

Pathogenesis of Plant Virus
MIC 204
(MSc Microbiology Semester II)

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Plant Virus Diseases and Symptoms



<https://www.bighaat.com/blogs/kb/viral-diseases-in-plants-and-its-control>

Virus taxonomy and nomenclature

- Modified binomial is used
- Taxonomy depends on particle properties, nucleic acid properties and especially sequence
- **Family** is the highest taxonomic level that is commonly used; ends in *viridae*, e.g., *Bromoviridae*
- **Genus** ends in suffix *virus*, e.g., *Bromovirus*
- **Species** is usually the commonly used virus name; it is italicized in formal usage, e.g., *Brome mosaic virus*
- Small genome sizes, gene shuffling make broad taxonomic schemes difficult (above Family level)

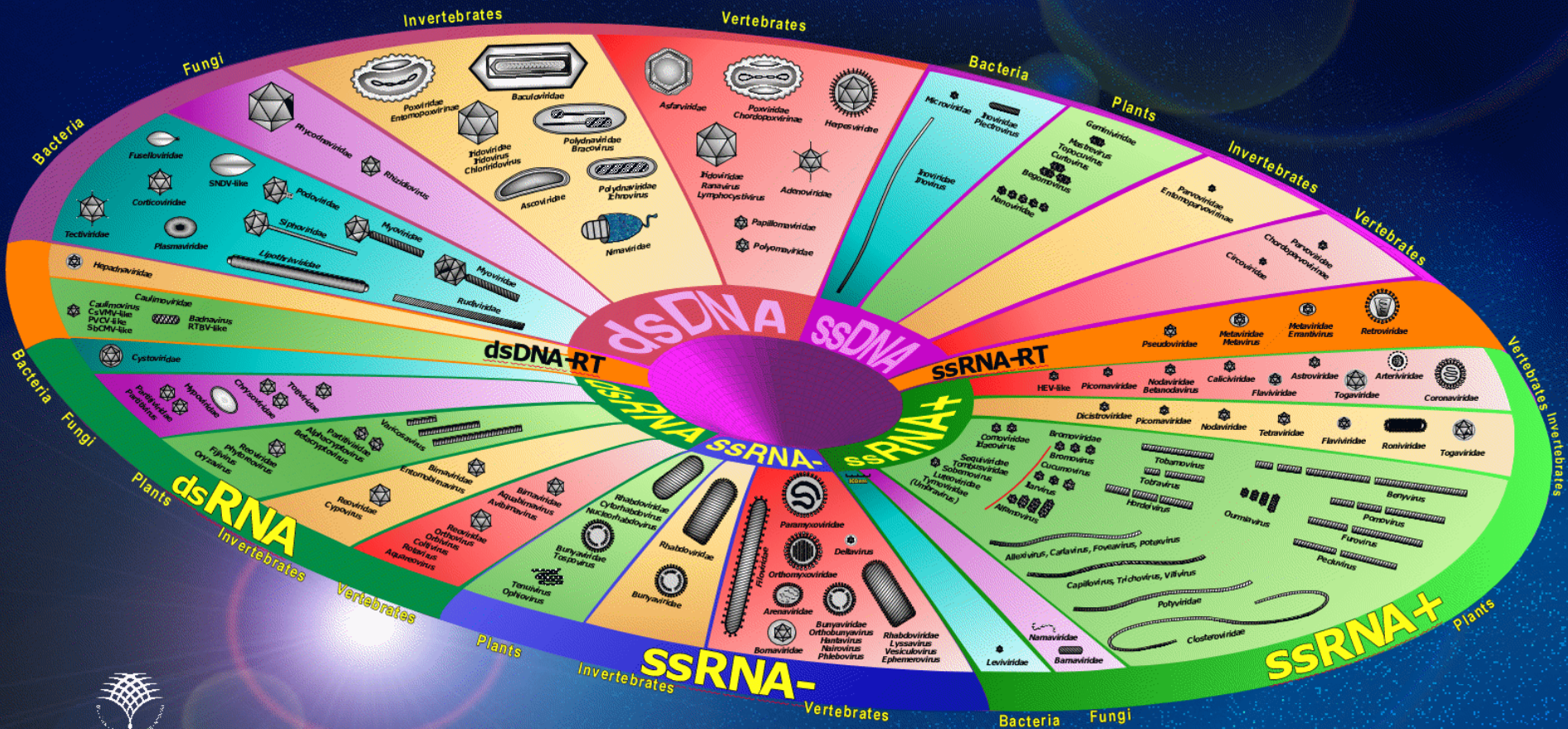
Virus properties: Plant viruses are often simpler than animal viruses

- Genome sizes 0.3 - 300 kb; **plant viruses 0.3-30 kb**
- May have single-stranded or double-stranded RNA or DNA genome; **most plant viruses ssRNA**
- If RNA, may be + or – sense; **most plant viruses + sense ssRNA**
- May have one or many proteins in particles; **most plant viruses have 1-2**
- May or may not have lipid envelope; **most plant viruses do not**

