

# DSC 680 Project 1 R

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```
#Set the working directory
setwd("C:/Users/Christine/Documents/Bellevue/DSC 680/Project 1")
```

Import data from file

```
#Import file
flu <- read.csv(file = "C:/Users/Christine/Documents/Bellevue/DSC 680/Project 1/fluprint_export.
csv", header = TRUE, na.strings = " ")
```

Display first five records of file

head(flu)

	donor_id	study_id	gen...	race	visit_id	visit_year	visit_day	visit_type_hai	visit
	<int>	<int>	<fctr>	<fctr>	<int>	<int>	<int>	<fctr>	
1	813	15	Female	Caucasian	2937	2014	0	pre	
2	813	15	Female	Caucasian	2937	2014	0	pre	
3	813	15	Female	Caucasian	2937	2014	0	pre	
4	813	15	Female	Caucasian	2937	2014	0	pre	
5	813	15	Female	Caucasian	2937	2014	0	pre	
6	813	15	Female	Caucasian	2937	2014	0	pre	

6 rows | 1-10 of 39 columns

Run stat.desc() function of file

stat.desc(flu)

	donor_id	study_id	gen...	r...	visit_id	visit_year	visit
	<dbl>	<dbl>	<lgl>	<lgl>	<dbl>	<dbl>	
nbr.val	1.561180e+05	1.561180e+05	NA	NA	1.561180e+05	1.561180e+05	1.561180
nbr.null	0.000000e+00	0.000000e+00	NA	NA	0.000000e+00	0.000000e+00	1.471340
nbr.na	0.000000e+00	0.000000e+00	NA	NA	0.000000e+00	0.000000e+00	0.000000
min	1.000000e+00	1.500000e+01	NA	NA	1.000000e+00	2.007000e+03	0.000000
max	8.130000e+02	3.000000e+01	NA	NA	2.937000e+03	2.015000e+03	7.000000

	donor_id <dbl>	study_id <dbl>	gen... <lgl>	r... <lgl>	visit_id <dbl>	visit_year <dbl>	visit
range	8.120000e+02	1.500000e+01	NA	NA	2.936000e+03	8.000000e+00	7.000000
sum	6.129445e+07	3.022399e+06	NA	NA	1.399764e+08	3.140742e+08	9.902000
median	4.020000e+02	1.800000e+01	NA	NA	8.710000e+02	2.013000e+03	0.000000
mean	3.926162e+02	1.935971e+01	NA	NA	8.966065e+02	2.011774e+03	6.342638
SE.mean	4.821530e-01	8.187006e-03	NA	NA	1.720132e+00	4.456957e-03	8.025971
1-10 of 14 rows   1-8 of 39 columns						Previous	1 2 Next

Run str() function of file

```
str(flu)
```

```

## 'data.frame': 156118 obs. of 38 variables:
## $ donor_id : int 813 813 813 813 813 813 813 813 813 813 ...
## $ study_id : int 15 15 15 15 15 15 15 15 15 15 ...
## $ gender : Factor w/ 2 levels "Female","Male": 1 1 1 1 1 1 1 1 1 1 ...
## $ race : Factor w/ 7 levels "American Indian or Alaska Native",...: 4 4
4 4 4 4 4 4 4 ...
## $ visit_id : int 2937 2937 2937 2937 2937 2937 2937 2937 2937 2937 ...
## $ visit_year : int 2014 2014 2014 2014 2014 2014 2014 2014 2014 2014 ...
## $ visit_day : int 0 0 0 0 0 0 0 0 0 0 ...
## $ visit_type_hai : Factor w/ 2 levels "pre","single": 1 1 1 1 1 1 1 1 1 1 ...
## $ visit_age : num 23 23 23 23 23 23 23 23 23 23 ...
## $ cmv_status : Factor w/ 3 levels "0","1","NULL": 1 1 1 1 1 1 1 1 1 1 ...
## $ ebv_status : Factor w/ 3 levels "0","1","NULL": 1 1 1 1 1 1 1 1 1 1 ...
## $ bmi : Factor w/ 401 levels "13.12","13.59",...: 401 401 401 401 401
401 401 401 401 401 ...
## $ vaccine : Factor w/ 7 levels "1","2","3","4",...: 4 4 4 4 4 4 4 4 4 4
...
## $ geo_mean : num 381 381 381 381 381 ...
## $ d_geo_mean : Factor w/ 40 levels "0","1","10","101",...: 2 2 2 2 2 2 2 2 2 2
2 ...
## $ vaccine_response : Factor w/ 3 levels "0","1","NULL": 1 1 1 1 1 1 1 1 1 1 ...
## $ mesurment_id : int 371121 371122 371123 371124 371125 371126 371127 371128
371129 371130 ...
## $ assay : int 4 4 4 4 4 4 4 4 4 4 ...
## $ name : Factor w/ 3283 levels "B cells","BASO %",...: 1 4 7 9 8 6 11 1
2 13 15 ...
## $ name_formatted : Factor w/ 3283 levels "B_cells","BASO_",...: 1 4 9 8 7 6 15 16
11 13 ...
## $ subset : Factor w/ 632 levels "Activated T: Lymph/CD3+",...: 36 30 103
102 101 45 32 98 62 66 ...
## $ units : Factor w/ 8 levels "% of Parent",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ data : num 34.4 1.45 5.85 1.55 4.46 3.79 65.6 62.6 11.4 1.66 ...
## $ statin_use : Factor w/ 3 levels "0","1","NULL": 1 1 1 1 1 1 1 1 1 1 ...
## $ flu_vaccination_history : Factor w/ 3 levels "0","1","NULL": 2 2 2 2 2 2 2 2 2 2 ...
## $ total_vaccines_received : Factor w/ 25 levels "0","1","10","11",...: 4 4 4 4 4 4 4 4 4 4
...
## $ vaccinated_1yr_prior : Factor w/ 4 levels "0","1","3","NULL": 2 2 2 2 2 2 2 2 2 2
...
## $ vaccine_type_1yr_prior : Factor w/ 4 levels "2","3","4","NULL": 1 1 1 1 1 1 1 1 1 1
...
## $ vaccinated_2yr_prior : Factor w/ 4 levels "0","1","3","NULL": 2 2 2 2 2 2 2 2 2 2
...
## $ vaccine_type_2yr_prior : Factor w/ 4 levels "2","3","4","NULL": 1 1 1 1 1 1 1 1 1 1
...
## $ vaccinated_3yr_prior : Factor w/ 4 levels "0","1","3","NULL": 2 2 2 2 2 2 2 2 2 2
...
## $ vaccine_type_3yr_prior : Factor w/ 4 levels "2","3","4","NULL": 1 1 1 1 1 1 1 1 1 1
...
## $ vaccinated_4yr_prior : Factor w/ 4 levels "0","1","3","NULL": 2 2 2 2 2 2 2 2 2 2
...
## $ vaccine_type_4yr_prior : Factor w/ 4 levels "2","3","4","NULL": 1 1 1 1 1 1 1 1 1 1
...
## $ vaccinated_5yr_prior : Factor w/ 4 levels "0","1","3","NULL": 2 2 2 2 2 2 2 2 2 2

```

```
...
## $ vaccine_type_5yr_prior      : Factor w/ 4 levels "2","3","4","NULL": 1 1 1 1 1 1 1 1 1 1
...
## $ influenza_infection_history: int  0 0 0 0 0 0 0 0 0 0 ...
## $ influenza_hospitalization  : int  0 0 0 0 0 0 0 0 0 0 ...
```

Count each variable

```
sapply(flu, function(x) length(unique(x)))
```

```
##           donor_id           study_id
##           740             8
##           gender           race
##           2             7
##           visit_id         visit_year
##           740             9
##           visit_day         visit_type_hai
##           3             2
##           visit_age         cmv_status
##           475             3
##           ebv_status        bmi
##           3             401
##           vaccine           geo_mean
##           7             76
##           d_geo_mean        vaccine_response
##           40             3
##           mesurment_id      assay
##           156118           13
##           name             name_formatted
##           3283             3283
##           subset           units
##           632             8
##           data             statin_use
##           5086             3
## flu_vaccination_history     total_vaccines_received
##           3             25
##           vaccinated_1yr_prior vaccine_type_1yr_prior
##           4             4
##           vaccinated_2yr_prior vaccine_type_2yr_prior
##           4             4
##           vaccinated_3yr_prior vaccine_type_3yr_prior
##           4             4
##           vaccinated_4yr_prior vaccine_type_4yr_prior
##           4             4
##           vaccinated_5yr_prior vaccine_type_5yr_prior
##           4             4
## influenza_infection_history influenza_hospitalization
##           2             2
```

Unique values of variable “subset”

```
table(flu$subset)
```

```
##
##
Activated T: Lymph/CD3+
##
26
##
Activated T: Lymph/CD3+/CD4+
##
26
##
Activated
T: Lymph/CD3+/CD4+/PD1+
##
26
##
Activated T: Lymph/CD
3+/CD4+/Q1: HLADR-CD38+
##
26
##
Activated T: Lymph/CD
3+/CD4+/Q2: HLADR+CD38+
##
26
##
Activated T: Lymph/CD
3+/CD4+/Q3: HLADR+CD38-
##
26
##
Activated
ated T: Lymph/CD3+/CD8+
##
26
##
Activated
T: Lymph/CD3+/CD8+/PD1+
##
26
##
Activated T: Lymph/CD
3+/CD8+/Q1: HLADR-CD38+
##
26
##
Activated T: Lymph/CD
3+/CD8+/Q2: HLADR+CD38+
##
26
##
Activated T: Lymph/CD
3+/CD8+/Q3: HLADR+CD38-
##
26
##
Activated T: Lymph/TCRgd+
##
26
##
B cell: Lymph/CD3-
##
26
```

##	B cel
1: Lymph/CD3-/CD19+CD20+	
##	
26	
##	B cell: Lymph/CD3
-/CD19+CD20+/CD24-/CD38+	
##	
26	
##	B cell: Lymph/CD3
-/CD19+CD20+/CD24+/CD38-	
##	
26	
##	B cell: Lymph/CD3
-/CD19+CD20+/CD24+/CD38+	
##	
26	
##	B cell: Lymph/CD3-/C
D19+CD20+/Q1: IgD-CD27+	
##	
26	
##	B cell: Lymph/CD3-/C
D19+CD20+/Q2: IgD+CD27+	
##	
26	
##	B cell: Lymph/CD3-/C
D19+CD20+/Q3: IgD+CD27-	
##	
26	
##	B cell: Lymph/CD3-/C
D19+CD20+/Q4: IgD-CD27-	
##	
26	
##	B
cell: Lymph/CD3-/CD20-	
##	
26	
##	B cell: Lymph
h/CD3-/CD20-/CD27+/CD38+	
##	
26	
##	
B cell: pSTAT1	
##	
1288	
##	
B cell: pSTAT3	
##	
1288	
##	
B cell: pSTAT5	
##	
1288	
##	
Basophil-%	

##	
50	
##	Basop
hil Absolute Count-K/uL	
##	
50	
##	
BDNF	
##	
162	
##	
CD123+HLADR-	
##	
186	
##	CD1
4-CD33-/CD3-/CD16+CD56+	
##	
308	
##	CD14-CD33
-/CD3-/CD16+CD56+/CD161+	
##	
308	
##	CD14-CD33
-/CD3-/CD16+CD56+/CD57+	
##	
186	
##	CD14-CD33
-/CD3-/CD16+CD56+/CD94+	
##	
308	
##	CD14-CD33
-/CD3-/CD16+CD56+/HLADR+	
##	
308	
##	CD1
4-CD33-/CD3-/CD19+CD20+	
##	
308	
##	CD14-CD33-/CD3
-/CD19+CD20+/CD24-CD38+	
##	
308	
##	CD14-CD33-/CD3
-/CD19+CD20+/CD24+CD38-	
##	
308	
##	CD14-CD33-/CD3
-/CD19+CD20+/CD24+CD38+	
##	
308	
##	CD14-CD33-/CD
3-/CD19+CD20+/IgD-CD27-	
##	
308	

##	CD14-CD33-/CD
3-/CD19+CD20+/IgD-CD27+	
##	
308	
##	CD14-CD33-/CD
3-/CD19+CD20+/IgD+CD27-	
##	
308	
##	CD14-CD33-/CD
3-/CD19+CD20+/IgD+CD27+	
##	
308	
##	CD14-CD33
-/CD3-/CD20-/CD27+CD38+	
##	
308	
##	CD14-CD33
-/CD3-/CD56bright CD16-	
##	
186	
##	
CD14-CD33-/CD3+	
##	
308	
##	CD14-CD
33-/CD3+/CD4-CD8+/CD57+	
##	
186	
##	CD14-CD
33-/CD3+/CD4-CD8+/ICOS+	
##	
186	
##	CD14-CD33-/CD3+/CD4-CD8+/Non-naive
CD8+/CXCR5-/CXCR3-CCR6-	
##	
186	
##	CD14-CD33-/CD3+/CD4-CD8+/Non-naive
CD8+/CXCR5-/CXCR3-CCR6+	
##	
186	
##	CD14-CD33-/CD3+/CD4-CD8+/Non-naive
CD8+/CXCR5-/CXCR3+CCR6-	
##	
186	
##	CD14-CD33-/CD3+/CD4-CD8
+ /Non-naive CD8+/CXCR5+	
##	
186	
##	CD14-CD33-/CD3+/CD4-CD8+/Non-naive
CD8+/CXCR5+/CXCR3-CCR6-	
##	
186	
##	CD14-CD33-/CD3+/CD4-CD8+/Non-naive
CD8+/CXCR5+/CXCR3-CCR6+	



##	
186	
##	CD14-CD33- /CD3+/CD4-CD8+/Non-naive
CD8+/CXCR5+/CXCR3+CCR6-	
##	
186	
##	CD14-C
D33- /CD3+/CD4-CD8+/PD1+	
##	
186	
##	
CD14-CD33- /CD3+/CD4+	
##	
308	
##	CD14-CD33
- /CD3+/CD4+/CCR7-CD45RA-	
##	
308	
##	CD14-CD33
- /CD3+/CD4+/CCR7-CD45RA+	
##	
308	
##	CD14-CD33
- /CD3+/CD4+/CCR7+CD45RA-	
##	
308	
##	CD14-CD33
- /CD3+/CD4+/CCR7+CD45RA+	
##	
308	
##	CD14
-CD33- /CD3+/CD4+/CD161+	
##	
308	
##	CD14-CD33- /C
D3+/CD4+/CD25hiCD127low	
##	
308	
##	CD14-CD33- /CD3+/CD4+/CD25h
iCD127low/CD161-CD45RA+	
##	
308	
##	CD14-CD33- /CD3+/CD4+/CD25h
iCD127low/CD161+CD45RA-	
##	
308	
##	CD14-CD33- /CD3+/CD4+/CD25h
iCD127low/CD161+CD45RA+	
##	
308	
##	CD1
4-CD33- /CD3+/CD4+/CD27+	
##	
308	

##	CD1
4-CD33-/CD3+/CD4+/CD28+	
##	
308	
##	CD14
-CD33-/CD3+/CD4+/CD85j+	
##	
308	
##	CD1
4-CD33-/CD3+/CD4+/CD94+	
##	
308	
##	CD14-CD33
-/CD3+/CD4+/HLADR-CD38+	
##	
308	
##	CD14-CD33
-/CD3+/CD4+/HLADR+CD38-	
##	
308	
##	CD14-CD33
-/CD3+/CD4+/HLADR+CD38+	
##	
308	
##	CD14-CD
33-/CD3+/CD4+CD8-/CD57+	
##	
186	
##	CD14-CD
33-/CD3+/CD4+CD8-/ICOS+	
##	
186	
##	CD14-CD33-/CD3+/CD4+CD8-/Non-naive
CD4+/CXCR5-/CXCR3-CCR6-	
##	
186	
##	CD14-CD33-/CD3+/CD4+CD8-/Non-naive
CD4+/CXCR5-/CXCR3-CCR6+	
##	
186	
##	CD14-CD33-/CD3+/CD4+CD8-/Non-naive
CD4+/CXCR5-/CXCR3+CCR6-	
##	
186	
##	CD14-CD33-/CD3+/CD4+CD8
-/Non-naive CD4+/CXCR5+	
##	
186	
##	CD14-CD33-/CD3+/CD4+CD8-/Non-naive
CD4+/CXCR5+/CXCR3-CCR6-	
##	
186	
##	CD14-CD33-/CD3+/CD4+CD8-/Non-naive
CD4+/CXCR5+/CXCR3-CCR6+	

##		
186		
##	CD14-CD33- /CD3+/CD4+CD8- /Non-naive	
CD4+/CXCR5+/CXCR3+CCR6-		
##		
186		
##		CD14-C
D33- /CD3+/CD4+CD8- /PD1+		
##		
186		
##		
CD14-CD33- /CD3+/CD56+		
##		
308		
##		
CD14-CD33- /CD3+/CD8+		
##		
308		
##		CD14-CD33
- /CD3+/CD8+/CCR7-CD45RA-		
##		
308		
##		CD14-CD33
- /CD3+/CD8+/CCR7-CD45RA+		
##		
308		
##		CD14-CD33
- /CD3+/CD8+/CCR7+CD45RA-		
##		
308		
##		CD14-CD33
- /CD3+/CD8+/CCR7+CD45RA+		
##		
308		
##		CD14
-CD33- /CD3+/CD8+/CD161+		
##		
308		
##		CD1
4-CD33- /CD3+/CD8+/CD27+		
##		
308		
##		CD1
4-CD33- /CD3+/CD8+/CD28+		
##		
308		
##		CD14
-CD33- /CD3+/CD8+/CD85j+		
##		
308		
##		CD1
4-CD33- /CD3+/CD8+/CD94+		
##		
308		

##	CD14-CD33
- /CD3+ /CD8+ /HLADR-CD38+	
##	
308	
##	CD14-CD33
- /CD3+ /CD8+ /HLADR+CD38-	
##	
308	
##	CD14-CD33
- /CD3+ /CD8+ /HLADR+CD38+	
##	
308	
##	CD14
-CD33- /CD3+CD56+ /CD161+	
##	
186	
##	
CD14-CD33- /TCRgd+	
##	
308	
##	
CD14+CD33+	
##	
308	
##	
CD14+CD33+ /CD14-CD16+	
##	
186	
##	
CD14+CD33+ /CD14+CD16+	
##	
186	
##	
CD14+CD33+ /CD16+	
##	
308	
##	
CD4+: pSTAT1	
##	
1288	
##	
CD4+: pSTAT3	
##	
1288	
##	
CD4+: pSTAT5	
##	
1288	
##	
CD4+CD45RA- : pSTAT1	
##	
1050	
##	
CD4+CD45RA- : pSTAT3	

```
##
1050
##
CD4+CD45RA-: pSTAT5
##
1050
##
CD4+CD45RA+: pSTAT1
##
1050
##
CD4+CD45RA+: pSTAT3
##
1050
##
CD4+CD45RA+: pSTAT5
##
1050
##
CD40L
##
650
##
CD45+ CD66+
##
192
##
CD45+ CD66+: IkBtot
##
192
##
CD45+ CD66+: Ki67
##
192
##
CD45+ CD66+: pCREB
##
192
##
CD45+ CD66+: pErk1_2
##
192
##
CD45+ CD66+: pp38
##
192
##
CD45+ CD66+: pPLCg2
##
192
##
CD45+ CD66+: pSTAT1
##
192
```

```

##
CD45+ CD66+: pSTAT3
##
192
##
CD45+ CD66+: pSTAT5
##
192
##
CD45++ CD66low/CD3-/CD123
-/BDCA-3 -/CD27++CD38++
##
192
##
CD45++ CD66low/CD3-/CD123-/BDCA-3 -/CD27++
CD38++/Q1: IgA- , Ki67+
##
192
##
CD45++ CD66low/CD3-/CD123-/BDCA-3 -/CD27++
CD38++/Q2: IgA+ , Ki67+
##
192
##
CD45++ CD66low/CD3-/CD123-/BDCA-3 -/CD27++
CD38++/Q3: IgA+ , Ki67-
##
192
##
CD45++ CD66low/CD3-/CD123-/BDCA-3 -/CD27++
CD38++/Q4: IgA- , Ki67-
##
192
##
CD45++ CD66low/CD3-/CD123-/BDCA-3
-/CD27++CD38++: IkBtot
##
187
##
CD45++ CD66low/CD3-/CD123-/BDCA
-3 -/CD27++CD38++: Ki67
##
187
##
CD45++ CD66low/CD3-/CD123-/BDCA-
3 -/CD27++CD38++: pCREB
##
187
##
CD45++ CD66low/CD3-/CD123-/BDCA-3
-/CD27++CD38++: pErk1_2
##
187
##
CD45++ CD66low/CD3-/CD123-/BDCA
-3 -/CD27++CD38++: pp38
##
187
##
CD45++ CD66low/CD3-/CD123-/BDCA-3
-/CD27++CD38++: pPLCg2
##
187
##
CD45++ CD66low/CD3-/CD123-/BDCA-3
-/CD27++CD38++: pSTAT1

```

```
##
187
##
- /CD27++CD38++: pSTAT3
##
187
##
- /CD27++CD38++: pSTAT5
##
187
##
D27++CD38++/CD7+ HLADR-
##
192
##
CD7+ HLADR- /CD56- CD16+
##
192
##
DR- /CD56- CD16+: IkBtot
##
187
##
LADR- /CD56- CD16+: Ki67
##
187
##
ADR- /CD56- CD16+: pCREB
##
187
##
R- /CD56- CD16+: pErk1_2
##
187
##
LADR- /CD56- CD16+: pp38
##
187
##
DR- /CD56- CD16+: pPLCg2
##
187
##
DR- /CD56- CD16+: pSTAT1
##
187
##
DR- /CD56- CD16+: pSTAT3
##
187
##
DR- /CD56- CD16+: pSTAT5
##
187
```

CD45++ CD66low/CD3- /CD123- /BDCA-3

CD45++ CD66low/CD3- /CD123- /BDCA-3

CD45++ CD66low/CD3- /CD123- /BDCA-3 - /Not C

CD45++ CD66low/CD3- /CD123- /BDCA-3 - /Not CD27++CD38++ /

CD45++ CD66low/CD3- /CD123- /BDCA-3 - /Not CD27++CD38++ /CD7+ HLA

CD45++ CD66low/CD3- /CD123- /BDCA-3 - /Not CD27++CD38++ /CD7+ H

CD45++ CD66low/CD3- /CD123- /BDCA-3 - /Not CD27++CD38++ /CD7+ HL

CD45++ CD66low/CD3- /CD123- /BDCA-3 - /Not CD27++CD38++ /CD7+ HLAD

CD45++ CD66low/CD3- /CD123- /BDCA-3 - /Not CD27++CD38++ /CD7+ H

CD45++ CD66low/CD3- /CD123- /BDCA-3 - /Not CD27++CD38++ /CD7+ HLA

CD45++ CD66low/CD3- /CD123- /BDCA-3 - /Not CD27++CD38++ /CD7+ HLA

CD45++ CD66low/CD3- /CD123- /BDCA-3 - /Not CD27++CD38++ /CD7+ HLA

```

##                                CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/CD7+
HLADR-/CD56bright CD16+
##
192
##                                CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/CD7+ HLADR-/C
D56bright CD16+: IkBtot
##
187
##                                CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/CD7+ HLADR
-/CD56bright CD16+: Ki67
##
187
##                                CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/CD7+ HLADR-/
CD56bright CD16+: pCREB
##
187
##                                CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/CD7+ HLADR-/CD
56bright CD16+: pErk1_2
##
187
##                                CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/CD7+ HLADR
-/CD56bright CD16+: pp38
##
187
##                                CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/CD7+ HLADR-/C
D56bright CD16+: pPLCg2
##
187
##                                CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/CD7+ HLADR-/C
D56bright CD16+: pSTAT1
##
187
##                                CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/CD7+ HLADR-/C
D56bright CD16+: pSTAT3
##
187
##                                CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/CD7+ HLADR-/C
D56bright CD16+: pSTAT5
##
187
##                                CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/CD
7+ HLADR-/CD56int CD16-
##
192
##                                CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/CD7+ HLADR
-/CD56int CD16-: IkBtot
##
187
##                                CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/CD7+ HLA
DR-/CD56int CD16-: Ki67
##
187
##                                CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/CD7+ HLAD
R-/CD56int CD16-: pCREB

```



```
##
187
##
CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/CD7+ HLADR
-/CD56int CD16-: pErk1_2
##
187
##
CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/CD7+ HLA
DR-/CD56int CD16-: pp38
##
187
##
CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/CD7+ HLADR
-/CD56int CD16-: pPLCg2
##
187
##
CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/CD7+ HLADR
-/CD56int CD16-: pSTAT1
##
187
##
CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/CD7+ HLADR
-/CD56int CD16-: pSTAT3
##
187
##
CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/CD7+ HLADR
-/CD56int CD16-: pSTAT5
##
187
##
CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/CD
7+ HLADR-/CD56int CD16+
##
192
##
CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/CD7+ HLADR
-/CD56int CD16+: IkBtot
##
187
##
CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/CD7+ HLA
DR-/CD56int CD16+: Ki67
##
187
##
CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/CD7+ HLAD
R-/CD56int CD16+: pCREB
##
187
##
CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/CD7+ HLADR
-/CD56int CD16+: pErk1_2
##
187
##
CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/CD7+ HLA
DR-/CD56int CD16+: pp38
##
187
##
CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/CD7+ HLADR
-/CD56int CD16+: pPLCg2
##
187
```

```

## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/CD7+ HLADR
-/CD56int CD16+: pSTAT1
##
187
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/CD7+ HLADR
-/CD56int CD16+: pSTAT3
##
187
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/CD7+ HLADR
-/CD56int CD16+: pSTAT5
##
187
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD3
8++/CD7+ HLADR-: IkBtot
##
187
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++C
D38++/CD7+ HLADR-: Ki67
##
187
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD
38++/CD7+ HLADR-: pCREB
##
187
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38
++/CD7+ HLADR-: pErk1_2
##
187
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++C
D38++/CD7+ HLADR-: pp38
##
187
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD3
8++/CD7+ HLADR-: pPLCg2
##
187
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD3
8++/CD7+ HLADR-: pSTAT1
##
187
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD3
8++/CD7+ HLADR-: pSTAT3
##
187
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD3
8++/CD7+ HLADR-: pSTAT5
##
187
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38+
+/HLADR+ CD7-/CD19+CD20+
##
192
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+
CD20+/CD24high CD38high

```

```
##
192
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/CD
24high CD38high: IkBtot
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/
CD24high CD38high: Ki67
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/C
D24high CD38high: pCREB
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/CD2
4high CD38high: pErk1_2
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/
CD24high CD38high: pp38
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/CD
24high CD38high: pPLCg2
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/CD
24high CD38high: pSTAT1
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/CD
24high CD38high: pSTAT3
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/CD
24high CD38high: pSTAT5
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD1
9+CD20+/IgA+ CD19+CD20+
##
192
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/
IgA+ CD19+CD20+: IkBtot
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20
+/IgA+ CD19+CD20+: Ki67
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20
+/IgA+ CD19+CD20+: pCREB
##
187
```

