-diphosphate (ADP)	17829829 19567207 19447017 16253400 19331734 19665441 16119902 19305993 14964916	
13 adenosine 5'-diphosphoribose (ADP-ribose)	5065001 5037979 8302545 4243314 6123915 5955234 3956941 4399186 3111079	
14 3-(4-hydroxyphenyl)lactate	3192734 7253383 2670469 2805752 4045479 4335856 2422332 3298828 4371642	
15 CoA-glutathione*	10177646 7299369 6994737 7620778 5361912 8924962 6956630 7482218 6810473	OPTION 1: Assays containing identified (named) metabolite species
16 coenzyme A	1089193 1006735 2265115 1164649 1383771 1647865 1147963 509139 1174901	
17 corticosterone	835316 568112 477274 518920 311460 606640 189446 690947 859007	Input Data in tab-delimited format in the text area below.
18 dehydroascorbate	32454432 21083647 32216119 19388828 36235128 32484331 29583050 22673380 23774351	First column must contain matchalite names
19 deoxycholate	626659 336491.4 282028 402366.3 235622.7 472615 310452 138040.7 854322.9	First column must contain metabolite names.
20 flavin adenine dinucleotide (FAD)	8808331 7452926 7844323 6840745 6582727 9071945 6816842 6943115 7798098	Subsequent columns must contain sample data with identical sample names as in Study Design submission.
21 guanosine 5'-monophosphate (5'-GMP)	5485435 6606740 5424499 7769851 6706212 5392608 6976651 7025423 5101963	First row must contain sample names
22 2'-deoxyguanosine	99267 135890 97149 97107 102892 98502 70779 92314 110304	
23 histidine	39813651 40862048 40067084 38140008 41099521 40907574 36531342 42120882 38346009	CHEMICAL_NAME WASH-02116 WASH-02117 WASH-02118 WASH-02119 WASH-02120
24 inosine	2.73E+08 2.29E+08 2.9E+08 2.3E+08 2.85E+08 2.74E+08 2.37E+08 2.62E+08 2.44E+08	WASH-02121 WASH-02122 WASH-02123 WASH-02125 WASH-02126 WASH-02127
25 malate	4.51E+08 5.03E+08 3.48E+08 5.29E+08 4.48E+08 3.77E+08 4E+08 5.5E+08 4.66E+08	WASH-02128 WASH-02129 WASH-02130 WASH-02131 WASH-02132 WASH-02133
26 palmitate (16:0)	6.97E+09 8.57E+09 9.49E+09 5.38E+09 4.9E+09 8.29E+09 7.04E+09 8.26E+09 4.22E+09	
27 stearate (18:0)	4.75E+09 6.45E+09 6.58E+09 3.52E+09 2.68E+09 5.35E+09 5.09E+09 5.95E+09 2.76E+09	WASH-02134 WASH-02135 WASH-02136 WASH-02137 WASH-02138 WASH-02139
28 palmitoleate (16:1n7)	4.75E+08 1.12E+09 1.01E+09 3.21E+08 4.56E+08 1.26E+09 6.75E+08 1.11E+09 3.45E+08	WASH-02140 WASH-02141 WASH-02142 WASH-02143 WASH-02144 WASH-02145
29 phosphate	4.66E+08 4.17E+08 3.84E+08 4.16E+08 3.57E+08 3.8E+08 3.8E+08 4.26E+08 4.14E+08	WASH-02146 WASH-02147
30 phosphoenolpyruvate (PEP)	1467211 1322880 747880 355454 1137711 2705133 583540 502348 876215	12.13-DİHOME 468682 161483 537652 318184 531691 126191 117547 84901 250094 278891
