

I. χ^2 -분포

II. t-분포

III. F-분포

1. 확률분포

- 표준정규분포의 제곱의 합 $X = \sum_{i=1}^n Z_i^2$ 이 자유도가 n 인 χ^2 -분포에 따름
- $X \sim \chi^2(n)$
- X 가 $X \sim \chi^2(n)$ 이라고 할 때, X 의 평균은 n , 분산은 $2n$

b1-ch4-10.R

```
set.seed(12345)
```

```
n<-10000;
```

```
z1<-rnorm(n,0,1)
```

```
z2<-rnorm(n,0,1)
```

```
z3<-rnorm(n,0,1)
```

```
z4<-rnorm(n,0,1)
```

```
z5<-rnorm(n,0,1)
```

```
chi5<-z1^2+z2^2+z3^2+z4^2+z5^2
```

```
(mean(chi5))
```

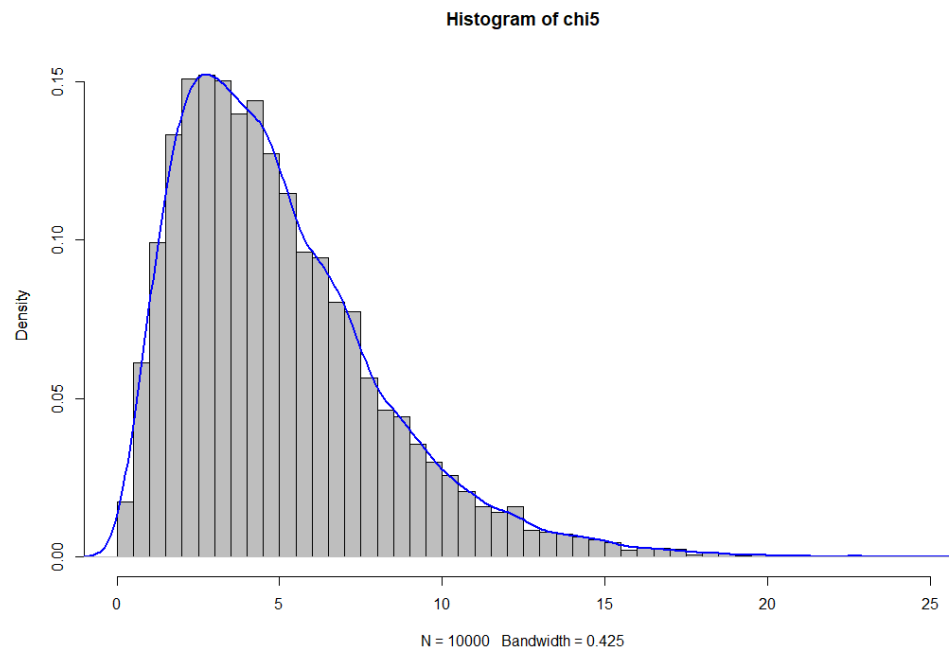
```
(var(chi5))
```

```
hist(chi5, freq=F, col="grey", xlab="", xlim=c(0, 25),  
breaks=100)
```

```
par(new=T)
```

```
plot(density(chi5), axes=F, main="", xlim=c(0, 25), l  
wd=2, col="blue")
```

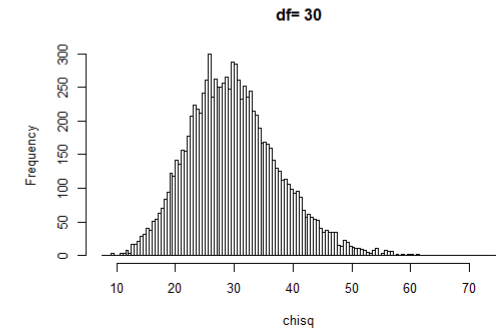
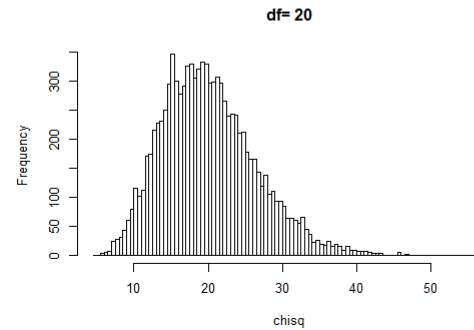
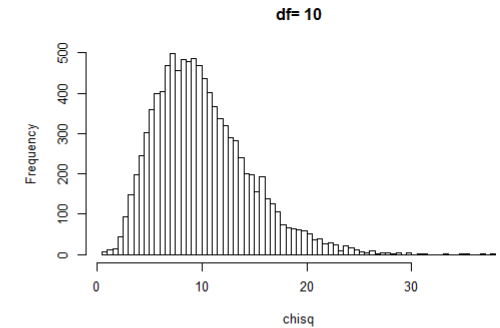
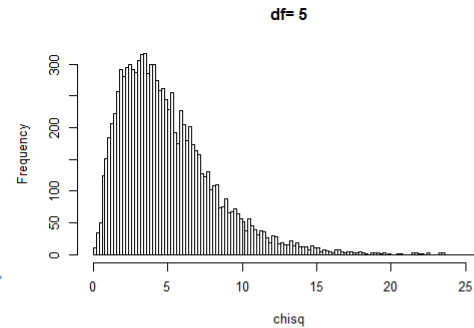
```
> (mean(chi5))  
[1] 4.975514  
> (var(chi5))  
[1] 10.05467  
> library(car)
```



