

Age	①TP/Cr	①PGE-MUM/Cr	①FABP/Cr	②TP/Cr	②PGE-MUM/Cr
17	0	10.2	2.07	0	12.5
17	0.04	9.71	3.16	0	7.98
17	0.01	13.2	5.54	0	15.8
17	0	11.9	0	0	11.4
17	0	39.5	0.77	0	49.3
17	0.22	34.8	3.01	0	31.4
17	0	25.1	4.16	0.02	18.4
17	0.01	9.23	1.5	0.01	9.37
17	0	18.2	0	0	13.1
17	0.02	25.3	3.51	0	22.9
17	0.01	11.5	2.56	0	11.4
17	0	10.5	2.51	0	12.9
17	0.02	31.7	0.87	0	17.1
17	0.02	21.9	1.32	0.02	17.9
17	0	13.1	0.3	0.02	16.8
17	0.03	11.3	4.66	0	11.2
17	0	17.3	1.7	0.05	10.2
17	0.01	20.3	1.31	0	18.1
17	0.02	14.6	1.16	0	12.3
17	0	10.1	2.95	0	14.9
17	0.02	18	3.25	0.02	8.91
17	0	14.9	1.65	0	19
18	0	10.9	0.94	0.01	9.03
18	0.02	111	1.11	0.01	10.8
18	0.01	10.1	2.15	0.03	96.7
18	0.02	16.4	0.7	0.01	11.6
18	0	9.95	0.84	0.02	20.7
18	0	16.6	1.56	0	6.88
18	0	20.1	2	0.02	18.5
18	0	19.7	2.75	0.01	23.3
18	0	16	0.78	0.02	24.9
18	0	16.1	2.08	0.03	15.3
18	0.05	39.9	1.73	0.01	16.7
18	0.03	16.9	2.57	0.04	37.8
18	0	13.4	1.45	0.02	16
18	0	23.4	2.15	0.08	11.9
18	0	14.1	2.28	0	22.5
18	0.01	15	4.77	0	16.9

18	0.02	40.9	1.79	0.01	10.1
18	0	41	0.71	0.03	28.9
18	0.02	29.2	1.89	0.01	52.4
18	0	20.5	0	0.01	29.6
18	0	24.2	0	0	14.3
18	0.04	11.5	1.98	0	17.6
16	0	48.4	1.86	0	16.5
16	0	11.9	0.96	0.03	9.54
16	0	9.33	1.65	0	60.4
16	0.01	15.9	2.19	0.03	13.7
16	0	12.6	0.62	0.01	11.5
16	0	27.6	0	0.03	17.5
16	0.02	10.6	2.37	0.02	13.6
16	0	12.2	1.47	0.1	26.3
16	0	12.5	1.15	0	10.7
16	0.03	9.64	3.05	0.04	14.6
16	0	22.1	0.45	0.02	13.7
16	0	22.1	4.99	0.04	12
16	0	19.7	0.54	0.02	19.2
16				0.03	17
16				0	19
Mean	17.11864	0.01246	20.41684211	1.8507018	0.01492
SD	0.789691	0.03066	15.40882391	1.3009528	0.0196
					19.50016949
					14.57853423

