Additional file 1 - This additional file summarizes the number and proportion of q-values, and p-value of 100 permutation results for different ranked score and q-value range under each of the 10 weighting score matrices.

## Weighting score matrix 1

Table 1. Number and proportion of q-values for each q-value range and ranked gene score under weighting score matrix 1.

Score	≥0.5	≥1	≥1.5	≥2	≥2.5	=3
All a value	91774	29001	8651	7516	6539	1094
All q value	(100%)	(100%)	(100%)	(100%)	(100%)	(100%)
~<0.0	6399	3000	202 (4.40/)	2012	710	161
q<0.9	(7.0%)	(10.3%)	382 (4.4%)	(26.8%)	(10.9%)	(14.7%)
q<0.8	469 (0.5%)	592 (2.0%)	151 (1.7%)	148 (2.0%)	144 (2.2%)	0
q < 0.7	8 (0.009%)	2 (0.007%)	5 (0.06%)	42 (0.6%)	67 (1.0%)	0
q<0.6	5 (0.005%)	2 (0.007%)	0	0	5 (0.08%)	0
q<0.5	5 (0.005%)	0	0	0	0	0
q<0.4	5 (0.005%)	0	0	0	0	0
q<0.3	0	0	0	0	0	0
q<0.2	0	0	0	0	0	0
q<0.1	0	0	0	0	0	0
q<0.05	0	0	0	0	0	0

Numbers in the parentheses are the proportion of number of q-values.

Table 2. P-value of 100 permutation result for different ranked gene score and q-value range under weighting score matrix 1.

Score	≥0.5	≥1	≥1.5	≥2	≥2.5	=3
q<0.9	0.34	0.31	0.63	0.15	0.44	0.57
q<0.8	0.5	0.32	0.48	0.48	0.49	N/A
q<0.7	0.47	0.68	0.61	0.4	0.37	N/A
q<0.6	0.25	0.5	N/A	N/A	0.5	N/A
q<0.5	0.11	N/A	N/A	N/A	N/A	N/A
q<0.4	0.02	N/A	N/A	N/A	N/A	N/A

N/A: not available due to absence of the real data at those categories.

Table 3. Number and proportion of q-values for each q-value range and ranked gene score under weighting score matrix 2.

Score	≥0.5	≥1	≥1.5	≥2	=2.5
All a realisa	91774	29001	8651	6539	1094
All q value	(100%)	(100%)	(100%)	(100%)	(100%)
q<0.9	6399	3000	382	710	161
q~0.9	(7.0%)	(10.3%)	(4.4%)	(10.9%)	(14.7%)
q<0.8	469	592	151	144	0
	(0.5%)	(2.0%)	(1.7%)	(2.2%)	U
q<0.7	8	2	5 (0.06%)	67 (1.0%)	0
	(0.009%)	(0.007%)	3 (0.0070)	07 (1.070)	O
q<0.6	5	2	0	5 (0.08%)	0
q 10.0	(0.005%)	(0.007%)	V	3 (0.0070)	O .
q<0.5	5	0	0	0	0
9 0.0	(0.005%)	V	V	· ·	· ·
q<0.4	5	0	0	0	0
	(0.005%)	v	v		· ·
q<0.3	0	0	0	0	0
q<0.2	0	0	0	0	0
q<0.1	0	0	0	0	0
q<0.05	0	0	0	0	0

Table 4. P-value of 100 permutation result for different ranked gene score and q-value range under weighting score matrix 2.

Score	≥0.5	≥1	≥1.5	≥2	=2.5
q<0.9	0.46	0.26	0.68	0.33	0.49
q<0.8	0.47	0.26	0.51	0.35	N/A
q<0.7	0.51	0.59	0.68	0.31	N/A
q<0.6	0.34	0.45	N/A	0.4	N/A
q<0.5	0.17	N/A	N/A	N/A	N/A
q<0.4	0.01	N/A	N/A	N/A	N/A

Table 5. Number and proportion of q-values for each q-value range and ranked gene score under weighting score matrix 3.