- 자유도에 따라 χ^2 -분포의 모양이 달라지는데 자유도가 클수록 정규분포와 근사

b1-ch4-11.R

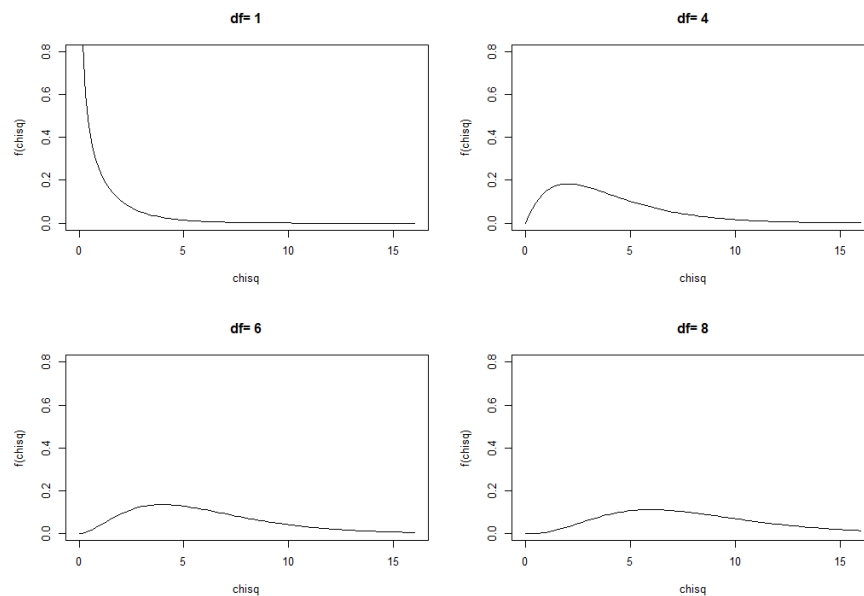
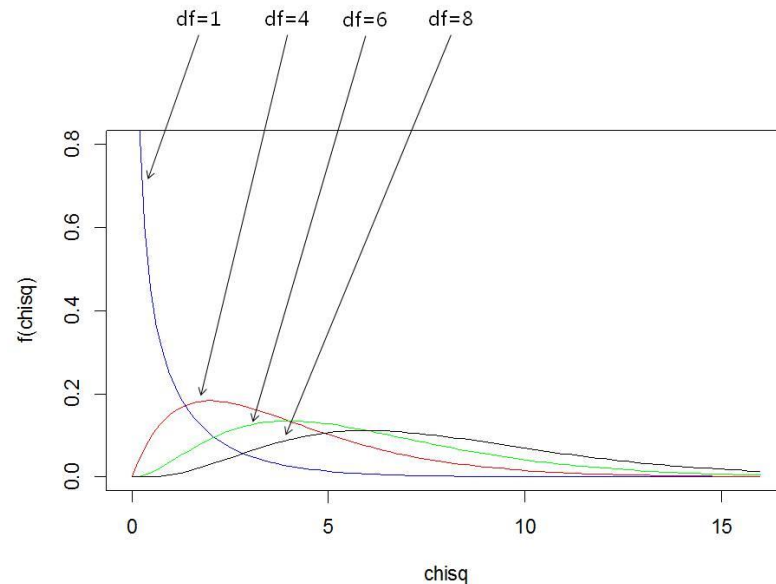
```
set.seed(12345)
n<-10000;
df_list<-c(5,10,20,30)
par(mfrow=c(2,2))
for (i in 1:length(df_list)) {
  hist(rchisq(n, df=df_list[i], ncp=0), breaks=100,
xlab="chisq", main=paste("df=", df_list[i]))
}
```



b1-ch4-12.R

```
n_list<-c(2,5,7,9) # 표본수(n)
df_list<-n_list-1 # 자유도
curve(dchisq(x, 1, ncp=0), col="blue", xlim=c(0, 16),
      ylim=c(0, 0.8), xlab="chisq", ylab="f(chisq)")
curve(dchisq(x, 4, ncp=0), add=T, col="red", xlim=c(0, 16),
      ylim=c(0, 0.8), xlab="chisq", ylab="f(chisq)")
curve(dchisq(x, 6, ncp=0), add=T, col="green", xlim=c(0, 16),
      ylim=c(0, 0.8), xlab="chisq", ylab="f(chisq)")
curve(dchisq(x, 8, ncp=0), add=T, col="black", xlim=c(0, 16),
      ylim=c(0, 0.8), xlab="chisq", ylab="f(chisq)")

par(mfrow=c(2,2))
for (i in 1:length(df_list)) {
  curve(dchisq(x, df_list[i], ncp=0), add=F, xlim=c(0, 16),
        ylim=c(0, 0.8), xlab="chisq", ylab="f(chisq)",
        main=paste("df=", df_list[i])) }
```



