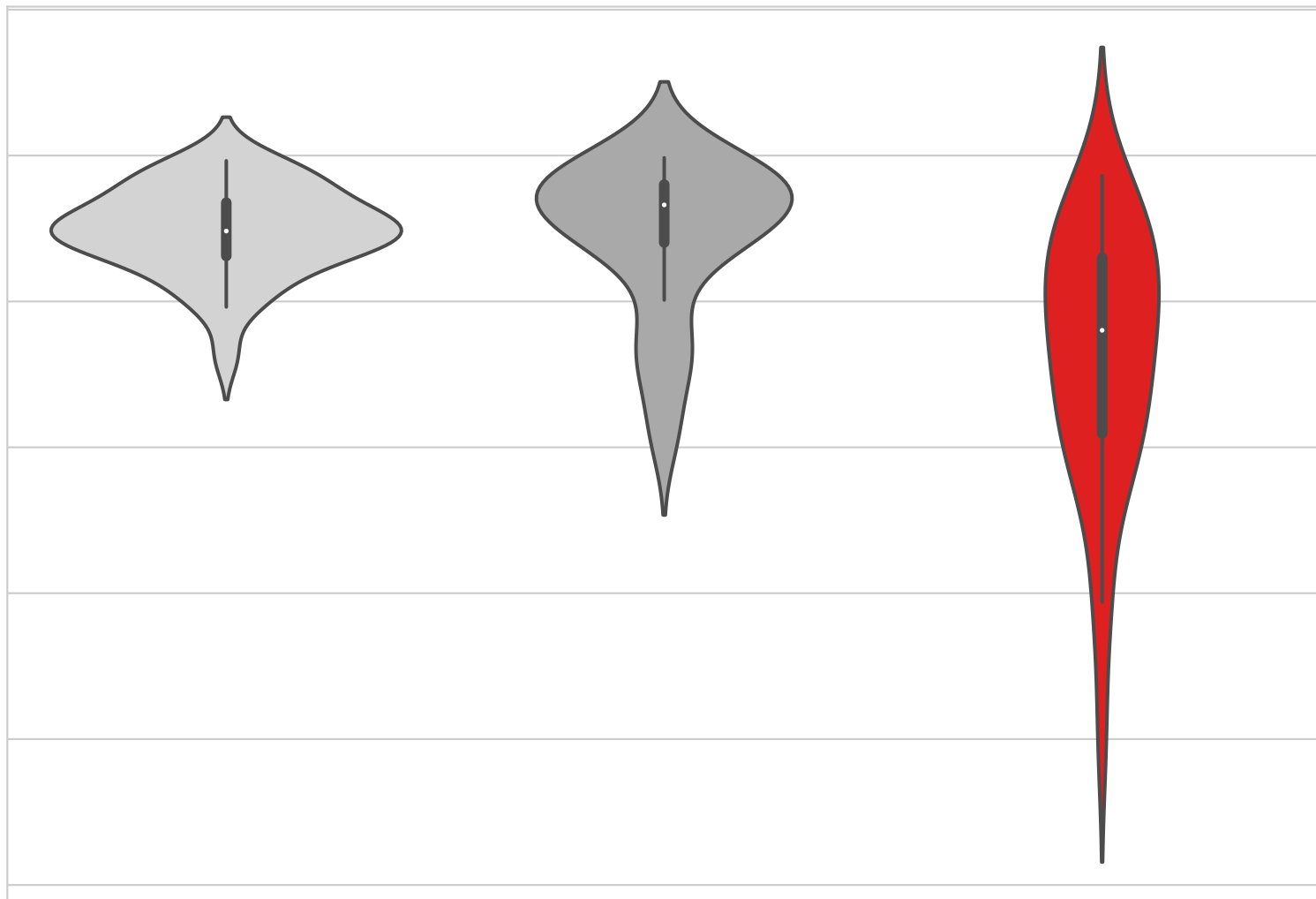


r2

1.1
1.0
0.9
0.8
0.7
0.6
0.5



LPMC

DATGAN

ciDATGAN