OKYL029_U_3_0.1vsOKYL029_AS_3_0.1 Biological Process – rankScore 50, p<0.01 <-- RankScore total number of genes --> base-excision repair 14 5 cellular metabolic process 5 regulation of catalytic activity 23 32 endonucleolytic cleavage to generate mature 5'-end of SSU-rRNA from (SSU-rRNA, 5...
O 10 3077 phospholipid catabolic process 2 protein folding 20 cellular localization 17 5 25% 50% 75% 0% 100% Genes Molecular Function - rankScore 50, p<0.01 <-- RankScore total number of genes --> transcription coactivator activity 12 12 9 guanyl-nucleotide exchange factor activity 13 enzyme regulator activity 20 6 motor activity 24 transaminase activity 21 5 chaperone binding 17 10 GO terms acid phosphatase activity 3 5 3077 unfolded protein binding 18 30 carbohydrate:proton symporter activity 16 9 transferase activity, transferring glycosyl groups 25 protein-macromolecule adaptor activity 9 19 oxidoreductase activity, acting on paired donors, with incorporation or reductio... 5 6 copper ion binding 7 heat shock protein binding 13 10 25% 50% 75% 0% 100% Genes Cellular Component – rankScore 50, p<0.01 <-- RankScore total number of genes --> peroxisomal membrane 12 integral component of mitochondrial inner membrane 23 cellular bud tip 21 cellular bud 3 mating projection tip 8 site of polarized growth 8 GO terms cellular bud neck 9 16 yeast-form cell wall 10 17 cell septum 7 5 mitotic spindle pole body 7 23 3077 ΑII extracellular vesicle 6 14 17 peroxisome 29 perinuclear region of cytoplasm 7 5 proteasome complex 15 9 50% 75% 0% 25% 100% Genes Up (p<0.01) Up (p>0.01) Down (p>0.01) Direction Down (p<0.01)