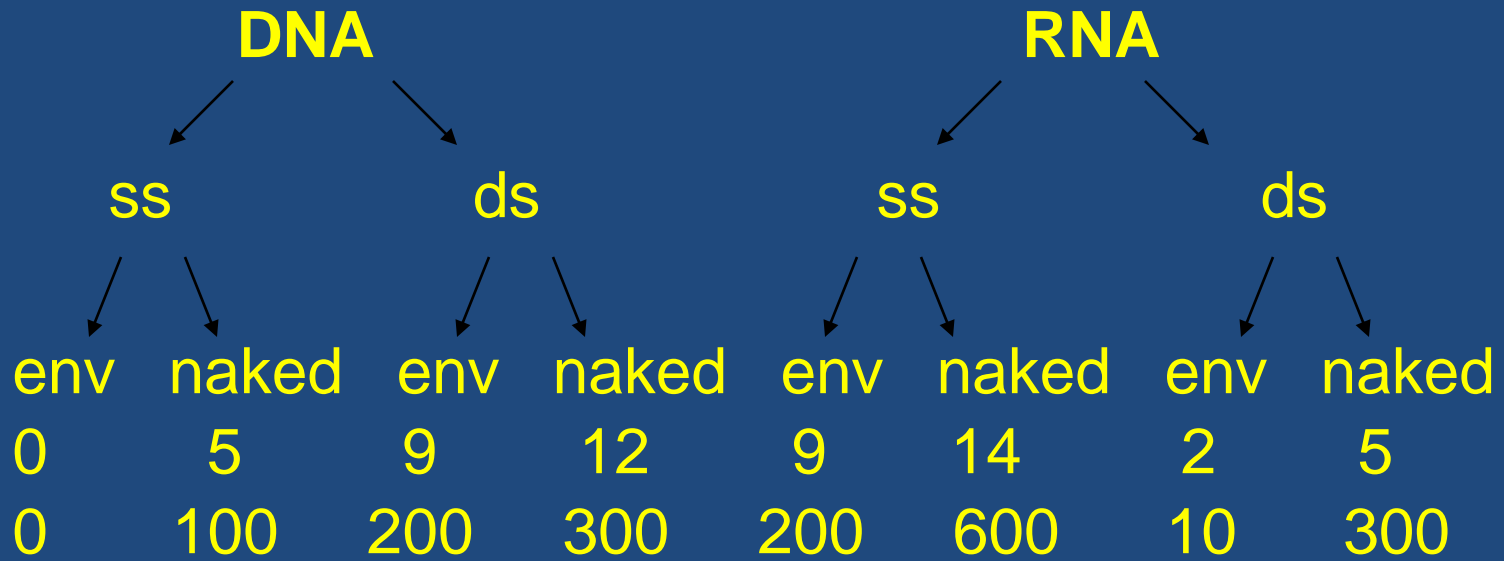
Types of plant virus genomes

- double-stranded (ds) DNA (rare)
- single-stranded (ss) DNA (rare)
- ssRNA, negative sense (rare)
- ssRNA, positive sense (common)
- dsRNA (rare)

Virosphere 2002



Virus types, by nucleic acid



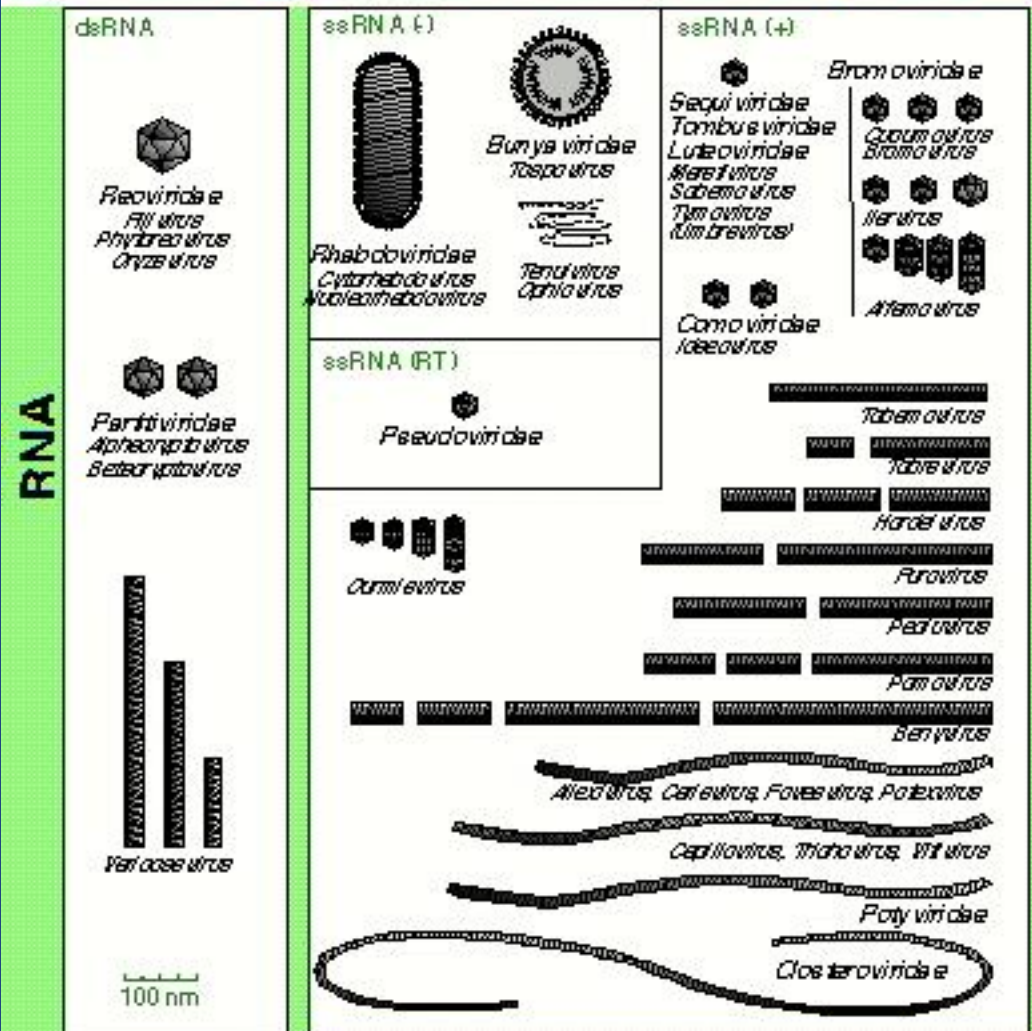
Host type

Vertebrate	-	+	++	++	++	++	-	++
Invertebrate	-	+	++	-	++	++	-	++
Plant	-	++	-	+	+	+++	-	+
Fungus	-	-	-	+	+	+	+	+++
Bacteria	-	+	+	+++	-	+	+	-