31 lactate	1.15E+09 9.75E+08 1.34E+09 7.83E+08 1.13E+09 1.17E+09 9.45E+08 9.66E+08 9.74E+08	239655 141466 374390 138535 238471 118147 226921 133784 267937 226393 289938
32 pyridoxal	263607 365083 331220 295021 258874 320411 137846 206092 304752	
33 pyridoxal phosphate	718694 599769 546396 509240 603413 606029 558852 584694 636620	30608/ 231444 13/668 162164 6465/6 242184 113506 14/438
34 riboflavin (Vitamin B2)	548423 637973 633848 585615 622532 625225 571184 601902 593938	kynurenate /5602 /8402 68/65 68644 50185 52258 85685 /2811
35 myristate (14:0)	1.66E+08 2.33E+08 2.54E+08 88022375 71566430 1.88E+08 1.56E+08 2.56E+08 61419774	41842 26471 81898 76429 76823 104724 80607 90058 116855 72911 68502
36 thiamin diphosphate	1659434 1598486 1619093 1247999 1637294 1759398 1480493 1791758 1595451	
37 thiamin monophosphate	1333949 999492 1032664 900805 1045031 1116207 1141045 1108135 1315156	Units of measurement (required): Peak area
38 uridine	78488682 76840706 84666589 64956103 73458404 71737826 66085122 74976658 65707592	Chits of measurement (required). I cak alca
39 2'-deoxyuridine	1394212 1042296 1642898 1036776 1287672 1651282 1024170 1458939 1250484	
40 pseudouridine	1341272 1435741 1096983 1327491 1633531 1596007 1359471 1769264 1582842	
41 thymidine	4096292 3627678 4421489 3945145 4199339 3738773 3489902 3788662 4008364	View/check metabolite data See examples of metabolite data layout Delete existing metabolite data (this analysis only)
42 thymine	90737 61397 148828 108020 181888 136305 161137	
43 caprate (10:0)	6644989 7723423 4727246 5184881 4538658 4252997 6246371 4952340 4148565	
44 margarate (17:0)	1.33E+08 1.78E+08 2.18E+08 64362010 57145008 1.64E+08 1.35E+08 2.14E+08 46456027	
45 nonadecanoate (19:0)	50970539 56255450 66852816 32521971 19582082 42244172 48098407 81195017 18990075	
46 arachidate (20:0)	1.89E+08 1.42E+08 2.98E+08 75300906 34697639 1.51E+08 1.65E+08 1.61E+08 68157111	OPTION 2: Untergoted assays not containing identified (named) motabolite species
47 maltose	2174658 2006083 3417160 2338841 3252445 2379888 2167201 2609217 1893566 10870	87034 1659934 1406851 668085 1463896 18107
48 caproate (6:0)	2333663 1822125 2153786 1127963 2647126 2312614 3707909 1420820 12056	356/5 94/010 121/1994 1458613 1666602 21101
49 caprylate (8:0)	3718685 4569903 2321915 3791914 1502843 2407206 3338909 3266454 1993363 26963	y6345 Z7/76848 3433998 4653494 1636852 55332/
50 pentadecanoate (15:0)	60510231 63473944 65505725 35437629 35033108 68701976 55108653 1.29E+08 34834407 557197	19/3/ 20484825 53309690 554130/ 7/2/91/3 84 /202
51 5-oxoproline	26775931 32137024 28/31914 27233688 33181674 30233363 27866915 32621677 37524593 248773	//388 2/99689/ 2261/784 30601890 28949121 301481