2. 확률분포표

자유도	P=0.99	0.95	0.90	0.10	0.05	0.01
1	0.000157	0.00393	0.0158	2.706	3.841	6.635
2	0.0201	0.103	0.211	4.605	5.991	9.210
3	0.115	0.352	0.584	6.251	7.815	11.341
4	0.297	0.711	1.064	7.779	9.488	13.277
5	0.554	1.145	1.610	9.236	11.070	15.086
6	0.872	1.635	2.204	10.645	12.592	16.812
7	1.239	2.167	2.833	12.017	14.067	18.475
8	1.646	2.733	3.490	13.362	15.507	20.090
9	2.088	3.325	4.168	14.684	16.919	21.666
10	2.558	3.940	4.865	15.987	18.307	23.209

b1-ch4-13.R

```
df<-10
chi1<-numeric(df);chi2<-numeric(df);chi3<-numeric(df)
chi4<-numeric(df);chi5<-numeric(df);chi6<-numeric(df)
for(j in 1:df) { chi1[j]<-qchisq(0.01,j) }
for(j in 1:df) { chi2[j]<-qchisq(0.05,j) }
for(j in 1:df) { chi3[j]<-qchisq(0.1,j) }
for(j in 1:df) { chi4[j]<-qchisq(0.9,j) }
for(j in 1:df) { chi5[j]<-qchisq(0.95,j) }
for(j in 1:df) { chi6[j]<-qchisq(0.99,j) }
round((chi<-cbind(chi1,chi2,chi3,chi4,chi5, chi6)),digits
=4)
```



```
> round((chi<-cbind(chi1,chi2,chi3,chi4,chi5, chi6)),digits=4)
      chi1    chi2    chi3    chi4    chi5    chi6
[1,] 0.0002 0.0039 0.0158 2.7055 3.8415 6.6349
[2,] 0.0201 0.1026 0.2107 4.6052 5.9915 9.2103
[3,] 0.1148 0.3518 0.5844 6.2514 7.8147 11.3449
[4,] 0.2971 0.7107 1.0636 7.7794 9.4877 13.2767
[5,] 0.5543 1.1455 1.6103 9.2364 11.0705 15.0863
[6,] 0.8721 1.6354 2.2041 10.6446 12.5916 16.8119
[7,] 1.2390 2.1673 2.8331 12.0170 14.0671 18.4753
[8,] 1.6465 2.7326 3.4895 13.3616 15.5073 20.0902
[9,] 2.0879 3.3251 4.1682 14.6837 16.9190 21.6660
[10,] 2.5582 3.9403 4.8652 15.9872 18.3070 23.2093
```

1. 확률분포

- 서로 독립적인 표준정규분포와 χ^2 -분포에 의해 t-분포가 도출
 $X \sim t(n-1)$

b1-ch4-14.R

```
set.seed(12345)
```

```
n<-10000;
```

```
z<-rnorm(n,0,1)
```

```
z1<-rnorm(n,0,1)
```

```
z2<-rnorm(n,0,1)
```

```
z3<-rnorm(n,0,1)
```

```
z4<-rnorm(n,0,1)
```

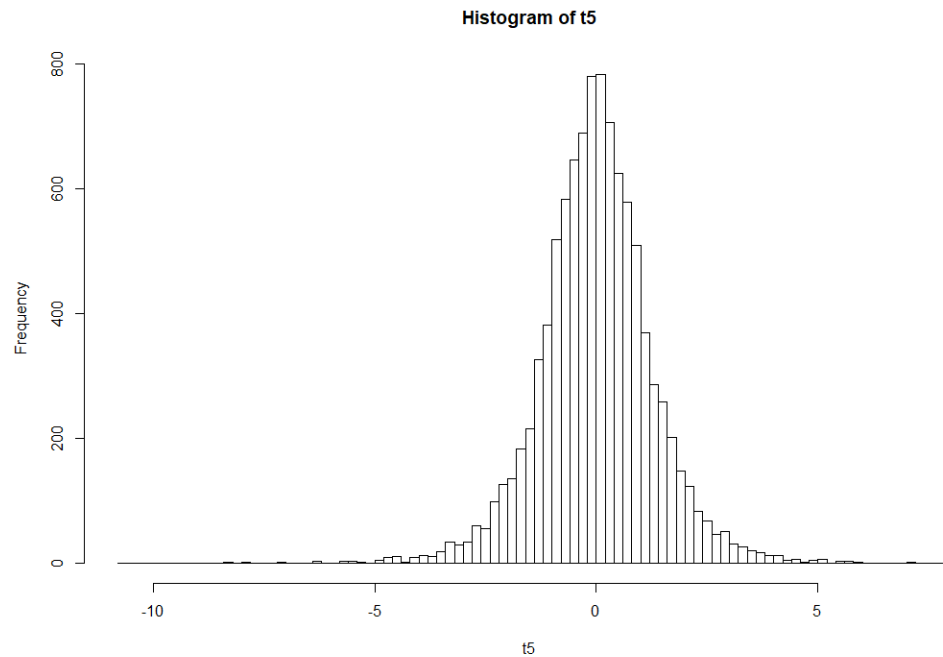
```
z5<-rnorm(n,0,1)
```

```
chi5<-z1^2+z2^2+z3^2+z4^2+z5^2
```

```
sqchi5<-sqrt(chi5/5)
```

```
t5<-z/sqchi5
```

```
hist(t5, breaks=100)
```



