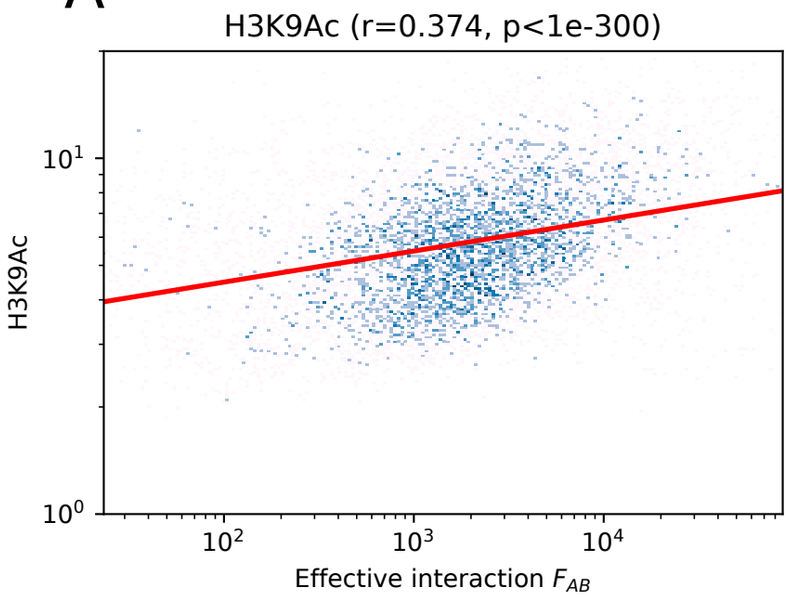
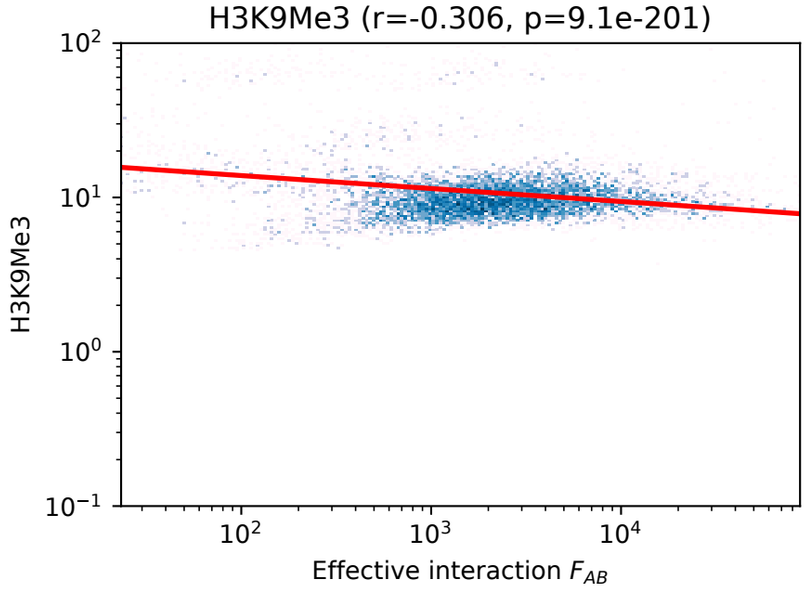
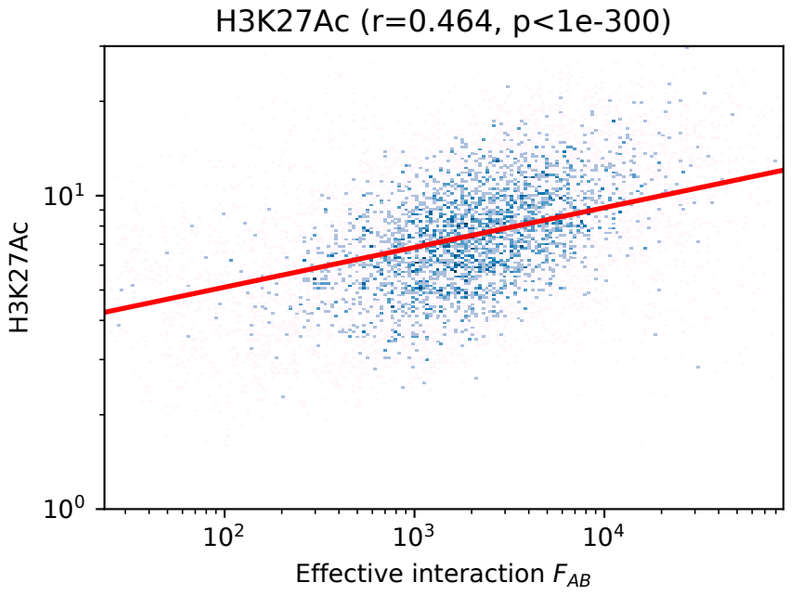
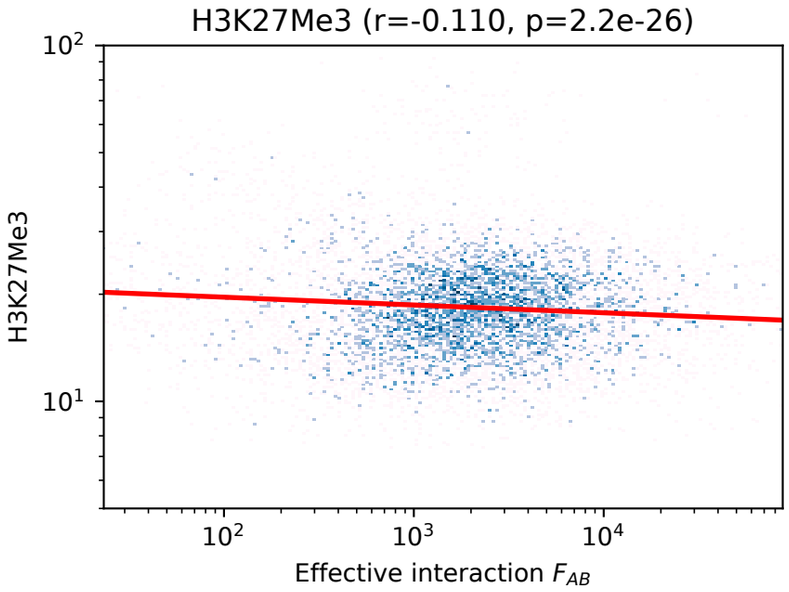
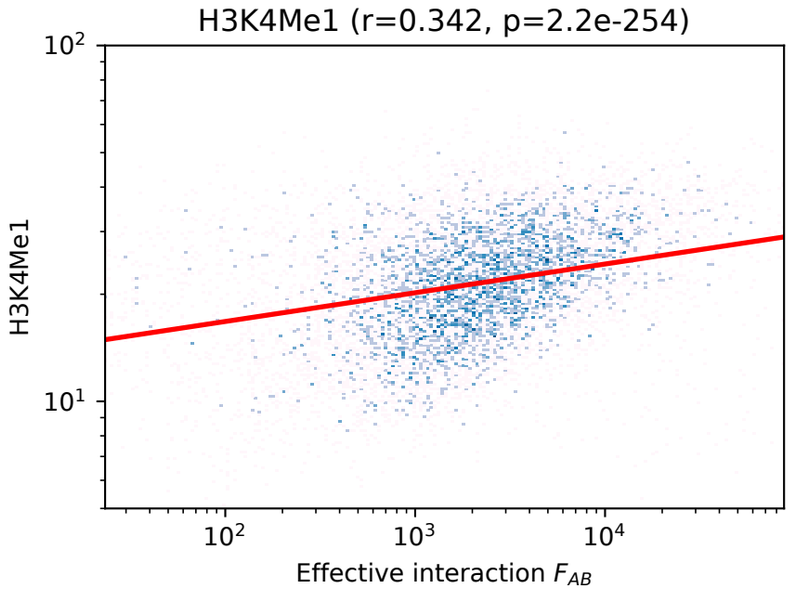
A

