

STA409

Answer to Assignment 3

12112627

李乐平

1. Solution.

(1).

列	diabetes	asthma	cancer	n
1	1	0	0	
2	0	0	0	
3	0	0	0	
4	0	0	0	
5	0	0	0	
6	0	0	0	
7	0	0	1	
8	0	0	0	
9	0	1	0	
10	0	0	0	
11	1	0	0	
12	0	0	0	
13	0	0	0	
14	0	1	0	
15	0	0	0	
16	0	0	0	

(2). The family-wise error rate (FWER) is computed by

$$\text{FWER} = 1 - (1 - 0.05)^{42} = 0.8840$$

That means we only have a chance of 11.6% that we do not make any Type I Error without any adjustment.

(3).

	Diabetes	Asthma	Cancer	Multiple Sclerosis	Thyroid Disease	Liver Disease	Arthritis
Leo	0.7953	0.8187	0.9528	0.1092	0.4827	0.1701	0.9060
Purple	0.6143	0.2577	0.4849	0.2487	0.8314	0.7887	0.1878
0:00-1:00	0.0628	0.4728	0.1874	0.0088	0.8947	0.0103	0.5419
Red Hair	0.2353	0.5705	0.7993	0.2556	0.3873	0.3861	0.7048
First Name C	0.4391	0.3479	0.3406	0.7265	0.2134	0.6235	0.8320
Summer	0.6329	0.3941	0.8162	0.3259	0.9229	0.3629	0.3189

In the table above, 2 combinations of disease and risk factor are determined to be dependent, which are actually Type I Errors. That implies it makes sense to do adjustment to reduce the chance of Type I Error happens under the increment of number of experiments.

2. Solution.

(1). The ANOVA table and coefficients table are given by

ANOVA Table					
Source	DF	Sum of Squares	Mean Square	F-value	p-value
Model	4	23665352	5916338	22.9782	5.0715×10^{-13}

Error	88	22657938	257476.5682		
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where $MS = \text{sum_of_squares} / DF$; $F\text{-value} = MS_{\text{Model}} / MS_{\text{Error}}$; $p\text{-value} = \Pr(F(DF_{\text{Model}}, DF_{\text{Error}}) > F)$.

Coefficients Table				
Parameter	Estimate	Standard Error	t-value	p-value
Intercept	3526.4	327.7	10.7610	0
Gender	722.5	117.8	6.1333	2.3917×10^{-8}
Education	90.02	24.69	3.6460	4.5033×10^{-4}
Experience	1.2690	0.5877	2.1593	3.3547×10^{-2}
Months	23.406	5.201	4.5003	2.0765×10^{-5}

where $t\text{-value} = \text{Estimate} / SE$; $p\text{-value} = \Pr(|t(DF_{\text{Error}})| > |t|)$

In this question, the model for test is given by

$$\text{Salary} = \beta_0 + \beta_1 \text{Gender} + \beta_2 \text{Education} + \beta_3 \text{Experience} + \beta_4 \text{Months} + \varepsilon$$

The overall F-test is rejected, indicating at least one of β s are significantly different from 0. And t-tests show that every predictor variable is significant.

(2). The R^2 and adjusted R^2 are computed by

$$R^2 = 1 - \frac{SSE}{SST} = 1 - \frac{23665652}{46323290} = 0.5109$$

$$\text{Adjusted } R^2 = 1 - \frac{SSE/(n-p-1)}{SST/(n-1)} = 1 - \frac{23665652/88}{46323290/92} = 0.4886$$

(3). A partial F-test is utilized to test which of the full model and the reduced model is better.

$$F = \frac{(SST_R - SST_F)/(p-k)}{SST_F/(n-p-1)} = \frac{(38460756 - 22657938)/3}{22657938/88} = 20.4586$$

Because $\Pr(F(3, 88) > F) = 3.8052 \times 10^{-10}$, hence the reduced model is rejected.