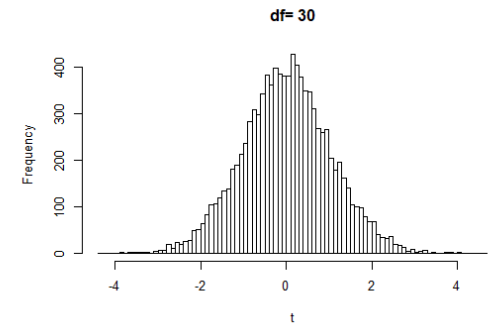
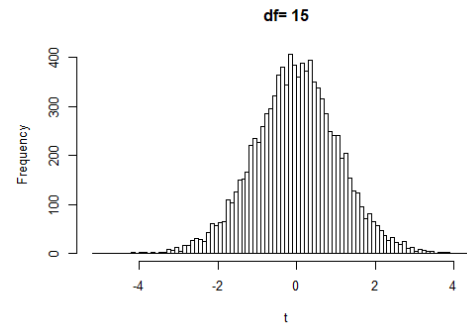
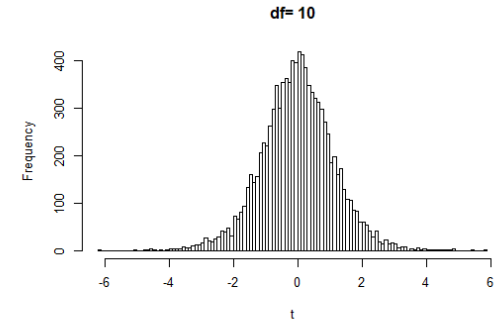
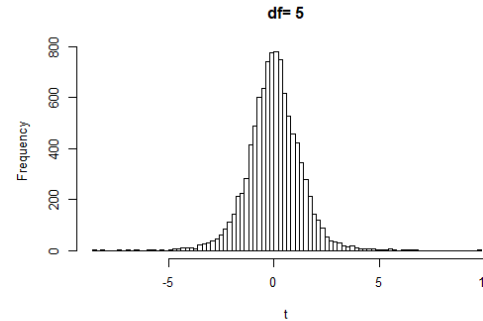
- 자유도에 따라 t-분포의 모양이 달라지는데 자유도가 30 이상이면 표준정규분포와 거의 같아짐

b1-ch4-15.R

```
set.seed(12345)

n<-10000;
df_list<-c(5,10,15,30)
par(mfrow=c(2,2))

for (i in 1:length(df_list)) {
  hist(rt(n, df=df_list[i], ncp=0), breaks=100, xlab
    ="t", main=paste("df=", df_list[i]))
}
```



b1-ch4-16.R

```
n_list<-c(2,5,10,30) # 표본수(n)
```

```
df_list<-n_list-1 # 자유도
```

```
curve(dt(x, 1, ncp=0), add=T, col="blue", xlim=c(-4, 4), ylim=c(0, 0.5), xlab="t", ylab="f(t)")
```

```
curve(dt(x, 4, ncp=0), add=T, col="red", xlim=c(-4, 4), ylim=c(0, 0.5), xlab="t", ylab="f(t)")
```

```
curve(dt(x, 9, ncp=0), add=T, col="green", xlim=c(-4, 4), ylim=c(0, 0.5), xlab="t", ylab="f(t)")
```

