

Quiz1_77931863

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Question 1 - Multiple Choice Questions

Put your answers for MC questions in the table below;

MC	answer
(a)	iv
(b)	iii
(c)	iv
(d)	iv
(e)	i

Question 2

```
sample_x <- c(78, 86, 97, 91, 83, 89, 92, 88, 79, 68)
sample_x
```

```
## [1] 78 86 97 91 83 89 92 88 79 68
```

```
set.seed(1413) #DO NOT REMOVE THIS

#your code here
mean_trim <- mean(sample_x, trim=0.2)
mean_trim
```

part a)

```
## [1] 86
```

```
# add your code here
library(boot)

num_bootstraps <- 1000
bootstrap_means <- numeric(num_bootstraps)
for (i in 1:num_bootstraps) {

  bootstrap_sample <- sample(sample_x, replace = TRUE)
  trimmed_mean <- mean(bootstrap_sample, trim = 0.2)
  bootstrap_means[i] <- trimmed_mean
}
standard_error <- sd(bootstrap_means)
standard_error
```

part b)

```
## [1] 2.683526
```

```
# add your code here
critical_value <- qnorm(1 - (1 - 0.9) / 2)
upper <- mean_trim + standard_error * critical_value
lower <- mean_trim - standard_error * critical_value
lower
```

part c)

```
## [1] 81.58599
```

```
upper
```

```
## [1] 90.41401
```

Question 3 Run this code before continue. note that the data is saved under the name `mydata`.

```
#install.package("mlbench") if needed
library(mlbench)
data("PimaIndiansDiabetes2", package = "mlbench")
mydata <- PimaIndiansDiabetes2
```

part a) Use binomial distribution as a link function since diabetes is a binary outcome.

```
#your code
str(mydata)
```

part b)

```
## 'data.frame': 768 obs. of 9 variables:
## $ pregnant: num 6 1 8 1 0 5 3 10 2 8 ...
## $ glucose : num 148 85 183 89 137 116 78 115 197 125 ...
## $ pressure: num 72 66 64 66 40 74 50 NA 70 96 ...
## $ triceps : num 35 29 NA 23 35 NA 32 NA 45 NA ...
## $ insulin : num NA NA NA 94 168 NA 88 NA 543 NA ...
## $ mass : num 33.6 26.6 23.3 28.1 43.1 25.6 31 35.3 30.5 NA ...
## $ pedigree: num 0.627 0.351 0.672 0.167 2.288 ...
## $ age : num 50 31 32 21 33 30 26 29 53 54 ...
## $ diabetes: Factor w/ 2 levels "neg","pos": 2 1 2 1 2 1 2 1 2 2 ...
```

```
newdata <- na.omit(mydata)
newdata
```

```
##      pregnant glucose pressure triceps insulin mass pedigree age diabetes
## 4           1      89       66      23      94 28.1    0.167 21      neg
## 5           0     137       40      35     168 43.1    2.288 33      pos
## 7           3      78       50      32      88 31.0    0.248 26      pos
## 9           2     197       70      45     543 30.5    0.158 53      pos
## 14          1     189       60      23     846 30.1    0.398 59      pos
## 15          5     166       72      19     175 25.8    0.587 51      pos
## 17          0     118       84      47     230 45.8    0.551 31      pos
```

