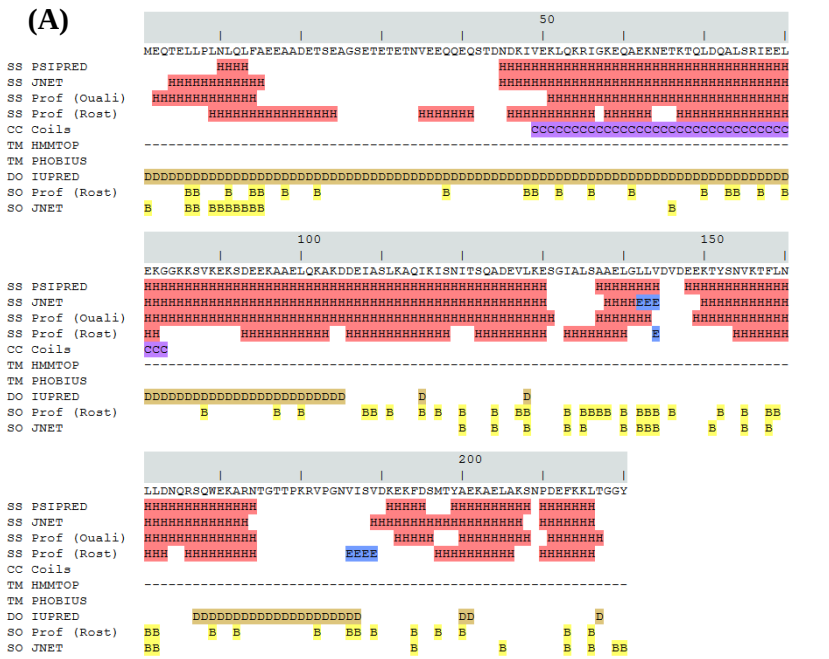
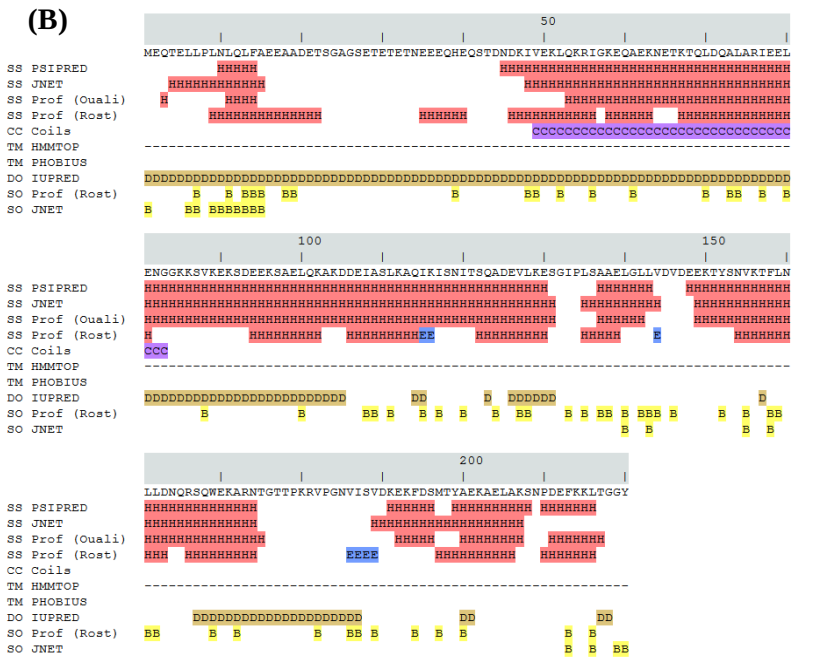
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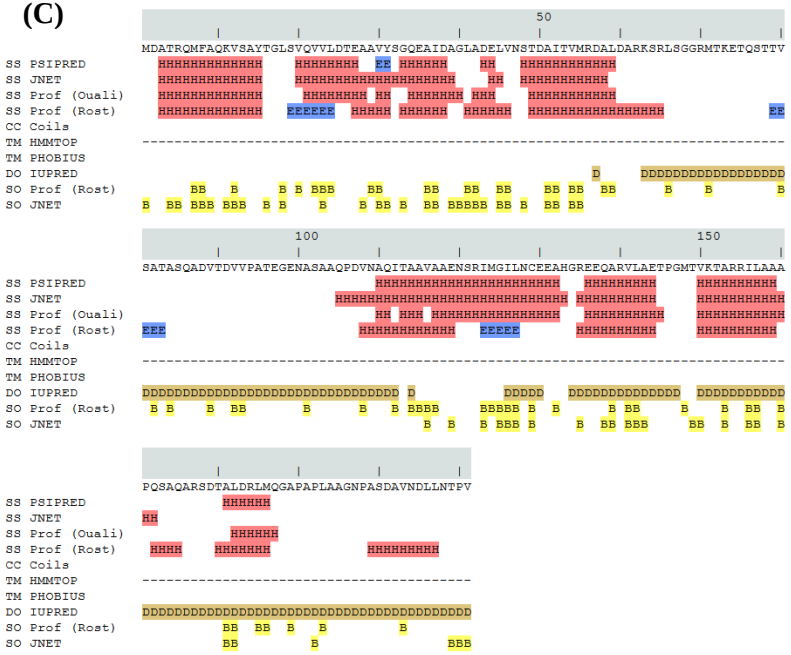
Title	Structure and assembly of TP901-1 virion unveiled by mutagenesis
Authors	Stockdale, Stephen R.;Collins, Barry;Spinelli, Silvia;Douillard, François P.;Mahony, Jennifer;Cambillau, Christian;van Sinderen, Douwe
Publication date	2015-01-01
Original Citation	Stockdale SR, Collins B, Spinelli S, Douillard FP, Mahony J, Cambillau C, et al. (2015) Structure and Assembly of TP901-1 Virion Unveiled by Mutagenesis. PLoS ONE 10(7): e0131676. doi:10.1371/journal.pone.0131676
Type of publication	Article (peer-reviewed)
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(A)
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SS PSIPRED
SS JNET
SS Prof (Ouali)
SS Prof (Rost)
CC Coils
TM HMMTOP
TM PHOBIUS
DO IUPRED
SO Prof (Rost)
SO JNET
SS = Alpha-Helix Beta-Sheet Secondary Structure
CC = Coiled Coils
TM = Transmembrane ('+'=outside, '-'=inside)
DO = Disorder
SP = Signal Peptide Cleavage Site
SO = Solvent accessibility (A burried residue has at most 25% of its surface exposed to the solvent.)