3. Solution.

Now we want to show that $\hat{\beta}^{\text{WLS}} = (X^T W X)^{-1} X^T W y$ is the best linear unbiased estimate of β , where $W = [\text{Var}(y)]^{-1}$.

The variance of $\hat{\beta}^{\text{WLS}}$ is

$$\begin{aligned} \text{Var}(\hat{\beta}^{\text{WLS}}) &= ((X^T W X)^{-1} X^T W) \text{Var}(y) ((X^T W X)^{-1} X^T W)^T \\ &= (X^T W X)^{-1} \end{aligned}$$

Assume there is another linear estimation of β : $\tilde{\beta} = Ay$, $A = (X^T W X)^{-1} X^T W + B$, then it must satisfies

$$E(\tilde{\beta}) = E[((X^T W X)^{-1} X^T W + B)(X\beta + \varepsilon)] = (I + BX)\beta = \beta,$$

i.e. $BX = 0$, or it would not be unbiased estimator of β .

Yet

$$\begin{aligned} \text{Var}(\tilde{\beta}) &= ((X^T W X)^{-1} X^T W + B) W^{-1} ((X^T W X)^{-1} X^T W + B)^T \\ &= (X^T W X)^{-1} + B W^{-1} B^T \\ &= \text{Var}(\hat{\beta}^{\text{WLS}}) + B W^{-1} B^T \\ &\geq \text{Var}(\hat{\beta}^{\text{WLS}}) \end{aligned}$$

then it is clear that $\hat{\beta}^{\text{WLS}}$ is the best linear unbiased estimate of β .

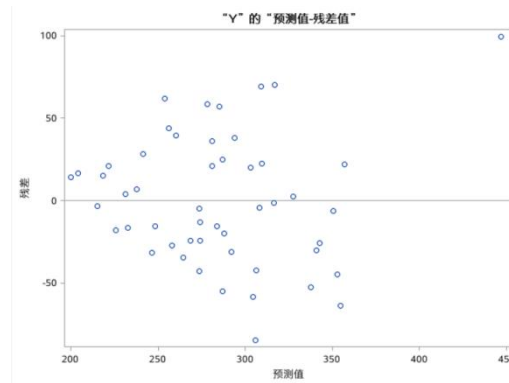
4. Solution.

(1). The fitted linear model is stated by

$$Y = -556.5680 + 0.0724X_1 + 1.5521X_2 - 0.0043X_3 + \varepsilon$$

And the assumptions are checked as follows.

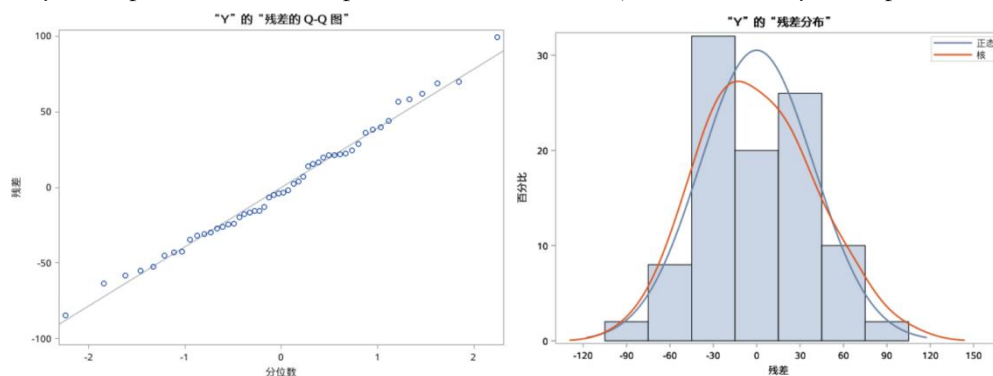
Linearity assumption: The predicted value - residual plot shows that the mean of the residuals are roughly 0 at any place, which implies the validity of linearity assumption.