```
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/I
gA+ CD19+CD20+: pErk1_2
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20
+/IgA+ CD19+CD20+: pp38
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/
IgA+ CD19+CD20+: pPLCg2
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/
IgA+ CD19+CD20+: pSTAT1
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/
IgA+ CD19+CD20+: pSTAT3
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/
IgA+ CD19+CD20+: pSTAT5
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19
+CD20+/Q1: IgD- , CD27+
##
192
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/Q
1: IgD- , CD27+: IkBtot
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20
+/Q1: IgD- , CD27+: Ki67
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/
Q1: IgD- , CD27+: pCREB
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/Q
1: IgD- , CD27+: pErk1_2
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20
+/Q1: IgD- , CD27+: pp38
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/Q
1: IgD- , CD27+: pPLCg2
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/Q
1: IgD- , CD27+: pSTAT1
```

```
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/Q
1: IgD- , CD27+: pSTAT3
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/Q
1: IgD- , CD27+: pSTAT5
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19
+CD20+/Q2: IgD+ , CD27+
##
192
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/Q
2: IgD+ , CD27+: IkBtot
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20
+/Q2: IgD+ , CD27+: Ki67
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/
Q2: IgD+ , CD27+: pCREB
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/Q
2: IgD+ , CD27+: pErk1_2
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20
+/Q2: IgD+ , CD27+: pp38
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/Q
2: IgD+ , CD27+: pPLCg2
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/Q
2: IgD+ , CD27+: pSTAT1
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/Q
2: IgD+ , CD27+: pSTAT3
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/Q
2: IgD+ , CD27+: pSTAT5
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19
+CD20+/Q3: IgD+ , CD27-
##
192
```

```
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/Q
3: IgD+ , CD27-: IkBtot
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20
+/Q3: IgD+ , CD27-: Ki67
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/
Q3: IgD+ , CD27-: pCREB
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/Q
3: IgD+ , CD27-: pErk1_2
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20
+/Q3: IgD+ , CD27-: pp38
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/Q
3: IgD+ , CD27-: pPLCg2
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/Q
3: IgD+ , CD27-: pSTAT1
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/Q
3: IgD+ , CD27-: pSTAT3
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/Q
3: IgD+ , CD27-: pSTAT5
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19
+CD20+/Q4: IgD- , CD27-
##
192
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/Q
4: IgD- , CD27-: IkBtot
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20
+/Q4: IgD- , CD27-: Ki67
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/
Q4: IgD- , CD27-: pCREB
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/Q
4: IgD- , CD27-: pErk1_2
```

```
##
187
##           CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20
+/Q4: IgD- , CD27-: pp38
##
187
##           CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/Q
4: IgD- , CD27-: pPLCg2
##
187
##           CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/Q
4: IgD- , CD27-: pSTAT1
##
187
##           CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/Q
4: IgD- , CD27-: pSTAT3
##
187
##           CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/CD19+CD20+/Q
4: IgD- , CD27-: pSTAT5
##
187
##           CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+
CD7-/CD19+CD20+: IkBtot
##
187
##           CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR
+ CD7-/CD19+CD20+: Ki67
##
187
##           CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+
CD7-/CD19+CD20+: pCREB
##
187
##           CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ C
D7-/CD19+CD20+: pErk1_2
##
187
##           CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR
+ CD7-/CD19+CD20+: pp38
##
187
##           CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+
CD7-/CD19+CD20+: pPLCg2
##
187
##           CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+
CD7-/CD19+CD20+: pSTAT1
##
187
##           CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+
CD7-/CD19+CD20+: pSTAT3
##
187
```

```
##                                CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+
CD7-/CD19+CD20+: pSTAT5
##
187
##                                CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non
CD19+CD20+/CD14+ Monos
##
192
##                                CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD
20+/CD14+ Monos: IkBtot
##
187
##                                CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+
CD20+/CD14+ Monos: Ki67
##
187
##                                CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+C
D20+/CD14+ Monos: pCREB
##
187
##                                CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD2
0+/CD14+ Monos: pErk1_2
##
187
##                                CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+
CD20+/CD14+ Monos: pp38
##
187
##                                CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD
20+/CD14+ Monos: pPLCg2
##
187
##                                CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD
20+/CD14+ Monos: pSTAT1
##
187
##                                CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD
20+/CD14+ Monos: pSTAT3
##
187
##                                CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD
20+/CD14+ Monos: pSTAT5
##
187
##                                CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non
CD19+CD20+/CD16+ Monos
##
192
##                                CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD
20+/CD16+ Monos: IkBtot
##
187
##                                CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+
CD20+/CD16+ Monos: Ki67
```



```
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+C
D20+/CD16+ Monos: pCREB
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD2
0+/CD16+ Monos: pErk1_2
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+
CD20+/CD16+ Monos: pp38
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD
20+/CD16+ Monos: pPLCg2
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD
20+/CD16+ Monos: pSTAT1
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD
20+/CD16+ Monos: pSTAT3
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD
20+/CD16+ Monos: pSTAT5
##
187
##          CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD20+/non
Monos/CD11c+ HLA-DRhigh
##
192
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD20+/non Monos/CD
11c+ HLA-DRhigh: IkBtot
##
187
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD20+/non Monos/
CD11c+ HLA-DRhigh: Ki67
##
187
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD20+/non Monos/C
D11c+ HLA-DRhigh: pCREB
##
187
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD20+/non Monos/CD1
1c+ HLA-DRhigh: pErk1_2
##
187
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD20+/non Monos/
CD11c+ HLA-DRhigh: pp38
##
187
```

```
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD20+/non Monos/CD
11c+ HLA-DRhigh: pPLCg2
##
187
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD20+/non Monos/CD
11c+ HLA-DRhigh: pSTAT1
##
187
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD20+/non Monos/CD
11c+ HLA-DRhigh: pSTAT3
##
187
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD20+/non Monos/CD
11c+ HLA-DRhigh: pSTAT5
##
187
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD20+/non Monos/Syk+ CD11c+
+/non Monos/Syk+ CD11c+
##
192
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD20+/non Mo
nos/Syk+ CD11c+: IkBtot
##
187
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD20+/non
Monos/Syk+ CD11c+: Ki67
##
187
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD20+/non M
onos/Syk+ CD11c+: pCREB
##
187
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD20+/non Mon
os/Syk+ CD11c+: pErk1_2
##
187
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD20+/non
Monos/Syk+ CD11c+: pp38
##
187
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD20+/non Mo
nos/Syk+ CD11c+: pPLCg2
##
187
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD20+/non Mo
nos/Syk+ CD11c+: pSTAT1
##
187
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD20+/non Mo
nos/Syk+ CD11c+: pSTAT3
##
187
## CD45++ CD66low/CD3-/CD123-/BDCA-3 -/Not CD27++CD38++/HLADR+ CD7-/non CD19+CD20+/non Mo
nos/Syk+ CD11c+: pSTAT5
```

```

##
187
##
w/CD3-/CD123-/type 2 mDC
##
192
##
123-/type 2 mDC: IkBtot
##
178
##
CD123-/type 2 mDC: Ki67
##
178
##
D123-/type 2 mDC: pCREB
##
178
##
23-/type 2 mDC: pErk1_2
##
178
##
CD123-/type 2 mDC: pp38
##
178
##
123-/type 2 mDC: pPLCg2
##
178
##
123-/type 2 mDC: pSTAT1
##
178
##
123-/type 2 mDC: pSTAT3
##
178
##
123-/type 2 mDC: pSTAT5
##
178
##
66low/CD3-/CD123+HLADR-
##
192
##
D3-/CD123+HLADR- IkBtot
##
187
##
CD3-/CD123+HLADR-: Ki67
##
187

```

CD45++ CD66lo

CD45++ CD66low/CD3-/CD

CD45++ CD66low/CD3-/

CD45++ CD66low/CD3-/C

CD45++ CD66low/CD3-/CD1

CD45++ CD66low/CD3-/

CD45++ CD66low/CD3-/CD

CD45++ CD66low/CD3-/CD

CD45++ CD66low/CD3-/CD

CD45++ CD66low/CD3-/CD

CD45++ CD

CD45++ CD66low/C

CD45++ CD66low/

##	CD45++ CD66low/C
D3-/CD123+HLADR-: pCREB	
##	
187	
##	CD45++ CD66low/CD3
-/CD123+HLADR-: pErk1_2	
##	
187	
##	CD45++ CD66low/
CD3-/CD123+HLADR-: pp38	
##	
187	
##	CD45++ CD66low/CD
3-/CD123+HLADR-: pPLCg2	
##	
187	
##	CD45++ CD66low/CD
3-/CD123+HLADR-: pSTAT1	
##	
187	
##	CD45++ CD66low/CD
3-/CD123+HLADR-: pSTAT3	
##	
187	
##	CD45++ CD66low/CD
3-/CD123+HLADR-: pSTAT5	
##	
187	
##	
CD45++ CD66low/CD3-/pDC	
##	
192	
##	CD45++ C
D66low/CD3-/pDC: IkBtot	
##	
187	
##	CD45++
CD66low/CD3-/pDC: Ki67	
##	
187	
##	CD45++
CD66low/CD3-/pDC: pCREB	
##	
187	
##	CD45++ CD
66low/CD3-/pDC: pErk1_2	
##	
187	
##	CD45++
CD66low/CD3-/pDC: pp38	
##	
187	
##	CD45++ C
D66low/CD3-/pDC: pPLCg2	

##		
187		
##		CD45++ C
D66low/CD3-/pDC: pSTAT1		
##		
187		
##		CD45++ C
D66low/CD3-/pDC: pSTAT3		
##		
187		
##		CD45++ C
D66low/CD3-/pDC: pSTAT5		
##		
187		
##		
CD45++ CD66low/CD3+		
##		
192		
##		C
D45++ CD66low/CD3+/CD4+		
##		
192		
##		CD45++
CD66low/CD3+/CD4+/CD25+		
##		
192		
##		CD45++ CD66low/
CD3+/CD4+/CD25+: IkBtot		
##		
187		
##		CD45++ CD66lo
w/CD3+/CD4+/CD25+: Ki67		
##		
187		
##		CD45++ CD66lo
w/CD3+/CD4+/CD25+: pCREB		
##		
187		
##		CD45++ CD66low/C
D3+/CD4+/CD25+: pErk1_2		
##		
187		
##		CD45++ CD66lo
w/CD3+/CD4+/CD25+: pp38		
##		
187		
##		CD45++ CD66low/
CD3+/CD4+/CD25+: pPLCg2		
##		
187		
##		CD45++ CD66low/
CD3+/CD4+/CD25+: pSTAT1		
##		
187		

##	CD45++ CD66low/
CD3+/CD4+/CD25+: pSTAT3	
##	
187	
##	CD45++ CD66low/
CD3+/CD4+/CD25+: pSTAT5	
##	
187	
##	CD45++ CD66lo
w/CD3+/CD4+/CD38+ HLADR+	
##	
192	
##	CD45++ CD66low/CD3+/CD
4+/CD38+ HLADR+: IkBtot	
##	
186	
##	CD45++ CD66low/CD3+/
CD4+/CD38+ HLADR+: Ki67	
##	
186	
##	CD45++ CD66low/CD3+/C
D4+/CD38+ HLADR+: pCREB	
##	
186	
##	CD45++ CD66low/CD3+/CD4
+ /CD38+ HLADR+: pErk1_2	
##	
186	
##	CD45++ CD66low/CD3+/
CD4+/CD38+ HLADR+: pp38	
##	
186	
##	CD45++ CD66low/CD3+/CD
4+/CD38+ HLADR+: pPLCg2	
##	
186	
##	CD45++ CD66low/CD3+/CD
4+/CD38+ HLADR+: pSTAT1	
##	
186	
##	CD45++ CD66low/CD3+/CD
4+/CD38+ HLADR+: pSTAT3	
##	
186	
##	CD45++ CD66low/CD3+/CD
4+/CD38+ HLADR+: pSTAT5	
##	
186	
##	CD45++ CD66low/CD3+/C
D4+/Q1: CD45RA- , CD27+	
##	
192	
##	CD45++ CD66low/CD3+/CD4+/Q1:
CD45RA- , CD27+: IkBtot	

```

##
187
##
1: CD45RA- , CD27+: Ki67
##
187
##
CD45RA- , CD27+: pCREB
##
187
##
D45RA- , CD27+: pErk1_2
##
187
##
1: CD45RA- , CD27+: pp38
##
187
##
CD45RA- , CD27+: pPLCg2
##
187
##
CD45RA- , CD27+: pSTAT1
##
187
##
CD45RA- , CD27+: pSTAT3
##
187
##
CD45RA- , CD27+: pSTAT5
##
187
##
D4+/Q2: CD45RA+ , CD27+
##
192
##
CD45RA+ , CD27+: IkBtot
##
187
##
2: CD45RA+ , CD27+: Ki67
##
187
##
CD45RA+ , CD27+: pCREB
##
187
##
D45RA+ , CD27+: pErk1_2
##
187

```

CD45++ CD66low/CD3+/CD4+/Q

CD45++ CD66low/CD3+/CD4+/Q1:

CD45++ CD66low/CD3+/CD4+/Q1: C

CD45++ CD66low/CD3+/CD4+/Q

CD45++ CD66low/CD3+/CD4+/Q1:

CD45++ CD66low/CD3+/CD4+/Q1:

CD45++ CD66low/CD3+/CD4+/Q1:

CD45++ CD66low/CD3+/CD4+/Q1:

CD45++ CD66low/CD3+/CD4+/Q1:

CD45++ CD66low/CD3+/C

CD45++ CD66low/CD3+/CD4+/Q2:

CD45++ CD66low/CD3+/CD4+/Q

CD45++ CD66low/CD3+/CD4+/Q2:

CD45++ CD66low/CD3+/CD4+/Q2: C

##	CD45++ CD66low/CD3+/CD4+/Q
2: CD45RA+ , CD27+: pp38	
##	
187	
##	CD45++ CD66low/CD3+/CD4+/Q2:
CD45RA+ , CD27+: pPLCg2	
##	
187	
##	CD45++ CD66low/CD3+/CD4+/Q2:
CD45RA+ , CD27+: pSTAT1	
##	
187	
##	CD45++ CD66low/CD3+/CD4+/Q2:
CD45RA+ , CD27+: pSTAT3	
##	
187	
##	CD45++ CD66low/CD3+/CD4+/Q2:
CD45RA+ , CD27+: pSTAT5	
##	
187	
##	CD45++ CD66low/CD3+/C
D4+/Q3: CD45RA+ , CD27-	
##	
192	
##	CD45++ CD66low/CD3+/CD4+/Q3:
CD45RA+ , CD27-: IkBtot	
##	
187	
##	CD45++ CD66low/CD3+/CD4+/Q
3: CD45RA+ , CD27-: Ki67	
##	
187	
##	CD45++ CD66low/CD3+/CD4+/Q3:
CD45RA+ , CD27-: pCREB	
##	
187	
##	CD45++ CD66low/CD3+/CD4+/Q3: C
D45RA+ , CD27-: pErk1_2	
##	
187	
##	CD45++ CD66low/CD3+/CD4+/Q
3: CD45RA+ , CD27-: pp38	
##	
187	
##	CD45++ CD66low/CD3+/CD4+/Q3:
CD45RA+ , CD27-: pPLCg2	
##	
187	
##	CD45++ CD66low/CD3+/CD4+/Q3:
CD45RA+ , CD27-: pSTAT1	
##	
187	
##	CD45++ CD66low/CD3+/CD4+/Q3:
CD45RA+ , CD27-: pSTAT3	



```

##
187
##
CD45RA+ , CD27-: pSTAT5
##
187
##
CD45++ CD66low/CD3+/CD4+/Q3:
D4+/Q4: CD45RA- , CD27-
##
192
##
CD45++ CD66low/CD3+/CD4+/Q4:
CD45RA- , CD27-: IkBtot
##
182
##
CD45++ CD66low/CD3+/CD4+/Q
4: CD45RA- , CD27-: Ki67
##
182
##
CD45++ CD66low/CD3+/CD4+/Q4:
CD45RA- , CD27-: pCREB
##
182
##
CD45++ CD66low/CD3+/CD4+/Q4: C
D45RA- , CD27-: pErk1_2
##
182
##
CD45++ CD66low/CD3+/CD4+/Q
4: CD45RA- , CD27-: pp38
##
182
##
CD45++ CD66low/CD3+/CD4+/Q4:
CD45RA- , CD27-: pPLCg2
##
182
##
CD45++ CD66low/CD3+/CD4+/Q4:
CD45RA- , CD27-: pSTAT1
##
182
##
CD45++ CD66low/CD3+/CD4+/Q4:
CD45RA- , CD27-: pSTAT3
##
182
##
CD45++ CD66low/CD3+/CD4+/Q4:
CD45RA- , CD27-: pSTAT5
##
182
##
CD45++ CD
66low/CD3+/CD4+: IkBtot
##
192
##
CD45++
CD66low/CD3+/CD4+: Ki67
##
192

```

##	CD45++ C
D66low/CD3+/CD4+: pCREB	
##	
192	
##	CD45++ CD6
6low/CD3+/CD4+: pErk1_2	
##	
192	
##	CD45++
CD66low/CD3+/CD4+: pp38	
##	
192	
##	CD45++ CD
66low/CD3+/CD4+: pPLCg2	
##	
192	
##	CD45++ CD
66low/CD3+/CD4+: pSTAT1	
##	
192	
##	CD45++ CD
66low/CD3+/CD4+: pSTAT3	
##	
192	
##	CD45++ CD
66low/CD3+/CD4+: pSTAT5	
##	
192	
##	C
D45++ CD66low/CD3+/CD8+	
##	
192	
##	CD45++ CD66lo
w/CD3+/CD8+/CD38+ HLADR+	
##	
192	
##	CD45++ CD66low/CD3+/CD
8+/CD38+ HLADR+: IkBtot	
##	
187	
##	CD45++ CD66low/CD3+/
CD8+/CD38+ HLADR+: Ki67	
##	
187	
##	CD45++ CD66low/CD3+/C
D8+/CD38+ HLADR+: pCREB	
##	
187	
##	CD45++ CD66low/CD3+/CD8
+/CD38+ HLADR+: pErk1_2	
##	
187	
##	CD45++ CD66low/CD3+/
CD8+/CD38+ HLADR+: pp38	

```

##
187
##
8+/CD38+ HLADR+: pPLCg2
##
187
##
8+/CD38+ HLADR+: pSTAT1
##
187
##
8+/CD38+ HLADR+: pSTAT3
##
187
##
8+/CD38+ HLADR+: pSTAT5
##
187
##
D8+/Q1: CD45RA- , CD27+
##
192
##
CD45RA- , CD27+: IkBtot
##
178
##
1: CD45RA- , CD27+: Ki67
##
178
##
CD45RA- , CD27+: pCREB
##
178
##
D45RA- , CD27+: pErk1_2
##
178
##
1: CD45RA- , CD27+: pp38
##
178
##
CD45RA- , CD27+: pPLCg2
##
178
##
CD45RA- , CD27+: pSTAT1
##
178
##
CD45RA- , CD27+: pSTAT3
##
178

```

CD45++ CD66low/CD3+/CD

CD45++ CD66low/CD3+/CD

CD45++ CD66low/CD3+/CD

CD45++ CD66low/CD3+/CD

CD45++ CD66low/CD3+/C

CD45++ CD66low/CD3+/CD8+/Q1:

CD45++ CD66low/CD3+/CD8+/Q

CD45++ CD66low/CD3+/CD8+/Q1:

CD45++ CD66low/CD3+/CD8+/Q1: C

CD45++ CD66low/CD3+/CD8+/Q

CD45++ CD66low/CD3+/CD8+/Q1:

CD45++ CD66low/CD3+/CD8+/Q1:

CD45++ CD66low/CD3+/CD8+/Q1:

```

##
CD45RA- , CD27+: pSTAT5
##
178
##
CD45++ CD66low/CD3+/CD8+/Q1:
D8+/Q2: CD45RA+ , CD27+
##
192
##
CD45++ CD66low/CD3+/CD8+/Q2:
CD45RA+ , CD27+: IkBtot
##
187
##
CD45++ CD66low/CD3+/CD8+/Q
2: CD45RA+ , CD27+: Ki67
##
187
##
CD45++ CD66low/CD3+/CD8+/Q2:
CD45RA+ , CD27+: pCREB
##
187
##
CD45++ CD66low/CD3+/CD8+/Q2: C
D45RA+ , CD27+: pErk1_2
##
187
##
CD45++ CD66low/CD3+/CD8+/Q
2: CD45RA+ , CD27+: pp38
##
187
##
CD45++ CD66low/CD3+/CD8+/Q2:
CD45RA+ , CD27+: pPLCg2
##
187
##
CD45++ CD66low/CD3+/CD8+/Q2:
CD45RA+ , CD27+: pSTAT1
##
187
##
CD45++ CD66low/CD3+/CD8+/Q2:
CD45RA+ , CD27+: pSTAT3
##
187
##
CD45++ CD66low/CD3+/CD8+/Q2:
CD45RA+ , CD27+: pSTAT5
##
187
##
CD45++ CD66low/CD3+/C
D8+/Q3: CD45RA+ , CD27-
##
192
##
CD45++ CD66low/CD3+/CD8+/Q3:
CD45RA+ , CD27-: IkBtot
##
187
##
CD45++ CD66low/CD3+/CD8+/Q
3: CD45RA+ , CD27-: Ki67

```

```

##
187
##
CD45RA+ , CD27-: pCREB
##
187
##
D45RA+ , CD27-: pErk1_2
##
187
##
3: CD45RA+ , CD27-: pp38
##
187
##
CD45RA+ , CD27-: pPLCg2
##
187
##
CD45RA+ , CD27-: pSTAT1
##
187
##
CD45RA+ , CD27-: pSTAT3
##
187
##
CD45RA+ , CD27-: pSTAT5
##
187
##
D8+/Q4: CD45RA- , CD27-
##
192
##
CD45RA- , CD27-: IkBtot
##
164
##
4: CD45RA- , CD27-: Ki67
##
164
##
CD45RA- , CD27-: pCREB
##
164
##
D45RA- , CD27-: pErk1_2
##
164
##
4: CD45RA- , CD27-: pp38
##
164

```

CD45++ CD66low/CD3+/CD8+/Q3:

CD45++ CD66low/CD3+/CD8+/Q3: C

CD45++ CD66low/CD3+/CD8+/Q

CD45++ CD66low/CD3+/CD8+/Q3:

CD45++ CD66low/CD3+/CD8+/Q3:

CD45++ CD66low/CD3+/CD8+/Q3:

CD45++ CD66low/CD3+/CD8+/Q3:

CD45++ CD66low/CD3+/CD8+/Q3:

CD45++ CD66low/CD3+/C

CD45++ CD66low/CD3+/CD8+/Q4:

CD45++ CD66low/CD3+/CD8+/Q

CD45++ CD66low/CD3+/CD8+/Q4:

CD45++ CD66low/CD3+/CD8+/Q4: C

CD45++ CD66low/CD3+/CD8+/Q

##	CD45RA- , CD27-: pPLCg2	CD45++ CD66low/CD3+/CD8+/Q4:
##		
164		
##	CD45RA- , CD27-: pSTAT1	CD45++ CD66low/CD3+/CD8+/Q4:
##		
164		
##	CD45RA- , CD27-: pSTAT3	CD45++ CD66low/CD3+/CD8+/Q4:
##		
164		
##	CD45RA- , CD27-: pSTAT5	CD45++ CD66low/CD3+/CD8+/Q4:
##		
164		
##		CD45++ CD
	66low/CD3+/CD8+: IkBtot	
##		
192		
##		CD45++
	CD66low/CD3+/CD8+: Ki67	
##		
192		
##		CD45++ C
	D66low/CD3+/CD8+: pCREB	
##		
192		
##		CD45++ CD6
	6low/CD3+/CD8+: pErk1_2	
##		
192		
##		CD45++
	CD66low/CD3+/CD8+: pp38	
##		
192		
##		CD45++ CD
	66low/CD3+/CD8+: pPLCg2	
##		
192		
##		CD45++ CD
	66low/CD3+/CD8+: pSTAT1	
##		
192		
##		CD45++ CD
	66low/CD3+/CD8+: pSTAT3	
##		
192		
##		CD45++ CD
	66low/CD3+/CD8+: pSTAT5	
##		
192		
##		
	CD8+: pSTAT1	

##	
1288	
##	
CD8+: pSTAT3	
##	
1288	
##	
CD8+: pSTAT5	
##	
1288	
##	
CD8+CD45RA-: pSTAT1	
##	
1050	
##	
CD8+CD45RA-: pSTAT3	
##	
1050	
##	
CD8+CD45RA-: pSTAT5	
##	
1050	
##	
CD8+CD45RA+: pSTAT1	
##	
1050	
##	
CD8+CD45RA+: pSTAT3	
##	
1050	
##	
CD8+CD45RA+: pSTAT5	
##	
1050	
##	
FM0: Lymph/CD16+/CD56+	
##	
26	
##	
CXCR3 FM0: Lymph/CD3-	
##	
26	
##	
0: Lymph/CD3-/CD19+CD20+	
##	
26	
##	
CXCR3 FM0: Lymph/CD3+	
##	
26	
##	
R3 FM0: Lymph/CD3+/CD4+	
##	
26	

CXCR3

CXCR3 FM

CXC

##	CXC
R3 FM0: Lymph/CD3+/CD8+	
##	
26	
##	
CXCR3 FM0: Mono	
##	
26	
##	
CXCR3 FM0: Mono/CD33+	
##	
26	
##	C
XCR3: Lymph/CD16+/CD56+	
##	
26	
##	CXCR3: L
ymph/CD16+/CD56+/CXCR3+	
##	
26	
##	
CXCR3: Lymph/CD3-	
##	
26	
##	CXCR
3: Lymph/CD3-/CD19+CD20+	
##	
26	
##	CXCR3: Lymph
h/CD3-/CD19+CD20+/CXCR3+	
##	
26	
##	
CXCR3: Lymph/CD3+	
##	
26	
##	
CXCR3: Lymph/CD3+/CD4+	
##	
26	
##	CXCR3:
Lymph/CD3+/CD4+/CXCR3+	
##	
26	
##	
CXCR3: Lymph/CD3+/CD8+	
##	
26	
##	CXCR3:
Lymph/CD3+/CD8+/CXCR3+	
##	
26	
##	
CXCR3: Mono	



##  
26  
##  
CXCR3: Mono/CD33+  
##  
26  
##  
XCR3: Mono/CD33+/CXCR3+  
##  
26  
##  
DHEA  
##  
26  
##  
EGF  
##  
191  
##  
ENA78  
##  
687  
##  
hil Absolute Count-K/uL  
##  
50  
##  
EOTAXIN  
##  
716  
##  
Erythrocyte-%  
##  
50  
##  
ESTRADIOL  
##  
26  
##  
FASL  
##  
375  
##  
FGFB  
##  
716  
##  
FLT3L  
##  
29  
##  
FRACTALKINE  
##  
29

C

Eosinop

##  
GCSF  
##  
716  
##  
GMCSF  
##  
734  
##  
GRO  
##  
29  
##  
GROA  
##  
687  
##  
Hematocrit-%  
##  
50  
##  
Hemoglobin-g/dL  
##  
50  
##  
HGF  
##  
687  
##  
ICAM1  
##  
687  
##  
IFNA  
##  
687  
##  
IFNA2  
##  
29  
##  
IFNB  
##  
687  
##  
IFNG  
##  
734  
##  
IL10  
##  
734  
##  
IL12P40

##  
716  
##  
IL12P70  
##  
734  
##  
IL13  
##  
716  
##  
IL15  
##  
716  
##  
IL17  
##  
554  
##  
IL17A  
##  
162  
##  
IL17F  
##  
687  
##  
IL18  
##  
228  
##  
IL1A  
##  
716  
##  
IL1B  
##  
800  
##  
IL1RA  
##  
716  
##  
IL2  
##  
734  
##  
IL21  
##  
162  
##  
IL22  
##  
162

##  
IL23  
##  
162  
##  
IL27  
##  
162  
##  
IL3  
##  
29  
##  
IL31  
##  
162  
##  
IL4  
##  
716  
##  
IL5  
##  
716  
##  
IL6  
##  
800  
##  
IL7  
##  
716  
##  
IL8  
##  
800  
##  
IL9  
##  
191  
##  
IP10  
##  
716  
##  
LEPTIN  
##  
716  
##  
LIF  
##  
687  
##  
Lymphocyte-%

##	
50	
##	Lymphoc
yte Absolute Count-K/uL	
##	
50	
##	
MCP1	
##	
716	
##	
MCP3	
##	
716	
##	
MCSF	
##	
687	
##	
MDC	
##	
29	
##	Mean
Cell Hemoglobin-pg/cell	
##	
50	
##	Mean Cell Hemogl
obin Concentration-g/dL	
##	
50	
##	Mean
Cell Volume-femtoliter	
##	
50	
##	
MIG	
##	
687	
##	
MIP1A	
##	
716	
##	
MIP1B	
##	
716	
##	
Mono: pSTAT1	
##	
1288	
##	
Mono: pSTAT3	
##	
1288	

