

Sequence alignment visualization showing five protein sequences (dISFV-NC_016997.1, dISFV-NC_017086.1, dISFV-NC_040610.1, dISFV-NC_024017.1, dISFV-NC_027999.1) aligned against a reference sequence. The alignment is displayed as a grid of colored blocks (yellow, green, red) representing different amino acid residues. The reference sequence is shown at the bottom, with positions 10 through 270 indicated. The alignment shows high similarity between the sequences, particularly in the regions corresponding to positions 10-20, 30-40, 50-60, 70-80, 90-100, 110-120, 130-140, 150-160, 170-180, 190-200, 210-220, 230-240, 250-260, and 270-280.