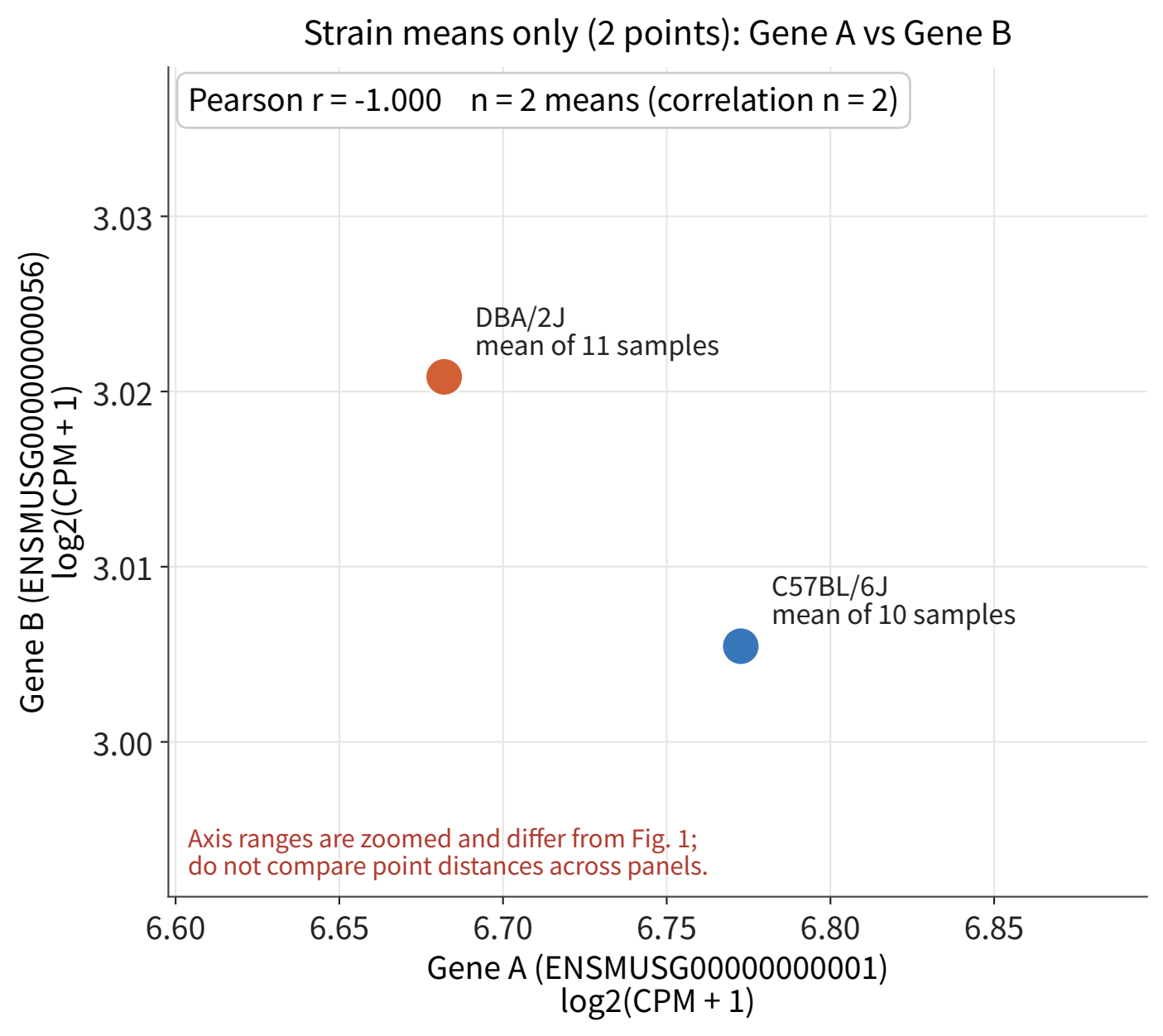

Strain means only (2 points): Gene A vs Gene B
Pearson r = -1.000 n = 2 means (correlation n = 2)
DBA/2J
mean of 11 samples
C57BL/6J
mean of 10 samples
Axis ranges are zoomed and differ from Fig. 1;
do not compare point distances across panels.
Gene B (ENSMUSG00000000056)
log2(CPM + 1)
3.03
3.02
3.01
3.00
6.60
6.65
6.70
6.75
6.80
6.85
Gene A (ENSMUSG00000000001)
log2(CPM + 1)