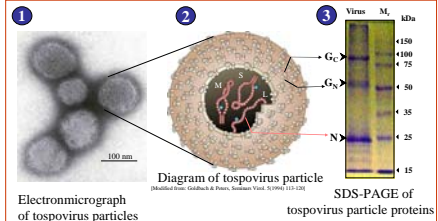

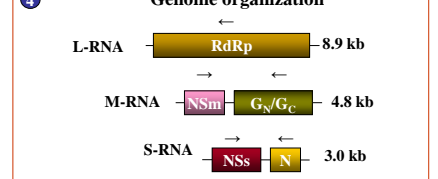



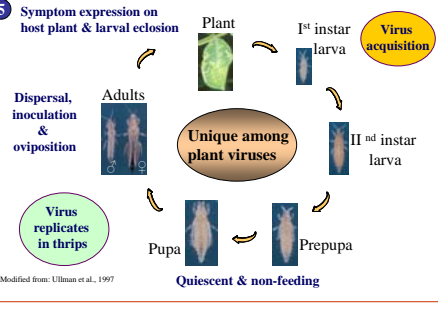


<h3>1. Tospoviruses</h3> <p>In recent years, tospoviruses (Type member: <i>Tomato spotted wilt virus</i>; Genus: <i>Tospovirus</i>; Family: <i>Bunyviridae</i>) have received international attention because they cause serious damage to agronomic crops, ornamentals, and vegetables worldwide. Tospoviruses replicate in two disparate hosts – plants and thrips. At least fifteen different tospoviruses have been characterized globally. There are about twelve species of thrips (<i>Thysanoptera: Thripidae</i>) that have been confirmed as vectors of one or more tospoviruses worldwide.</p> <p>Tospoviruses have complex genome and intricate replication strategy. The virus particles (Fig.1 & 2) contain three single-stranded genomic segments. Each segment is encapsidated separately with the nucleocapsid (N) protein to form helical structures and is associated with polymerase. The virus genome is contained within a host-derived lipid envelope containing virus-encoded glycoproteins (GPs), G_N and G_C. The two GPs appear as spike-like projections on the surface of the virions (Fig. 2). The three genomic segments codes for three structural and three non-structural proteins (Fig. 3 & 4).</p>	<h3>3. Tospoviruses: a global view</h3> <p>Phylogenetic tree based on N protein sequences</p> <p>Cluster I Primarily distributed in Americas</p> <p>Cluster II Primarily distributed in Eurasia</p> <p>TSWV = <i>Tomato spotted wilt virus</i> GRSV = <i>Groundnut ringspot virus</i> TCSV = <i>Tomato chlorotic spot virus</i> CSMV = <i>Chrysanthemum stem necrosis virus</i> ZLCV = <i>Zucchini leaflet chlorosis virus</i> INSV = <i>Impatiens necrotic spot virus</i></p> <p>PBNV = <i>Peanut bud necrosis virus</i> WSMV = <i>Watermelon silver mottle virus</i> WBNV = <i>Watermelon bud necrosis virus</i> CaCV = <i>Capiscum chlorosis virus</i> MYSV = <i>Melon yellow spot virus</i> IYSV = <i>Iris yellow spot virus</i> TYRV = <i>Tomato yellow fruit ring virus</i> PYSV = <i>Peanut yellow spot virus</i> PCFV = <i>Peanut chlorotic fupspot virus</i></p>	<h3>6. South & Southeast Asia region: a biological ‘Cuisinart’ for tospoviruses ?</h3> <p>Thrips-borne tospoviruses are emerging as a significant limiting factor in the sustainable production of vegetables and other economically important crops in smallholder farming systems in the region</p> <p>South Asia</p> <ul style="list-style-type: none"> 60% of ‘global’ tospoviruses occur in the region Four distinct tospoviruses documented in India: PBNV, WBNV, IYSV and PYSV Three distinct tospoviruses documented in Thailand: CaCV, WSMV and MYSV Tospoviruses are more diverse in India Status of tospoviruses in other countries not known <p>Southeast Asia</p> <p>50% of known vector thrips species present in the region</p> <p><i>Thrips palmi</i> <i>T. tabaci</i> <i>Scirtothrips dorsalis</i> <i>Frankliniella occidentalis</i> <i>F. schultzei</i> <i>Ceratothrips claratris</i></p>
 <p>Electronmicrograph of tospovirus particles</p> <p>Diagram of tospovirus particle</p> <p>SDS-PAGE of tospovirus particle proteins</p>	<p>(Source: Hassani-Mehraban et al 2005 - Phytopathology 95: 852-858)</p> <p>All tospoviruses characterized so far fall into two major groups indicating geographic clustering.</p> <p>However, there are certain exceptions: TSWV is present outside Americas – in Europe, Africa, Australia and Japan. IYSV, reported initially from Europe and Israel, is present in the Americas, Australia and India.</p> <p>Tospoviruses are ‘generalists’ because they have wide host range. They are mostly ‘opportunists’ because they survive by infecting many species that include crops and weeds, invade new geographic areas and quickly exploit suitable ecological ‘niches’.