②FABP/Cr	③TP/Cr	③PGE-MUM/Cr	③FABP/Cr	④TP/Cr	④PGE-MUM/Cr	④FABP/Cr
0.54	0.04	15.5	2.15	0.01	17	1.39
0.3	0.05	9.93	3.92	0.01	10.3	1.48
1.85	0.02	18.1	2.51	0.02	23.3	2.1
0.44	0.38	30.6	13.85	0.03	21.6	5.2
1.06	0.13	95.4	1.02	0.04	85.6	0.96
1.53	0.01	38	3.32	0	42.1	2.56
3.08	0.04	30.2	2.67	0.03	35.2	2.36
0	0.03	11.9	1.79	0.04	12.4	1.27
0	0.04	18.2	0.9	0.01	17.7	0.83
0.99	0.02	29.2	1.95	0	28.2	0
3.08	0.01	18.9	2.61	0.02	17.1	2.1
1.53	0.01	19.1	2.6	0	19.2	0.89
2.83	0.03	16.6	2.83	0.02	20.6	1.5
2.4	0.05	29	2.7	0.02	25.4	2.84
0.98	0.03	17.8	0.97	0.01	19.9	1.11
1.37	0.01	15.9	1.18	0.03	20.9	0.87
4.14	0.09	15.1	5.07	0.01	11.9	2.03
1.01	0.17	15.5	1.38	0.06	13.6	1.03
1.66	0.28	23.6	2.8	0.05	14.5	2.08
1.87	0.12	23.8	5.62	0.05	15.7	1.3
3.28	0.02	12	4.61	0.02	17.2	4.04
1.94	0.05	32.1	2.57	0.02	24.6	1.68
1.38	0.05	15.2	1.69	0.01	11.4	0.6
1.08	0.06	27.3	1.93	0.02	14.9	1.24
0.87	0.16	82.7	4.19	0	97.5	1.24
1.22	0.02	12.5	1.22	0	13.3	0.89
0.8	0.12	28.9	1.2	0.01	23	0.99
1.24	0.02	8.94	0.92	0	23.5	0.94
1.32	0.03	17.2	1.31	0.33	19.6	0.75
1.37	0.2	33.5	10	0.01	23.1	3.14
1.74	0.07	25.1	3.52	0.02	22.1	2
0.85	0.1	26.5	2.69	0.03	28.7	8.36
1.32	0.47	28.9	3.41	0.01	21.2	1.81
1.54	0.17	40.1	1.78	0.04	38.5	1.84
2.65	0.15	23.4	6.41	0.04	28.5	8.58
2.22	0.05	19.4	5.93	0.01	14.2	2.72
0	0.06	36.7	2.61	0.02	26	2.63
1.24	0.13	23.8	2.39	0.01	20.8	2.05

1.09	0.24	29.7	3.96	0.01	15.9	4.37
1.21	0.07	29.8	2.32	0.03	42.1	1.64
1.91	0.02	44.9	4.72	0	38.4	2.03
1.77	0.05	28.8	0.88	0.03	27.1	0.55
2.68	0	12.9	0.66	0	15.3	1.16
0.49	0.02	29.4	1.39	0.04	24.7	0.77
0.33	0.08	21.8	6.43	0.03	22.9	1.24
2.77	0.18	16	3.51	0.13	19.7	2.35
4.63	0.04	65.1	10.2	0	62.3	1.12
1.61	0.38	25	2.88	0.13	23.8	4.42
2.81	0.01	19.1	3.2	0.02	15.8	3.23
3.67	0.22	18	2.03	0.03	22.9	3.23
0.79	0.36	17.2	2.92	0.15	19.7	1.5
3.62	0.59	58.7	11.96	0.11	37.6	5.75
1.25	0.04	11.1	1.08	0.02	8.67	0.89
2.52	0.04	18.9	3.93	0.04	14.3	3.01
1.89	0.1	22.5	1.45	0.09	15.9	6.2
5.71	0.03	17.5	2.4	0.04	11.7	2.66
0.57	0.11	24.9	16.56	0	18.8	0
3.76	0.19	25.3	17.11	0.08	21.6	22.08
1.21	0.06	23.9	2.16	0.03	20.2	1.76
1.7459322	0.10712	26.22152542	3.7622034	0.035085	24.40118644	2.5315254
1.186775	0.12257	15.92345546	3.6281576	0.051306	15.77647355	3.1294688

⑤TP/Cr	⑤PGE-MUM/Cr	⑤FABP/Cr	①PGE-MUM/Cr	②PGE-MUM/Cr
0.02	17	0.94	10.2	12.5
0.01	8.62	1.19	9.71	7.98
0	13.9	1.2	13.2	15.8
0	13	0.34	11.9	11.4
0	54.2	0.77	39.5	49.3
0.03	32.1	2.75	34.8	31.4
0	12.8	0.92	25.1	18.4
0	10.7	0.28	9.23	9.37
0	14	0	18.2	13.1
0	30.8	0.34	25.3	22.9
0.02	12.6	4.08	11.5	11.4
0.51	20.3	0.75	10.5	12.9
0.02	16.6	1.27	31.7	17.1
0.02	16	2.07	21.9	17.9
0	23	0	13.1	16.8
0	12.4	0.75	11.3	11.2
0	10.8	0.39	17.3	10.2
0	9.8	0.42	20.3	18.1
0.03	14.7	1.01	14.6	12.3
0.02	13.3	1.93	10.1	14.9
0.02	9	1.47	18	8.91
0.02	16.7	1.24	14.9	19
0.02	22	1.21	10.9	9.03
0.02	11.1	0.88	111	10.8
0.05	102	0.58	10.1	96.7
0.02	11.2	1.13	16.4	11.6
0.03	12.4	0.47	9.95	20.7
0.03	11.6	1.19	16.6	6.88
0.02	18	0.46	20.1	18.5
0.01	16.4	0.24	19.7	23.3
0.04	17.4	0.86	16	24.9
0.03	12.1	0.72	16.1	15.3
0	18.1	0.34	39.9	16.7
0	29.3	0	16.9	37.8
0.04	17.5	3.3	13.4	16
0.02	12.1	0.86	23.4	11.9
0.01	24.7	0.91	14.1	22.5
0.02	16.9	1.48	15	16.9

