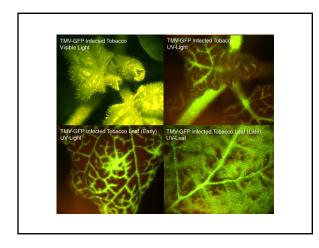
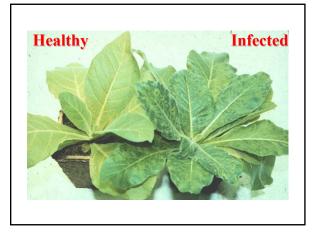
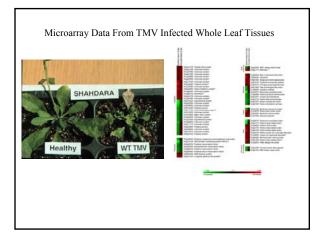
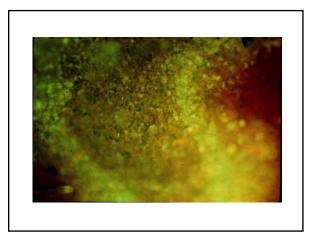


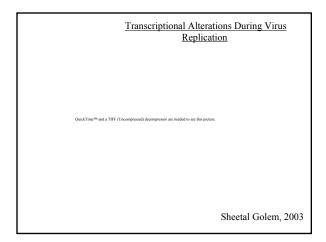
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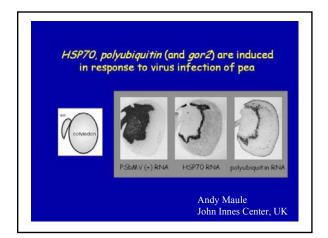


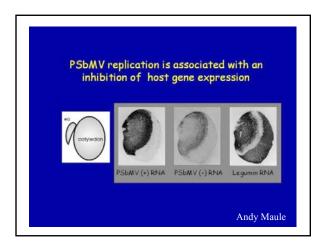


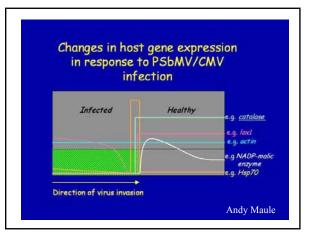


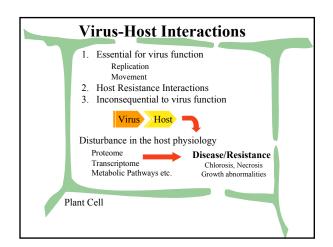


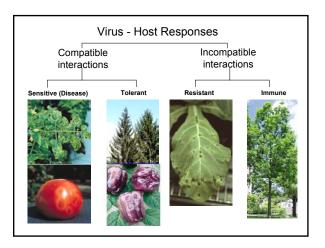


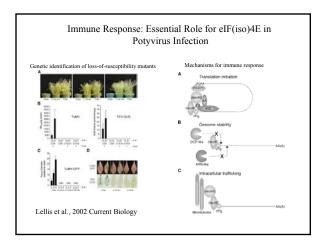


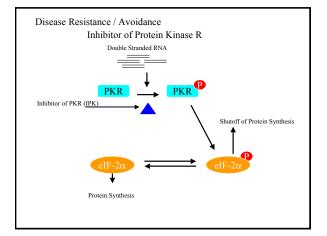


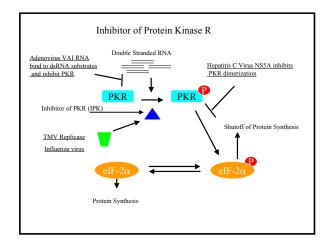


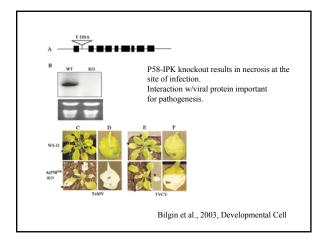


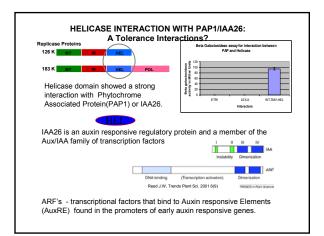


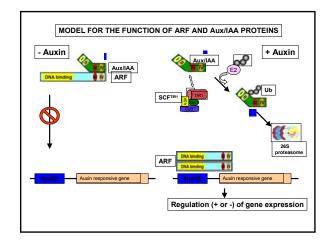


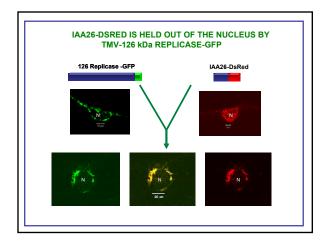




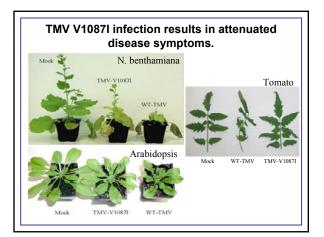


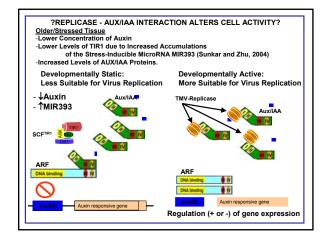


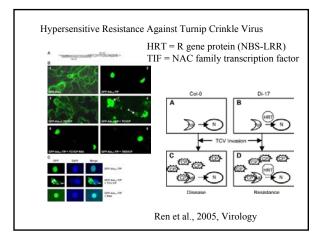




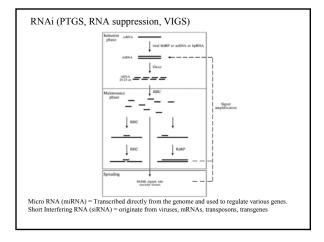
(Golem et	/ data identified 68 Ara al. 2003)	bidopsis gene	s altered in infect	ed tissue				
 Promoter analysis of 2 kb region upstream of each gene to identify Aux RE's TGTCTC element – ARF binding site 								
 20 genes contained 2 or more AuxRE's. ~ 30% of the genes displaying transcriptional alterations to TMV infection may be linked to auxin response system Fold expression 								
	Protein name	Number of	In TMV infection	+50µM IAA				
		AuxRE's	(from microarray)	(qRT-PCR derived)				
	Put protein	3		(qRT-PCR derived) -4.09				
Model ID	Put protein Unk protein		(from microarray)	,				
Model ID At5g02160		3	(from microarray) -1.9	-4.09				
Model ID At5g02160 At1g19350	Unk protein	3 3	(from microarray) -1.9 -2.0	-4.09 -1.8				
At1g19350 At3g17790	Unk protein Acid Phos'tase type 5	3 3 2	(from microarray) -1.9 -2.0 -2.3	-4.09 -1.8 -6.6				

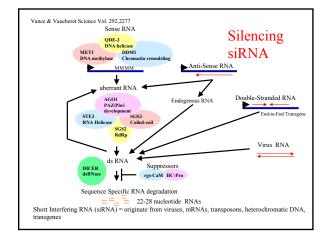












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