Table S2: Average number of proteins (network size) comprised in each network and the number of seeds that are re-discovered during cross-validation when considering different data for generating the specific HIV receptor network.

HIV Network Type	Average Network Size	Number of recovered receptors
PPI Network	89 ($\pm$ 6)	2 of 13
PPI+GO Network	418 ($\pm$ 8)	11 of 13
PPI+ GO$_{enrich}$ Network	726 ($\pm$ 16)	12 of 13