



S3 Fig. Statistical significance of overlaps between sets of PPIs (network edges). (A) The overlap between HCV-VIPs_{direct} PPIs and PPIs in PHISTO that were determined as direct based on their experimental methods (see S4 Table). (B) The overlap between HCV-VIPs_{indirect} PPIs and PPIs in PHISTO that could not be determined as direct. Note that in PHISTO, other than core, NS3-4A and NS5A, the identity of the individual HCV protein(s) involving in the interaction with host proteins is not known; consequently, HCV was considered as a single node in the PPI network from PHISTO, and, therefore, the number of virus-host PPIs (i.e. network edges) is the same as that of the host proteins in this enrichment test.