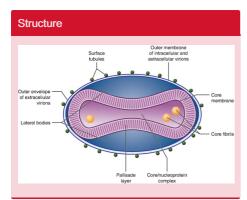


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Structure		
Spherical Shaped		
Enveloped virus	One lipid membrane	
Outer layer	Corrugated due to protrusions	
Internal Core	Dumbbell shaped	
Lateral bodies	Two present in the concavities between outer layer and Core	
Outer core wall	Palisade structure with T- shaped spike proteins	
Inner Core wall	Smooth	

Proteins	
Attachment	A17, A25, A26, A27, D8, H3
Entry	Entry Fusion Complex-A16, A21, A28, F9, G3, G9, H2, I2, J5, L1, L5, O3
Redox active proteins	E10, A2.5, and G4
Phosphory- lation	Kinase- F10 & B1
Dephos- phorylation	Phosphotase - H1
Proteolytic processing	proteinase - I7

Bommana	Sushma Bhandarkar
Chanakya	
213501003	213501004
Poster By	
Poster By Bommana	Sushma Bhandarkar

213501004

Poster By

213501003

Genom	e Organisation
	Terrical loop (20 75 55 50 50 50 50 50 50 50 50 50 50 50 50
	Terion reports Freend students Freend
	C TROMS

Has ITRs with secondary nairpin loop structures that connect two DNA strands and
contains the variable genes that
are responsible for host cell
nteractions
Conserved genes responsible or replication and morpho-

Proteins	
Attachment	A17, A25, A26, A27, D8, H3
Entry	Entry Fusion Complex - A16, A21, A28, F9, G3, G9, H2, I2, J5, L1, L5, O3
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Protein	
Early Gene Proteins	Inducing proliferation of neighbouring cells
	Counteracting host immune defences
	Some proteins can induce a second uncoating reaction thereby releasing the viral genome from the core
	Mediating replication of the genome
Protiens In Replication	Viral DNA polymerase
	Helicase-primase
	Uracil DNA glycosylase
	Single-stranded DNA-binding protein
	DNA ligase
	Holliday junction resolvase
Proteins In Transc- ription	RNA polymerase
	ETF
	Capping enzyme
	VITF-3
	VLTF-1, VLTF-2, VLTF-3, VLTF-

Protein (cont) Release factor Elongation factor Poly(A) polymerase DNA topoisomerase Replication Cycle Replication process Attachment Entry



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Early Gene Transcription

Late Gene Transription

Intermediate Gene Transcription

DNA Replication

Assembly Release

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