

Table S8. Robustness of RGC to Thresholds.

	Fold Change Threshold						
	log2(6/5)	log2(3/2)	log2(5/3)	log2(2)	log2(5/2)	log2(3)	log2(4)
0.001	0.9862	0.9862	0.9862	0.9889	0.9938	0.9900	0.9714
0.0025	0.9916	0.9916	0.9916	0.9954	0.9954	0.9898	0.9684
0.005	0.9938	0.9938	0.9938	0.9985	0.9955	0.9885	0.9665
0.01	0.9950	0.9950	0.9960	1.0000	0.9949	0.9875	0.9659
0.025	0.9929	0.9929	0.9966	0.9978	0.9922	0.9862	0.9633
0.05	0.9915	0.9934	0.9960	0.9949	0.9890	0.9846	0.9633
0.1	0.9899	0.9923	0.9913	0.9841	0.9834	0.9809	0.9629

For a range of fold change and p-value thresholds the correlation of the RGC with the RGC calculated using the chosen threshold values are shown. Cell values are the Pearson Correlation between the used thresholds and other thresholds on the number of identified hits.