```

##
Mono: pSTAT5
##
1288
##
Monocyte-%
##
50
##
yte Absolute Count-K/uL
##
50
##
Neutrophil-%
##
50
##
hil Absolute Count-K/uL
##
50
##
NGF
##
716
##
NK-NKT: Lymph/CD3-
##
26
##
Lymph/CD3-/CD16+/CD56+
##
26
##
CD3-/CD16+/CD56+/HLADR+
##
26
##
+/CD56+/Q1: CD314-CD94+
##
26
##
+/CD56+/Q2: CD314+CD94+
##
26
##
+/CD56+/Q3: CD314+CD94-
##
26
##
+/CD56+/Q4: CD314-CD94-
##
26
##
NK-NKT: Lymph/CD3+

```

Monoc

Neutrop

NK-NKT:

NK-NKT: Lymph/

NK-NKT: Lymph/CD3-/CD16

NK-NKT: Lymph/CD3-/CD16

NK-NKT: Lymph/CD3-/CD16

NK-NKT: Lymph/CD3-/CD16

```
##
26
##
NK-NKT: Lymph/CD3+/CD8-
##
26
##
NK-NKT: Lymph/CD
3+/CD8-/Q1: CD314-CD94+
##
26
##
NK-NKT: Lymph/CD
3+/CD8-/Q2: CD314+CD94+
##
26
##
NK-NKT: Lymph/CD
3+/CD8-/Q3: CD314+CD94-
##
26
##
NK-NKT: Lymph/CD
3+/CD8-/Q4: CD314-CD94-
##
26
##
NK-NKT: Lymph/CD3+/CD8+
##
26
##
NK-NKT: Lymph/CD
3+/CD8+/Q1: CD314-CD94+
##
26
##
NK-NKT: Lymph/CD
3+/CD8+/Q2: CD314+CD94+
##
26
##
NK-NKT: Lymph/CD
3+/CD8+/Q3: CD314+CD94-
##
26
##
NK-NKT: Lymph/CD
3+/CD8+/Q4: CD314-CD94-
##
26
##
NK-NKT: Lymph/CD3+CD56+
##
26
##
Non BT: pSTAT1
##
526
##
Non BT: pSTAT3
##
526
```

```
##
Non BT: pSTAT5
##
526
## Nonbasophils/Nonbasophil CD14-CD33-/Nonbasophil CD3-CD20-/Nonb
asophil CD56-CD16-/mDCs
##
186
## Nonbasophils/Nonbasophil CD14-CD33-/Nonbasophil CD3-CD20-/Nonb
asophil CD56-CD16-/pDCs
##
186
##
PAI1
##
687
##
PDGFAA
##
29
##
PDGFABBB
##
29
##
PDGFBB
##
716
##
PIGF1
##
119
##
Platelets-K/uL
##
50
##
PROGESTERONE
##
26
##
RANTES
##
716
##
RBC-MIL/uL
##
50
##
BC Distribution Width-%
##
50
##
RESISTIN
```

R



```
##
687
##
SCF
##
687
##
SDF1A
##
162
##
SFASL
##
342
##
SIL2RA
##
29
##
T cell: Lymph/CD3+
##
26
##
  T cell: Lymph/CD3+/CD4+
##
26
##
T cell: Lymph/CD3+/CD4+/CD28+
##
26
##
T cell: Lymph/CD3+/CD4+/CD85j+
##
26
##
T cell: Lymph/CD3
+/CD4+/Q1: CD45RA-CD27+
##
26
##
T cell: Lymph/CD3
+/CD4+/Q2: CD45RA+CD27+
##
26
##
T cell: Lymph/CD3
+/CD4+/Q3: CD45RA+CD27-
##
26
##
T cell: Lymph/CD3
+/CD4+/Q4: CD45RA-CD27-
##
26
##
  T cell: Lymph/CD3+/CD8+
##
26
```

##	T cel
1: Lymph/CD3+/CD8+/CD28+	
##	
26	
##	T cell:
Lymph/CD3+/CD8+/CD85j+	
##	
26	
##	T cell: Lymph/CD3
+ /CD8+/Q1: CD45RA-CD27+	
##	
26	
##	T cell: Lymph/CD3
+ /CD8+/Q2: CD45RA+CD27+	
##	
26	
##	T cell: Lymph/CD3
+ /CD8+/Q3: CD45RA+CD27-	
##	
26	
##	T cell: Lymph/CD3
+ /CD8+/Q4: CD45RA-CD27-	
##	
26	
##	
TESTOSTERONE	
##	
147	
##	
TGFA	
##	
716	
##	
TGFB	
##	
687	
##	
TNFA	
##	
800	
##	
TNFB	
##	
716	
##	
TRAIL	
##	
621	
##	
Treg: Lymph/CD3+	
##	
26	
##	
Treg: Lymph/CD3+/CD4+	

```
##
26
##
Lymph/CD3+/CD4+/CD161+
##
26
##
D3+/CD4+/CD25hiCD127low
##
26
##
27low/Q1: CD161-CD45RA+
##
26
##
27low/Q2: CD161+CD45RA+
##
26
##
27low/Q3: CD161+CD45RA-
##
26
##
Treg: Lymph/CD3+/CD8+
##
26
##
Lymph/CD3+/CD8+/CD161+
##
26
##
VCAM1
##
687
##
VEGF
##
716
##
VEGFD
##
162
##
WBC-K/uL
##
50
```

Treg:

Treg: Lymph/C

Treg: Lymph/CD3+/CD4+/CD25hiCD1

Treg: Lymph/CD3+/CD4+/CD25hiCD1

Treg: Lymph/CD3+/CD4+/CD25hiCD1

Treg:

Unique values of variable "data"