## 19	1	103	30	38	83	43.3	0.183	33	neg
## 20	1	115	70	30	96	34.6	0.529	32	pos
## 21	3	126	88	41	235	39.3	0.704	27	neg
## 25	11	143	94	33	146	36.6	0.254	51	pos
## 26	10	125	70	26	115	31.1	0.205	41	pos
## 28	1	97	66	15	140	23.2	0.487	22	neg
## 29	13	145	82	19	110	22.2	0.245	57	neg
## 32	3	158	76	36	245	31.6	0.851	28	pos
## 33	3	88	58	11	54	24.8	0.267	22	neg
## 36	4	103	60	33	192	24.0	0.966	33	neg
## 40	4	111	72	47	207	37.1	1.390	56	pos
## 41	3	180	64	25	70	34.0	0.271	26	neg
## 44	9	171	110	24	240	45.4	0.721	54	pos
## 51	1	103	80	11	82	19.4	0.491	22	neg
## 52	1	101	50	15	36	24.2	0.526	26	neg
## 53	5	88	66	21	23	24.4	0.342	30	neg
## 54	8	176	90	34	300	33.7	0.467	58	pos
## 55	7	150	66	42	342	34.7	0.718	42	neg
## 57	7	187	68	39	304	37.7	0.254	41	pos
## 58	0	100	88	60	110	46.8	0.962	31	neg
## 60	0	105	64	41	142	41.5	0.173	22	neg
## 64	2	141	58	34	128	25.4	0.699	24	neg
## 69	1	95	66	13	38	19.6	0.334	25	neg
## 70	4	146	85	27	100	28.9	0.189	27	neg
## 71	2	100	66	20	90	32.9	0.867	28	pos
## 72	5	139	64	35	140	28.6	0.411	26	neg
## 74	4	129	86	20	270	35.1	0.231	23	neg
## 83	7	83	78	26	71	29.3	0.767	36	neg
## 86	2	110	74	29	125	32.4	0.698	27	neg
## 88	2	100	68	25	71	38.5	0.324	26	neg
## 89	15	136	70	32	110	37.1	0.153	43	pos
## 92	4	123	80	15	176	32.0	0.443	34	neg
## 93	7	81	78	40	48	46.7	0.261	42	neg
## 95	2	142	82	18	64	24.7	0.761	21	neg
## 96	6	144	72	27	228	33.9	0.255	40	neg
## 98	1	71	48	18	76	20.4	0.323	22	neg
## 99	6	93	50	30	64	28.7	0.356	23	neg
## 100	1	122	90	51	220	49.7	0.325	31	pos
## 104	1	81	72	18	40	26.6	0.283	24	neg
## 106	1	126	56	29	152	28.7	0.801	21	neg
## 108	4	144	58	28	140	29.5	0.287	37	neg
## 109	3	83	58	31	18	34.3	0.336	25	neg
## 110	0	95	85	25	36	37.4	0.247	24	pos
## 111	3	171	72	33	135	33.3	0.199	24	pos
## 112	8	155	62	26	495	34.0	0.543	46	pos
## 113	1	89	76	34	37	31.2	0.192	23	neg
## 115	7	160	54	32	175	30.5	0.588	39	pos
## 120	4	99	76	15	51	23.2	0.223	21	neg
## 121	0	162	76	56	100	53.2	0.759	25	pos
## 123	2	107	74	30	100	33.6	0.404	23	neg
## 126	1	88	30	42	99	55.0	0.496	26	pos
## 127	3	120	70	30	135	42.9	0.452	30	neg
## 128	1	118	58	36	94	33.3	0.261	23	neg
## 129	1	117	88	24	145	34.5	0.403	40	pos

## 131	4	173	70	14	168	29.7	0.361	33	pos
## 133	3	170	64	37	225	34.5	0.356	30	pos
## 135	2	96	68	13	49	21.1	0.647	26	neg
## 136	2	125	60	20	140	33.8	0.088	31	neg
## 137	0	100	70	26	50	30.8	0.597	21	neg
## 138	0	93	60	25	92	28.7	0.532	22	neg
## 140	5	105	72	29	325	36.9	0.159	28	neg
## 143	2	108	52	26	63	32.5	0.318	22	neg
## 145	4	154	62	31	284	32.8	0.237	23	neg
## 148	2	106	64	35	119	30.5	1.400	34	neg
## 151	1	136	74	50	204	37.4	0.399	24	neg
## 153	9	156	86	28	155	34.3	1.189	42	pos
## 154	1	153	82	42	485	40.6	0.687	23	neg
## 157	2	99	52	15	94	24.6	0.637	21	neg
## 158	1	109	56	21	135	25.2	0.833	23	neg
## 159	2	88	74	19	53	29.0	0.229	22	neg
## 160	17	163	72	41	114	40.9	0.817	47	pos
## 162	7	102	74	40	105	37.2	0.204	45	neg
## 163	0	114	80	34	285	44.2	0.167	27	neg
## 166	6	104	74	18	156	29.9	0.722	41	pos
## 170	3	111	90	12	78	28.4	0.495	29	neg
## 172	6	134	70	23	130	35.4	0.542	29	pos
## 174	1	79	60	42	48	43.5	0.678	23	neg
## 175	2	75	64	24	55	29.7	0.370	33	neg
## 176	8	179	72	42	130	32.7	0.719	36	pos
## 178	0	129	110	46	130	67.1	0.319	26	pos
## 182	0	119	64	18	92	34.9	0.725	23	neg
## 187	8	181	68	36	495	30.1	0.615	60	pos
## 188	1	128	98	41	58	32.0	1.321	33	pos
## 189	8	109	76	39	114	27.9	0.640	31	pos
## 190	5	139	80	35	160	31.6	0.361	25	pos
## 192	9	123	70	44	94	33.1	0.374	40	neg
## 196	5	158	84	41	210	39.4	0.395	29	pos
## 198	3	107	62	13	48	22.9	0.678	23	pos
## 199	4	109	64	44	99	34.8	0.905	26	pos
## 200	4	148	60	27	318	30.9	0.150	29	pos
## 204	2	99	70	16	44	20.4	0.235	27	neg
## 205	6	103	72	32	190	37.7	0.324	55	neg
## 207	8	196	76	29	280	37.5	0.605	57	pos
## 209	1	96	64	27	87	33.2	0.289	21	neg
## 214	0	140	65	26	130	42.6	0.431	24	pos
## 215	9	112	82	32	175	34.2	0.260	36	pos
## 216	12	151	70	40	271	41.8	0.742	38	pos
## 217	5	109	62	41	129	35.8	0.514	25	pos
## 218	6	125	68	30	120	30.0	0.464	32	neg
## 221	0	177	60	29	478	34.6	1.072	21	pos
## 224	7	142	60	33	190	28.8	0.687	61	neg
## 225	1	100	66	15	56	23.6	0.666	26	neg
## 226	1	87	78	27	32	34.6	0.101	22	neg
## 229	4	197	70	39	744	36.7	2.329	31	neg
## 230	0	117	80	31	53	45.2	0.089	24	neg
## 232	6	134	80	37	370	46.2	0.238	46	pos
## 233	1	79	80	25	37	25.4	0.583	22	neg
## 235	3	74	68	28	45	29.7	0.293	23	neg

