

Assignment 3

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```
library(ggplot2)
library(ggfortify)
library(factoextra)
```

```
## Welcome! Want to learn more? See two factoextra-related books at https://goo.gl/ve3WBa
```

```
library(Rtsne)
library(dplyr)
```

```
##
## Attaching package: 'dplyr'

## The following objects are masked from 'package:stats':
##
##   filter, lag

## The following objects are masked from 'package:base':
##
##   intersect, setdiff, setequal, union
```

Question 1

```
#Read data
setwd("C:/Users/aless/Desktop/Bioinformatics/Erasmus BCN/Data Visualisation/homeassignment")
P3<- read.delim("P3.tsv.gz", row.names=1)
P4<- read.delim("P4.tsv.gz", row.names=1)
P5<- read.delim("P5.tsv.gz", row.names=1)

#Check data
head(P3, 5)
```

```
##               P2769_N701.S502 P2769_N701.S503 P2769_N701.S505
## ENSG000000000003.14           0                0                0
## ENSG000000000005.5           0                0                0
## ENSG000000000419.12          39               68               16
## ENSG000000000457.13           0                8               17
## ENSG000000000460.16           0               46                4
##               P2769_N701.S506 P2769_N701.S507 P2769_N701.S508
## ENSG000000000003.14           0                0                0
```

##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	0	0	0
##	ENSG000000000457.13	0	0	0
##	ENSG000000000460.16	0	0	0
##	P2769_N701.S510	P2769_N701.S511	P2769_N702.S502	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	23	2	24
##	ENSG000000000457.13	0	0	0
##	ENSG000000000460.16	0	11	7
##	P2769_N702.S503	P2769_N702.S505	P2769_N702.S506	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	62	76	0
##	ENSG000000000457.13	0	0	0
##	ENSG000000000460.16	7	0	0
##	P2769_N702.S507	P2769_N702.S508	P2769_N702.S510	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	0	0	39
##	ENSG000000000457.13	0	0	0
##	ENSG000000000460.16	0	0	170
##	P2769_N702.S511	P2769_N703.S502	P2769_N703.S503	
##	ENSG000000000003.14	0	0	0.00
##	ENSG000000000005.5	0	0	0.00
##	ENSG000000000419.12	88	29	92.00
##	ENSG000000000457.13	0	0	28.79
##	ENSG000000000460.16	0	0	187.21
##	P2769_N703.S505	P2769_N703.S506	P2769_N703.S507	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	39	69	0
##	ENSG000000000457.13	15	0	0
##	ENSG000000000460.16	0	0	0
##	P2769_N703.S508	P2769_N703.S510	P2769_N703.S511	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	0	59	79
##	ENSG000000000457.13	0	0	0
##	ENSG000000000460.16	0	0	3
##	P2769_N704.S502	P2769_N704.S503	P2769_N704.S505	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	19	111	9
##	ENSG000000000457.13	0	1	0
##	ENSG000000000460.16	45	30	0
##	P2769_N704.S506	P2769_N704.S507	P2769_N704.S508	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	162	0	0
##	ENSG000000000457.13	0	0	0
##	ENSG000000000460.16	0	0	0
##	P2769_N704.S510	P2769_N704.S511	P2769_N705.S502	
##	ENSG000000000003.14	0	0	0

##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	25	0	110
##	ENSG000000000457.13	0	0	33
##	ENSG000000000460.16	0	110	79
##	P2769_N705.S503	P2769_N705.S505	P2769_N705.S506	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	74	9	16
##	ENSG000000000457.13	0	0	0
##	ENSG000000000460.16	185	0	240
##	P2769_N705.S507	P2769_N705.S508	P2769_N705.S510	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	79	45	0
##	ENSG000000000457.13	0	6	0
##	ENSG000000000460.16	0	0	1
##	P2769_N705.S511	P2769_N706.S502	P2769_N706.S503	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	17	53	182
##	ENSG000000000457.13	0	0	0
##	ENSG000000000460.16	0	0	0
##	P2769_N706.S505	P2769_N706.S506	P2769_N706.S507	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	49	125	0
##	ENSG000000000457.13	0	0	0
##	ENSG000000000460.16	0	0	0
##	P2769_N706.S508	P2769_N706.S510	P2769_N706.S511	
##	ENSG000000000003.14	0	0.00	0
##	ENSG000000000005.5	0	0.00	0
##	ENSG000000000419.12	175	110.00	29
##	ENSG000000000457.13	0	185.82	0
##	ENSG000000000460.16	0	63.18	0
##	P2769_N707.S502	P2769_N707.S503	P2769_N707.S505	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	157	73	47
##	ENSG000000000457.13	0	0	0
##	ENSG000000000460.16	47	186	0
##	P2769_N707.S506	P2769_N707.S507	P2769_N707.S508	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	207	114	84
##	ENSG000000000457.13	0	0	0
##	ENSG000000000460.16	0	170	152
##	P2769_N707.S510	P2769_N707.S511	P2769_N710.S502	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	27	58	152
##	ENSG000000000457.13	0	91	0
##	ENSG000000000460.16	94	26	0
##	P2769_N710.S503	P2769_N710.S505	P2769_N710.S506	
##	ENSG000000000003.14	0	0	0

##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	107	0	50
##	ENSG000000000457.13	21	0	0
##	ENSG000000000460.16	0	0	0
##	P2769_N710.S507	P2769_N710.S508	P2769_N710.S510	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	82	176	68
##	ENSG000000000457.13	134	0	0
##	ENSG000000000460.16	55	0	87
##	P2769_N710.S511	P2769_N711.S502	P2769_N711.S503	
##	ENSG000000000003.14	0	0.00	0
##	ENSG000000000005.5	0	0.00	0
##	ENSG000000000419.12	0	139.00	69
##	ENSG000000000457.13	0	28.19	0
##	ENSG000000000460.16	8	99.81	0
##	P2769_N711.S505	P2769_N711.S506	P2769_N711.S507	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	73	81	0
##	ENSG000000000457.13	79	0	8
##	ENSG000000000460.16	17	0	147
##	P2769_N711.S508	P2769_N711.S510	P2769_N711.S511	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	285	84	60
##	ENSG000000000457.13	0	0	0
##	ENSG000000000460.16	62	91	32
##	P2769_N712.S502	P2769_N712.S503	P2769_N712.S505	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	240	32	33
##	ENSG000000000457.13	0	0	0
##	ENSG000000000460.16	118	100	0
##	P2769_N712.S506	P2769_N712.S507	P2769_N712.S508	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	53	178	0
##	ENSG000000000457.13	0	0	0
##	ENSG000000000460.16	0	0	0
##	P2769_N712.S510	P2769_N712.S511	P2769_N714.S502	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	48	0	103
##	ENSG000000000457.13	0	0	0
##	ENSG000000000460.16	0	0	123
##	P2769_N714.S503	P2769_N714.S505	P2769_N714.S506	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	58	35	0
##	ENSG000000000457.13	0	0	0
##	ENSG000000000460.16	0	259	0
##	P2769_N714.S507	P2769_N714.S508	P2769_N714.S510	
##	ENSG000000000003.14	0	0	0

##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	86	114	0
##	ENSG000000000457.13	0	0	10
##	ENSG000000000460.16	0	0	0
##	P2769_N714.S511	P2769_N715.S502	P2769_N715.S503	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	44	351	83
##	ENSG000000000457.13	16	0	0
##	ENSG000000000460.16	0	0	325
##	P2769_N715.S505	P2769_N715.S506	P2769_N715.S507	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	292	110	110
##	ENSG000000000457.13	0	0	136
##	ENSG000000000460.16	0	0	2
##	P2769_N715.S508	P2769_N715.S510	P2769_N715.S511	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	0	56	0
##	ENSG000000000457.13	0	0	0
##	ENSG000000000460.16	0	0	14

```
head(P4, 5)
```

##	P2770_N701.S513	P2770_N701.S515	P2770_N701.S516	
##	ENSG000000000003.14	0	0.00	0
##	ENSG000000000005.5	0	0.00	0
##	ENSG000000000419.12	49	79.00	144
##	ENSG000000000457.13	0	68.42	0
##	ENSG000000000460.16	0	127.58	0
##	P2770_N701.S517	P2770_N701.S518	P2770_N701.S520	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	21	0	87
##	ENSG000000000457.13	2	0	0
##	ENSG000000000460.16	13	0	0
##	P2770_N701.S521	P2770_N701.S522	P2770_N702.S513	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	0	0	75
##	ENSG000000000457.13	0	0	0
##	ENSG000000000460.16	202	0	82
##	P2770_N702.S515	P2770_N702.S516	P2770_N702.S517	
##	ENSG000000000003.14	0.00	0.00	0
##	ENSG000000000005.5	0.00	0.00	0
##	ENSG000000000419.12	151.00	28.00	83
##	ENSG000000000457.13	16.55	30.68	0
##	ENSG000000000460.16	114.45	12.32	0
##	P2770_N702.S518	P2770_N702.S520	P2770_N702.S521	
##	ENSG000000000003.14	0	0	0.00
##	ENSG000000000005.5	0	0	0.00
##	ENSG000000000419.12	33	185	95.00
##	ENSG000000000457.13	0	0	11.64

## ENSG00000000460.16	0	105	208.36
##	P2770_N702.S522	P2770_N703.S513	P2770_N703.S515
## ENSG00000000003.14	0	0	0
## ENSG00000000005.5	0	0	0
## ENSG00000000419.12	1	88	39
## ENSG00000000457.13	0	0	9
## ENSG00000000460.16	0	10	51
##	P2770_N703.S516	P2770_N703.S517	P2770_N703.S518
## ENSG00000000003.14	0	0	0
## ENSG00000000005.5	0	0	0
## ENSG00000000419.12	84	60	137
## ENSG00000000457.13	0	0	0
## ENSG00000000460.16	0	46	70
##	P2770_N703.S520	P2770_N703.S521	P2770_N703.S522
## ENSG00000000003.14	0	0	0
## ENSG00000000005.5	0	0	0
## ENSG00000000419.12	0	103	34
## ENSG00000000457.13	0	0	0
## ENSG00000000460.16	0	72	0
##	P2770_N704.S513	P2770_N704.S515	P2770_N704.S516
## ENSG00000000003.14	0	0	0
## ENSG00000000005.5	0	0	0
## ENSG00000000419.12	53	116	88
## ENSG00000000457.13	0	0	0
## ENSG00000000460.16	17	42	0
##	P2770_N704.S517	P2770_N704.S518	P2770_N704.S520
## ENSG00000000003.14	0	0	0.00
## ENSG00000000005.5	0	0	0.00
## ENSG00000000419.12	59	0	138.00
## ENSG00000000457.13	33	0	35.02
## ENSG00000000460.16	0	0	78.98
##	P2770_N704.S521	P2770_N704.S522	P2770_N705.S513
## ENSG00000000003.14	0	0	0
## ENSG00000000005.5	0	0	0
## ENSG00000000419.12	0	0	59
## ENSG00000000457.13	0	0	112
## ENSG00000000460.16	14	0	0
##	P2770_N705.S515	P2770_N705.S516	P2770_N705.S517
## ENSG00000000003.14	0	0	0
## ENSG00000000005.5	0	0	0
## ENSG00000000419.12	95	89	29
## ENSG00000000457.13	0	0	0
## ENSG00000000460.16	56	41	87
##	P2770_N705.S518	P2770_N705.S520	P2770_N705.S521
## ENSG00000000003.14	0	0	0
## ENSG00000000005.5	0	0	0
## ENSG00000000419.12	22	134	83
## ENSG00000000457.13	0	0	0
## ENSG00000000460.16	23	0	0
##	P2770_N705.S522	P2770_N706.S513	P2770_N706.S515
## ENSG00000000003.14	0	0	0.00
## ENSG00000000005.5	0	0	0.00
## ENSG00000000419.12	28	44	64.00
## ENSG00000000457.13	0	0	22.64

## ENSG00000000460.16	0	0	68.36
##	P2770_N706.S516	P2770_N706.S517	P2770_N706.S518
## ENSG00000000003.14	0	0.00	0
## ENSG00000000005.5	0	0.00	0
## ENSG00000000419.12	71	64.00	0
## ENSG00000000457.13	0	117.38	0
## ENSG00000000460.16	0	227.62	0
##	P2770_N706.S520	P2770_N706.S521	P2770_N706.S522
## ENSG00000000003.14	0	0	0
## ENSG00000000005.5	0	0	0
## ENSG00000000419.12	96	59	46
## ENSG00000000457.13	0	29	5
## ENSG00000000460.16	0	0	0
##	P2770_N707.S513	P2770_N707.S515	P2770_N707.S516
## ENSG00000000003.14	0	0	0.00
## ENSG00000000005.5	0	0	0.00
## ENSG00000000419.12	92	99	33.00
## ENSG00000000457.13	1	99	1.02
## ENSG00000000460.16	141	0	202.98
##	P2770_N707.S517	P2770_N707.S518	P2770_N707.S520
## ENSG00000000003.14	0	0	0
## ENSG00000000005.5	0	0	0
## ENSG00000000419.12	103	37	87
## ENSG00000000457.13	0	125	0
## ENSG00000000460.16	0	0	20
##	P2770_N707.S521	P2770_N707.S522	P2770_N710.S513
## ENSG00000000003.14	0	0	0.0
## ENSG00000000005.5	0	0	0.0
## ENSG00000000419.12	104	97	81.0
## ENSG00000000457.13	0	0	120.8
## ENSG00000000460.16	0	87	64.2
##	P2770_N710.S515	P2770_N710.S516	P2770_N710.S517
## ENSG00000000003.14	0	0	0
## ENSG00000000005.5	0	0	0
## ENSG00000000419.12	52	18	13
## ENSG00000000457.13	0	2	0
## ENSG00000000460.16	0	0	6
##	P2770_N710.S518	P2770_N710.S520	P2770_N710.S521
## ENSG00000000003.14	0	0	0
## ENSG00000000005.5	0	0	0
## ENSG00000000419.12	23	58	79
## ENSG00000000457.13	0	0	1
## ENSG00000000460.16	51	0	0
##	P2770_N710.S522	P2770_N711.S513	P2770_N711.S515
## ENSG00000000003.14	0	0.00	0
## ENSG00000000005.5	0	0.00	0
## ENSG00000000419.12	111	75.00	5
## ENSG00000000457.13	0	44.33	0
## ENSG00000000460.16	0	183.67	0
##	P2770_N711.S516	P2770_N711.S517	P2770_N711.S518
## ENSG00000000003.14	0	0	0
## ENSG00000000005.5	0	0	0
## ENSG00000000419.12	41	44	0
## ENSG00000000457.13	0	0	0

##	ENSG000000000460.16	0	0	28
##		P2770_N711.S520	P2770_N711.S521	P2770_N711.S522
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	10	0	19
##	ENSG000000000457.13	0	0	0
##	ENSG000000000460.16	1	3	0
##		P2770_N712.S513	P2770_N712.S515	P2770_N712.S516
##	ENSG000000000003.14	0.00	0.00	0
##	ENSG000000000005.5	0.00	0.00	0
##	ENSG000000000419.12	78.00	65.00	17
##	ENSG000000000457.13	14.69	30.62	0
##	ENSG000000000460.16	77.31	2.38	0
##		P2770_N712.S517	P2770_N712.S518	P2770_N712.S520
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	0	37	41
##	ENSG000000000457.13	0	0	0
##	ENSG000000000460.16	0	92	0
##		P2770_N712.S521	P2770_N712.S522	P2770_N714.S513
##	ENSG000000000003.14	0.00	0	0
##	ENSG000000000005.5	0.00	0	0
##	ENSG000000000419.12	24.00	50	60
##	ENSG000000000457.13	15.88	0	18
##	ENSG000000000460.16	8.12	0	110
##		P2770_N714.S515	P2770_N714.S516	P2770_N714.S517
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	20	25	0
##	ENSG000000000457.13	22	4	0
##	ENSG000000000460.16	0	0	84
##		P2770_N714.S518	P2770_N714.S520	P2770_N714.S521
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	161	13	78
##	ENSG000000000457.13	0	0	0
##	ENSG000000000460.16	95	70	0
##		P2770_N714.S522	P2770_N715.S513	P2770_N715.S515
##	ENSG000000000003.14	0.00	0	0
##	ENSG000000000005.5	0.00	0	0
##	ENSG000000000419.12	21.00	92	50
##	ENSG000000000457.13	22.92	40	20
##	ENSG000000000460.16	127.08	0	0
##		P2770_N715.S516	P2770_N715.S517	P2770_N715.S518
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	61	43	3
##	ENSG000000000457.13	0	0	0
##	ENSG000000000460.16	0	118	92
##		P2770_N715.S520	P2770_N715.S521	P2770_N715.S522
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	32	75	46
##	ENSG000000000457.13	0	0	0


