```
*Problem 1;
data gi;
input center $ treat $ response count @@;
n response=(response='1');
datalines;
a test 1 22
              a test 0 13
a placebo 1 33 a placebo 0 19
b test 1 27
              b test 0 7
b placebo 1 18 b placebo 0 15
run;
proc freq data=gi order=data;
weight count;
tables center*treat*response / chisq cmh(mf) nocol nopct;
*Problem 2;
data rashd;
input treat $ sex $ rash count @@;
datalines;
placebo m 0 6 placebo m 1 10
                                  placebo m 2 12
                                                      placebo m 3 20
placebo f 0 7 placebo f 1 14 placebo f 2 19 placebo f 3 18
low m 0 9
                   low m 1 7
                                  low m 2 30
                                                low m 3 19
low f 0 10
                   low f 1 17
                                  low f 2 11
                                                 low f 3 16
high m 0 19
                   high
                             m 1 15
                                       high
                                                 m 2 17
                                                          high
                                                                    m 3 5
high f 0 21
                      f 1 18
                                            f 2 10
                                                               f 3 5
              high
                                  high
                                                      high
run;
data rashp;
input treat sex rash count;
datalines;
     1
         1
1
              44
         1
    0
              56
         1
              33
    0
         1
              37
1
         0
1
     1
              10
1
    0
         0
              9
1
    1
         0
              21
    0
         0
              19
1
0
    0
         1
              42
0
         1
              51
     1
0
    0
         0
              6
0
         0
              7
run;
proc freq data=rashp order=data;
weight count;
tables sex*treat*rash / chisq cmh(mf) nocol nopct;
```

```
run;
proc logistic data=rashp order=data;
freq count;
model rash= treat sex;
oddsratio sex:
run:
proc logistic data=rashp order=data;
freq count;
class treat(ref="0") sex(ref="0") / param=ref;
model rash= treat|sex / aggregate;
oddsratio sex;
run;
*Problem 3;
data teeth;
input toothex $ jaw $ insurance count @@;
datalines;
                                            none upper 2 373
                        none upper 1 193
none
         upper 0 279
                        none lower 1 83
                                            none lower 2 137
         lower 0 149
none
                   one upper 121
one upper 0 69
                                            one upper 281
one lower 0 29
                   one lower 1 44
                                       one lower 2 75
twoplus upper 0 45 twoplus upper 1 33 twoplus upper 2 24
twoplus lower 0 21 twoplus lower 1 12 twoplus lower 2 19
run;
data teeth1;
input toothex jaw $ insurance count @@;
datalines;
    upper 0 279
                   0 upper 1 193 0 upper 2 373
0
    lower 0 149
                   0 lower 1 83 0 lower 2 137
1
    upper 0 69
                   1 upper 1 21 1 upper 2 81
1
                   1 lower 1 44 1 lower 2 75
    lower 0 29
1
    upper 0 45
                   1 upper 1 33 1 upper 2 24
                   1 lower 1 12 1 lower 2 19
1
    lower 0 21
run;
*Part a;
proc freq data=teeth1 order=data;
weight count;
tables jaw*toothex*insurance / all chisq cmh(mf) nocol nopct scores=modridit;
run;
*Part c:
data teeth2;
input toothex high count @@;
datalines;
0
    1 510
              0 0 704
1
     1 156
              1 0 163
2
    1 43 2 0 111
```

```
run;
proc freq data=teeth2 order=data;
weight count;
tables high*toothex / all chisq cmh(mf) nocol nopct;
*Part d:
data teeth3;
input jaw $ toothex high count @@;
datalines;
lower 0 1 137 lower 0 0 232
lower 1 1 75 lower 1 0 73
lower 2 1 19 lower 2 0 33
upper 0 1 373 upper 0 0 472
upper 1 1 81 upper 1 0 90
upper 2 1 24 upper 2 0 78
run;
proc freq data=teeth3 order=data;
weight count;
tables jaw*high*toothex / all chisq cmh(mf) nocol nopct;
*Problem 4;
data drug;
input druga $ drugb $ drugc $ count;
datalines;
FFF6
F F U 16
FUF2
FUU4
UFF2
UFU4
UUF6
UUU6
run;
data drug2; set drug;
keep patient drug response;
retain patient 0;
do i=1 to count;
patient=patient+1;
drug='A'; response=druga; output;
drug='B'; response=drugb; output;
drug='C'; response=drugc; output;
end;
proc print data=drug2;
run;
```

```
proc logistic data=drug2;
class drug(ref="A") /param=ref;
strata patient;
model response(event="F")= drug / aggregate alpha=.01;
oddsratio drug;
run;

*Part c;
proc logistic data=drug2;
class drug(ref="A") /param=ref;
strata patient;
model response(event="F")= drug / alpha=.05;
contrast "b vs c" drug 1 -1;
run;
```