## 237	7	181	84	21	192	35.9	0.586	51	pos
## 242	4	91	70	32	88	33.1	0.446	22	neg
## 244	6	119	50	22	176	27.1	1.318	33	pos
## 245	2	146	76	35	194	38.2	0.329	29	neg
## 248	0	165	90	33	680	52.3	0.427	23	neg
## 249	9	124	70	33	402	35.4	0.282	34	neg
## 253	2	90	80	14	55	24.4	0.249	24	neg
## 255	12	92	62	7	258	27.6	0.926	44	pos
## 259	1	193	50	16	375	25.9	0.655	24	neg
## 260	11	155	76	28	150	33.3	1.353	51	pos
## 261	3	191	68	15	130	30.9	0.299	34	neg
## 266	5	96	74	18	67	33.6	0.997	43	neg
## 272	2	108	62	32	56	25.2	0.128	21	neg
## 274	1	71	78	50	45	33.2	0.422	21	neg
## 276	2	100	70	52	57	40.5	0.677	25	neg
## 278	0	104	64	23	116	27.8	0.454	23	neg
## 280	2	108	62	10	278	25.3	0.881	22	neg
## 282	10	129	76	28	122	35.9	0.280	39	neg
## 283	7	133	88	15	155	32.4	0.262	37	neg
## 286	7	136	74	26	135	26.0	0.647	51	neg
## 287	5	155	84	44	545	38.7	0.619	34	neg
## 288	1	119	86	39	220	45.6	0.808	29	pos
## 289	4	96	56	17	49	20.8	0.340	26	neg
## 290	5	108	72	43	75	36.1	0.263	33	neg
## 291	0	78	88	29	40	36.9	0.434	21	neg
## 292	0	107	62	30	74	36.6	0.757	25	pos
## 293	2	128	78	37	182	43.3	1.224	31	pos
## 294	1	128	48	45	194	40.5	0.613	24	pos
## 296	6	151	62	31	120	35.5	0.692	28	neg
## 297	2	146	70	38	360	28.0	0.337	29	pos
## 298	0	126	84	29	215	30.7	0.520	24	neg
## 299	14	100	78	25	184	36.6	0.412	46	pos
## 302	2	144	58	33	135	31.6	0.422	25	pos
## 303	5	77	82	41	42	35.8	0.156	35	neg
## 306	2	120	76	37	105	39.7	0.215	29	neg
## 307	10	161	68	23	132	25.5	0.326	47	pos
## 308	0	137	68	14	148	24.8	0.143	21	neg
## 309	0	128	68	19	180	30.5	1.391	25	pos
## 310	2	124	68	28	205	32.9	0.875	30	pos
## 312	0	106	70	37	148	39.4	0.605	22	neg
## 313	2	155	74	17	96	26.6	0.433	27	pos
## 314	3	113	50	10	85	29.5	0.626	25	neg
## 316	2	112	68	22	94	34.1	0.315	26	neg
## 317	3	99	80	11	64	19.3	0.284	30	neg
## 319	3	115	66	39	140	38.1	0.150	28	neg
## 321	4	129	60	12	231	27.5	0.527	31	neg
## 324	13	152	90	33	29	26.8	0.731	43	pos
## 326	1	157	72	21	168	25.6	0.123	24	neg
## 327	1	122	64	32	156	35.1	0.692	30	pos
## 329	2	102	86	36	120	45.5	0.127	23	pos
## 330	6	105	70	32	68	30.8	0.122	37	neg
## 332	2	87	58	16	52	32.7	0.166	25	neg
## 335	1	95	60	18	58	23.9	0.260	22	neg
## 336	0	165	76	43	255	47.9	0.259	26	neg