```
## ENSG00000000460.16          5          0          0
```

```
head(P5, 5)
```

```
##          P2771_N701.S502 P2771_N701.S503 P2771_N701.S505
## ENSG00000000003.14          0          0          0
## ENSG00000000005.5          0          0          0
## ENSG000000000419.12        41         127         113
## ENSG000000000457.13        60          2          16
## ENSG000000000460.16         0         59          48
##          P2771_N701.S506 P2771_N701.S507 P2771_N701.S508
## ENSG00000000003.14          0          0          0
## ENSG00000000005.5          0          0          0
## ENSG000000000419.12        60          0         197
## ENSG000000000457.13         0          0          0
## ENSG000000000460.16         0          0          0
##          P2771_N701.S510 P2771_N701.S511 P2771_N702.S502
## ENSG00000000003.14          0          0          0
## ENSG00000000005.5          0          0          0
## ENSG000000000419.12        14         35          84
## ENSG000000000457.13        55          0          9
## ENSG000000000460.16         0          0         65
##          P2771_N702.S503 P2771_N702.S505 P2771_N702.S506
## ENSG00000000003.14         0.00         0.00          0
## ENSG00000000005.5         0.00         0.00          0
## ENSG000000000419.12       134.00        12.00         13
## ENSG000000000457.13        34.04        46.47          0
## ENSG000000000460.16        66.96        46.53          0
##          P2771_N702.S507 P2771_N702.S508 P2771_N702.S510
## ENSG00000000003.14          0          0          0
## ENSG00000000005.5          0          0          0
## ENSG000000000419.12          0        151          57
## ENSG000000000457.13          0          0          0
## ENSG000000000460.16          0          0         102
##          P2771_N702.S511 P2771_N703.S502 P2771_N703.S503
## ENSG00000000003.14          0          0         0.00
## ENSG00000000005.5          0          0         0.00
## ENSG000000000419.12          0         71        35.00
## ENSG000000000457.13          0          0        31.61
## ENSG000000000460.16          0          0        51.39
##          P2771_N703.S505 P2771_N703.S506 P2771_N703.S507
## ENSG00000000003.14          0          0          0
## ENSG00000000005.5          0          0          0
## ENSG000000000419.12        24         11        129
## ENSG000000000457.13          0          0          0
## ENSG000000000460.16         6         24          0
##          P2771_N703.S508 P2771_N703.S510 P2771_N703.S511
## ENSG00000000003.14          0          0          0
## ENSG00000000005.5          0          0          0
## ENSG000000000419.12        61          0         24
## ENSG000000000457.13          0          0          0
## ENSG000000000460.16        55          0          0
##          P2771_N704.S502 P2771_N704.S503 P2771_N704.S505
## ENSG00000000003.14          0         0.00          0
```

##	ENSG000000000005.5	0	0.00	0
##	ENSG000000000419.12	74	151.00	51
##	ENSG000000000457.13	0	26.41	0
##	ENSG000000000460.16	0	33.59	0
##	P2771_N704.S506	P2771_N704.S507	P2771_N704.S508	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	0	50	60
##	ENSG000000000457.13	0	0	0
##	ENSG000000000460.16	0	46	0
##	P2771_N704.S510	P2771_N704.S511	P2771_N705.S502	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	21	13	41
##	ENSG000000000457.13	0	0	0
##	ENSG000000000460.16	0	40	89
##	P2771_N705.S503	P2771_N705.S505	P2771_N705.S506	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	127	31	22
##	ENSG000000000457.13	0	0	0
##	ENSG000000000460.16	51	0	169
##	P2771_N705.S507	P2771_N705.S508	P2771_N705.S510	
##	ENSG000000000003.14	0	0.00	0.00
##	ENSG000000000005.5	0	0.00	0.00
##	ENSG000000000419.12	96	3.00	53.00
##	ENSG000000000457.13	111	133.87	112.67
##	ENSG000000000460.16	0	4.13	195.33
##	P2771_N705.S511	P2771_N706.S502	P2771_N706.S503	
##	ENSG000000000003.14	0	0	0.00
##	ENSG000000000005.5	0	0	0.00
##	ENSG000000000419.12	51	25	37.00
##	ENSG000000000457.13	0	0	21.93
##	ENSG000000000460.16	90	0	20.07
##	P2771_N706.S505	P2771_N706.S506	P2771_N706.S507	
##	ENSG000000000003.14	0.00	0	0
##	ENSG000000000005.5	0.00	0	0
##	ENSG000000000419.12	67.00	34	7
##	ENSG000000000457.13	26.23	0	0
##	ENSG000000000460.16	97.77	0	0
##	P2771_N706.S508	P2771_N706.S510	P2771_N706.S511	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	0	13	19
##	ENSG000000000457.13	0	0	0
##	ENSG000000000460.16	0	11	0
##	P2771_N707.S502	P2771_N707.S503	P2771_N707.S505	
##	ENSG000000000003.14	0.00	0	0
##	ENSG000000000005.5	0.00	0	0
##	ENSG000000000419.12	114.00	69	82
##	ENSG000000000457.13	57.74	42	0
##	ENSG000000000460.16	46.26	0	31
##	P2771_N707.S506	P2771_N707.S507	P2771_N707.S508	
##	ENSG000000000003.14	0	0	0

##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	28	177	52
##	ENSG000000000457.13	0	0	0
##	ENSG000000000460.16	0	4	93
##	P2771_N707.S510	P2771_N707.S511	P2771_N710.S502	
##	ENSG000000000003.14	0	0	0.00
##	ENSG000000000005.5	0	0	0.00
##	ENSG000000000419.12	90	98	122.00
##	ENSG000000000457.13	0	0	15.76
##	ENSG000000000460.16	22	25	44.24
##	P2771_N710.S503	P2771_N710.S505	P2771_N710.S506	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	10	0	0
##	ENSG000000000457.13	0	18	0
##	ENSG000000000460.16	0	64	0
##	P2771_N710.S507	P2771_N710.S508	P2771_N710.S510	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	76	88	113
##	ENSG000000000457.13	0	0	0
##	ENSG000000000460.16	1	0	9
##	P2771_N710.S511	P2771_N711.S502	P2771_N711.S503	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	74	98	61
##	ENSG000000000457.13	0	12	55
##	ENSG000000000460.16	33	10	0
##	P2771_N711.S505	P2771_N711.S506	P2771_N711.S507	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	73	101	165
##	ENSG000000000457.13	0	0	0
##	ENSG000000000460.16	106	56	40
##	P2771_N711.S508	P2771_N711.S510	P2771_N711.S511	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	0	48	27
##	ENSG000000000457.13	0	59	10
##	ENSG000000000460.16	0	0	65
##	P2771_N712.S502	P2771_N712.S503	P2771_N712.S505	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	44	42	0
##	ENSG000000000457.13	24	0	0
##	ENSG000000000460.16	68	0	0
##	P2771_N712.S506	P2771_N712.S507	P2771_N712.S508	
##	ENSG000000000003.14	0	0	0
##	ENSG000000000005.5	0	0	0
##	ENSG000000000419.12	98	136	85
##	ENSG000000000457.13	0	0	0
##	ENSG000000000460.16	0	150	54
##	P2771_N712.S510	P2771_N712.S511	P2771_N714.S502	
##	ENSG000000000003.14	0	0	0

```

## ENSG000000000005.5      0      0      0
## ENSG0000000000419.12    23     12    108
## ENSG0000000000457.13     0      0      0
## ENSG0000000000460.16    135     9     28
##      P2771_N714.S503 P2771_N714.S505 P2771_N714.S506
## ENSG0000000000003.14     0.00      0      0
## ENSG0000000000005.5     0.00      0      0
## ENSG0000000000419.12    60.00     93     75
## ENSG0000000000457.13     8.12      0      0
## ENSG0000000000460.16    94.88      0      0
##      P2771_N714.S507 P2771_N714.S508 P2771_N714.S510
## ENSG0000000000003.14     0      0      0
## ENSG0000000000005.5     0      0      0
## ENSG0000000000419.12    85     39      0
## ENSG0000000000457.13     0      0     43
## ENSG0000000000460.16    37      0      0
##      P2771_N714.S511 P2771_N715.S502 P2771_N715.S503
## ENSG0000000000003.14     0      0      0
## ENSG0000000000005.5     0      0      0
## ENSG0000000000419.12    77      0     66
## ENSG0000000000457.13     0      3      0
## ENSG0000000000460.16     5     41      3
##      P2771_N715.S505 P2771_N715.S506 P2771_N715.S507
## ENSG0000000000003.14     0      0      0
## ENSG0000000000005.5     0      0      0
## ENSG0000000000419.12    32    101    235
## ENSG0000000000457.13     0      0      0
## ENSG0000000000460.16    57     15     51
##      P2771_N715.S508 P2771_N715.S510 P2771_N715.S511
## ENSG0000000000003.14     0      0      0
## ENSG0000000000005.5     0      0      0
## ENSG0000000000419.12    59     59     27
## ENSG0000000000457.13     0      0      0
## ENSG0000000000460.16     0      0      0

```

#Alternatives

