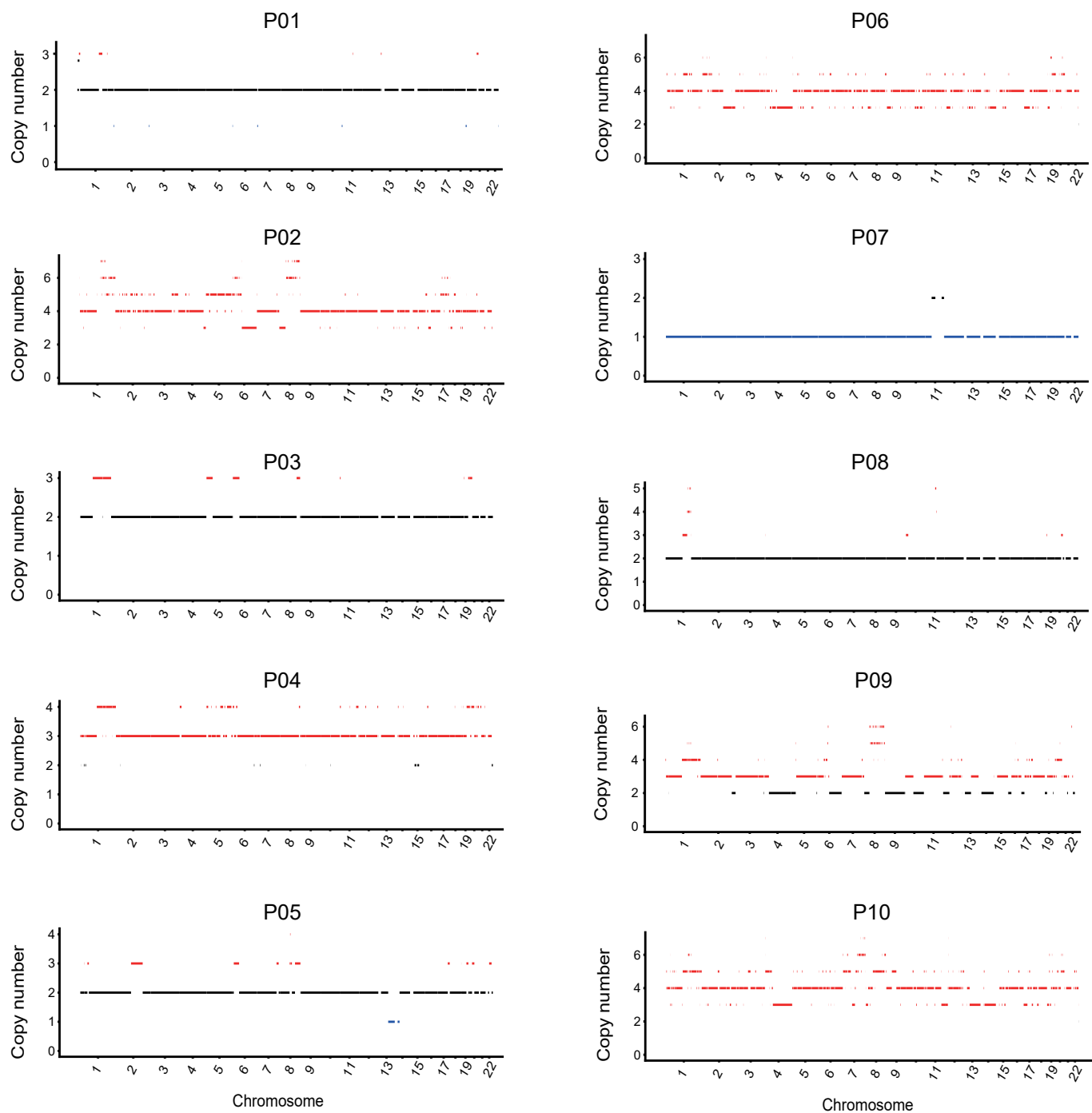


Figure S8



**Figure S8. CNA profiles of bulk tissue samples for each corresponding single-cell samples.**

Based on the raw data of single cells of each sample, the corresponding bulk tissue data is generated, and then the copy number is extracted. Black lines represent normal genome, blue lines represent deletions, and red lines represent amplifications.