## 339	9	152	78	34	171	34.2	0.893	33	pos
## 341	1	130	70	13	105	25.9	0.472	22	neg
## 342	1	95	74	21	73	25.9	0.673	36	neg
## 346	8	126	88	36	108	38.5	0.349	49	neg
## 347	1	139	46	19	83	28.7	0.654	22	neg
## 349	3	99	62	19	74	21.8	0.279	26	neg
## 354	1	90	62	12	43	27.2	0.580	24	neg
## 357	1	125	50	40	167	33.3	0.962	28	pos
## 359	12	88	74	40	54	35.3	0.378	48	neg
## 360	1	196	76	36	249	36.5	0.875	29	pos
## 361	5	189	64	33	325	31.2	0.583	29	pos
## 365	4	147	74	25	293	34.9	0.385	30	neg
## 366	5	99	54	28	83	34.0	0.499	30	neg
## 369	3	81	86	16	66	27.5	0.306	22	neg
## 370	1	133	102	28	140	32.8	0.234	45	pos
## 371	3	173	82	48	465	38.4	2.137	25	pos
## 373	0	84	64	22	66	35.8	0.545	21	neg
## 374	2	105	58	40	94	34.9	0.225	25	neg
## 375	2	122	52	43	158	36.2	0.816	28	neg
## 376	12	140	82	43	325	39.2	0.528	58	pos
## 377	0	98	82	15	84	25.2	0.299	22	neg
## 378	1	87	60	37	75	37.2	0.509	22	neg
## 380	0	93	100	39	72	43.4	1.021	35	neg
## 381	1	107	72	30	82	30.8	0.821	24	neg
## 383	1	109	60	8	182	25.4	0.947	21	neg
## 384	1	90	62	18	59	25.1	1.268	25	neg
## 385	1	125	70	24	110	24.3	0.221	25	neg
## 386	1	119	54	13	50	22.3	0.205	24	neg
## 389	5	144	82	26	285	32.0	0.452	58	pos
## 390	3	100	68	23	81	31.6	0.949	28	neg
## 391	1	100	66	29	196	32.0	0.444	42	neg
## 393	1	131	64	14	415	23.7	0.389	21	neg
## 394	4	116	72	12	87	22.1	0.463	37	neg
## 396	2	127	58	24	275	27.7	1.600	25	neg
## 397	3	96	56	34	115	24.7	0.944	39	neg
## 403	5	136	84	41	88	35.0	0.286	35	pos
## 406	2	123	48	32	165	42.1	0.520	26	neg
## 410	1	172	68	49	579	42.4	0.702	28	pos
## 412	1	112	72	30	176	34.4	0.528	25	neg
## 413	1	143	84	23	310	42.4	1.076	22	neg
## 414	1	143	74	22	61	26.2	0.256	21	neg
## 415	0	138	60	35	167	34.6	0.534	21	pos
## 416	3	173	84	33	474	35.7	0.258	22	pos
## 420	3	129	64	29	115	26.4	0.219	28	pos
## 421	1	119	88	41	170	45.3	0.507	26	neg
## 422	2	94	68	18	76	26.0	0.561	21	neg
## 423	0	102	64	46	78	40.6	0.496	21	neg
## 425	8	151	78	32	210	42.9	0.516	36	pos
## 426	4	184	78	39	277	37.0	0.264	31	pos
## 428	1	181	64	30	180	34.1	0.328	38	pos
## 429	0	135	94	46	145	40.6	0.284	26	neg
## 430	1	95	82	25	180	35.0	0.233	43	pos
## 432	3	89	74	16	85	30.4	0.551	38	neg
## 433	1	80	74	11	60	30.0	0.527	22	neg