```

#P3<- read.table(gzfile("P3.tsv.gz", "rt"), header= TRUE)
#P4<- read.table(gzfile("P4.tsv.gz", "rt"), header= TRUE)
#P5<- read.table(gzfile("P5.tsv.gz", "rt"), header= TRUE)

```

Question 2

```

treat<- read.delim("samplesheet.tsv", row.names=1)
#treat <- as.factor(readLines("samplesheet.tsv"))
#Alternatives
#treat<- read.table(("samplesheet.tsv", "rt"), header= TRUE))

table(treat)

```

```

## treat
## J-LatA2+DMSO  J-LatA2+PMA  J-LatA2+SAHA  Jurkat+DMSO  Jurkat+PMA  Jurkat+SAHA
##           36           81           81           18           36           36

```

There are 2 kinds of cells (one infected with HIV and one with a latent virus), each one is treated with DMSO, SAHA and PMA. This creates 6 different combinations, the table above shows the number of cells in each condition.

Question 3

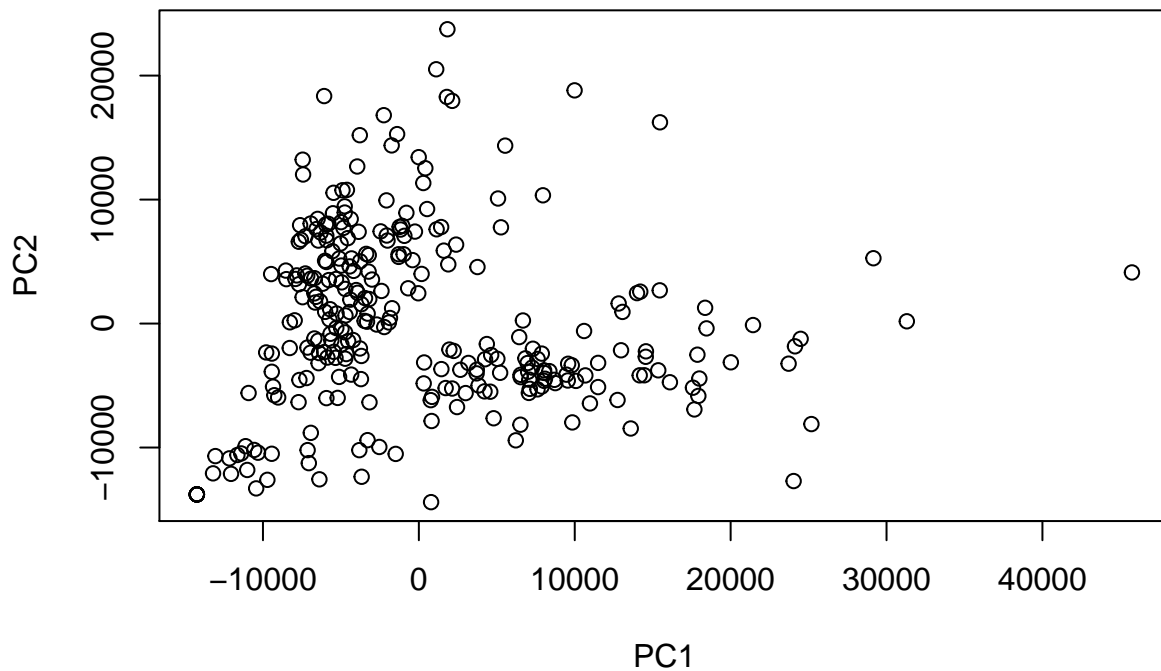
```
X3 <- t(P3)
X4 <- t(P4)
X5 <- t(P5)

X1 = rbind(X3, X4, X5)

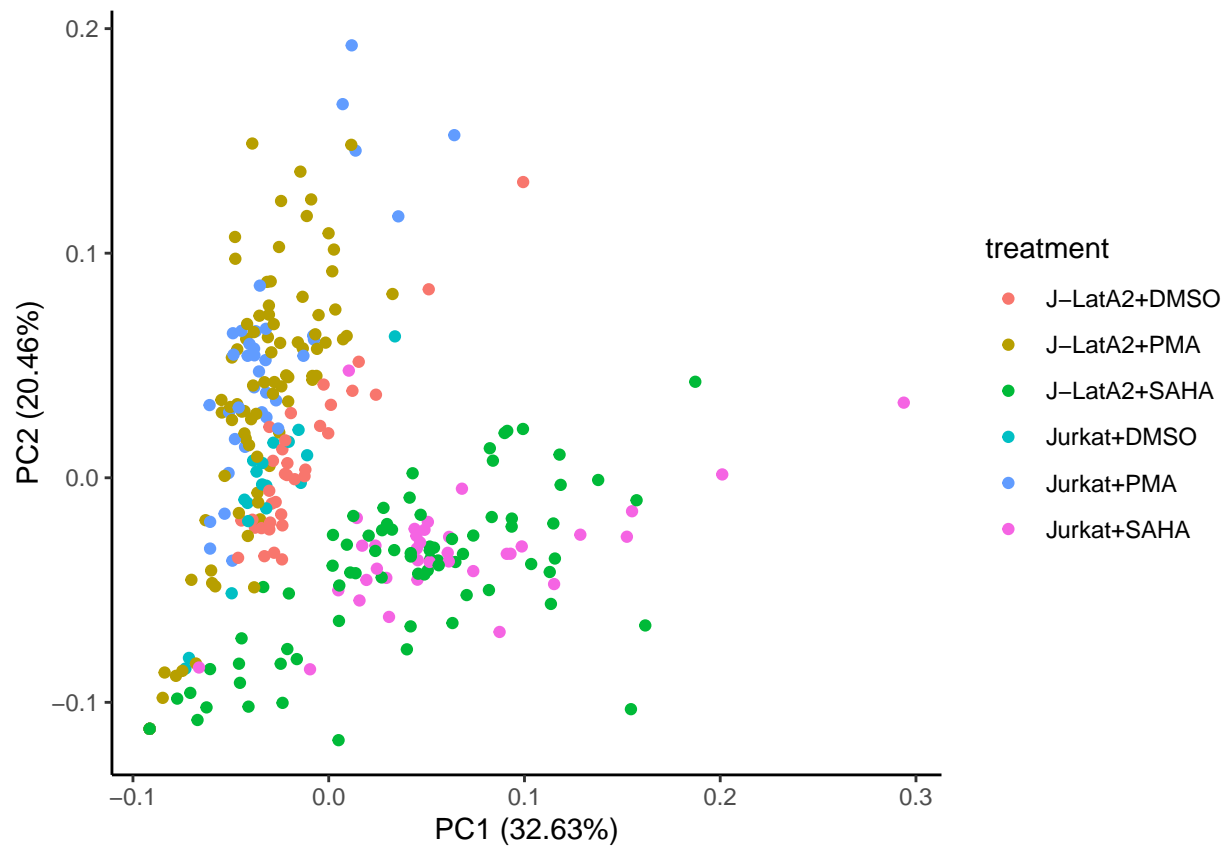
data <- data.frame(row.names= rownames(X1), experiment= as.factor(rep(c("A", "B", "C"), each=96)), treat= rep(c("DMSO", "SAHA", "PMA"), each=32))

colnames(data) <- c("experiment", "treatment")

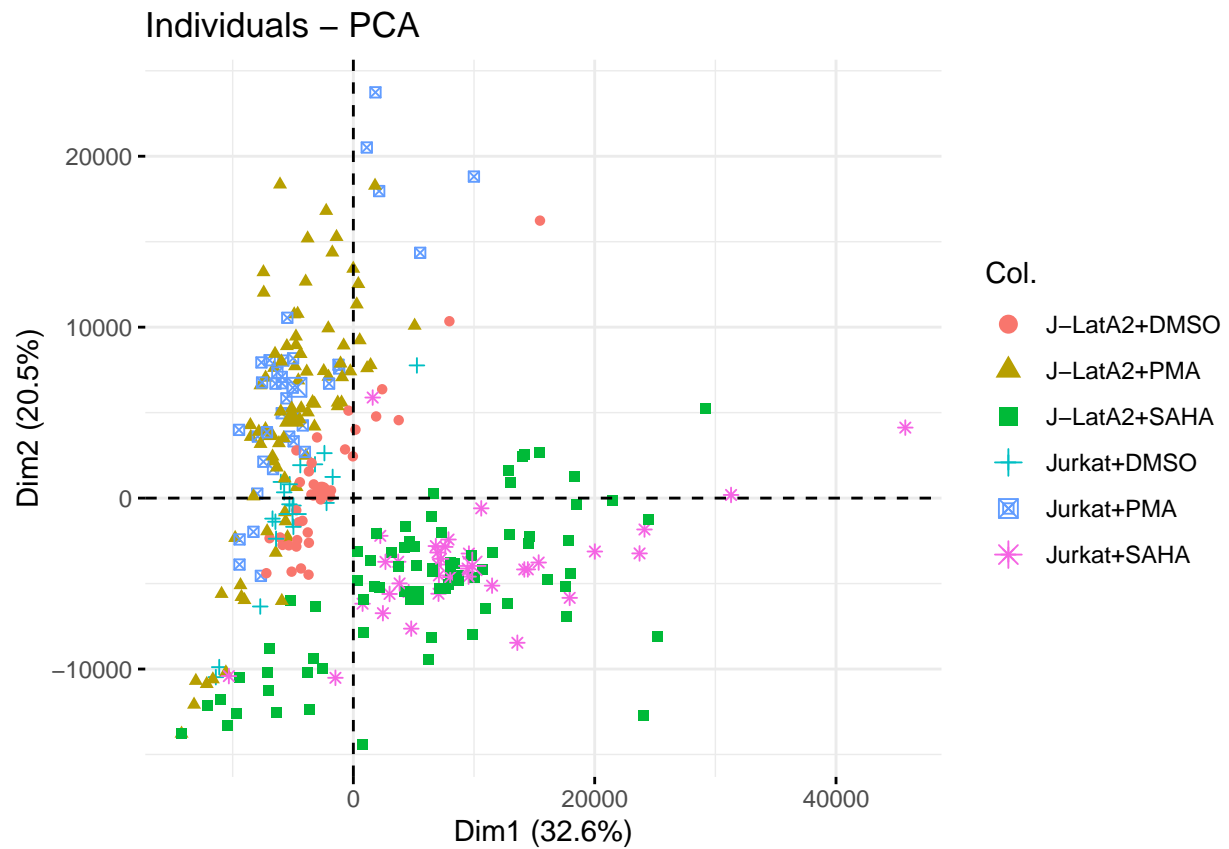
#PCA
PCA = stats::prcomp(X1)
plot(PCA$x)
```



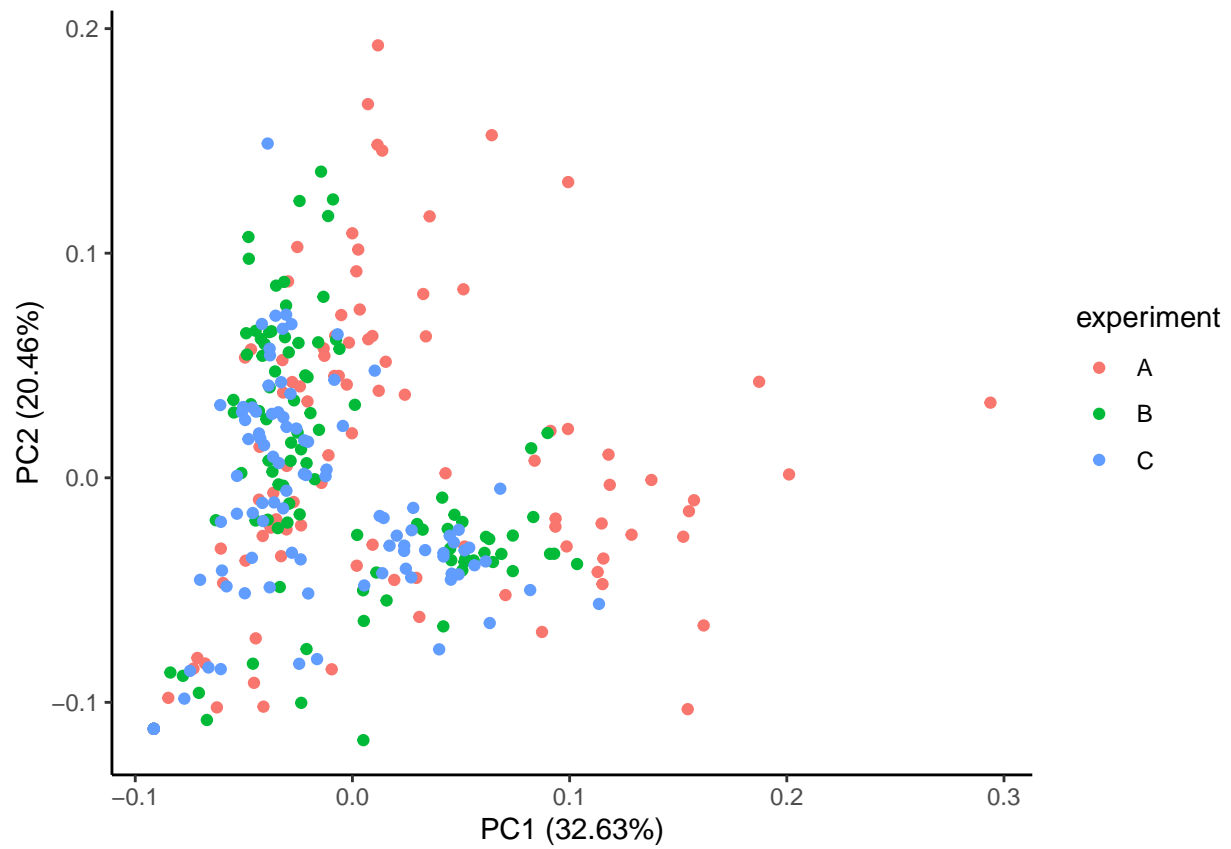
```
#coloring by treatment
#using autoplot
ggplot2::autoplot(PCA, data = data, colour = "treatment") + theme_classic()
```



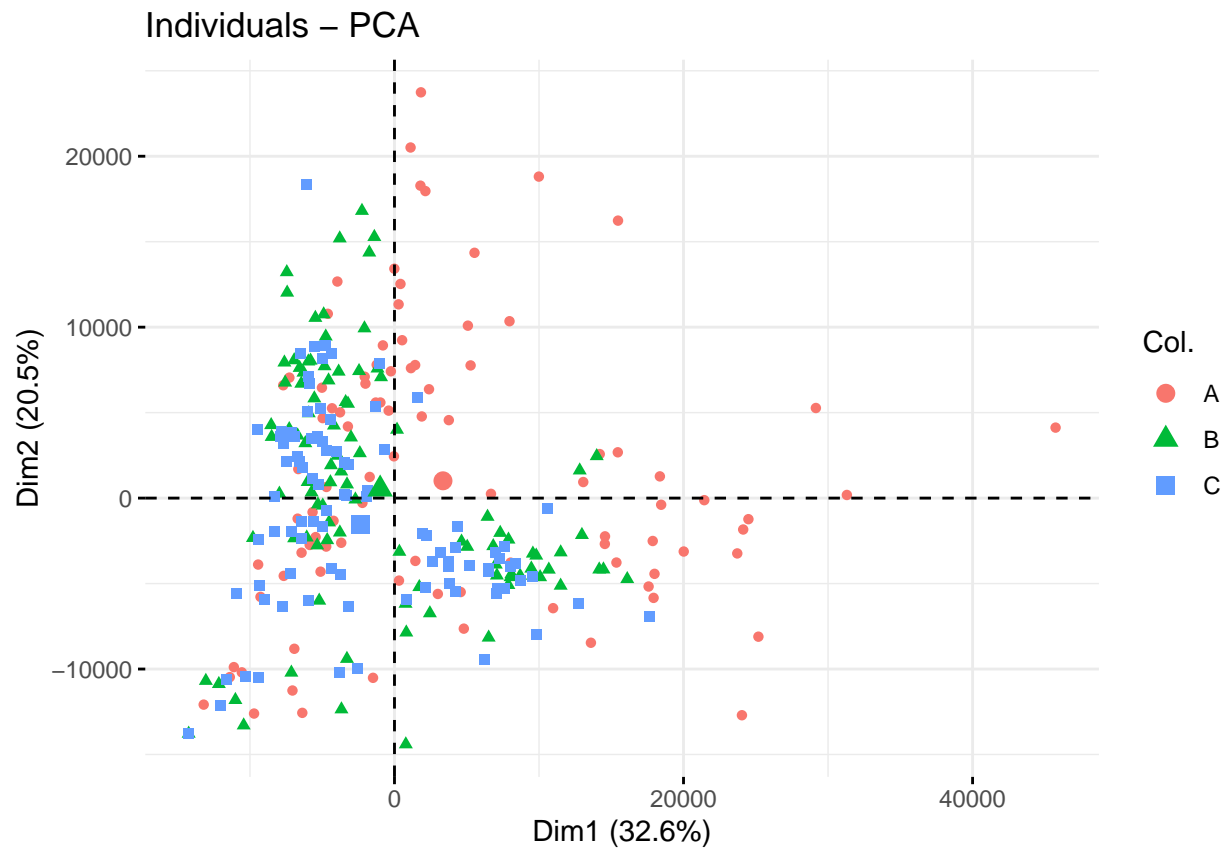
```
#using factoextra  
factoextra::fviz_pca_ind(PCA, geom = "point", col.ind = data$treatment)
```



