mES | Mean value of the gene body coverage Full length sequence 5'-end sequence 3'-end sequence 1.00 -0.75 protocol Gene coverage C1 CAGE:5'-end-Seq CEL-seq:3'-end-Seq Quartz-seq:Full-length-Seq Ram-DA:Full-length-Seq SMARTer:Full-length-Seq STRT:5'-end-Seq SUPer-seq:Full-length-Seq 0.25 0.00 -10 20 30 40 50 60 70 80 90100 10 20 30 40 50 60 70 80 90100 10 20 30 40 50 60 70 80 90100

 μ of the gene body percentile (5'-> 3')