## 442	2	83	66	23	50	32.2	0.497	22	neg
## 443	4	117	64	27	120	33.2	0.230	24	neg
## 446	0	180	78	63	14	59.4	2.420	25	pos
## 447	1	100	72	12	70	25.3	0.658	28	neg
## 448	0	95	80	45	92	36.5	0.330	26	neg
## 449	0	104	64	37	64	33.6	0.510	22	pos
## 450	0	120	74	18	63	30.5	0.285	26	neg
## 451	1	82	64	13	95	21.2	0.415	23	neg
## 453	0	91	68	32	210	39.9	0.381	25	neg
## 455	2	100	54	28	105	37.8	0.498	24	neg
## 458	5	86	68	28	71	30.2	0.364	24	neg
## 459	10	148	84	48	237	37.6	1.001	51	pos
## 460	9	134	74	33	60	25.9	0.460	81	neg
## 461	9	120	72	22	56	20.8	0.733	48	neg
## 463	8	74	70	40	49	35.3	0.705	39	neg
## 466	0	124	56	13	105	21.8	0.452	21	neg
## 467	0	74	52	10	36	27.8	0.269	22	neg
## 468	0	97	64	36	100	36.8	0.600	25	neg
## 470	6	154	78	41	140	46.1	0.571	27	neg
## 477	2	105	80	45	191	33.7	0.711	29	pos
## 478	7	114	76	17	110	23.8	0.466	31	neg
## 479	8	126	74	38	75	25.9	0.162	39	neg
## 481	3	158	70	30	328	35.5	0.344	35	pos
## 483	4	85	58	22	49	27.8	0.306	28	neg
## 484	0	84	82	31	125	38.2	0.233	23	neg
## 486	0	135	68	42	250	42.3	0.365	24	pos
## 487	1	139	62	41	480	40.7	0.536	21	neg
## 488	0	173	78	32	265	46.5	1.159	58	neg
## 491	2	83	65	28	66	36.8	0.629	24	neg
## 494	4	125	70	18	122	28.9	1.144	45	pos
## 498	2	81	72	15	76	30.1	0.547	25	neg
## 499	7	195	70	33	145	25.1	0.163	55	pos
## 500	6	154	74	32	193	29.3	0.839	39	neg
## 501	2	117	90	19	71	25.2	0.313	21	neg
## 504	7	94	64	25	79	33.3	0.738	41	neg
## 507	0	180	90	26	90	36.5	0.314	35	pos
## 508	1	130	60	23	170	28.6	0.692	21	neg
## 509	2	84	50	23	76	30.4	0.968	21	neg
## 512	0	139	62	17	210	22.1	0.207	21	neg
## 515	3	99	54	19	86	25.6	0.154	24	neg
## 516	3	163	70	18	105	31.6	0.268	28	pos
## 517	9	145	88	34	165	30.3	0.771	53	pos
## 520	6	129	90	7	326	19.6	0.582	60	neg
## 521	2	68	70	32	66	25.0	0.187	25	neg
## 522	3	124	80	33	130	33.2	0.305	26	neg
## 527	1	97	64	19	82	18.2	0.299	21	neg
## 528	3	116	74	15	105	26.3	0.107	24	neg
## 529	0	117	66	31	188	30.8	0.493	22	neg
## 531	2	122	60	18	106	29.8	0.717	22	neg
## 533	1	86	66	52	65	41.3	0.917	29	neg
## 535	1	77	56	30	56	33.3	1.251	24	neg
## 539	0	127	80	37	210	36.3	0.804	23	neg
## 540	3	129	92	49	155	36.4	0.968	32	pos
## 541	8	100	74	40	215	39.4	0.661	43	pos

## 542	3	128	72	25	190	32.4	0.549	27	pos
## 544	4	84	90	23	56	39.5	0.159	25	neg
## 545	1	88	78	29	76	32.0	0.365	29	neg
## 546	8	186	90	35	225	34.5	0.423	37	pos
## 547	5	187	76	27	207	43.6	1.034	53	pos
## 548	4	131	68	21	166	33.1	0.160	28	neg
## 549	1	164	82	43	67	32.8	0.341	50	neg
## 552	3	84	68	30	106	31.9	0.591	25	neg
## 554	1	88	62	24	44	29.9	0.422	23	neg
## 555	1	84	64	23	115	36.9	0.471	28	neg
## 556	7	124	70	33	215	25.5	0.161	37	neg
## 562	0	198	66	32	274	41.3	0.502	28	pos
## 563	1	87	68	34	77	37.6	0.401	24	neg
## 564	6	99	60	19	54	26.9	0.497	32	neg
## 566	2	95	54	14	88	26.1	0.748	22	neg
## 567	1	99	72	30	18	38.6	0.412	21	neg
## 568	6	92	62	32	126	32.0	0.085	46	neg
## 569	4	154	72	29	126	31.3	0.338	37	neg
## 570	0	121	66	30	165	34.3	0.203	33	pos
## 573	3	111	58	31	44	29.5	0.430	22	neg
## 574	2	98	60	17	120	34.7	0.198	22	neg
## 575	1	143	86	30	330	30.1	0.892	23	neg
## 576	1	119	44	47	63	35.5	0.280	25	neg
## 577	6	108	44	20	130	24.0	0.813	35	neg
## 585	8	124	76	24	600	28.7	0.687	52	pos
## 589	3	176	86	27	156	33.3	1.154	52	pos
## 592	2	112	78	50	140	39.4	0.175	24	neg
## 594	2	82	52	22	115	28.5	1.699	25	neg
## 595	6	123	72	45	230	33.6	0.733	34	neg
## 596	0	188	82	14	185	32.0	0.682	22	pos
## 598	1	89	24	19	25	27.8	0.559	21	neg
## 600	1	109	38	18	120	23.1	0.407	26	neg
## 604	7	150	78	29	126	35.2	0.692	54	pos
## 607	1	181	78	42	293	40.0	1.258	22	pos
## 608	1	92	62	25	41	19.5	0.482	25	neg
## 609	0	152	82	39	272	41.5	0.270	27	neg
## 610	1	111	62	13	182	24.0	0.138	23	neg
## 611	3	106	54	21	158	30.9	0.292	24	neg
## 612	3	174	58	22	194	32.9	0.593	36	pos
## 613	7	168	88	42	321	38.2	0.787	40	pos
## 615	11	138	74	26	144	36.1	0.557	50	pos
## 618	2	68	62	13	15	20.1	0.257	23	neg
## 621	2	112	86	42	160	38.4	0.246	28	neg
## 624	0	94	70	27	115	43.5	0.347	21	neg
## 626	4	90	88	47	54	37.7	0.362	29	neg
## 632	0	102	78	40	90	34.5	0.238	24	neg
## 634	1	128	82	17	183	27.5	0.115	22	neg
## 638	2	94	76	18	66	31.6	0.649	23	neg
## 639	7	97	76	32	91	40.9	0.871	32	pos
## 640	1	100	74	12	46	19.5	0.149	28	neg
## 641	0	102	86	17	105	29.3	0.695	27	neg
## 645	3	103	72	30	152	27.6	0.730	27	neg
## 646	2	157	74	35	440	39.4	0.134	30	neg
## 647	1	167	74	17	144	23.4	0.447	33	pos