```
#coloring by experiment
#using autoplot
ggplot2::autoplot(PCA, data = data, colour = "experiment") + theme_classic()
```



```
#using factoextra  
factoextra::fviz_pca_ind(PCA, geom = "point", col.ind = data$experiment)
```

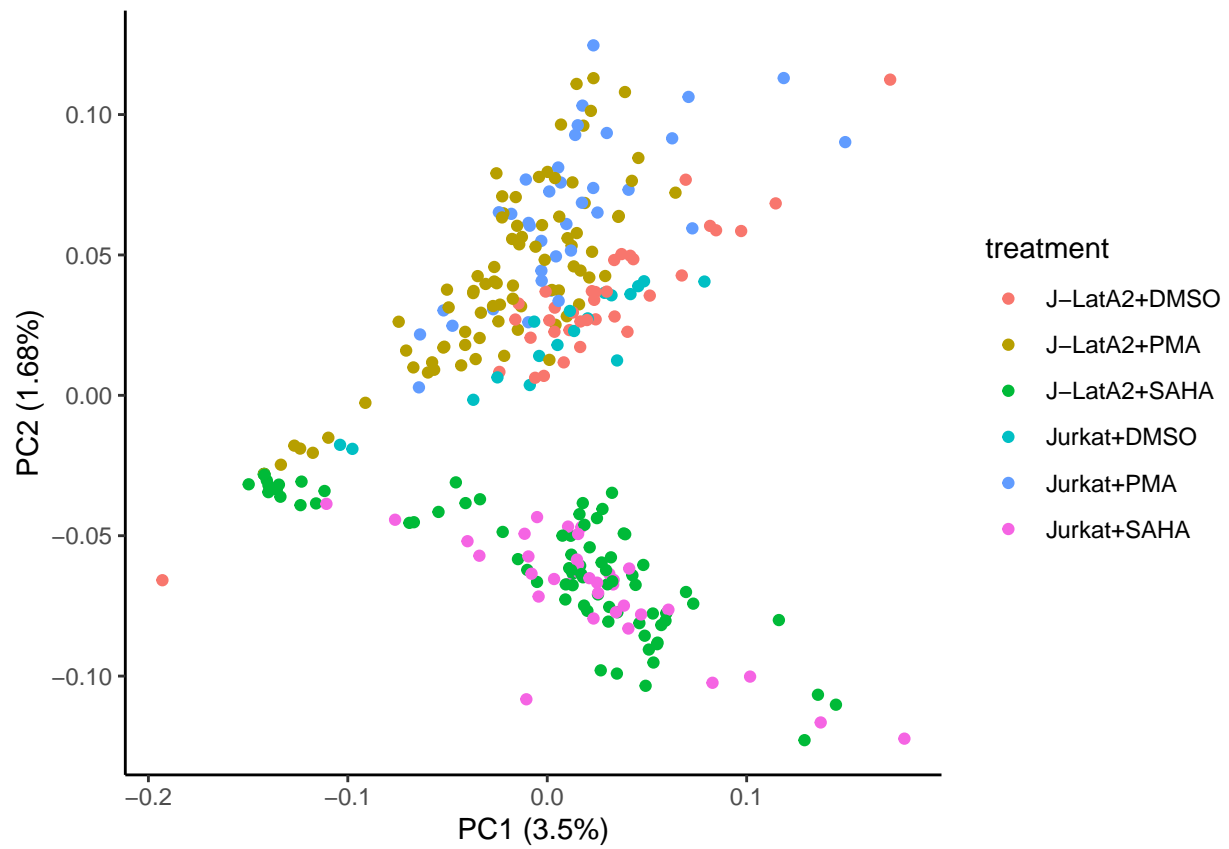
```
#creating scaled PCA
print(dim(X1))
```

```
## [1] 288 58780
```

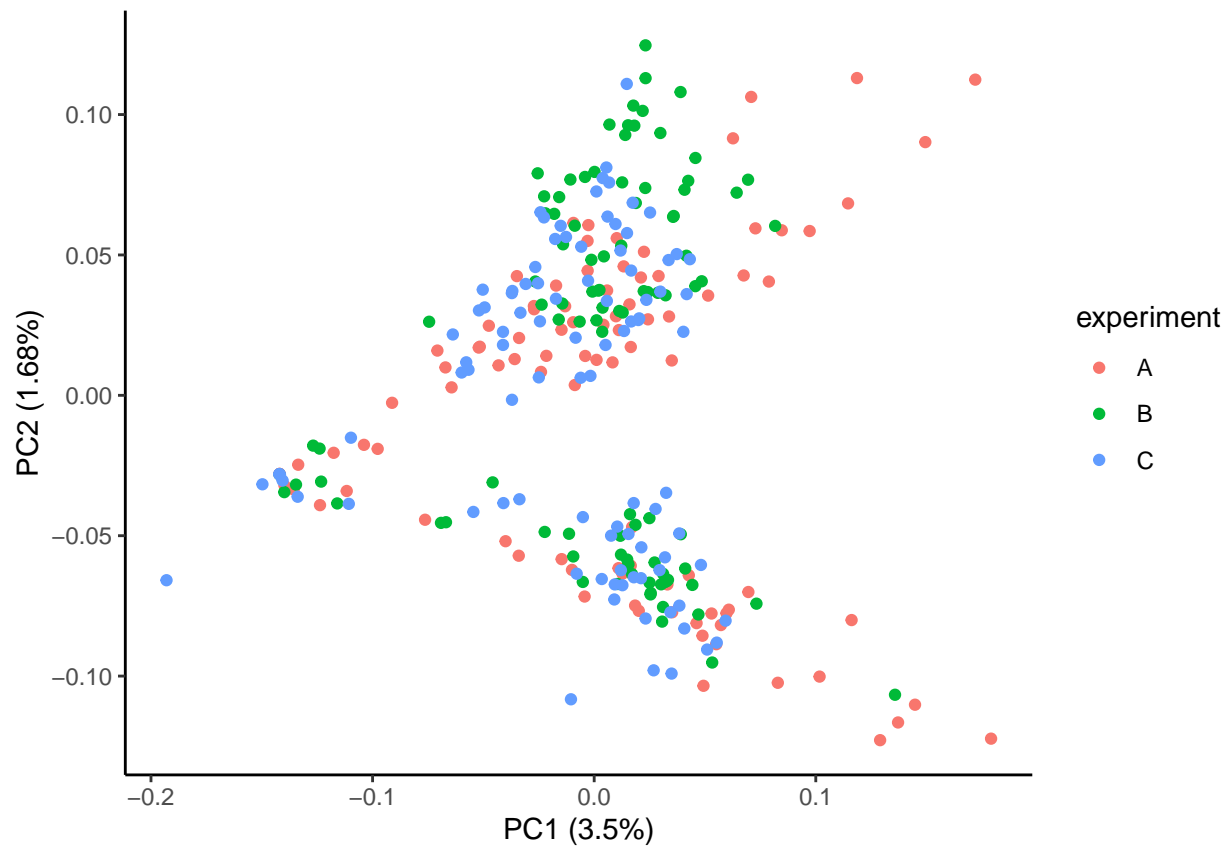
```
X1=X1[, colSums(X1)>0] ## tip: rowSums > 0
print(dim(X1))
```