0.02	10.6	1.59	40.9	10.1
0.01	17.8	0.7	41	28.9
0	34.3	0.99	29.2	52.4
0.01	17.1	1.14	20.5	29.6
0.02	11.9	3	24.2	14.3
0	24.2	0.58	11.5	17.6
0	15.4	0	48.4	16.5
0.05	7.98	2.03	11.9	9.54
0.01	53.7	2.69	9.33	60.4
0.04	10.5	0.72	15.9	13.7
0	12.3	0.39	12.6	11.5
0.03	18.2	3.19	27.6	17.5
0.03	11.7	0.82	10.6	13.6
0.02	22.2	3.28	12.2	26.3
0.01	8.3	0.89	12.5	10.7
0.02	10.9	2.37	9.64	14.6
0.03	12.3	2.01	22.1	13.7
0.03	9.96	2.65	22.1	12
0.05	14.3	0.78	19.7	19.2
0.07	17.4	5.65		17
0.01	13	0.94		19
0.026102	18.46033898	1.27372881 ####	20.41684211	19.50016949
0.06602	14.32948708	1.1041144 ####	15.40882391	14.57853423

③PGE-MUM/Cr	④PGE-MUM/Cr	⑤PGE-MUM/Cr	①FABP/Cr	②FABP/Cr	③FABP/Cr
15.5	17	17	2.07	0.54	2.15
9.93	10.3	8.62	3.16	0.3	3.92
18.1	23.3	13.9	5.54	1.85	2.51
30.6	21.6	13	0	0.44	13.85
95.4	85.6	54.2	0.77	1.06	1.02
38	42.1	32.1	3.01	1.53	3.32
30.2	35.2	12.8	4.16	3.08	2.67
11.9	12.4	10.7	1.5	0	1.79
18.2	17.7	14	0	0	0.9
29.2	28.2	30.8	3.51	0.99	1.95
18.9	17.1	12.6	2.56	3.08	2.61
19.1	19.2	20.3	2.51	1.53	2.6
16.6	20.6	16.6	0.87	2.83	2.83
29	25.4	16	1.32	2.4	2.7
17.8	19.9	23	0.3	0.98	0.97
15.9	20.9	12.4	4.66	1.37	1.18
15.1	11.9	10.8	1.7	4.14	5.07
15.5	13.6	9.8	1.31	1.01	1.38
23.6	14.5	14.7	1.16	1.66	2.8
23.8	15.7	13.3	2.95	1.87	5.62
12	17.2	9	3.25	3.28	4.61
32.1	24.6	16.7	1.65	1.94	2.57
15.2	11.4	22	0.94	1.38	1.69
27.3	14.9	11.1	1.11	1.08	1.93
82.7	97.5	102	2.15	0.87	4.19
12.5	13.3	11.2	0.7	1.22	1.22
28.9	23	12.4	0.84	0.8	1.2
8.94	23.5	11.6	1.56	1.24	0.92
17.2	19.6	18	2	1.32	1.31
33.5	23.1	16.4	2.75	1.37	10
25.1	22.1	17.4	0.78	1.74	3.52
26.5	28.7	12.1	2.08	0.85	2.69
28.9	21.2	18.1	1.73	1.32	3.41
40.1	38.5	29.3	2.57	1.54	1.78
23.4	28.5	17.5	1.45	2.65	6.41
19.4	14.2	12.1	2.15	2.22	5.93
36.7	26	24.7	2.28	0	2.61
23.8	20.8	16.9	4.77	1.24	2.39

