			positive regulation of muscle cell apoptotic process (GO:0010861) positive regulation of receptor biosynthetic process (GO:0010870) histone IPZA bubquithation (GO:0033522) regulation of cofactor metabolic process (GO:0051193) regulation of coenzyme metabolic process (GO:0051196) regulation of acetyl—CoA biosynthetic process (GO:00512) regulation of acetyl—CoA biosynthetic process (GO:0043268) histone ubiquithation (GO:0016574) granulocyte differentiation (GO:0030851) amino acid transmembrane transport (GO:0003333) face development (GO:0030324) proteolysis involved in cellular protein catabolic process (GO:0043632) modification—dependent macromolecule catabolic process (GO:0043632) modification—dependent macromolecule catabolic process (GO:0043632) modification—dependent macromolecule catabolic process (GO:0043632) regulation of microtubule polymerization or depolymerization (GO:0031111) tube morphogenesis (GO:0035239) regulation of microtubule depolymerization (GO:0031114) negative regulation of microtubule depolymerization (GO:0007026) body morphogenesis (GO:0010171) regithelial tube morphogenesis (GO:000562) cellular protein complex assembly (GO:0043623) regulation of peptidase activity (GO:0052547) positive regulation of growth (GO:0071705) actin cytoskeleton organization (GO:003036) regulation of small GTPase mediated signal transduction (GO:0051056) negative regulation of protein depolymerization (GO:1901880) coponition (GO:005689) negative regulation of protein processing (GO:001055) negative regulation of rotein saturation (GO:0003414) RNA splicing (GO:000887) negative regulation of cell—matrix adhesion (GO:001689) negative	
ls10ppm	IsCdHet	IsCdHet10ppm		