```
## [1] 288 23104
```

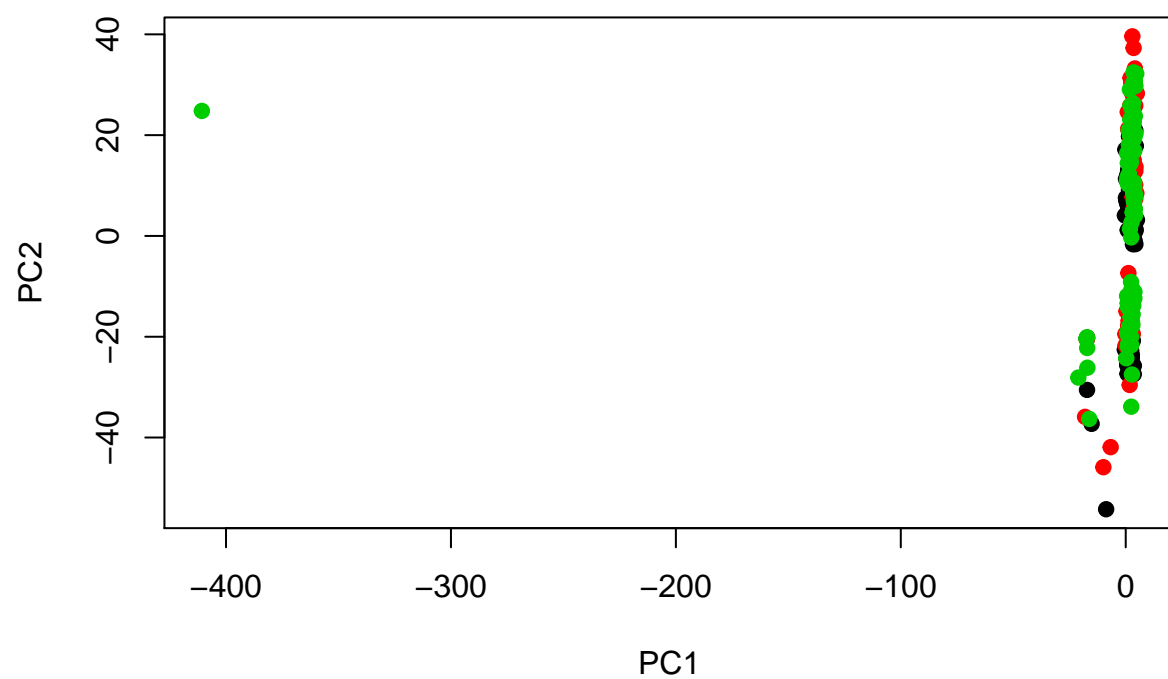
```
pca_res_scaled <- stats::prcomp(X1, scale=TRUE)
#colored by treatment
ggplot2::autoplot(pca_res_scaled, data = data, colour = "treatment") + theme_classic()
```



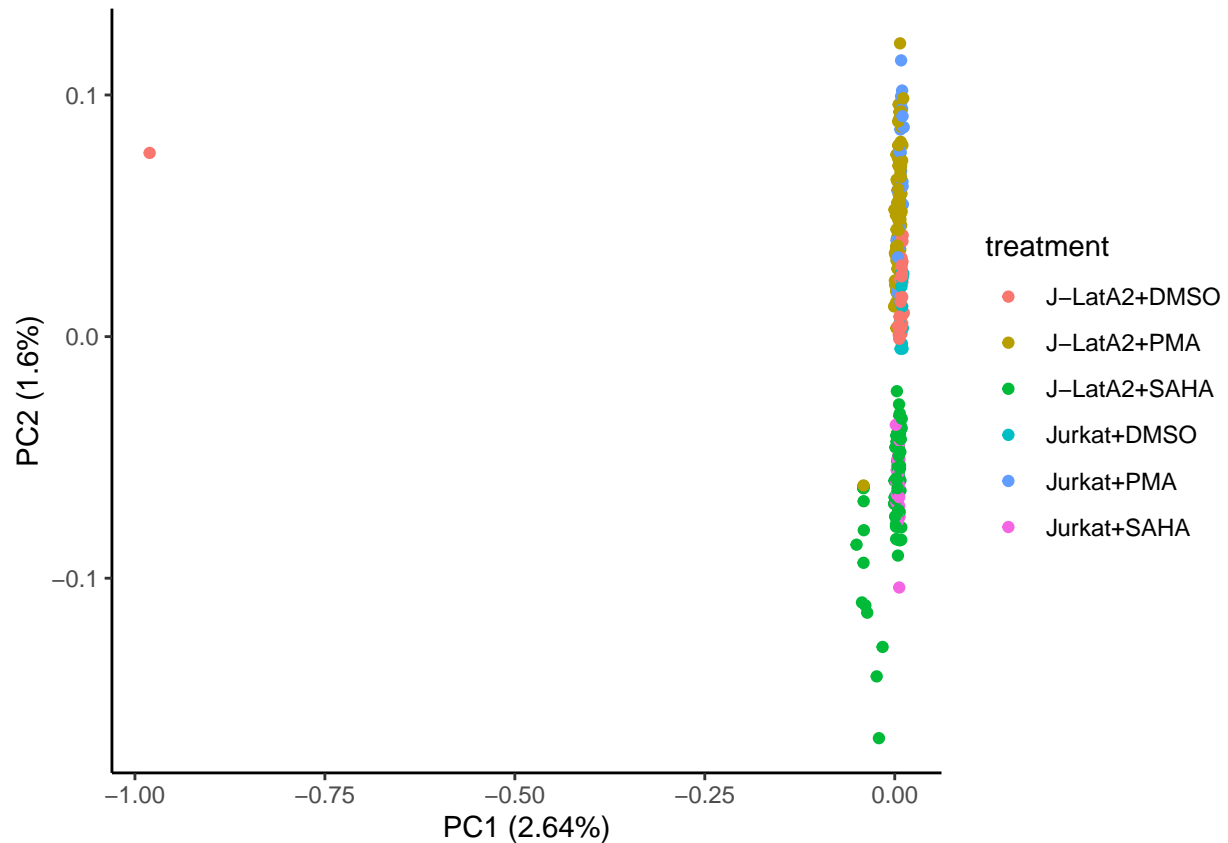
```
#colored by experiment
ggplot2::autoplot(pca_res_scaled, data = data, colour = "experiment") + theme_classic()
```



```
#creating normalized PCA  
Y1 = X1/rowSums(X1)  
normpca <- stats::prcomp(Y1, scale=TRUE)  
#colored by experiment  
plot(normpca$x, pch=19, col=data$experiment)
```



```
#colored by treatment, using autoplot  
ggplot2::autoplot(normpca, data=data, colour="treatment") + theme_classic()
```



DMSO is the control and it's very similar to PMA, meaning that PMA doesn't have a significant effect. On the other hand, SAHA is very different from both of them, meaning that it has a significant effect. There is no batch effect, because in the graph colored by experiment the samples are not clustered based on the experiment that they're part of. However, we scaled the PCA and the scaled PCAs are more precise

```
#tSNE
tsne1 <- Rtsne(X1)

## Check results generated
names(tsne1)
```

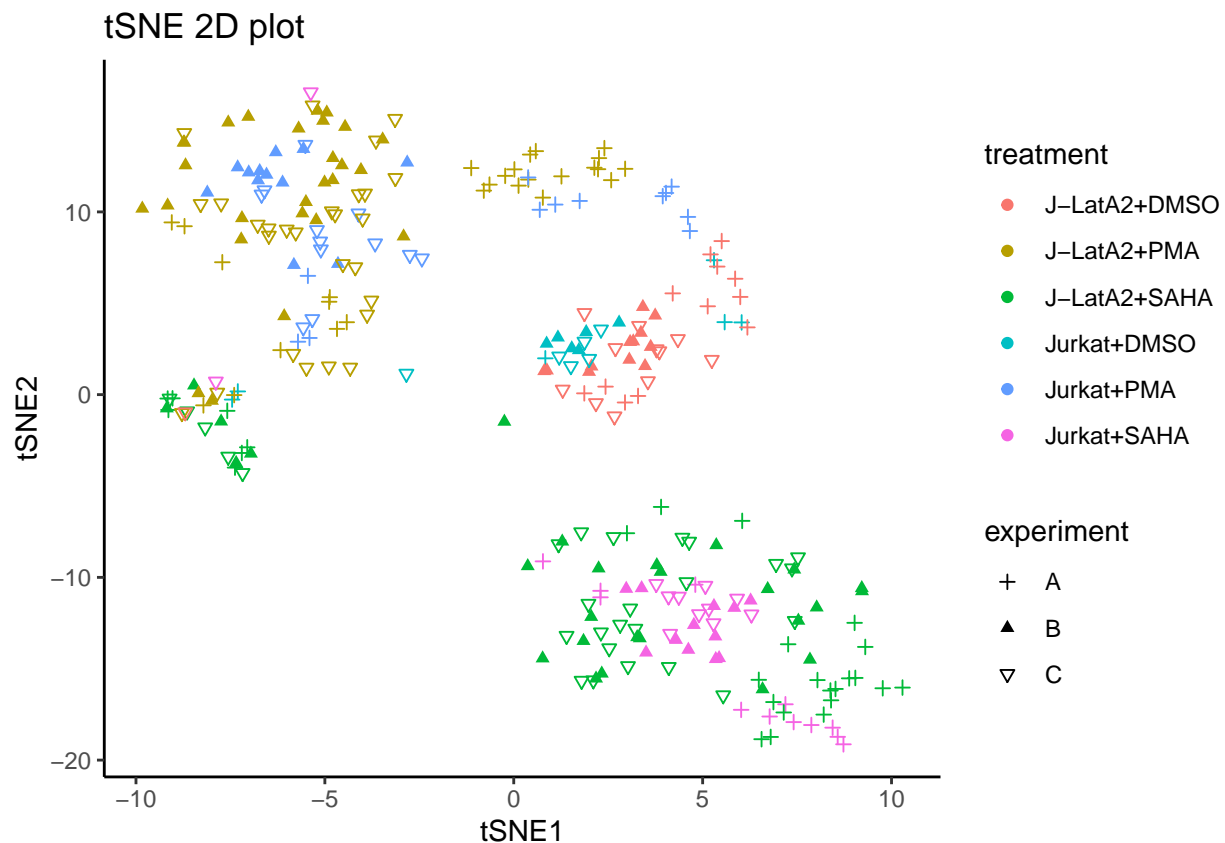
```
## [1] "N" "Y" "costs"
## [4] "itercosts" "origD" "perplexity"
## [7] "theta" "max_iter" "stop_lying_iter"
## [10] "mom_switch_iter" "momentum" "final_momentum"
## [13] "eta" "exaggeration_factor"
```

```
head(tsne1$Y)
```

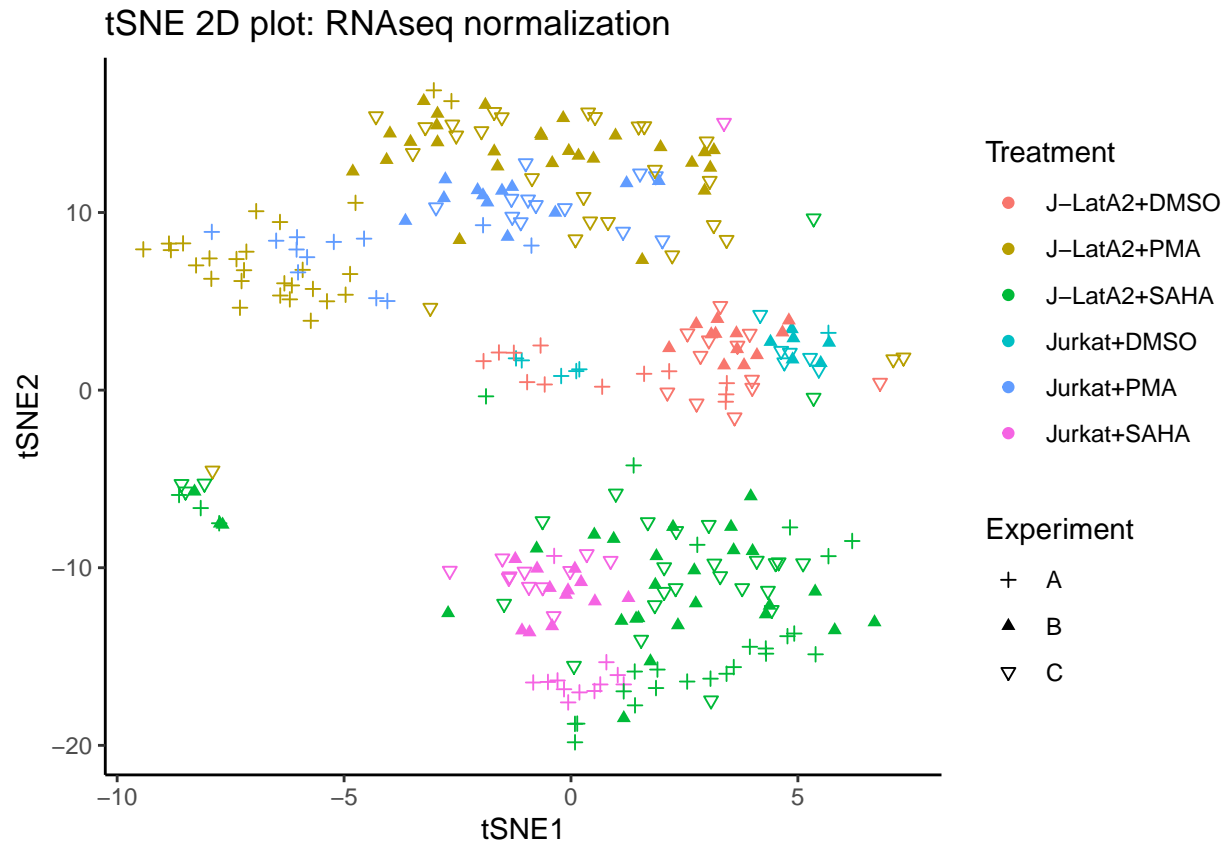
```
##           [,1]           [,2]
## [1,] -7.304639  0.17519535
## [2,]  2.295887 -11.09239025
## [3,] -5.711962  2.89508730
## [4,]  1.867787  0.07082732
## [5,] -7.055208 -2.88328641
## [6,] -7.583680 -0.88662495
```