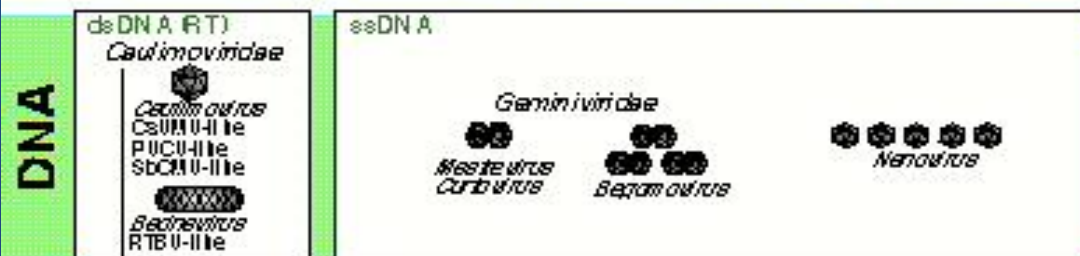
Plant viruses are diverse, but not as diverse as animal viruses – probably because of size constraints imposed by requirement to move cell-to-cell through plasmodesmata of host plants

Plant viruses often contain divided genomes spread among several particles

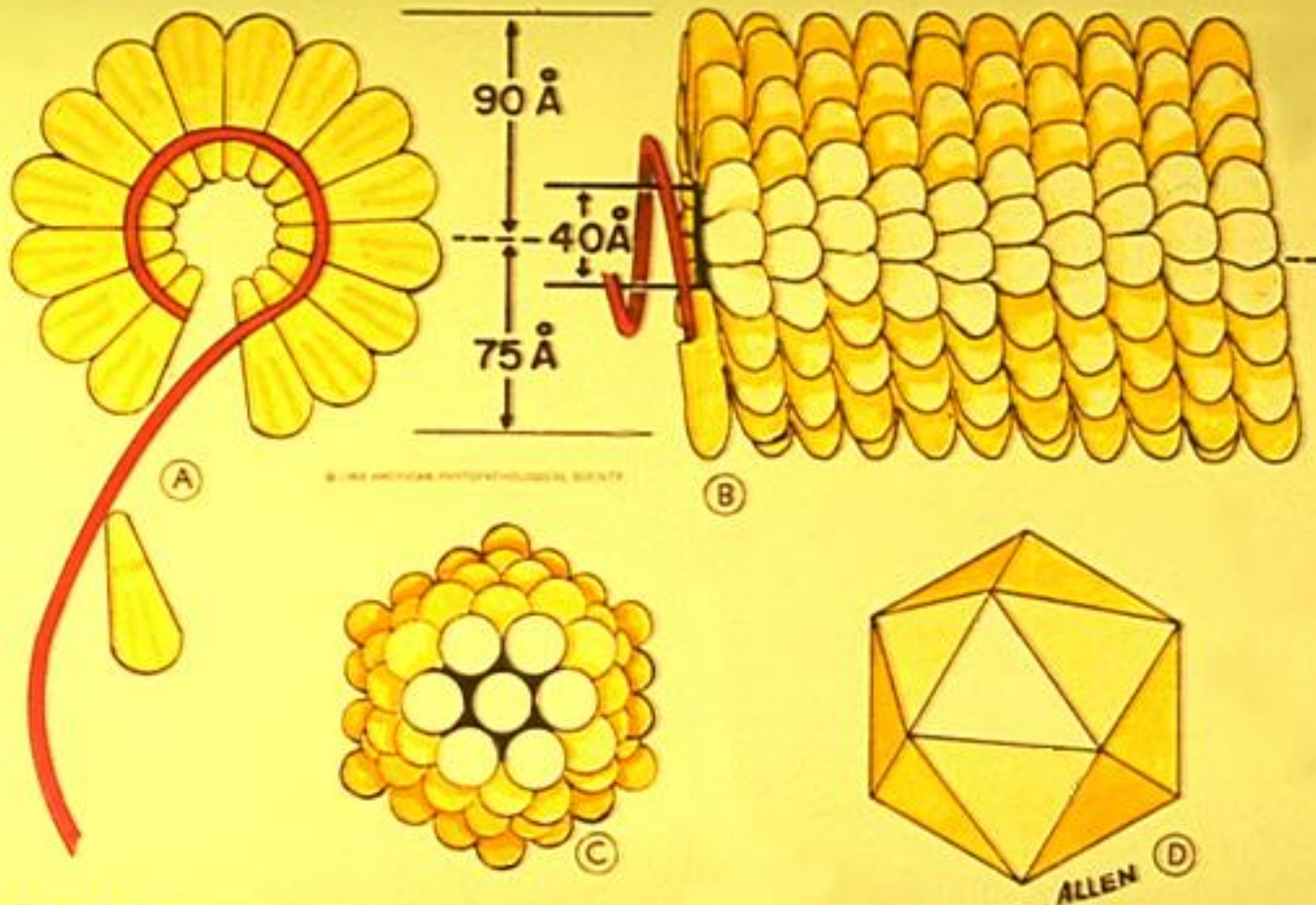
Families and Genera of Viruses Infecting Plants