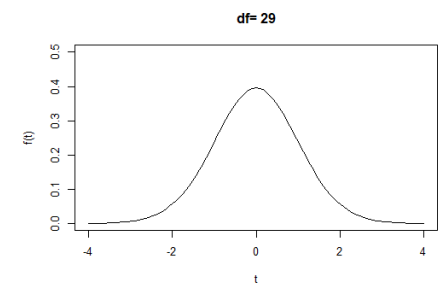
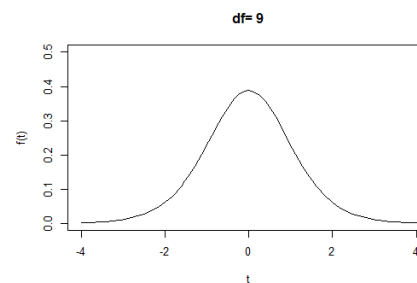
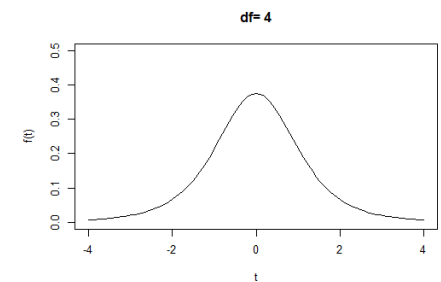
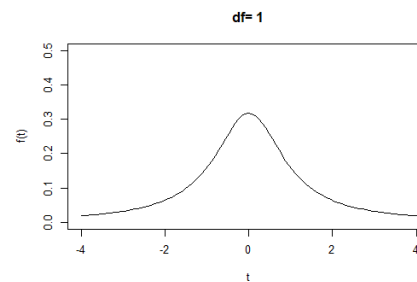
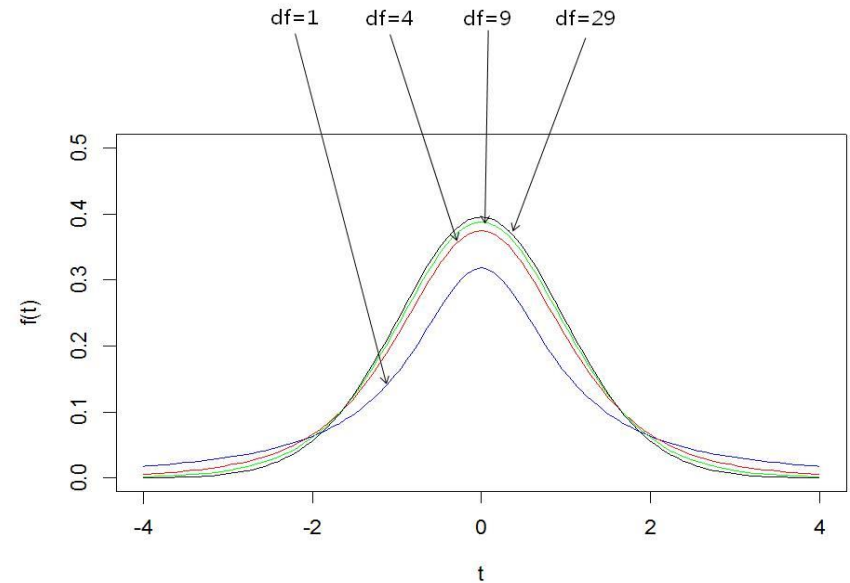
```
curve(dt(x, 29, ncp=0), add=T, col="black", xlim=c(-4, 4), ylim=c(0, 0.5), xlab="t", ylab="f(t)")
```

```
par(mfrow=c(2,2))
```

```
for (i in 1:length(df_list)) {
```

```
  curve(dt(x, df_list[i], ncp=0), xlim=c(-4, 4), ylim=c(0, 0.5), xlab="t", ylab="f(t)", main=paste("df=", df_list[i]))
```

```
}
```



2. 확률분포표

$\begin{matrix} p \\ \backslash \\ v \end{matrix}$	0.1	0.05	0.025	0.01	0.005
1	3.078	6.314	12.706	31.821	63.657
2	1.886	2.920	4.303	6.965	9.923
3	0.1638	2.353	3.182	4.541	5.841
4	1.533	2.132	2.776	3.747	4.604
5	1.476	2.015	2.571	3.365	4.032
6	1.440	1.943	2.447	3.143	3.707
7	1.415	1.895	2.365	2.998	3.499
8	1.397	1.860	2.306	2.896	3.355
9	1.383	1.833	2.262	2.821	3.250

b1-ch4-17.R

```
t11<-rep(NA,9)
t12<-rep(NA,9)
t13<-rep(NA,9)
t14<-rep(NA,9)
t15<-rep(NA,9)
```

```
for(i in 1:9) { t11[i]<-qt(0.9, i) }
for(i in 1:9) { t12[i]<-qt(0.95, i) }
for(i in 1:9) { t13[i]<-qt(0.975,i) }
for(i in 1:9) { t14[i]<-qt(0.99, i) }
for(i in 1:9) { t15[i]<-qt(0.995, i) }
```

```
round((poi<-cbind(t11,t12,t13,t14,t15)), digits=3)
```



```
> round((poi<-cbind(t11,t12,t13,t14,t15)), digits=3)
      t11  t12  t13  t14  t15
[1,] 3.078 6.314 12.706 31.821 63.657
[2,] 1.886 2.920 4.303 6.965 9.925
[3,] 1.638 2.353 3.182 4.541 5.841
[4,] 1.533 2.132 2.776 3.747 4.604
[5,] 1.476 2.015 2.571 3.365 4.032
[6,] 1.440 1.943 2.447 3.143 3.707
[7,] 1.415 1.895 2.365 2.998 3.499
[8,] 1.397 1.860 2.306 2.896 3.355
[9,] 1.383 1.833 2.262 2.821 3.250
```