Homoscedasticity assumption: The predicted value - residual plot does not show homoscedasticity since it is narrow at left side and wide at right side. Also, the White test and Breusch-Pagan test reject the homoscedasticity assumption with p-values less than 0.01.

异方差性检验					
方程	检验	统计量	自由度	Pr > 卡方	变量
Y	White 检验	22.68	9	0.0070	所有变量的叉积
	Breusch-Pagan	15.59	3	0.0014	1, X1, X2, X3

Normality: The Q-Q plot as well as distribution of residuals does not show violation of the normality assumption. Also, the Shapiro-Wilk test does not reject the normality assumption.

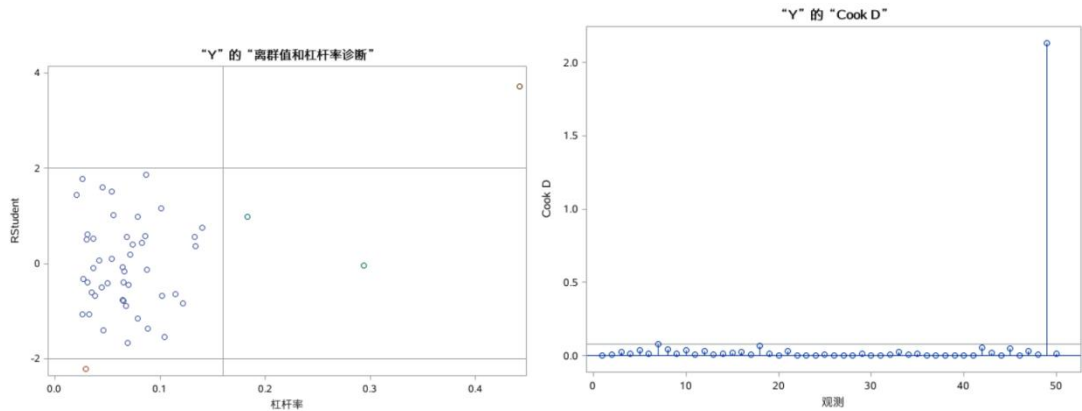


正态性检验				
检验	统计量		p 值	
Shapiro-Wilk	W	0.962619	Pr < W	0.1145
Kolmogorov-Smirnov	D	0.077587	Pr > D	>0.1500
Cramer-von Mises	W-Sq	0.0368	Pr > W-Sq	>0.2500
Anderson-Darling	A-Sq	0.295345	Pr > A-Sq	>0.2500

Independence: The data are collected from 50 states independently, hence we can assume the independence of the data.

(2). Notice that the state Alaska has both high leverage, Cook's distance and studentized residual, which shows Alaska is an outlier and influential observation. Also shown in the plot.

	State	leverage	cookD	rstudent
1	AK	0.4419099163	2.1327949114	3.7099223829
2	UT	0.2941338295	0.0002247544	-0.04594205
3	NM	0.1829094976	0.054304552	0.9847410811
4	VT	0.1400044576	0.0235191903	0.7566427892
5	WV	0.1334181929	0.0053186159	0.3682198446
6	FL	0.1329380512	0.0121187746	0.5580649664
7	MS	0.121715957	0.024280392	-0.834377609
8	LA	0.1150154266	0.0134134888	-0.638374691
9	ND	0.1046879429	0.0682663128	-1.551362037
10	CT	0.101921737	0.01292351	-0.670860476
11	RI	0.1011891856	0.0370314221	1.1510950224
12	NJ	0.088828501	0.0452504612	-1.375748431
13	TX	0.0872943074	0.0003731912	-0.123585963
14	NY	0.0864580927	0.077778339	1.8610093979
15	CA	0.0861878909	0.0078594908	0.57311144



Considering Alaska is away from the mainland of U.S., it may be dropped from the analysis.