Families and Genera of Viruses Infecting Plants



Basic Plant Virus Structures



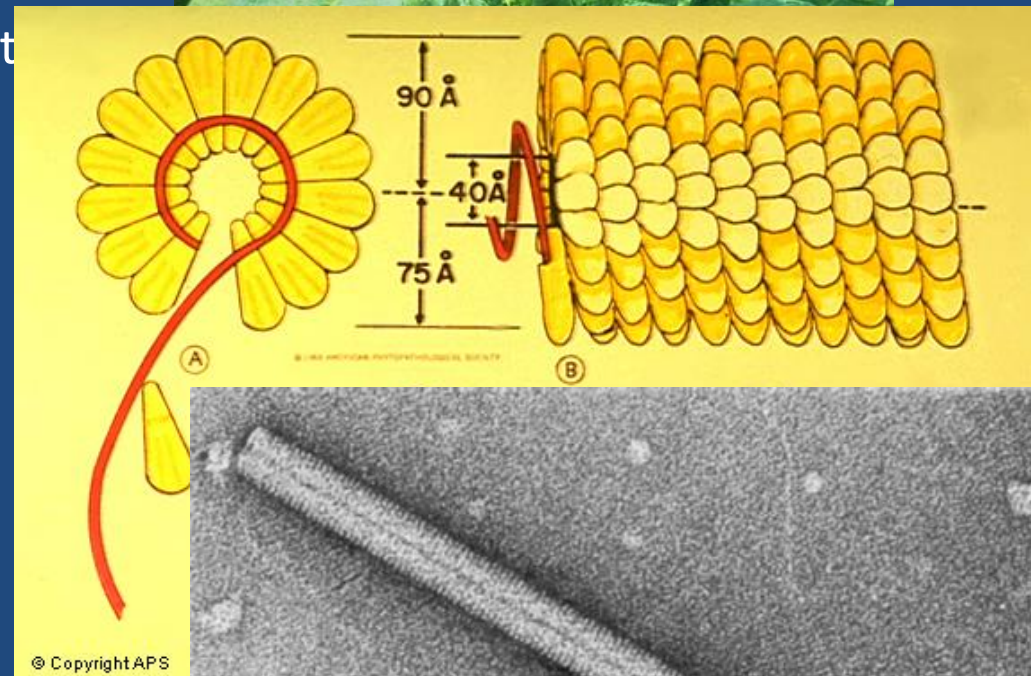
Helix (rod)
e.g., **TMV**

Icosahedron
(sphere)
e.g., **BMV**

Helical symmetry



- *Tobacco mosaic virus* is typical, well-studied example
- Each particle contains only a single molecule of RNA (6395 nucleotide residues) and 2130 copies of the coat protein subunit (158 amino acid residues; 17.3 kilodaltons)
 - 3 nt/subunit
 - 16.33 subunits/turn
 - 49 subunits/3 turns
- TMV protein subunits + nucleic acid will self-assemble *in vitro* in an energy-independent fashion
- Self-assembly also occurs in the absence of RNA