```
tsne_data <- as.data.frame(tsne1$Y)
tsne_data$experiment <- data$experiment
tsne_data$treatment <- data$treatment

## ggplot
ggplot(tsne_data, aes(x=V1, y=V2)) +
  geom_point(aes(color=treatment, shape=experiment)) +
  scale_shape_manual(values = c(3,17, 6)) +
  ggtitle("tSNE 2D plot") + xlab("tSNE1") + ylab("tSNE2") +
  theme_classic()
```



```
## Normalized tSNE
X1=X1[, colSums(X1)>0]
Y = X1/rowSums(X1)
tSNE_norm <- Rtsne(Y, pca=TRUE)
tsne_data2 <- as.data.frame(tSNE_norm$Y)
tsne_data2$Experiment <- data$experiment
tsne_data2$Treatment <- data$treatment
## ggplot
tSNE2 <- ggplot(tsne_data2, aes(x=V1, y=V2)) +
  geom_point(aes(color=Treatment, shape=Experiment)) +
  scale_shape_manual(values = c(3,17,6)) +
  ggtitle("tSNE 2D plot: RNAseq normalization") + xlab("tSNE1") + ylab("tSNE2") + theme_classic()
print(tSNE2)
```



Question 4

```
#changing perplexity
set.seed(123) ## allows us to reproduce results

list_plots <- list()
for (p_value in c(1, 5, 10, 25, 50)) {
  print(paste0("tSNE perplexity=", p_value))

  ## Default parameters for Rtsne
  tsne_p <- Rtsne(X1, perplexity = p_value)

  tsne_data2 <- as.data.frame(tsne_p$Y)
  tsne_data2$Treatment <- data$treatment
  tsne_data2$Experiment <- data$experiment

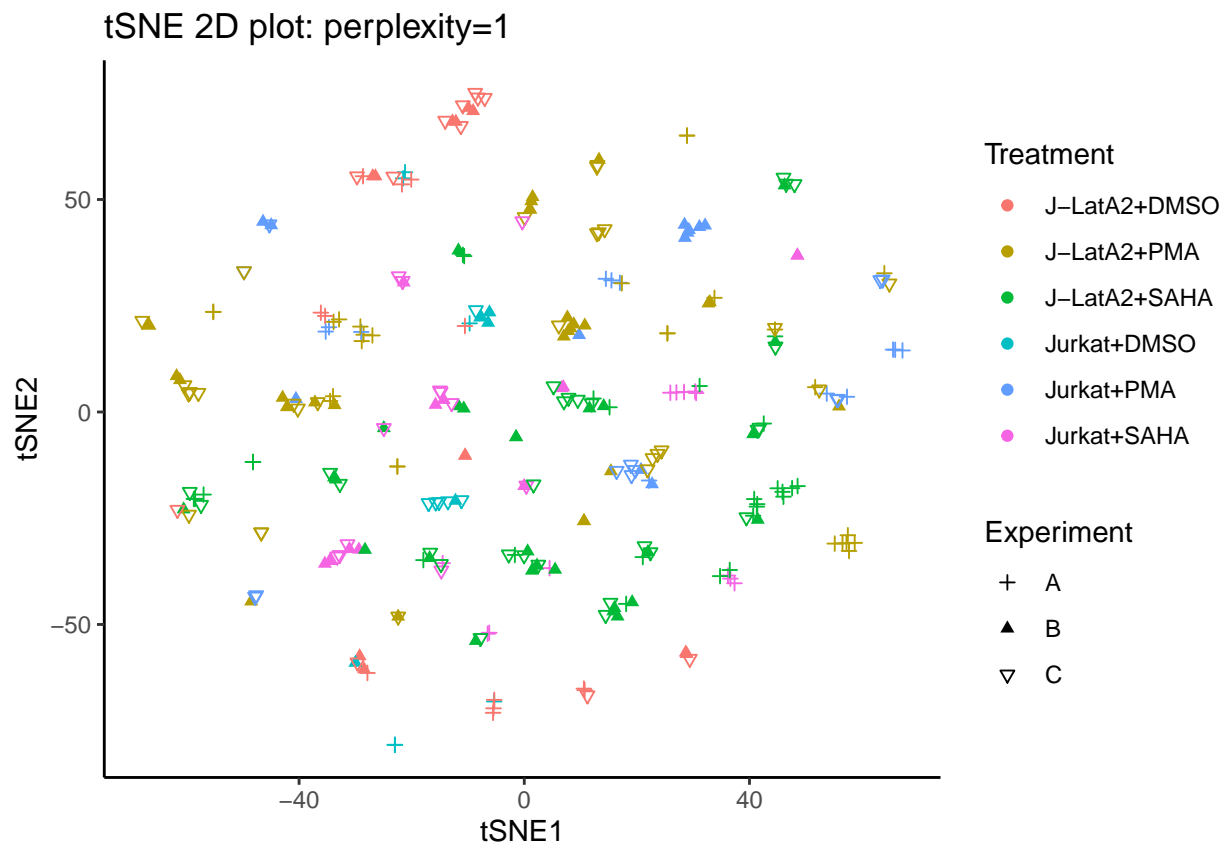
  ## ggplot
  plot_p <- ggplot(tsne_data2, aes(x=V1, y=V2)) +
    geom_point(aes(color=Treatment, shape=Experiment)) +
    scale_shape_manual(values = c(3,17,6)) +
    ggtitle(paste0("tSNE 2D plot: perplexity=", p_value)) + xlab("tSNE1") + ylab("tSNE2") +
    theme_classic()

  print(plot_p)

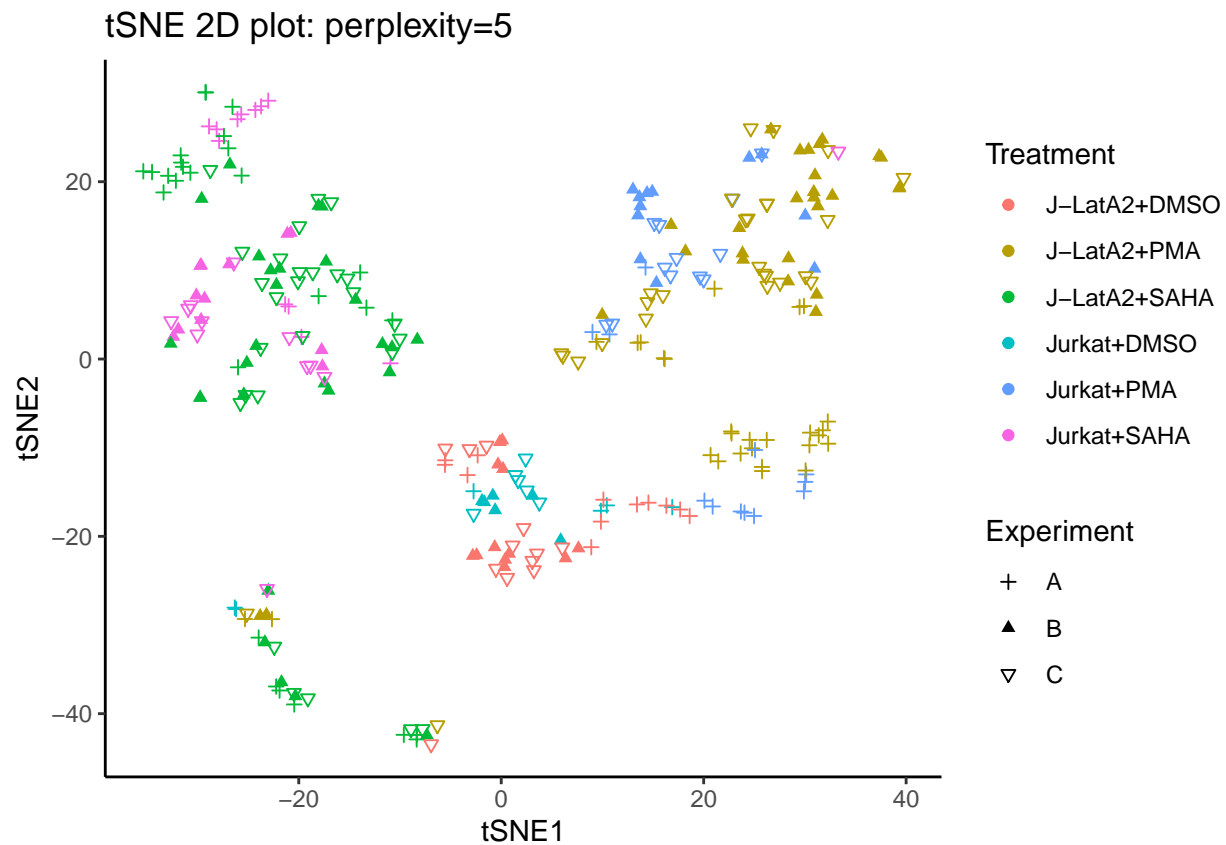
  # alternative: save plots in list_plots
}
```