(3). If we only refit the model using weighted GLM without dropping outlier to solve the heteroscedasticity, the refitted model is stated as

$$Y = -423.0778 + 0.0649X_1 + 1.1954X_2 + 0.0223X_3 + \varepsilon$$

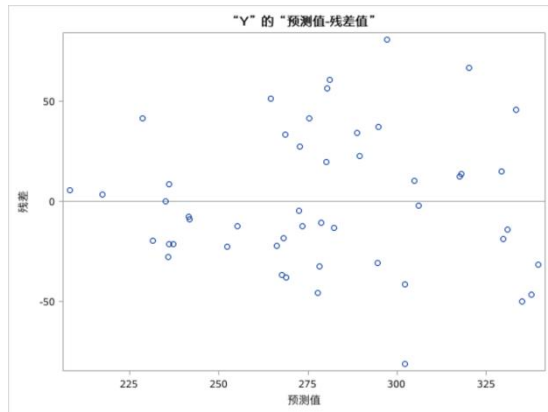
参数	估计	标准误差	t 值	Pr > t
截距	-423.0777720	116.1533088	-3.64	0.0007
X1	0.0648833	0.0114096	5.69	<.0001
X2	1.1953871	0.2978225	4.01	0.0002
X3	0.0222879	0.0468463	0.48	0.6365

However, after dropping the data of Alaska, the fitted linear model is stated as

$$Y = -277.5773 + 0.0483X_1 + 0.8869X_2 + 0.0668X_3 + \varepsilon$$

This time the homoscedasticity assumption is accepted. And the other 3 assumptions are also valid (omitted).

参数估计						
变量	标签	自由度	参数估计	标准误差	t 值	Pr > t
Intercept	Intercept	1	-277.57731	132.42286	-2.10	0.0417
X1	X1	1	0.04829	0.01215	3.98	0.0003
X2	X2	1	0.88693	0.33114	2.68	0.0103
X3	X3	1	0.06679	0.04934	1.35	0.1826



异方差性检验					
方程	检验	统计量	自由度	Pr > 卡方	变量
Y	White 检验	8.69	9	0.4665	所有变量的叉积
	Breusch-Pagan	5.03	3	0.1698	1, X1, X2, X3

Compared with the former fitted model,

$$Y = -556.5680 + 0.0724X_1 + 1.5521X_2 - 0.0043X_3 + \varepsilon$$

all the assumptions are accepted, and the fitted coefficients differs significantly, which indicates the Alaska row is indeed an outlier.

5. Solution.

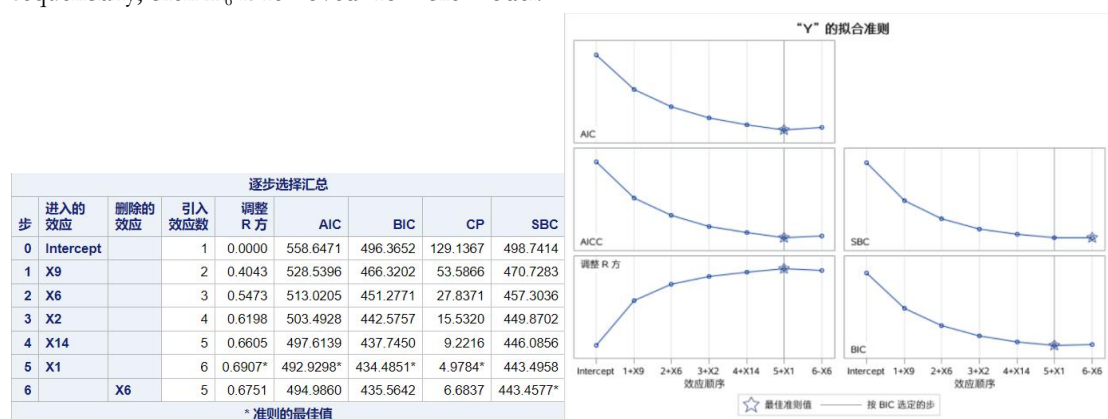
(1). By the ranked table of pairwise Pearson correlations, we can observe that X_{12} and X_{13} are highly correlated, whose correlation coefficient is 0.98384.