</p>	<h3>7. Tospoviruses are a threat to sustainable production of vegetables in India</h3> <p>PBNV infects a broad range of crops</p>  <ul style="list-style-type: none"> Second largest producer of vegetables in the world - estimated production of about 50.09 million tons from an area of 4.5 million hectares at an average yield of 11.3 tons/hectare With 2 % of the world’s geographical area, India is home for 16 % of global human population and over a quarter of the world’s poor people Vegetables constitute the most important food next only to cereals and milk The demand for vegetables has been rapidly increasing due to urban population growth at more than 3% per annum Consequently, peri-urban farming of vegetables, which account for nearly 95% of peri-urban vegetable production, has intensified Smallholder farmers cultivate their land throughout the year to produce vegetables Intensive and widespread use of pesticides is the predominant tactic farmers deploy to manage virus diseases The number and extent of vegetable crops affected by tospoviruses increasing
<h3>4. Genome organization</h3>  <p>L-RNA → RdRp → 8.9 kb</p> <p>M-RNA → NSm, G_N/G_C → 4.8 kb</p> <p>S-RNA → NSs, N → 3.0 kb</p>	<h3>4. Expansion of geographic range of vector thrips</h3> <p>Western flower thrips (<i>Frankliniella occidentalis</i>)</p>  <ul style="list-style-type: none"> A native to the southwestern USA Spread through global trade in ornamental greenhouse plants around the world since 1980s. <p>Melon thrips (<i>Thrips palmi</i>)</p>  <ul style="list-style-type: none"> A native to Southeast Asia Expanded its geographic range in 1970s and 1980s through trade and commerce ? <p>(Source: www.epi.org)</p>	<h3>Yield losses due to PBNV</h3>  <p>Healthy Infected</p> <p>Crop-specific problems</p> <p>WBNV in watermelon</p> <p>IYSV in onion</p>
<h3>2. Tospoviruses: vector transmission</h3> <p>Thrips-mediated transmission of tospoviruses is closely linked to the developmental stages of the thrips on plants (Fig. 5). Successful transmission by adult thrips occurs only when the virus is acquired at the first-instar larval stage of the thrips life cycle. Since virus replicate in the vector, adult thrips, that acquire a tospovirus during the larval stage, remain viruliferous throughout their life. Thus, adult thrips contribute to short- and long-distance spread of the virus. Males exhibit a higher transmission rate than females.</p>	<h3>5. Challenges in controlling tospovirus diseases</h3> <p>Tospoviruses</p> <ul style="list-style-type: none"> broad host range multiple vector species evolution of new strains ability to overcome host plant resistance <p>Vector thrips</p> <ul style="list-style-type: none"> polyphagous & show habitat infidelity has superior reproductive output has propensity to ‘overwinter’ on a broad range of plant species develop resistance against pesticides 	<h3>8. Management of tospoviruses requires a multidisciplinary team effort</h3> <ul style="list-style-type: none"> Tospoviruses have complex genome, replicate in plants and vectors and produce a range of symptoms The life cycle of a tospovirus involves transmission from plant to plant by several species of polyphagous thrips Efforts to control vector thrips with insecticides have been mostly unsuccessful Durable resistance difficult due to rapid evolution of virulent and resistance-breaking strains of tospoviruses A comprehensive understanding of tospovirus pathosystem is critical Multidisciplinary team effort can bring long-lasting solutions for the management of diseases caused by tospoviruses
<h3>5. Symptom expression on host plant & larval eclosion</h3>  <p>Dispersal, inoculation & oviposition</p> <p>Adults</p> <p>Plant</p> <p>Ist instar larva</p> <p>Virus acquisition</p> <p>Unique among plant viruses</p> <p>IInd instar larva</p> <p>Pupa</p> <p>Prepupa</p> <p>Quiescent & non-feeding</p> <p>Virus replicates in thrips</p> <p>Modified from: Ullman et al., 1997</p>	<h3>The way forward</h3> <p>Objectives of the project funded by IPM-CRSP of USAID</p> <ul style="list-style-type: none"> Conduct strategic research on tospoviruses and thrips vectors Carryout applied and adaptive research to deploy ‘eco-friendly’ IPM strategies to control tospovirus diseases Develop strategies for strengthening institutional capacities within host countries to conduct problem-oriented research on virus diseases 	