52 pantothenate	1.99E+U8 2.34E+U8 1.8E+08 1.8E+08 2.23E+08 2.25E+08 1.81E+08 1.98E+08 1.95E+08 1.82E+	2Ł+U8 2.409±U8 1.31±U8 1.54±U8 1.74±U8 1.55±H
53 phosphoethanolamine	11512288 12288800 11411184 11695989 12441503 12859684 11190477 12811819 11019309 101754	/54/2 9/52482 8/84453 12589995 11/US0U1 115458
D4 D-KEIE	3805022 0013588 4881925 1114816 1068496 5908906 49/3058 13304036 836400 17002	1/2/1 - 2/2/1/2 - 1532/1/3 - 2/1/801 - 2/1/8/3 - 2/058
Data Key & Explanation Chemical Annotation	Sample Meta Data Peak Area Data Transposed Peak area data Polar Pos ea	seany Postate patch-normalized Lata batch

Make a copy of the "Chemical annotation" tab and put "Chemical name" on the left This column must EXACTLY match that of results datasets with regard to metabolite names Note that several columns may be deleted for the purpose of submission upload (Metabolon identifiers, etc) Split this table into 4 sub-tables corresponding to Pos Early, Pos Late, Neg and Polar metadata

	B CUDED DATUMAN		D	E CANUES	F	G	Н	KECC	DUDCUT	K
1 CHEMICAL_NAME	SUPER_PATHWAY	SUB_PATHWAY	INCHIKEY	SMILES	CAS	CHEMSPIDER	HMDB	KEGG	PUBCHEM	PLATFORM
2 12,13-DIHOME	Lipid	Fatty Acid, Dihydroxy	CQSLIKIXAJIQGA-FLIBIINWBI		263399-3	58412123	HMDB000 C1	14829	10236635	Neg
3 kynurenate	Amino Acid	Tryptophan Metabolism	HCZHHEIFKROPDY-UHFFFAOYAN	OC1=C(C=CC=C2)C2=NC(C	492-27-3	3/12	HMDB000 CC	01/1/	3845	Neg
4 3-phosphoglycerate	Carbohydrate	Glycolysis, Gluconeogenesis, and Pyruvate Metabolish	n IACAHGYCKIUESG-REWHXWOFAW	OC(C(O)=O)COP(O)(O)=O	80731-10	- 10669764	HMDB000 CC	00597	724	Neg
5 cholate	Lipid	Primary Bile Acid Metabolism	BHQCQFFYRZLCQQ-OELDTZBJBZ	O[C@H]1[C@@]2([H])[C@	81-25-4		HMDB000 CC	00695	221493	Neg
6 4-hydroxynonenal	Lipid	Fatty Acid, Oxidized	JVJFIQYAHPMBBX-UHFFFAOYAI	CCCCCC(/C=C/C=O)O	128946-6	51630	<u>HMDB000 C2</u>	21642	5283344	Neg
7 hypoxanthine	Nucleotide	Purine Metabolism, (Hypo)Xanthine/Inosine containir	FDGQSTZJBFJUBT-UHFFFAOYAJ	O=C1C(N=CN2)=C2N=CN1	68-94-0	768	HMDB000 CC	00262	790,135398638	Neg
8 9,10-DiHOME	Lipid	Fatty Acid, Dihydroxy	XEBKSQSGNGRGDW-YFHOEESVBC	ccccc/c=c\cc(c(cccccc	263399-3	48142232	HMDB000 C1	14828	9966640	Neg
9 linoleate (18:2n6)	Lipid	Long Chain Polyunsaturated Fatty Acid (n3 and n6)	OYHQOLUKZRVURQ-HZJYTTRNBX	OC(CCCCCCC/C=C\C/C=C\	60-33-3	4444105	HMDB000 CC	01595	5280450	Neg
10 S-adenosylhomocysteine (SAH)	Amino Acid	Methionine, Cysteine, SAM and Taurine Metabolism		OC1C(N2C=NC3=C2N=CN=	979-92-0	16788012	HMDB009 CC	00021	439155	Neg
11 adenosine 3',5'-cyclic monophosphate (c/	Al Nucleotide	Purine Metabolism, Adenine containing	IVOMOUWHDPKRLL-UHFFFAOYAA	NC1=NC=NC2=C1N=CN2C	60-92-4	268	HMDB000 CC	00575	6076	Neg