Pearson 相关系数, N = 60 Prob > r , H0: Rho=0																
Y	Y	X9	X6	X1	X7	X14	X11	X5	X10	X3	X8	X12	X4	X15	X13	X2
Y	1.00000	0.64374 <.0001	-0.51099 <.0001	0.50950 <.0001	-0.42682 0.0007	0.42590 0.0007	0.41049 0.0011	0.35731 0.0051	-0.28480 0.0274	0.27702 0.0321	0.26550 0.0403	-0.17724 0.1755	-0.17459 0.1821	-0.08850 0.5013	-0.07738 0.5568	-0.03002 0.8199
X1	X1	X12	Y	X11	X3	X7	X6	X13	X9	X10	X5	X14	X4	X2	X15	X8
X1	1.00000	-0.53176 <.0001	0.50950 <.0001	0.50659 <.0001	0.50327 <.0001	-0.49076 <.0001	-0.49043 <.0001	-0.48732 <.0001	0.41320 0.0010	-0.29729 0.0211	0.26344 0.0420	-0.10692 0.4161	0.10111 0.4421	0.09221 0.4835	-0.07734 0.5570	-0.00352 0.9787
X2	X2	X11	X9	X4	X12	X3	X13	X10	X5	X6	X14	X8	X1	X15	Y	X7
X2	1.00000	0.56531 <.0001	0.45377 0.0003	-0.39810 0.0016	0.35081 0.0060	0.34628 0.0067	0.32101 0.0124	0.23799 0.0671	-0.20921 0.1087	0.11628 0.3763	-0.10781 0.4123	-0.10005 0.4469	0.09221 0.4835	0.06787 0.6064	-0.03002 0.8199	0.01485 0.9103
X3	X3	X11	X9	X1	X15	X4	X7	X12	X2	X13	Y	X5	X6	X14	X8	X10
X3	1.00000	0.61931 <.0001	0.57531 <.0001	0.50327 <.0001	-0.45281 0.0003	-0.43404 0.0005	-0.41503 0.0010	-0.35649 0.0052	0.34628 0.0067	-0.33767 0.0083	0.27702 0.0321	0.26228 0.0429	-0.23854 0.0664	-0.09935 0.4501	-0.06099 0.6434	-0.02141 0.8710
X4	X4	X9	X5	X3	X2	X11	Y	X8	X6	X10	X15	X1	X7	X12	X14	X13
X4	1.00000	-0.63782 <.0001	-0.50909 <.0001	-0.43404 0.0005	-0.39810 0.0016	-0.30977 0.0160	-0.17459 0.1821	0.16199 0.2162	-0.13886 0.2900	-0.11771 0.3704	0.11243 0.3924	0.10111 0.4421	0.06501 0.6217	-0.02049 0.8765	0.01725 0.8959	-0.00208 0.9874
X5	X5	X4	X10	X9	X7	X6	X12	X13	Y	X1	X3	X11	X2	X8	X15	X14
X5	1.00000	-0.50909 <.0001	-0.42572 0.0007	0.41941 0.0009	-0.41059 0.0011	-0.39507 0.0018	-0.38821 0.0022	-0.35843 0.0049	0.35731 0.0051	0.26344 0.0420	0.26228 0.0429	0.25990 0.0449	-0.20921 0.1087	-0.18433 0.1586	-0.13574 0.3011	-0.00408 0.9753
X6	X6	X10	X7	Y	X1	X11	X5	X12	X8	X3	X14	X13	X15	X9	X4	X2
X6	1.00000	0.70320 <.0001	0.55224 <.0001	-0.51099 <.0001	-0.49043 <.0001	-0.40334 0.0014	-0.39507 0.0018	0.28683 0.0263	-0.24388 0.0604	-0.23854 0.0664	-0.23435 0.0715	0.22440 0.0848	-0.20877 0.1094	0.17649 0.1774	-0.13886 0.2900	0.11628 0.3763
X7	X7	X11	X6	X1	Y	X3	X5	X9	X12	X13	X10	X8	X15	X14	X4	X2
X7	1.00000	-0.68068 <.0001	0.55224 <.0001	-0.49076 <.0001	-0.42682 0.0007	-0.41503 0.0010	-0.41059 0.0011	0.38677 0.0023	0.34825 0.0064	0.33875 0.0081	0.33875 0.0081	0.18188 0.1643	0.12190 0.3535	0.11795 0.3694	0.06501 0.6217	0.01485 0.9103