Score	≥0.5	≥1	≥1.5	≥2	=2.5
All a value	91774	18988	7948	2293	210
All q value	(100%)	(100%)	(100%)	(100%)	(100%)
a<0.0	6399	1863	399	866	178
q<0.9	(7.0%)	(9.8%)	(5.0%)	(37.8%)	(84.8%)
q<0.8	469	415	163	199	108
<b>q</b> <0.8	(0.5%)	(2.2%)	(2.1%)	(8.7%)	(51.4%)
q<0.7	8	164	5 (0.069/)	117 (5 10/)	72
	(0.009%)	(0.9%)	5 (0.06%)	117 (5.1%)	(34.3%)
q<0.6	5	0	0	42 (1.8%)	53
<b>q</b> <0.0	(0.005%)	U	U	42 (1.070)	(25.2%)
q<0.5	5	0	0	39 (1.7%)	0
q <0.5	(0.005%)	U	U	39 (1.770)	U
q<0.4	5	0	0	27 (1.2%)	0
q<0.4	(0.005%)	U	U	27 (1.270)	U
q<0.3	0	0	0	0	0
q<0.2	0	0	0	0	0
q<0.1	0	0	0	0	0
q<0.05	0	0	0	0	0

Table 6. P-value of 100 permutation result for different ranked gene score and q-value range under weighting score matrix 3.

Score	≥0.5	≥1	≥1.5	≥2	=2.5
q<0.9	0.39	0.42	0.68	0.18	0.05
q<0.8	0.49	0.35	0.53	0.28	0.16
q<0.7	0.5	0.23	0.67	0.21	0.09
q<0.6	0.24	N/A	N/A	0.26	0.05
q<0.5	0.07	N/A	N/A	0.16	N/A
q<0.4	0.01	N/A	N/A	0.13	N/A

Table 7. Number and proportion of q-values for each q-value range and ranked gene score under weighting score matrix 4.

Score	≥0.5	≥1	≥1.5	≥2	≥2.5	=3
All a value	91774	29001	15768	7674	1191	570
All q value	(100%)	(100%)	(100%)	(100%)	(100%)	(100%)
a<0.0	6399	3000	602 (4.49/)	240 (4.50/)	2 (0.29/)	114 (200/)
q<0.9	(7.0%)	(10.3%)	692 (4.4%)	349 (4.5%)	2 (0.2%)	114 (20%)
q<0.8	469 (0.5%)	592 (2.0%)	305 (1.9%)	143 (1.9%)	0	76 (13.3%)
q < 0.7	8 (0.009%)	2 (0.007%)	86 (0.5%)	5 (0.07%)	0	0
q<0.6	5 (0.005%)	2 (0.007%)	0	0	0	0
q<0.5	5 (0.005%)	0	0	0	0	0
q<0.4	5 (0.005%)	0	0	0	0	0
q<0.3	0	0	0	0	0	0
q<0.2	0	0	0	0	0	0
q<0.1	0	0	0	0	0	0
q<0.05	0	0	0	0	0	0

Table 8. P-value of 100 permutation result for different ranked gene score and q-value range under weighting score matrix 4.

Score	≥0.5	≥1	≥1.5	≥2	≥2.5	=3
q<0.9	0.41	0.36	0.59	0.58	0.83	0.51
q<0.8	0.52	0.29	0.36	0.43	N/A	0.34
q<0.7	0.57	0.63	0.29	0.58	N/A	N/A
q<0.6	0.31	0.4	N/A	N/A	N/A	N/A
q<0.5	0.12	N/A	N/A	N/A	N/A	N/A
q<0.4	0.03	N/A	N/A	N/A	N/A	N/A

Table 9. Number and proportion of q-values for each q-value range and ranked gene score under weighting score matrix 5.

Score	≥1	≥1.5	≥2	≥2.5	≥3	=3.5
All a value	91774	18988	11168	6686	1171	266
All q value	(100%)	(100%)	(100%)	(100%)	(100%)	(100%)
a<0.0	6399	1863	2920	294 (4.4%)	310	132
q<0.9	(7.0%)	(9.8%)	(26.1%)	294 (4.4%)	(26.5%)	(50.0%)
q<0.8	469 (0.5%)	415 (2.2%)	397 (3.6%)	126 (1.9%)	0	5 (1.9%)
q < 0.7	8 (0.009%)	164 (0.9%)	109 (1.0%)	0	0	4 (1.5%)
q<0.6	5 (0.005%)	0	0	0	0	2 (0.8%)
q < 0.5	5 (0.005%)	0	0	0	0	0
q<0.4	5 (0.005%)	0	0	0	0	0
q<0.3	0	0	0	0	0	0
q<0.2	0	0	0	0	0	0
q<0.1	0	0	0	0	0	0
q<0.05	0	0	0	0	0	0

Table 10. P-value of 100 permutation result for different ranked gene score and q-value range under weighting score matrix 5.