```
table(flu$data)
```

##										
##	-4.19	-4.16	-4.15	-3.98	-3.92	-3.75	-3.69	-3.64	-3.63	-3.6
##	1	1	2	1	1	1	1	1	1	1
##	-3.58	-3.48	-3.47	-3.45	-3.43	-3.41	-3.39	-3.37	-3.35	-3.33
##	1	1	3	2	2	1	1	1	1	1
##	-3.28	-3.26	-3.25	-3.24	-3.22	-3.21	-3.2	-3.15	-3.14	-3.13
##	2	1	1	1	1	1	2	1	1	2
##	-3.11	-3.1	-3.09	-3.07	-3.06	-3.05	-3.04	-2.99	-2.98	-2.97
##	2	1	1	1	2	1	1	1	2	2
##	-2.95	-2.94	-2.93	-2.92	-2.89	-2.88	-2.85	-2.84	-2.83	-2.82
##	5	1	2	4	2	1	4	1	1	1
##	-2.81	-2.8	-2.79	-2.78	-2.77	-2.76	-2.75	-2.74	-2.71	-2.7
##	1	1	4	2	1	1	4	1	1	2
##	-2.69	-2.68	-2.67	-2.66	-2.65	-2.64	-2.63	-2.62	-2.61	-2.6
##	3	2	1	2	4	2	3	1	2	3
##	-2.59	-2.58	-2.57	-2.56	-2.55	-2.54	-2.53	-2.52	-2.51	-2.5
##	2	4	3	1	2	3	3	4	2	4
##	-2.49	-2.48	-2.47	-2.46	-2.45	-2.44	-2.43	-2.42	-2.41	-2.4
##	5	6	4	5	10	5	8	4	4	4
##	-2.39	-2.38	-2.37	-2.36	-2.35	-2.34	-2.33	-2.32	-2.31	-2.3
##	6	3	5	6	3	6	4	8	9	4
##	-2.29	-2.28	-2.27	-2.26	-2.25	-2.24	-2.23	-2.22	-2.21	-2.2
##	5	5	9	5	4	5	6	10	7	8
##	-2.19	-2.18	-2.17	-2.16	-2.15	-2.14	-2.13	-2.12	-2.11	-2.1
##	6	9	6	8	6	2	7	7	10	10
##	-2.09	-2.08	-2.07	-2.06	-2.05	-2.04	-2.03	-2.02	-2.01	-2
##	6	8	3	14	11	7	11	6	7	12
##	-1.99	-1.98	-1.97	-1.96	-1.95	-1.94	-1.93	-1.92	-1.91	-1.9
##	5	10	12	8	11	9	8	17	14	12
##	-1.89	-1.88	-1.87	-1.86	-1.85	-1.84	-1.83	-1.82	-1.81	-1.8
##	8	14	16	15	20	10	10	14	11	16
##	-1.79	-1.78	-1.77	-1.76	-1.75	-1.74	-1.73	-1.72	-1.71	-1.7
##	11	16	11	15	17	19	20	16	16	11
##	-1.69	-1.68	-1.67	-1.66	-1.65	-1.64	-1.63	-1.62	-1.61	-1.6
##	24	15	19	23	14	18	24	31	18	30
##	-1.59	-1.58	-1.57	-1.56	-1.55	-1.54	-1.53	-1.52	-1.51	-1.5
##	30	31	27	37	26	24	30	30	24	36
##	-1.49	-1.48	-1.47	-1.46	-1.45	-1.44	-1.43	-1.42	-1.41	-1.4
##	31	33	25	38	27	31	37	30	41	43
##	-1.39	-1.38	-1.37	-1.36	-1.35	-1.34	-1.33	-1.32	-1.31	-1.3
##	30	35	38	43	40	42	43	45	43	39
##	-1.29	-1.28	-1.27	-1.26	-1.25	-1.24	-1.23	-1.22	-1.21	-1.2
##	52	48	27	41	56	46	50	48	62	51
##	-1.19	-1.18	-1.17	-1.16	-1.15	-1.14	-1.13	-1.12	-1.11	-1.1
##	57	72	70	68	63	61	78	71	74	62
##	-1.09	-1.08	-1.07	-1.06	-1.05	-1.04	-1.03	-1.02	-1.01	-1
##	72	82	80	89	76	81	82	91	83	108
##	-0.99	-0.98	-0.97	-0.96	-0.95	-0.94	-0.93	-0.92	-0.91	-0.9
##	101	84	73	87	107	93	98	106	124	115
##	-0.89	-0.88	-0.87	-0.86	-0.85	-0.84	-0.83	-0.82	-0.81	-0.8
##	110	102	97	111	105	110	134	124	117	137
##	-0.79	-0.78	-0.77	-0.76	-0.75	-0.74	-0.73	-0.72	-0.71	-0.7
##	120	143	123	132	120	114	150	149	137	152

##	-0.69	-0.68	-0.67	-0.66	-0.65	-0.64	-0.63	-0.62	-0.61	-0.6
##	158	149	158	153	138	132	134	142	189	159
##	-0.59	-0.58	-0.57	-0.56	-0.55	-0.54	-0.53	-0.52	-0.51	-0.5
##	156	155	152	171	146	177	146	136	169	163
##	-0.49	-0.48	-0.47	-0.46	-0.45	-0.44	-0.43	-0.42	-0.41	-0.4
##	187	160	168	174	164	170	200	180	168	186
##	-0.39	-0.38	-0.37	-0.36	-0.35	-0.34	-0.33	-0.32	-0.31	-0.3
##	161	173	166	187	179	194	176	178	181	170
##	-0.29	-0.28	-0.27	-0.26	-0.25	-0.24	-0.23	-0.22	-0.21	-0.2
##	172	193	165	169	201	149	167	172	215	186
##	-0.19	-0.18	-0.17	-0.16	-0.15	-0.14	-0.13	-0.12	-0.11	-0.1
##	169	182	214	227	223	226	216	221	280	324
##	-0.09	-0.08	-0.07	-0.06	-0.05	-0.04	-0.03	-0.02	-0.01	0
##	479	1037	2454	3306	3310	3147	3257	2625	2235	1525
##	0.01	0.02	0.03	0.04	0.05	0.06	0.07	0.08	0.09	0.1
##	654	593	646	652	652	659	650	602	674	657
##	0.11	0.12	0.13	0.14	0.15	0.16	0.17	0.18	0.19	0.2
##	616	601	605	614	680	604	644	724	629	672
##	0.21	0.22	0.23	0.24	0.25	0.26	0.27	0.28	0.29	0.3
##	746	693	645	758	679	711	639	606	680	685
##	0.31	0.32	0.33	0.34	0.35	0.36	0.37	0.38	0.39	0.4
##	635	606	647	601	576	653	556	505	513	538
##	0.41	0.42	0.43	0.44	0.45	0.46	0.47	0.48	0.49	0.5
##	487	468	431	373	447	417	361	397	382	393
##	0.51	0.52	0.53	0.54	0.55	0.56	0.57	0.58	0.59	0.6
##	358	381	400	340	387	343	338	346	316	321
##	0.61	0.62	0.63	0.64	0.65	0.66	0.67	0.68	0.69	0.7
##	318	308	340	346	328	293	316	348	311	285
##	0.71	0.72	0.73	0.74	0.75	0.76	0.77	0.78	0.79	0.8
##	307	266	269	334	265	283	256	273	237	271
##	0.81	0.82	0.83	0.84	0.85	0.86	0.87	0.88	0.89	0.9
##	256	277	263	238	244	257	237	255	224	273
##	0.91	0.92	0.93	0.94	0.95	0.96	0.97	0.98	0.99	1
##	246	239	254	219	211	231	216	229	218	189
##	1.01	1.02	1.03	1.04	1.05	1.06	1.07	1.08	1.09	1.1
##	231	207	219	197	208	185	188	198	179	188
##	1.11	1.12	1.13	1.14	1.15	1.16	1.17	1.18	1.19	1.2
##	193	156	190	178	162	199	174	167	204	151
##	1.21	1.22	1.23	1.24	1.25	1.26	1.27	1.28	1.29	1.3
##	157	165	151	133	163	172	141	176	150	164
##	1.31	1.32	1.33	1.34	1.35	1.36	1.37	1.38	1.39	1.4
##	182	159	153	151	156	150	142	167	150	134
##	1.41	1.42	1.43	1.44	1.45	1.46	1.47	1.48	1.49	1.5
##	138	132	135	125	139	123	153	102	135	133
##	1.51	1.52	1.53	1.54	1.55	1.56	1.57	1.58	1.59	1.6
##	134	135	134	113	122	102	106	133	95	113
##	1.61	1.62	1.63	1.64	1.65	1.66	1.67	1.68	1.69	1.7
##	131	113	111	109	114	105	91	118	111	98
##	1.71	1.72	1.73	1.74	1.75	1.76	1.77	1.78	1.79	1.8
##	88	92	117	84	90	89	85	95	91	93
##	1.81	1.82	1.83	1.84	1.85	1.86	1.87	1.88	1.89	1.9
##	75	77	79	83	84	99	111	87	90	68
##	1.91	1.92	1.93	1.94	1.95	1.96	1.97	1.98	1.99	2
##	73	82	82	79	81	69	65	61	79	65

##	2.01	2.02	2.03	2.04	2.05	2.06	2.07	2.08	2.09	2.1
##	72	57	71	46	60	73	70	57	53	58
##	2.11	2.12	2.13	2.14	2.15	2.16	2.17	2.18	2.19	2.2
##	63	57	58	59	59	67	63	54	58	51
##	2.21	2.22	2.23	2.24	2.25	2.26	2.27	2.28	2.29	2.3
##	60	53	72	67	63	38	39	56	51	50
##	2.31	2.32	2.33	2.34	2.35	2.36	2.37	2.38	2.39	2.4
##	51	41	53	58	46	34	62	47	57	42
##	2.41	2.42	2.43	2.44	2.45	2.46	2.47	2.48	2.49	2.5
##	51	46	49	51	48	49	40	32	35	31
##	2.51	2.52	2.53	2.54	2.55	2.56	2.57	2.58	2.59	2.6
##	26	33	32	41	45	34	37	36	24	43
##	2.61	2.62	2.63	2.64	2.65	2.66	2.67	2.68	2.69	2.7
##	31	34	33	22	29	39	39	32	24	39
##	2.71	2.72	2.73	2.74	2.75	2.76	2.77	2.78	2.79	2.8
##	26	21	28	35	24	27	35	21	25	31
##	2.81	2.82	2.83	2.84	2.85	2.86	2.87	2.88	2.89	2.9
##	30	26	18	26	35	26	23	26	33	21
##	2.91	2.92	2.93	2.94	2.95	2.96	2.97	2.98	2.99	3
##	23	25	33	26	24	29	27	32	17	20
##	3.01	3.02	3.03	3.04	3.05	3.06	3.07	3.08	3.09	3.1
##	23	28	23	24	21	26	22	20	21	24
##	3.11	3.12	3.13	3.14	3.15	3.16	3.17	3.18	3.19	3.2
##	15	22	20	23	23	22	19	21	27	22
##	3.21	3.22	3.23	3.24	3.25	3.26	3.27	3.28	3.29	3.3
##	21	21	20	21	20	20	19	24	26	19
##	3.31	3.32	3.33	3.34	3.35	3.36	3.37	3.38	3.39	3.4
##	24	21	16	9	25	22	19	19	22	14
##	3.41	3.42	3.43	3.44	3.45	3.46	3.47	3.48	3.49	3.5
##	18	27	24	17	27	13	14	22	14	11
##	3.51	3.52	3.53	3.54	3.55	3.56	3.57	3.58	3.59	3.6
##	15	14	15	17	14	14	18	26	14	15
##	3.61	3.62	3.63	3.64	3.65	3.66	3.67	3.68	3.69	3.7
##	18	15	16	12	12	26	12	8	10	24
##	3.71	3.72	3.73	3.74	3.75	3.76	3.77	3.78	3.79	3.8
##	15	14	10	16	8	15	12	14	17	13
##	3.81	3.82	3.83	3.84	3.85	3.86	3.87	3.88	3.89	3.9
##	7	12	12	13	14	14	13	15	15	10
##	3.91	3.92	3.93	3.94	3.95	3.96	3.97	3.98	3.99	4
##	11	11	15	19	13	14	10	13	17	12
##	4.01	4.02	4.03	4.04	4.05	4.06	4.07	4.08	4.09	4.1
##	7	12	17	8	11	7	8	9	15	14
##	4.11	4.12	4.13	4.14	4.15	4.16	4.17	4.18	4.19	4.2
##	9	13	12	11	11	10	18	14	13	15
##	4.21	4.22	4.23	4.24	4.25	4.26	4.27	4.28	4.29	4.3
##	9	8	14	13	10	17	12	13	8	12
##	4.31	4.32	4.33	4.34	4.35	4.36	4.37	4.38	4.39	4.4
##	13	18	11	12	17	13	19	15	9	12
##	4.41	4.42	4.43	4.44	4.45	4.46	4.47	4.48	4.49	4.5
##	14	13	9	14	9	19	14	10	20	17
##	4.51	4.52	4.53	4.54	4.55	4.56	4.57	4.58	4.59	4.6
##	15	13	17	15	13	10	17	9	15	11
##	4.61	4.62	4.63	4.64	4.65	4.66	4.67	4.68	4.69	4.7
##	6	18	6	7	13	14	12	10	8	13

##	4.71	4.72	4.73	4.74	4.75	4.76	4.77	4.78	4.79	4.8
##	11	10	12	15	6	11	10	14	12	15
##	4.81	4.82	4.83	4.84	4.85	4.86	4.87	4.88	4.89	4.9
##	7	6	5	14	5	9	10	11	10	14
##	4.91	4.92	4.93	4.94	4.95	4.96	4.97	4.98	4.99	5
##	13	6	10	14	12	13	14	9	17	11
##	5.01	5.02	5.03	5.04	5.05	5.06	5.07	5.08	5.09	5.1
##	8	13	10	8	8	14	11	9	9	14
##	5.11	5.12	5.13	5.14	5.15	5.16	5.17	5.18	5.19	5.2
##	8	13	9	8	13	9	13	10	7	6
##	5.21	5.22	5.23	5.24	5.25	5.26	5.27	5.28	5.29	5.3
##	4	6	9	8	7	11	9	8	6	11
##	5.31	5.32	5.33	5.34	5.35	5.36	5.37	5.38	5.39	5.4
##	8	5	8	10	14	12	9	9	9	11
##	5.41	5.42	5.43	5.44	5.45	5.46	5.47	5.48	5.49	5.5
##	11	11	8	8	7	17	8	4	10	8
##	5.51	5.52	5.53	5.54	5.55	5.56	5.57	5.58	5.59	5.6
##	8	7	12	7	13	11	12	6	11	17
##	5.61	5.62	5.63	5.64	5.65	5.66	5.67	5.68	5.69	5.7
##	6	10	5	12	10	13	9	10	5	11
##	5.71	5.72	5.73	5.74	5.75	5.76	5.77	5.78	5.79	5.8
##	13	13	5	8	17	11	12	11	12	10
##	5.81	5.82	5.83	5.84	5.85	5.86	5.87	5.88	5.89	5.9
##	9	7	14	7	7	4	12	12	9	12
##	5.91	5.92	5.93	5.94	5.95	5.96	5.97	5.98	5.99	6
##	6	12	4	7	9	7	12	11	7	19
##	6.01	6.02	6.03	6.04	6.05	6.06	6.07	6.08	6.09	6.1
##	9	10	7	9	13	8	10	6	9	6
##	6.11	6.12	6.13	6.14	6.15	6.16	6.17	6.18	6.19	6.2
##	6	11	6	7	12	6	3	11	7	6
##	6.21	6.22	6.23	6.24	6.25	6.26	6.27	6.28	6.29	6.3
##	9	10	2	12	11	5	7	6	6	8
##	6.31	6.32	6.33	6.34	6.35	6.36	6.37	6.38	6.39	6.4
##	13	11	15	12	4	5	7	5	11	14
##	6.41	6.42	6.43	6.44	6.45	6.46	6.47	6.48	6.49	6.5
##	11	7	7	8	7	7	10	6	9	14
##	6.51	6.52	6.53	6.54	6.55	6.56	6.57	6.58	6.59	6.6
##	8	10	4	7	5	10	7	5	6	7
##	6.61	6.62	6.63	6.64	6.65	6.66	6.67	6.68	6.69	6.7
##	5	6	4	9	10	10	8	7	9	6
##	6.71	6.72	6.73	6.74	6.75	6.76	6.77	6.78	6.79	6.8
##	9	7	6	6	5	7	6	9	8	10
##	6.81	6.82	6.83	6.84	6.85	6.86	6.87	6.88	6.89	6.9
##	8	10	6	6	5	11	9	11	3	5
##	6.91	6.92	6.93	6.94	6.95	6.96	6.97	6.98	6.99	7
##	8	7	9	7	5	5	9	12	9	12
##	7.01	7.02	7.03	7.04	7.05	7.06	7.07	7.08	7.09	7.1
##	11	16	9	5	11	7	9	6	6	10
##	7.11	7.12	7.13	7.14	7.15	7.16	7.17	7.18	7.19	7.2
##	12	8	5	9	10	6	9	4	3	8
##	7.21	7.22	7.23	7.24	7.25	7.26	7.27	7.28	7.29	7.3
##	7	10	8	6	6	11	13	8	8	10
##	7.31	7.32	7.33	7.34	7.35	7.36	7.37	7.38	7.39	7.4
##	9	7	8	10	4	6	7	5	8	8

##	7.41	7.42	7.43	7.44	7.45	7.46	7.47	7.48	7.49	7.5
##	9	10	10	12	7	8	4	7	11	20
##	7.51	7.52	7.53	7.54	7.55	7.56	7.57	7.58	7.59	7.6
##	12	4	11	7	9	9	5	9	13	16
##	7.61	7.62	7.63	7.64	7.65	7.66	7.67	7.68	7.69	7.7
##	5	8	3	8	5	9	6	10	5	9
##	7.71	7.72	7.73	7.74	7.75	7.76	7.77	7.78	7.79	7.8
##	14	10	6	9	5	8	8	4	5	5
##	7.81	7.82	7.83	7.84	7.85	7.86	7.87	7.88	7.89	7.9
##	6	7	11	4	4	8	7	11	3	10
##	7.91	7.92	7.93	7.94	7.95	7.96	7.97	7.98	7.99	8
##	3	5	7	6	12	6	9	8	11	13
##	8.01	8.02	8.03	8.04	8.05	8.06	8.07	8.08	8.09	8.1
##	6	10	3	7	6	7	6	7	9	14
##	8.11	8.12	8.13	8.14	8.15	8.16	8.17	8.18	8.19	8.2
##	6	3	10	6	6	5	7	4	11	11
##	8.21	8.22	8.23	8.24	8.25	8.26	8.27	8.28	8.29	8.3
##	8	6	5	6	7	9	11	8	7	8
##	8.31	8.32	8.33	8.34	8.35	8.36	8.37	8.38	8.39	8.4
##	5	10	13	8	12	6	5	4	8	5
##	8.41	8.42	8.43	8.44	8.45	8.46	8.47	8.48	8.49	8.5
##	5	4	10	5	5	5	5	10	8	10
##	8.51	8.52	8.53	8.54	8.55	8.56	8.57	8.58	8.59	8.6
##	8	8	7	5	3	11	8	6	4	10
##	8.61	8.62	8.63	8.64	8.65	8.66	8.67	8.68	8.69	8.7
##	3	5	1	6	6	1	8	4	7	10
##	8.71	8.72	8.73	8.74	8.75	8.76	8.77	8.78	8.79	8.8
##	2	7	3	11	7	4	9	8	4	14
##	8.81	8.82	8.83	8.84	8.85	8.86	8.87	8.88	8.89	8.9
##	3	5	7	7	8	6	9	4	5	11
##	8.91	8.92	8.93	8.94	8.95	8.96	8.97	8.98	8.99	9
##	2	5	8	4	9	4	8	7	8	17
##	9.01	9.02	9.03	9.04	9.05	9.06	9.07	9.08	9.09	9.1
##	10	7	5	7	4	8	9	8	10	8
##	9.11	9.12	9.13	9.14	9.15	9.16	9.17	9.18	9.19	9.2
##	5	4	9	8	6	9	6	4	7	7
##	9.21	9.22	9.23	9.24	9.25	9.26	9.27	9.28	9.29	9.3
##	10	4	4	13	6	5	9	1	4	7
##	9.31	9.32	9.33	9.34	9.35	9.36	9.37	9.38	9.39	9.4
##	8	6	7	10	3	6	3	10	2	12
##	9.41	9.42	9.43	9.44	9.45	9.46	9.47	9.48	9.49	9.5
##	5	9	5	6	2	5	4	7	9	11
##	9.51	9.52	9.53	9.54	9.55	9.56	9.57	9.58	9.59	9.6
##	2	5	3	4	5	3	5	4	6	6
##	9.61	9.62	9.63	9.64	9.65	9.66	9.67	9.68	9.69	9.7
##	5	7	9	5	5	2	2	9	7	7
##	9.71	9.72	9.73	9.74	9.75	9.76	9.77	9.78	9.79	9.8
##	7	6	4	10	5	2	4	1	4	3
##	9.81	9.82	9.83	9.84	9.85	9.86	9.87	9.88	9.89	9.9
##	8	5	1	8	7	6	3	7	6	5
##	9.91	9.92	9.93	9.94	9.95	9.96	9.97	9.98	9.99	10
##	1	7	3	4	5	4	7	7	11	37
##	10.1	10.2	10.3	10.4	10.5	10.6	10.7	10.75	10.8	10.9
##	57	52	55	51	54	61	50	1	44	50



##	11	11.1	11.2	11.3	11.4	11.5	11.6	11.7	11.75	11.8
##	60	43	44	40	43	48	38	45	1	38
##	11.9	12	12.1	12.2	12.25	12.3	12.4	12.5	12.6	12.7
##	46	39	48	44	2	44	38	51	36	54
##	12.75	12.8	12.9	13	13.1	13.2	13.25	13.3	13.4	13.5
##	2	43	42	55	50	48	1	48	44	54
##	13.6	13.7	13.8	13.9	14	14.1	14.2	14.3	14.4	14.5
##	40	48	35	51	43	43	27	41	39	50
##	14.6	14.7	14.8	14.9	15	15.1	15.2	15.3	15.4	15.5
##	43	40	47	29	38	40	45	38	41	29
##	15.6	15.7	15.8	15.9	16	16.1	16.2	16.25	16.3	16.4
##	32	39	41	28	42	34	39	1	29	27
##	16.5	16.6	16.7	16.8	16.9	17	17.1	17.2	17.3	17.4
##	35	35	41	31	30	33	36	23	21	30
##	17.5	17.6	17.7	17.8	17.9	18	18.1	18.2	18.25	18.3
##	29	34	33	32	18	42	27	42	2	28
##	18.4	18.5	18.6	18.7	18.8	18.9	19	19.1	19.2	19.3
##	29	30	27	22	26	26	27	38	23	25
##	19.4	19.5	19.6	19.7	19.8	19.9	20	20.1	20.2	20.3
##	25	25	23	27	24	25	34	29	32	25
##	20.4	20.5	20.6	20.7	20.8	20.9	21	21.1	21.2	21.25
##	14	25	19	25	23	25	24	23	24	1
##	21.3	21.4	21.5	21.6	21.7	21.75	21.8	21.9	22	22.1
##	29	25	28	21	19	1	23	27	17	31
##	22.2	22.3	22.4	22.5	22.6	22.7	22.8	22.9	23	23.1
##	27	11	18	16	21	14	20	17	27	23
##	23.2	23.25	23.3	23.4	23.5	23.6	23.7	23.8	23.9	24
##	24	1	26	22	18	19	24	21	17	32
##	24.1	24.2	24.3	24.4	24.5	24.6	24.7	24.8	24.9	25
##	25	16	25	22	25	24	23	21	26	32
##	25.1	25.2	25.25	25.3	25.4	25.5	25.6	25.7	25.8	25.9
##	16	19	2	26	16	24	20	22	30	26
##	26	26.1	26.2	26.3	26.4	26.5	26.6	26.7	26.8	26.9
##	15	30	25	10	14	16	16	16	28	15
##	27	27.1	27.2	27.3	27.4	27.5	27.6	27.7	27.8	27.9
##	24	19	16	18	21	23	12	10	21	18
##	28	28.1	28.2	28.25	28.3	28.4	28.5	28.6	28.7	28.8
##	18	20	32	1	24	22	35	24	18	34
##	28.9	29	29.1	29.2	29.3	29.4	29.5	29.6	29.7	29.8
##	25	25	18	22	13	20	17	24	20	22
##	29.9	30	30.1	30.2	30.25	30.3	30.4	30.5	30.6	30.7
##	10	27	15	19	1	17	26	16	19	31
##	30.75	30.8	30.9	31	31.1	31.2	31.3	31.4	31.5	31.6
##	1	20	17	20	24	22	22	26	15	25
##	31.7	31.75	31.8	31.9	32	32.1	32.2	32.3	32.4	32.5
##	20	1	27	18	27	18	18	21	18	19
##	32.6	32.7	32.75	32.8	32.9	33	33.1	33.2	33.25	33.3
##	20	28	2	23	20	23	15	20	3	27
##	33.4	33.5	33.6	33.7	33.75	33.8	33.9	34	34.1	34.2
##	15	23	24	21	2	20	23	22	20	22
##	34.25	34.3	34.4	34.5	34.6	34.7	34.8	34.9	35	35.1
##	1	25	25	25	32	25	19	19	27	26
##	35.2	35.3	35.4	35.5	35.6	35.7	35.8	35.9	36	36.1
##	22	22	19	24	16	22	16	15	18	19

##	36.2	36.25	36.3	36.4	36.5	36.6	36.7	36.8	36.9	37
##	22	1	17	22	20	18	18	18	19	16
##	37.1	37.2	37.3	37.4	37.5	37.6	37.7	37.75	37.8	37.9
##	26	25	24	18	24	15	17	1	21	15
##	38	38.1	38.2	38.25	38.3	38.4	38.5	38.6	38.7	38.8
##	31	18	15	1	19	17	15	30	23	16
##	38.9	39	39.1	39.2	39.3	39.4	39.5	39.6	39.7	39.75
##	15	23	23	20	10	22	15	21	25	1
##	39.8	39.9	40	40.1	40.2	40.3	40.4	40.5	40.6	40.7
##	23	15	32	23	20	24	18	22	11	24
##	40.75	40.8	40.9	41	41.1	41.2	41.25	41.3	41.4	41.5
##	1	17	25	19	20	16	1	22	11	16
##	41.6	41.7	41.8	41.9	42	42.1	42.2	42.3	42.4	42.5
##	18	19	16	13	20	18	14	15	17	15
##	42.6	42.7	42.8	42.9	43	43.1	43.2	43.3	43.4	43.5
##	17	14	16	15	18	11	13	13	14	14
##	43.6	43.7	43.8	43.9	44	44.1	44.2	44.3	44.4	44.5
##	15	28	20	10	17	20	11	16	19	21
##	44.6	44.7	44.8	44.9	45	45.1	45.2	45.3	45.4	45.5
##	11	15	18	17	8	9	18	15	14	16
##	45.6	45.7	45.8	45.9	46	46.1	46.2	46.3	46.4	46.5
##	19	12	9	10	13	12	6	17	7	11
##	46.6	46.7	46.8	46.9	47	47.1	47.2	47.3	47.4	47.5
##	11	14	18	15	14	16	16	13	13	19
##	47.6	47.7	47.75	47.8	47.9	48	48.1	48.2	48.3	48.4
##	13	12	1	17	9	16	11	9	26	12
##	48.5	48.6	48.7	48.8	48.9	49	49.1	49.2	49.3	49.4
##	17	15	14	11	12	21	19	13	14	10
##	49.5	49.6	49.7	49.8	49.9	50	50.1	50.2	50.3	50.4
##	16	26	14	12	13	28	11	21	4	14
##	50.5	50.6	50.7	50.75	50.8	50.9	51	51.1	51.2	51.25
##	16	10	18	1	15	13	10	11	9	1
##	51.3	51.4	51.5	51.6	51.7	51.75	51.8	51.9	52	52.1
##	6	17	13	15	10	1	13	18	9	16
##	52.2	52.3	52.4	52.5	52.6	52.7	52.8	52.9	53	53.1
##	7	11	17	14	11	14	15	19	11	17
##	53.2	53.3	53.4	53.5	53.6	53.7	53.8	53.9	54	54.1
##	12	12	16	19	17	10	10	12	16	11
##	54.2	54.3	54.4	54.5	54.6	54.7	54.75	54.8	54.9	55
##	17	15	14	19	13	22	1	15	18	9
##	55.1	55.2	55.3	55.4	55.5	55.6	55.7	55.8	55.9	56
##	18	13	16	18	21	18	13	18	12	16
##	56.1	56.2	56.3	56.4	56.5	56.6	56.7	56.75	56.8	56.9
##	19	19	9	18	13	22	17	1	10	23
##	57	57.1	57.2	57.25	57.3	57.4	57.5	57.6	57.7	57.8
##	13	17	15	1	11	18	21	15	16	13
##	57.9	58	58.1	58.2	58.3	58.4	58.5	58.6	58.7	58.8
##	18	23	11	17	20	20	17	13	12	16
##	58.9	59	59.1	59.2	59.3	59.4	59.5	59.6	59.7	59.8
##	12	15	20	20	13	11	17	11	13	15
##	59.9	60	60.1	60.2	60.3	60.4	60.5	60.6	60.7	60.75
##	15	17	15	14	10	11	14	18	10	1
##	60.8	60.9	61	61.1	61.2	61.25	61.3	61.4	61.5	61.6
##	13	15	14	14	22	1	19	19	19	22

##	61.7	61.75	61.8	61.9	62	62.1	62.2	62.25	62.3	62.4
##	21	1	14	24	20	17	16	1	12	25
##	62.5	62.6	62.7	62.8	62.9	63	63.1	63.2	63.25	63.3
##	19	20	21	19	23	24	18	16	1	21
##	63.4	63.5	63.6	63.7	63.8	63.9	64	64.1	64.2	64.3
##	13	16	18	14	19	11	17	13	16	14
##	64.4	64.5	64.6	64.7	64.75	64.8	64.9	65	65.1	65.2
##	10	13	18	17	1	16	12	10	14	19
##	65.25	65.3	65.4	65.5	65.6	65.7	65.8	65.9	66	66.1
##	1	10	22	16	17	18	12	14	19	22
##	66.2	66.3	66.4	66.5	66.6	66.7	66.75	66.8	66.9	67
##	13	11	12	17	14	17	1	10	12	25
##	67.1	67.2	67.3	67.4	67.5	67.6	67.7	67.8	67.9	68
##	20	21	14	18	22	15	18	16	14	14
##	68.1	68.2	68.3	68.4	68.5	68.6	68.7	68.8	68.9	69
##	14	18	14	22	12	19	9	16	9	15
##	69.1	69.2	69.3	69.4	69.5	69.6	69.7	69.8	69.9	70
##	15	10	11	12	19	18	14	11	16	10
##	70.1	70.2	70.3	70.4	70.5	70.6	70.7	70.8	70.9	71
##	12	7	15	14	9	15	12	9	13	17
##	71.1	71.2	71.3	71.4	71.5	71.6	71.7	71.8	71.9	72
##	14	11	10	16	13	17	7	15	8	15
##	72.1	72.2	72.3	72.4	72.5	72.6	72.7	72.8	72.9	73
##	11	8	14	9	11	12	18	16	10	6
##	73.1	73.2	73.3	73.4	73.5	73.6	73.7	73.8	73.9	74
##	19	11	12	15	12	17	5	12	8	17
##	74.1	74.2	74.3	74.4	74.5	74.6	74.7	74.8	74.9	75
##	5	16	15	13	12	16	11	14	16	14
##	75.1	75.2	75.3	75.4	75.5	75.6	75.7	75.8	75.9	76
##	11	16	15	7	14	9	8	6	6	8
##	76.1	76.2	76.3	76.4	76.5	76.6	76.7	76.8	76.9	77
##	8	12	8	13	4	13	14	12	12	16
##	77.1	77.2	77.3	77.4	77.5	77.6	77.7	77.8	77.9	78
##	11	17	13	9	16	10	6	8	11	15
##	78.1	78.2	78.3	78.4	78.5	78.6	78.7	78.8	78.9	79
##	12	9	15	10	12	11	13	13	15	14
##	79.1	79.2	79.3	79.4	79.5	79.6	79.7	79.8	79.9	80
##	10	18	9	18	12	15	10	19	9	15
##	80.1	80.2	80.3	80.4	80.5	80.6	80.7	80.8	80.9	81
##	16	15	8	13	5	16	13	17	13	13
##	81.1	81.2	81.25	81.3	81.4	81.5	81.6	81.7	81.8	81.9
##	19	19	1	8	17	9	13	16	8	8
##	82	82.1	82.2	82.3	82.4	82.5	82.6	82.7	82.8	82.9
##	10	11	9	7	10	13	7	11	8	9
##	83	83.1	83.2	83.3	83.4	83.5	83.6	83.7	83.8	83.9
##	5	6	11	13	10	10	12	6	9	11
##	84	84.1	84.2	84.3	84.4	84.5	84.6	84.7	84.75	84.8
##	6	20	6	12	4	15	11	5	1	9
##	84.9	85	85.1	85.2	85.3	85.4	85.5	85.6	85.7	85.8
##	8	13	8	10	10	8	14	8	8	12
##	85.9	86	86.1	86.2	86.3	86.4	86.5	86.6	86.7	86.8
##	11	9	7	4	12	8	13	10	5	6
##	86.9	87	87.1	87.2	87.3	87.4	87.5	87.6	87.7	87.8
##	11	5	7	8	10	6	7	8	11	8

##	87.9	88	88.1	88.2	88.3	88.4	88.5	88.6	88.7	88.8
##	9	14	13	10	9	11	7	11	10	8
##	88.9	89	89.1	89.2	89.3	89.4	89.5	89.6	89.7	89.8
##	10	6	15	13	11	16	9	8	12	19
##	89.9	90	90.1	90.2	90.3	90.4	90.5	90.6	90.7	90.8
##	11	11	15	14	10	10	11	16	8	12
##	90.9	91	91.1	91.2	91.3	91.4	91.5	91.6	91.7	91.8
##	9	11	6	12	8	11	8	8	13	4
##	91.9	92	92.1	92.2	92.3	92.4	92.5	92.6	92.7	92.8
##	9	12	16	12	10	3	11	15	11	15
##	92.9	93	93.1	93.2	93.3	93.4	93.5	93.6	93.7	93.8
##	14	9	13	11	8	9	7	10	8	8
##	93.9	94	94.1	94.2	94.3	94.4	94.5	94.6	94.7	94.8
##	6	9	10	6	18	11	7	11	15	20
##	94.9	95	95.1	95.2	95.3	95.4	95.5	95.6	95.7	95.8
##	14	13	9	18	9	15	15	14	13	21
##	95.9	96	96.1	96.2	96.3	96.4	96.5	96.6	96.7	96.8
##	13	15	9	12	8	10	11	10	6	12
##	96.9	97	97.1	97.2	97.3	97.4	97.5	97.6	97.7	97.8
##	13	13	8	7	5	10	14	8	9	13
##	97.9	98	98.1	98.2	98.3	98.4	98.5	98.6	98.7	98.8
##	8	17	18	10	12	12	11	17	12	15
##	98.9	99	99.1	99.2	99.3	99.4	99.5	99.6	99.7	99.8
##	18	19	11	6	7	6	6	8	11	14
##	99.9	100	101	102	103	104	105	106	106.5	107
##	18	142	3	2	2	2	1	2	1	11
##	108	109	110	111	112	112.5	113	114	115	115.25
##	2	5	2	5	1	1	9	3	8	1
##	116	117	118	119	120	121	122	123	124	125
##	3	5	7	2	5	10	10	9	10	6
##	126	127	128	128.25	129	130	131	131.5	132	132.5
##	8	8	15	1	16	14	17	1	18	1
##	133	133.5	134	135	136	136.5	137	138	139	140
##	34	1	26	31	20	1	21	19	28	25
##	140.5	141	142	142.75	143	143.5	144	145	146	147
##	1	19	33	1	19	1	23	25	21	22
##	148	149	150	151	152	153	154	155	155.5	156
##	32	29	28	27	31	28	24	27	1	20
##	156.5	157	157.5	158	158.5	159	160	161	161.25	162
##	2	21	1	26	1	12	28	25	1	33
##	163	164	165	165.5	166	167	168	169	169.5	170
##	29	24	25	1	39	39	40	43	1	31
##	171	171.25	172	172.5	173	173.5	174	175	176	176.5
##	62	1	45	1	56	1	38	55	51	1
##	177	178	179	180	181	181.5	182	182.5	183	183.5
##	58	52	38	39	47	1	53	1	48	1
##	184	185	186	186.5	187	187.5	188	189	190	190.5
##	64	48	61	1	52	1	56	74	38	1
##	191	192	192.5	193	193.25	193.5	194	195	196	197
##	48	69	1	69	1	4	49	40	50	54
##	198	199	200	201	202	203	204	204.5	205	206
##	52	69	53	61	60	51	39	1	68	62
##	207	208	208.5	209	209.5	210	211	212	213	214
##	54	82	2	59	1	58	78	66	75	79

##	214.5	215	215.5	216	216.5	217	217.5	218	219	220
##	1	87	1	84	1	78	2	76	62	73
##	221	221.5	222	222.5	223	224	224.5	225	225.5	226
##	60	1	71	2	64	73	1	70	1	64
##	226.5	227	228	229	229.5	230	231	232	233	234
##	1	61	61	58	1	62	82	77	69	73
##	234.5	235	236	237	238	239	240	240.5	241	242
##	2	76	64	83	61	67	72	1	78	72
##	242.5	243	244	245	246	246.5	247	247.5	248	249
##	2	71	65	68	66	1	63	1	55	65
##	250	251	252	252.5	253	253.25	253.5	254	255	256
##	59	65	61	1	49	1	1	58	56	62
##	256.5	257	257.5	258	259	260	261	261.5	262	263
##	1	71	1	66	57	62	62	1	47	73
##	264	265	266	267	268	269	270	270.5	271	271.5
##	66	56	70	58	63	70	59	1	52	1
##	272	273	274	274.5	275	276	277	278	278.5	279
##	55	61	57	1	38	68	52	57	1	61
##	279.75	280	280.5	281	281.5	282	282.5	283	283.5	284
##	1	63	1	69	1	47	1	56	1	50
##	285	286	286.5	287	288	289	290	291	292	293
##	59	50	1	62	61	62	61	73	70	63
##	294	295	296	297	297.5	298	299	299.5	300	301
##	68	58	46	70	1	59	53	1	58	46
##	301.5	302	303	303.5	304	305	306	306.5	307	308
##	1	71	58	1	57	67	67	1	61	73
##	309	309.25	309.5	310	311	312	313	314	315	316
##	46	1	1	75	64	52	64	77	72	54
##	316.5	317	318	318.5	319	320	321	321.5	322	323
##	1	76	75	1	72	55	51	1	63	53
##	324	325	325.5	325.75	326	327	328	329	330	330.5
##	71	56	1	1	74	67	49	62	65	1
##	331	332	333	334	334.5	335	336	336.5	337	338
##	49	72	51	62	1	70	70	1	59	57
##	338.5	339	339.5	339.75	340	341	342	343	343.5	344
##	1	55	1	1	45	54	46	55	1	49
##	345	346	346.5	347	348	348.5	349	350	351	352
##	57	63	1	71	55	1	56	61	66	74
##	353	354	355	356	357	357.75	358	358.5	359	360
##	44	71	62	58	47	1	68	1	67	57
##	361	362	362.5	363	364	365	366	366.5	367	367.5
##	48	62	1	62	61	71	72	2	58	1
##	368	368.5	369	369.5	369.75	370	371	372	373	374
##	72	1	49	1	1	68	67	64	50	60
##	375	375.5	376	377	377.5	378	379	380	380.5	381
##	48	1	55	58	1	59	55	60	1	54
##	382	383	384	385	385.75	386	387	388	389	390
##	62	46	56	61	1	62	49	44	57	63
##	391	391.5	392	393	394	394.5	395	395.5	396	397
##	40	1	63	54	41	1	51	1	42	49
##	398	398.5	399	400	401	402	403	403.5	404	405
##	54	1	37	55	48	48	53	1	52	49
##	406	407	408	409	410	411	412	412.5	413	414
##	45	40	48	51	44	45	42	1	34	34

##	415	415.75	416	417	418	419	419.5	420	420.5	421
##	45	1	28	44	43	44	1	36	1	49
##	422	422.5	423	424	424.5	425	426	427	428	429
##	27	1	58	33	1	47	30	50	35	48
##	430	431	431.5	432	433	434	435	435.5	436	437
##	35	53	1	40	47	37	33	1	45	18
##	438	439	440	441	442	442.5	443	444	444.5	445
##	39	33	49	31	44	1	28	39	1	36
##	446	446.5	447	447.5	448	448.25	449	450	451	451.5
##	55	1	35	1	47	1	36	39	42	3
##	452	453	454	455	455.5	456	457	458	459	460
##	40	43	45	33	1	46	32	36	31	31
##	461	461.5	462	463	464	465	466	466.5	467	468
##	36	1	40	41	44	29	33	1	27	30
##	469	470	470.5	471	472	473	474	475	476	477
##	28	21	1	27	34	30	27	43	30	36
##	478	479	480	481	482	483	484	485	486	487
##	25	45	26	32	33	34	43	30	28	28
##	487.5	488	489	490	491	492	493	494	494.5	495
##	1	21	30	28	29	21	38	27	1	31
##	496	497	497.25	498	499	499.5	500	501	502	503
##	31	35	1	33	29	1	23	25	38	29
##	504	504.5	505	506	506.5	507	508	509	510	510.5
##	22	1	35	25	2	34	32	20	31	2
##	511	512	513	514	515	516	517	518	519	520
##	31	28	23	24	26	30	26	29	33	31
##	521	522	523	524	524.5	525	526	527	528	529
##	26	23	31	21	1	27	27	33	31	27
##	530	531	532	532.5	533	533.75	534	535	536	536.5
##	29	26	33	1	18	1	27	33	15	1
##	537	538	539	540	541	542	543	544	545	546
##	39	26	31	17	21	25	29	21	31	20
##	547	548	548.5	549	550	551	552	552.75	553	554
##	29	25	1	29	25	22	22	1	11	28
##	555	556	557	557.5	558	559	560	561	562	562.25
##	21	30	25	1	25	21	25	30	23	1
##	563	564	565	566	566.5	567	568	569	570	571
##	20	23	13	25	1	29	20	27	21	28
##	572	573	574	575	576	577	578	579	580	581
##	22	31	30	31	24	31	18	15	31	33
##	582	583	583.5	584	585	586	586.5	587	588	589
##	11	31	1	25	22	11	1	19	25	25
##	590	591	591.5	592	593	594	594.5	595	596	597
##	33	22	1	32	22	25	2	20	23	22
##	598	598.25	599	600	601	602	603	604	605	606
##	14	1	21	21	20	16	18	20	21	27
##	607	607.5	608	609	610	611	612	613	614	615
##	16	1	26	29	28	24	18	25	21	11
##	616	617	618	619	620	621	622	623	624	625
##	16	16	31	16	21	20	19	18	28	16
##	626	627	628	629	629.25	630	630.5	631	632	633
##	17	13	18	17	1	25	1	18	20	22
##	634	635	635.5	636	637	638	638.5	639	640	641
##	20	15	1	24	10	15	1	15	11	17

##	642	643	644	645	646	647	648	649	650	650.5
##	22	26	18	16	22	15	16	24	15	1
##	651	652	653	654	655	656	657	658	659	660
##	19	16	18	12	26	12	17	17	19	13
##	661	661.5	662	663	664	665	666	667	668	669
##	11	1	18	21	7	21	22	20	19	9
##	670	671	672	673	673.25	674	675	676	677	678
##	22	15	17	19	1	12	10	15	11	17
##	679	680	681	682	683	684	685	686	687	688
##	13	21	12	14	18	9	17	17	13	25
##	689	690	691	692	693	694	694.5	695	696	697
##	13	13	19	11	17	16	1	8	22	11
##	698	699	699.5	700	700.5	701	702	703	704	705
##	15	12	1	16	1	10	20	20	9	13
##	706	707	708	709	710	711	711.5	712	712.75	713
##	7	15	18	10	11	12	1	13	1	12
##	714	715	716	717	718	719	720	721	722	723
##	8	13	20	15	18	12	7	17	5	9
##	724	724.5	725	726	727	728	729	729.5	730	731
##	11	1	19	16	12	14	16	1	7	16
##	732	733	734	735	736	737	738	739	740	740.5
##	16	8	20	13	12	15	8	16	12	1
##	741	742	743	744	744.25	745	746	747	748	749
##	8	16	9	20	1	13	4	15	10	8
##	750	751	752	753	754	755	756	757	758	759
##	5	12	17	19	10	14	10	16	12	16
##	759.5	760	761	761.5	762	763	764	765	766	766.5
##	1	11	12	2	12	16	16	14	18	1
##	767	768	769	770	770.5	771	772	773	774	775
##	9	17	10	8	1	14	14	12	6	16
##	776	777	778	779	780	781	782	783	784	785
##	15	20	11	18	12	11	17	13	11	12
##	786	787	788	789	790	791	792	792.5	793	794
##	15	12	15	18	14	11	9	1	10	9
##	795	796	797	798	799	800	801	801.5	802	803
##	11	8	16	9	15	6	14	1	17	14
##	804	805	806	806.5	807	808	809	810	811	811.5
##	8	9	5	1	9	5	16	8	21	1
##	812	813	814	815	816	817	818	819	820	821
##	7	13	14	18	8	10	6	16	13	15
##	822	823	824	825	826	827	828	829	830	831
##	15	10	9	15	14	15	11	11	9	11
##	831.5	832	833	834	835	836	837	838	839	840
##	1	8	8	19	12	11	4	10	11	10
##	841	842	843	844	845	846	847	848	849	850
##	15	15	15	18	7	14	12	17	8	13
##	851	852	853	854	855	856	857	857.5	858	859
##	15	10	12	13	14	12	13	1	11	13
##	860	861	861.5	862	863	864	865	866	867	868
##	12	13	1	9	2	8	16	13	14	9
##	869	870	871	872	873	874	875	876	877	878
##	6	14	7	12	10	12	12	14	11	6
##	879	879.5	880	880.5	881	882	883	884	885	886
##	12	1	4	1	16	11	12	8	6	9

##	887	888	889	889.5	890	891	892	892.5	893	894
##	13	4	16	1	8	9	6	1	12	8
##	894.5	895	896	897	898	899	899.5	900	901	902
##	1	15	7	14	8	13	1	10	13	11
##	903	904	905	906	907	908	908.5	909	910	911
##	12	2	9	6	9	9	1	8	18	5
##	912	913	914	915	915.5	916	916.5	917	918	919
##	9	5	11	3	1	7	1	8	11	9
##	920	921	921.5	922	922.25	922.5	923	924	925	926
##	13	8	1	10	1	1	13	7	8	12
##	927	927.25	928	929	930	931	931.5	932	933	934
##	10	1	7	14	9	7	1	4	9	3
##	935	935.25	935.5	936	937	938	939	940	941	942
##	14	1	1	5	14	6	9	4	11	11
##	943	944	945	946	947	948	949	950	951	951.5
##	8	6	5	10	6	8	7	10	6	1
##	952	953	954	955	956	957	958	959	960	961
##	9	4	6	10	11	6	9	5	6	11
##	962	962.5	963	963.5	964	965	966	966.5	967	968
##	15	1	7	1	14	11	6	1	10	6
##	969	970	971	972	973	974	975	976	977	978
##	8	8	7	3	11	7	6	11	4	11
##	979	980	981	982	983	984	985	986	987	988
##	3	19	4	7	8	8	9	5	7	8
##	989	990	991	992	993	994	995	996	996.5	997
##	6	8	5	9	5	11	5	10	1	12
##	998	999	999.5	1000	1001	1002	1002.5	1003	1004	1004.5
##	5	6	2	8	12	13	1	7	3	1
##	1005	1006	1007	1008	1009	1010	1010.5	1011	1012	1013
##	2	11	8	9	9	4	1	5	8	7
##	1014	1015	1016	1016.5	1017	1018	1019	1020	1021	1022
##	12	5	6	2	9	10	11	2	4	11
##	1023	1024	1025	1026	1027	1028	1028.5	1029	1030	1031
##	8	6	6	5	6	8	1	8	6	8
##	1032	1033	1034	1035	1036	1037	1037.5	1038	1038.5	1039
##	5	5	5	7	4	6	1	6	1	3
##	1040	1041	1042	1043	1044	1045	1046	1047	1048	1049
##	12	5	9	5	9	6	4	8	2	7
##	1050	1051	1051.5	1052	1053	1054	1055	1055.5	1056	1057
##	5	10	1	6	4	5	1	1	8	3
##	1057.5	1058	1059	1060	1061	1062	1063	1064	1065	1065.5
##	2	2	4	7	10	4	7	7	4	1
##	1066	1067	1067.5	1068	1069	1070	1071	1072	1073	1074
##	4	7	1	5	3	12	7	6	5	5
##	1075	1076	1077	1077.5	1078	1078.5	1079	1080	1081	1082
##	6	1	4	1	4	1	7	2	2	6
##	1083	1083.75	1084	1085	1085.5	1086	1086.5	1087	1088	1088.5
##	8	1	2	11	1	3	2	3	10	1
##	1089	1090	1091	1092	1093	1093.5	1094	1095	1096	1097
##	5	7	5	6	6	1	4	6	3	2
##	1098	1099	1100	1101	1102	1103	1104	1105	1105.5	1106
##	10	8	5	7	3	5	1	9	1	4
##	1107	1108	1109	1110	1111	1112	1113	1114	1115	1116
##	2	9	5	8	3	4	4	4	7	6



##	1117	1117.5	1118	1119	1119.5	1120	1121	1122	1123	1124
##	6	1	6	5	1	9	12	7	11	6
##	1125	1126	1127	1128	1128.5	1129	1130	1131	1132	1133
##	5	6	3	9	1	3	1	6	2	8
##	1134	1135	1136	1137	1138	1139	1140	1141	1142	1143
##	11	4	3	5	7	3	7	7	5	8
##	1144	1145	1145.25	1146	1147	1148	1149	1150	1151	1152
##	9	2	1	2	7	2	6	7	1	6
##	1153	1154	1155	1156	1157	1158	1159	1160	1161	1162
##	9	6	6	8	6	5	6	5	2	5
##	1163	1164	1165	1166	1167	1168	1168.5	1169	1170	1171
##	5	4	5	4	3	6	1	8	2	5
##	1172	1173	1174	1175	1176	1177	1178	1179	1180	1181
##	4	1	4	2	10	6	5	6	1	4
##	1182	1183	1184	1185	1186	1187	1188	1189	1190	1191
##	5	2	11	7	3	6	4	6	4	4
##	1192	1193	1194	1195	1196	1197	1198	1199	1200	1201
##	3	8	7	2	8	5	3	1	4	6
##	1202	1203	1204	1205	1206	1207	1208	1209	1210	1211
##	7	7	1	8	4	2	5	3	6	5
##	1212	1213	1214	1215	1216	1217	1218	1219	1220	1221
##	5	5	5	2	3	4	1	4	2	1
##	1222	1223	1225	1226	1227	1228	1229	1230	1231	1232
##	5	3	3	4	2	6	4	3	4	1
##	1233	1234	1235	1236	1237	1237.5	1238	1239	1240	1241
##	2	5	2	2	7	1	3	3	2	3
##	1241.5	1242	1243	1244	1245	1246	1247	1248	1249	1250
##	1	3	4	2	3	1	3	6	5	4
##	1251	1252	1253	1254	1255	1257	1258	1259	1260	1261
##	7	2	2	4	2	5	3	4	1	4
##	1262	1263	1264	1265	1266	1267	1268	1269	1270	1271
##	11	4	1	5	2	7	5	4	3	2
##	1271.5	1272	1272.5	1273	1273.5	1274	1275	1276	1276.5	1277
##	1	3	1	4	1	5	4	4	1	5
##	1278	1279	1280	1281	1282	1283	1284	1284.5	1284.75	1285
##	2	3	3	2	4	3	1	2	1	1
##	1286	1287	1288	1289	1290	1291	1292	1293	1294	1295
##	6	4	2	7	2	3	4	5	1	2
##	1296	1297	1298	1299	1300	1301	1302	1303	1304	1305
##	4	3	4	2	2	8	3	2	3	2
##	1306	1307	1308	1309	1310	1310.5	1311	1312	1312.5	1313
##	1	4	2	5	4	1	1	8	1	3
##	1314	1314.5	1316	1317	1318	1319	1320	1320.5	1321	1322
##	3	1	1	2	2	3	3	1	6	5
##	1323	1324	1325	1325.5	1326	1327	1328	1329	1330	1331
##	1	2	2	1	3	4	2	3	3	3
##	1331.5	1332	1333	1334	1335	1336	1337	1338	1339	1340
##	1	5	3	3	1	4	1	3	1	2
##	1340.75	1341	1342	1343	1344	1345	1346	1347	1348	1348.25
##	1	3	1	1	4	1	5	5	1	1
##	1349	1350	1352	1353	1355	1356	1357	1358	1359	1360
##	1	5	1	3	3	1	3	1	3	2
##	1361	1362	1363	1364	1365	1366	1367	1368	1369	1371
##	5	2	2	7	5	7	3	1	5	2

##	1372	1373	1373.5	1374	1375	1376	1377	1379	1380	1381
##	3	3	1	2	1	2	2	5	1	2
##	1382	1383	1384	1386	1387	1389	1390	1391	1392	1393
##	2	5	3	3	1	4	4	3	4	2
##	1394	1394.5	1395	1396	1397	1398	1399	1401	1402	1403
##	2	1	5	4	1	8	1	6	1	4
##	1404	1405	1406	1408	1409	1410	1411	1412	1413	1414
##	1	3	2	3	1	2	1	2	2	4
##	1415	1416	1417	1418	1418.5	1419	1420	1420.5	1421	1422
##	4	3	3	4	1	1	2	1	3	2
##	1424	1425	1426	1427	1427.5	1428	1429	1430	1431.5	1432
##	2	1	2	3	1	1	2	3	1	3
##	1433	1434	1434.5	1435	1436	1437	1438	1439	1440	1441
##	1	6	2	1	1	2	8	4	2	1
##	1442	1444	1446	1447	1448	1449	1450	1450.5	1451	1451.5
##	1	3	5	1	1	1	2	1	4	2
##	1453	1454	1455	1456	1457	1458	1459	1460	1461	1462
##	3	1	2	1	1	5	3	1	6	2
##	1463	1465	1466	1467	1467.5	1468	1469	1470	1471	1472
##	3	2	1	4	1	4	2	1	5	3
##	1473	1474.5	1475	1476	1478	1479	1482	1483	1484	1485
##	4	1	3	2	5	2	6	2	1	5
##	1486	1486.5	1487	1488	1489	1491	1492	1493	1494	1494.5
##	1	1	4	2	3	1	3	1	1	1
##	1495	1496	1497	1499	1500	1501	1502	1503	1504	1506
##	1	2	2	1	1	1	2	4	5	7
##	1507	1508	1509	1510	1511	1512	1513	1514	1515	1516
##	1	2	3	2	2	2	2	2	1	2
##	1516.5	1517	1518	1519	1520	1521	1522	1523	1524.5	1525
##	1	2	1	1	3	3	4	5	1	2
##	1528	1528.5	1529	1531	1532	1533	1534	1535	1536	1537
##	1	1	2	1	1	2	8	1	4	3
##	1538	1539	1540	1541	1542	1543	1544	1545	1546	1546.5
##	3	1	2	2	1	1	2	2	4	1
##	1547	1548	1549	1550	1552	1553	1553.5	1554	1555	1556
##	2	1	3	1	3	2	1	1	2	4
##	1559	1560	1562	1563	1564	1565	1566	1567	1568	1570
##	2	2	3	2	2	3	3	1	1	2
##	1570.5	1572	1573	1574	1576	1578	1579	1581	1582	1583
##	1	2	3	1	1	1	1	2	1	5
##	1585	1587	1589	1590	1594	1595	1596	1597	1598	1599
##	2	3	2	3	1	1	1	7	2	4
##	1600	1601	1602	1602.25	1605	1606	1608	1609	1613	1614
##	1	1	1	1	1	4	3	1	1	2
##	1615	1616	1617	1618	1619	1620	1621	1623	1624	1627
##	1	2	1	1	2	3	1	4	2	3
##	1628	1629	1630	1631	1632	1633	1634	1634.5	1635	1638
##	1	2	1	1	1	1	1	1	3	1
##	1640	1641	1642	1643	1644	1645	1646	1647	1647.75	1648
##	3	2	2	3	1	2	3	2	1	1
##	1649	1650	1652	1653	1654	1654.5	1656	1657	1658	1659
##	1	2	2	2	4	1	3	3	3	2
##	1660	1664	1666	1667	1668	1669	1670	1672	1673	1674
##	1	2	1	1	2	1	2	3	4	1

##	1675	1676	1678	1679	1681	1683	1684	1685	1686	1688
##	3	4	2	1	2	3	4	3	3	1
##	1689	1690	1691	1693	1693.25	1694	1696	1697	1697.5	1698
##	4	2	2	1	1	1	1	1	1	2
##	1699	1700	1701	1703	1704	1705	1706	1708	1709	1712
##	2	1	2	1	2	3	2	1	1	1
##	1713	1714	1715	1717	1717.5	1718	1719	1721	1725	1726
##	2	1	1	2	1	2	2	4	2	1
##	1727	1729	1730	1731	1732	1733	1734	1735	1736	1737
##	2	3	3	1	2	2	5	1	1	2
##	1738	1741	1742	1743	1744	1745	1746	1747	1750	1753
##	4	1	3	1	1	1	3	1	4	1
##	1755	1758	1760	1761	1762	1763	1764	1765	1766	1767
##	1	3	1	2	1	1	2	1	5	1
##	1768	1770	1772	1773	1775	1778	1779	1782	1783	1783.5
##	1	2	1	2	5	1	4	2	3	1
##	1784	1785	1786	1787.5	1790	1791	1792	1793	1794	1795
##	5	1	3	1	3	1	2	1	2	1
##	1796	1798	1800	1801	1802	1803	1804	1806	1807	1809
##	1	1	2	4	2	1	5	4	3	1
##	1811	1812	1813	1814	1815	1816	1816.75	1817	1818	1819
##	2	1	1	2	2	2	1	2	2	1
##	1820	1821	1822	1823	1826	1827	1829	1830	1831	1832
##	1	1	4	2	1	3	1	1	1	1
##	1834	1835	1836	1839	1840	1840.5	1840.75	1841	1844	1845
##	2	1	3	1	3	1	1	1	4	2
##	1846	1847	1849	1850	1851	1853	1854	1856	1857	1858
##	1	2	1	3	1	1	3	1	2	1
##	1859.5	1860	1860.75	1861	1862	1863	1864	1865	1866	1868
##	1	1	1	1	1	3	1	1	1	3
##	1870	1871	1874	1875	1876	1876.5	1878	1882	1883	1885
##	1	3	2	1	1	1	2	1	1	1
##	1885.5	1887	1890	1891	1892	1894	1896	1898	1899	1902
##	1	2	1	1	2	1	1	1	1	4
##	1905	1907	1909	1910	1911	1912	1913	1917	1918	1920
##	2	2	4	1	1	2	2	1	2	1
##	1922	1923	1924	1925	1926	1927	1928	1930	1931	1932
##	1	1	3	1	1	2	1	2	2	1
##	1934	1935	1938	1941	1942	1943	1945	1947	1949	1951
##	1	3	1	1	1	1	2	1	2	4
##	1952	1954	1955	1959	1960	1962	1963	1966	1967	1968
##	1	1	1	1	2	1	2	1	1	2
##	1969	1970	1972	1976	1978	1978.5	1979	1980	1981	1982
##	4	1	1	1	1	1	2	1	4	2
##	1984	1985	1988	1989	1991	1992	1994	1995	1996	1997
##	1	1	1	2	1	1	1	1	2	2
##	1998	2001	2004	2005	2006	2008	2009	2010	2011	2015
##	1	2	3	2	1	2	1	2	1	2
##	2017	2017.5	2019	2021	2029	2030	2032	2033	2034	2035
##	1	1	2	2	2	1	5	1	1	4
##	2036	2038	2040	2041	2042	2044	2045	2046	2047	2048
##	4	2	1	1	1	1	3	3	1	1
##	2049	2050	2051	2052	2054	2057	2059	2060	2062	2063
##	1	1	2	1	5	2	2	1	1	1

##	2064	2065	2067	2069	2070	2072	2074	2075	2076	2077
##	3	1	1	1	1	1	3	1	1	2
##	2081	2084	2085	2086	2088	2089	2091	2094	2095	2097
##	1	3	2	1	2	2	1	4	1	2
##	2098	2099	2100.5	2103	2107	2107.5	2108	2112	2114	2115
##	1	1	1	1	1	1	2	2	1	1
##	2118	2120	2121	2124	2126	2128	2129	2130	2133	2135.5
##	4	4	1	1	1	3	1	4	1	1
##	2138	2140	2143	2144	2145	2146	2147	2152	2154	2155
##	1	2	1	2	1	1	1	1	2	1
##	2160	2161	2162	2163.5	2165	2166	2168	2169	2170	2173
##	1	1	2	1	1	2	2	1	1	1
##	2175	2178	2180	2181	2182	2183	2186	2187	2188	2189
##	1	1	2	2	2	1	1	1	3	1
##	2192	2192.5	2194	2197	2198	2202	2203	2204.5	2208	2210
##	1	1	2	3	1	4	1	1	2	2
##	2213	2215	2217	2218	2221	2225	2226	2228	2235	2239
##	3	3	1	1	2	1	2	1	3	1
##	2240	2241	2244	2246	2252	2253	2254	2255	2257	2260.5
##	2	1	1	1	1	1	1	2	2	1
##	2262	2266	2267	2268	2269	2270	2271	2271.5	2272.5	2274
##	1	2	1	3	1	3	1	1	1	1
##	2276	2279	2280	2281	2282	2283	2284.75	2285	2286	2288
##	3	3	1	2	1	1	1	1	1	2
##	2290	2292	2297	2298	2302	2303	2304	2305	2307	2308
##	3	2	1	2	1	1	1	1	1	2
##	2309	2309.5	2314	2318	2320	2323	2324	2324.5	2326	2330
##	1	1	1	1	1	1	1	1	1	1
##	2330.5	2332	2333	2334	2335	2342	2343	2345	2346	2351
##	1	2	1	1	1	1	1	2	1	1
##	2354	2356	2357	2359	2361	2364	2366.75	2371	2372	2374
##	1	1	2	3	1	1	1	1	1	1
##	2376	2377	2379	2382	2384	2385	2393	2394	2395	2396
##	1	2	2	2	1	1	1	1	2	2
##	2400	2402	2407	2408	2414	2417	2419	2423	2425	2428
##	2	2	2	1	1	1	1	1	1	1
##	2430	2432	2434	2435	2436	2447	2449	2456	2459	2460
##	1	1	1	1	1	1	1	1	4	2
##	2464	2465	2466	2467	2470	2471	2473	2474	2479	2482
##	1	1	2	1	1	2	1	1	2	2
##	2487	2489	2493	2497	2498	2499	2502	2504	2508	2509
##	1	1	1	2	1	1	1	1	2	1
##	2518	2521	2522	2527	2528	2533	2534	2535	2540	2541
##	1	2	1	1	1	1	1	1	1	2
##	2542	2545	2545.25	2551	2554	2558	2567	2569	2570	2574
##	1	1	1	1	2	2	1	1	1	1
##	2579	2581	2584	2586	2590	2592	2593	2595.5	2596	2602
##	3	1	1	1	1	1	1	1	1	1
##	2603	2604	2605	2606	2609	2611	2618	2619	2621	2622
##	1	2	1	1	1	1	1	1	1	1
##	2624	2625.5	2627	2632	2637	2638	2641	2645	2648	2656
##	1	1	1	1	1	1	1	1	1	1
##	2660	2670	2673	2675	2676	2677	2678	2688	2689	2690
##	3	1	1	1	1	1	1	2	1	1

##	2696	2698	2700	2703	2709	2713	2714	2716	2718	2720
##	1	1	1	2	1	1	1	1	1	1
##	2721	2722	2723	2726	2729	2736	2737	2739	2740	2743
##	1	1	2	1	2	1	1	1	2	1
##	2751	2760	2764	2765	2767	2778	2784	2788	2791	2794
##	1	2	1	1	2	1	2	1	1	2
##	2801	2802	2804	2808	2823	2824	2832	2834.75	2838	2843
##	2	1	3	2	1	1	1	1	1	1
##	2846	2851	2854	2857	2859	2860	2861	2864	2866	2868
##	1	1	1	2	1	1	1	2	2	1
##	2870	2875	2881	2884	2885	2890	2892	2895	2896	2897
##	1	1	2	1	2	2	1	1	1	1
##	2898	2906	2911	2914	2920	2924	2933	2934	2939	2944
##	1	1	1	1	1	1	1	1	1	1
##	2946	2947	2957	2962	2970	2974	2976	2989	2993	2994
##	1	1	1	1	1	1	1	1	1	1
##	2996	3001	3002	3008	3010	3016	3019	3020	3021	3023
##	1	1	1	2	1	1	1	1	1	1
##	3025	3029	3035	3041	3042	3046	3050	3051	3061	3062
##	1	1	1	1	1	1	1	1	1	1
##	3068	3069	3074	3076	3087	3089	3095	3098	3104	3105
##	3	1	1	1	2	1	1	2	1	1
##	3110	3120	3123.25	3131	3143	3145	3151	3158	3161	3166
##	1	1	1	1	1	1	1	1	1	1
##	3170	3175	3182	3183	3184	3191	3207	3208	3209	3215
##	1	1	1	1	1	2	2	1	1	1
##	3216	3222	3230	3231	3236	3238	3247	3248	3249	3255
##	1	1	1	1	1	1	1	1	1	2
##	3264	3268	3279	3280	3295	3297	3303	3304	3309	3326
##	1	1	2	1	4	1	1	1	2	1
##	3333	3338	3340	3342	3352	3355	3357	3361	3369.75	3375
##	2	1	1	1	1	1	1	1	1	1
##	3377	3383	3391	3392	3400	3402	3416	3417	3425	3426
##	2	1	1	1	1	2	1	1	1	1
##	3437	3438	3445	3448	3453	3454	3459	3459.5	3462	3468
##	1	1	1	1	1	1	1	1	1	1
##	3470	3474	3479	3480	3485	3488	3490	3502	3505	3516
##	1	1	2	1	1	1	1	2	1	1
##	3519	3537	3538	3542	3545	3548	3550	3552	3555	3559
##	1	1	1	1	1	1	1	2	1	1
##	3573	3574	3575	3583	3586	3593	3594	3601	3602	3603
##	1	1	1	1	1	2	1	1	1	1
##	3608	3612	3617	3619	3624	3627.75	3628	3632	3635	3637
##	1	1	1	1	1	1	1	1	2	1
##	3644	3646	3680	3682	3683	3692	3708	3710	3715	3718
##	1	1	1	1	1	1	1	1	1	1
##	3723.5	3726	3728	3735	3738	3740	3745	3746	3755	3756
##	1	1	1	1	1	1	1	1	1	1
##	3762	3765	3774	3774.5	3781	3784	3786	3787	3791	3806
##	1	1	1	1	1	1	1	1	1	1
##	3812	3820	3831	3833	3849	3858.5	3862	3865.75	3888	3898
##	2	1	2	1	1	1	1	1	1	1
##	3908	3910.5	3915	3921	3925	3927	3930	3933	3935	3952
##	1	1	1	1	2	1	1	1	2	1

##	3954	3966	3974	3978	3979.75	3980	3984	4000	4001	4002
##	1	1	1	1	1	1	1	1	1	1
##	4004	4012	4017	4036	4044	4053	4059	4060	4066	4075
##	2	1	1	1	1	1	1	1	1	2
##	4083	4084	4093	4094	4101	4115	4116	4125	4135	4146
##	1	2	1	1	1	1	1	1	1	2
##	4148	4156	4166	4167	4172	4177	4208	4224	4228	4236
##	1	1	1	1	1	2	1	1	1	1
##	4239	4250	4278	4282	4293	4300	4301	4303	4308	4314
##	1	1	1	2	1	1	1	1	1	1
##	4315	4321	4325	4332	4357	4398	4401	4406	4408	4411
##	1	1	1	1	1	1	1	1	1	1
##	4422	4433	4444	4461	4480	4489	4495	4499	4515	4524
##	1	1	1	1	1	1	1	1	1	1
##	4526	4526.5	4532	4567	4590	4592	4604	4613	4636	4643
##	1	1	1	1	1	1	1	1	1	1
##	4659	4694	4704	4709	4711	4717	4727	4776	4824	4825
##	2	1	1	1	1	1	1	1	1	1
##	4836	4841	4874	4894	4895	4926	4927	4945	4984	4987
##	1	1	1	1	1	1	1	1	1	1
##	4998	5007	5017	5045	5056	5057	5058	5070	5076.75	5082
##	1	1	1	1	1	1	2	3	1	1
##	5097	5111	5159	5201.5	5209	5220.5	5242	5255	5263	5275
##	1	1	1	1	1	1	1	1	1	1
##	5360	5374	5384	5387	5430.5	5461	5465	5468.5	5486.5	5512
##	1	1	1	1	1	1	1	1	1	1
##	5636	5652.5	5665	5669	5718	5728	5814	5837	5846	5919.5
##	1	1	1	1	1	1	1	1	1	1
##	5964	5975	5996.5	6008.5	6069	6081	6089	6121.5	6130	6161
##	1	1	1	1	1	1	1	1	1	1
##	6175	6178.5	6220.5	6284	6315	6337	6378	6380	6396	6399
##	1	1	1	1	1	1	1	1	1	1
##	6441	6467.5	6524	6545	6658	6684.5	6690	6720	6729	6734
##	1	1	1	1	1	1	1	1	1	1
##	6787.5	6809	6831	7396	7410	7559.5	7577	7681.75	7824	7831.5
##	1	1	1	1	1	1	1	1	1	1
##	7950.5	8083	8120.25	8125	8158.5	8479	8896	8917	8987.75	9000
##	1	1	1	1	1	1	1	1	1	1
##	9015.5	9028.5	9065	9133	9536	9688.5	9710	9849.5	9874	9991.25
##	1	1	1	1	1	1	1	1	1	1
##	10258	10295	10484	11300	11461.5	11697	11826.5	11925.5	12006	12361
##	1	1	1	1	1	1	1	1	1	1
##	12431.5	12438	12667.5	12792	12895	13151.2	13176	13270.5	13283.8	13581.5
##	1	1	1	1	1	1	1	1	1	1
##	13750.5	13771.5	13776	13889	14202.5	14267.5	14322.5	14339	14456	15094
##	1	1	1	1	1	1	1	1	1	1
##	15108.5	15340.5	15486.2	15675	15699	17483.5				
##	1	1	1	1	1	1				

Create categorical variables for analysis Variable are already factored, don't need this section