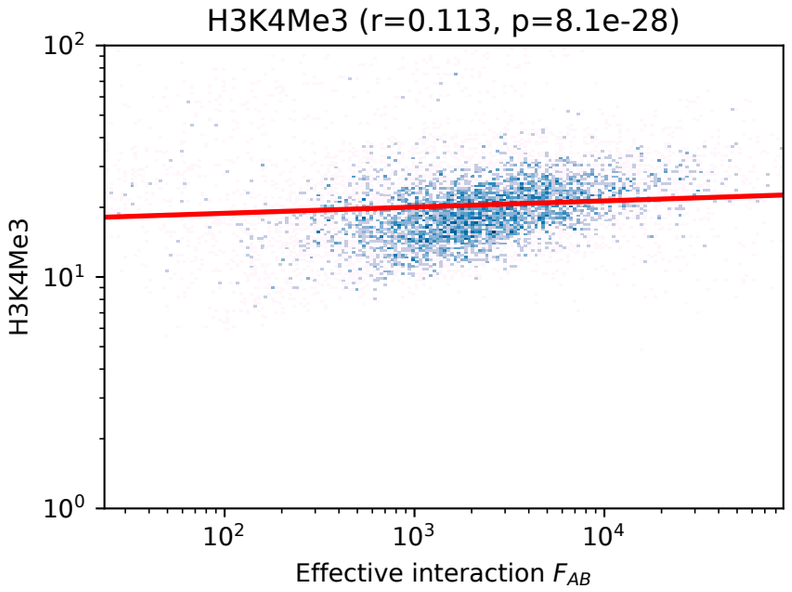

H3K9Ac (r=0.374, p<1e-300)
H3K9Ac
Effective interaction F_{AB}

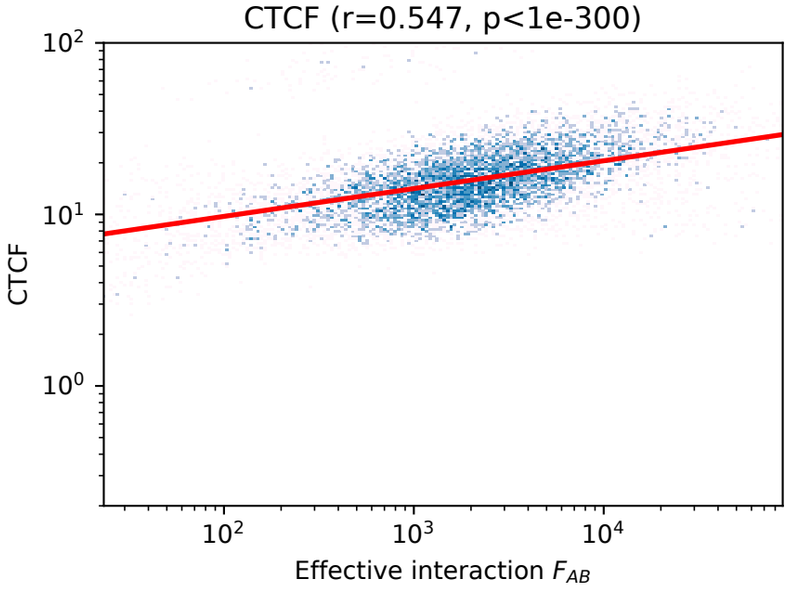

H3K9Me3 (r=-0.306, p=9.1e-201)
H3K9Me3
Effective interaction F_{AB}

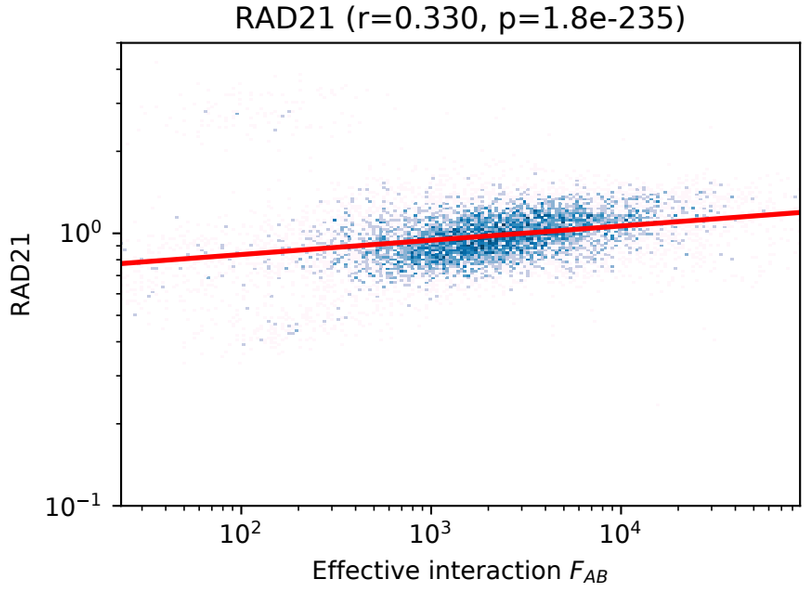

H3K27Ac (r=0.464, p<1e-300)
H3K27Ac
Effective interaction F_{AB}

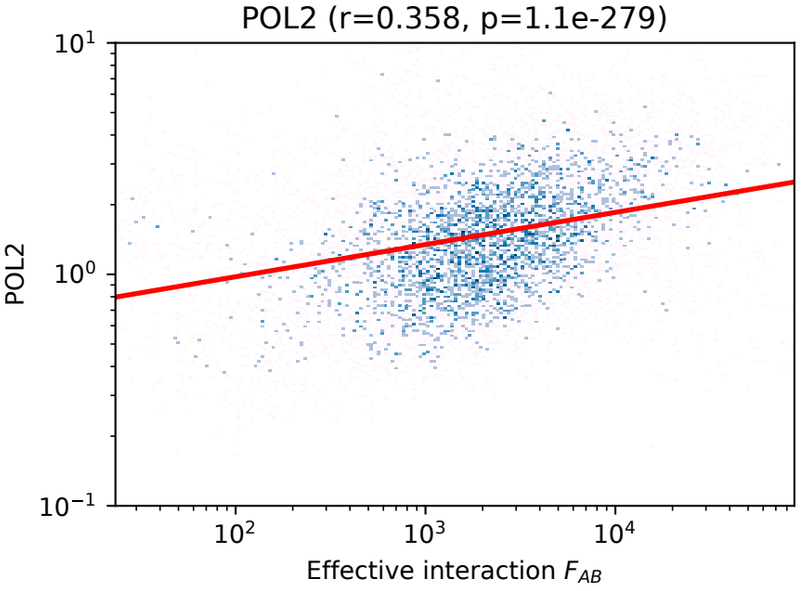

H3K27Me3 (r=-0.110, p=2.2e-26)
H3K27Me3
Effective interaction F_{AB}

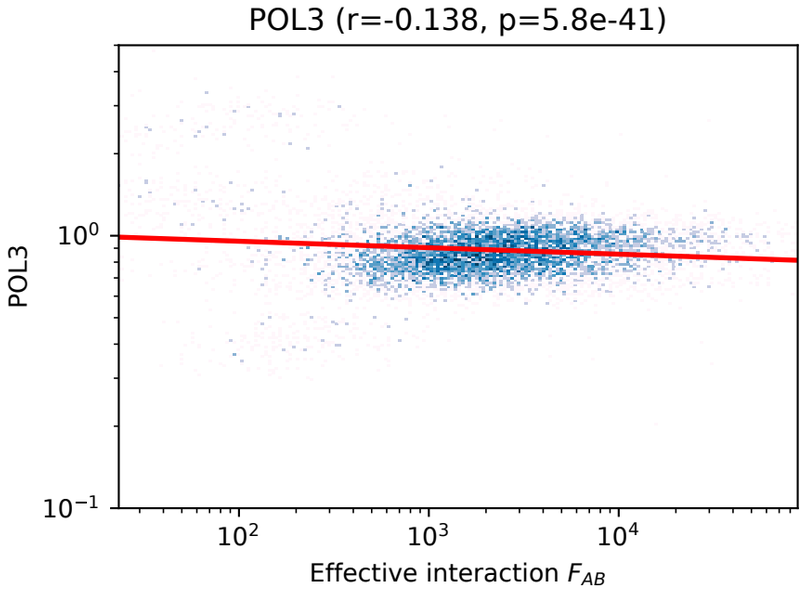

H3K4Me1 (r=0.342, p=2.2e-254)
H3K4Me1
Effective interaction F_{AB}

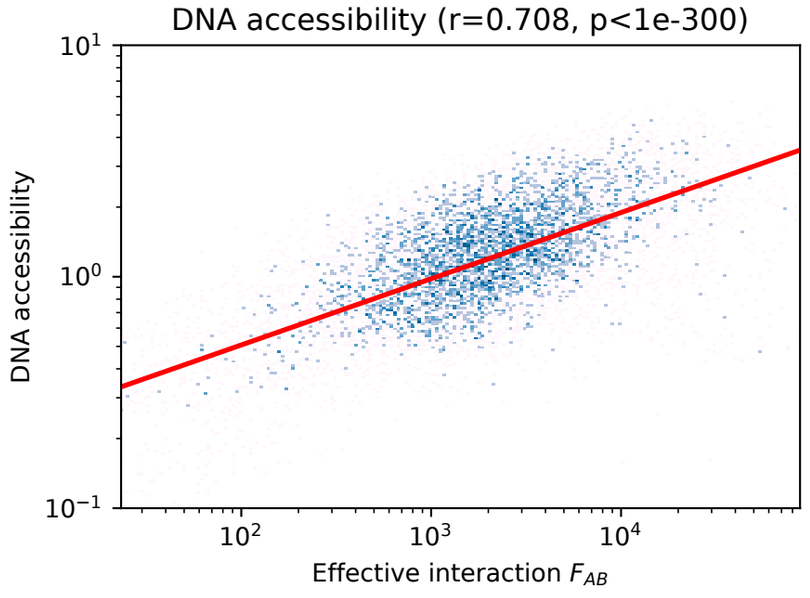

