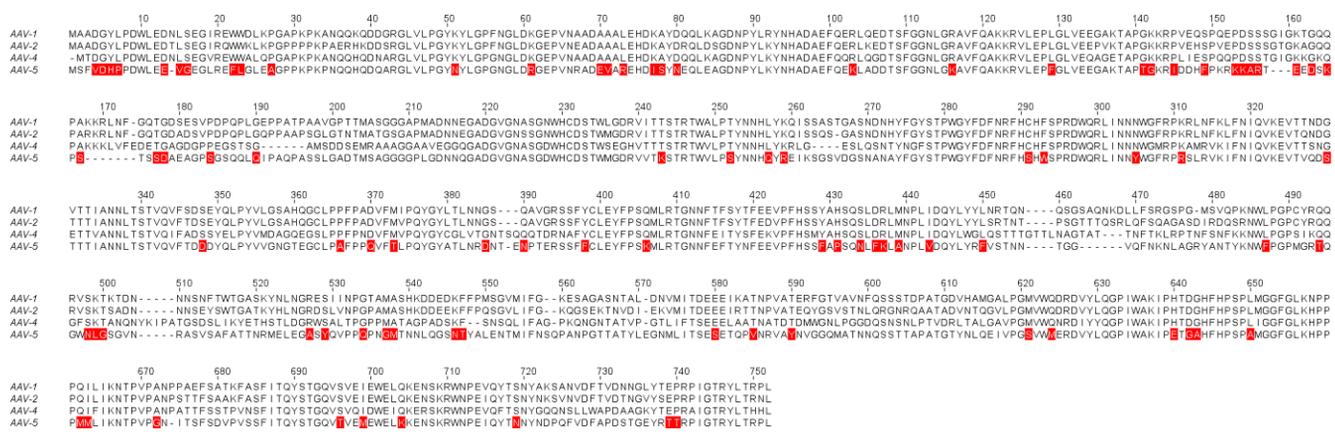Fig. S3. Multiple Amino Acid Sequence Alignment for the Detection of Sequence Variation in AAV5. (A) VP1. (B) MAAP. (C) AAP. Amino acid sequence mutations of AAV5 within the consensus sequence are marked in red.