



Figure S1. Genome comparison between China/GD/2019 and China/2018/AnhuiXCGQ: A deletion of approximately 1kb was found in China/GD/2019 genome which located at EP153R and EP402R genes in comparison to China/2018/AnhuiXCGQ strain. (A) Collinearity analysis of China/GD/2019 with China/2018/AnhuiXCGQ. (B) Comparative genomic analysis of China/GD/2019 and China/2018/AnhuiXCGQ.