29.7	15.9	10.6	1.79	1.09	3.96
29.8	42.1	17.8	0.71	1.21	2.32
44.9	38.4	34.3	1.89	1.91	4.72
28.8	27.1	17.1	0	1.77	0.88
12.9	15.3	11.9	0	2.68	0.66
29.4	24.7	24.2	1.98	0.49	1.39
21.8	22.9	15.4	1.86	0.33	6.43
16	19.7	7.98	0.96	2.77	3.51
65.1	62.3	53.7	1.65	4.63	10.2
25	23.8	10.5	2.19	1.61	2.88
19.1	15.8	12.3	0.62	2.81	3.2
18	22.9	18.2	0	3.67	2.03
17.2	19.7	11.7	2.37	0.79	2.92
58.7	37.6	22.2	1.47	3.62	11.96
11.1	8.67	8.3	1.15	1.25	1.08
18.9	14.3	10.9	3.05	2.52	3.93
22.5	15.9	12.3	0.45	1.89	1.45
17.5	11.7	9.96	4.99	5.71	2.4
24.9	18.8	14.3	0.54	0.57	16.56
25.3	21.6	17.4		3.76	17.11
23.9	20.2	13		1.21	2.16
26.22152542	24.40118644	18.46033898	1.8507018	1.7459322	3.76220339
15.92345546	15.77647355	14.32948708	1.3009528	1.186775	3.62815763

④FABP/Cr	⑤FABP/Cr	①TP/Cr	②TP/Cr	③TP/Cr	④TP/Cr	⑤TP/Cr
1.39	0.94	0	0	0.04	0.01	0.02
1.48	1.19	0.04	0	0.05	0.01	0.01
2.1	1.2	0.01	0	0.02	0.02	0
5.2	0.34	0	0	0.38	0.03	0
0.96	0.77	0	0	0.13	0.04	0
2.56	2.75	0.22	0	0.01	0	0.03
2.36	0.92	0	0.02	0.04	0.03	0
1.27	0.28	0.01	0.01	0.03	0.04	0
0.83	0	0	0	0.04	0.01	0
0	0.34	0.02	0	0.02	0	0
2.1	4.08	0.01	0	0.01	0.02	0.02
0.89	0.75	0	0	0.01	0	0.51
1.5	1.27	0.02	0	0.03	0.02	0.02
2.84	2.07	0.02	0.02	0.05	0.02	0.02
1.11	0	0	0.02	0.03	0.01	0
0.87	0.75	0.03	0	0.01	0.03	0
2.03	0.39	0	0.05	0.09	0.01	0
1.03	0.42	0.01	0	0.17	0.06	0
2.08	1.01	0.02	0	0.28	0.05	0.03
1.3	1.93	0	0	0.12	0.05	0.02
4.04	1.47	0.02	0.02	0.02	0.02	0.02
1.68	1.24	0	0	0.05	0.02	0.02
0.6	1.21	0	0.01	0.05	0.01	0.02
1.24	0.88	0.02	0.01	0.06	0.02	0.02
1.24	0.58	0.01	0.03	0.16	0	0.05
0.89	1.13	0.02	0.01	0.02	0	0.02
0.99	0.47	0	0.02	0.12	0.01	0.03
0.94	1.19	0	0	0.02	0	0.03
0.75	0.46	0	0.02	0.03	0.33	0.02
3.14	0.24	0	0.01	0.2	0.01	0.01
2	0.86	0	0.02	0.07	0.02	0.04
8.36	0.72	0	0.03	0.1	0.03	0.03
1.81	0.34	0.05	0.01	0.47	0.01	0
1.84	0	0.03	0.04	0.17	0.04	0
8.58	3.3	0	0.02	0.15	0.04	0.04
2.72	0.86	0	0.08	0.05	0.01	0.02
2.63	0.91	0	0	0.06	0.02	0.01
2.05	1.48	0.01	0	0.13	0.01	0.02