H3K4Me3 (r=0.113, p=8.1e-28)
H3K4Me3
Effective interaction F_{AB}


CTCF (r=0.547, p<1e-300)
CTCF
Effective interaction F_{AB}

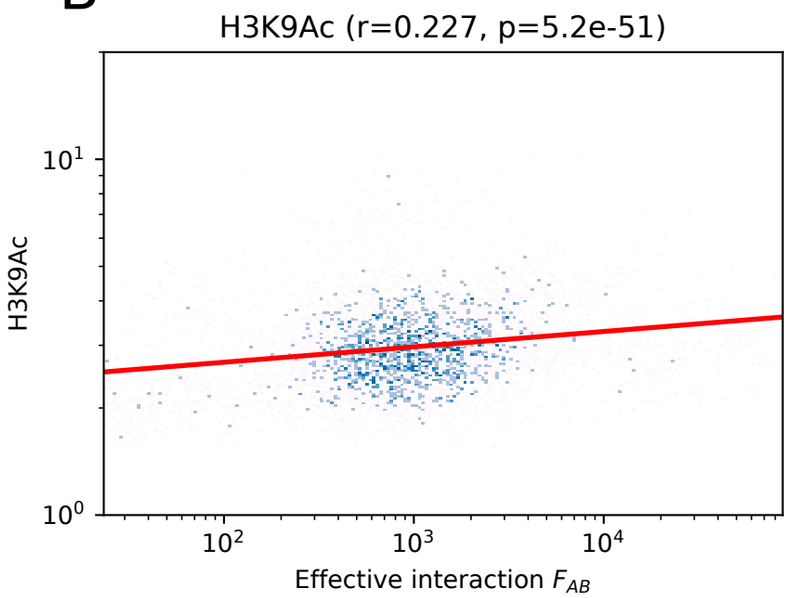

RAD21 (r=0.330, p=1.8e-235)
RAD21
Effective interaction F_{AB}

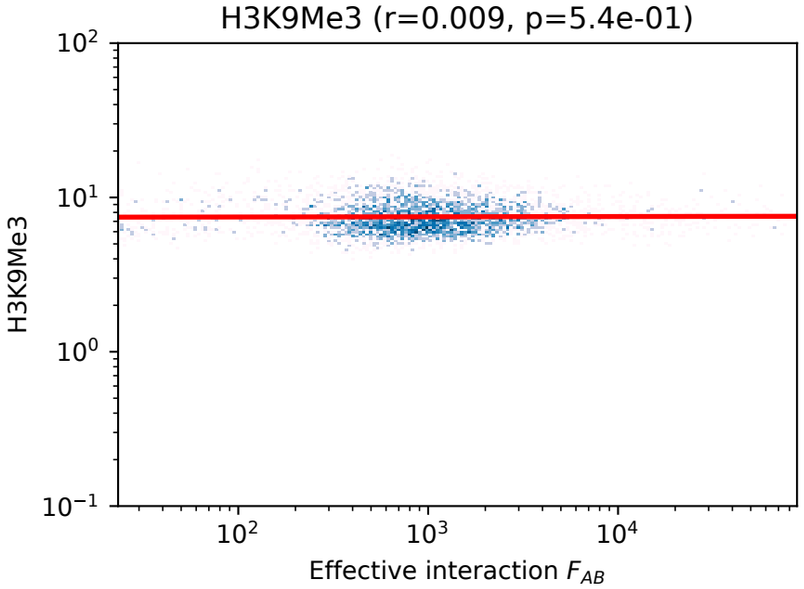

POL2 (r=0.358, p=1.1e-279)
POL2
Effective interaction F_{AB}

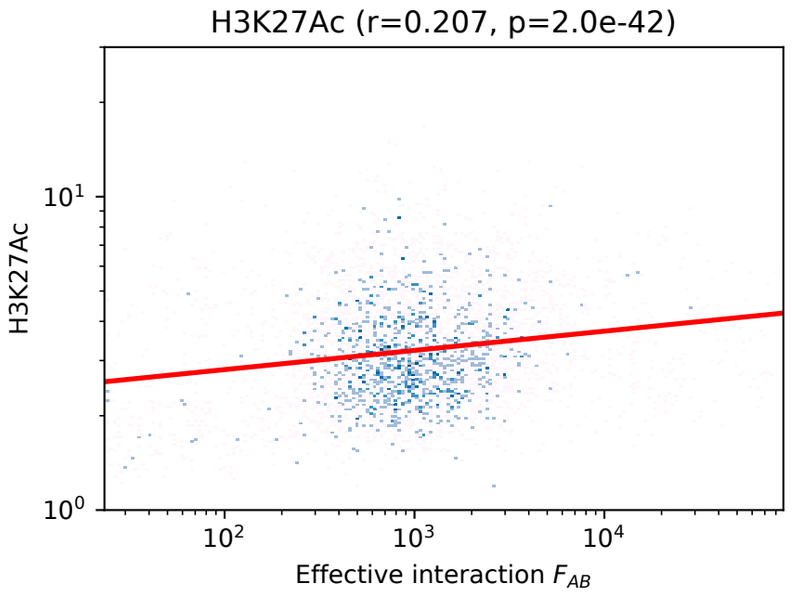

POL3 (r=-0.138, p=5.8e-41)
POL3
Effective interaction F_{AB}

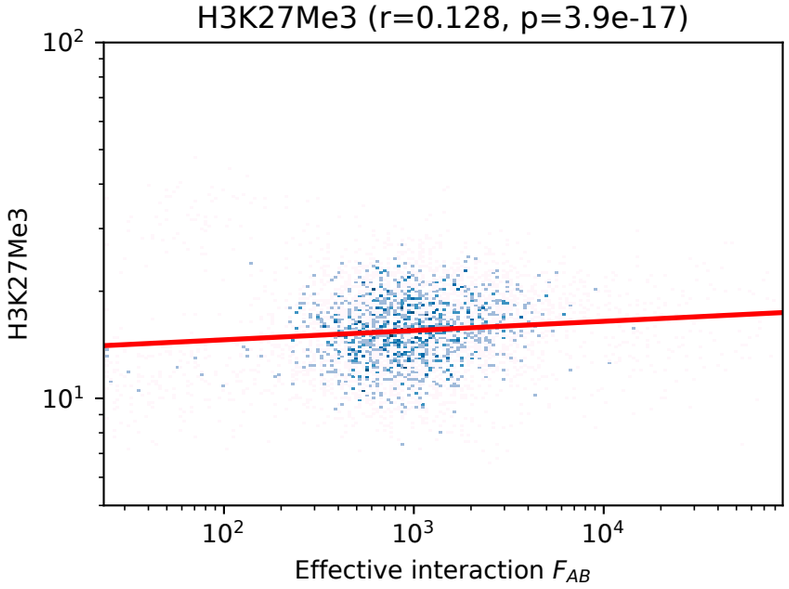

DNA accessibility (r=0.708, p<1e-300)
DNA accessibility
Effective interaction F_{AB}

