

Measured Gene Expression from Mixture (RPKM)

**S10 Fig. Linearity assumption test using 32 experimental mixtures.** We tested the linearity assumption employed by csSAM and other deconvolution methods by comparing the measured RPKMs with the reconstituted RPKMs using known mixing proportions and four pure cell lines. *x*-axis represents the measured GEP of mixtures (in RPKM) and the y-axis denotes the RPKMs of the reconstituted mixtures. *R* denotes the correlation between measured mixtures RPKM and the reconstituted RPKM.