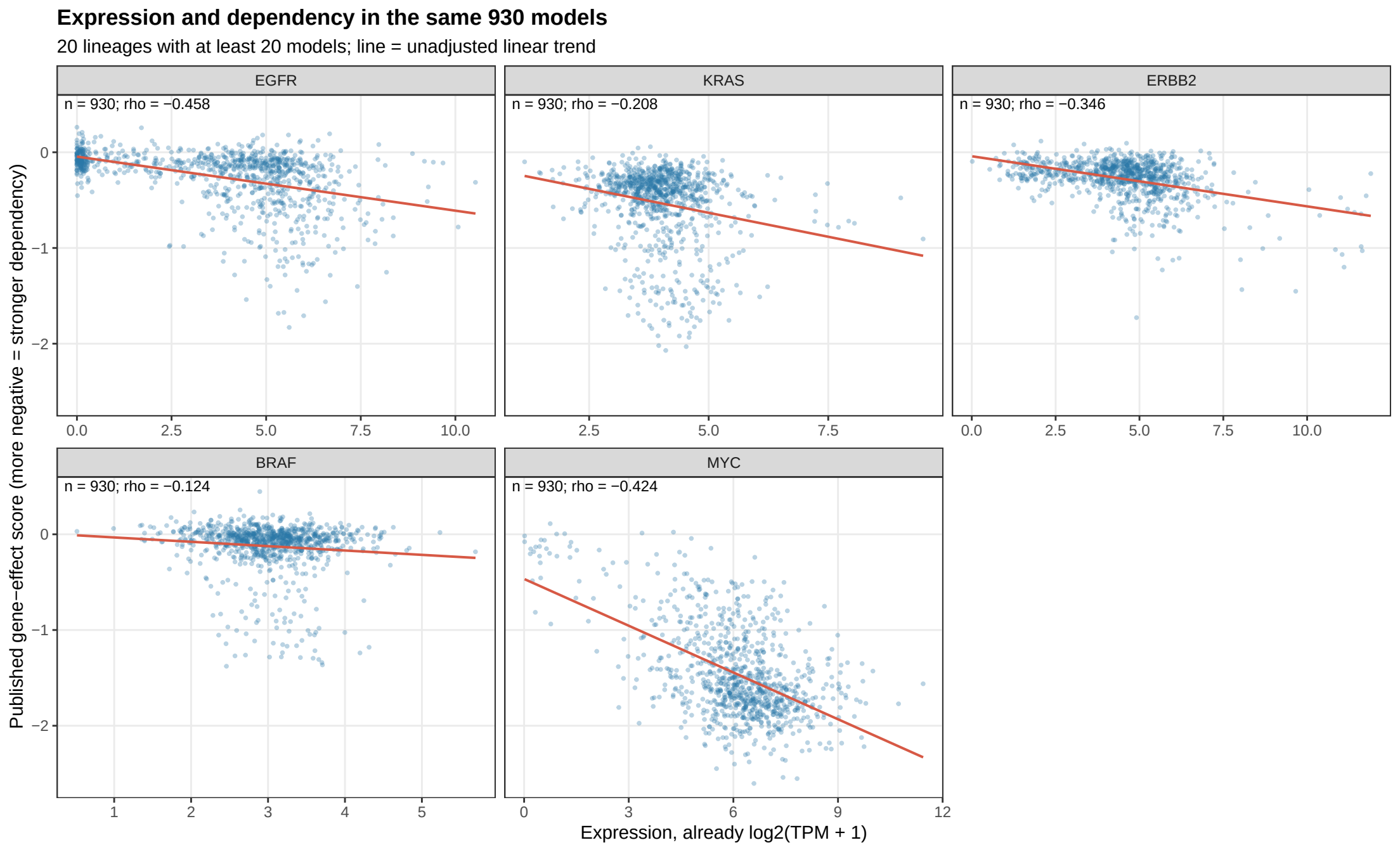
Expression and dependency in the same 930 models
20 lineages with at least 20 models; line = unadjusted linear trend
EGFR
n = 930; rho = −0.458
KRAS
n = 930; rho = −0.208
ERBB2
n = 930; rho = −0.346
BRAF
n = 930; rho = −0.124
MYC
n = 930; rho = −0.424
Published gene−effect score (more negative = stronger dependency)
Expression, already log2(TPM + 1)