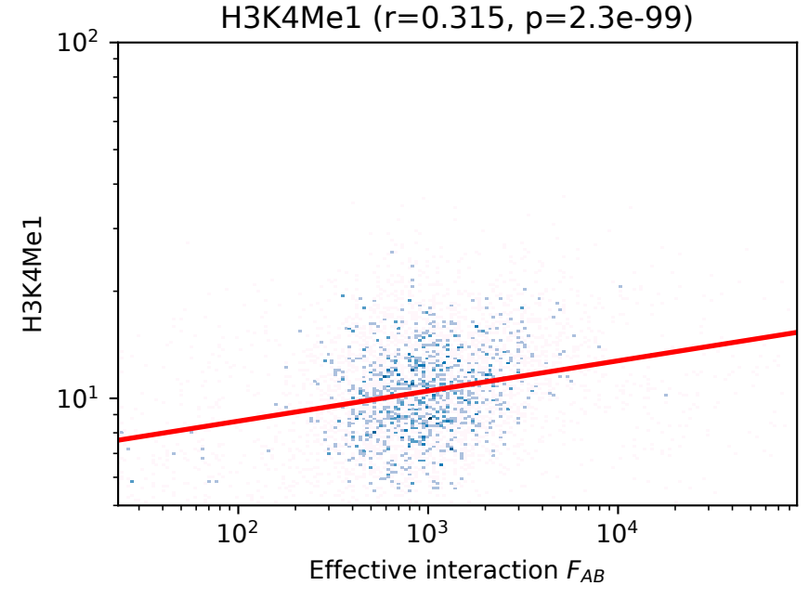
B

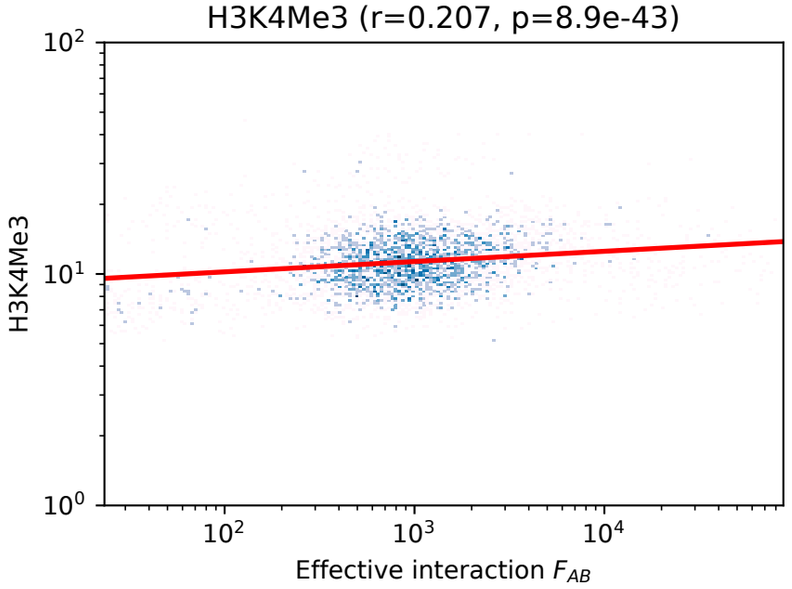

H3K9Ac (r=0.227, p=5.2e-51)
H3K9Ac
Effective interaction F_{AB}

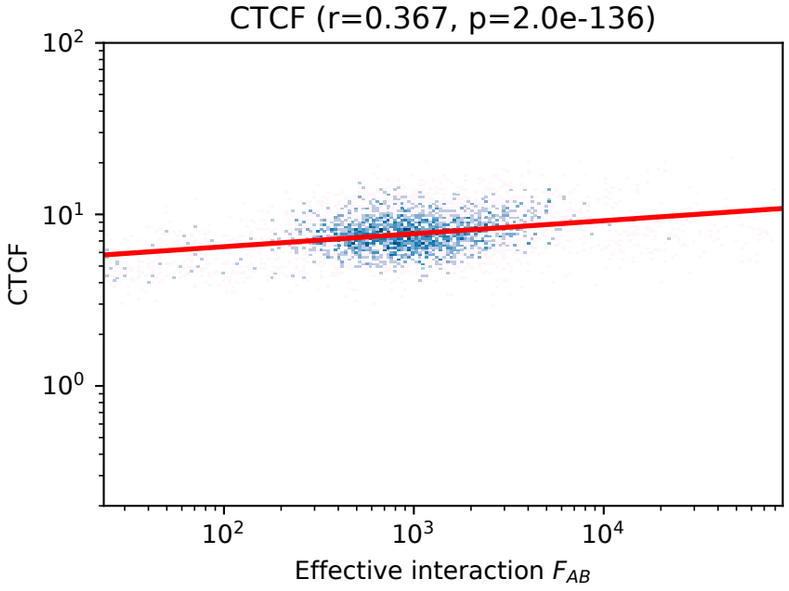

H3K9Me3 (r=0.009, p=5.4e-01)
H3K9Me3
Effective interaction F_{AB}

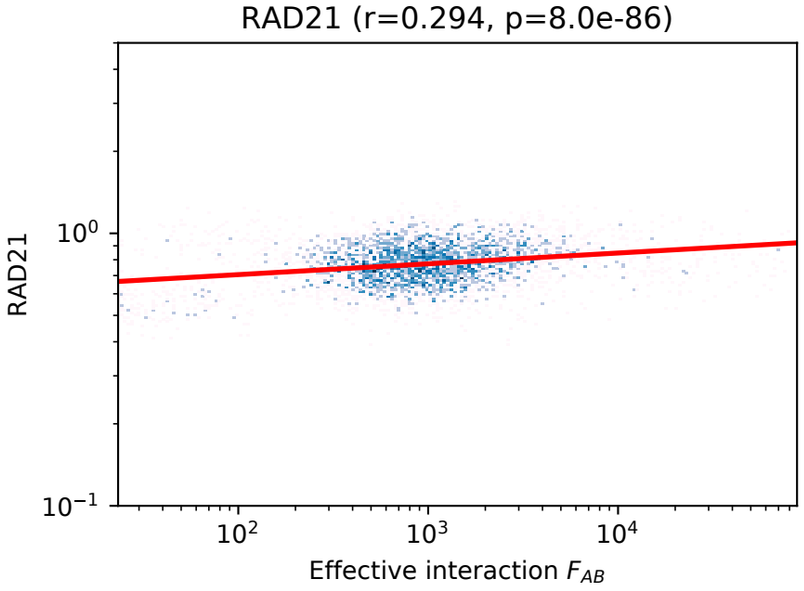

H3K27Ac (r=0.207, p=2.0e-42)
H3K27Ac
Effective interaction F_{AB}

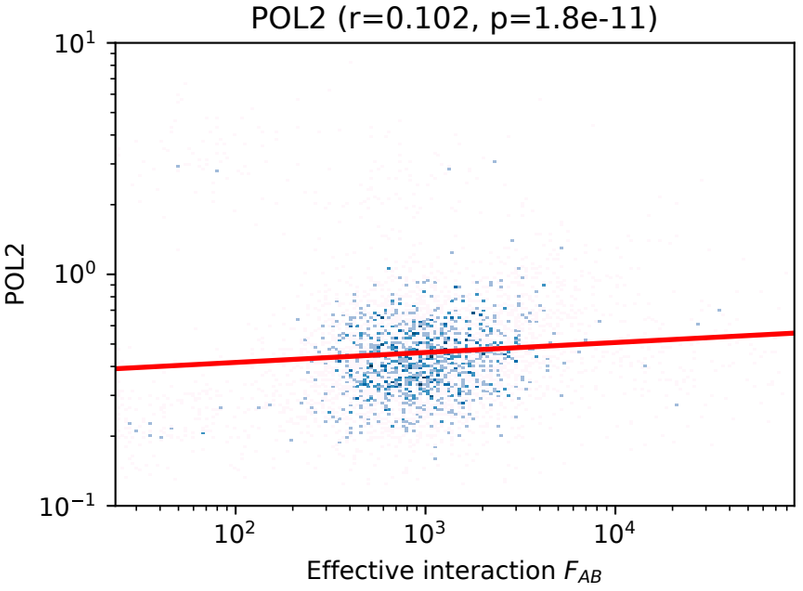

