

Figure S1. Phylogenetic trees constructed by PhyML 3.0 software (GTR model) based on concatenated sequences of the *Rickettsia*, *Ehrlichia*, and *Anaplasma* strains

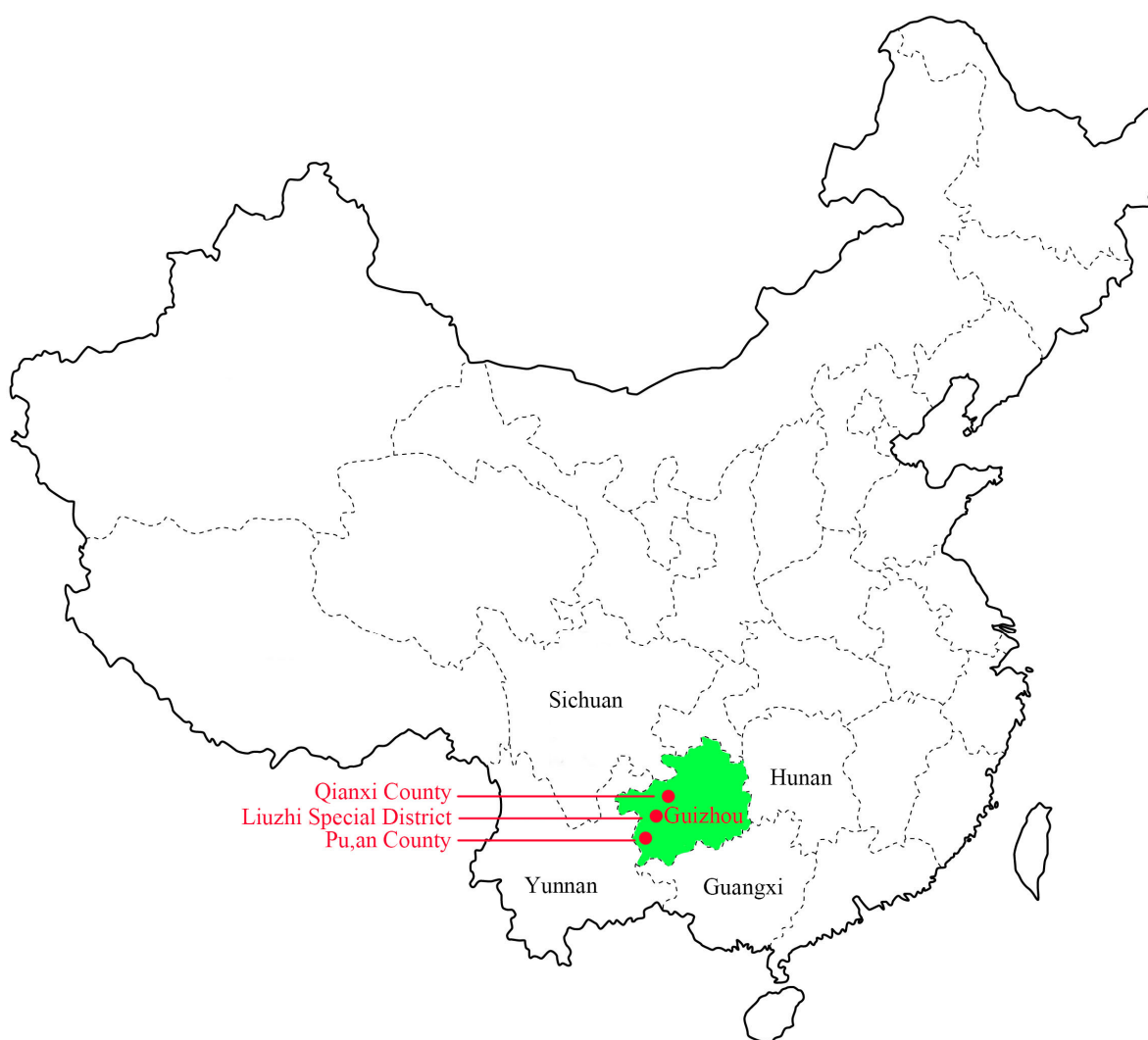


Figure S2. Map showing the locations where the samples were collected: Puan County of Qianxinan Bouyei-and-Miao Autonomous Prefecture; Liuzhi Special District of Liupanshui City; and Qianxi County of Bijie City