12 adenosine 5'-diphosphate (ADP)	Nucleotide	Purine Metabolism, Adenine containing	XTWYTFMLZFPYCI-KQYNXXCUBP	NC1=NC=NC2=C1N=CN2[C	20398-34	- 5800	HMDB000 CC	00008	6022	Neg
13 adenosine 5'-diphosphoribose (ADP-ribo	seCofactors and Vitamins	Nicotinate and Nicotinamide Metabolism	SDMADEZQMYCSNO-UHFFFAOYAX	NC1=NC=NC2=C1N=CN2C	68414-18	3674071	HMDB000 CC	00301	192	Neg
14 3-(4-hydroxyphenyl)lactate	Amino Acid	Tyrosine Metabolism	JVGVDSSUAVXRDY-UHFFFAOYAV	OC(C(O)=O)CC1=CC=C(O)	(6482-98-0	9010	HMDB000 CC	03672	9378	Neg
15 CoA-glutathione*	Amino Acid	Glutathione Metabolism	JYKWMJBUIXNJOG-JHKACSFWBF	O[C@H]1[C@H](N2C(N=C	6477-52-	10128089	CC	00920	46873828	Neg
16 coenzyme A	Cofactors and Vitamins	Pantothenate and CoA Metabolism	RGJOEKWQDUBAIZ-UHFFFAOYAQ	CC(COP(O)(OP(O)(OCC1O	85-61-0,1	8311	HMDB000 CC	00010	87642	Neg
17 corticosterone	Lipid	Corticosteroids	OMFXVFTZEKFJBZ-HJTSIMOOBD	CC(C(CC1)=CC2=O)(CC2)C	50-22-6	5550	HMDB000 CC	02140	5753	Neg
18 dehydroascorbate	Cofactors and Vitamins	Ascorbate and Aldarate Metabolism	SBJKKFFYIZUCET-UHFFFAOYAZ	OCC(C1OC(C(C1=O)=O)=O	490-83-5	812	HMDB000 CC	05422	440667	Neg
19 deoxycholate	Lipid	Secondary Bile Acid Metabolism	KXGVEGMKQFWNSR-LLQZFEROBK	CC12[C@@]([C@]3([H])C	(83-44-3	389318	HMDB000 CC	04483	222528,440355	Neg
20 flavin adenine dinucleotide (FAD)	Cofactors and Vitamins	Riboflavin Metabolism	VWWQXMAJTJZDQX-UYBVJOGSBL	OC(C(O)C(O)COP(OP(OC[0	(146-14-5,	\$559059	HMDB000 CC	00016	643975	Neg
21 guanosine 5'-monophosphate (5'-GMP)	Nucleotide	Purine Metabolism, Guanine containing	RQFCJASXJCIDSX-UHFFFAOYAQ	NC1=NC2=C(C(N1)=O)N=C	5550-12-9	741	HMDB000 CC	00144	6804	Neg
22 2'-deoxyguanosine	Nucleotide	Purine Metabolism, Guanine containing	YKBGVTZYEHREMT-KVQBGUIXBI	OC1=C(N=CN2C(CC3O)OC	961-07-9	163230	HMDB000 CC	00330	135398592	Neg
23 histidine	Amino Acid	Histidine Metabolism	HNDVDQJCIGZPNO-UHFFFAOYAG	NC(C(O)=O)CC1=CN=CN1	5934-29-3	752	HMDB000 CC	00135	6274	Neg
24 inosine	Nucleotide	Purine Metabolism, (Hypo)Xanthine/Inosine containir	UGQMRVRMYYASKQ-DGPXGRDGBU	O[C@H]1[C@H](N2C(N=C	58-63-9	21241953	HMDB000 CC	00294	6021	Neg
25 malate	Energy	TCA Cycle	BJEPYKJPYRNKOW-UHFFFAOYAM	OC(C(O)=O)CC(O)=O	6915-15-	510	HMDB003 CC	00149,C	C 525	Neg
26 palmitate (16:0)	Lipid	Long Chain Saturated Fatty Acid	IPCSVZSSVZVIGE-UHFFFAOYAJ	oc(cccccccccccccc)=	57-10-3	960	HMDB000 CC	00249	985	Neg
27 stearate (18:0)	Lipid	Long Chain Saturated Fatty Acid	QIQXTHQIDYTERH-UHEEFAQYAB		57-11-4	5091	HMDB000 CC	01530	5281	Neg
28 palmitoleate (16:1n7)	Lipid	Long Chain Monounsaturated Fatty Acid	SECPZKHBENQXJG-FPLPWBNLBE	cccccc\c=c/cccccccc	373-49-9	393216	HMDB000 CC	08362	445638	Neg
29 phosphate	Energy	Oxidative Phosphorylation	NBIIXXVUZAFLBC-DFZHHIFOAA	O=P(O)(O)O	7664-38-3	1032	HMDB000 CC	00009	1061	Neg
