Copy-number correlation attenuation enrichment (Copy-number~Transcriptomics - Copy-number~Proteomics) integrator complex type CC mediator complex BP organellar small ribosomal subunit PTM mitochondrial small ribosomal subunit small ribosomal subunit u12 dependent spliceosome mitochondrial ribosome organellar ribosome ribosomal subunit transcription factor tfiid complex rna elongation rrna processing rrna metabolic process small nuclear ribonucleoprotein complex eukaryotic translation initiation factor 3 complex androgen receptor signaling pathway dna directed rna polymeraseii holoenzyme nucleolar part response to carbohydrate stimulus dna fragmentation during apoptosis amino acid derivative metabolic process integrin complex dystrophin associated glycoprotein complex innate immune response sulfation negative regulation of hydrolase activity anion transport protein tetramerization regulation of gene specific transcription tube morphogenesis tube development collagen acute inflammatory response morphogenesis of an epithelium dna damage responsesignal transduction resulting in induction of apoptosis calcium mediated signaling regulation of peptidyl tyrosine phosphorylation icosanoid metabolic process biogenic amine metabolic process -1.0-0.50.0 0.5 1.0 Enrichment score (GSEA)