```

flu$Gender <- factor(flu$gender)

flu$Vaccine <- factor(flu$vaccine)

flu$Vaccine_Response <- factor(flu$vaccine_response)

flu$Influenza_Infection <- factor(flu$influenza_infection_history)

flu$Influenza_Hospitalization <- factor(flu$influenza_hospitalization)

flu$subset <- factor(flu$subset)

head(flu)

```

	donor_id <int>	study_id <int>	gen... <fctr>	race <fctr>	visit_id <int>	visit_year <int>	visit_day <int>	visit_type_hai <fctr>	visit
1	813	15	Female	Caucasian	2937	2014	0	pre	
2	813	15	Female	Caucasian	2937	2014	0	pre	
3	813	15	Female	Caucasian	2937	2014	0	pre	
4	813	15	Female	Caucasian	2937	2014	0	pre	
5	813	15	Female	Caucasian	2937	2014	0	pre	
6	813	15	Female	Caucasian	2937	2014	0	pre	

6 rows | 1-10 of 45 columns

### Import data from file

```

#Import file
result <- read.csv(file = "C:/Users/Christine/Documents/Bellevue/DSC 680/Project 1/result_flu.csv", header = TRUE, na.strings = " ")

```

### Display first five records of file

```
head(result)
```

	donor_id <int>	gen... <fctr>	race <fctr>	visit_id <int>	visit_year <int>	visit_day <int>	visit_age <dbl>	cmv_status <dbl>	ebv_stat <dbl>
1	813	Female	Caucasian	2937	2014	0	23	0	
2	812	Male	Caucasian	2936	2014	0	28	1	
3	811	Male	Caucasian	2935	2014	0	23	0	
4	810	Male	Caucasian	2934	2014	0	27	1	
5	809	Female	Asian	2933	2014	0	27	1	

donor_id	gen...	race	visit_id	visit_year	visit_day	visit_age	cmv_status	ebv_stat
<int>	<fctr>	<fctr>	<int>	<int>	<int>	<dbl>	<dbl>	<dbl>
6	808	Female	Other	2932	2014	0	29	0

6 rows | 1-10 of 16 columns

Create categorical variables for analysis Variable are already factored, don't need this section

```
result$Gender <- factor(result$gender)

result$Race <- factor(result$race)

result$Vaccine <- factor(result$vaccine)

result$Vaccine_Response <- factor(result$vaccine_response)

