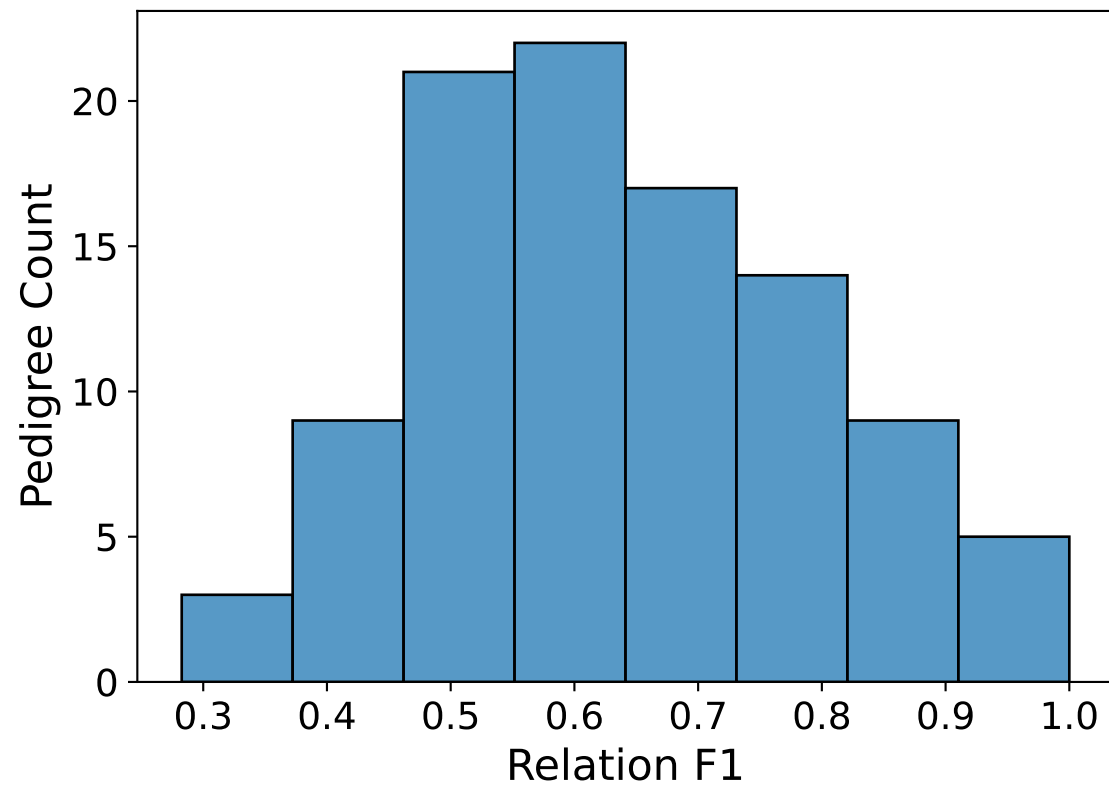Pedigree-Level Reconstruction Performance Distribution
p(Mask Node)=0.4, Simulated Sequence Coverage=0.1x