## 648	0	179	50	36	159	37.8	0.455	22	pos
## 649	11	136	84	35	130	28.3	0.260	42	pos
## 651	1	91	54	25	100	25.2	0.234	23	neg
## 652	1	117	60	23	106	33.8	0.466	27	neg
## 653	5	123	74	40	77	34.1	0.269	28	neg
## 655	1	106	70	28	135	34.2	0.142	22	neg
## 656	2	155	52	27	540	38.7	0.240	25	pos
## 657	2	101	58	35	90	21.8	0.155	22	neg
## 658	1	120	80	48	200	38.9	1.162	41	neg
## 660	3	80	82	31	70	34.2	1.292	27	pos
## 663	8	167	106	46	231	37.6	0.165	43	pos
## 664	9	145	80	46	130	37.9	0.637	40	pos
## 666	1	112	80	45	132	34.8	0.217	24	neg
## 669	6	98	58	33	190	34.0	0.430	43	neg
## 670	9	154	78	30	100	30.9	0.164	45	neg
## 671	6	165	68	26	168	33.6	0.631	49	neg
## 673	10	68	106	23	49	35.5	0.285	47	neg
## 674	3	123	100	35	240	57.3	0.880	22	neg
## 680	2	101	58	17	265	24.2	0.614	23	neg
## 681	2	56	56	28	45	24.2	0.332	22	neg
## 683	0	95	64	39	105	44.6	0.366	22	neg
## 686	2	129	74	26	205	33.2	0.591	25	neg
## 689	1	140	74	26	180	24.1	0.828	23	neg
## 690	1	144	82	46	180	46.1	0.335	46	pos
## 693	2	121	70	32	95	39.1	0.886	23	neg
## 694	7	129	68	49	125	38.5	0.439	43	pos
## 696	7	142	90	24	480	30.4	0.128	43	pos
## 697	3	169	74	19	125	29.9	0.268	31	pos
## 699	4	127	88	11	155	34.5	0.598	28	neg
## 701	2	122	76	27	200	35.9	0.483	26	neg
## 705	4	110	76	20	100	28.4	0.118	27	neg
## 708	2	127	46	21	335	34.4	0.176	22	neg
## 710	2	93	64	32	160	38.0	0.674	23	pos
## 711	3	158	64	13	387	31.2	0.295	24	neg
## 712	5	126	78	27	22	29.6	0.439	40	neg
## 714	0	134	58	20	291	26.4	0.352	21	neg
## 716	7	187	50	33	392	33.9	0.826	34	pos
## 717	3	173	78	39	185	33.8	0.970	31	pos
## 719	1	108	60	46	178	35.5	0.415	24	neg
## 722	1	114	66	36	200	38.1	0.289	21	neg
## 723	1	149	68	29	127	29.3	0.349	42	pos
## 724	5	117	86	30	105	39.1	0.251	42	neg
## 727	1	116	78	29	180	36.1	0.496	25	neg
## 731	3	130	78	23	79	28.4	0.323	34	pos
## 733	2	174	88	37	120	44.5	0.646	24	pos
## 734	2	106	56	27	165	29.0	0.426	22	neg
## 737	0	126	86	27	120	27.4	0.515	21	neg
## 739	2	99	60	17	160	36.6	0.453	21	neg
## 741	11	120	80	37	150	42.3	0.785	48	pos
## 742	3	102	44	20	94	30.8	0.400	26	neg
## 743	1	109	58	18	116	28.5	0.219	22	neg
## 745	13	153	88	37	140	40.6	1.174	39	neg
## 746	12	100	84	33	105	30.0	0.488	46	neg
## 748	1	81	74	41	57	46.3	1.096	32	neg

```
## 749      3      187      70      22      200 36.4      0.408 36      pos
## 752      1      121      78      39       74 39.0      0.261 28      neg
## 754      0      181      88      44      510 43.3      0.222 26      pos
## 756      1      128      88      39      110 36.5      1.057 37      pos
## 761      2       88      58      26       16 28.4      0.766 22      neg
## 764     10      101      76      48      180 32.9      0.171 63      neg
## 766      5      121      72      23      112 26.2      0.245 30      neg
```

