

Supplementary Figure 2. Overall viral composition in NAT and OSCC patient samples: Sunburst plots showing inferred viral composition in NAT (left) and OSCC (right) patient samples. Viruses were inferred by at least 2 proteins, with those showing  $log_2$  fold-changes of  $\leq$  -1 and  $\geq$  1 being allocated to NAT and OSCC conditions, respectively. Species inferred by proteins within this range were considered present in both conditions. Classification level from inner to outermost radials: Kingdom, Phylum, Class. Colours represent species' phylum while percentage values represent proportion within a given classification level.