В GO over-representation analysis - biological process **GO** over-representation analysis - molecular function heme bindingcarboxylic acid transport tetrapyrrole bindingmonocarboxylic acid binding p.adjust p.adjust urea transmembrane 0.01 0.0356 transporter activity 0.0360 organic acid transport 0.02 amide transmembrane 0.0364 0.03 transporter activity 0.0368 0.04 0.0372 monooxygenase activityorganic acid bindingkeratinocyte differentiationfatty acid binding -10 15 10 15 5 Count Count