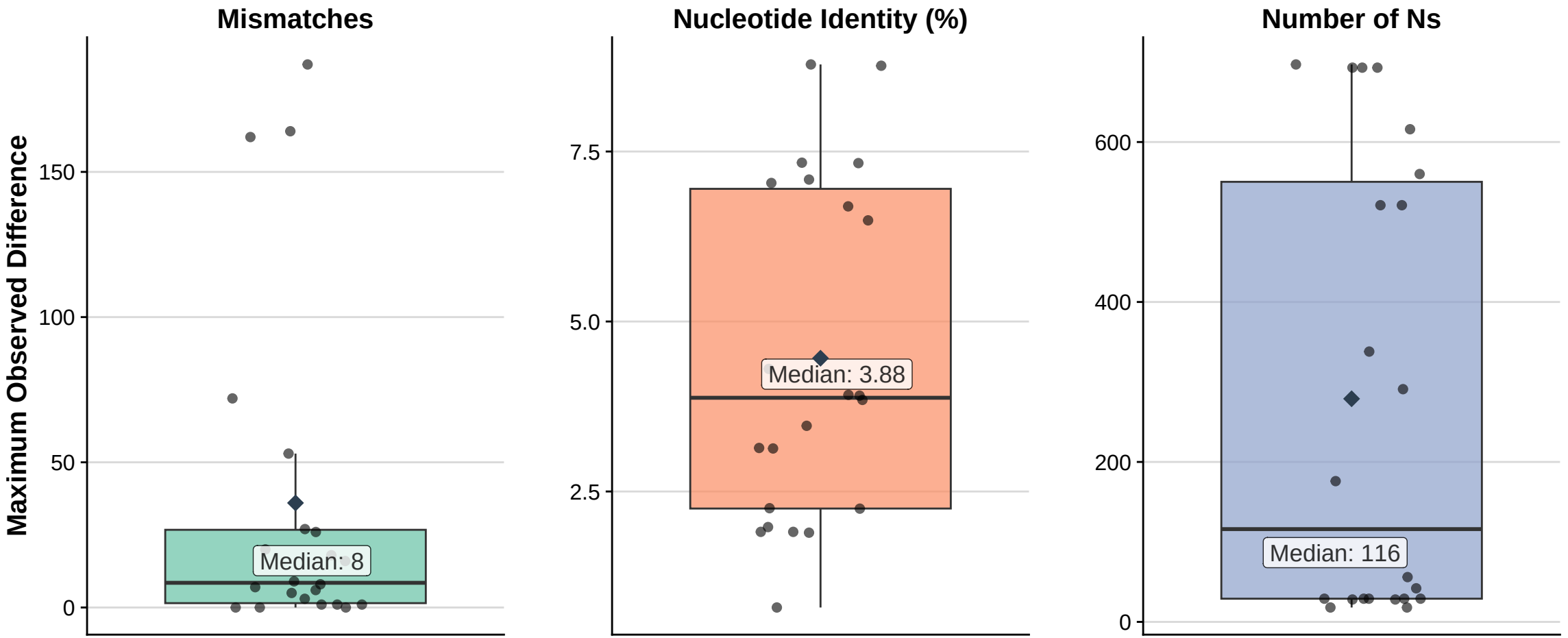Impact of Reference Genome on Consensus Quality