

experiment 3

1.

```
d25 <- conf.design(rbind(c(A = 1, B = 1, C = 1, D = 0, E = 0),
                        c(A = 1, B = 0, C = 0, D = 1, E = 1)), p = 2)

d25

d25 <- d25 %>% mutate(e1=ifelse(A==1, "a", ""),
                    e2=ifelse(B==1, "b", ""),
                    e3=ifelse(C==1, "c", ""),
                    e4=ifelse(D==1, "d", ""),
                    e5=ifelse(E==1, "e", ""),
                    effect=paste(e1,e2,e3,e4,e5, sep="")) %>% select(-c(e1,e2,e3,e4,e5))

d25
```

	Blocks	A	B	C	D	E	effect
1	00	0	0	0	0	0	
2	00	0	1	1	0	0	bc
3	00	1	1	0	1	0	abd
4	00	1	0	1	1	0	acd
5	00	1	1	0	0	1	abe
6	00	1	0	1	0	1	ace
7	00	0	0	0	1	1	de
8	00	0	1	1	1	1	bcde
9	01	1	1	0	0	0	ab
10	01	1	0	1	0	0	ac
11	01	0	0	0	1	0	d
12	01	0	1	1	1	0	bcd
13	01	0	0	0	0	1	e
14	01	0	1	1	0	1	bce
15	01	1	1	0	1	1	abde
16	01	1	0	1	1	1	acde
17	10	0	1	0	0	0	b
18	10	0	0	1	0	0	c
19	10	1	0	0	1	0	ad
20	10	1	1	1	1	0	abcd
21	10	1	0	0	0	1	ae
22	10	1	1	1	0	1	abce
23	10	0	1	0	1	1	bde
24	10	0	0	1	1	1	cde
25	11	1	0	0	0	0	a
26	11	1	1	1	0	0	abc
27	11	0	1	0	1	0	bd
28	11	0	0	1	1	0	cd
29	11	0	1	0	0	1	be
30	11	0	0	1	0	1	ce
31	11	1	0	0	1	1	ade
32	11	1	1	1	1	1	abcde

```

decimals <- 0:31
m <- sapply(decimals,function(x){ as.integer(intToBits(x))})
m[1:5,]

m1<-as.data.frame(t(m[1:5,]))
m1

m2<-m1 %>% mutate(L1=V1+V4+V5, L2=V2+V3+V5, m1=mod(L1,2), m2=mod(L2,2),
  block=ifelse(m1==0 & m2==0, 1,
    ifelse(m1==0 & m2==1, 2,
      ifelse(m1==1 & m2==0, 3,4))))

```

```
m2 <- m2 %>% arrange(block)
m2

library(readr)
group_response <- read_csv("group project/group_response.csv")
y2<- group_response["y2"][complete.cases(group_response["y2"]), ]
y2

df<- cbind(d25, y2)
df
```

	Blocks	A	B	C	D	E	effect	y2
1	00	0	0	0	0	0		1.68
2	00	0	1	1	0	0	bc	1.98
3	00	1	1	0	1	0	abd	4.98
4	00	1	0	1	1	0	acd	5.70
5	00	1	1	0	0	1	abe	3.24
6	00	1	0	1	0	1	ace	3.44
7	00	0	0	0	1	1	de	9.97
8	00	0	1	1	1	1	bcde	9.07
9	01	1	1	0	0	0	ab	2.07
10	01	1	0	1	0	0	ac	2.44
11	01	0	0	0	1	0	d	7.77
12	01	0	1	1	1	0	bcd	9.43
13	01	0	0	0	0	1	e	4.09
14	01	0	1	1	0	1	bce	4.53
15	01	1	1	0	1	1	abde	11.75
16	01	1	0	1	1	1	acde	16.30
17	10	0	1	0	0	0	b	2.69
18	10	0	0	1	0	0	c	2.30
19	10	1	0	0	1	0	ad	6.35
20	10	1	1	1	1	0	abcd	6.75
21	10	1	0	0	0	1	ae	4.34
22	10	1	1	1	0	1	abce	4.20
23	10	0	1	0	1	1	bde	10.06
24	10	0	0	1	1	1	cde	9.35
25	11	1	0	0	0	0	a	3.22
26	11	1	1	1	0	0	abc	3.55
27	11	0	1	0	1	0	bd	9.33
28	11	0	0	1	1	0	cd	9.52
29	11	0	1	0	0	1	be	5.91
30	11	0	0	1	0	1	ce	5.83
31	11	1	0	0	1	1	ade	13.23
32	11	1	1	1	1	1	abcde	16.40

