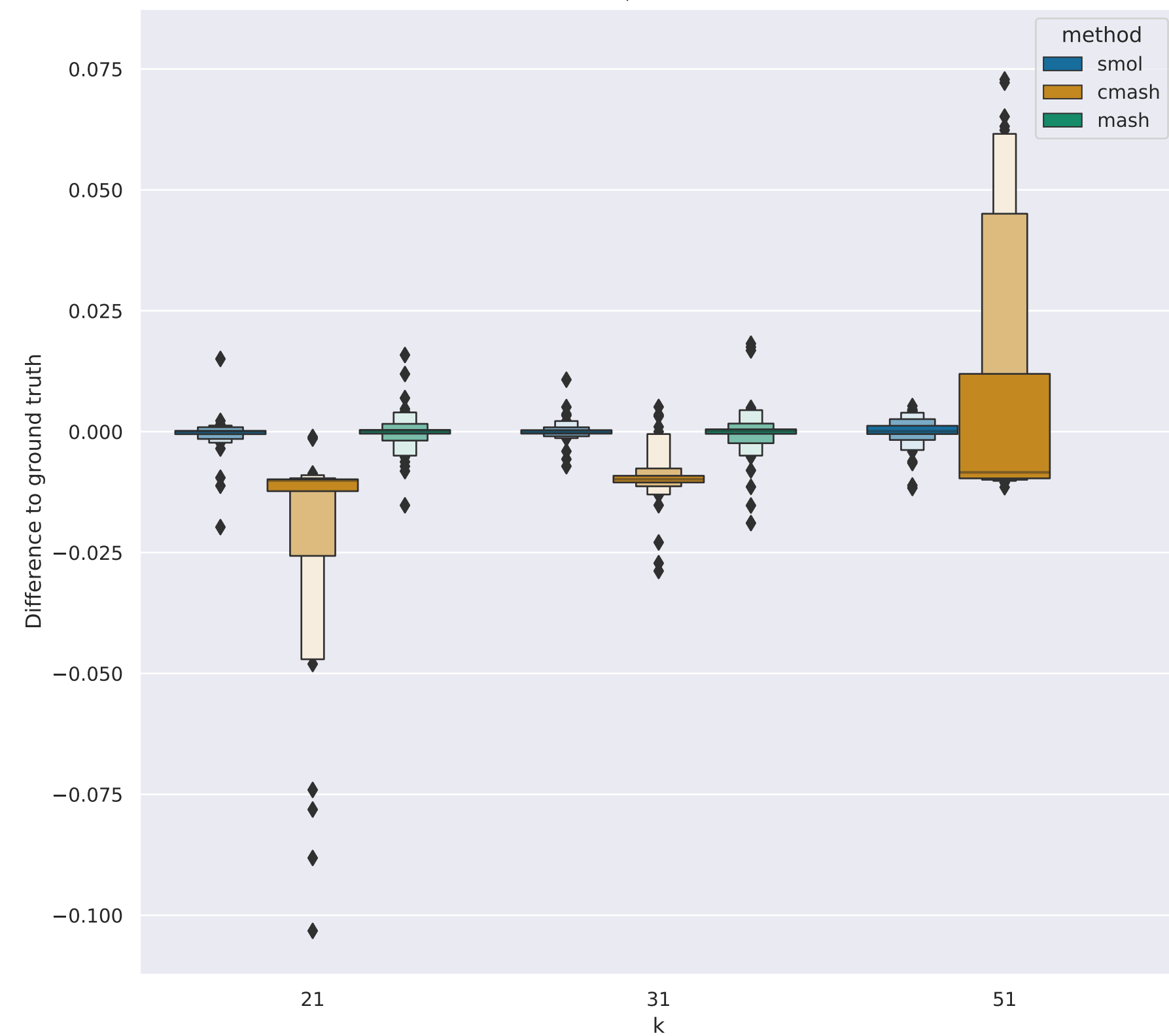


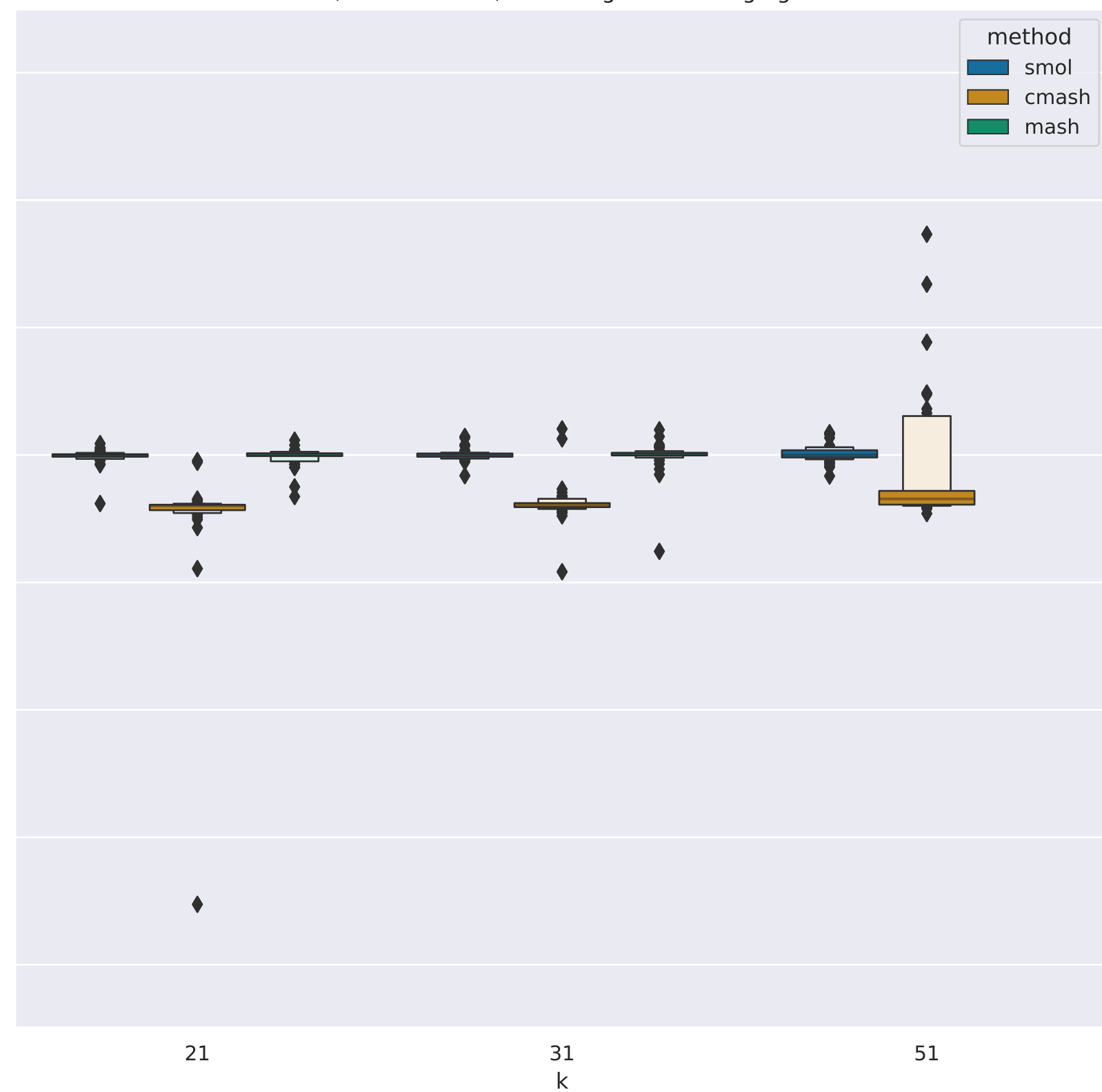
A