X8	X8	X14	Y	X6	X5	X7	X13	X11	X4	X15	X12	X2	X3	X10	X9	X1
X8	1.00000	0.43209 0.0006	0.26550 0.0403	-0.24388 0.0604	-0.18433 0.1586	0.18188 0.1643	0.16531 0.2069	-0.16295 0.2135	0.16199 0.2162	-0.12498 0.3414	0.12028 0.3600	-0.10005 0.4469	-0.06099 0.6434	-0.03177 0.8096	-0.00568 0.9657	-0.00352 0.9787
X9	X9	X11	Y	X4	X3	X2	X5	X1	X7	X6	X14	X15	X12	X13	X8	X10
X9	1.00000	0.70492 <.0001	0.64374 <.0001	-0.63782 <.0001	0.57531 <.0001	0.45377 0.0003	0.41941 0.0009	0.41320 0.0010	-0.41033 0.0011	-0.20877 0.1094	0.15929 0.2241	-0.11796 0.3694	-0.02586 0.8445	0.01839 0.8891	-0.00568 0.9657	-0.00439 0.9735
X10	X10	X6	X5	X7	X1	Y	X2	X12	X11	X13	X4	X15	X8	X3	X9	X1
X10	1.00000	0.70320 <.0001	-0.42572 0.0007	0.33875 0.0081	-0.29729 0.0211	-0.28480 0.0274	0.23799 0.0671	0.20367 0.1186	-0.18516 0.1567	0.16003 0.2219	-0.11771 0.3704	-0.06846 0.6032	0.06071 0.6449	-0.03177 0.8096	-0.02141 0.8710	-0.00439 0.9735
X11	X11	X9	X7	X3	X2	X1	Y	X6	X4	X5	X10	X8	X15	X12	X13	X14
X11	1.00000	0.70492 <.0001	-0.68068 <.0001	0.61931 <.0001	0.56531 <.0001	0.50659 <.0001	0.41049 0.0011	-0.40334 0.0014	-0.30977 0.0160	0.25990 0.0449	-0.18516 0.1567	-0.16295 0.2135	-0.15222 0.2456	-0.12978 0.4356	-0.10254 0.8891	-0.09648 0.9874
X12	X12	X13	X1	X5	X7	X3	X2	X6	X14	X10	Y	X11	X8	X9	X4	X15
X12	1.00000	0.98384 <.0001	-0.53176 <.0001	-0.38821 0.0022	0.38677 0.0023	-0.35649 0.0052	0.35081 0.0060	0.28683 0.0263	0.28230 0.0289	0.20367 0.1186	-0.17724 0.1755	-0.12978 0.3230	0.12028 0.3600	-0.02586 0.8445	-0.02049 0.8765	-0.02018 0.8784
X13	X13	X12	X1	X14	X5	X7	X3	X2	X6	X8	X10	X11	Y	X15	X9	X4
X13	1.00000	0.98384 <.0001	-0.48732 <.0001	0.40939 0.0012	-0.35843 0.0064	0.34825 0.0064	-0.33767 0.0083	0.32101 0.0124	0.22440 0.0848	0.16531 0.2069	0.16003 0.2219	-0.10254 0.4356	-0.07738 0.5568	-0.04591 0.7276	0.01839 0.8891	-0.00208 0.9874
X14	X14	X8	Y	X13	X12	X6	X9	X7	X2	X1	X15	X3	X11	X10	X4	X5
X14	1.00000	0.43209 0.0006	0.26550 0.0007	0.40939 0.0012	0.28230 0.0289	-0.23435 0.0715	0.15929 0.2241	0.11795 0.3694	-0.10781 0.4123	-0.10692 0.4161	-0.10255 0.4356	-0.09935 0.4501	-0.09648 0.4633	-0.06846 0.6032	0.01725 0.8959	-0.00408 0.9753
X15	X15	X3	X6	X11	X5	X8	X7	X9	X4	X14	Y	X1	X2	X10	X13	X12
X15	1.00000	-0.45281 0.0003	0.17649 0.1774	-0.15222 0.2456	-0.13574 0.3011	-0.12498 0.3414	0.12190 0.3535	-0.11796 0.3694	0.11243 0.3924	-0.10255 0.4356	-0.08850 0.5013	-0.07734 0.5570	0.06787 0.6064	0.06071 0.6449	-0.04591 0.7276	-0.02018 0.8784