1. 확률분포

- 두 개의 독립적인 χ^2 -분포에 의해 F-분포가 도출

$$X \sim F(n_1-1, n_2-1)$$

b1-ch4-18.R

```
set.seed(12345)
```

```
n<-10000;
```

```
z1<-rnorm(n,0,1)
```

```
z2<-rnorm(n,0,1)
```

```
z3<-rnorm(n,0,1)
```

```
z4<-rnorm(n,0,1)
```

```
z5<-rnorm(n,0,1)
```

```
z6<-rnorm(n,0,1)
```

```
z7<-rnorm(n,0,1)
```

```
z8<-rnorm(n,0,1)
```

```
z9<-rnorm(n,0,1)
```

```
z10<-rnorm(n,0,1)
```

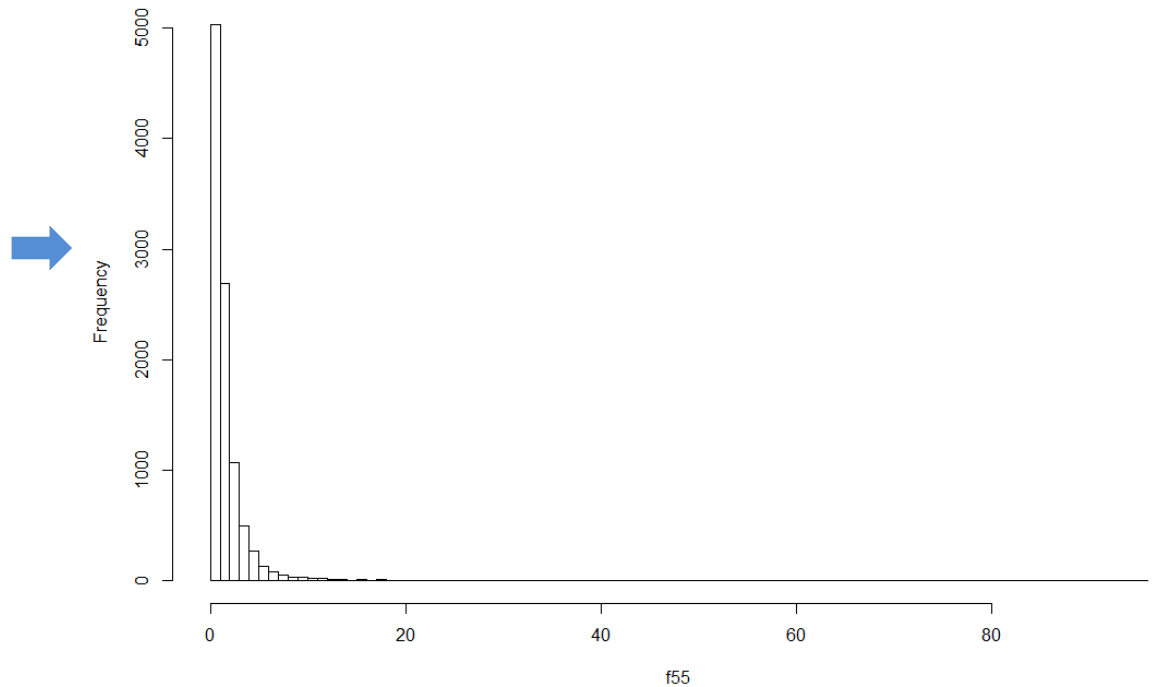
```
chi15<-z1^2+z2^2+z3^2+z4^2+z5^2
```

```
chi25<-z6^2+z7^2+z8^2+z9^2+z10^2
```

```
f55<-(chi15/5)/(chi25/5)
```

```
hist(f55, breaks=100)
```

Histogram of f55



- 분자 및 분모의 자유도에 같아지면서 커질수록 좌우대칭 분포와 비슷하게 됨

b1-ch4-19.R

```
set.seed(12345)
```

```
n<-10000;
```

```
df1_list<-c(5,9,15,38)
```

```
df2_list<-c(10,10,20,40)
```

```
par(mfrow=c(2,2))
```