n=1000, scaled=1000



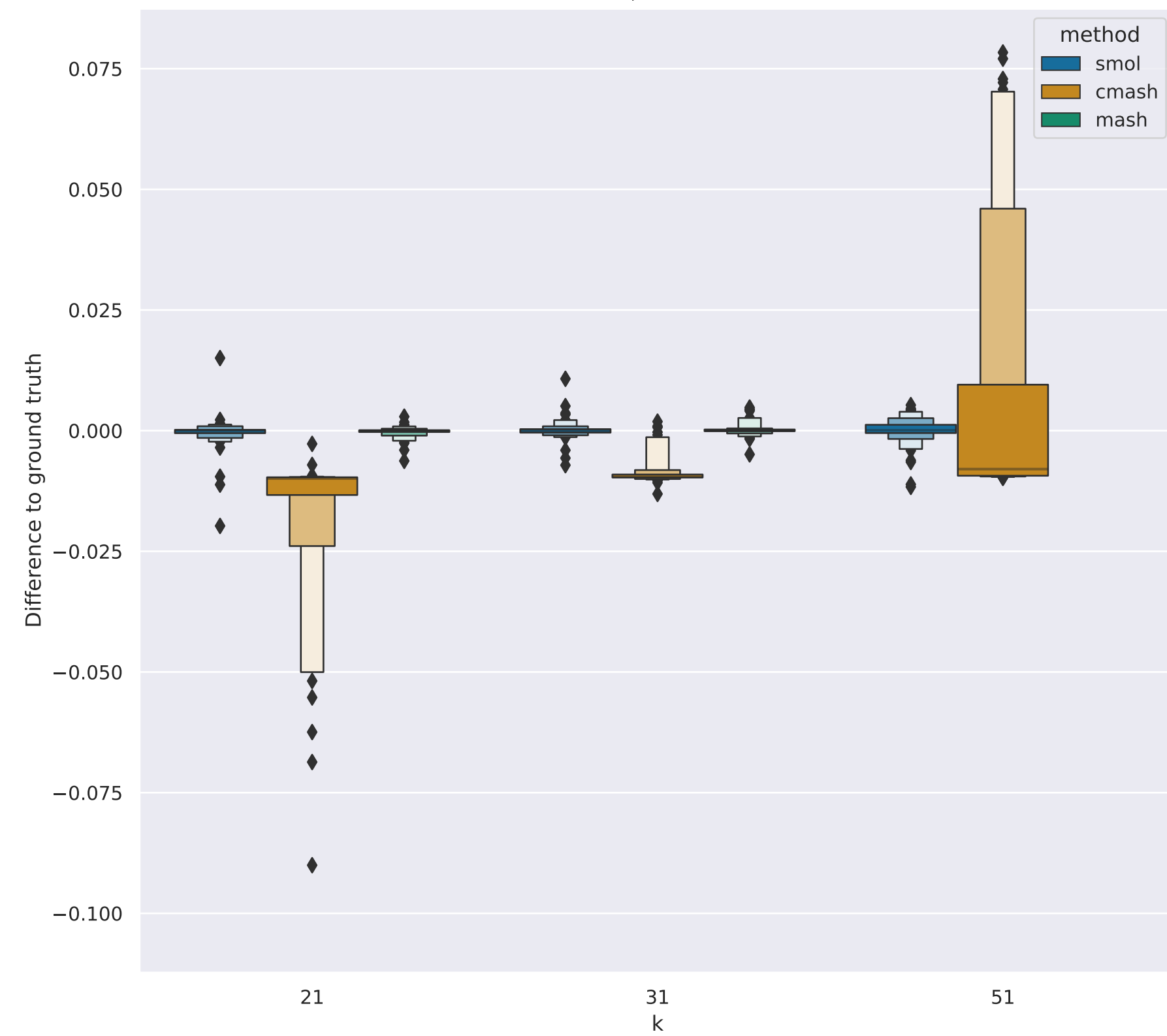
B

n=1000, scaled=1000, excluding low coverage genomes



C

n=10000, scaled=1000



D

n=10000, scaled=1000, excluding low coverage genomes

