FISH: fast and accurate diploid genotype imputation via segmental hidden Markov model

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The author would like to note an affiliation correction. The first affiliation address was mistakenly stated as ‘School of Public Health, Xi’an Jiaotong University, Shaanxi, China’, while the correct one should be ‘Center for Genetic Epidemiology and Genomics, School of Public Health, Soochow University, Jiangsu, P. R. China’.