your codes here

```
glm_diabetes <- glm(diabetes ~ pregnant + glucose + pressure + triceps + insulin + mass + pedigree + age,
  family = binomial, data = newdata)
glm_diabetes
```

part c)

```
##
## Call:  glm(formula = diabetes ~ pregnant + glucose + pressure + triceps +
##        insulin + mass + pedigree + age, family = binomial, data = newdata)
##
## Coefficients:
## (Intercept)      pregnant      glucose      pressure      triceps      insulin
## -1.004e+01    8.216e-02    3.827e-02   -1.420e-03    1.122e-02   -8.253e-04
##      mass      pedigree      age
##  7.054e-02    1.141e+00    3.395e-02
##
## Degrees of Freedom: 391 Total (i.e. Null);  383 Residual
## Null Deviance:      498.1
## Residual Deviance: 344    AIC: 362
```

part d) The coefficient for glucose is 0.03827, this means that for every one unit increase in the glucose variable, holding all other variables constant, the diabetes increases by 0.03827.

```
summary(glm_diabetes)
```

part e)

```
##
## Call:
## glm(formula = diabetes ~ pregnant + glucose + pressure + triceps +
##      insulin + mass + pedigree + age, family = binomial, data = newdata)
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept) -1.004e+01  1.218e+00  -8.246  < 2e-16 ***
## pregnant      8.216e-02  5.543e-02   1.482  0.13825
## glucose      3.827e-02  5.768e-03   6.635 3.24e-11 ***
## pressure    -1.420e-03  1.183e-02  -0.120  0.90446
## triceps      1.122e-02  1.708e-02   0.657  0.51128
## insulin     -8.253e-04  1.306e-03  -0.632  0.52757
## mass         7.054e-02  2.734e-02   2.580  0.00989 **
## pedigree     1.141e+00  4.274e-01   2.669  0.00760 **
## age          3.395e-02  1.838e-02   1.847  0.06474 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
##
## (Dispersion parameter for binomial family taken to be 1)
##
## Null deviance: 498.10 on 391 degrees of freedom
## Residual deviance: 344.02 on 383 degrees of freedom
## AIC: 362.02
##
## Number of Fisher Scoring iterations: 5
```

Add your explanation here: Variables of glucose, mass, pedigree are significant since their p-value is very small.

```
#your code
step(glm_diabetes, direction = "backward")
```

part f)

```
## Start: AIC=362.02
## diabetes ~ pregnant + glucose + pressure + triceps + insulin +
## mass + pedigree + age
##
## Df Deviance AIC
## - pressure 1 344.04 360.04
## - insulin 1 344.42 360.42
## - triceps 1 344.45 360.45
## <none> 344.02 362.02
## - pregnant 1 346.24 362.24
## - age 1 347.55 363.55
## - mass 1 350.89 366.89
## - pedigree 1 351.58 367.58
## - glucose 1 396.95 412.95
##
## Step: AIC=360.04
## diabetes ~ pregnant + glucose + triceps + insulin + mass + pedigree +
## age
##
## Df Deviance AIC
## - insulin 1 344.42 358.42
## - triceps 1 344.46 358.46
## <none> 344.04 360.04
## - pregnant 1 346.24 360.24
## - age 1 347.60 361.60
## - mass 1 351.28 365.28
## - pedigree 1 351.67 365.67
## - glucose 1 397.31 411.31
##
## Step: AIC=358.42
## diabetes ~ pregnant + glucose + triceps + mass + pedigree + age
##
## Df Deviance AIC
## - triceps 1 344.89 356.89
## <none> 344.42 358.42
## - pregnant 1 346.74 358.74
## - age 1 347.87 359.87
```

```
## - mass      1    351.32 363.32
## - pedigree  1    351.90 363.90
## - glucose   1    411.11 423.11
##
## Step: AIC=356.89
## diabetes ~ pregnant + glucose + mass + pedigree + age
##
##           Df Deviance    AIC
## <none>          344.89 356.89
## - pregnant  1    347.23 357.23
## - age       1    348.72 358.72
## - pedigree  1    352.72 362.72
## - mass      1    360.44 370.44
## - glucose   1    411.85 421.85
##
## Call: glm(formula = diabetes ~ pregnant + glucose + mass + pedigree +
##           age, family = binomial, data = newdata)
##
## Coefficients:
## (Intercept)    pregnant    glucose      mass    pedigree      age
##      -9.99208      0.08395      0.03646      0.07814      1.15091      0.03436
##
## Degrees of Freedom: 391 Total (i.e. Null); 386 Residual
## Null Deviance:      498.1
## Residual Deviance: 344.9    AIC: 356.9
```