ABC, ADE are confounded with blocks => it means BCDE = (ABC)(ADE) is also confounded with blocks

2.

```
m4<-aov(y2 ~ A*B*C*D*E + Error(Blocks), data=df)
summary(m4)
```

```

meanall<-mean(df$y2)
meanall

summ<-df %>% group_by(Blocks) %>% summarise(meanblk=mean(y), nobs=n()) %>% mutate(ssbl=nobs*(meanblk-meanall)^2)
summ

sum(summ$ssbl)

```

Error: Blocks

	Df	Sum Sq	Mean Sq
A:B:C	1	6.65	6.65
A:D:E	1	48.19	48.19
B:C:D:E	1	0.22	0.22

Error: Within

	Df	Sum Sq	Mean Sq
A	1	0.62	0.62
B	1	0.01	0.01
C	1	3.19	3.19
D	1	315.32	315.32
E	1	84.34	84.34
A:B	1	0.65	0.65
A:C	1	2.58	2.58
B:C	1	0.09	0.09
A:D	1	2.80	2.80
B:D	1	0.05	0.05
C:D	1	2.03	2.03
A:E	1	17.60	17.60
B:E	1	0.32	0.32
C:E	1	0.27	0.27
D:E	1	13.33	13.33
A:B:D	1	0.06	0.06
A:C:D	1	2.06	2.06
B:C:D	1	0.17	0.17
A:B:E	1	0.01	0.01
A:C:E	1	2.51	2.51
B:C:E	1	0.10	0.10
B:D:E	1	0.16	0.16
C:D:E	1	0.35	0.35
A:B:C:D	1	0.20	0.20
A:B:C:E	1	0.02	0.02
A:B:D:E	1	0.02	0.02
A:C:D:E	1	3.89	3.89
A:B:C:D:E	1	0.39	0.39

```

> summ
# A tibble: 4 × 4
  Blocks meanblk nobs ssbl
  <fct>   <dbl> <int> <dbl>
1 00      0.593     8 3.08
2 01     -0.151     8 0.124
3 10     -0.224     8 0.312
4 11     -0.326     8 0.713
> sum(summ$ssbl)
[1] 4.227558
> summ<-df %>% group_by(Blocks) %>% summarise(meanblk=mean(y2), nobs=n()) %>% mutate(ssbl=nobs*(meanblk-meanall)^2)
> summ
# A tibble: 4 × 4
  Blocks meanblk nobs ssbl
  <fct>   <dbl> <int> <dbl>
1 00      5.01     8 20.5
2 01      7.30     8  3.80
3 10      5.76     8  5.83
4 11      8.37     8 24.9
> sum(summ$ssbl)
[1] 55.05993

```

3. based on anova table and SSBL, factors D, E are significant, while blocks are not significant.

A is confounded with BCDE and ABE.

B is confounded with ACDE and ABE.

C is confounded with ABDE.

D is confounded with ABCE and ADE.