H3K27Me3 (r=0.128, p=3.9e-17)
H3K27Me3
Effective interaction F_{AB}

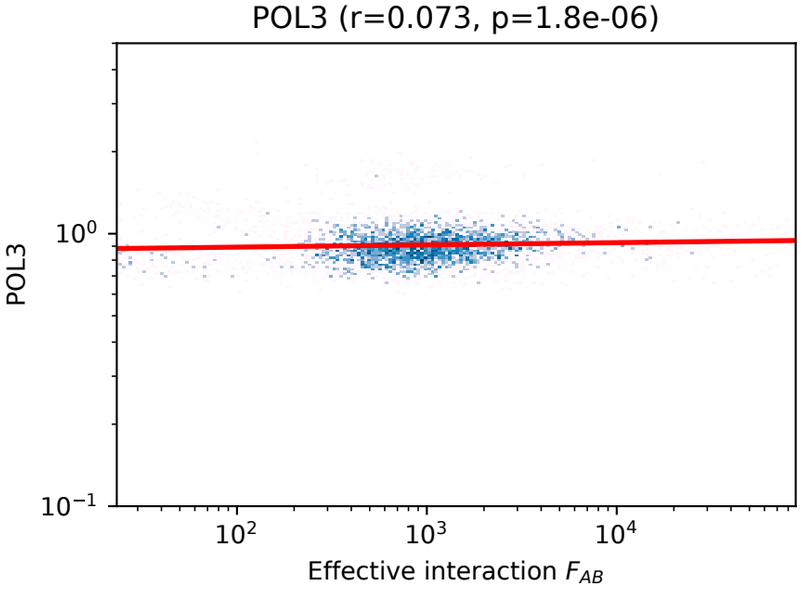

H3K4Me1 (r=0.315, p=2.3e-99)
H3K4Me1
Effective interaction F_{AB}

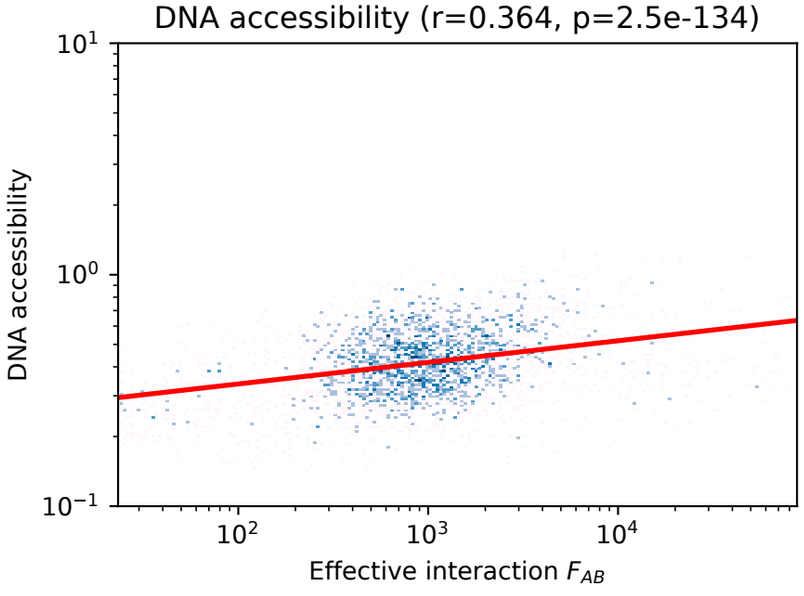

H3K4Me3 (r=0.207, p=8.9e-43)
H3K4Me3
Effective interaction F_{AB}


CTCF (r=0.367, p=2.0e-136)
CTCF
Effective interaction F_{AB}

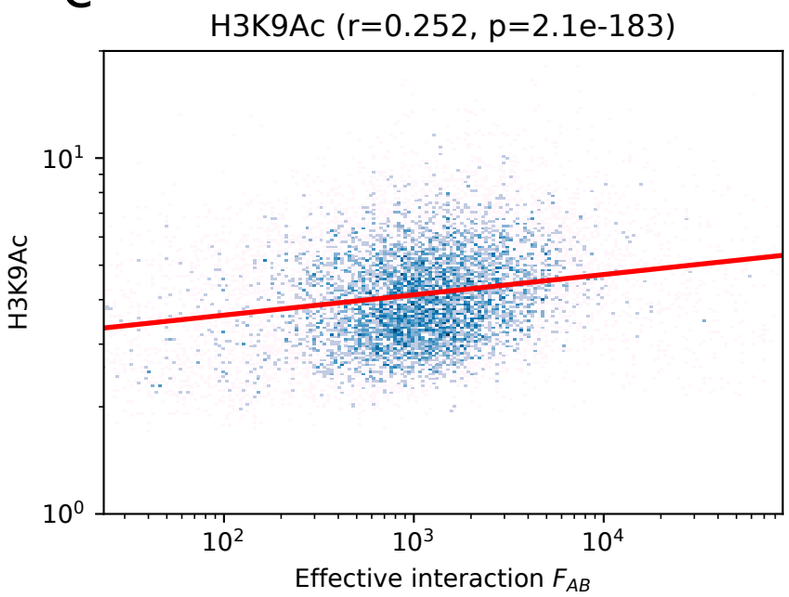

