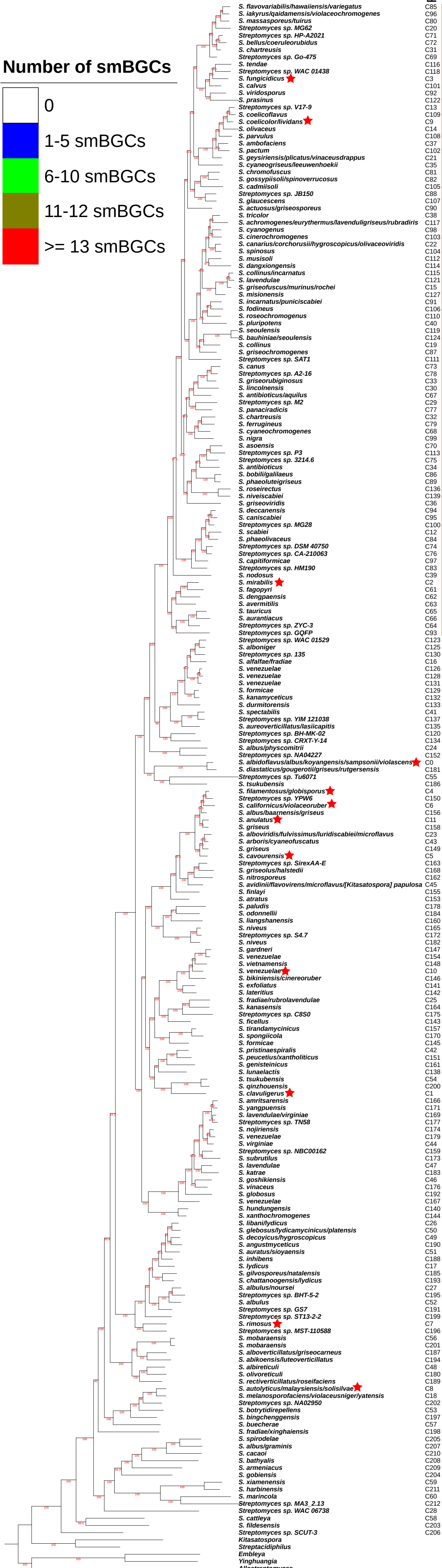
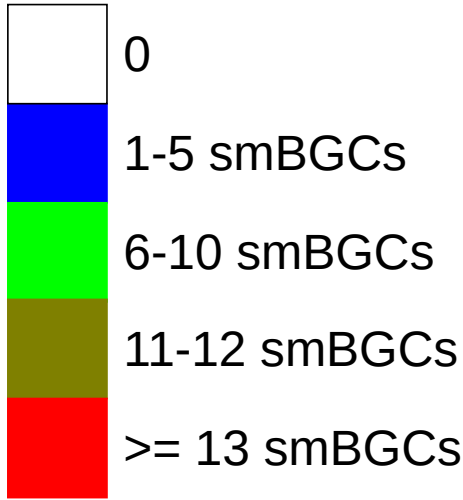
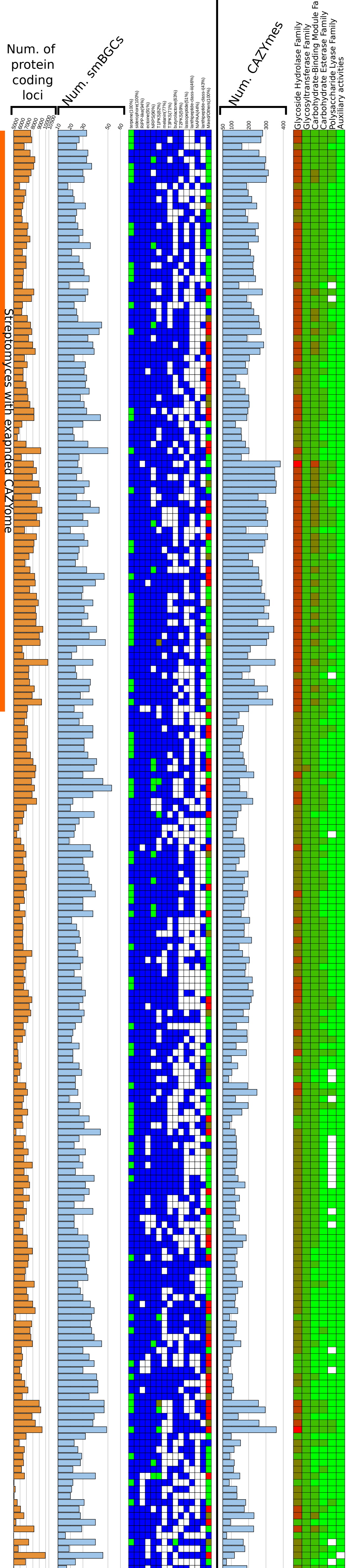


Number of smBGCs



A)

FastANI species cluster



B)

Number of CAZymes per family

