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1 2Department Of Molecular Biosciences,

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Et Al. 2008, Conrad Et Al. 2010, Cridland Et Al. 2013, Rogers Et Al. 2015). Accurate And Gap-free Genomes Provide A Direct And Reliable Path To Comprehensive Identification Of SVs (Alkan Et Al. 2011, Huddleston Et Al. 2017, Chakraborty Et Al. 2018). To Achieve This Goal, We Assembled Reference-quality Genomes For Fourteen Geographically Diverse 3th, 2024

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