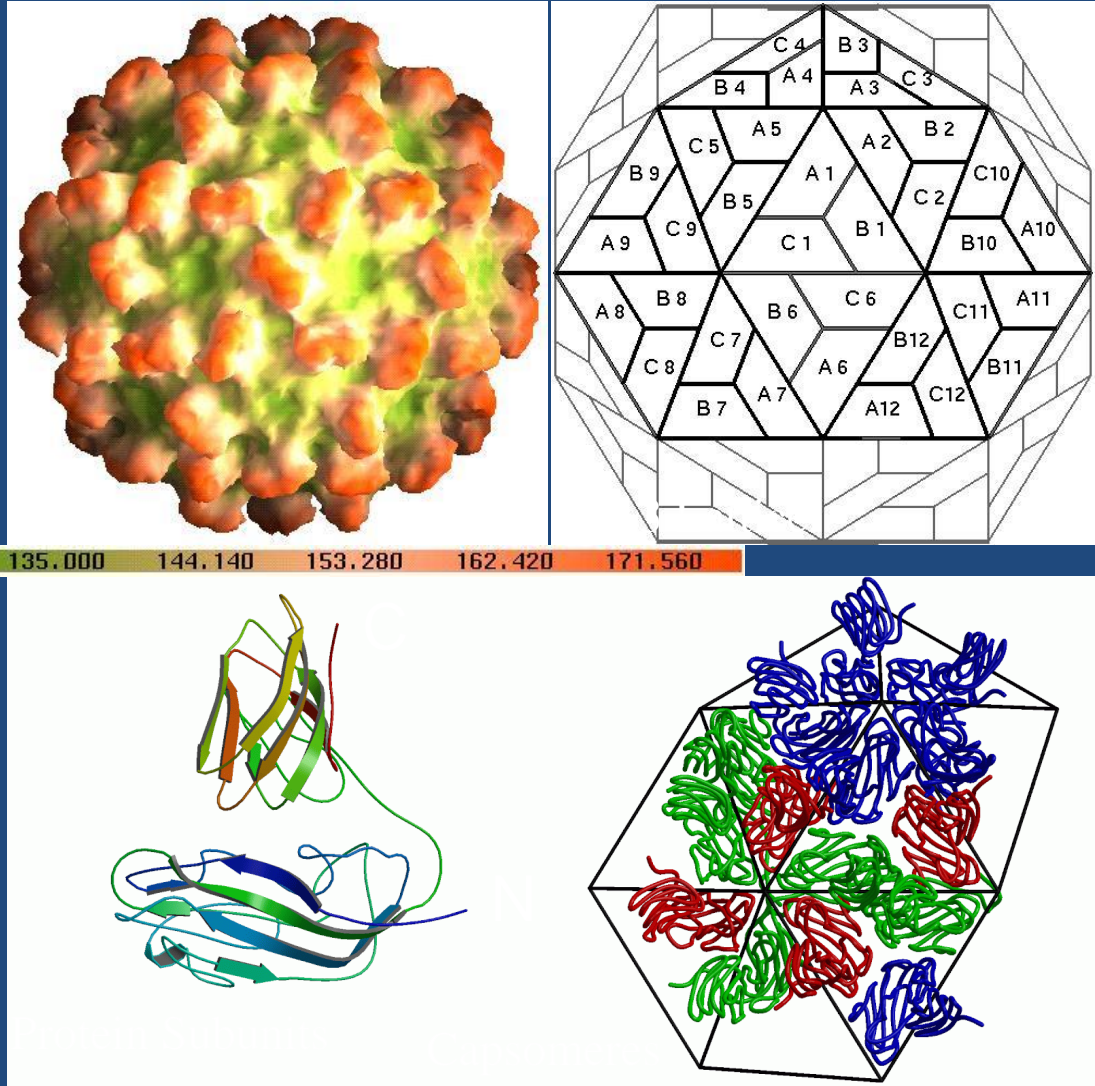


**TMV rod is 18 nanometers (nm)
X 300 nm**

Cubic (icosahedral) symmetry

TBSV icosahedron is 35.4 nm in diameter

- *Tomato bushy stunt virus* is typical, well-studied example
- Each particle contains only a single molecule of RNA (4800 nt) and 180 copies of the coat protein subunit (387 aa; 41 kd)
- Viruses similar to TBSV will self-assemble *in vitro* from protein subunits + nucleic acid in an energy-independent fashion



GENOME ORGANIZATIONS

Plant virus genome organizations

- Very compact
- Most are +sense RNA viruses, so translation regulation very important
- Use various strategies for genome expression
- Only a few genes **absolutely** required:
 - Replicase
 - Coat protein
 - Cell-to-cell movement protein
- Other genes present in some viruses

Genome expression of + strand RNA viruses

- **Most use more than one strategy**
 - Polyprotein processing
 - Subgenomic RNA
 - Segmented genome
 - Translational readthrough
 - Frameshift
 - Internal initiation of translation (without scanning)
 - Scanning to alternative start site (truncated product)
 - Alternative reading frame (gene-within-a-gene)

Polyprotein processing

- Post-translational cleavage of viral proteins may occur *in cis* or *in trans*
- Some viruses use polyprotein processing exclusively to regulate gene expression
- Many viruses use polyprotein processing as one of several regulation mechanisms
- Examples:
 - Potyviruses*
 - Comoviruses*
 - Closteroviruses
 - Carlaviruses

Subgenomic RNA

- Similar to traditional mRNA, but synthesized from an RNA template
- Many viruses use polyprotein processing as one of several regulation mechanisms
- Examples:
 - Tobamoviruses (TMV)*
 - Bromoviruses (BMV)*
 - Tombusviruses (TBSV)
 - Potexviruses (PVX)

Segmented genome

- Positive sense RNA genomes are usually encapsidated in separate particles
- Segmented genomes lend themselves to recombination
- Examples:
 - Bromoviruses (*Brome mosaic virus*, BMV)*
 - Dianthoviruses (*Red clover necrotic mosaic virus*, RCNMV)*
 - Hordeiviruses (*Barley stripe mosaic virus*, BSMV)

Translational readthrough

- Usually UAG codon is read through using suppressor tyrosine tRNA
- Common mechanism in plant viruses
- Examples:
 - Tobamoviruses (*Tobacco mosaic virus*, TMV)*
 - Dianthoviruses (*Red clover necrotic mosaic virus*, RCNMV)*
 - Hordeiviruses (*Barley stripe mosaic virus*, BSMV)