RAD21 (r=0.294, p=8.0e-86)
RAD21
Effective interaction F_{AB}

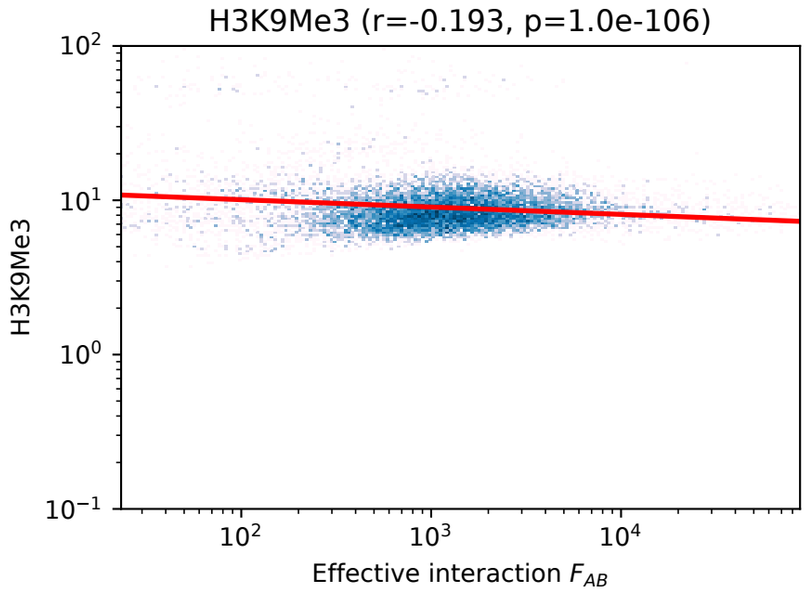

POL2 (r=0.102, p=1.8e-11)
POL2
Effective interaction F_{AB}

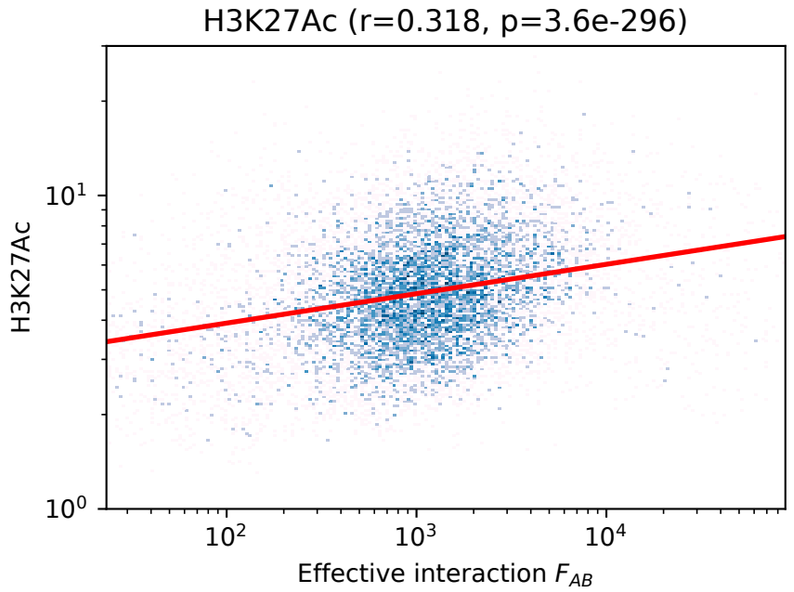

POL3 (r=0.073, p=1.8e-06)
POL3
Effective interaction F_{AB}

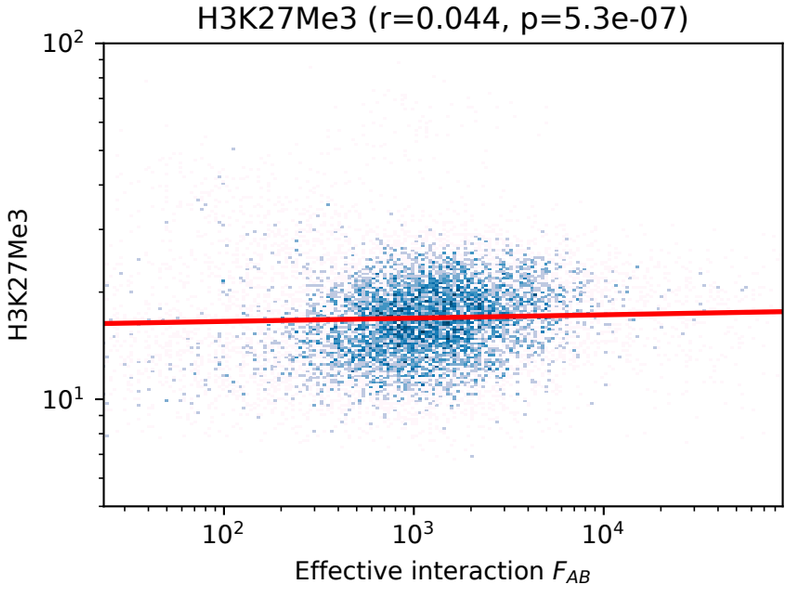

DNA accessibility (r=0.364, p=2.5e-134)
DNA accessibility
Effective interaction F_{AB}

