
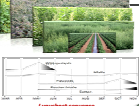

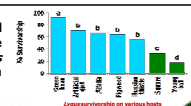

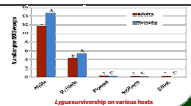

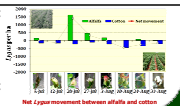

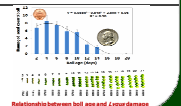

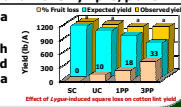

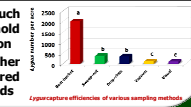

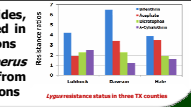

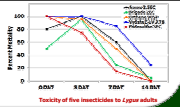

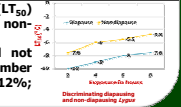

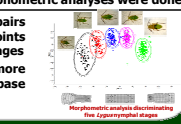
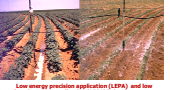
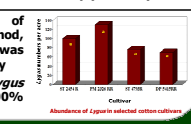

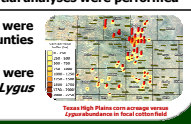

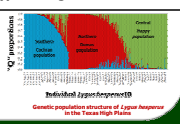

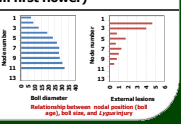




Overview of *Lygus* Research and Outreach Program in the Texas High Plains: Serving the Clientele of the World's Largest Cotton Patch

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Introduction: Texas A&M Cotton Entomology research in the Texas High Plains began in 1937, but the focus on *Lygus* research was started in 2002. Our program has used *Lygus* as the model insect to answer various ecological questions in cotton pest management. In the past eleven years (2002-2012), more than 20 research projects were conducted on various aspects of *Lygus* biology, behavior, and ecology. Experiments were conducted in the laboratory, greenhouse, research farms, and growers' fields. *Lygus* research conducted in our program includes host-plant survey, life table analysis, host preference, intercrop movement, feeding biology, cotton plant/*Lygus* interactions, sampling, insecticide resistance, pesticide evaluations, overwintering biology, morphology, molecular ecology, cultural control, landscape structure, and economic threshold development.