4.37	1.59	0.02	0.01	0.24	0.01	0.02
1.64	0.7	0	0.03	0.07	0.03	0.01
2.03	0.99	0.02	0.01	0.02	0	0
0.55	1.14	0	0.01	0.05	0.03	0.01
1.16	3	0	0	0	0	0.02
0.77	0.58	0.04	0	0.02	0.04	0
1.24	0	0	0	0.08	0.03	0
2.35	2.03	0	0.03	0.18	0.13	0.05
1.12	2.69	0	0	0.04	0	0.01
4.42	0.72	0.01	0.03	0.38	0.13	0.04
3.23	0.39	0	0.01	0.01	0.02	0
3.23	3.19	0	0.03	0.22	0.03	0.03
1.5	0.82	0.02	0.02	0.36	0.15	0.03
5.75	3.28	0	0.1	0.59	0.11	0.02
0.89	0.89	0	0	0.04	0.02	0.01
3.01	2.37	0.03	0.04	0.04	0.04	0.02
6.2	2.01	0	0.02	0.1	0.09	0.03
2.66	2.65	0	0.04	0.03	0.04	0.03
0	0.78	0	0.02	0.11	0	0.05
22.08	5.65		0.03	0.19	0.08	0.07
1.76	0.94		0	0.06	0.03	0.01
2.53152542	1.2737288	0.01246	0.01492	0.10712	0.03508	0.0261
3.12946884	1.1041144	0.03066	0.0196	0.12257	0.05131	0.06602

①Total protein(mg/dL)	②Total pr	③Total pr	④Total pr	⑤Total protein	L-FABP①(ng/mL)
0	0	13.2	3.2	4	2.35
8.1	0	17.6	2.2	2.5	6.47
2.6	0	8.5	5.8	0	14
0	0	143.3	6.6	0	0
0	0	43.7	11.3	0	1.04
38.9	0	3.7	0	6.1	5.31
0	3.8	14.7	7.3	0	5.88
3.1	2.2	8	13	0	3.76
0	0	13.8	3.3	0	0
4.6	0	7.1	0	0	8.49
2.2	0	3.1	8	3.2	4.91
0	0	7.7	0	111.2	2.7
3	0	10.9	6.4	4.7	1.46
3.6	2.9	23	8	3.5	2.7
0	2.7	10.6	2	0	0.53
6.7	0	4.4	13.4	0	9.97
0	7.2	31	3.6	0	2.76
2.1	0	64	19.1	0	1.85
2.3	0	142.6	22.1	10	1.32
0	0	31.8	11.4	3.6	5.22
4.3	4.9	10.1	5.9	7.5	8.21
0	0	20.7	4.7	6.7	3.64
0	2.1	17.6	2.2	2.7	1.17
2	3.8	28.7	8.1	5.8	1.41
2.9	4.6	66.9	0	10.7	4.4
2.2	4	7.9	0	3.5	0.86
0	4.7	43.3	3	8.1	1.06
0	0	12.6	0	10	2.91
0	3.3	7	90.2	3.9	2.76
0	2.7	68.6	3.5	2.5	3.13
0	5	35.5	7.1	8	0.93
0	5.6	38.8	16.1	8.4	3.97
13.6	3	215.5	3.8	0	4.63
3.1	13.6	76.4	13.5	0	2.65
0	3.7	50	10.8	9.5	2.02
0	16.7	18.7	2.3	3.1	1.75
0	0	27	5.6	3.2	2.04
2.1	0	58.3	2.7	7.4	9.69

2.7	2	204.9	2.5	6.9	2.26
0	6.1	38.7	11.8	3.7	1.17
3.2	2.7	8.6	0	0	3.3
0	2.4	17.9	10.3	2.8	0
0	0	0	0	2.4	0
4.6	0	5.9	10.1	0	2.23
0	0	19.1	5.6	0	2.06
0	5.7	56.8	25.2	16.2	1.25
0	0	13	0	2.3	2.02
2.8	8.2	199.9	53.4	16	5.4
0	2	4.8	6.3	0	0.71
0	6.6	73.5	7.7	8	0
3.8	2.6	97.9	38.1	5.6	5.24
0	30.3	261.2	40.6	5.1	3.09
0	0	12.7	5.8	2.7	0.58
4.5	11.8	19.7	15.5	4.6	5.06
0	2.9	49.3	30.7	6.3	0.77
0	7	11.7	11.3	9.6	6.78
0	2.8	23.7	0	10.6	1.11
	2.5	44.6	11.7	13.4	
	0	19.6	12.5	2.54	
2.263157895	3.255932	43.89492	10.59831	6.246440678	3.175087719
5.54483774	5.020107	57.08988	14.8753	14.51179565	2.845389832