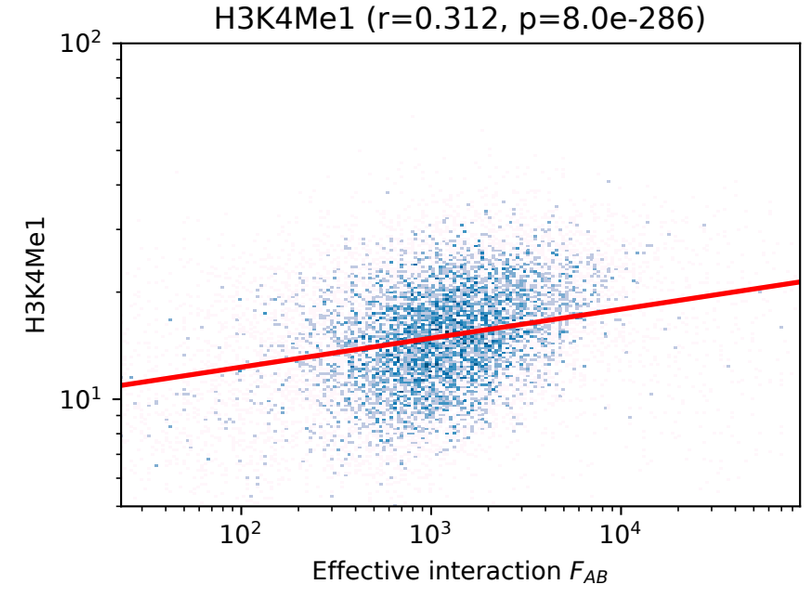
C

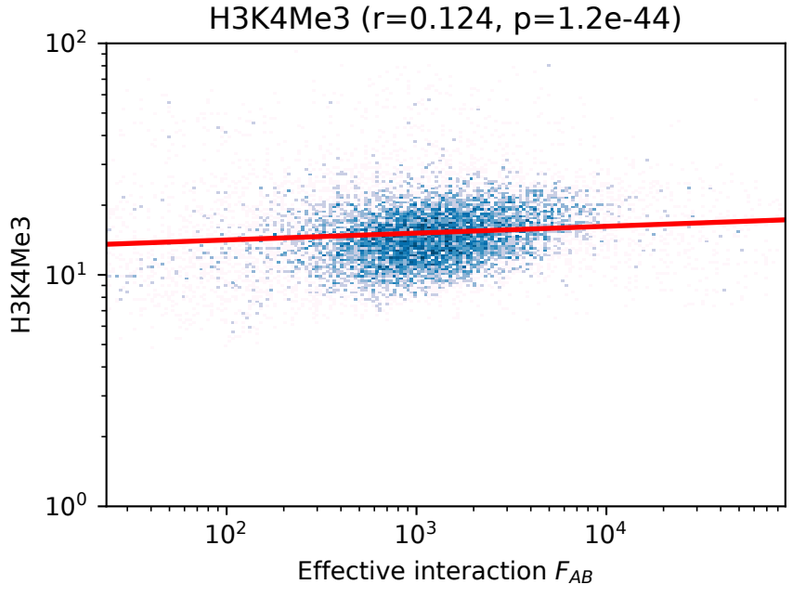

H3K9Ac (r=0.252, p=2.1e-183)
H3K9Ac
Effective interaction F_{AB}

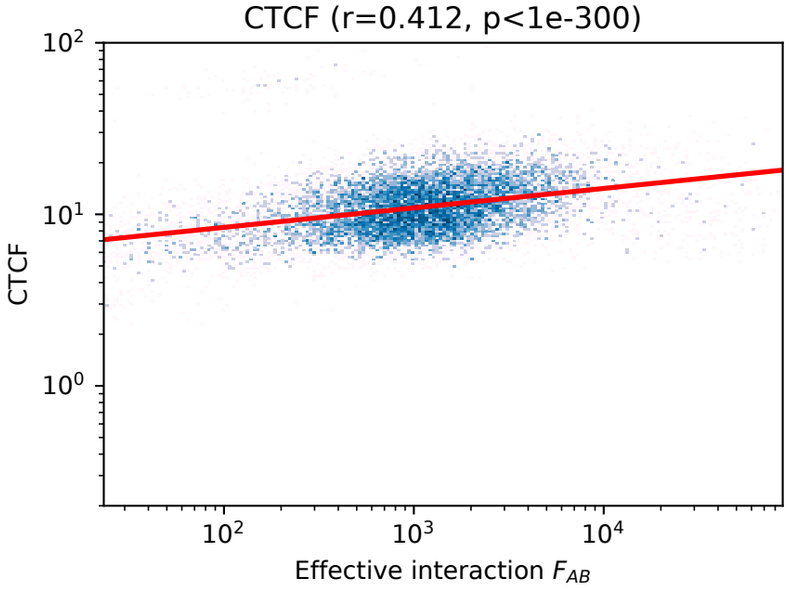

H3K9Me3 (r=-0.193, p=1.0e-106)
H3K9Me3
Effective interaction F_{AB}

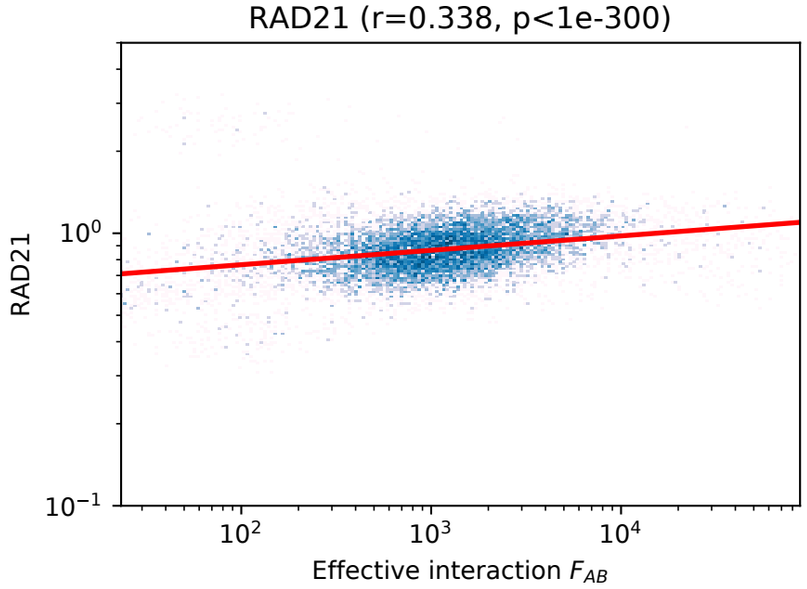

H3K27Ac (r=0.318, p=3.6e-296)
H3K27Ac
Effective interaction F_{AB}

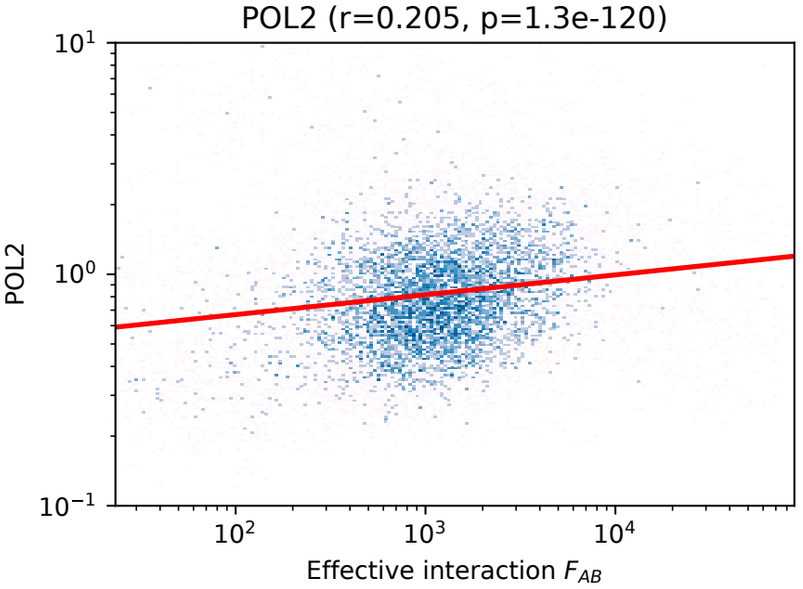

H3K27Me3 (r=0.044, p=5.3e-07)
H3K27Me3
Effective interaction F_{AB}

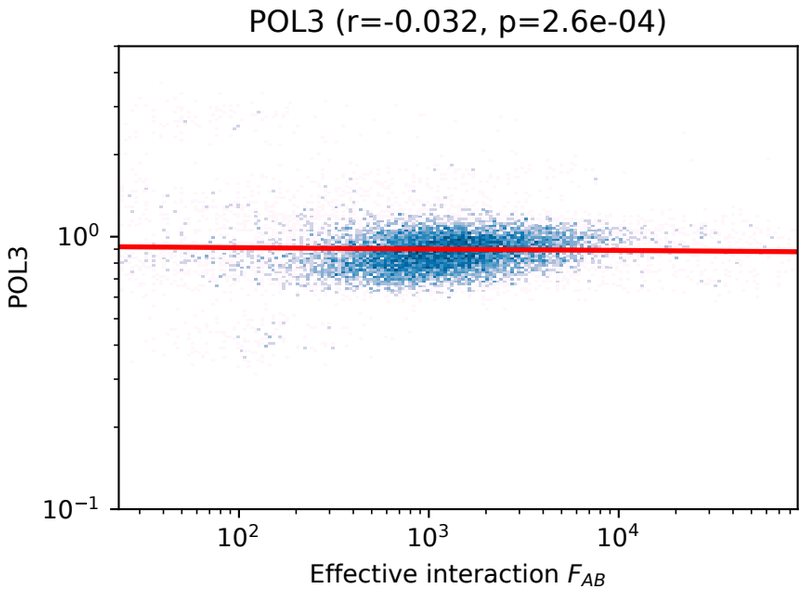