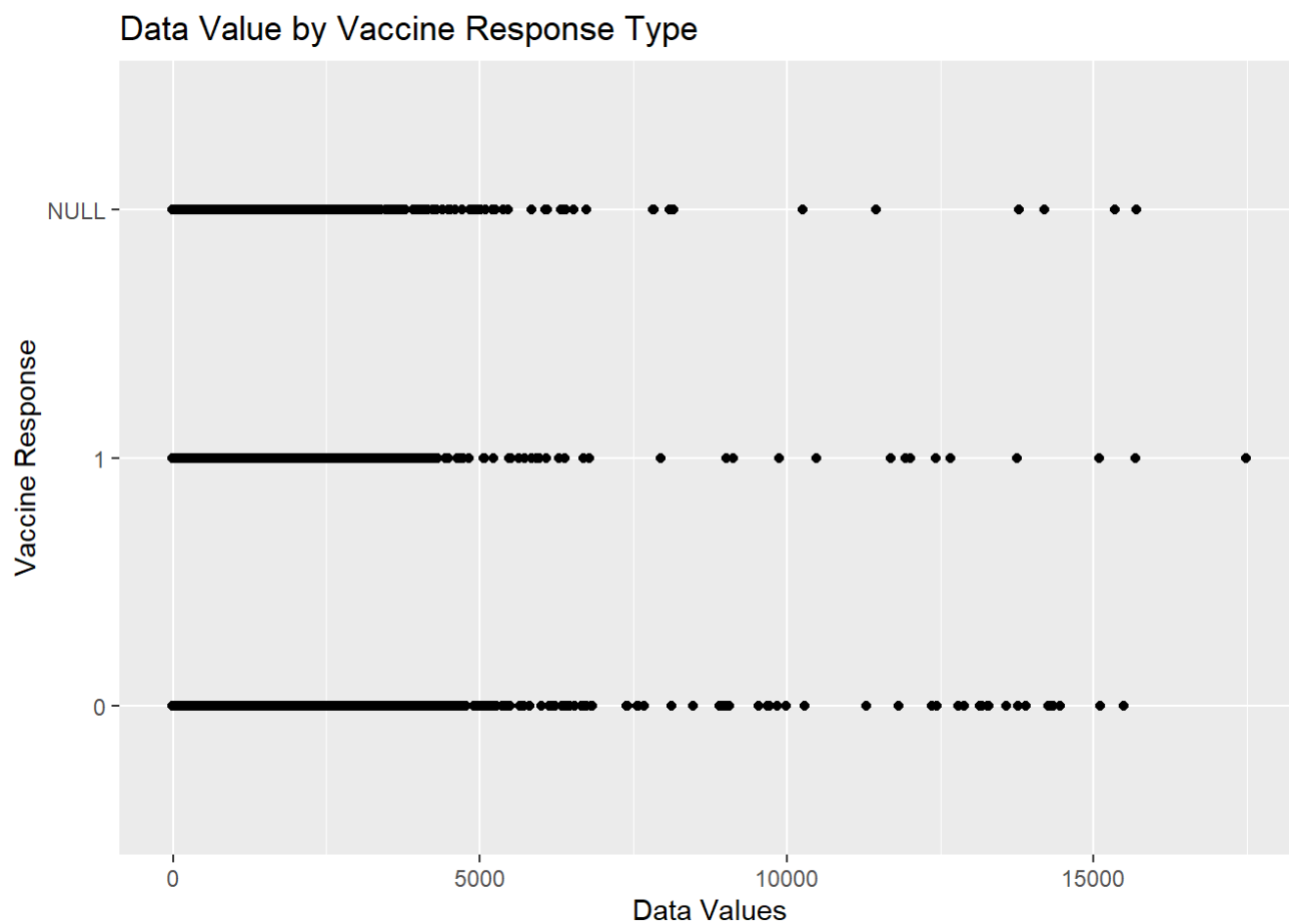
result$Influenza_Infection <- factor(result$influenza_infection_history)

result$Influenza_Hospitalization <- factor(result$influenza_hospitalization)
```

Scatterplot

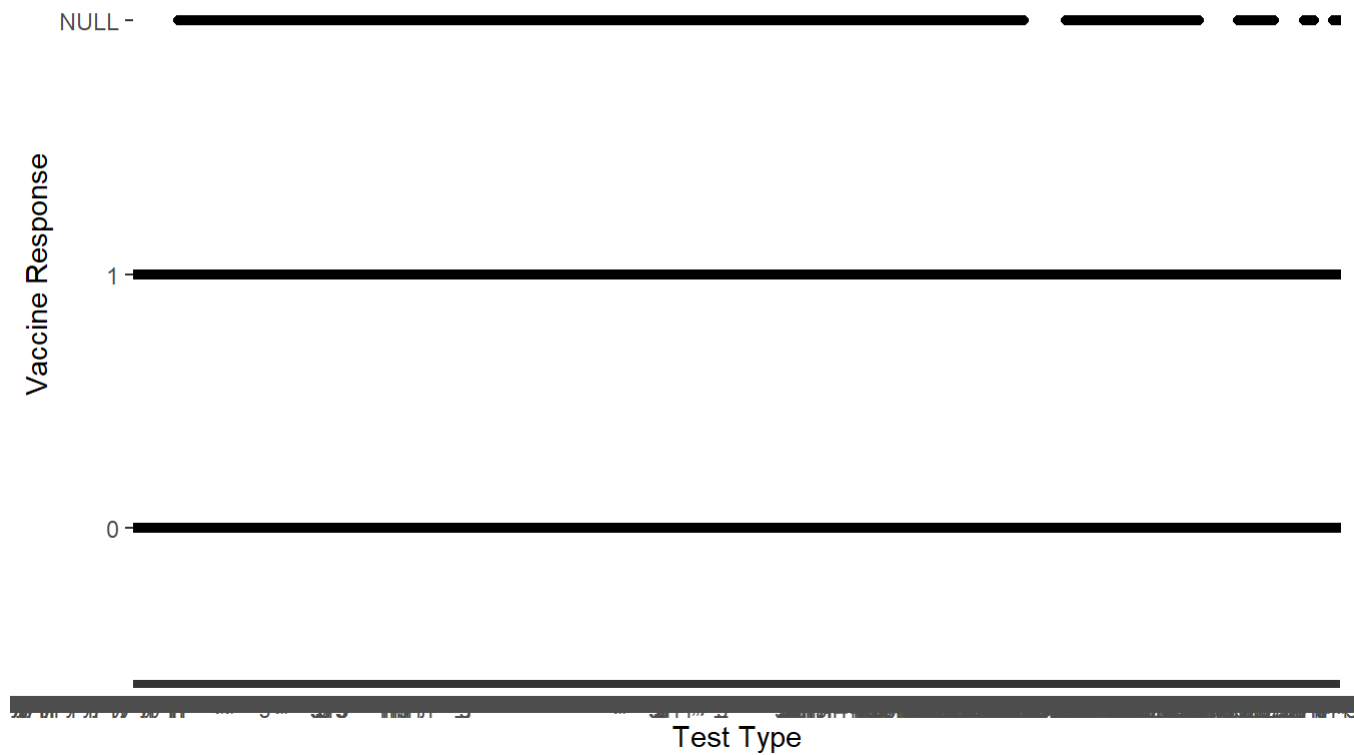
```
ggplot(flu, aes(x=data, y=vaccine_response)) + geom_point() + ggtitle("Data Value by Vaccine Res
ponse Type") +
  xlab("Data Values") + ylab("Vaccine Response")
```





```
ggplot(flu, aes(x=subset, y=vaccine_response)) + geom_point()+ ggtitle("Test Types by Vaccine Response Type") +  
  xlab("Test Type") + ylab("Vaccine Response")
```

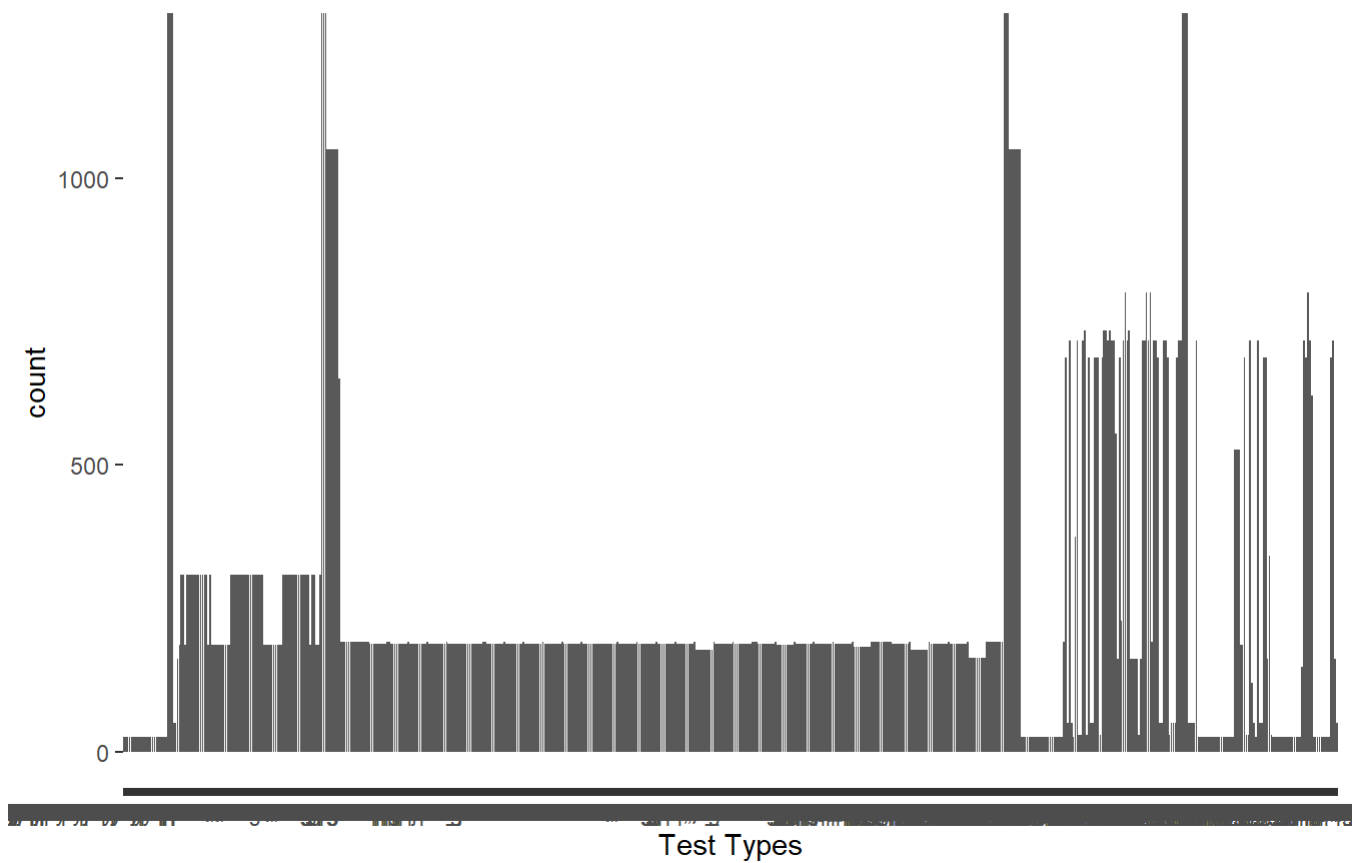
## Test Types by Vaccine Response Type



### Histograms

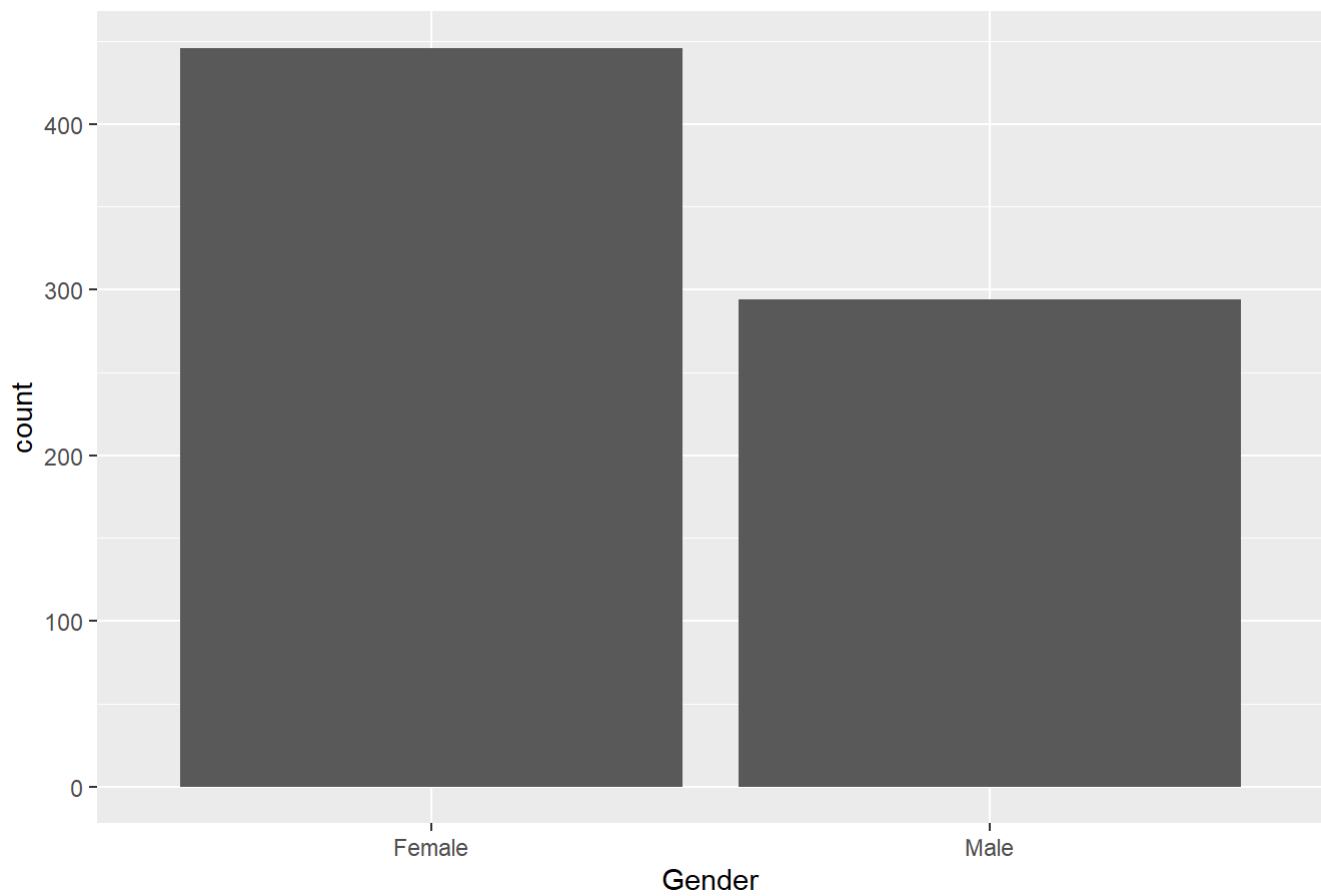
```
ggplot(flu, aes(x=subset)) + geom_bar()+ ggtitle("Count of Various Test Types") +  
  xlab("Test Types")
```

## Count of Various Test Types



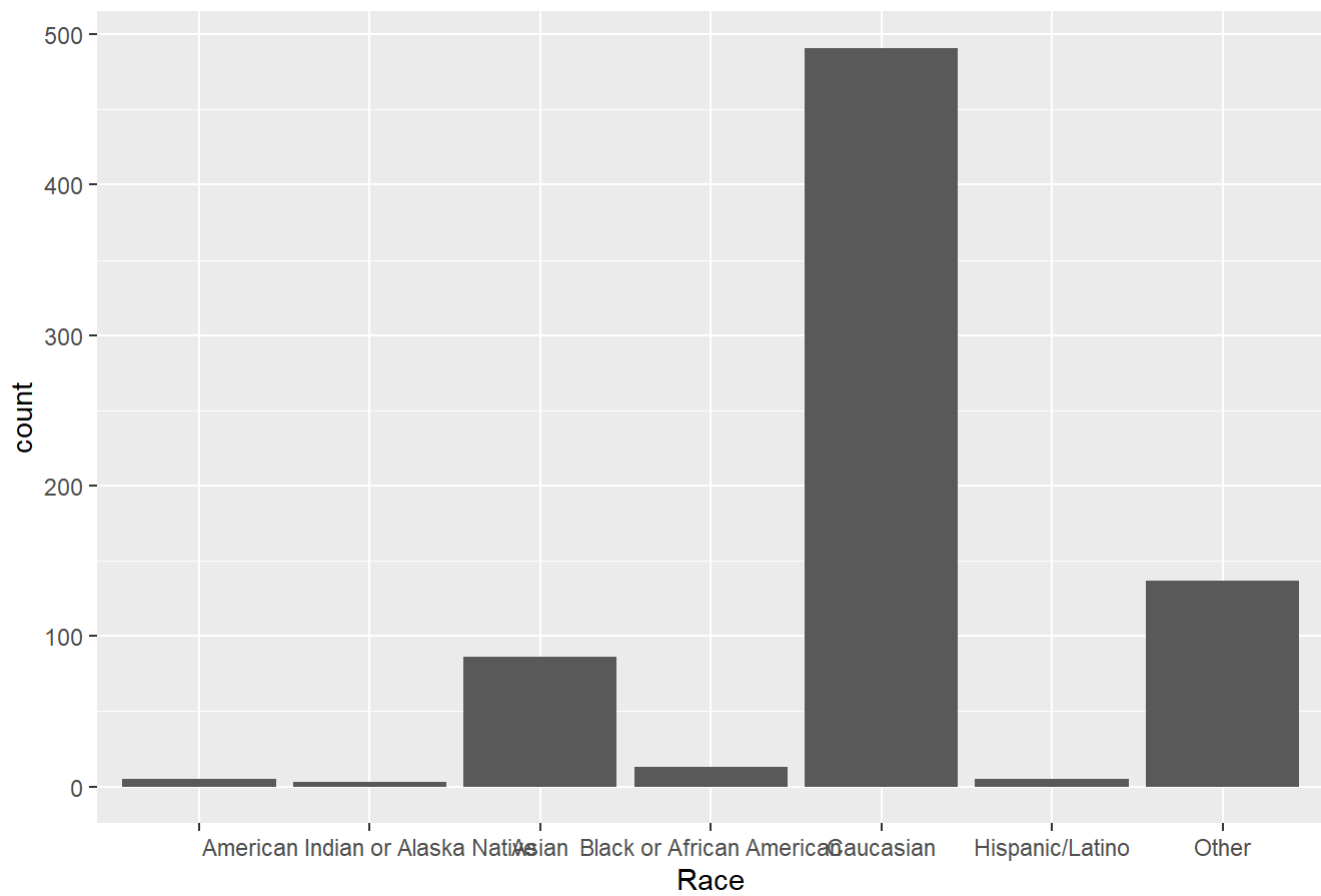
```
ggplot(result, aes(x=Gender)) + geom_bar()+ ggtitle("Donors by Gender") +  
  xlab("Gender")
```

## Donors by Gender



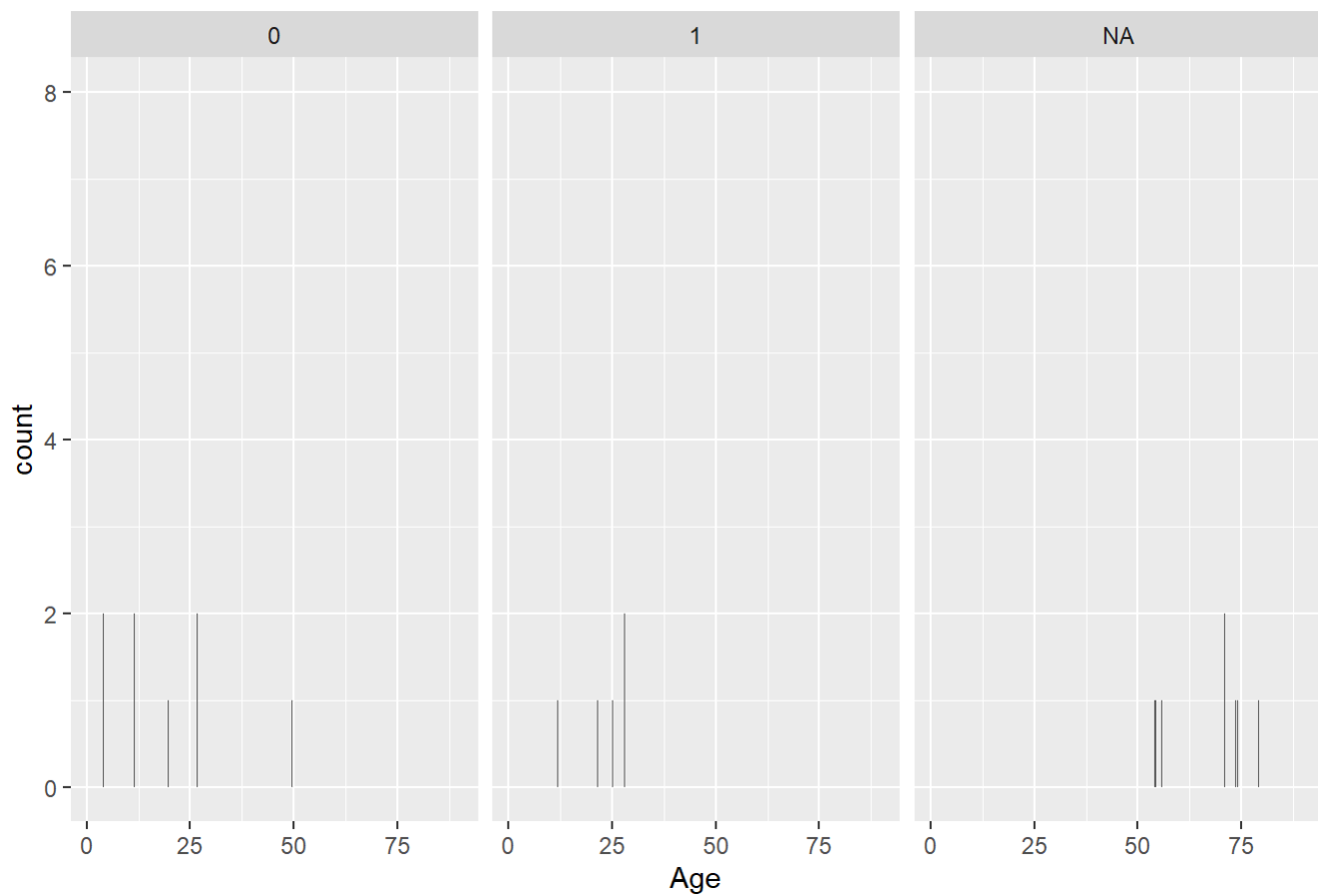
```
ggplot(result, aes(x=Race)) + geom_bar()+ ggtitle("Donors by Race") +  
  xlab("Race")
```

Donors by Race



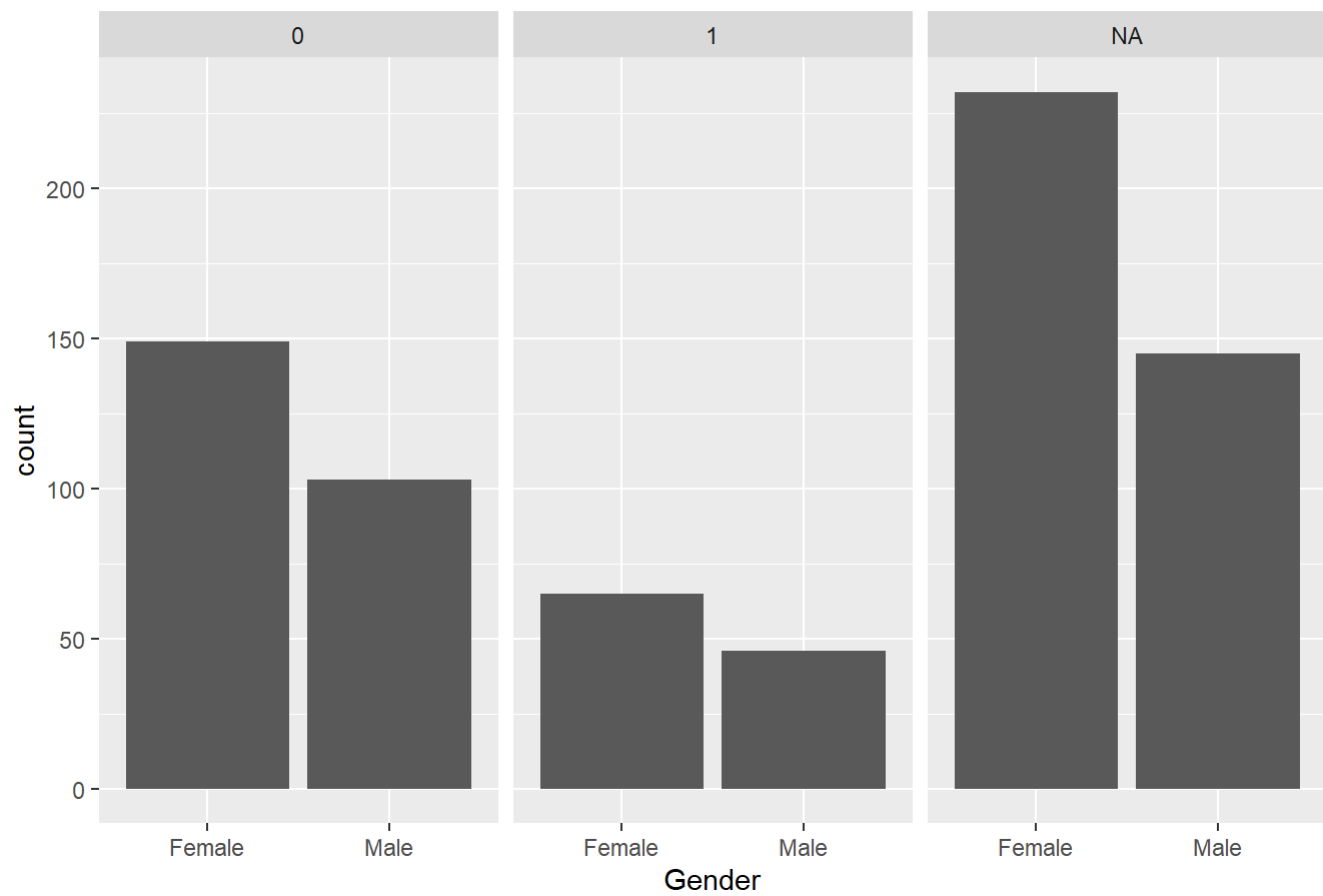
```
ggplot(result, aes(x=visit_age)) + geom_bar()+facet_wrap(~Vaccine_Response) + ggtitle("Donor Age  
by Vaccine Response") +  
  xlab("Age")
```

## Donor Age by Vaccine Response



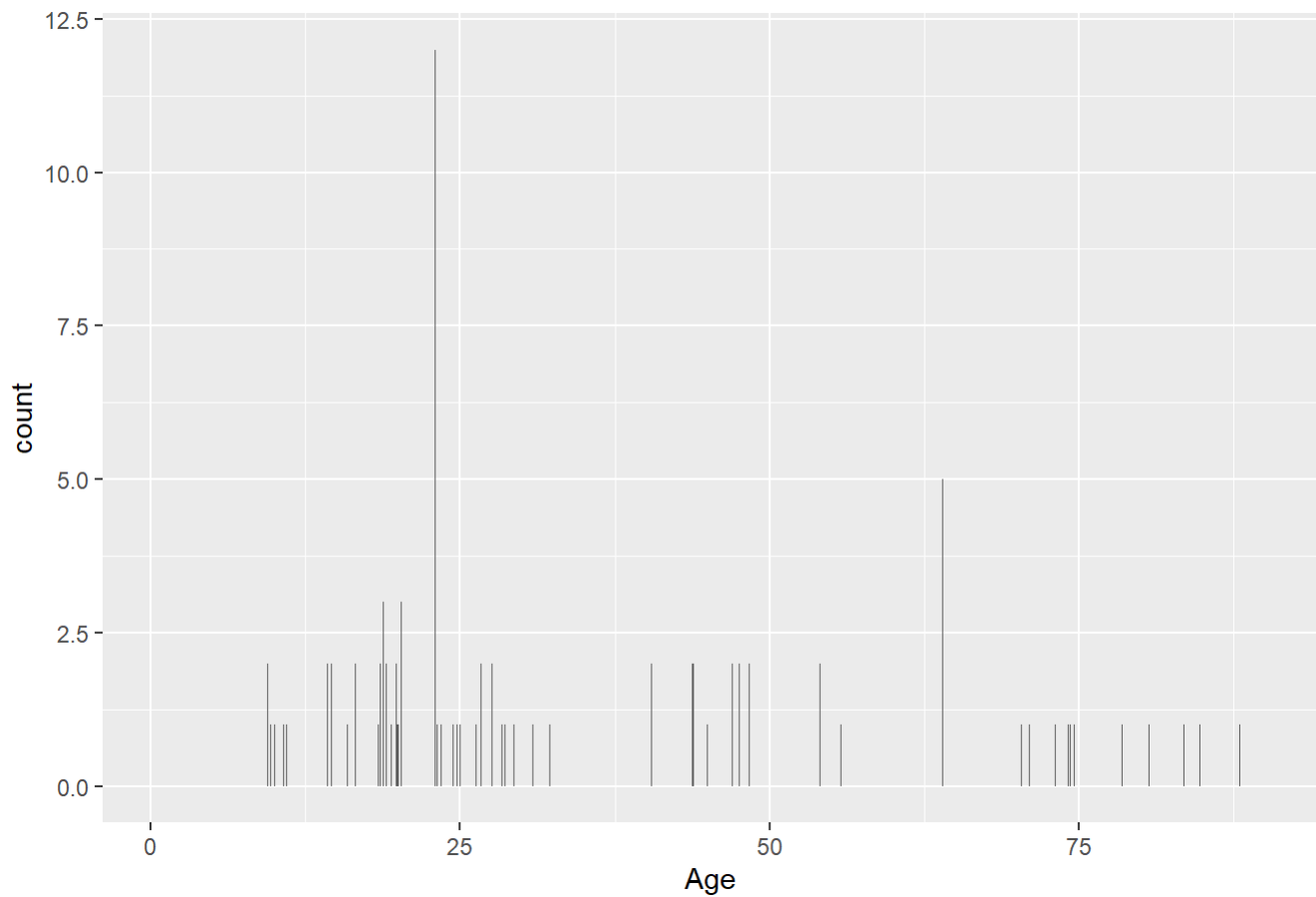
```
ggplot(result, aes(x=Gender)) + geom_bar()+facet_wrap(~Vaccine_Response)+ ggtitle("Gender by Vaccine Response Type") +  
  xlab("Gender")
```

## Gender by Vaccine Response Type