```
list_plots[[paste0("perp_", p_value)]] = plot_p
}
```

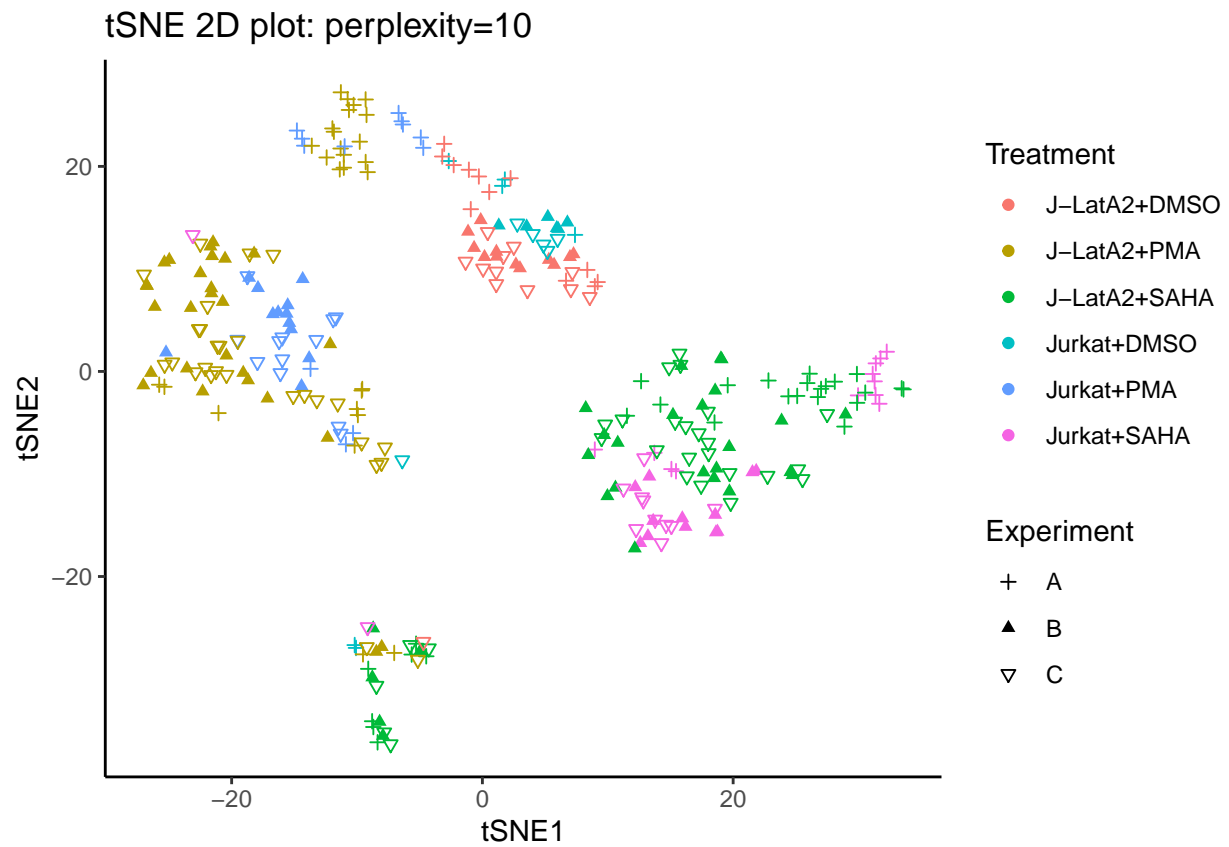
```
## [1] "tSNE perplexity=1"
```



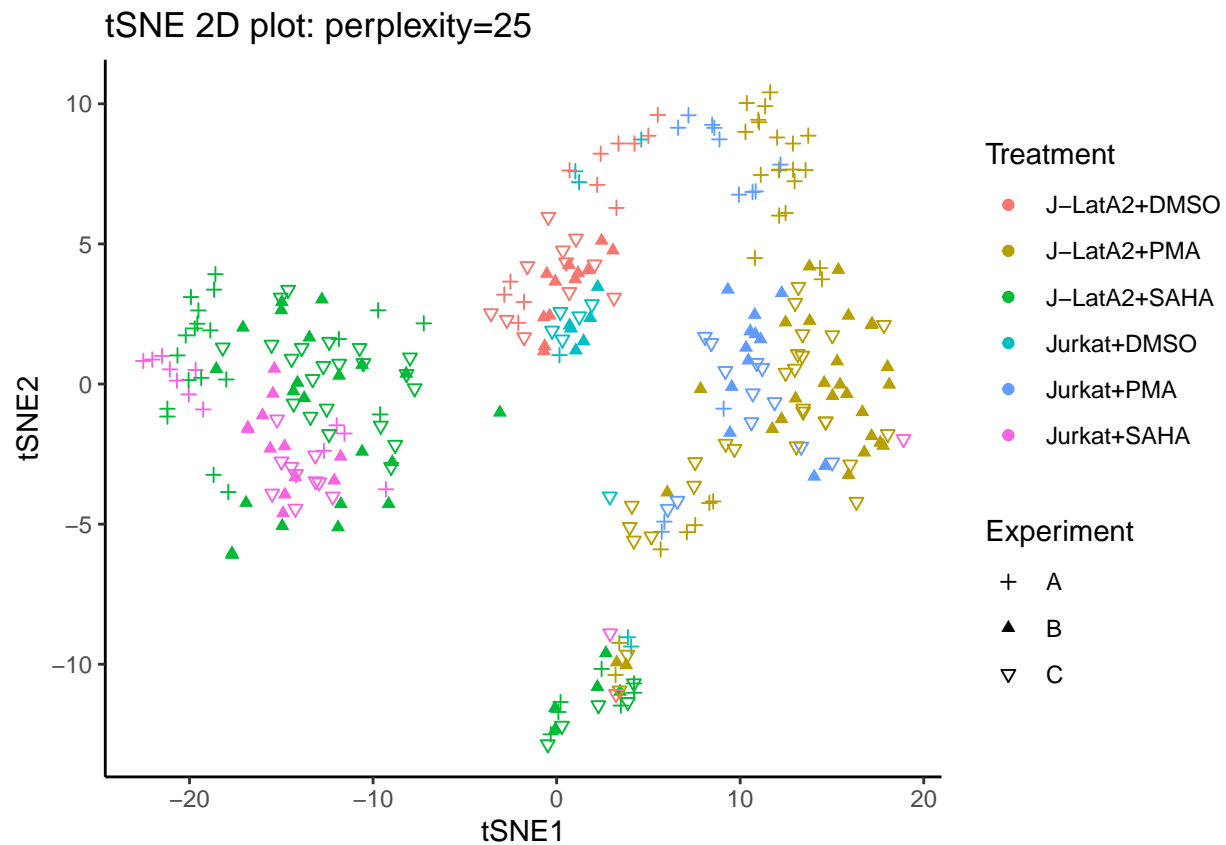
```
## [1] "tSNE perplexity=5"
```

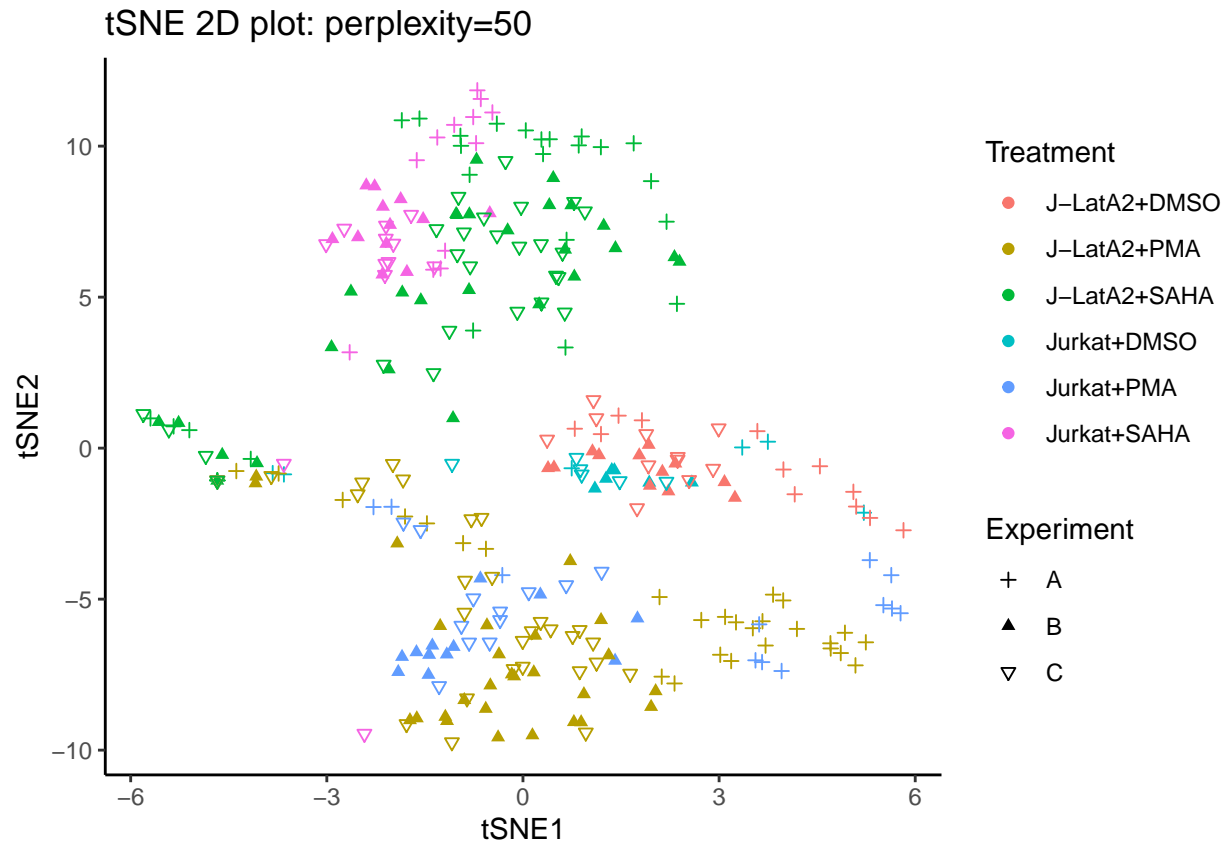
```
## [1] "tSNE perplexity=10"
```



[1] "tSNE perplexity=25"



```
## [1] "tSNE perplexity=50"
```

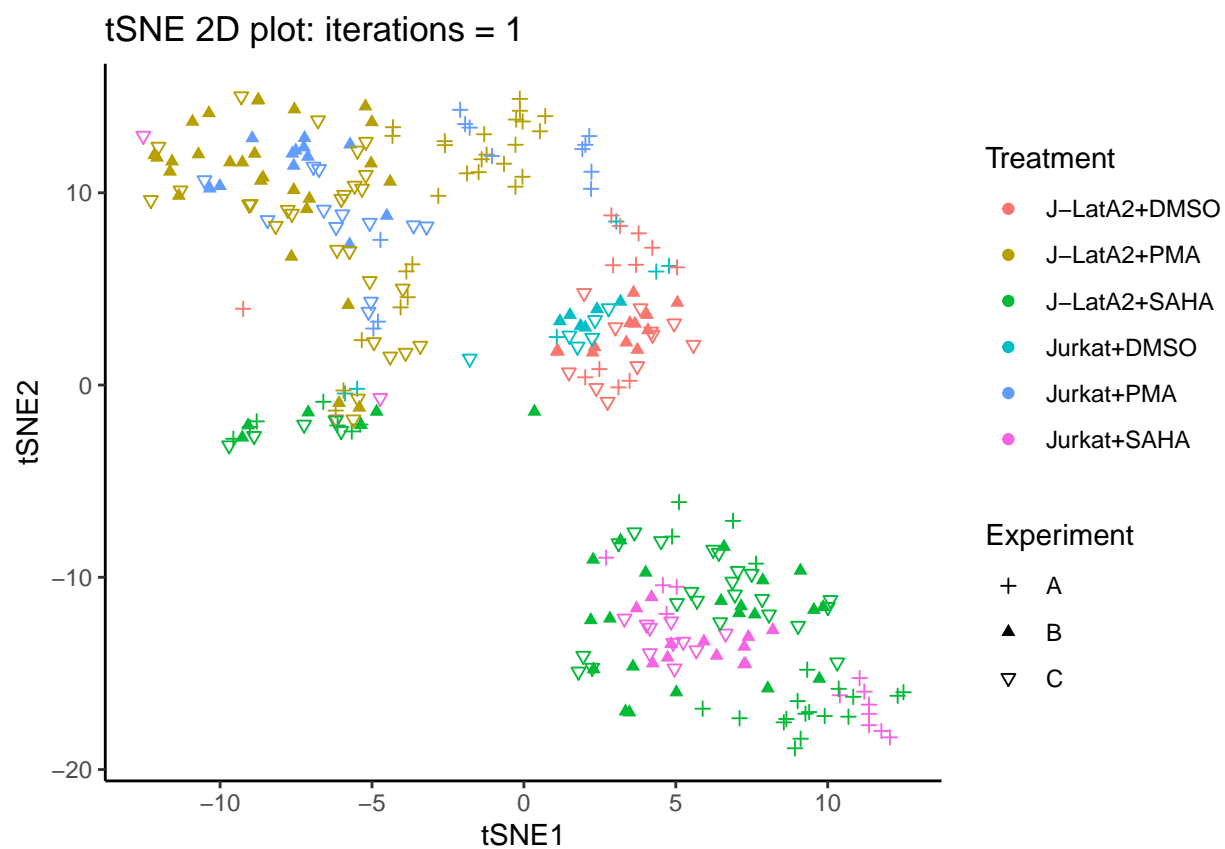


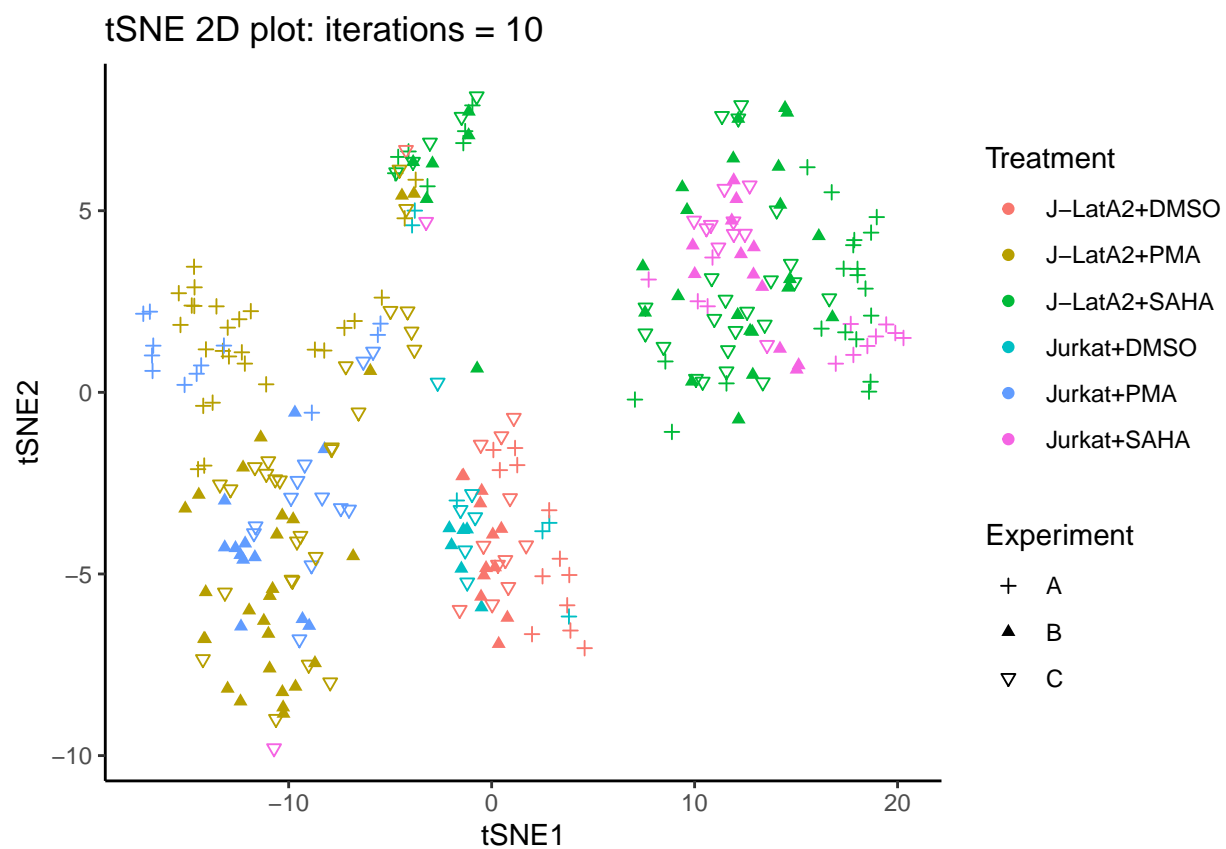
```
#changing max_iterations
set.seed(123) ## allows us to reproduce results

