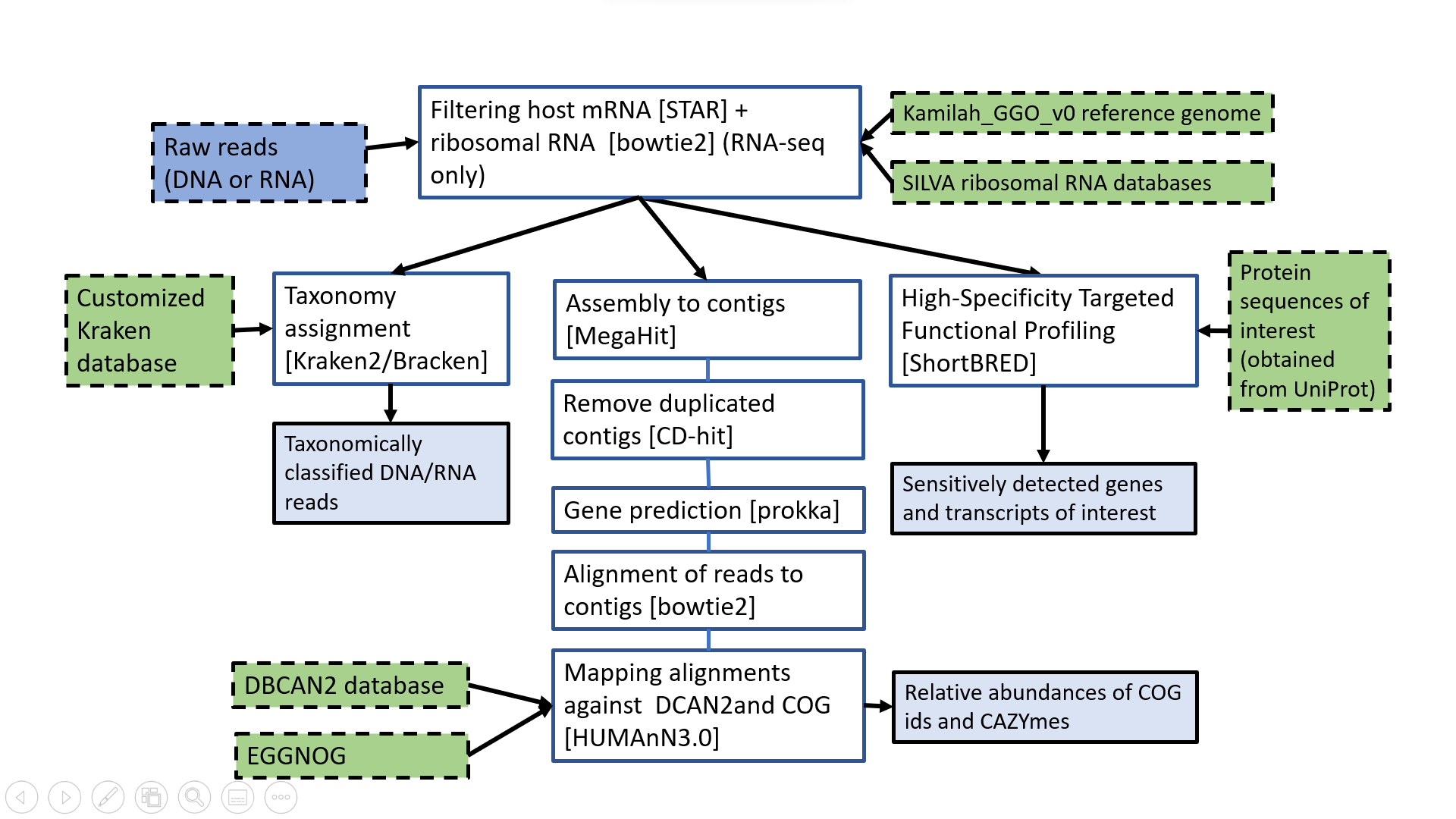
**Supplementary Figure S2**

**Manuscript:**

Houtkamp, I., Van Zijll Langhout, M., Bessem, M., Pirovano, W., & Kort, R. (2023). Multiomics characterization of the of the zoo-housed gorilla gut microbiome reveals bacterial community compositions shifts, fungal cellulose-degrading, and archaeal methanogenic activity. *Gut Microbiome,* 1-25. doi:10.1017/gmb.2023.11



**Figure S2: Shotgun metagenomics/RNA-seq analysis workflow.** The workflow summarizes the analysis of DNA and RNA reads obtained through shotgun metagenomics and RNA-seq, respectively, as described in detail in the methods section.