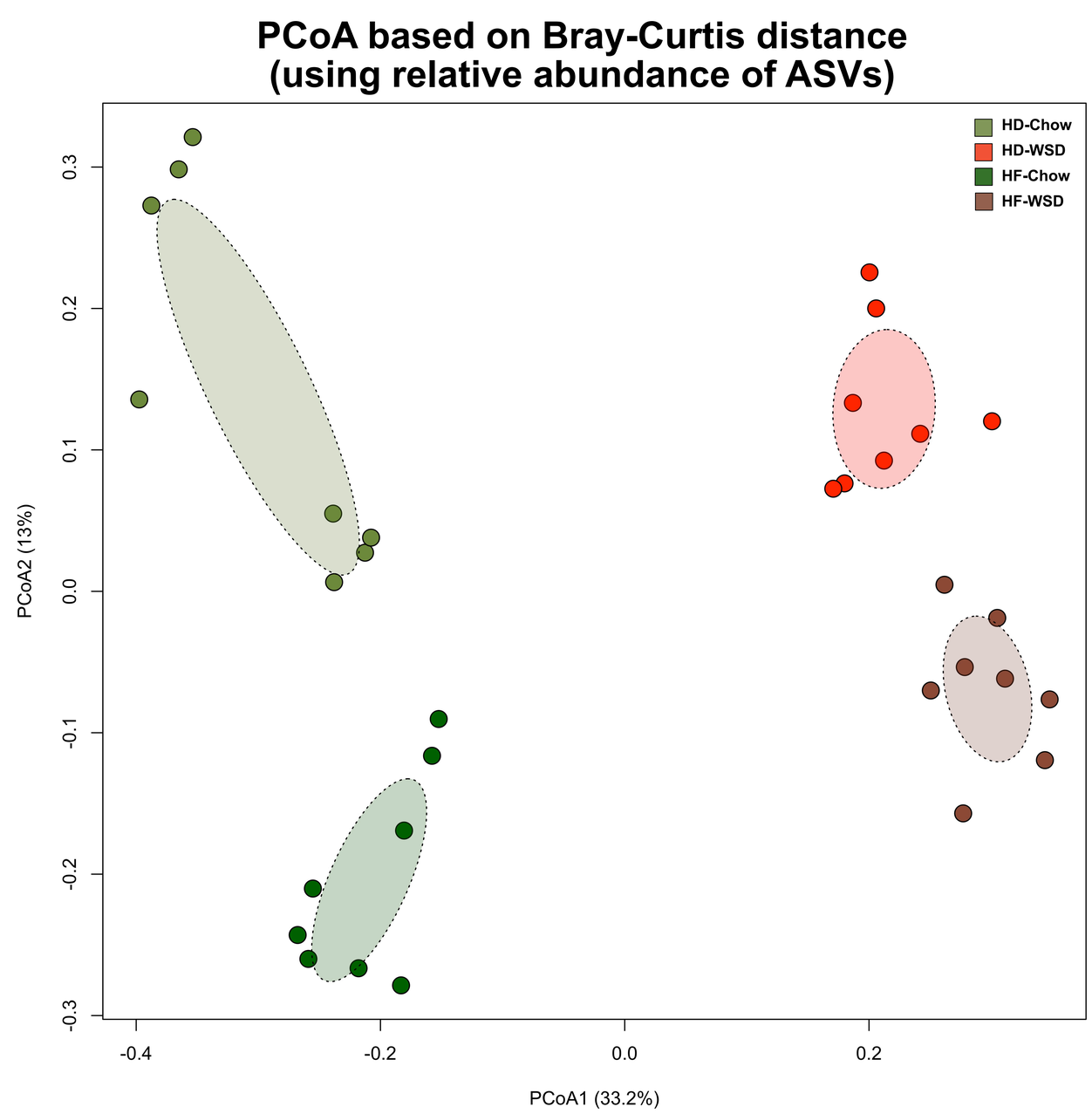
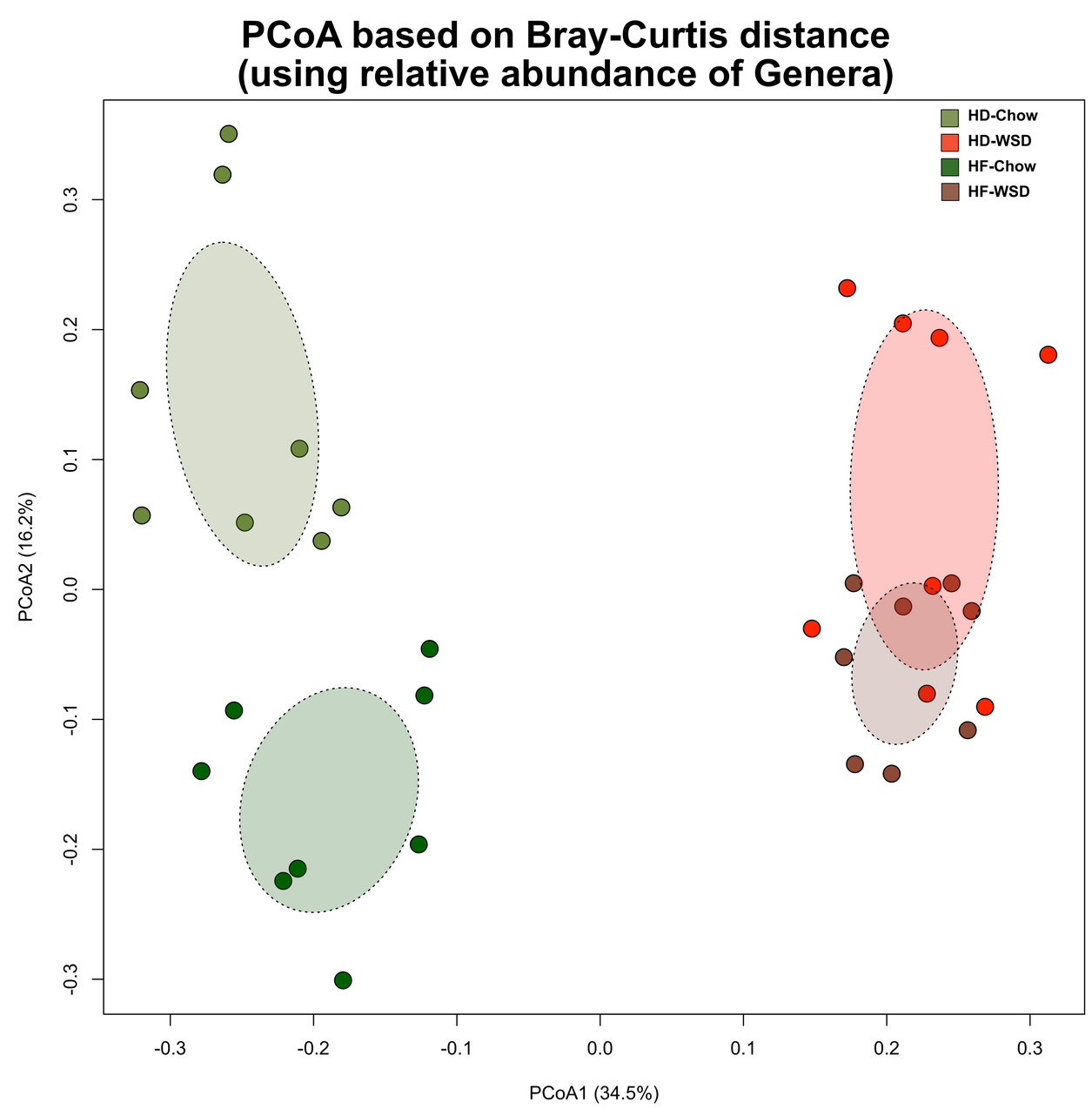
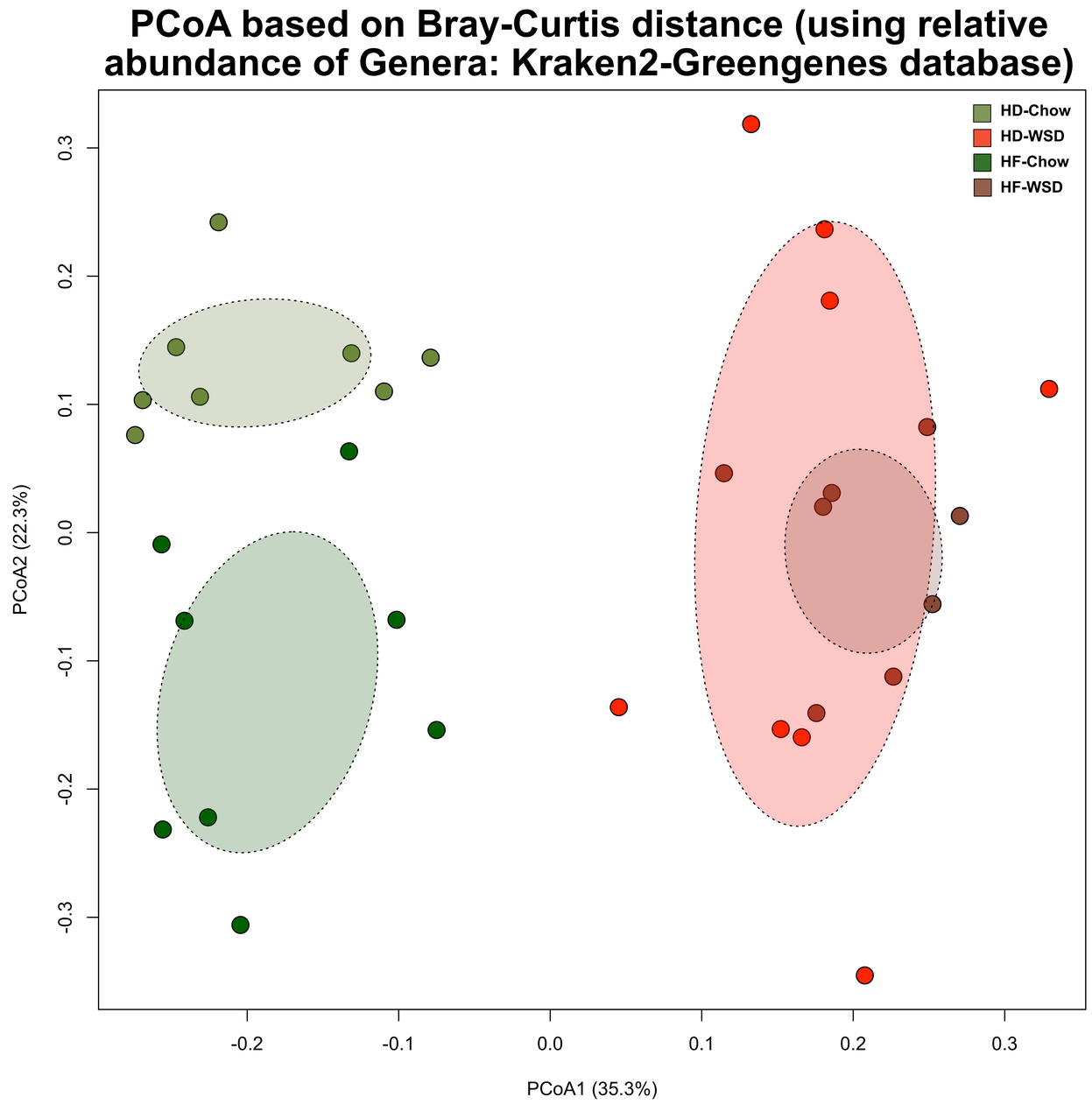
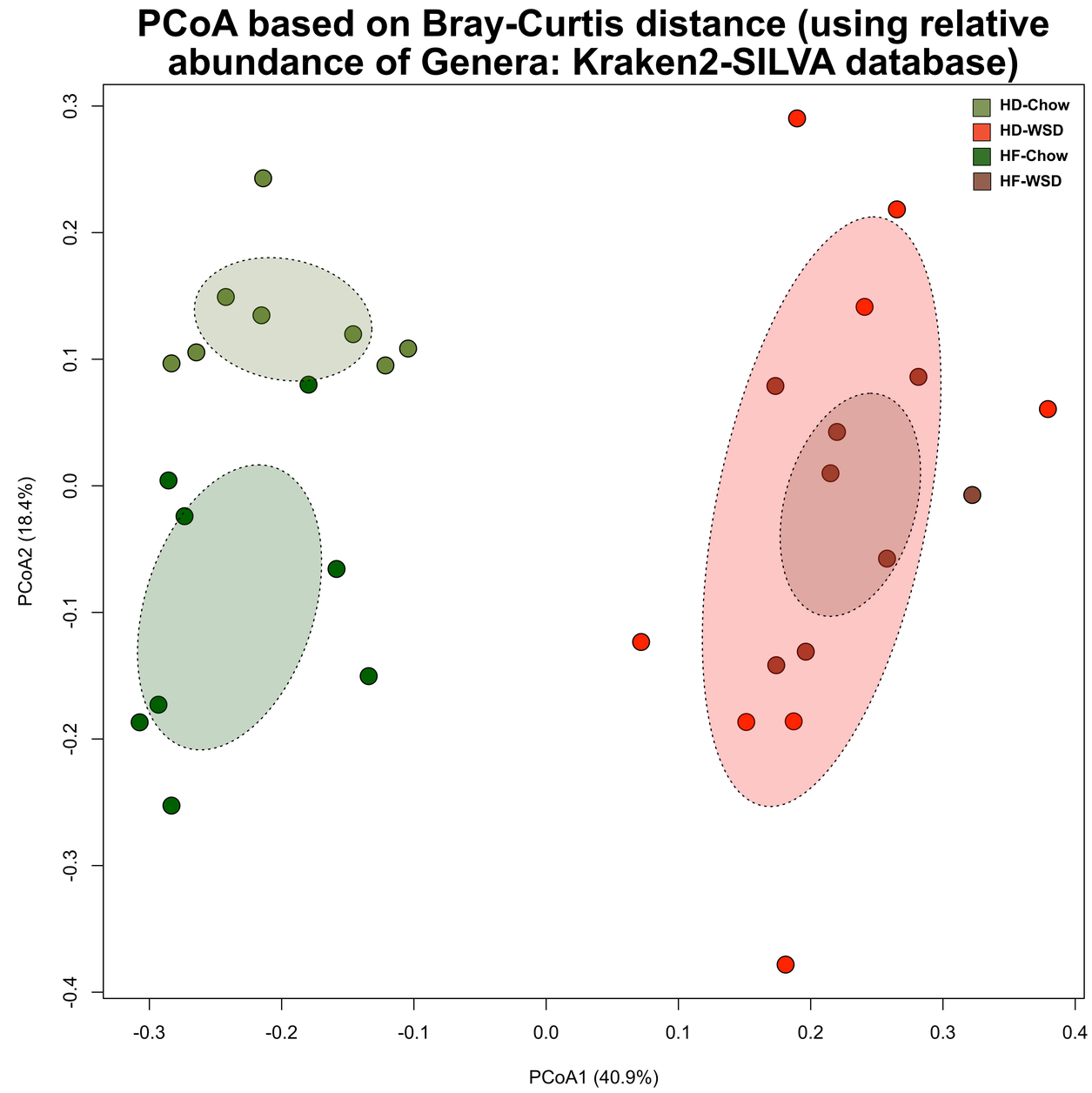

PCoA based on Bray-Curtis distance
(using relative abundance of ASVs)
HD-Chow
HD-WSD
HF-Chow
HF-WSD
PCoA2 (13%)
PCoA1 (33.2%)


PCoA based on Bray-Curtis distance
(using relative abundance of Genera)
HD-Chow
HD-WSD
HF-Chow
HF-WSD
PCoA2 (16.2%)
PCoA1 (34.5%)


PCoA based on Bray-Curtis distance (using relative
abundance of Genera: Kraken2-Greengenes database)
HD-Chow
HD-WSD
HF-Chow
HF-WSD
PCoA2 (22.3%)
PCoA1 (35.3%)


PCoA based on Bray-Curtis distance (using relative
abundance of Genera: Kraken2-SILVA database)
HD-Chow
HD-WSD
HF-Chow
HF-WSD
PCoA2 (18.4%)
PCoA1 (40.9%)