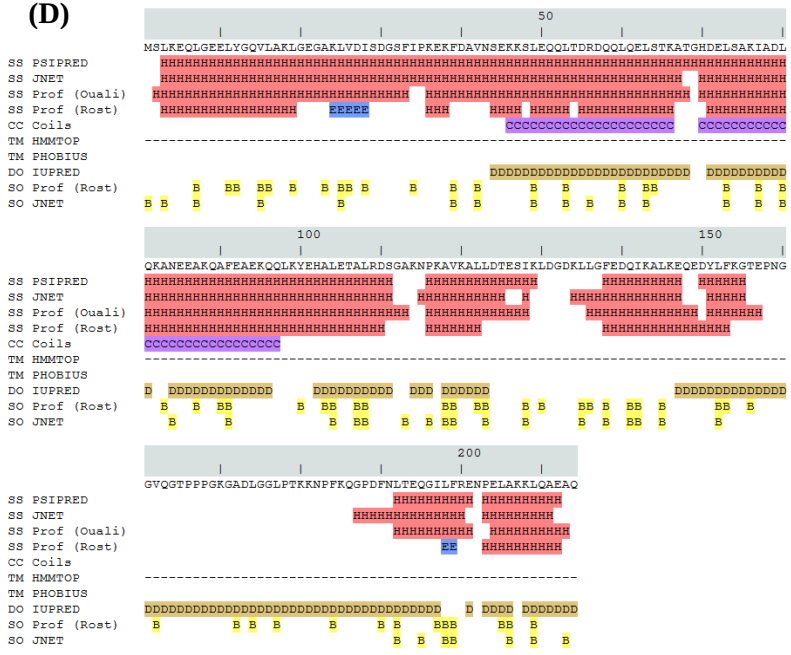

(B)
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LLDNQRSQWEKARNTGTTPKRVPGNVISVDKEKFDSMTYAEKAELAKSNPDEFKKLTGGY
SS PSIPRED
SS JNET
SS Prof (Ouali)
SS Prof (Rost)
CC Coils
TM HMMTOP
TM PHOBIUS
DO IUPRED
SO Prof (Rost)
SO JNET
SS = Alpha-Helix Beta-Sheet Secondary Structure
CC = Coiled Coils
TM = Transmembrane ('+'=outside, '-'=inside)
DO = Disorder
SP = Signal Peptide Cleavage Site
SO = Solvent accessibility (A burried residue has at most 25% of its surface exposed to the solvent.)

(C)



SS = Alpha-Helix Beta-Sheet Secondary Structure
CC = Coiled Coils
TM = Transmembrane ('+'=outside, '-'=inside)
DO = Disorder
SP = Signal Peptide Cleavage Site
SO = Solvent accessibility (A burried residue has at most 25% of its surface exposed to the solvent.)

(D)



SS = Alpha-Helix Beta-Sheet Secondary Structure
CC = Coiled Coils
TM = Transmembrane ('+'=outside, '-'=inside)
DO = Disorder
SP = Signal Peptide Cleavage Site
SO = Solvent accessibility (A burried residue has at most 25% of its surface exposed to the solvent.)