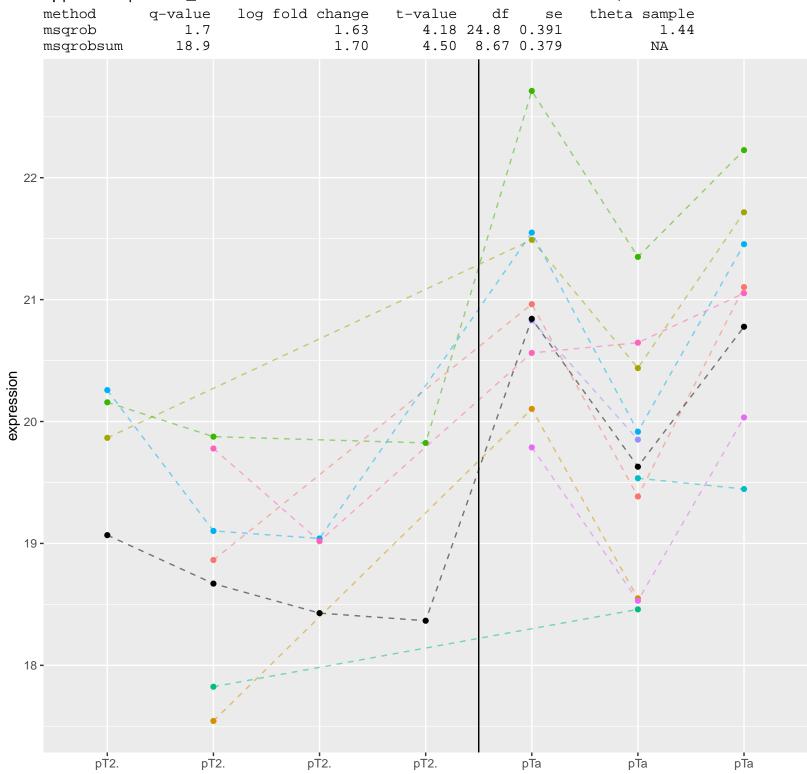
## sp|O15382|BCAT2\_HUMANBranched-chain-amino-acidaminotransferase,mitochondrial



sp|O75369|FLNB\_HUMANFilamin-B method q-value log fold change t-value theta sample df se 0.900 msqrob 1.48 3.89 287. 0.379 1.19 20.7 1.54 4.18 9.68 0.368 msqrobsum NA 22 -20 expression 18**-**16**-**

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sp|O76070|SYUG\_HUMANGamma-synuclein q-value 1 14.1 log fold change 2.19 method t-value theta sample df se msqrob 4.11 49.5 0.533 1.36 msqrobsum 2.19 4.70 NA 9.67 0.467 22.5 expression 20.0 -17.5 pT2. pT2. pT2. рТа рТа pT2. рТа рТа

sp|P00390|GSHR\_HUMANGlutathionereductase,mitochondrial method log fold change t-value q-value df se theta sample 0.900 1.22 msqrob 4.70 22.2 0.26 1.11 1.34 4.82 9.65 0.277 msqrobsum 14.1 NA 22 -21 -20 expression 19 -18 **-**17 p**T**2. pT2. p**T**2. pT2. рТа рТа рТа рТа

## sp|P08758|ANXA5\_HUMANAnnexinA5 log fold change -1.29 method q-value t-value theta sample df se 0.427 msqrob -5.50 83.5 0.235 20.7 -1.27 msqrobsum -4.21 9.66 0.301 NA 24 -22 -20 -18**-**

expression

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sp|P09488|GSTM1\_HUMANGlutathioneS-transferaseMu1 log fold change 1.91 2.08 method q-value t-value theta sample df se 0 9.4 msqrob 8.69 50.2 0.22 msqrobsum 5.94 8.66 0.35 NA 26 -24 expression 22 -20 -18 pT2. pT2. pT2. рТа рТа рТа рТа

sp|P13798|ACPH\_HUMANAcylamino-acid-releasingenzyme log fold change
1.23
1.20 method q-value t-value df theta sample 0.5 msqrob 4.65 33.0 0.265 1.01 4.00 9.66 0.301 msqrobsum NA 21 -20 expression 19**-**18 pT2. pT2. pT2. рТа рТа pT2. рТа рТа

sp|P15924|DESP\_HUMANDesmoplakin method q-value log fold change t-value theta sample df se 2.5 -2.05 msqrob -3.57 155. 0.576 1.93 -2.23 msqrobsum 10.7 -5.23 9.67 0.427 NA 22 -20 expression 18 **-**16-

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sp|P25815|S100P\_HUMANProteinS100-P method log fold change q-value t-value theta sample df se 0.7 1.91 0.614 msqrob 4.49 33.2 0.425 18.9 1.80 4.32 NA msqrobsum 9.68 0.417 26 -24 expression 22 -20 pT2. pT2. p**T**2. pT2. рТа рТа рТа рТа

## $sp|P48735|IDHP\_HUMANI soci trated e hydrogenase [NADP], mitochondrial$ log fold change -1.17 method t-value df theta sample q-value se 0.3 -4.48 96.5 0.693 msqrob 0.261 23.8 -1.15 -3.88 9.67 0.296 msqrobsum NA 24 -22 expression 50 -

18 **-**

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sp|Q5VW32|BROX\_HUMANBRO1domain-containingproteinBROX log fold change 1.46 1.48 method q-value t-value df theta sample 0 14.1 7.99 15.8 5.12 8.63 msqrob 0.108 0.183 8.63 0.290 msqrobsum NA 21 expression 20-19 pT2. pT2. pT2. рТа рТа рТа рТа

## $sp|Q6ZVX7|FBX50\_HUMANF-boxonly protein 50$ log fold change -1.94 method q-value t-value theta sample df se 0 -7.56 14.6 0.257 msqrob -2.32 msqrobsum -5.82 7.65 0.399 NA 22 -20 expression 18-

16 -

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