(2). By checking the tolerances and VIFs of X_{12} and X_{13} , with the criterion Tolerance < 0.1 and VIF > 10, multicollinearity does exist between X_{12} and X_{13} .

参数估计								
变量	标签	自由度	参数估计	标准误差	t 值	Pr > t	容差	方差膨胀
Intercept	Intercept	1	1763.99793	437.33031	4.03	0.0002	.	0
X1	X1	1	1.90536	0.92374	2.06	0.0451	0.24308	4.11389
X2	X2	1	-1.93762	1.10839	-1.75	0.0874	0.16277	6.14355
X3	X3	1	-3.10040	1.90167	-1.63	0.1102	0.25203	3.96777
X4	X4	1	-9.06517	8.48622	-1.07	0.2912	0.13387	7.47004
X5	X5	1	-106.83103	69.78007	-1.53	0.1329	0.23215	4.30762
X6	X6	1	-17.15689	11.86012	-1.45	0.1551	0.20574	4.86054
X7	X7	1	-0.65111	1.76777	-0.37	0.7144	0.25033	3.99478
X8	X8	1	0.00360	0.00403	0.89	0.3761	0.60303	1.65828
X9	X9	1	4.45958	1.32721	3.36	0.0016	0.14750	6.77960
X10	X10	1	-0.18715	1.66169	-0.11	0.9108	0.35192	2.84158
X11	X11	1	-0.16741	3.22730	-0.05	0.9589	0.11472	8.71707
X12	X12	1	-0.67216	0.49102	-1.37	0.1780	0.01014	98.63993
X13	X13	1	1.34010	1.00559	1.33	0.1895	0.00953	104.98240
X14	X14	1	0.08626	0.14752	0.58	0.5617	0.23647	4.22893
X15	X15	1	0.10674	1.16943	0.09	0.9277	0.52436	1.90709

(3). The process of model selection is shown as follows. X₉, X₆, X₂, X₁₄ and X₁ are added to model sequentially, then X₆ is removed from the model.



The final model is stated as

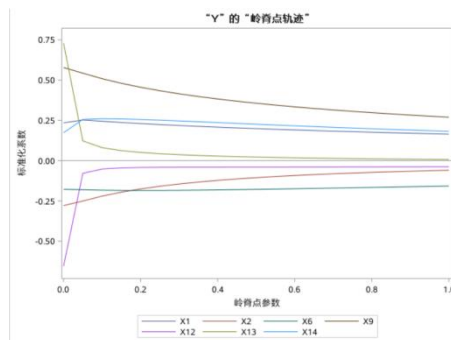
$$Y = 857.43 + 2.06 X_1 - 1.77 X_2 + 4.08 X_9 + 0.33 X_{14} + \varepsilon$$

参数估计				
参数	自由度	估计	标准误差	t 值
Intercept	1	857.431124	26.234966	32.68
X1	1	2.059246	0.524393	3.93
X2	1	-1.771640	0.527540	-3.36
X9	1	4.078669	0.671468	6.07
X14	1	0.330551	0.077299	4.28