Add your explanation here: The new model is diabetes ~ pregnant + glucose + mass + pedigree + age

Question 4:

Run this code before continue. note that the data is saved under the name `mydata`.

```
attach(mtcars)
mtcars
```

```
##           mpg  cyl  disp  hp  drat    wt  qsec vs  am  gear  carb
## Mazda RX4      21.0   6  160.0 110  3.90  2.620 16.46  0   1    4     4
## Mazda RX4 Wag  21.0   6  160.0 110  3.90  2.875 17.02  0   1    4     4
## Datsun 710     22.8   4  108.0  93  3.85  2.320 18.61  1   1    4     1
## Hornet 4 Drive  21.4   6  258.0 110  3.08  3.215 19.44  1   0    3     1
## Hornet Sportabout 18.7   8  360.0 175  3.15  3.440 17.02  0   0    3     2
## Valiant        18.1   6  225.0 105  2.76  3.460 20.22  1   0    3     1
## Duster 360     14.3   8  360.0 245  3.21  3.570 15.84  0   0    3     4
## Merc 240D      24.4   4  146.7  62  3.69  3.190 20.00  1   0    4     2
## Merc 230       22.8   4  140.8  95  3.92  3.150 22.90  1   0    4     2
## Merc 280       19.2   6  167.6 123  3.92  3.440 18.30  1   0    4     4
## Merc 280C      17.8   6  167.6 123  3.92  3.440 18.90  1   0    4     4
## Merc 450SE     16.4   8  275.8 180  3.07  4.070 17.40  0   0    3     3
## Merc 450SL     17.3   8  275.8 180  3.07  3.730 17.60  0   0    3     3
## Merc 450SLC    15.2   8  275.8 180  3.07  3.780 18.00  0   0    3     3
## Cadillac Fleetwood 10.4   8  472.0 205  2.93  5.250 17.98  0   0    3     4
## Lincoln Continental 10.4   8  460.0 215  3.00  5.424 17.82  0   0    3     4
## Chrysler Imperial 14.7   8  440.0 230  3.23  5.345 17.42  0   0    3     4
## Fiat 128       32.4   4   78.7  66  4.08  2.200 19.47  1   1    4     1
## Honda Civic    30.4   4   75.7  52  4.93  1.615 18.52  1   1    4     2
```

```
## Toyota Corolla      33.9   4  71.1   65 4.22 1.835 19.90   1   1    4    1
## Toyota Corona      21.5   4 120.1   97 3.70 2.465 20.01   1   0    3    1
## Dodge Challenger    15.5   8 318.0  150 2.76 3.520 16.87   0   0    3    2
## AMC Javelin         15.2   8 304.0  150 3.15 3.435 17.30   0   0    3    2
## Camaro Z28          13.3   8 350.0  245 3.73 3.840 15.41   0   0    3    4
## Pontiac Firebird    19.2   8 400.0  175 3.08 3.845 17.05   0   0    3    2
## Fiat X1-9           27.3   4   79.0   66 4.08 1.935 18.90   1   1    4    1
## Porsche 914-2       26.0   4  120.3   91 4.43 2.140 16.70   0   1    5    2
## Lotus Europa        30.4   4   95.1  113 3.77 1.513 16.90   1   1    5    2
## Ford Pantera L      15.8   8 351.0  264 4.22 3.170 14.50   0   1    5    4
## Ferrari Dino        19.7   6 145.0  175 3.62 2.770 15.50   0   1    5    6
## Maserati Bora       15.0   8 301.0  335 3.54 3.570 14.60   0   1    5    8
## Volvo 142E         21.4   4 121.0  109 4.11 2.780 18.60   1   1    4    2
```

```
# you can select a column of a dataset by $ operator, for example
# wt_col <- mtcars$wt
```

part a the matrix size is 32x3

```
#your code
wt_col <- mtcars$wt
hp_col <- mtcars$hp
X <- cbind(1, wt_col, hp_col)
y <- mtcars$mpg
p <- 3
X.QR <- qr(X)
R <- qr.R(X.QR, complete = TRUE)
Q <- qr.Q(X.QR, complete = TRUE)
U <- qr.R(X.QR)
U_inverse <- solve(U)
Q1 <- Q[, 1:p]
Q1y <- t(Q1) %*% y
betahat <- solve(U, Q1y)
betahat
```

part b

```
##           [,1]
##      37.22727012
## wt_col -3.87783074
## hp_col -0.03177295
```

```
Q2 <- Q[, -(1:3)]
Q2y <- t(Q2) %*% y
SSE <- t(Q2y) %*% Q2y
SSE
```

part c

```
##           [,1]
## [1,] 195.0478
```

```
# Part d error variance
MSE <- SSE/(32-3)
```

MSE

```
##           [,1]  
## [1,] 6.725785
```