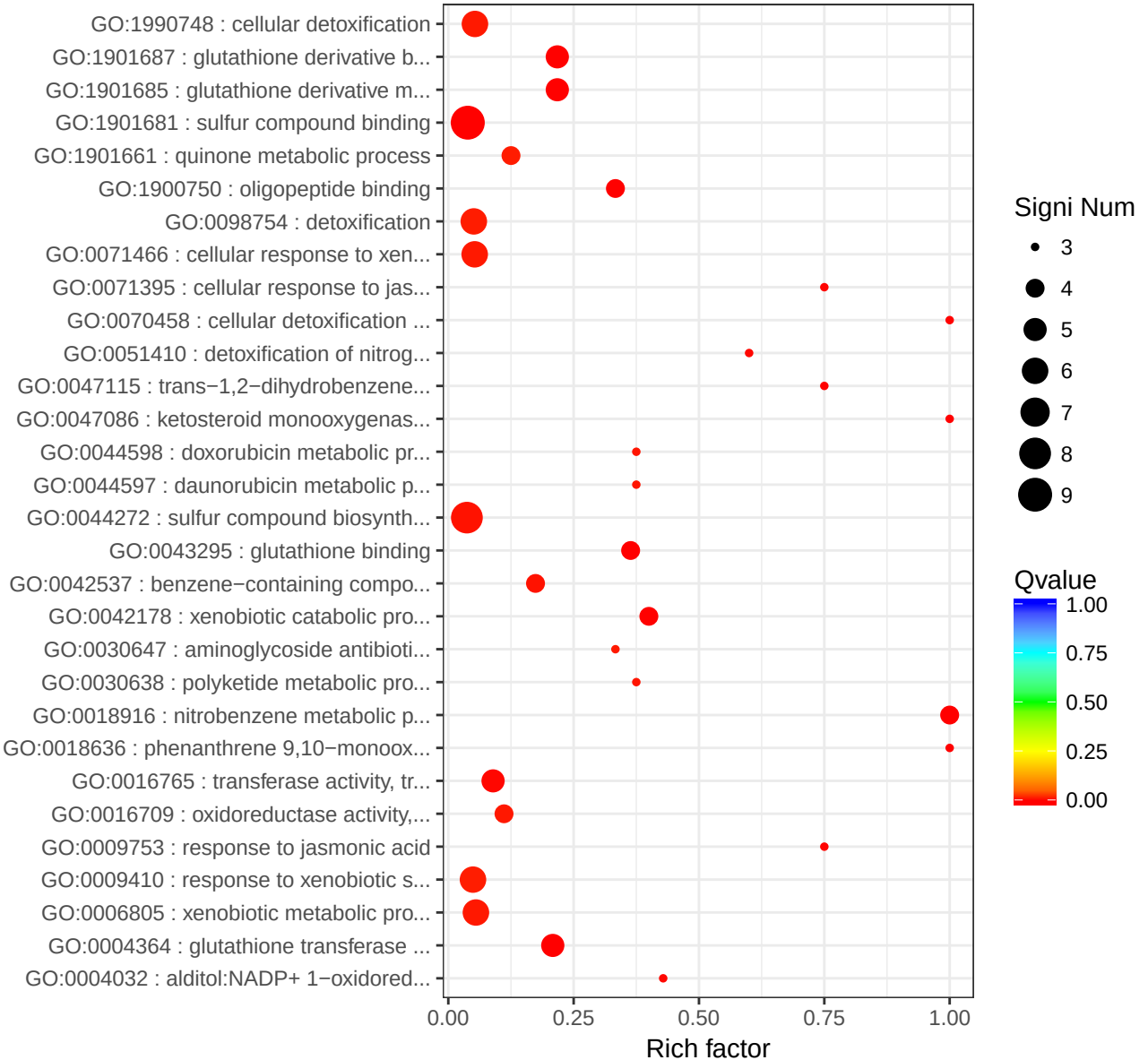
1 **Figure S1. GO analysis results for the differentially expressed gene between Group A and C.**

2


GO:1990748 : cellular detoxification
GO:1901687 : glutathione derivative b...
GO:1901685 : glutathione derivative m...
GO:1901681 : sulfur compound binding
GO:1901661 : quinone metabolic process
GO:1900750 : oligopeptide binding
GO:0098754 : detoxification
GO:0071466 : cellular response to xen...
GO:0071395 : cellular response to jas...
GO:0070458 : cellular detoxification ...
GO:0051410 : detoxification of nitrog...
GO:0047115 : trans−1,2−dihydrobenzene...
GO:0047086 : ketosteroid monooxygenas...
GO:0044598 : doxorubicin metabolic pr...
GO:0044597 : daunorubicin metabolic p...
GO:0044272 : sulfur compound biosynth...
GO:0043295 : glutathione binding
GO:0042537 : benzene−containing compo...
GO:0042178 : xenobiotic catabolic pro...
GO:0030647 : aminoglycoside antibioti...
GO:0030638 : polyketide metabolic pro...
GO:0018916 : nitrobenzene metabolic p...
GO:0018636 : phenanthrene 9,10−monoox...
GO:0016765 : transferase activity, tr...
GO:0016709 : oxidoreductase activity,...
GO:0009753 : response to jasmonic acid
GO:0009410 : response to xenobiotic s...
GO:0006805 : xenobiotic metabolic pro...
GO:0004364 : glutathione transferase ...
GO:0004032 : alditol:NADP+ 1−oxidored...
0.00
0.25
0.50
0.75
1.00
Rich factor
Signi Num
3
4
5
6
7
8
9
Qvalue
1.00
0.75
0.50
0.25
0.00