This linear model describes the relationship between the total age-adjusted mortality rate (per 100,000) and five explanatory variables. The intercept term of 857.43 indicates that when all explanatory variables are zero, the expected total age-adjusted mortality rate is 857.43. For the coefficients of each explanatory variable, the interpretations are as follows: The coefficient of X₁ is 2.06, meaning that when the mean annual precipitation increases by one unit (in inches), the total age-adjusted mortality rate is expected to increase by 2.06 units. The coefficient of X₂ is -1.77, indicating that when the mean January temperature increases by one unit (in degrees Fahrenheit), the

total age-adjusted mortality rate is expected to decrease by 1.77 units. The coefficient of X_9 is 4.08, suggesting that when the percent of nonwhite population increases by one unit, the total age-adjusted mortality rate is expected to increase by 4.08 units. Lastly, the coefficient of X_{14} is 0.33, signifying that for each unit increase in the relative pollution potential of sulfur dioxide, the total age-adjusted mortality rate is expected to increase by 0.33 units. ϵ represents the error term, which accounts for the unexplained portion of the model.

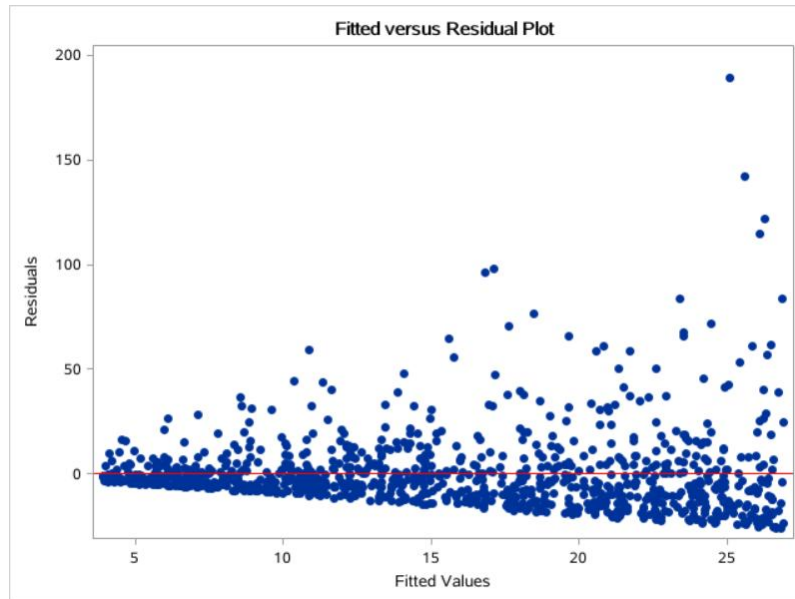
(4). The lines showing the parameter estimates of X_1 , X_2 , X_6 , X_9 , X_{12} , X_{13} and X_{14} against λ is plotted as follows.



From the plot, it's evident that as λ increases slightly, the coefficients of X_{12} and X_{13} experience rapid absolute value decreases, which aligns with their high sample correlation coefficient of 0.98 as noted in question 5.(1). These coefficients quickly converge towards zero, exhibiting a nearly symmetrical pattern about the zero line. Additionally, the effects of X_1 , X_2 , X_6 , and X_9 seem to be initially overestimated, with their absolute values decreasing as λ increases and stabilizing at non-zero values. Conversely, the effect of X_{14} appears to be initially underestimated, with its coefficient slightly increasing as λ increases. Notably, coefficients tend to stabilize around $\lambda = 0.2$, suggesting that estimates at this level of regularization are more suitable for assessing the explanatory variables' effects.

6. Solution.

(1). The Fitted versus Residual plot is shown below. Obviously it does not follow the homoscedasticity assumption.



(2). After the transformation of $Y_i' = \log Y_i$, the transformed plot is as follows. And now it seems more like a distribution of homoscedasticity.