H3K4Me1 (r=0.312, p=8.0e-286)
H3K4Me1
Effective interaction F_{AB}

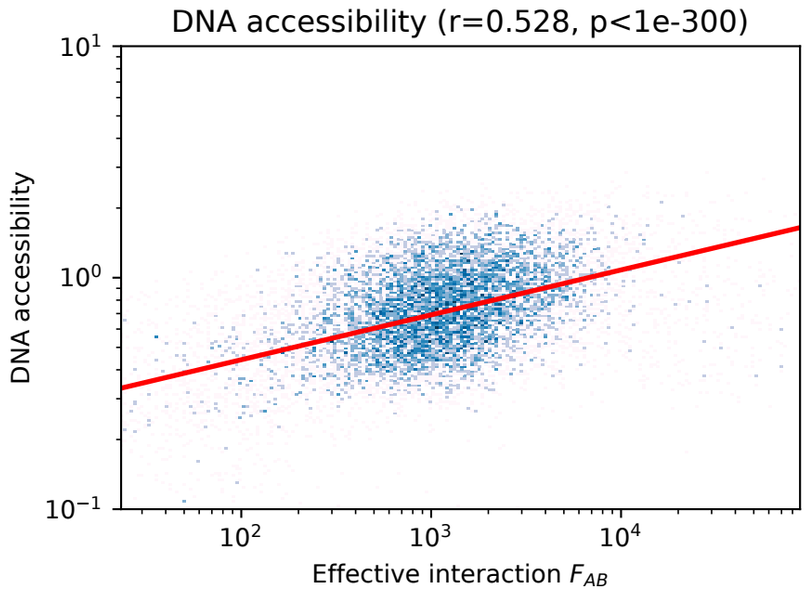

H3K4Me3 (r=0.124, p=1.2e-44)
H3K4Me3
Effective interaction F_{AB}


CTCF (r=0.412, p<1e-300)
CTCF
Effective interaction F_{AB}


RAD21 (r=0.338, p<1e-300)
RAD21
Effective interaction F_{AB}


POL2 (r=0.205, p=1.3e-120)
POL2
Effective interaction F_{AB}


POL3 (r=-0.032, p=2.6e-04)
POL3
Effective interaction F_{AB}


DNA accessibility (r=0.528, p<1e-300)
DNA accessibility
Effective interaction F_{AB}