Score	≥1	≥1.5	≥2	≥2.5	≥3	=3.5
q<0.9	0.42	0.33	0.06	0.64	0.36	0.54
q<0.8	0.59	0.33	0.18	0.44	N/A	0.76
q<0.7	0.58	0.27	0.2	N/A	N/A	0.68
q<0.6	0.29	N/A	N/A	N/A	N/A	0.54
q<0.5	0.12	N/A	N/A	N/A	N/A	N/A
q<0.4	0	N/A	N/A	N/A	N/A	N/A

Table 11. Number and proportion of q-values for each q-value range and ranked gene score under weighting score matrix 6.

Score	≥0.5	≥1	≥1.5	≥2	≥2.5	=3
All a volue	91774	81761	18285	10883	1700	405
All q value	(100%)	(100%)	(100%)	(100%)	(100%)	(100%)
q<0.9	6399	5600	1902	728	0	224
q<0.9	(7.0%)	(6.8%)	(10.4%)	(6.7%)	U	(55.3%)
a<0.9	469	463	507	272	0	117
q<0.8	(0.5%)	(0.6%)	(2.8%)	(2.5%)	U	(28.9%)
a<0.7	8	26	179	50 (0.50/)	0	6 (1.50/)
q<0.7	(0.009%)	(0.03%)	(1.0%)	59 (0.5%)	U	6 (1.5%)
a<0.6	5	4	0 0	0	6 (1.50/)	
q<0.6	(0.005%)	(0.005%)	U	U	U	6 (1.5%)
a<0.5	5	4	0	0	0	0
q<0.5	(0.005%)	(0.005%)	U	U	U	U
a<0.4	5	0	0	0	0	0
q<0.4	(0.005%)	U	U	U	U	U
q<0.3	0	0	0	0	0	0
q<0.2	0	0	0	0	0	0
q<0.1	0	0	0	0	0	0
q<0.05	0	0	0	0	0	0

Table 12. P-value of 100 permutation result for different ranked gene score and q-value range under weighting score matrix 6.

Score	≥0.5	≥1	≥1.5	≥2	≥2.5	=3
q<0.9	0.46	0.39	0.38	0.61	N/A	0.29
q<0.8	0.5	0.59	0.23	0.39	N/A	0.26
q<0.7	0.51	0.21	0.24	0.38	N/A	0.57
q<0.6	0.32	0.11	N/A	N/A	N/A	0.46
q<0.5	0.2	0.02	N/A	N/A	N/A	N/A
q<0.4	0.03	N/A	N/A	N/A	N/A	N/A

Table 13. Number and proportion of q-values for each q-value range and ranked gene score under weighting score matrix 7.

Score	≥1	≥2	≥2.5	≥3	=3.5
All a volue	91774	18988	6686	1762	774 (100%)
All q value	(100%)	(100%)	(100%)	(100%)	774 (100%)
q<0.9	6399 (7.0%)	1863 (9.8%)	294 (4.4%)	0	2 (0.3%)
q<0.8	469 (0.5%)	415 (2.2%)	126 (1.9%)	0	2 (0.3%)
q<0.7	8 (0.009%)	164 (0.09%)	0	0	0
q<0.6	5 (0.005%)	0	0	0	0
q<0.5	5 (0.005%)	0	0	0	0
q<0.4	5 (0.005%)	0	0	0	0
q<0.3	0	0	0	0	0
q<0.2	0	0	0	0	0
q<0.1	0	0	0	0	0
q<0.05	0	0	0	0	0

Table 14. P-value of 100 permutation result for different ranked gene score and q-value range under weighting score matrix 7.

Score	≥1	≥2	≥2.5	≥3	=3.5
q<0.9	0.4	0.53	0.59	N/A	0.8
q<0.8	0.46	0.33	0.47	N/A	0.65
q<0.7	0.56	0.24	N/A	N/A	N/A
q<0.6	0.33	N/A	N/A	N/A	N/A
q<0.5	0.14	N/A	N/A	N/A	N/A

Table 15. Number and proportion of q-values for each q-value range and ranked gene score under weighting score matrix 8.