30 phosphoenolpyruvate (PEP)	Carbohydrate	Glycolysis, Gluconeogenesis, and Pyruvate Metabolish	DTBNBXWJWCWCIK-UHFFFAOYAU	O=P(O)(OC(C(O)=O)=C)O	10526-80	980	HMDB000 CC	00074	1005	Neg
31 lactate	Carbohydrate	Glycolysis, Glyconeogenesis, and Pyruvate Metabolish	JVTAAEKCZENVCJ-UHEFFAOYAX	CC(0)C(0)=0	79-33-4	592	HMDB000 CC	00186	612	Neg
32 pyridoxal	Cofactors and Vitamins	Vitamin B6 Metabolism	RADKZDMEGJYCBB-UHEEEAOYAP	OC1=C(C)N=CC(CO)=C1C=	65-22-5	1021	HMDB000 CC	00250	1050	Neg
33 pyridoxal phosphate	Cofactors and Vitamins	Vitamin B6 Metabolism	NGVDGCNEYWLIEO-UHEEEAOYAW	O=P(O)(O)OCC1=CN=C(C)	(41468-25	1022	HMDB000 CC	00018	1051	Neg
34 riboflavin (Vitamin B2)	Cofactors and Vitamins	Biboflavin Metabolism	AUNGANR7IHBGPY-SCRDCRAPBG	OC(C(0)C(0)C0)CN1C(C=	(83-88-5	431981	HMDB000 CC	00255	493570	Neg
35 myristate (14:0)	Linid	Long Chain Saturated Fatty Acid	TUNESRHWOTWDNC-UHEFEAOYA7		544-63-8	10539	HMDB000 CC	06424	11005	Neg
36 thiamin dinhosphate	Cofectors and Vitamins	Thiamine Metabolism	YXVCI PIOTZXII H-LIHEEEAOYAW	CC1=NC=C(C(N)=N1)C[N+]	154-87-0	8715	HMDB000 CC	00068	1132	Neg
37 thiamin monophosphate	Cofactors and Vitamins	Thiamine Metabolism	HZSAIDV/WZBBGIE-LIHEEEAOVAO	CC1=NC=C(C[N+]2=CSC(C	532-40-1	2627905	HMDB000 CC	01081	1131 3382778	Neg
38 uridine	Nucleotide	Purimidine Metabolism Uracil containing		OC1C(N(C=CC2=O)C(N2)=	(58-06-8	5807	HMDB000 CC	01001	6020	Neg
39 2'-deoxyuridine	Nucleotide	Pyrimidine Metabolism, Uracil containing	MXHRCPNRIAMMIM-SHYZELIOEBK	O = C(NC1 = O)N(C = C1)C(CC)	951-78-0	13118	HMDB000 CC	00526	13712	Neg
40 pseudouridine	Nucleotide	Pyrimidine Metabolism, Uracil containing		0[c@]1([H])[c@@](0)([H	1445-07-	21403010	HMDB000 CC	02067	15047	Neg
41 thymidine	Nucleotide	Pyrimidine Metabolism, Drack containing		0=c(Nc(c(c)=c1)=0)N1[c	50-89-5	5585	HMDB000 CC	00214	5789	Neg
42 thymine	Nucleotide	Pyrimidine Metabolism, Thymine containing	RWONBRDOKXIBIV-LIHEEEAOYAL	0=C1C(C)=CNC(N1)=0	65-71-4	1103	HMDB000 CC	00178	1135	Neg
43 caprate (10:0)	Lipid	Medium Chain Fatty Acid	GHVNEZECNZKVNT-LIHEEEAOVAC		334-48-5	2863	HMDB000 CC	01571	2969	Neg
44 margarate (17:0)	Lipid	Long Chain Saturated Eatty Acid	KEMOGTRYUADDNZ-UHEEEAOVAT	0-010)000000000000000000000000000000000	506-12-7	10033	HMDB00022	50	10465	Neg
45 popodecopoote (19:0)	Lipid	Long Chain Saturated Fatty Acid			646-20-0	12071	HMDB00022	16525	12501	Neg
46 prochidate (20:0)	Lipid	Long Chain Saturated Fatty Acid			COG 20 0	10025	HMDB000 C	06425	10467	Neg
40 arachidate (20.0)	Carbobudrate	Glysogon Motobolism			6363 53	20035	HMDB000 CC	00423	10001480	Neg
47 marcose 48 contracto (6:0)	Linid	Modium Chain Eatty Asid			142 62 1	0550	HMDB000 CC	01595	10991409	Neg
40 caproate (0:0)	Lipia Lipia	Medium Chain Fatty Acid			142-02-1	0002		06433	0092	Neg
