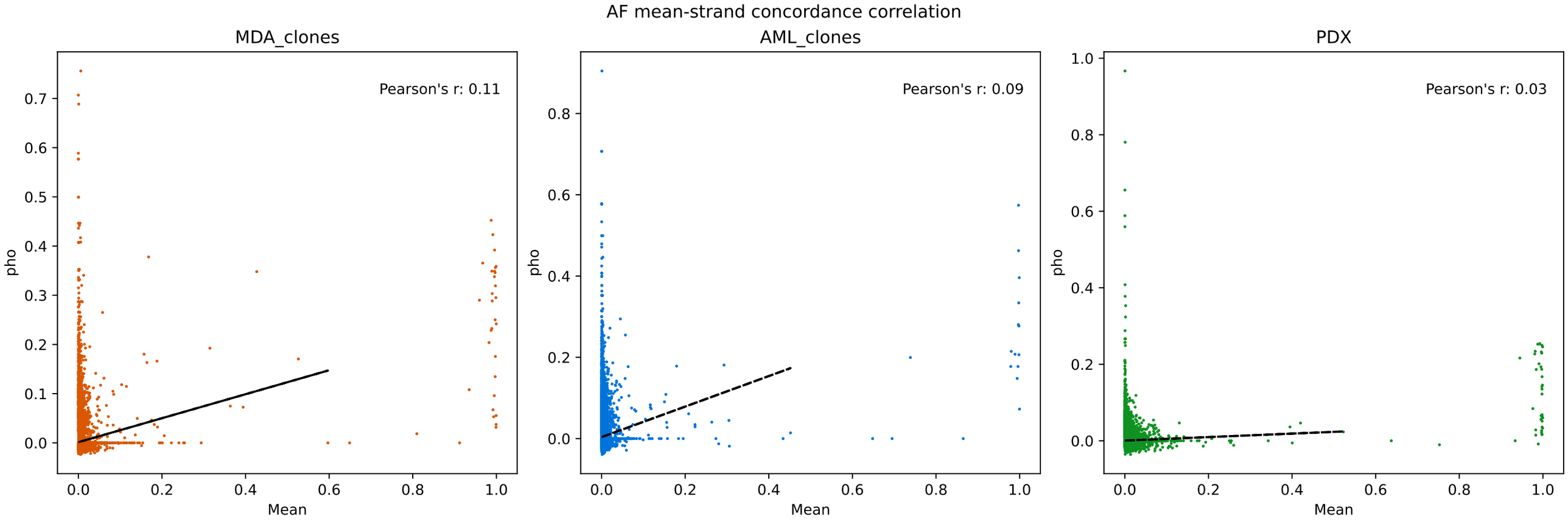
AF mean-strand concordance correlation
MDA_clones
Pearson's r: 0.11
AML_clones
Pearson's r: 0.09
PDX
Pearson's r: 0.03
pho
Mean
0.0
0.1
0.2
0.3
0.4
0.5
0.6
0.7
0.8
1.0