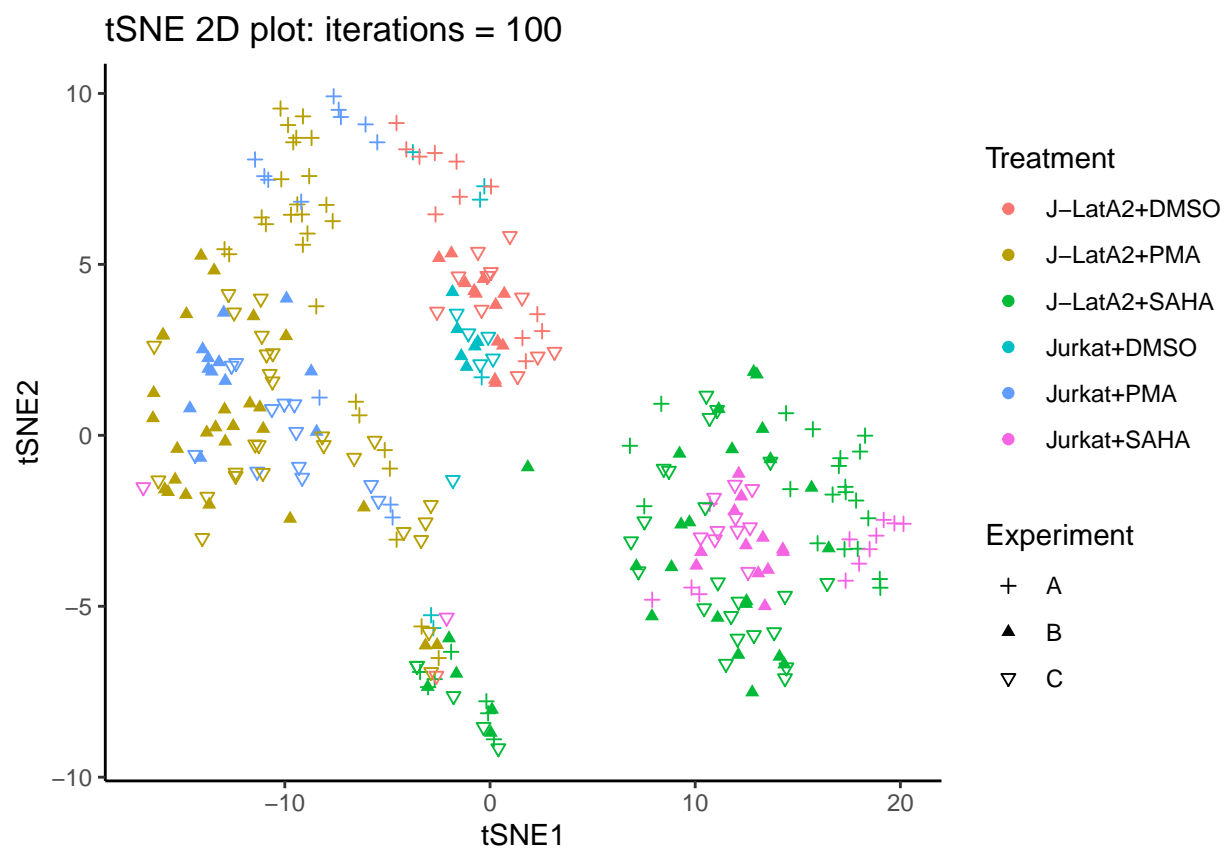
for (max_iter in c(1, 10, 100, 500, 1000)) {
  tsne <- Rtsne(X1)

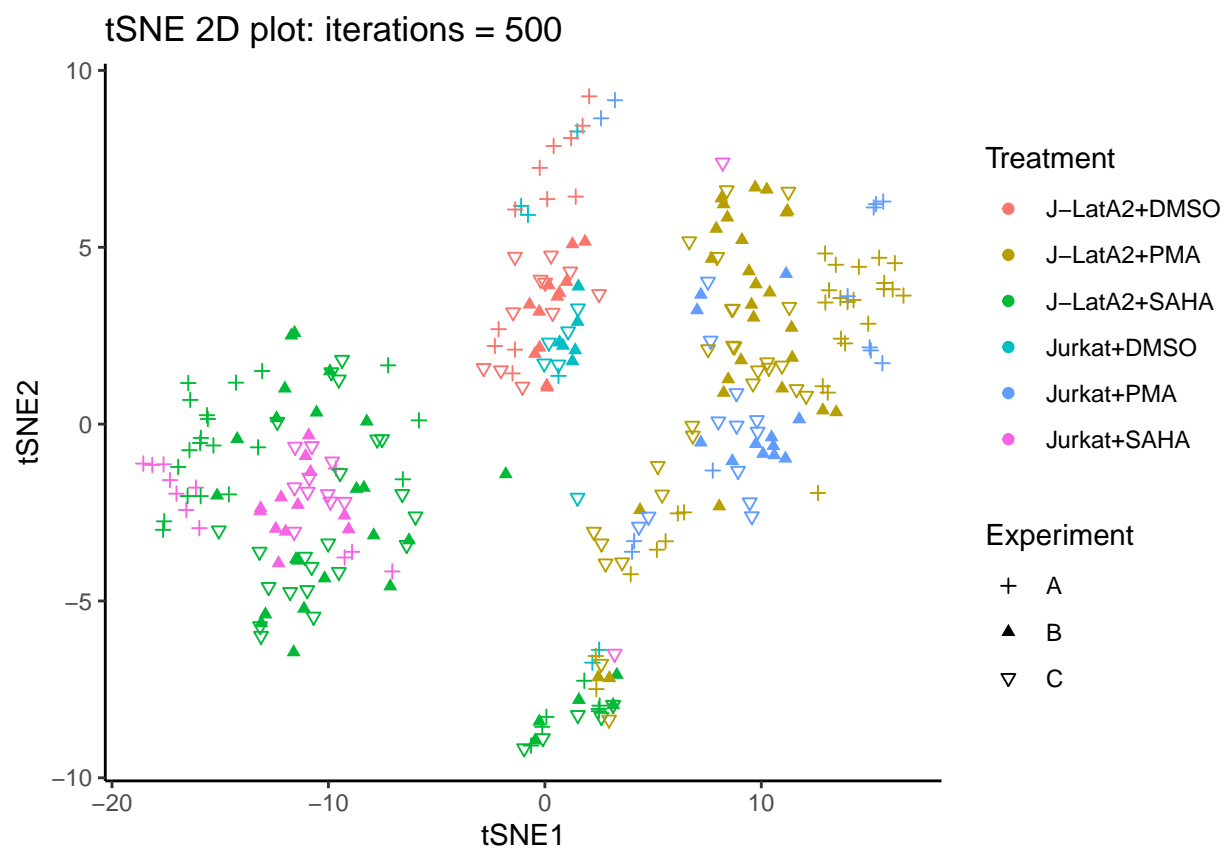
  tsne_data <- as.data.frame(tsne$Y)
  tsne_data$Experiment <- data$experiment
  tsne_data$Treatment <- data$treatment

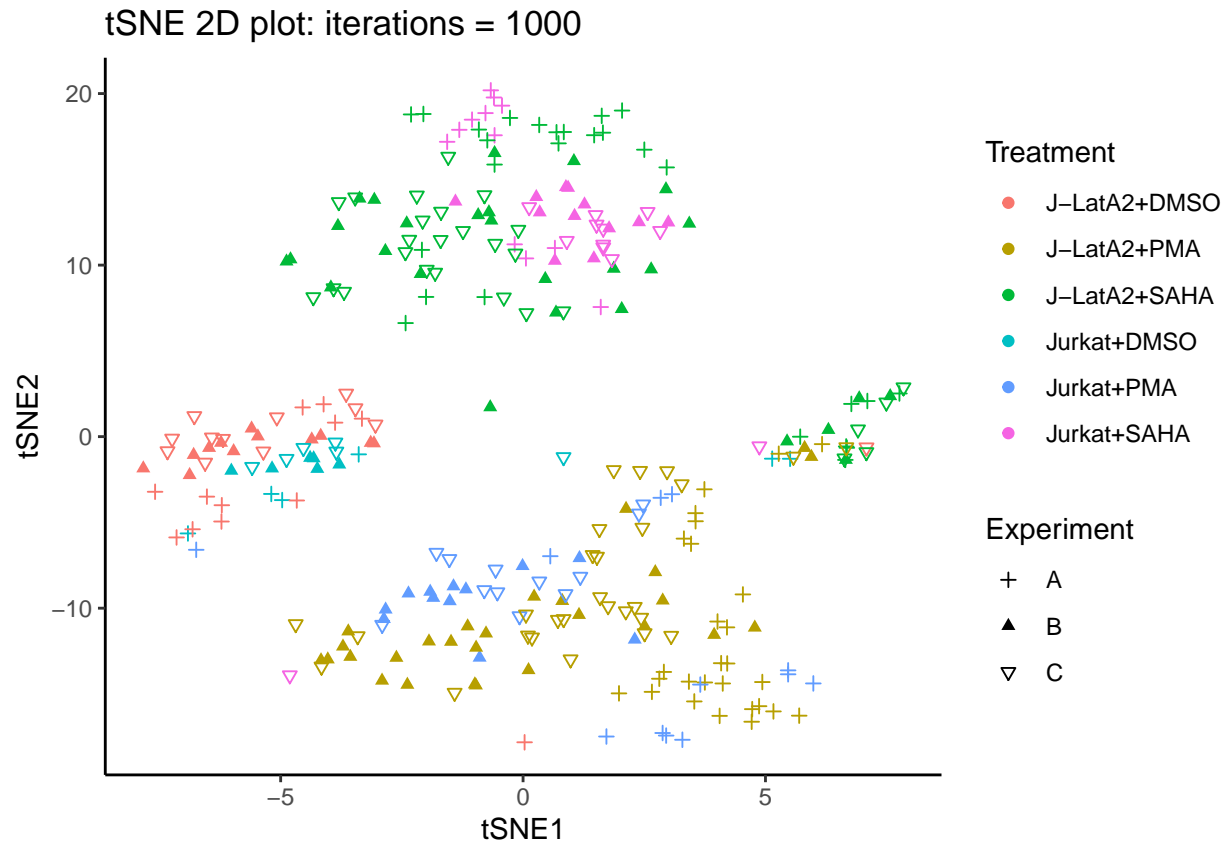
  library(ggplot2)
  p <- ggplot(tsne_data, aes(x=V1, y=V2)) +
    geom_point(aes(color=Treatment, shape=Experiment)) +
    scale_shape_manual(values = c(3,17,6)) +
    ggtitle(paste("tSNE 2D plot: iterations =", max_iter)) + xlab("tSNE1") + ylab("tSNE2") +
    theme_classic()
  print(p)
}
```











In the previous plots we show how increasing the perplexity or the iterations numbers for a tSNE can generate more dense and defined and more distinct clusters, respectively.

QUESTION 5

No, it is pretty clear from both the PCA and the tSNE graphs that SAHA and PMA have a different effect on the cells as the points associated to each drug are grouped very distinctly in two separate clusters. In particular, from the PCA plot we see that PMA treated cells are observed to be closer in similarity to DMSO (control) cells, so we might assume that PMA has a smaller effect than SAHA.

QUESTION 6

The PC1 separates dead from alive cells, meanwhile PC2 separates differently treated cells. On the right we have cells with low expression of genes, on the left with high expression.

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