L-FABP②	L-FABP③	L-FABP④	L-FABP⑤	①Creatini	②Creatini	③Creatini	④Creatini
1.14	7.5	3.5	2.42	113.58	210.83	349.35	251.76
0.55	13.43	3.9	2.3	204.94	181.63	342.45	264.05
3.59	9.71	6.87	2.76	252.79	194.07	386.95	326.65
0.6	52.42	13.26	0.89	116.21	136.29	378.55	254.76
2.68	3.57	3.03	1.63	135.01	253.7	349.05	315.32
2.15	11.19	5.72	6.11	176.36	140.78	336.82	223.77
7.44	11.14	6.57	2.58	141.27	241.64	417.7	278.27
0	4.17	4.43	0.54	251.25	186.83	233.43	349.9
0	2.87	2.1	0	166.67	154.47	320.13	254.26
2.03	7.01	0	0.58	241.83	205.4	359.05	259.76
8.99	13.2	9.57	7.42	191.71	291.61	504.8	455.6
1.91	13.64	1.49	1.63	107.46	124.71	523.75	167.82
4.16	9.54	4.56	3.63	167.71	147.11	336.85	304.15
3.97	11.93	11.76	3.95	203.91	165.27	441.4	414
1.73	3.26	2.2	0	175.7	177.43	335.11	199.03
2	4.11	3.49	1.31	214.12	146.26	347.55	399.6
5.87	17.41	5.99	0.79	162.82	141.84	343.15	294.78
1.67	5.2	3.07	0.87	140.9	165.59	375.75	297.29
2.23	14.12	9.56	3.29	114.23	134.5	504.5	458.8
2.29	15.02	3.05	3.16	176.83	122.53	267.46	234.01
8.88	18.68	15.29	5.18	252.8	270.52	405.2	378.75
2.89	10.98	4.63	3.62	219.99	149.1	427	275.9
2.21	5.81	1.09	2.07	124.82	160.55	343.68	180.76
3.17	9.99	4.33	2.94	127.42	292.49	517.3	349.97
1.42	17.06	2.5	1.14	205.01	163.41	407.35	202.12
3.4	4.04	1.15	1.86	123.53	279.6	330	128.96
1.93	4.46	2.22	1.33	126.64	240.01	373.15	224.3
4.24	4.89	2.63	4.31	186.98	341.54	533.55	279.22
2.7	3.56	2.06	1.17	137.85	205.18	271.64	275.63
3.09	33.77	9.71	0.51	113.75	225.8	337.6	309.51
4.04	17.01	6.68	1.8	118.67	231.56	482.8	333.26
1.4	10.37	39.06	1.77	190.88	165.6	384.95	466.95
3.18	15.69	8.26	0.62	268.3	240.25	460.75	457
5.31	7.85	6.22	0	103	344.03	441.15	338.05
4.02	21.49	21.81	7.66	139.15	151.66	335.05	254.06
4.4	22.32	7.95	1.57	81.54	198.11	376.2	292.28
0	12.73	7.56	1.98	89.38	215.31	487.7	286.99
2.81	10.43	6.23	4.44	203.22	226.01	436.55	303.86

3.43	33.73	12.85	5.68	126.04	315.15	852.65	293.79
2.81	13.59	6.38	1.79	165.65	232.45	584.8	389.25
5.32	17.33	8.18	3.36	174.98	278.45	367.15	403.7
2.84	3.17	1.8	2.49	177.67	160.22	360.9	325.42
4.21	1.29	3.33	4.8	194.95	156.84	196.5	288.13
0.88	4.48	1.99	1.13	112.66	178.84	321.52	258.11
0.41	15.31	2.62	0	110.52	151.23	238.15	210.73
4.82	11.21	4.72	6.39	130.28	173.93	319.05	200.65
10.88	34.14	3.26	8.77	122.17	234.91	334.65	290.36
4.18	15.32	17.6	2.6	247.04	260.02	532.4	398.4
4.52	11.2	11.08	0.86	115.22	160.66	350.05	342.81
8.22	6.71	7.74	7.59	143.7	223.68	330.85	239.88
0.91	7.92	3.73	1.41	220.83	114.78	271.06	248.38
10.69	53.03	21.28	7.94	210.89	295.06	443.25	369.9
2.86	3.59	2.55	2.56	50.29	228.86	331.26	286.17
7.73	17.74	12.08	4.69	165.95	306.82	451.3	401.35
2.28	7.26	20.57	4.78	170.1	120.78	501.2	331.8
10.55	10.51	8.15	8.1	135.97	184.85	437.05	306.71
0.99	36.8	0	1.82	204.48	174.3	222.25	103.08
3.46	39.39	33.48	10.84		92.11	230.19	151.65
1.77	6.92	7.51	2.39		146.75	320.96	427.77
3.52288136	13.76627	7.532203	3.047797	162.2389	200.2358	386.4849	298.4608
2.68311454	11.41553	7.477127	2.522315	49.39664	60.35375	106.0856	83.09996

