```
Call:
qlm(formula = fram_1$cvd ~ ., data = fram_1)
Deviance Residuals:
    Min
               10
                     Median
                                   30
                                            Max
-0.78619 -0.03438 -0.01520
                              0.01024
                                        0.60653
Coefficients: (1 not defined because of singularities)
              Estimate Std. Error t value Pr(>|t|)
                                    1.629 0.103498
             1.000e-01 6.142e-02
(Intercept)
sex
            -1.238e-02
                        7.227e-03 -1.713 0.086791 .
            -5.455e-05 7.497e-05 -0.728 0.466917
totchol
age
             8.560e-06 1.205e-03
                                    0.007 0.994334
sysbp
            -3.612e-04
                        2.757e-04 -1.310 0.190304
             4.367e-04 4.503e-04
                                    0.970 0.332225
diabp
             2.783e-02 9.940e-03
                                    2.800 0.005153 **
cursmoke
cigpday
            -9.670e-04 4.349e-04 -2.224 0.026274 *
bmi
             5.775e-04
                        1.485e-03
                                    0.389 0.697342
             4.168e-02 2.496e-02
                                    1.670 0.095069 .
diabetes
            -7.511e-02 1.899e-02 -3.955 7.88e-05 ***
bpmeds
            -2.429e-04 2.731e-04 -0.890 0.373824
heartrte
glucose
             2.715e-04 1.685e-04
                                   1.612 0.107148
prevchd
            -1.977e-01 4.899e-02 -4.036 5.61e-05 ***
             8.609e-02 4.571e-02
                                   1.883 0.059781 .
prevap
             1.649e-01 3.890e-02
                                    4.239 2.34e-05 ***
prevmi
            -2.415e-01 4.410e-02 -5.477 4.83e-08 ***
prevstrk
prevhyp
            -3.271e-03 1.099e-02 -0.298 0.765884
death
            -1.545e-02 1.189e-02 -1.299 0.194080
            -2.410e-01 1.833e-02 -13.146 < 2e-16 ***
angina
            -5.401e-02 2.095e-02 -2.578 0.010000 **
hospmi
             3.173e-01 2.210e-02 14.358 < 2e-16 ***
mi fchd
             5.844e-01 1.908e-02 30.626 < 2e-16 ***
anychd
stroke
             6.456e-01 1.386e-02 46.593 < 2e-16 ***
             4.766e-03 1.074e-02
                                    0.444 0.657322
hyperten
            -2.093e-05 5.663e-06 -3.697 0.000224 ***
timeap
                        7.250e-06 -4.218 2.56e-05 ***
timemi
            -3.058e-05
                                   7.721 1.74e-14 ***
timemifc
             5.835e-05 7.557e-06
timechd
             8.633e-05 6.429e-06 13.427 < 2e-16 ***
timestrk
             8.841e-05 5.403e-06 16.364 < 2e-16 ***
                        3.871e-06 -37.744 < 2e-16 ***
            -1.461e-04
timecvd
            -4.096e-05 5.209e-06 -7.864 5.79e-15 ***
timedth
timehvp
            -3.520e-06 1.853e-06 -1.899 0.057673 .
bmicat
            -6.516e-03 8.540e-03 -0.763 0.445522
aaecat
             7.474e-03
                        1.084e-02
                                    0.689 0.490612
obese_normal
                    NA
                               NA
                                       NA
Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' '1
(Dispersion parameter for gaussian family taken to be 0.02140781)
Null deviance: 408.31 on 2222 degrees of freedom
Residual deviance: 46.84 on 2188 degrees of freedom
  (2211 observations deleted due to missingness)
AIC: -2199.9
```

Number of Fisher Scoring iterations: 2