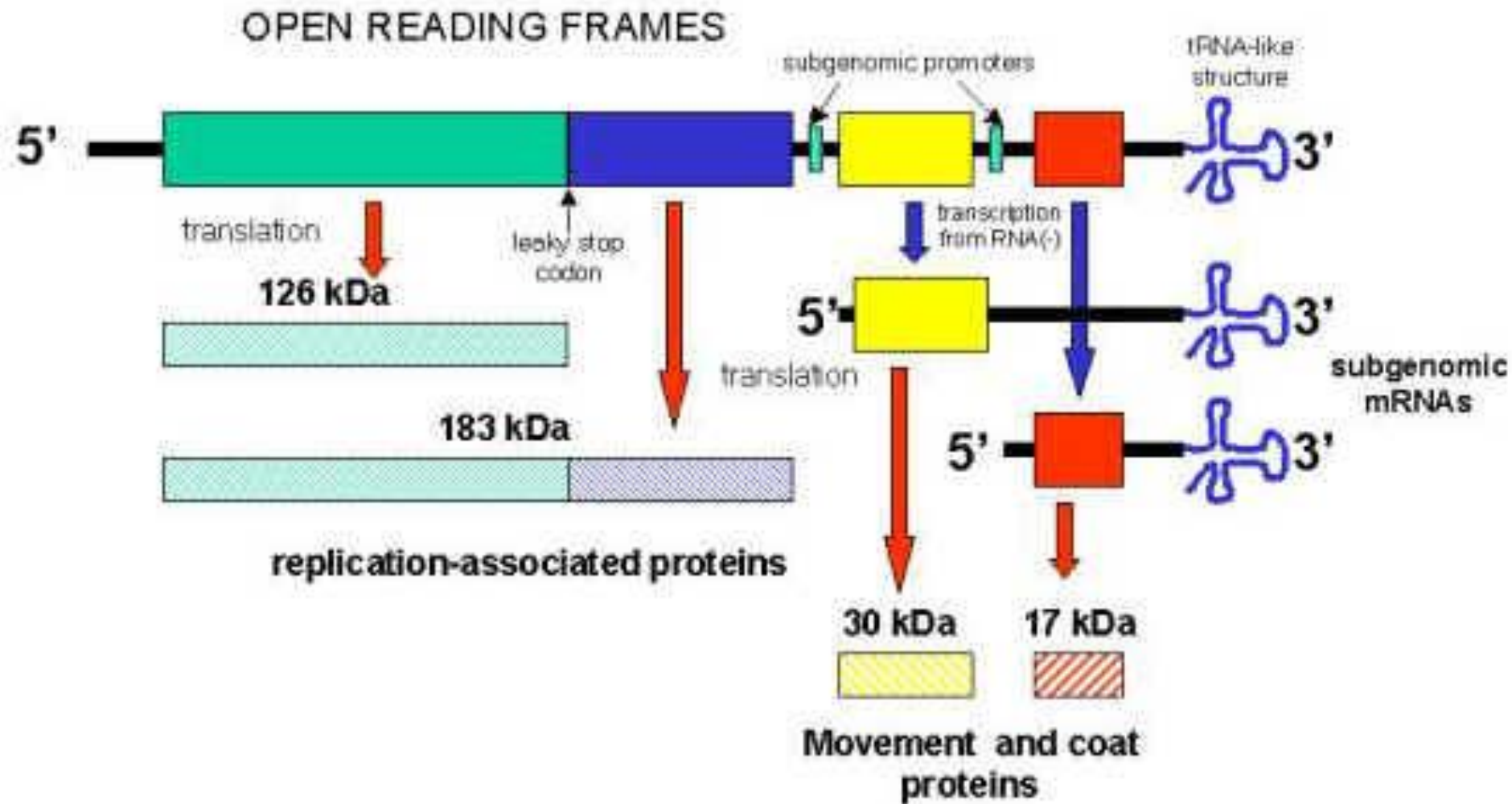
Translational frameshift

- Typically +1 or -1
- Common mechanism in plant viruses
- Examples:
 - Luteoviruses (*Barley yellow dwarf virus*, BYBV)*
 - Dianthoviruses (*Red clover necrotic mosaic virus*, RCNMV)*
 - Closteroviruses (*Beet yellow vein virus*, BYVV)

Internal initiation

- Cap-free translation
- Less complex in plant viruses than in animal viruses
- Examples:
 - Potyviruses (*Tobacco etch virus*, TEV)*
 - Sobemoviruses (*Southern bean mosaic virus*, (SBMV))*

Tobacco mosaic virus is a typical positive-sense RNA plant virus with a 6.4 kilobase genome



INFECTION CYCLE

Plant Virus Life Cycle

- Virus entry into host
 - no attachment step with plant viruses
 - by vector, mechanical, etc. – must be forced
 - requires healable wound – delivery into cell
- Uncoating of viral nucleic acid
 - may be co-translational for + sense RNA viruses
 - poorly understood for many
- Replication
 - replication is a complex, multistep process
 - viruses encode their own replication enzymes

Plant Virus Life Cycle 2

- Cell-to-cell movement
 - cell-to-cell movement through plasmodesmata
 - move as whole particles or as protein/nucleic acid complex (no coat protein required)
- Long distance movement in plant
 - through phloem
 - as particles or protein/nucleic acid complex (coat protein required)
- Transmission from plant to plant
 - requires whole particles