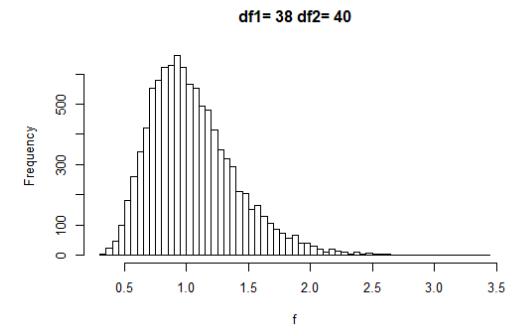
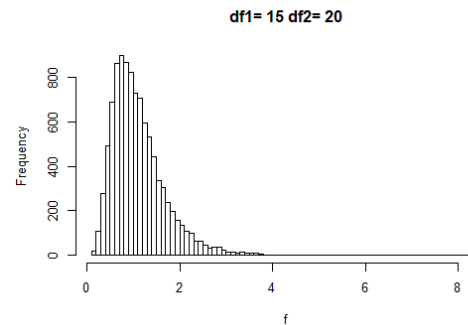
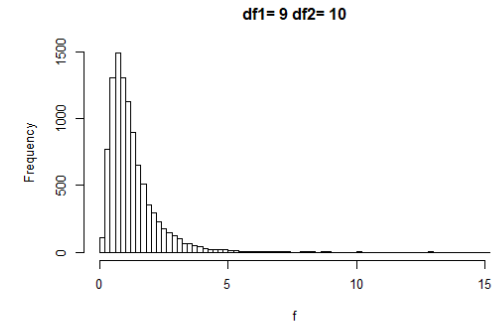
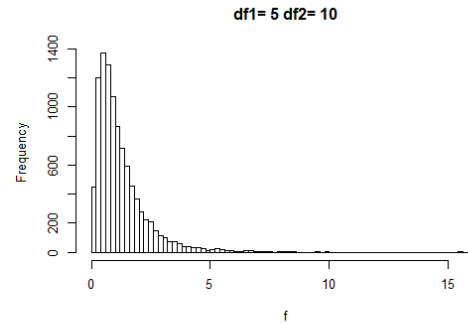
```
for (i in 1:length(df_list)) {
```

```
  hist(rf(n, df1=df1_list[i], df2=df2_list[i], ncp=0),
```

```
    breaks=100, xlab="f", main=paste("df1=", df1_list[
```

```
    i], "df2=", df2_list[i]))
```

```
}
```

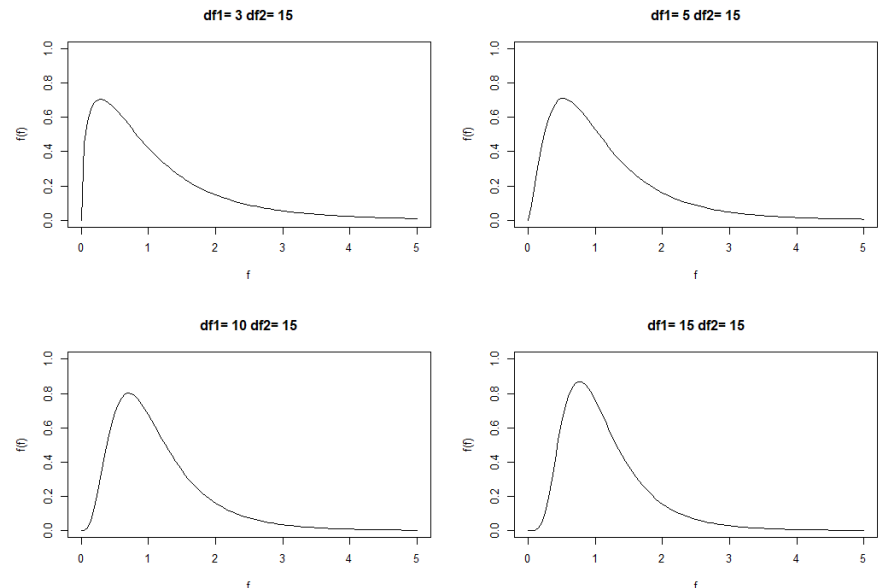
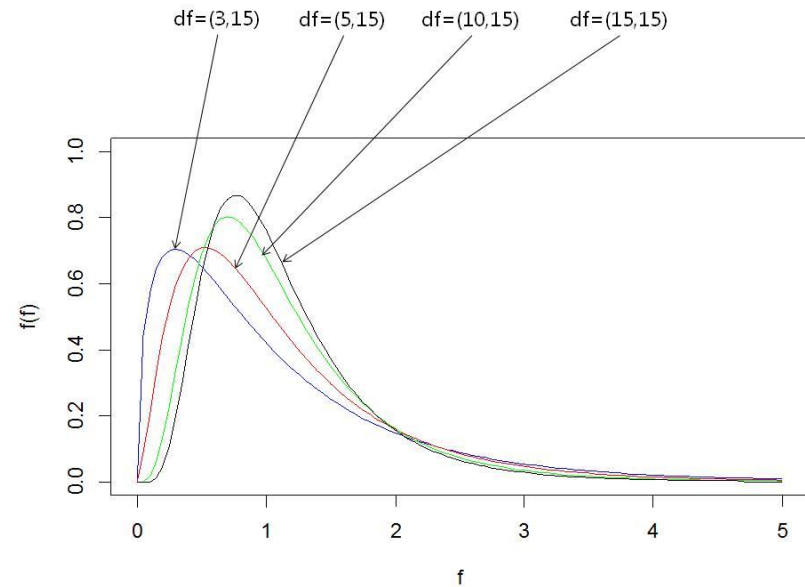


