vag: Sum of absolute difference to normalized reference 'counts' Seq depth nknown function 10000 100000 1000000 Unclassified 10000000 binding proteins Transcription nal transduction ulatory functions and nucleotides Protein synthesis Protein fate ement functions thetical proteins lipid metabolism ergy metabolism DNA metabolism liary metabolism ellular processes Cell envelope ups, and carriers acid biosynthesis 0.000 0.025 0.050 0.075 0.100 0.125 0.150 0.175 0.200