```
ggplot(result, aes(x=visit_age)) + geom_bar() + ggtitle("Age Distribution") +  
  xlab("Age")
```

## Age Distribution

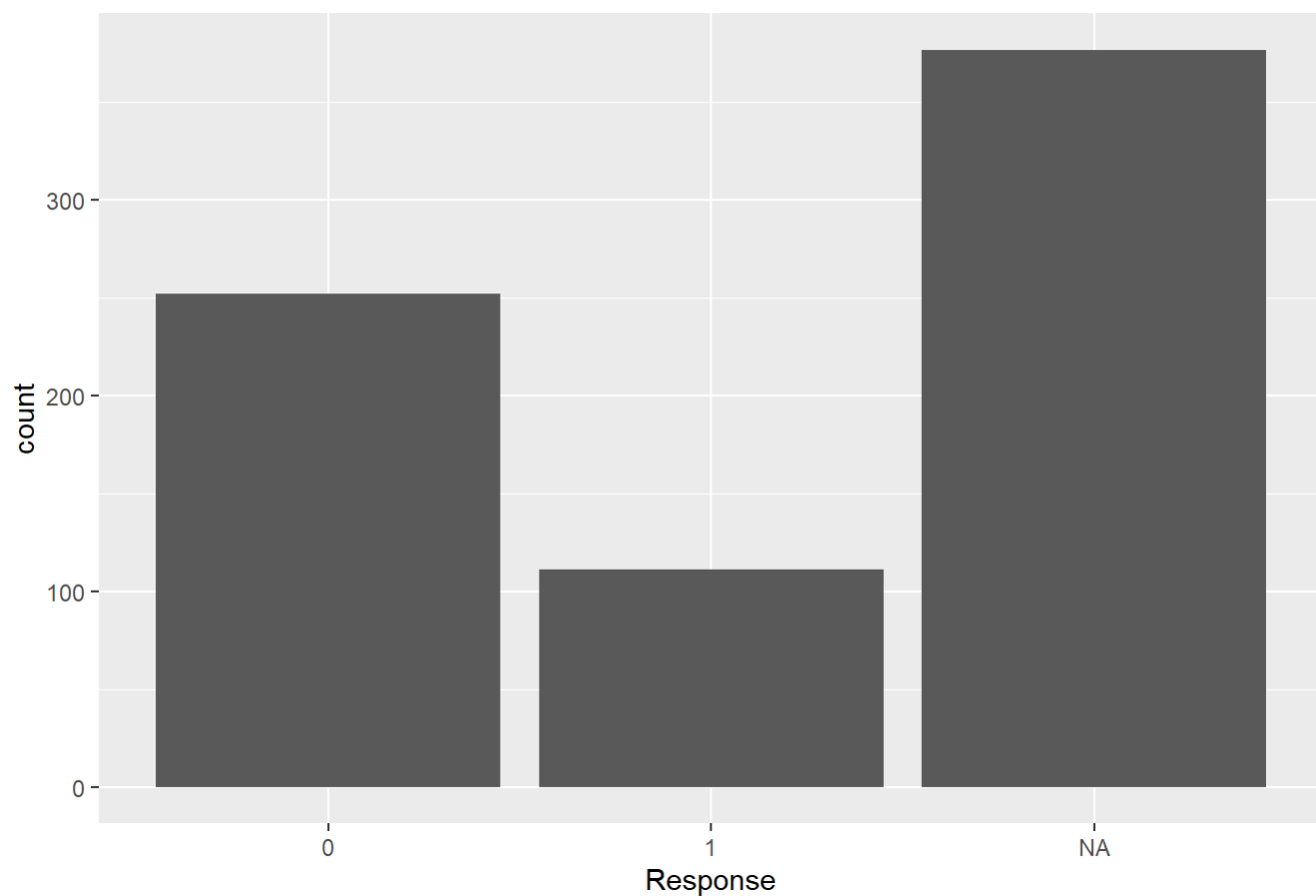


## Histograms

```
ggplot(result, aes(x=Vaccine_Response)) + geom_bar() + ggtitle("Vaccine Response Distribution")  
+  
  xlab("Response")
```



## Vaccine Response Distribution



Create regression model for influenza infection based on vaccine type

```
model1 <- glm(Influenza_Infection~Vaccine+Gender+visit_age+Race, data = result, family = "binomial")
```

Output model results

```
summary(model1)
```

```
##
## Call:
## glm(formula = Influenza_Infection ~ Vaccine + Gender + visit_age +
##      Race, family = "binomial", data = result)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -0.6409  -0.4781  -0.4585  -0.4360   2.4153
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)    -1.660e+01  1.697e+03  -0.010    0.992
## Vaccine2         1.843e-01  8.296e-01   0.222    0.824
## Vaccine3        -1.441e+01  1.385e+03  -0.010    0.992
## Vaccine4        -1.845e-02  3.989e-01  -0.046    0.963
## Vaccine5        -1.437e+01  1.385e+03  -0.010    0.992
## Vaccine6         5.951e-01  5.898e-01   1.009    0.313
## GenderMale      -1.341e-01  2.762e-01  -0.486    0.627
## visit_age        1.118e-03  6.789e-03   0.165    0.869
## RaceAmerican Indian or Alaska Native -1.902e-01  2.185e+03   0.000    1.000
## RaceAsian         1.386e+01  1.697e+03   0.008    0.993
## RaceBlack or African American    1.439e+01  1.697e+03   0.008    0.993
## RaceCaucasian     1.442e+01  1.697e+03   0.009    0.993
## RaceHispanic/Latino  -9.349e-03  2.189e+03   0.000    1.000
## RaceOther         1.449e+01  1.697e+03   0.009    0.993
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 407.27  on 618  degrees of freedom
## Residual deviance: 399.57  on 605  degrees of freedom
## (121 observations deleted due to missingness)
## AIC: 427.57
##
## Number of Fisher Scoring iterations: 15
```

Create regression model for vaccine response based on vaccine type

```
model2 <- glm(Vaccine_Response~Vaccine+Gender+visit_age+Race, data = result, family = "binomial"
)
```

Output model results

```
summary(model2)
```

```
##
## Call:
## glm(formula = Vaccine_Response ~ Vaccine + Gender + visit_age +
##      Race, family = "binomial", data = result)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -1.3538  -0.9118  -0.7107   1.2705   2.1781
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)    -17.06254  1455.39768  -0.012   0.9906
## Vaccine2         1.11051    0.55320   2.007   0.0447 *
## Vaccine4         0.65104    0.30951   2.103   0.0354 *
## GenderMale     -0.07348    0.26914  -0.273   0.7848
## visit_age       0.03335    0.02324   1.435   0.1513
## RaceAsian      14.50095  1455.39759   0.010   0.9921
## RaceBlack or African American 14.71753  1455.39801   0.010   0.9919
## RaceCaucasian   14.98843  1455.39756   0.010   0.9918
## RaceHispanic/Latino 31.40365  1664.83303   0.019   0.9850
## RaceOther       15.55404  1455.39757   0.011   0.9915
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 369.19  on 292  degrees of freedom
## Residual deviance: 345.38  on 283  degrees of freedom
## (447 observations deleted due to missingness)
## AIC: 365.38
##
## Number of Fisher Scoring iterations: 14
```

Create regression model for influenza infection based on donor factors

```
model12 <- glm(Influenza_Infection~Gender+visit_age+Race + bmi + statin_use, data = result, fami
ly = "binomial")
```

Output model results

```
summary(model12)
```

```
##
## Call:
## glm(formula = Influenza_Infection ~ Gender + visit_age + Race +
##      bmi + statin_use, family = "binomial", data = result)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -0.7678  -0.4840  -0.4194  -0.3313   2.7170
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)    -1.725e+01  1.693e+03  -0.010    0.992
## GenderMale      -3.863e-01  3.656e-01  -1.057    0.291
## visit_age       -1.486e-02  9.749e-03  -1.524    0.128
## RaceAmerican Indian or Alaska Native  5.472e-01  2.394e+03   0.000    1.000
## RaceAsian        1.317e+01  1.693e+03   0.008    0.994
## RaceBlack or African American    1.470e+01  1.693e+03   0.009    0.993
## RaceCaucasian    1.483e+01  1.693e+03   0.009    0.993
## RaceHispanic/Latino  2.531e-01  2.186e+03   0.000    1.000
## RaceOther        1.484e+01  1.693e+03   0.009    0.993
## bmi              3.510e-02  2.965e-02   1.184    0.236
## statin_use       3.054e-01  6.488e-01   0.471    0.638
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 256.00  on 424  degrees of freedom
## Residual deviance: 246.89  on 414  degrees of freedom
## (315 observations deleted due to missingness)
## AIC: 268.89
##
## Number of Fisher Scoring iterations: 15
```

Create regression model for vaccine response based on donor factors

```
model13 <- glm(Vaccine_Response~Gender+visit_age+Race + bmi + statin_use, data = result, family
= "binomial")
```

Output model results

```
summary(model13)
```

```
##
## Call:
## glm(formula = Vaccine_Response ~ Gender + visit_age + Race +
##      bmi + statin_use, family = "binomial", data = result)
##
## Deviance Residuals:
##      Min        1Q    Median        3Q        Max
## -1.4918  -0.8577  -0.7401   1.2363   1.9507
##
## Coefficients: (1 not defined because of singularities)
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)    -19.03908  2399.54496  -0.008    0.994
## GenderMale         0.05579    0.30182   0.185    0.853
## visit_age         0.03408    0.02843   1.199    0.231
## RaceAsian        16.18334  2399.54478   0.007    0.995
## RaceBlack or African American -0.03774  2769.09463   0.000    1.000
## RaceCaucasian    16.22066  2399.54475   0.007    0.995
## RaceHispanic/Latino 33.71846  2766.12814   0.012    0.990
## RaceOther        16.87326  2399.54476   0.007    0.994
## bmi              0.04670    0.03022   1.545    0.122
## statin_use              NA          NA      NA      NA
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 293.46  on 231  degrees of freedom
## Residual deviance: 271.80  on 223  degrees of freedom
## (508 observations deleted due to missingness)
## AIC: 289.8
##
## Number of Fisher Scoring iterations: 15
```

Regression model showing correlation between data values and vaccine response is significant

```
#Too big to run
model3 <- glm(vaccine_response~ data +gender+visit_age+race, data = flu, family = "binomial")
```

Output model results

```
summary(model3)
```

```
##
## Call:
## glm(formula = vaccine_response ~ data + gender + visit_age +
##      race, family = "binomial", data = flu)
##
## Deviance Residuals:
##      Min        1Q    Median        3Q        Max
## -1.9610  -1.0190  -0.7537   1.1855   3.0467
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)      1.155e+01  2.960e+01   0.390   0.696
## data             -2.801e-04  1.556e-05 -17.998 <2e-16 ***
## genderMale        1.325e-01  1.135e-02  11.679 <2e-16 ***
## visit_age         5.605e-02  5.477e-04 102.342 <2e-16 ***
## raceAsian        -1.298e+01  2.960e+01  -0.438   0.661
## raceBlack or African American -1.288e+01  2.960e+01  -0.435   0.663
## raceCaucasian     -1.250e+01  2.960e+01  -0.422   0.673
## raceHispanic/Latino -1.102e+01  2.960e+01  -0.372   0.710
## raceNULL          -1.447e+01  2.960e+01  -0.489   0.625
## raceOther         -1.278e+01  2.960e+01  -0.432   0.666
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 216418  on 156117  degrees of freedom
## Residual deviance: 190792  on 156108  degrees of freedom
## AIC: 190812
##
## Number of Fisher Scoring iterations: 12
```

Names of tests with most frequent count of high vaccine response

```
fluyes <- subset(flu, flu$vaccine_response == 1)
#head(fluyes)
rank <- sort(table(fluyes$name), decreasing = T)
head(rank)
```

```
##
## CD8+ T cells CD4+ T cells      T cells      B cells      NK cells      L50_CD40L
##              96              94              94              90              90              88
```


Create subset of data based on CD8+ T cells

```
tcell18 <- subset(flu, name == 'CD8+ T cells')
head(tcell18)
```

donor_id	study_id	gen...	race	visit_id	visit_year	visit_day	visit_type_hai	vi
<int>	<int>	<fctr>	<fctr>	<int>	<int>	<int>	<fctr>	

	donor_id <int>	study_id <int>	gen... <fctr>	race <fctr>	visit_id <int>	visit_year <int>	visit_day <int>	visit_type_hai <fctr>	vi
22	813	15	Female	Caucasian	2937	2014	0	pre	
162	812	15	Male	Caucasian	2936	2014	0	pre	
302	811	15	Male	Caucasian	2935	2014	0	pre	
441	810	15	Male	Caucasian	2934	2014	0	pre	
582	809	15	Female	Asian	2933	2014	0	pre	
722	808	15	Female	Other	2932	2014	0	pre	

6 rows | 1-10 of 45 columns



```
str(tcell18)
```

```
## 'data.frame': 464 obs. of 44 variables:
## $ donor_id : int 813 812 811 810 809 808 807 806 805 804 ...
## $ study_id : int 15 15 15 15 15 15 15 15 15 15 ...
## $ gender : Factor w/ 2 levels "Female","Male": 1 2 2 2 1 1 1 2 1 1 ...
## $ race : Factor w/ 7 levels "American Indian or Alaska Native",...: 4 4
4 4 2 7 4 7 4 4 ...
## $ visit_id : int 2937 2936 2935 2934 2933 2932 2931 2930 2929 2927 ...
## $ visit_year : int 2014 2014 2014 2014 2014 2014 2014 2014 2014 2014 ...
## $ visit_day : int 0 0 0 0 0 0 0 0 0 0 ...
## $ visit_type_hai : Factor w/ 2 levels "pre","single": 1 1 1 1 1 1 1 1 1 1 ...
## $ visit_age : num 23 28 23 27 27 29 24 27 23 26 ...
## $ cmv_status : Factor w/ 3 levels "0","1","NULL": 1 2 1 2 2 1 2 2 2 2 ...
## $ ebv_status : Factor w/ 3 levels "0","1","NULL": 1 2 1 2 2 2 1 2 2 2 ...
## $ bmi : Factor w/ 401 levels "13.12","13.59",...: 401 401 401 401 401
401 401 401 401 401 ...
## $ vaccine : Factor w/ 7 levels "1","2","3","4",...: 4 4 4 4 4 4 4 4 4 4
...
## $ geo_mean : num 380.6 47.6 95.1 190.3 56.6 ...
## $ d_geo_mean : Factor w/ 40 levels "0","1","10","101",...: 2 2 2 23 16 2 16 1
6 34 9 ...
## $ vaccine_response : Factor w/ 3 levels "0","1","NULL": 1 1 1 1 1 1 1 2 3 ...
## $ mesurment_id : int 371142 371002 370862 370721 370582 370442 370302 370162
370022 369882 ...
## $ assay : int 4 4 4 4 4 4 4 4 4 4 ...
## $ name : Factor w/ 3283 levels "B cells","BASO %",...: 44 44 44 44 44 4
4 44 44 44 44 ...
## $ name_formatted : Factor w/ 3283 levels "B_cells","BASO_",...: 44 44 44 44 44 44
44 44 44 44 ...
## $ subset : Factor w/ 632 levels "Activated T: Lymph/CD3+",...: 85 85 85 8
5 85 85 85 85 85 ...
## $ units : Factor w/ 8 levels "% of Parent",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ data : num 21.3 18.6 26.8 39.7 23.1 16.5 18.9 51.8 35.7 30 ...
## $ statin_use : Factor w/ 3 levels "0","1","NULL": 1 1 1 1 1 1 1 1 1 1 ...
## $ flu_vaccination_history : Factor w/ 3 levels "0","1","NULL": 2 2 2 2 2 2 2 2 2 2 ...
## $ total_vaccines_received : Factor w/ 25 levels "0","1","10","11",...: 4 8 6 2 20 24 14 11
8 20 ...
## $ vaccinated_1yr_prior : Factor w/ 4 levels "0","1","3","NULL": 2 2 2 2 2 2 2 2 2 2
...
## $ vaccine_type_1yr_prior : Factor w/ 4 levels "2","3","4","NULL": 1 1 1 1 1 1 1 1 1 1
...
## $ vaccinated_2yr_prior : Factor w/ 4 levels "0","1","3","NULL": 2 2 2 1 2 2 2 2 1 3
...
## $ vaccine_type_2yr_prior : Factor w/ 4 levels "2","3","4","NULL": 1 1 1 2 1 1 1 3 2 2
...
## $ vaccinated_3yr_prior : Factor w/ 4 levels "0","1","3","NULL": 2 2 3 1 2 2 2 2 1 3
...
## $ vaccine_type_3yr_prior : Factor w/ 4 levels "2","3","4","NULL": 1 1 2 2 1 1 1 1 2 2
...
## $ vaccinated_4yr_prior : Factor w/ 4 levels "0","1","3","NULL": 2 2 3 1 2 2 2 2 1 3
...
## $ vaccine_type_4yr_prior : Factor w/ 4 levels "2","3","4","NULL": 1 1 2 2 1 1 3 1 2 2
...
## $ vaccinated_5yr_prior : Factor w/ 4 levels "0","1","3","NULL": 2 2 2 1 2 2 2 1 2 3
```



```
...  
## $ vaccine_type_5yr_prior      : Factor w/ 4 levels "2","3","4","NULL": 1 1 1 2 1 1 3 2 1 2  
...  
## $ influenza_infection_history: int  0 0 0 0 0 1 0 0 0 0 ...  
## $ influenza_hospitalization  : int  0 0 0 0 0 0 0 0 0 0 ...  
## $ Gender                     : Factor w/ 2 levels "Female","Male": 1 2 2 2 1 1 1 2 1 1 ...  
## $ Vaccine                    : Factor w/ 7 levels "1","2","3","4",...: 4 4 4 4 4 4 4 4 4 4  
...  
## $ Vaccine_Response           : Factor w/ 3 levels "0","1","NULL": 1 1 1 1 1 1 1 1 2 3 ...  
## $ Influenza_Infection        : Factor w/ 2 levels "0","1": 1 1 1 1 1 2 1 1 1 1 ...  
## $ Influenza_Hospitalization  : Factor w/ 2 levels "0","1": 1 1 1 1 1 1 1 1 1 1 ...  
## $ subtest                    : Factor w/ 632 levels "Activated T: Lymph/CD3+",...: 85 85 85 8  
5 85 85 85 85 85 85 ...
```

Create regression model for vaccine response based on Test CD8+ T cells

```
model5 <- glm(vaccine_response~ data +gender+visit_age+race, data = tcell8, family = "binomial")
```

Output model results

```
summary(model5)
```

```
##
## Call:
## glm(formula = vaccine_response ~ data + gender + visit_age +
##      race, family = "binomial", data = tcell18)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -1.4059  -0.8508  -0.6871   1.2752   1.9617
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)    16.78574  1455.39762   0.012 0.990798
## data           -0.05305   0.01418  -3.742 0.000183 ***
## genderMale     -0.04434   0.21838  -0.203 0.839117
## visit_age       0.01039   0.01360   0.764 0.444768
## raceAsian      -15.68529  1455.39757  -0.011 0.991401
## raceBlack or African American -31.33409  1674.45202  -0.019 0.985070
## raceCaucasian  -16.47059  1455.39755  -0.011 0.990971
## raceHispanic/Latino    0.15736  1669.24508   0.000 0.999925
## raceNULL       -30.32341  2058.24295  -0.015 0.988245
## raceOther      -15.93886  1455.39756  -0.011 0.991262
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 561.35  on 463  degrees of freedom
## Residual deviance: 523.76  on 454  degrees of freedom
## AIC: 543.76
##
## Number of Fisher Scoring iterations: 14
```

Create subset of data based on CD4+ T cells

```
tcell14 <- subset(flu, name == 'CD4+ T cells')
head(tcell14)
```

	donor_id	study_id	gen...	race	visit_id	visit_year	visit_day	visit_type_hai	v
	<int>	<int>	<fctr>	<fctr>	<int>	<int>	<int>	<fctr>	
16	813	15	Female	Caucasian	2937	2014	0	pre	
156	812	15	Male	Caucasian	2936	2014	0	pre	
296	811	15	Male	Caucasian	2935	2014	0	pre	
435	810	15	Male	Caucasian	2934	2014	0	pre	
576	809	15	Female	Asian	2933	2014	0	pre	
716	808	15	Female	Other	2932	2014	0	pre	

6 rows | 1-10 of 45 columns

```
str(tcell14)
```

```

## 'data.frame': 438 obs. of 44 variables:
## $ donor_id : int 813 812 811 810 809 808 807 806 805 804 ...
## $ study_id : int 15 15 15 15 15 15 15 15 15 15 ...
## $ gender : Factor w/ 2 levels "Female","Male": 1 2 2 2 1 1 1 2 1 1 ...
## $ race : Factor w/ 7 levels "American Indian or Alaska Native",...: 4 4
4 4 2 7 4 7 4 4 ...
## $ visit_id : int 2937 2936 2935 2934 2933 2932 2931 2930 2929 2927 ...
## $ visit_year : int 2014 2014 2014 2014 2014 2014 2014 2014 2014 2014 ...
## $ visit_day : int 0 0 0 0 0 0 0 0 0 0 ...
## $ visit_type_hai : Factor w/ 2 levels "pre","single": 1 1 1 1 1 1 1 1 1 1 ...
## $ visit_age : num 23 28 23 27 27 29 24 27 23 26 ...
## $ cmv_status : Factor w/ 3 levels "0","1","NULL": 1 2 1 2 2 1 2 2 2 2 ...
## $ ebv_status : Factor w/ 3 levels "0","1","NULL": 1 2 1 2 2 2 1 2 2 2 ...
## $ bmi : Factor w/ 401 levels "13.12","13.59",...: 401 401 401 401 401
401 401 401 401 401 ...
## $ vaccine : Factor w/ 7 levels "1","2","3","4",...: 4 4 4 4 4 4 4 4 4 4
...
## $ geo_mean : num 380.6 47.6 95.1 190.3 56.6 ...
## $ d_geo_mean : Factor w/ 40 levels "0","1","10","101",...: 2 2 2 23 16 2 16 1
6 34 9 ...
## $ vaccine_response : Factor w/ 3 levels "0","1","NULL": 1 1 1 1 1 1 1 2 3 ...
## $ mesurment_id : int 371136 370996 370856 370715 370576 370436 370296 370156
370016 369876 ...
## $ assay : int 4 4 4 4 4 4 4 4 4 4 ...
## $ name : Factor w/ 3283 levels "B cells","BASO %",...: 36 36 36 36 36 3
6 36 36 36 36 ...
## $ name_formatted : Factor w/ 3283 levels "B_cells","BASO_",...: 38 38 38 38 38 38
38 38 38 38 ...
## $ subset : Factor w/ 632 levels "Activated T: Lymph/CD3+",...: 57 57 57 5
7 57 57 57 57 57 57 ...
## $ units : Factor w/ 8 levels "% of Parent",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ data : num 70 70.7 63.1 45.4 49.1 71.3 50.8 42.6 53.8 48.4 ...
## $ statin_use : Factor w/ 3 levels "0","1","NULL": 1 1 1 1 1 1 1 1 1 1 ...
## $ flu_vaccination_history : Factor w/ 3 levels "0","1","NULL": 2 2 2 2 2 2 2 2 2 2 ...
## $ total_vaccines_received : Factor w/ 25 levels "0","1","10","11",...: 4 8 6 2 20 24 14 11
8 20 ...
## $ vaccinated_1yr_prior : Factor w/ 4 levels "0","1","3","NULL": 2 2 2 2 2 2 2 2 2 2
...
## $ vaccine_type_1yr_prior : Factor w/ 4 levels "2","3","4","NULL": 1 1 1 1 1 1 1 1 1 1
...
## $ vaccinated_2yr_prior : Factor w/ 4 levels "0","1","3","NULL": 2 2 2 1 2 2 2 2 1 3
...
## $ vaccine_type_2yr_prior : Factor w/ 4 levels "2","3","4","NULL": 1 1 1 2 1 1 1 3 2 2
...
## $ vaccinated_3yr_prior : Factor w/ 4 levels "0","1","3","NULL": 2 2 3 1 2 2 2 2 1 3
...
## $ vaccine_type_3yr_prior : Factor w/ 4 levels "2","3","4","NULL": 1 1 2 2 1 1 1 1 2 2
...
## $ vaccinated_4yr_prior : Factor w/ 4 levels "0","1","3","NULL": 2 2 3 1 2 2 2 2 1 3
...
## $ vaccine_type_4yr_prior : Factor w/ 4 levels "2","3","4","NULL": 1 1 2 2 1 1 3 1 2 2
...
## $ vaccinated_5yr_prior : Factor w/ 4 levels "0","1","3","NULL": 2 2 2 1 2 2 2 1 2 3

```

