

Table S5. Significant Gene Ontology terms associated with genes in group 1.

<i>P</i> -value ^a	GO ID, name
3.6E-03	GO:0016567, protein ubiquitination
6.5E-03	GO:0016311, Dephosphorylation
7.9E-03	GO:0006397, mRNA processing
1.2E-02	GO:0006396, RNA processing
3.5E-02	GO:0009968, negative regulation of signal transduction
4.9E-02	GO:0022402, cell cycle process

^aBased on Fisher's exact test.