E is confounded with ABCD and ADE.

```

m1<-lm(data=df, y2 ~ D+E)

points<-get_regression_points(m1)
points

shapiro.test(points$residual)

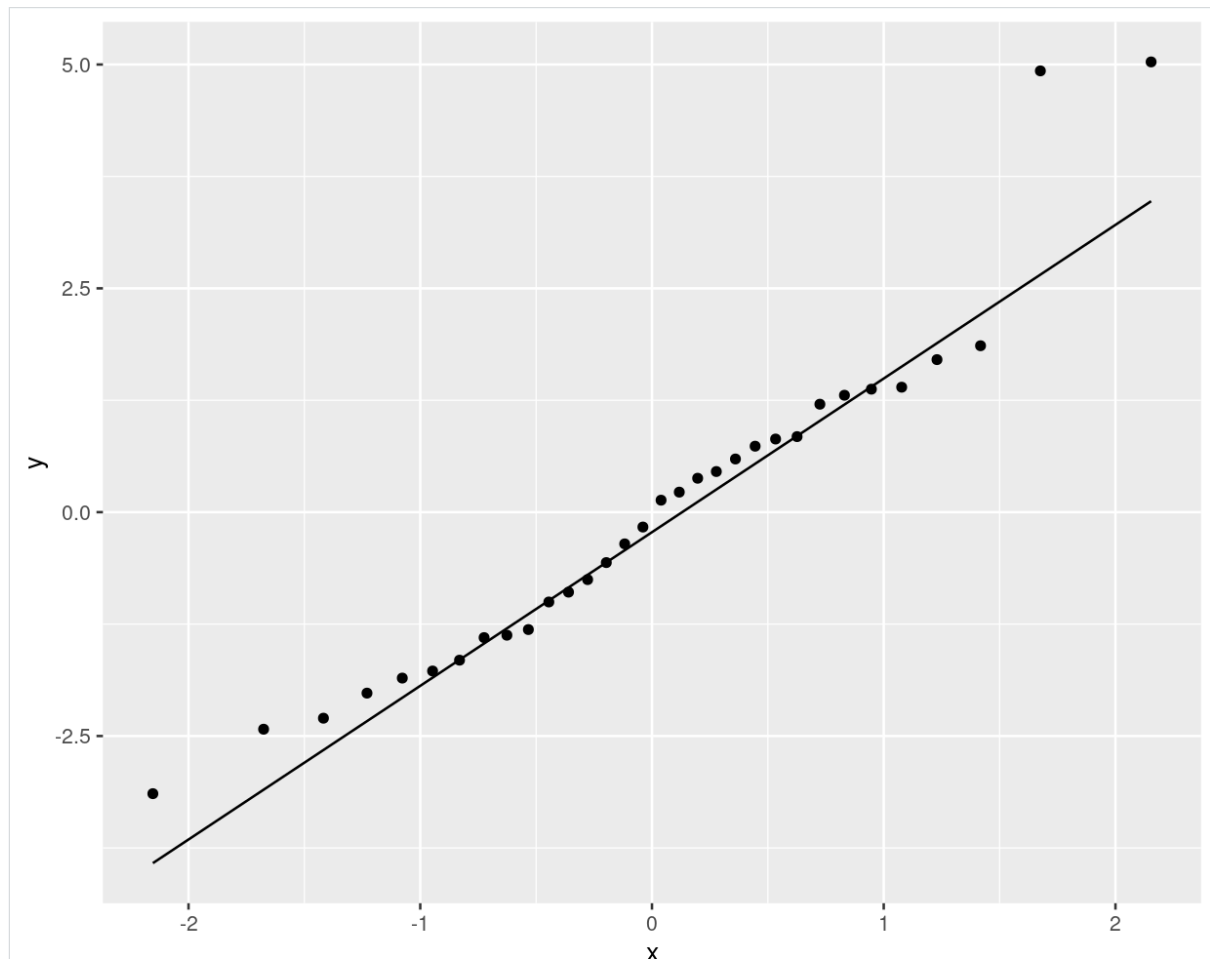
ggplot(data=points, aes(sample=residual)) + stat_qq() + stat_qq_line()

```

Shapiro-Wilk normality test

data: points\$residual

W = 0.93009, p-value = 0.0394



no obvious violation of assumptions.