Typical RNA-containing plant virus replication cycle

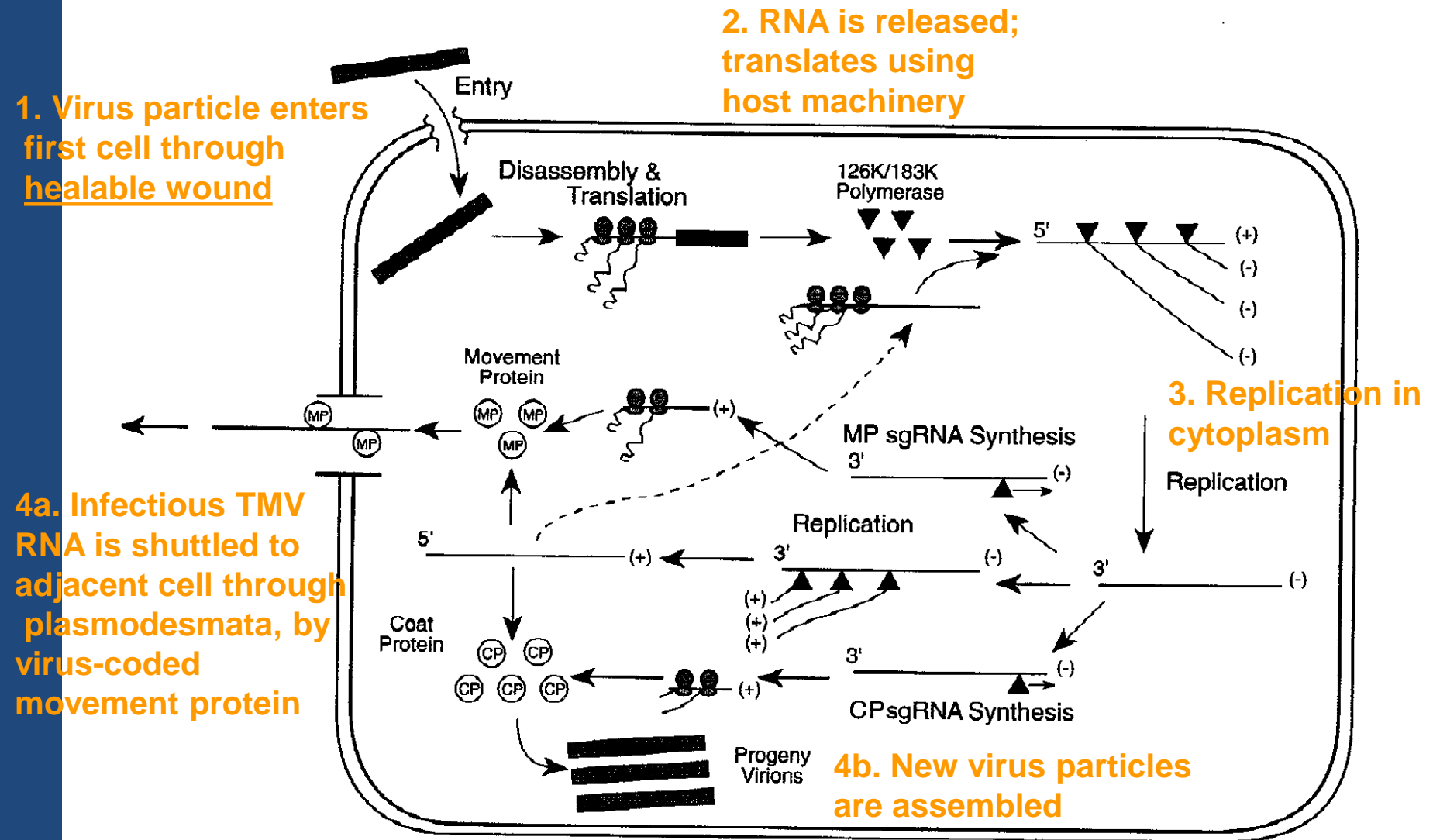


FIG. 8. Diagram of stages in TMV infections. All the events shown are presumed to occur in the cytoplasm of infected cells.

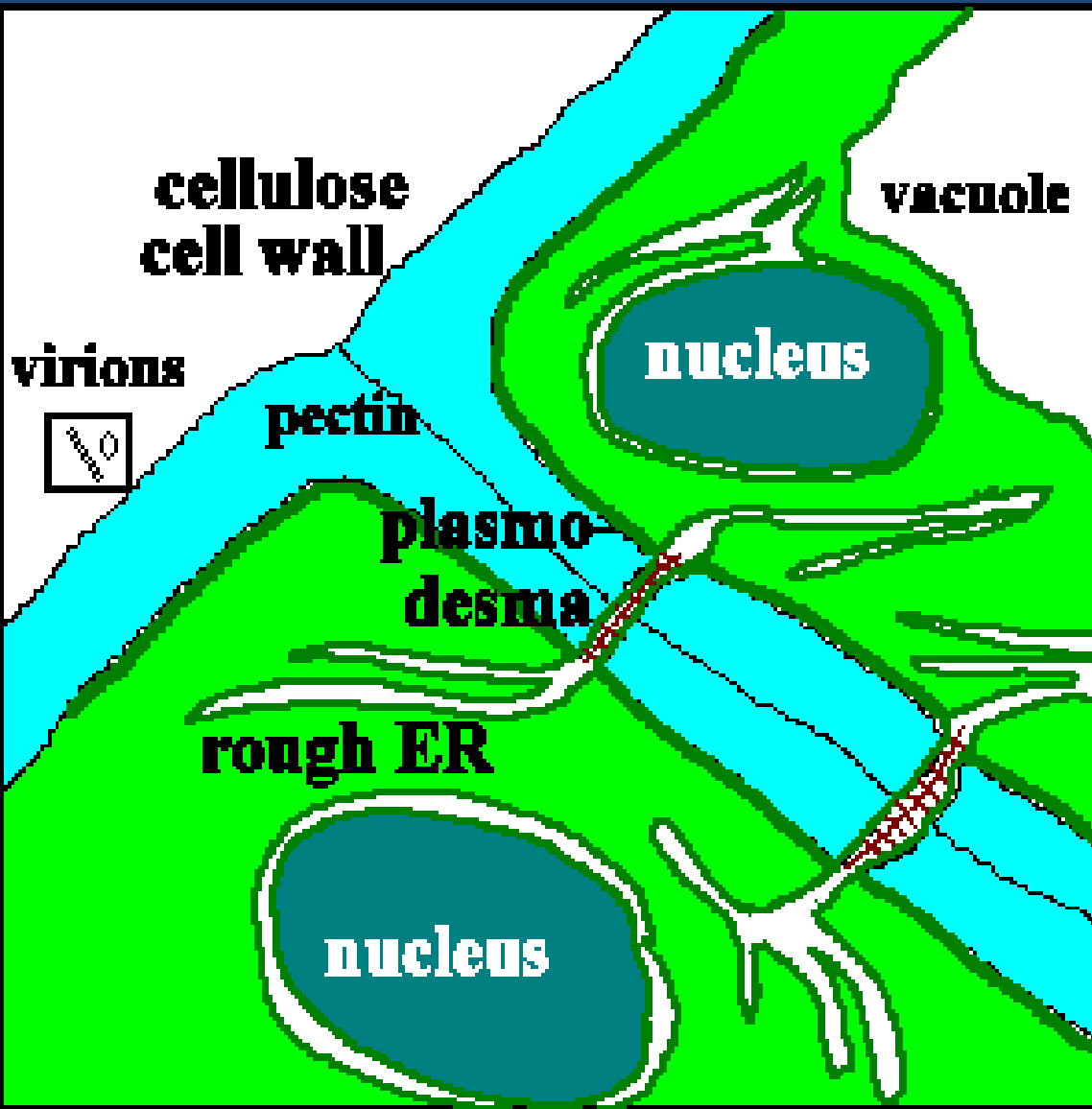
Cell-to-Cell Movement of Plant Viruses

- Plant viruses move cell-to-cell slowly through **plasmodesmata**
- Most plant viruses move cell-to-cell as complexes of non-structural protein and genomic RNA
- The viral protein that facilitates movement is called the “**movement protein**” (MP)
- Coat protein is often dispensable for cell-to-cell movement

