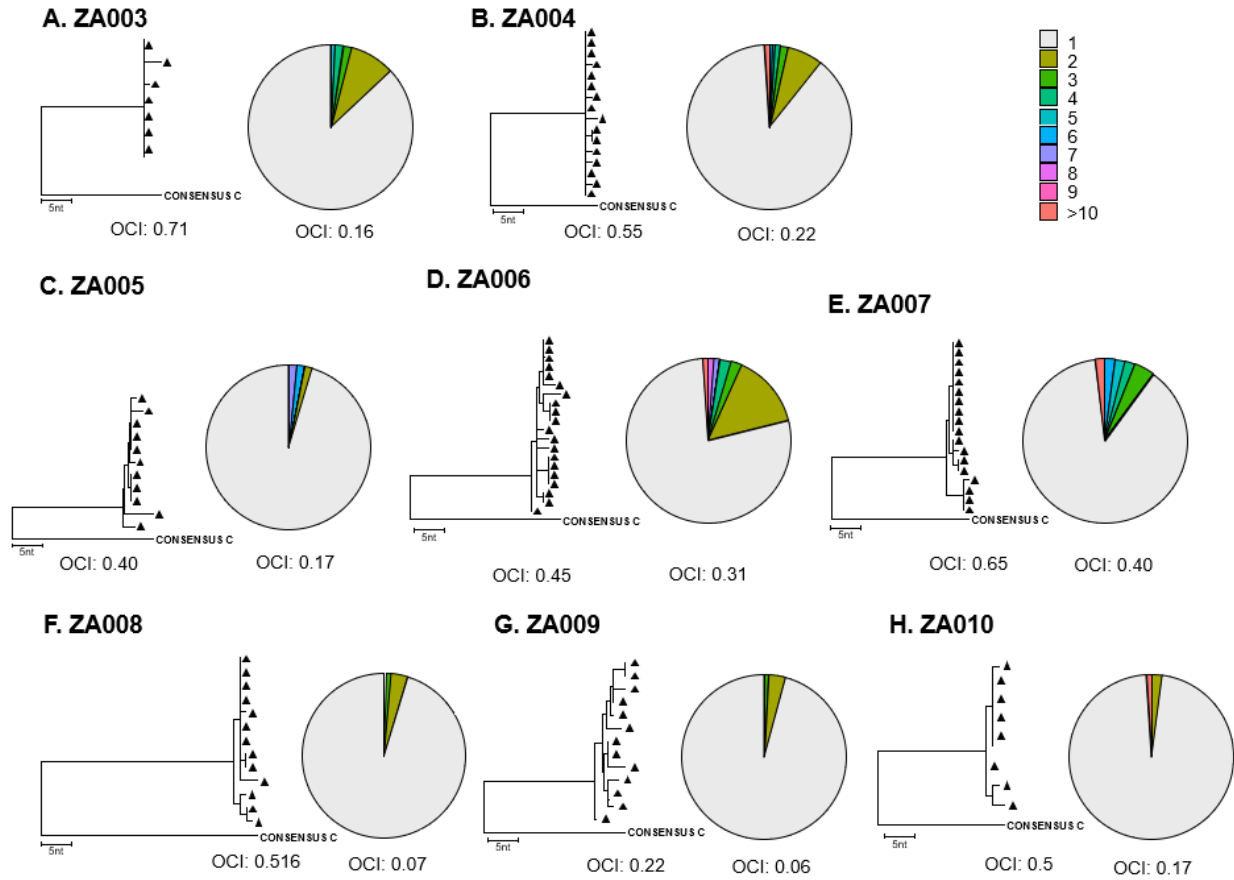# Figure S1. Number of Detections of Integration Sites.

For each study participant, a neighbor-joining phylogenetic tree representing *gag-pol* single genome sequences with its respective OCI value is shown on the left; on the right, a pie chart representing the number of detections of integrations sites by ISA and the respective OCI value.