Package	Source	Version	Note
	T	R ver	sion
Base packages	CRAN	4.1.2	-
1 4 4 11	CDAN	Common	
data.table	CRAN	1.14.2	-
dplyr	CRAN	1.0.9	-
stringr	CRAN	1.4.0	-
doParallel	CRAN	1.0.17	-
BiocParallel	BC	1.28.3	-
ggplot2	CRAN	3.3.6 Random	- imption
tunlo	CRAN	0.4-02	lization
tuple randomizr	CRAN	0.4-02	-
randonnizi	CKAN	Integr	etion
ino	BC	1.20.0	
ipo Autotuner	BC	1.7.0	-
MetaboAnalystR	GH	3.0.2	_
xcms	BC	3.0.2	_
MSnbase	BC	2.20.1	_
MsFeatures	GH	1.2.0	_
	GH	0.1.0.9001	-
warpgroup ncGTW	BC	1.8.0	_
	GH	0.1.0	_
срс	UII	Corre	
nmn	BC	1.6.0	-
pmp StatTarget	BC	1.24.0	_
MetNormalizer	GH	1.3.02	_
batchCorr	GL	0.2.5	_
BatchCorrMetabolomics	GH	0.2.3	_
TIGERr	CRAN	1.0.0	-
notame	GH	0.0.5	_
openxlsx	CRAN	4.2.5	-
pbapply	CRAN	1.5-0	_
caret	CRAN	6.0-92	-
rpart	CRAN	4.1.16	-
ipred	CRAN	0.9-13	-
catboost	GH	0.20	-
xgboost	CRAN	1.6.0.1	-
dbnorm	GH	0.2.2	-
ProteoMM	BC	1.12.0	-
WaveICA	GH	0.1.0	-
WaveICA2.0	GH	0.1.0	-
NormalizeMets	CRAN	0.25	-
RUVSeq	BC	1.28.0	-
tibble	CRAN	3.1.7	-
fpc	CRAN	2.2-9	-
FactoMineR	CRAN	2.4	-
cluster	CRAN	2.1.3	-
proBatch	BC	1.10.0	-
gPCA	CRAN	1.0	-

pcpr2	GH	0.0.0.1	_		
MetabolomicsBasics	CRAN	1.2			
factoextra	CRAN	1.0.7	-		
rafalib	CRAN	1.0.7	-		
		4.1-1	-		
RSEIS	CRAN		-		
ggsci	CRAN	2.9	-		
corrplot	CRAN	0.92	tation		
CAMERA	ВС	1.50.0	tation		
RAMClustR	CRAN	1.30.0	-		
	+	1.2.2	- Image Imag		
xMSannotator	GH	1.3.2	From kuppal2 repo (https://github.com/kuppal2/xMSannotator). Fix for R 4.0 fun string 95 to n2<-as.data.frame(n1, stringsAsFactors = T). According to yufree repo (https://github.com/yufree/xMSannotator). Install manually (tar.gz), type = "source". Also see instruction from GH and SF.		
mWISE	GH	0.1.0	-		
metid	GH	1.2.13	Databases from		
			(https://github.com/jaspershen/demoData/tree/master/inst/ms2_databada) should be copied in "ms2_database" folder in metID library folder.		
tidyverse	CRAN	1.3.1	-		
batchCorr	GL	0.2.5	-		
curl	CRAN	4.3.2	-		
piggyback	CRAN	0.1.3	-		
MetaboAnnotation	BC	1.1.1	Database from <i>OUKS</i> release 1.10.1		
CompoundDb	BC	0.99.8	-		
	•	Norma	lization		
affy	BC	1.72.0	-		
ABCstats	GH	1.0.0	-		
clusterSim	CRAN	0.50-1	-		
DiffCorr	CRAN	0.4.2	-		
lmm2met	GH	1.0	-		
MetabolomicsBasics	CRAN	1.2	-		
lme4	CRAN	1.1-29	-		
lmerTest	CRAN	3.1-3	-		
mgcv	CRAN	1.8-40	-		
gamm4	CRAN	0.2-6	-		
NormalizeMets	CRAN	0.25	-		
RColorBrewer	CRAN	1.1-3	-		
limma	BC	3.50.3	-		
Imputation					
MetProc	CRAN	1.0.1	-		
MAI	BC	1.0.0	-		
impute	BC	1.68.0	-		
mice	CRAN	3.14.0	-		
missForest	CRAN	1.5	-		
StatTools	GL	0.0.915	-		
pcaMethods	BC	1.86.0	-		