```
...  
## $ vaccine_type_5yr_prior      : Factor w/ 4 levels "2","3","4","NULL": 1 1 1 2 1 1 3 2 1 2  
...  
## $ influenza_infection_history: int  0 0 0 0 0 1 0 0 0 0 ...  
## $ influenza_hospitalization  : int  0 0 0 0 0 0 0 0 0 0 ...  
## $ Gender                     : Factor w/ 2 levels "Female","Male": 1 2 2 2 1 1 1 2 1 1 ...  
## $ Vaccine                    : Factor w/ 7 levels "1","2","3","4",...: 4 4 4 4 4 4 4 4 4 4  
...  
## $ Vaccine_Response           : Factor w/ 3 levels "0","1","NULL": 1 1 1 1 1 1 1 1 2 3 ...  
## $ Influenza_Infection        : Factor w/ 2 levels "0","1": 1 1 1 1 1 2 1 1 1 1 ...  
## $ Influenza_Hospitalization  : Factor w/ 2 levels "0","1": 1 1 1 1 1 1 1 1 1 1 ...  
## $ subtest                    : Factor w/ 632 levels "Activated T: Lymph/CD3+",...: 57 57 57 5  
7 57 57 57 57 57 57 ...
```

Create regression model for vaccine response based on Test CD4+ T cells

```
model6 <- glm(vaccine_response~ data +gender+visit_age+race, data = tcell4, family = "binomial")
```

Output model results

```
summary(model6)
```

```
##
## Call:
## glm(formula = vaccine_response ~ data + gender + visit_age +
##      race, family = "binomial", data = tcell4)
##
## Deviance Residuals:
##      Min        1Q    Median        3Q        Max
## -1.2665  -0.8617  -0.7370   1.3151   1.8344
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)      1.371e+01  1.455e+03   0.009   0.9925
## data              3.192e-02  1.323e-02   2.413   0.0158 *
## genderMale       -2.877e-02  2.208e-01  -0.130   0.8963
## visit_age         8.833e-03  1.340e-02   0.659   0.5098
## raceAsian        -1.592e+01  1.455e+03  -0.011   0.9913
## raceBlack or African American -3.161e+01  1.678e+03  -0.019   0.9850
## raceCaucasian     -1.677e+01  1.455e+03  -0.012   0.9908
## raceHispanic/Latino -2.441e-01  1.675e+03   0.000   0.9999
## raceNULL          -3.092e+01  2.058e+03  -0.015   0.9880
## raceOther        -1.626e+01  1.455e+03  -0.011   0.9911
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 539.45  on 437  degrees of freedom
## Residual deviance: 512.76  on 428  degrees of freedom
## AIC: 532.76
##
## Number of Fisher Scoring iterations: 14
```

Create subset of data based on Tcells

```
tcell <- subset(flu, name == 'T cells')
head(tcell)
```

	donor_id	study_id	gen...	race	visit_id	visit_year	visit_day	visit_type_hai	v
	<int>	<int>	<fctr>	<fctr>	<int>	<int>	<int>	<fctr>	
60	813	15	Female	Caucasian	2937	2014	0	pre	
200	812	15	Male	Caucasian	2936	2014	0	pre	
339	811	15	Male	Caucasian	2935	2014	0	pre	
478	810	15	Male	Caucasian	2934	2014	0	pre	
620	809	15	Female	Asian	2933	2014	0	pre	
760	808	15	Female	Other	2932	2014	0	pre	

6 rows | 1-10 of 45 columns

```
str(tcell)
```

```

## 'data.frame':   438 obs. of  44 variables:
## $ donor_id      : int  813 812 811 810 809 808 807 806 805 804 ...
## $ study_id      : int  15 15 15 15 15 15 15 15 15 15 ...
## $ gender        : Factor w/ 2 levels "Female","Male": 1 2 2 2 1 1 1 2 1 1 ...
## $ race          : Factor w/ 7 levels "American Indian or Alaska Native",...: 4 4
4 4 2 7 4 7 4 4 ...
## $ visit_id      : int  2937 2936 2935 2934 2933 2932 2931 2930 2929 2927 ...
## $ visit_year    : int  2014 2014 2014 2014 2014 2014 2014 2014 2014 2014 ...
## $ visit_day     : int   0 0 0 0 0 0 0 0 0 0 ...
## $ visit_type_hai : Factor w/ 2 levels "pre","single": 1 1 1 1 1 1 1 1 1 1 ...
## $ visit_age     : num  23 28 23 27 27 29 24 27 23 26 ...
## $ cmv_status    : Factor w/ 3 levels "0","1","NULL": 1 2 1 2 2 1 2 2 2 2 ...
## $ ebv_status    : Factor w/ 3 levels "0","1","NULL": 1 2 1 2 2 2 1 2 2 2 ...
## $ bmi          : Factor w/ 401 levels "13.12","13.59",...: 401 401 401 401 401
401 401 401 401 401 ...
## $ vaccine      : Factor w/ 7 levels "1","2","3","4",...: 4 4 4 4 4 4 4 4 4 4
...
## $ geo_mean     : num  380.6 47.6 95.1 190.3 56.6 ...
## $ d_geo_mean   : Factor w/ 40 levels "0","1","10","101",...: 2 2 2 23 16 2 16 1
6 34 9 ...
## $ vaccine_response : Factor w/ 3 levels "0","1","NULL": 1 1 1 1 1 1 1 1 2 3 ...
## $ mesurment_id  : int  371180 371040 370899 370758 370620 370480 370340 370199
370060 369920 ...
## $ assay        : int   4 4 4 4 4 4 4 4 4 4 ...
## $ name         : Factor w/ 3283 levels "B cells","BASO %",...: 2885 2885 2885 2
885 2885 2885 2885 2885 2885 ...
## $ name_formatted : Factor w/ 3283 levels "B_cells","BASO_",...: 2885 2885 2885 28
85 2885 2885 2885 2885 2885 ...
## $ subset       : Factor w/ 632 levels "Activated T: Lymph/CD3+",...: 46 46 46 4
6 46 46 46 46 46 46 ...
## $ units        : Factor w/ 8 levels "% of Parent",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ data         : num  56.3 51.8 53.2 56.2 59.8 52.7 69.9 67 71.9 43.1 ...
## $ statin_use    : Factor w/ 3 levels "0","1","NULL": 1 1 1 1 1 1 1 1 1 1 ...
## $ flu_vaccination_history : Factor w/ 3 levels "0","1","NULL": 2 2 2 2 2 2 2 2 2 2 ...
## $ total_vaccines_received : Factor w/ 25 levels "0","1","10","11",...: 4 8 6 2 20 24 14 11
8 20 ...
## $ vaccinated_1yr_prior : Factor w/ 4 levels "0","1","3","NULL": 2 2 2 2 2 2 2 2 2 2
...
## $ vaccine_type_1yr_prior : Factor w/ 4 levels "2","3","4","NULL": 1 1 1 1 1 1 1 1 1 1
...
## $ vaccinated_2yr_prior : Factor w/ 4 levels "0","1","3","NULL": 2 2 2 1 2 2 2 2 1 3
...
## $ vaccine_type_2yr_prior : Factor w/ 4 levels "2","3","4","NULL": 1 1 1 2 1 1 1 3 2 2
...
## $ vaccinated_3yr_prior : Factor w/ 4 levels "0","1","3","NULL": 2 2 3 1 2 2 2 2 1 3
...
## $ vaccine_type_3yr_prior : Factor w/ 4 levels "2","3","4","NULL": 1 1 2 2 1 1 1 1 2 2
...
## $ vaccinated_4yr_prior : Factor w/ 4 levels "0","1","3","NULL": 2 2 3 1 2 2 2 2 1 3
...
## $ vaccine_type_4yr_prior : Factor w/ 4 levels "2","3","4","NULL": 1 1 2 2 1 1 3 1 2 2
...
## $ vaccinated_5yr_prior : Factor w/ 4 levels "0","1","3","NULL": 2 2 2 1 2 2 2 1 2 3

```



```
...  
## $ vaccine_type_5yr_prior      : Factor w/ 4 levels "2","3","4","NULL": 1 1 1 2 1 1 3 2 1 2  
...  
## $ influenza_infection_history: int  0 0 0 0 0 1 0 0 0 0 ...  
## $ influenza_hospitalization  : int  0 0 0 0 0 0 0 0 0 0 ...  
## $ Gender                     : Factor w/ 2 levels "Female","Male": 1 2 2 2 1 1 1 2 1 1 ...  
## $ Vaccine                    : Factor w/ 7 levels "1","2","3","4",...: 4 4 4 4 4 4 4 4 4 4  
...  
## $ Vaccine_Response           : Factor w/ 3 levels "0","1","NULL": 1 1 1 1 1 1 1 1 2 3 ...  
## $ Influenza_Infection        : Factor w/ 2 levels "0","1": 1 1 1 1 1 2 1 1 1 1 ...  
## $ Influenza_Hospitalization  : Factor w/ 2 levels "0","1": 1 1 1 1 1 1 1 1 1 1 ...  
## $ subtest                    : Factor w/ 632 levels "Activated T: Lymph/CD3+",...: 46 46 46 4  
6 46 46 46 46 46 46 ...
```

Create regression model for vaccine response based on Test Tcells

```
model7 <- glm(vaccine_response~ data +gender+visit_age+race, data = tcell, family = "binomial")
```

Output model results

```
summary(model7)
```

```
##
## Call:
## glm(formula = vaccine_response ~ data + gender + visit_age +
##      race, family = "binomial", data = tcell)
##
## Deviance Residuals:
##      Min        1Q    Median        3Q        Max
## -1.1656  -0.8618  -0.7429   1.3000   2.0072
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)    1.430e+01  1.455e+03   0.010   0.992
## data           1.974e-02  9.416e-03   2.097   0.036 *
## genderMale     -5.265e-02  2.195e-01  -0.240   0.810
## visit_age       1.125e-02  1.354e-02   0.831   0.406
## raceAsian      -1.618e+01  1.455e+03  -0.011   0.991
## raceBlack or African American -3.140e+01  1.680e+03  -0.019   0.985
## raceCaucasian  -1.692e+01  1.455e+03  -0.012   0.991
## raceHispanic/Latino -2.979e-01  1.680e+03   0.000   1.000
## raceNULL       -3.147e+01  2.058e+03  -0.015   0.988
## raceOther      -1.642e+01  1.455e+03  -0.011   0.991
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 539.45  on 437  degrees of freedom
## Residual deviance: 514.19  on 428  degrees of freedom
## AIC: 534.19
##
## Number of Fisher Scoring iterations: 14
```

Create subset of data based on B Cells

```
bcells <- subset(flu, name == 'B cells')
head(bcells)
```

	donor_id	study_id	gen...	race	visit_id	visit_year	visit_day	visit_type_hai	v
	<int>	<int>	<fctr>	<fctr>	<int>	<int>	<int>	<fctr>	
1	813	15	Female	Caucasian	2937	2014	0	pre	
141	812	15	Male	Caucasian	2936	2014	0	pre	
281	811	15	Male	Caucasian	2935	2014	0	pre	
421	810	15	Male	Caucasian	2934	2014	0	pre	
561	809	15	Female	Asian	2933	2014	0	pre	
701	808	15	Female	Other	2932	2014	0	pre	

6 rows | 1-10 of 45 columns

```
str(bcells)
```

```

## 'data.frame':   386 obs. of  44 variables:
## $ donor_id      : int  813 812 811 810 809 808 807 806 805 804 ...
## $ study_id      : int  15 15 15 15 15 15 15 15 15 15 ...
## $ gender        : Factor w/ 2 levels "Female","Male": 1 2 2 2 1 1 1 2 1 1 ...
## $ race          : Factor w/ 7 levels "American Indian or Alaska Native",...: 4 4
4 4 2 7 4 7 4 4 ...
## $ visit_id      : int  2937 2936 2935 2934 2933 2932 2931 2930 2929 2927 ...
## $ visit_year    : int  2014 2014 2014 2014 2014 2014 2014 2014 2014 2014 ...
## $ visit_day     : int   0 0 0 0 0 0 0 0 0 0 ...
## $ visit_type_hai : Factor w/ 2 levels "pre","single": 1 1 1 1 1 1 1 1 1 1 ...
## $ visit_age     : num  23 28 23 27 27 29 24 27 23 26 ...
## $ cmv_status    : Factor w/ 3 levels "0","1","NULL": 1 2 1 2 2 1 2 2 2 2 ...
## $ ebv_status    : Factor w/ 3 levels "0","1","NULL": 1 2 1 2 2 2 1 2 2 2 ...
## $ bmi          : Factor w/ 401 levels "13.12","13.59",...: 401 401 401 401 401
401 401 401 401 401 ...
## $ vaccine      : Factor w/ 7 levels "1","2","3","4",...: 4 4 4 4 4 4 4 4 4 4
...
## $ geo_mean     : num  380.6 47.6 95.1 190.3 56.6 ...
## $ d_geo_mean   : Factor w/ 40 levels "0","1","10","101",...: 2 2 2 23 16 2 16 1
6 34 9 ...
## $ vaccine_response : Factor w/ 3 levels "0","1","NULL": 1 1 1 1 1 1 1 1 2 3 ...
## $ mesurment_id  : int  371121 370981 370841 370701 370561 370421 370281 370141
370001 369861 ...
## $ assay        : int   4 4 4 4 4 4 4 4 4 4 ...
## $ name         : Factor w/ 3283 levels "B cells","BASO %",...: 1 1 1 1 1 1 1 1 1
1 1 ...
## $ name_formatted : Factor w/ 3283 levels "B_cells","BASO_",...: 1 1 1 1 1 1 1 1 1
1 ...
## $ subset       : Factor w/ 632 levels "Activated T: Lymph/CD3+",...: 36 36 36 3
6 36 36 36 36 36 ...
## $ units        : Factor w/ 8 levels "% of Parent",...: 1 1 1 1 1 1 1 1 1 ...
## $ data         : num  34.4 41 34.6 34.8 40.8 72.5 44.8 48.1 61.6 45.7 ...
## $ statin_use    : Factor w/ 3 levels "0","1","NULL": 1 1 1 1 1 1 1 1 1 ...
## $ flu_vaccination_history : Factor w/ 3 levels "0","1","NULL": 2 2 2 2 2 2 2 2 2 ...
## $ total_vaccines_received : Factor w/ 25 levels "0","1","10","11",...: 4 8 6 2 20 24 14 11
8 20 ...
## $ vaccinated_1yr_prior : Factor w/ 4 levels "0","1","3","NULL": 2 2 2 2 2 2 2 2 2
...
## $ vaccine_type_1yr_prior : Factor w/ 4 levels "2","3","4","NULL": 1 1 1 1 1 1 1 1 1
...
## $ vaccinated_2yr_prior : Factor w/ 4 levels "0","1","3","NULL": 2 2 2 1 2 2 2 2 1 3
...
## $ vaccine_type_2yr_prior : Factor w/ 4 levels "2","3","4","NULL": 1 1 1 2 1 1 1 3 2 2
...
## $ vaccinated_3yr_prior : Factor w/ 4 levels "0","1","3","NULL": 2 2 3 1 2 2 2 2 1 3
...
## $ vaccine_type_3yr_prior : Factor w/ 4 levels "2","3","4","NULL": 1 1 2 2 1 1 1 1 2 2
...
## $ vaccinated_4yr_prior : Factor w/ 4 levels "0","1","3","NULL": 2 2 3 1 2 2 2 2 1 3
...
## $ vaccine_type_4yr_prior : Factor w/ 4 levels "2","3","4","NULL": 1 1 2 2 1 1 3 1 2 2
...
## $ vaccinated_5yr_prior : Factor w/ 4 levels "0","1","3","NULL": 2 2 2 1 2 2 2 1 2 3

```

```

...
## $ vaccine_type_5yr_prior      : Factor w/ 4 levels "2","3","4","NULL": 1 1 1 2 1 1 3 2 1 2
...
## $ influenza_infection_history: int  0 0 0 0 0 1 0 0 0 0 ...
## $ influenza_hospitalization : int  0 0 0 0 0 0 0 0 0 0 ...
## $ Gender                     : Factor w/ 2 levels "Female","Male": 1 2 2 2 1 1 1 2 1 1 ...
## $ Vaccine                    : Factor w/ 7 levels "1","2","3","4",...: 4 4 4 4 4 4 4 4 4 4
...
## $ Vaccine_Response           : Factor w/ 3 levels "0","1","NULL": 1 1 1 1 1 1 1 1 2 3 ...
## $ Influenza_Infection        : Factor w/ 2 levels "0","1": 1 1 1 1 1 2 1 1 1 1 ...
## $ Influenza_Hospitalization  : Factor w/ 2 levels "0","1": 1 1 1 1 1 1 1 1 1 1 ...
## $ subtest                    : Factor w/ 632 levels "Activated T: Lymph/CD3+",...: 36 36 36 3
6 36 36 36 36 36 36 ...

```

Create regression model for vaccine response based on Test Bcells

```
model8 <- glm(vaccine_response~ data +gender+visit_age+race, data = bcells, family = "binomial")
```

Output model results

```
summary(model8)
```

```

##
## Call:
## glm(formula = vaccine_response ~ data + gender + visit_age +
##      race, family = "binomial", data = bcells)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -1.2420  -0.8981  -0.8004   1.3486   1.7106
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)    1.549e+01  2.400e+03   0.006   0.995
## data            1.114e-02  7.752e-03   1.438   0.151
## genderMale     -3.714e-02  2.270e-01  -0.164   0.870
## visit_age       1.153e-02  1.365e-02   0.844   0.398
## raceAsian      -1.666e+01  2.400e+03  -0.007   0.994
## raceBlack or African American -3.289e+01  2.768e+03  -0.012   0.991
## raceCaucasian  -1.714e+01  2.400e+03  -0.007   0.994
## raceHispanic/Latino    2.525e-01  2.771e+03   0.000   1.000
## raceNULL       -3.294e+01  3.393e+03  -0.010   0.992
## raceOther      -1.667e+01  2.400e+03  -0.007   0.994
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 493.22  on 385  degrees of freedom
## Residual deviance: 473.61  on 376  degrees of freedom
## AIC: 493.61
##
## Number of Fisher Scoring iterations: 15

```

## Create subset of data based on NK cells

```
nkcell <- subset(flu, name == 'NK cells')
head(nkcell)
```

	<b>donor_id</b> <int>	<b>study_id</b> <int>	<b>gen...</b> <fctr>	<b>race</b> <fctr>	<b>visit_id</b> <int>	<b>visit_year</b> <int>	<b>visit_day</b> <int>	<b>visit_type_hai</b> <fctr>	<b>vi</b>
54	813	15	Female	Caucasian	2937	2014	0	pre	
194	812	15	Male	Caucasian	2936	2014	0	pre	
334	811	15	Male	Caucasian	2935	2014	0	pre	
473	810	15	Male	Caucasian	2934	2014	0	pre	
614	809	15	Female	Asian	2933	2014	0	pre	
754	808	15	Female	Other	2932	2014	0	pre	

6 rows | 1-10 of 45 columns

```
str(nkcell)
```

```

## 'data.frame':   386 obs. of  44 variables:
## $ donor_id      : int  813 812 811 810 809 808 807 806 805 804 ...
## $ study_id      : int  15 15 15 15 15 15 15 15 15 15 ...
## $ gender        : Factor w/ 2 levels "Female","Male": 1 2 2 2 1 1 1 2 1 1 ...
## $ race          : Factor w/ 7 levels "American Indian or Alaska Native",...: 4 4
4 4 2 7 4 7 4 4 ...
## $ visit_id      : int  2937 2936 2935 2934 2933 2932 2931 2930 2929 2927 ...
## $ visit_year    : int  2014 2014 2014 2014 2014 2014 2014 2014 2014 2014 ...
## $ visit_day     : int   0 0 0 0 0 0 0 0 0 0 ...
## $ visit_type_hai : Factor w/ 2 levels "pre","single": 1 1 1 1 1 1 1 1 1 1 ...
## $ visit_age     : num  23 28 23 27 27 29 24 27 23 26 ...
## $ cmv_status    : Factor w/ 3 levels "0","1","NULL": 1 2 1 2 2 1 2 2 2 2 ...
## $ ebv_status    : Factor w/ 3 levels "0","1","NULL": 1 2 1 2 2 2 1 2 2 2 ...
## $ bmi          : Factor w/ 401 levels "13.12","13.59",...: 401 401 401 401 401
401 401 401 401 401 ...
## $ vaccine      : Factor w/ 7 levels "1","2","3","4",...: 4 4 4 4 4 4 4 4 4 4
...
## $ geo_mean     : num  380.6 47.6 95.1 190.3 56.6 ...
## $ d_geo_mean   : Factor w/ 40 levels "0","1","10","101",...: 2 2 2 23 16 2 16 1
6 34 9 ...
## $ vaccine_response : Factor w/ 3 levels "0","1","NULL": 1 1 1 1 1 1 1 1 2 3 ...
## $ mesurment_id  : int  371174 371034 370894 370753 370614 370474 370334 370194
370054 369914 ...
## $ assay        : int   4 4 4 4 4 4 4 4 4 4 ...
## $ name         : Factor w/ 3283 levels "B cells","BASO %",...: 2520 2520 2520 2
520 2520 2520 2520 2520 2520 ...
## $ name_formatted : Factor w/ 3283 levels "B_cells","BASO_",...: 2520 2520 2520 25
20 2520 2520 2520 2520 2520 ...
## $ subset       : Factor w/ 632 levels "Activated T: Lymph/CD3+",...: 31 31 31 3
1 31 31 31 31 31 31 ...
## $ units        : Factor w/ 8 levels "% of Parent",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ data         : num  26.5 20.2 10.6 34.9 5.4 13.7 6.53 39.4 15.7 1.37 ...
## $ statin_use    : Factor w/ 3 levels "0","1","NULL": 1 1 1 1 1 1 1 1 1 1 ...
## $ flu_vaccination_history : Factor w/ 3 levels "0","1","NULL": 2 2 2 2 2 2 2 2 2 2 ...
## $ total_vaccines_received : Factor w/ 25 levels "0","1","10","11",...: 4 8 6 2 20 24 14 11
8 20 ...
## $ vaccinated_1yr_prior : Factor w/ 4 levels "0","1","3","NULL": 2 2 2 2 2 2 2 2 2 2
...
## $ vaccine_type_1yr_prior : Factor w/ 4 levels "2","3","4","NULL": 1 1 1 1 1 1 1 1 1 1
...
## $ vaccinated_2yr_prior : Factor w/ 4 levels "0","1","3","NULL": 2 2 2 1 2 2 2 2 1 3
...
## $ vaccine_type_2yr_prior : Factor w/ 4 levels "2","3","4","NULL": 1 1 1 2 1 1 1 3 2 2
...
## $ vaccinated_3yr_prior : Factor w/ 4 levels "0","1","3","NULL": 2 2 3 1 2 2 2 2 1 3
...
## $ vaccine_type_3yr_prior : Factor w/ 4 levels "2","3","4","NULL": 1 1 2 2 1 1 1 1 2 2
...
## $ vaccinated_4yr_prior : Factor w/ 4 levels "0","1","3","NULL": 2 2 3 1 2 2 2 2 1 3
...
## $ vaccine_type_4yr_prior : Factor w/ 4 levels "2","3","4","NULL": 1 1 2 2 1 1 3 1 2 2
...
## $ vaccinated_5yr_prior : Factor w/ 4 levels "0","1","3","NULL": 2 2 2 1 2 2 2 1 2 3

```

```
...  
## $ vaccine_type_5yr_prior      : Factor w/ 4 levels "2","3","4","NULL": 1 1 1 2 1 1 3 2 1 2  
...  
## $ influenza_infection_history: int  0 0 0 0 0 1 0 0 0 0 ...  
## $ influenza_hospitalization  : int  0 0 0 0 0 0 0 0 0 0 ...  
## $ Gender                     : Factor w/ 2 levels "Female","Male": 1 2 2 2 1 1 1 2 1 1 ...  
## $ Vaccine                    : Factor w/ 7 levels "1","2","3","4",...: 4 4 4 4 4 4 4 4 4 4  
...  
## $ Vaccine_Response           : Factor w/ 3 levels "0","1","NULL": 1 1 1 1 1 1 1 1 2 3 ...  
## $ Influenza_Infection        : Factor w/ 2 levels "0","1": 1 1 1 1 1 2 1 1 1 1 ...  
## $ Influenza_Hospitalization  : Factor w/ 2 levels "0","1": 1 1 1 1 1 1 1 1 1 1 ...  
## $ subtest                    : Factor w/ 632 levels "Activated T: Lymph/CD3+",...: 31 31 31 3  
1 31 31 31 31 31 31 ...
```

Create regression model for vaccine response based on Test NK cells

```
model9 <- glm(vaccine_response~ data +gender+visit_age+race, data = nkcell, family = "binomial")
```

Output model results

```
summary(model9)
```



```
##
## Call:
## glm(formula = vaccine_response ~ data + gender + visit_age +
##      race, family = "binomial", data = nkcell)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -1.2726  -0.9109  -0.7901   1.3503   1.8293
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)    16.40939  2399.54475   0.007  0.9945
## data           -0.02101   0.01036  -2.028  0.0425 *
## genderMale     -0.01220   0.22811  -0.053  0.9574
## visit_age       0.01133   0.01370   0.827  0.4080
## raceAsian     -16.66358  2399.54475  -0.007  0.9945
## raceBlack or African American -32.98700  2764.99387  -0.012  0.9905
## raceCaucasian  -17.21805  2399.54474  -0.007  0.9943
## raceHispanic/Latino    0.21349  2767.31887   0.000  0.9999
## raceNULL       -33.05937  3393.46870  -0.010  0.9922
## raceOther      -16.66640  2399.54475  -0.007  0.9945
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 493.22  on 385  degrees of freedom
## Residual deviance: 471.46  on 376  degrees of freedom
## AIC: 491.46
##
## Number of Fisher Scoring iterations: 15
```