49 caprylate (8:0)	Lipia	International Contraction Contraction		0-0/0/000000000000000000000000000000000	124-07-2	370		16527	3/9	iveg
50 pentadecanoate (15:0)		cong chain Saturated Fatty Acid	WQEPLOUGTEDZJT-OHFFFAOTAK		10002-84	- 13249	HIVIDBOODCI	10537	13649	Neg
51 S-oxoproline	Amino Acid	Giutatnione Metabolism	CUCKINGTUZIEAOD ZETCOVENING	O=CICC[C@@H]([C@@](98-79-3	/12/	HIVIDBUU0 CC	UZ237,C	439685,7405	iveg
52 pantothenate	Coractors and Vitamins	Pantotnenate and CoA Metabolism	GHUKWGTUZJEAQD-ZETCQYMHBS	U=C(NCCC(U)=U)[C@H](C	.137-08-6	0301	HIVIDBUU0 CC	00864	6100	iveg
53 phosphoethanolamine		Phospholipid Metabolism	SUHOUTKUPISOBE-UHFFFAOYAT	NCCOP(O)(O)=O	10/1-23-4	1 990	HMDB000 CC	00346	1015	Neg
54 5-KETE	Lipid	Elcosanoid	4000	LLLLC\C=C/C\C=C/C\C=C	. 106154-1	81/65	HMDB001C1	14732	5353355	Neg
Data Key & Explanation	Chemical Annotation	nical Annotation (2) Sample Meta Data Peak Are	a Data Transposed Peak area dat	a Neg Polar Pos	early	Pos late Batc	h-normalized	Data	Batch-norm In	nputed Data
- · · · · · · · · ·										

These sub-tables are then ready to be copied/pasted into the online submission form at the "metabolite metadata" steps Order of analyses: 1:Pos Early, 2:Pos Late, 3:Neg 4:Polar

Start/Edit Data Submission Examples of study design and data layouts Online Study Submission Tutorial(pdf)
mwTab Identifier: Javier_Munoz_Briones_24_20230930_161810 Return to start Analysis: Reversed phase POSITIVE(1)
Metabolite metadata in tab-delimited format. First column must contain metabolite names. Subsequent columns should contain KEGG,PubChem identifiers, retention index,quantitated m/z, etc. First row must contain headings.
<pre>metabolite_name SUPER_PATHWAY SUB_PATHWAY TYPE INCHIKEY SMILES CAS CHEMSPIDER HMDB KEGG PUBCHEM S-1-pyrroline-5-carboxylate Amino Acid Glutamate Metabolism NAMED DWAKNKKXGALPNW- UHFFFAOYSA-N OC(C1CCC=N1)=0 2906-39-0 10140206 HMDB0001301 C04322 11966181 1,3-diaminopropane Amino Acid Polyamine Metabolism NAMED XFNJVJPLKCPIBV-UHFFFAOYSA-N NCCCN 109-76-2 415 HMDB0000002 C00986 428 putrescine Amino Acid Polyamine Metabolism NAMED KIDHWZJUCRJVML-UHFFFAOYSA-N NCCCCN 110-60-1 13837702 HMDB0001414 C00134 1045 spermidine Amino Acid Polyamine Metabolism NAMED ATHGHOPFGPMSJY-UHFFFAOYSA-N NCCCCN 124-20-9 1071 HMDB0001257 C00315 1102 1-methyladenine Nucleotide Purine Metabolism, Adenine containing NAMED SATCOUWSAZBIJO-</pre>
View/check metabolite metadata See examples of metabolite metadata layout

Data upload is complete when you have copied/pasted results and metadata tables for each of the 4 Metabolon analyses Order of analyses: 1:Pos Early, 2:Pos Late, 3:Neg 4:Polar

Click the "Finalize" button in the form when complete