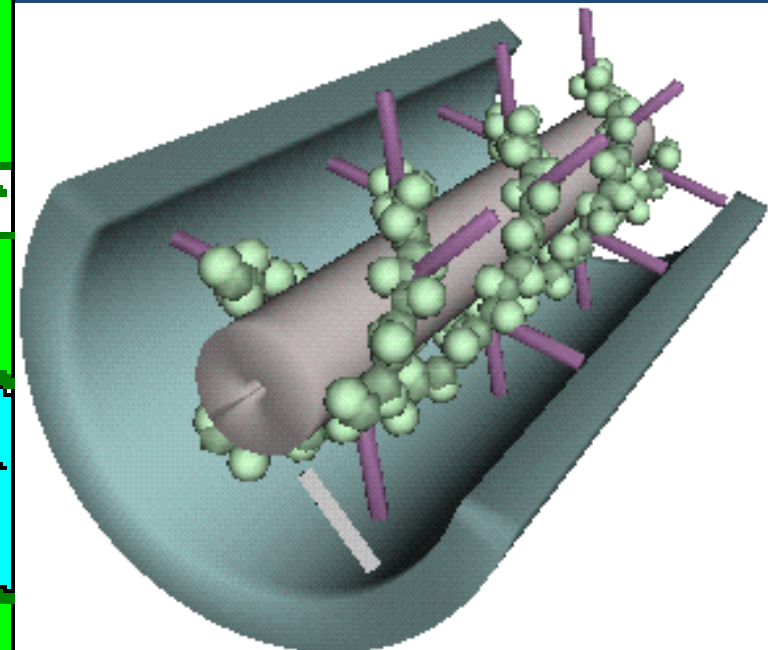
Cell-to-Cell Movement of Plant Viruses

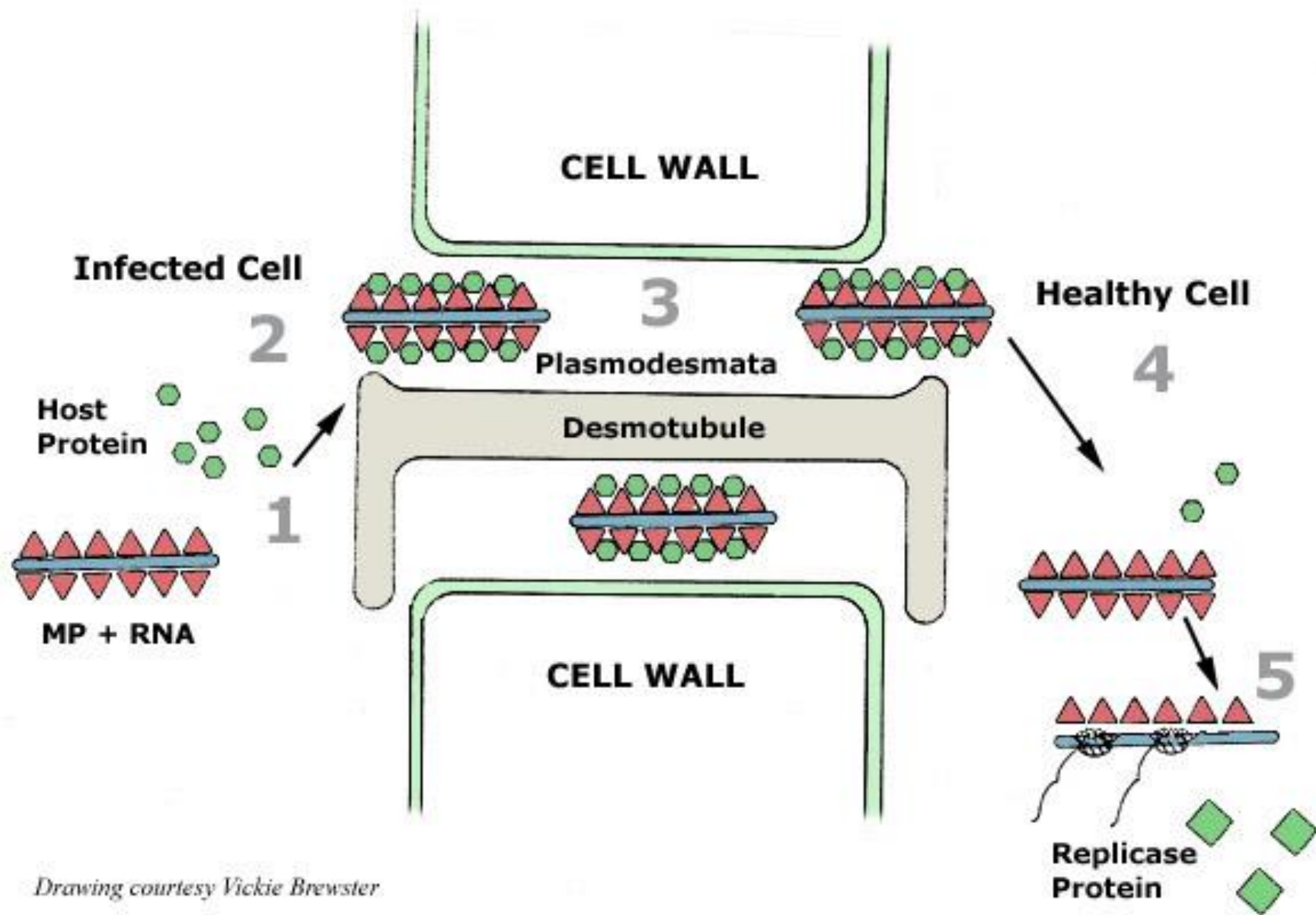
- Several unrelated lineages of MP proteins have been described
- MPs act as host range determinants
- MP alone causes expansion of normally constricted plasmodesmata pores; MPs then traffic through rapidly
- MPs are homologs of proteins that naturally traffic mRNAs between cells
- MPs may act as suppressors of gene silencing

Plant cells are bound by rigid cell walls and are interconnected by plasmodesmata, which are too small to allow passage of whole virus particles.



Plasmodesma





Drawing courtesy Vickie Brewster

Systemic spread of plant viruses is primarily through vascular tissue, especially phloem