b1-ch4-20.R

```
curve(df(x, 3, 15, ncp=0), col="blue", xlim=c(0,5), y
lim=c(0,1), xlab="f", ylab="f(f)")
curve(df(x, 5, 15, ncp=0), add=T, col="red", xlim=c
(0,5), ylim=c(0,1), xlab="f", ylab="f(f)")
curve(df(x, 10, 15, ncp=0), add=T, col="green", xli
m=c(0,5), ylim=c(0,1), xlab="f", ylab="f(f)")
curve(df(x, 15, 15, ncp=0), add=T, col="black", xli
m=c(0,5), ylim=c(0,1), xlab="f", ylab="f(f)")
```

```
df1_list<-c(3,5,10,15)
df2_list<-c(15,15,15,15)
par(mfrow=c(2,2))
```

```
for (i in 1:length(df1_list)) {
  curve(df(x, df1=df1_list[i], df2=df2_list[i], ncp=0)
, xlim=c(0,5), ylim=c(0,1), xlab="f", ylab="f(f)", mai
n=paste("df1=", df1_list[i], "df2=", df2_list[i]))
}
```



2. 확률분포표

b1-ch4-21.R

```
11<-rep(NA,10);f12<-rep(NA,10);f13<-rep(NA,10);
f14<-rep(NA,10);f15<-rep(NA,10);f16<-rep(NA,10);
f17<-rep(NA,10);f18<-rep(NA,10);f19<-rep(NA,10);
f110<-rep(NA,10)

for(i in 1:10) { f11[i]<-qf(0.95, 1, i)}
for(i in 1:10) { f12[i]<-qf(0.95, 2, i)}
for(i in 1:10) { f13[i]<-qf(0.95, 3, i)}
for(i in 1:10) { f14[i]<-qf(0.95, 4, i)}
for(i in 1:10) { f15[i]<-qf(0.95, 5, i)}
for(i in 1:10) { f16[i]<-qf(0.95, 6, i)}
for(i in 1:10) { f17[i]<-qf(0.95, 7, i)}
for(i in 1:10) { f18[i]<-qf(0.95, 8, i)}
for(i in 1:10) { f19[i]<-qf(0.95, 9, i)}
for(i in 1:10) { f110[i]<-qf(0.95, 10, i)}

round((poi<-cbind(f11,f12,f13,f14,f15,f16,f17,f18,f19,f110)), digits=2)
```

$v_1 \backslash v_2$	1	2	3	4	5	6	7	8	9	10
1	161.45	199.50	215.71	224.58	230.16	233.99	236.77	238.88	240.54	241.88
2	18.51	19.00	19.16	19.25	19.30	19.33	19.35	19.37	19.38	19.40
3	10.13	9.55	9.28	9.12	9.01	8.94	8.89	8.85	8.81	8.76
4	7.71	6.94	6.59	6.39	6.26	6.16	6.09	6.04	6.00	5.96
5	6.61	5.79	5.41	5.19	5.05	4.95	4.48	4.82	4.77	4.74
6	5.99	4.74	7.35	4.12	3.94	3.87	3.79	3.73	3.68	3.64
7	5.59	4.74	4.35	4.12	3.97	3.87	3.79	3.73	3.68	3.64
8	5.32	4.46	4.07	3.84	3.69	3.58	3.50	3.44	3.39	3.35
9	5.12	4.26	3.86	3.63	3.48	3.37	3.29	3.23	3.18	3.14
10	4.96	4.10	3.71	3.48	3.33	3.22	3.14	3.07	3.02	2.98



```
> round((poi<-cbind(f11,f12,f13,f14,f15,f16,f17,f18,f19,f110)), digits=2)
      f11    f12    f13    f14    f15    f16    f17    f18    f19    f110
[1,] 161.45 199.50 215.71 224.58 230.16 233.99 236.77 238.88 240.54 241.88
[2,]  18.51  19.00  19.16  19.25  19.30  19.33  19.35  19.37  19.38  19.40
[3,]  10.13   9.55   9.28   9.12   9.01   8.94   8.89   8.85   8.81   8.79
[4,]   7.71   6.94   6.59   6.39   6.26   6.16   6.09   6.04   6.00   5.96
[5,]   6.61   5.79   5.41   5.19   5.05   4.95   4.88   4.82   4.77   4.74
[6,]   5.99   5.14   4.76   4.53   4.39   4.28   4.21   4.15   4.10   4.06
[7,]   5.59   4.74   4.35   4.12   3.97   3.87   3.79   3.73   3.68   3.64
[8,]   5.32   4.46   4.07   3.84   3.69   3.58   3.50   3.44   3.39   3.35
[9,]   5.12   4.26   3.86   3.63   3.48   3.37   3.29   3.23   3.18   3.14
[10,]  4.96   4.10   3.71   3.48   3.33   3.22   3.14   3.07   3.02   2.98
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