<h3>Host Plant Survey</h3>  <ul style="list-style-type: none"> Two years (2002-03) of general survey of 25 counties using sweepnet Two additional years (2004-05) of thorough three-county survey <p><i>L. hesperus</i> was most dominant <i>Lygus</i> species (>95%)</p> <ul style="list-style-type: none"> Twenty-five reproductive plant hosts were identified, and Texas High Plains <i>Lygus</i> host-plant sequences were established 	<h3>Life Table Study</h3>  <ul style="list-style-type: none"> <i>L. hesperus</i> were reared on six food sources from egg to adult in the laboratory  <ul style="list-style-type: none"> Lowest survivorship and longest egg-to-adult life cycle were found for <i>Lygus</i> reared on 1-day old cotton bolls 	<h3>Host Preference Study</h3>  <ul style="list-style-type: none"> Two-year (2004-05) field study was conducted evaluating <i>Lygus</i> host preference in a mosaic of four prevalent weed hosts (alfalfa, Russian thistle, pigweed, sunflower) plus cotton  <ul style="list-style-type: none"> <i>Lygus</i> colonization was highest in alfalfa followed by Russian thistle Reproduction correlated with colonization Cotton was not the preferred host for <i>Lygus</i> 	<h3>Intercrop Movement Study</h3>  <ul style="list-style-type: none"> Two-protein mark-capture study, using milk and egg, examined <i>Lygus</i> intercrop movement in an alfalfa-cotton system (2008-09) Marker proteins were detected using ELISA  <ul style="list-style-type: none"> <i>Lygus</i> bidirectional movement suggests alfalfa serves both as source and sink depending on host phenological stages Increased alfalfa to cotton movement was observed at cotton blooming stage 	<h3>Damage Potential Study</h3>  <ul style="list-style-type: none"> Cup-cage studies quantified <i>Lygus</i> damage potential to age-specific cotton bolls (2008-11) Single <i>Lygus</i> adults were allowed to feed 48 h on cup-caged bolls at various maturity levels  <ul style="list-style-type: none"> <i>Lygus</i> injury caused external lesions in bolls at all maturity levels Bolls >25 mm dia or >350 heat units were safe from <i>Lygus</i> injury to cotton seed
<h3>Cotton Crop Response</h3>  <ul style="list-style-type: none"> Numerous studies quantified cotton yield compensation of <i>Lygus</i>-induced fruit loss under various production scenarios Four treatments were spray control, untreated control, and 1 or 3 nymphs/plant  <ul style="list-style-type: none"> <i>Lygus</i> treatments caused a "stair-case" injury effect Plants compensated as high as 33% <i>Lygus</i>-induced square loss, typically via adding more lateral nodes 	<h3>Sampling Method</h3>  <ul style="list-style-type: none"> Five <i>Lygus</i> sampling methods were compared (2002-2003) <i>Lygus</i> capture rates and sampling efficiencies were established for each method  <ul style="list-style-type: none"> <i>Lygus</i> densities were much below economic threshold levels throughout the season Beat bucket captured higher numbers of <i>Lygus</i> compared with other sampling methods 	<h3>Insecticide Resistance</h3>  <ul style="list-style-type: none"> Spatial and temporal variations in pesticide resistance levels in <i>Lygus</i> were studied (2005-06) Four insecticides were evaluated using 20 ml glass vial bioassay  <ul style="list-style-type: none"> Among the tested insecticides, resistance was not detected in Texas High Plains populations Acephate LC₅₀ for <i>L. hesperus</i> increased continuously from April to November populations 	<h3>Pesticide Evaluation</h3>  <ul style="list-style-type: none"> Five insecticides were evaluated at high doses <i>Lygus</i> were exposed to treated leaves for 48 h <i>Lygus</i> mortalities were determined by Petri dish bioassay  <ul style="list-style-type: none"> Among five insecticides evaluated, Vydate and Endosulfan were most and least toxic to <i>Lygus</i> adults Vydate had the longest residual activity 	<h3>Overwintering Biology</h3>  <ul style="list-style-type: none"> Dig-up cages and cold-chambers were used to determine the <i>Lygus</i> winter survivorship and spring emergence profile Field collected and laboratory reared <i>Lygus</i> were used for cold tolerance (cold-bath) study  <ul style="list-style-type: none"> Lethal temperatures (LT₅₀) discriminated diapausing and non-diapausing <i>Lygus</i> October released cohorts did not survive the winter, but November released cohorts survived at 12%; emergence peaked in February
<h3>Morphometric Study</h3>  <ul style="list-style-type: none"> <i>Lygus hesperus</i> were reared on green beans from egg to adult Ontogenic digital images of each live insect were taken daily Images were digitized and morphometric analyses were done  <ul style="list-style-type: none"> Measurements between 4 pairs of specific landmark points differentiated 5 nymphal stages Landmark technique was more efficient than outline base ontogenic shape analysis 	<h3>Cultural Control</h3>  <ul style="list-style-type: none"> Five cotton cultivars, two irrigation methods (LEPA and LES), and three water levels (50, 75, and 100% ET) were evaluated to assess <i>Lygus</i> field activity (2002-03)  <ul style="list-style-type: none"> No significant effect of cultivar, irrigation method, or irrigation level was detected on <i>Lygus</i> activity Marginally higher <i>Lygus</i> detected in LEPA and 100% ET replacement plots 	<h3>Landscape Level Study</h3>  <ul style="list-style-type: none"> <i>Lygus</i> were monitored in 50 focal cotton fields in the Texas High Plains for three years (2008-10) Habitats within 2-mile radius of each focal field were mapped and spatial analyses were performed  <ul style="list-style-type: none"> <i>Lygus</i> abundances in cotton were higher in northern counties (Castro, Swisher, and Hale) Corn and alfalfa acreages were positively correlated with <i>Lygus</i> abundance in cotton 	<h3>Genetic Diversity Study</h3>  <ul style="list-style-type: none"> <i>Lygus hesperus</i> microsatellite markers were developed by sequencing SSR enriched partial genomic DNA library Eight <i>Lygus</i> populations were genotyped using 10 SSR markers  <ul style="list-style-type: none"> Ten SSR markers were identified and characterized Molecular marker analysis (SSR) identified 3 distinct genetic populations of Texas High Plains <i>Lygus</i> 	<h3>Economic Threshold</h3>  <ul style="list-style-type: none"> Four <i>Lygus</i> densities (0, 1, 2, 4 bugs/plant) were evaluated ET is being developed for two distinct cotton phenological stages (0-350 and 0-550 HU from first flower)  <ul style="list-style-type: none"> <i>Lygus</i> adults displayed preference for smaller bolls (<20 mm diameter) and the top 3-4 mainstem nodes This is the first year study of a multi-year ongoing project

Summary: Our programmatic effort on *Lygus* research has resulted in some significant outcomes for further scientific investigation and for greater adoption, including alternate host identification, characterization of feeding and movement biology, pesticide spray initiation and termination rules, molecular marker development, determination of genetic structure, pesticide resistance monitoring, morphological characterization, life table investigation, and crop protection product evaluation. The Cotton Entomology Program has contributed significantly to assisting Texas cotton producers, crop consultants, Extension agents, and the scientific community by expanding our cotton pest management knowledge and skills through research and outreach.

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