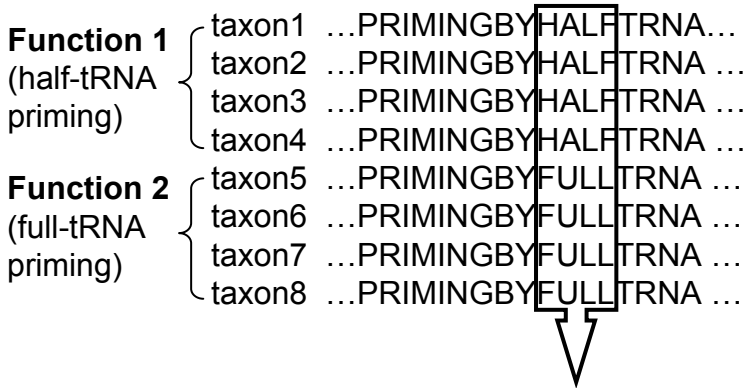
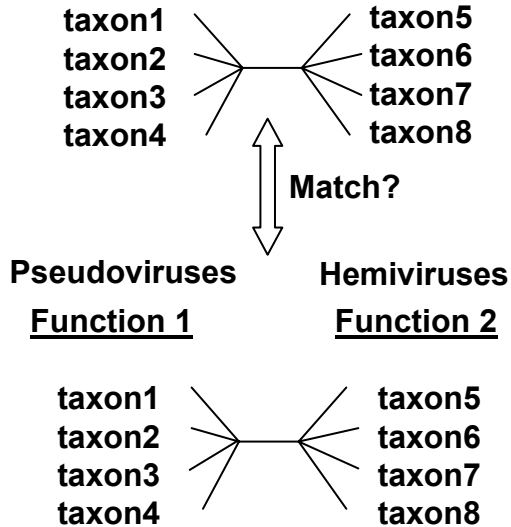


*Amino Acid Alignment*



*Tree from aligned sequences (split cluster)*



*Predefined functional split*