imputeLCMD	CRAN	2.1	-			
tWLSA	GH	1.0	-			
imputeR	CRAN	2.2	-			
vegan	CRAN	2.6-2	-			
missMethods	CRAN	0.3.0	-			
	CDAN	Grou	ping			
corrr	CRAN	0.4.3	-			
gplots tidyverse	CRAN CRAN	3.1.3	-			
igraph	CRAN	1.3.1	_			
ggraph	CRAN	2.0.5	-			
robCompositions	CRAN	2.3.1	-			
compositions	CRAN	2.0-4	-			
gridExtra	CRAN	2.3	-			
stringi	CRAN	1.7.6	-			
notame	GH	0.0.5	-			
pmd	CRAN	0.2.1	-			
	1	Stati	stics			
pcaMethods	BC	1.86.0	-			
OutlierDetection	GH	0.1.1	Should be installed from			
G1 51	GD 137	2.1.0	"rubak/spatstat.revdep"			
ClassDiscovery	CRAN	3.4.0	-			
ggforce	CRAN	0.3.3	-			
HotellingEllipse	CRAN	1.1.0	-			
caret	CRAN	6.0-92	Other methods may require additional packages			
klaR	CRAN	0.6-14	The klaR package should be exactly version 0.6-14 for RFS			
cwhmisc	CRAN	6.6	The shapiro.wilk.test function should be used if some error with normality test in UVF.			
tuple	CRAN	0.4-02	-			
pamr	CRAN	1.56.1	For ML model in caret			
pls	CRAN	2.8-0	For ML model in caret			
kernlab	CRAN	0.9-31	For ML model in caret			
randomForest	CRAN	4.7-1.1	For ML model in caret			
rsample	CRAN	0.1.1	-			
ropls	BC	1.26.4	-			
tdfdr	CRAN	0.1	-			
MetabolomicsBasics	CRAN	1.2	-			
lme4	CRAN	1.1-29	-			
lmerTest	CRAN	3.1-3	-			
mgcv	CRAN	1.8-40	-			
gamm4	CRAN	0.2-6	-			
drc	CRAN	3.0-1	-			
limma	BC	3.50.3	-			
NormalizeMets	CRAN	0.25	-			

Boruta	CRAN	7.0.0	-
permimp	CRAN	1.0-2	-
party	CRAN	1.3-10	-
glmnet	CRAN	4.1-4	-
MASS	CRAN	7.3-57	-
pROC	CRAN	1.18.0	-
reshape2	CRAN	1.4.4	-
leaps	CRAN	3.1	-
gWQS	CRAN	3.0.4	-
vegan	CRAN	2.6-2	-
pairwiseAdonis	GH	0.0.1	-
multcomp	CRAN	1.4-19	-
MWASTools	BC	1.18.0	-
glmmsr	CRAN	0.2.3	-
MetStaT	CRAN	1.0	-
mixOmics	BC	6.18.1	-
DRomics	CRAN	2.4-0	-
TOXcms	GH	1.0.3	Require dr4pl and hashmap packages
timecourse	BC	1.66.0	-
timeOmics	BC	1.6.0	-
lmms	CRAN	1.3.3	-
polyPK	CRAN	3.1.0	-
openxlsx	CRAN	4.2.5	-
tidyverse	CRAN	1.3.1	-
factoextra	CRAN	1.0.7	-
FactoMineR	CRAN	2.4	-
dendextend	CRAN	1.15.2	-
rafalib	CRAN	1.0.0	-
cluster	CRAN	2.1.3	-
dbscan	CRAN	1.1-10	-
umap	CRAN	0.2.8.0	-
NMF	CRAN	0.24.0	-
Rdimtools	CRAN	1.0.9	-
dimRed	CRAN	0.2.5	-
RSEIS	CRAN	4.1-1	-
ggsci	CRAN	2.9	-
pheatmap	CRAN	1.0.12	-
Rtsne	CRAN	0.16	-
NbClust	CRAN	3.0.1	-
clustertend	CRAN	1.6	-
mclust	CRAN	5.4.10	-
clValid	CRAN	0.7	-

pvclust	CRAN	2.2-0	-	
fpc	CRAN	2.2-9	-	
Hmisc	CRAN	4.7-0	-	
corrplot	CRAN	0.92	-	
psych	CRAN	2.2.5	-	
pwr	CRAN	1.3-0	-	
effectsize	CRAN	0.7.0	-	
Directly used packages: 14				