⑤Creatini ①PGE-MI ②PGE-MI ③PGE-MI ④PGE-MI ⑤PGE-MUM

257.17	11.6	26.4	54.3	42.8	43.7
193.77	19.9	14.5	34	27.1	16.7
230.25	33.3	30.6	70.1	76	31.9
265.57	13.8	15.5	116	55	34.6
210.5	53.3	125	333	270	114
222.41	61.4	44.2	128	94.1	71.3
280.22	35.5	44.4	126	98	35.8
194.25	23.2	17.5	27.7	43.5	20.8
180.63	30.4	20.3	58.4	45.1	25.3
168.99	61.2	47.1	105	73.3	52.1
182.06	22.1	33.2	95.3	77.7	22.9
216.47	11.3	16.1	99.9	32.2	44
286.55	53.1	25.1	56	62.6	47.5
190.7	44.6	29.6	128	105	30.5
195.44	23	29.8	59.5	39.6	45
175.81	24.2	16.4	55.3	83.5	21.8
201.74	28.2	14.4	51.7	35.2	21.8
209.16	28.6	30	58.4	40.5	20.5
324.24	16.7	16.6	119	66.7	47.8
163.68	17.8	18.3	63.6	36.7	21.8
352.08	45.4	24.1	48.7	65.3	31.7
292.43	32.8	28.3	137	68	48.7
170.99	13.6	14.5	52.3	20.6	37.6
332.48	141	31.6	141	52.2	36.8
197.28	20.7	158	337	197	202
165.31	20.2	32.3	41.3	17.1	18.5
282.21	12.6	49.8	108	51.5	35
362.19	31	23.5	47.7	65.6	42
256.58	27.7	37.9	46.6	54	46.2
208.33	22.4	52.6	113	71.6	34.2
208.13	19	57.6	121	73.6	36.2
245.92	30.8	25.4	102	134	29.7
180.35	107	40.2	133	96.9	32.6
183.2	17.4	130	177	130	53.6
232	18.7	24.2	78.5	72.5	40.7
182.27	19.1	23.6	72.9	41.5	22
218.55	12.6	48.4	179	74.7	53.9
299.19	30.4	38.2	104	63.1	50.7

357.54	51.6	31.8	253	46.8	37.9
255.51	67.9	67.2	174	164	45.5
338.11	51.1	146	165	155	116
218.58	36.5	47.4	104	88.2	37.3
159.85	47.1	22.4	25.4	44.1	19
194.42	12.9	31.4	94.5	63.7	47
144.25	53.5	25	51.9	48.2	22.2
314.5	15.5	16.6	51.2	39.5	25.1
325.84	11.4	142	218	181	175
358.71	39.4	35.7	133	94.8	37.8
222.93	14.5	18.5	66.8	54.3	27.5
238.15	39.7	39.1	59.4	55	43.4
172.8	23.4	15.6	46.7	49	20.3
241.83	25.8	77.5	260	139	53.6
286.75	6.3	24.6	36.7	24.8	23.8
197.79	16	44.9	85.4	57.3	21.5
237.46	37.6	16.5	113	52.6	29.1
305.2	30.1	22.2	76.7	35.8	30.4
234.33	40.3	33.5	55.4	19.4	33.5
191.76		15.7	58.3	32.8	33.3
254.62		27.9	76.6	86.6	33
236.7802	32.56491	39.94407	103.122	72.63898	42.95085
58.66686	23.19274	33.73429	68.10461	47.23526	33.21467