High K562 correlation perturbations

GO Term	
Pos. Reg. Of DNA-templated Transcription (GO:0045893)	
Histone Acetylation (GO:0016573)	
Histone H3 Acetylation (GO:0043966)	
Reg. Of DNA Repair (GO:0006282)	
Histone Deubiquitination (GO:0016578)	
Pos. Reg. Of Nucleic Acid-Templated Transcription (GO:1903508)	
Monoubiquitinated Histone H2A Deubiquitination (GO:0035522)	
Monoubiquitinated Histone Deubiquitination (GO:0035521)	
RNA Polyadenylation (GO:0043631)	
mRNA Polyadenylation (GO:0006378)	
Reg. Of Transcription By RNA Polymerase II (GO:0006357)	
Reg. Of DNA-templated Transcription (GO:0006355)	
Chromatin Remodeling (GO:0006338)	
Chromatin Organization (GO:0006325)	
Reg. Of Cell Cycle (GO:0051726)	
Reg. Of Telomere Maintenance In Response To DNA Damage (GO:1904505)	
Reg. Of Nuclear-Transcribed mRNA Catabolic Process (GO:1900151)	
Pos. Reg. Of Telomere Maintenance In Response To DNA Damage (GO:1904507)	
Reg. Of Cellular Response To Stress (GO:0080135)	
protein-RNA Complex Assembly (GO:0022618)	
Pos. Reg. Of Cytoplasmic Translation (GO:2000767)	
Monoubiquitinated Protein Deubiquitination (GO:0035520)	
Reg. Of DNA Metabolic Process (GO:0051052)	

-log10(Adjusted P-value)