Score	≥0.5	≥1	≥1.5	≥2	≥2.5	=3
All q	91774	18988	7948	7516	2293	210
value	(100%)	(100%)	(100%)	(100%)	(100%)	(100%)
a<0.0	6200 (7.09/)	1962 (0.90/)	200 (5.00/)	2012	866	178
q<0.9	6399 (7.0%)	1863 (9.8%)	399 (5.0%)	(26.7%)	(37.8%)	(84.8%)
a<0.9	460 (0.50/)	415 (2.20/)	162 (2 10/)	149 (2.00/)	199 (8.7%)	108
q<0.8	469 (0.5%)	415 (2.2%)	163 (2.1%)	148 (2.0%)		(51.4%)
q<0.7	8 (0.009%)	164 (0.09%)	5 (0.06%)	42 (0.6%)	117 (5.1%)	72 (34.3%)
q<0.6	5 (0.005%)	0	0	0	42 (1.8%)	53 (25.2%)
q<0.5	5 (0.005%)	0	0	0	39 (1.7%)	0
q<0.4	5 (0.005%)	0	0	0	27 (1.2%)	0
q<0.3	0	0	0	0	0	0
q<0.2	0	0	0	0	0	0
q<0.1	0	0	0	0	0	0
q<0.05	0	0	0	0	0	0

Table 16. P-value of 100 permutation result for different ranked gene score and q-value range under weighting score matrix 8.

Score	≥0.5	≥1	≥1.5	≥2	≥2.5	=3
q<0.9	0.36	0.4	0.55	0.13	0.2	0.09
q<0.8	0.48	0.34	0.36	0.38	0.24	0.13
q<0.7	0.53	0.29	0.63	0.35	0.16	0.13
q<0.6	0.33	N/A	N/A	N/A	0.21	0.08
q<0.5	0.15	N/A	N/A	N/A	0.1	N/A
q<0.4	0.01	N/A	N/A	N/A	0.08	N/A

Table 17. Number and proportion of q-values for each q-value range and ranked gene score under weighting score matrix 9.

Score	≥1	≥2	≥3	=4
All a volue	91774	18988	7948	1330
All q value	(100%)	(100%)	(100%)	(100%)
q<0.9	6399 (7.0%)	1863 (9.8%)	399 (5.0%)	0
q<0.8	469 (0.5%)	415 (2.2%)	163 (2.1%)	0
q<0.7	8 (0.009%)	164 (0.09%)	5 (0.06%)	0
q<0.6	5 (0.005%)	0	0	0
q<0.5	5 (0.005%)	0	0	0
q<0.4	5 (0.005%)	0	0	0
q<0.3	0	0	0	0
q<0.2	0	0	0	0
q<0.1	0	0	0	0
q<0.05	0	0	0	0

Table 18. P-value of 100 permutation result for different ranked gene score and q-value range under weighting score matrix 9.

Score	≥1	≥2	≥3	=4
q<0.9	0.4	0.36	0.55	N/A
q<0.8	0.49	0.34	0.43	N/A
q<0.7	0.56	0.21	0.6	N/A
q<0.6	0.26	N/A	N/A	N/A
q<0.5	0.08	N/A	N/A	N/A
q<0.4	0.01	N/A	N/A	N/A

Table 19. Number and proportion of q-values for each q-value range and ranked gene score under weighting score matrix 10.

Score	≥1	≥2	≥2.5	≥3	=3.5
A 11	91774	18988	7948	1762	1330
All q value	(100%)	(100%)	(100%)	(100%)	(100%)
a<0.0	6399	1863	200 (5.0%)	0	0
q<0.9	(7.0%)	(9.8%)	399 (5.0%)	U	U
q<0.8	469 (0.5%)	415 (2.2%)	163 (2.1%)	0	0
q<0.7	8 (0.009%)	164	5 (0.06%)	0	0
q 30.7	0 (0.00570)	(0.09%)		O	O
q<0.6	5 (0.005%)	0	0	0	0
q<0.5	5 (0.005%)	0	0	0	0
q<0.4	5 (0.005%)	0	0	0	0
q<0.3	0	0	0	0	0
q<0.2	0	0	0	0	0
q<0.1	0	0	0	0	0
q<0.05	0	0	0	0	0

Table 20. P-value of 100 permutation result for different ranked gene score and q-value range under weighting score matrix 10.

Score	≥1	≥2	≥2.5	≥3	=3.5
q<0.9	0.48	0.35	0.65	N/A	N/A
q<0.8	0.56	0.32	0.54	N/A	N/A
q<0.7	0.56	0.24	0.69	N/A	N/A
q<0.6	0.41	N/A	N/A	N/A	N/A
q<0.5	0.18	N/A	N/A	N/A	N/A
q<0.4	0.02	N/A	N/A	N/A	N/A