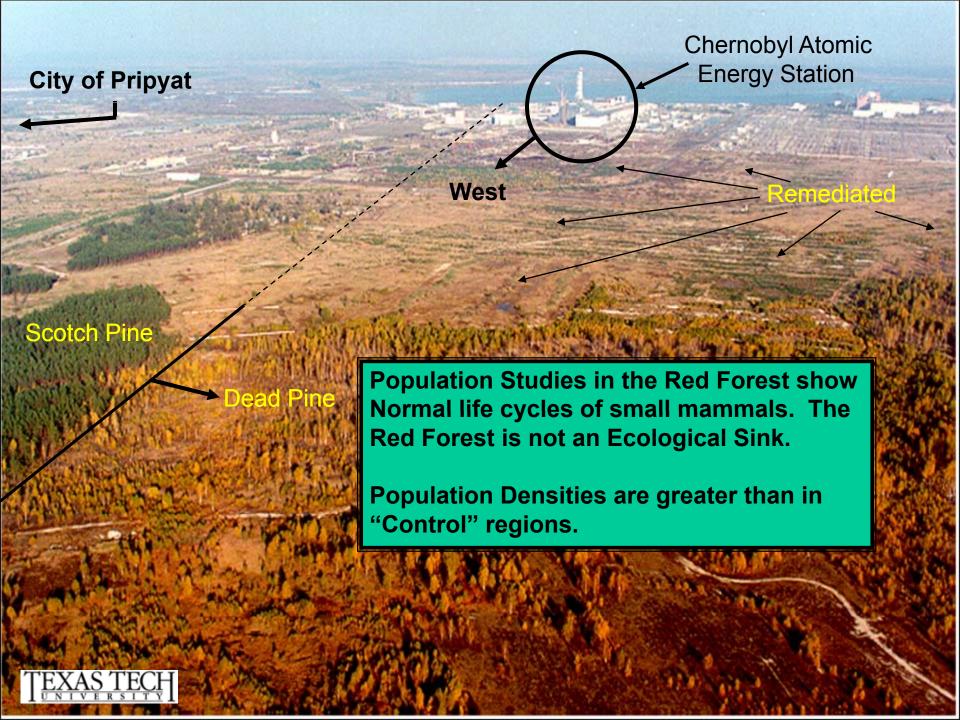


#### Overview

- Brief History of Studies at Chernobyl
- Specific Aims of Radioadaptive Experiments
- Experimental Design
- Results
- New Directions

#### Bank vole – the All-time radiation winner!





#### Overview

- Brief History of Studies at Chernobyl
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 To characterize transcriptional responses as a function of dose, dose rate and duration of exposure

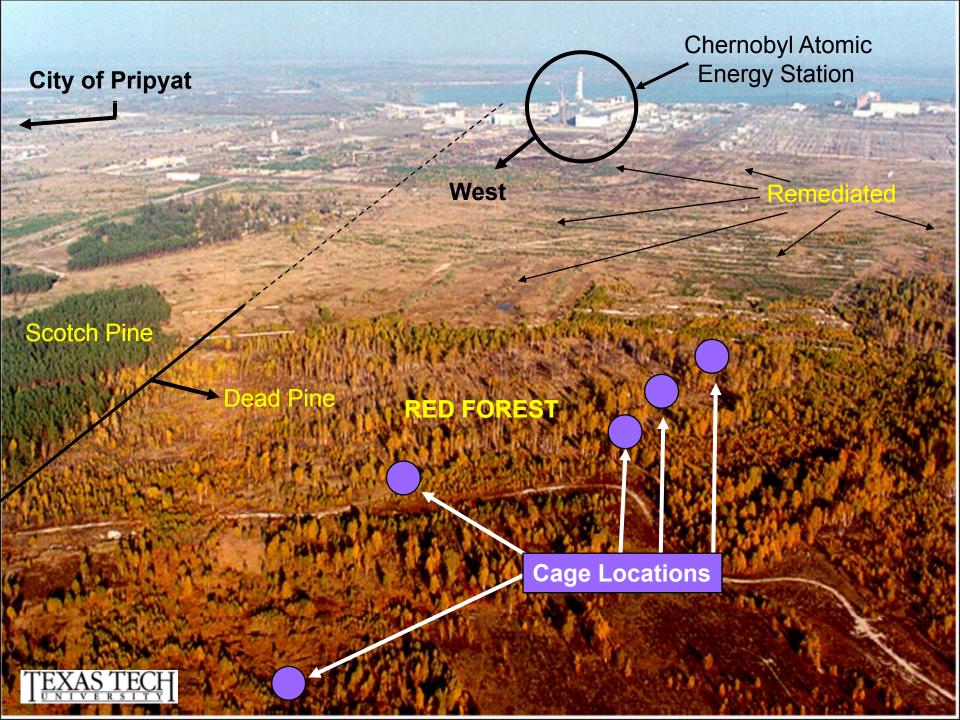
 To investigate radio-adaptive response in inbred strains of mice continuously exposed to the Chernobyl environment

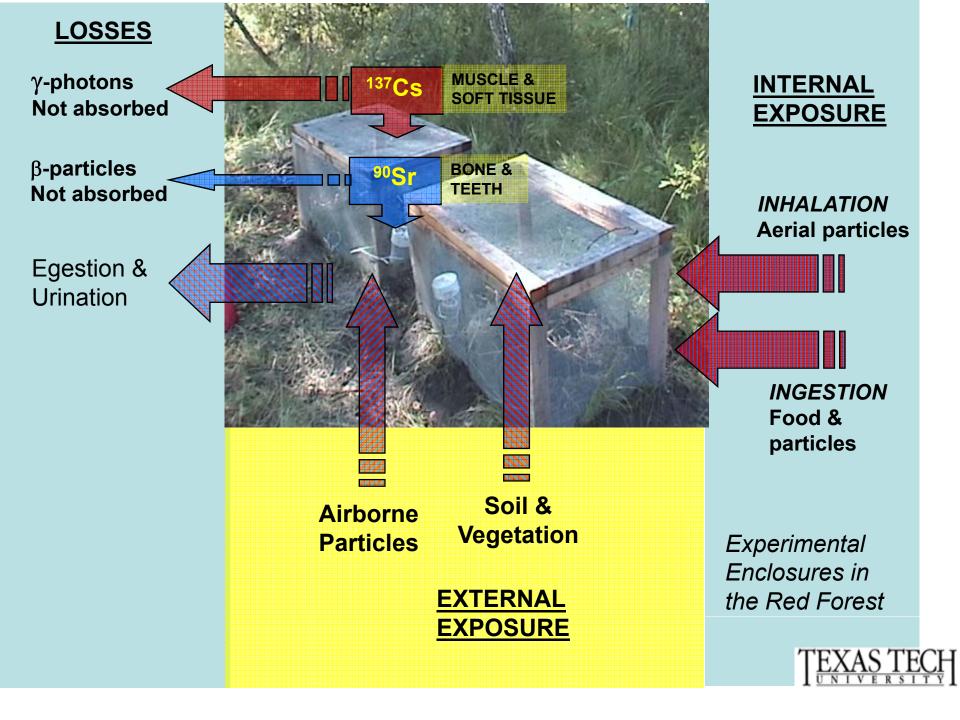
 To investigate variation in response as a function of priming dose, dose rate and duration of exposure

To archive tissues and nucleic acids as a resource for future studies

#### Overview

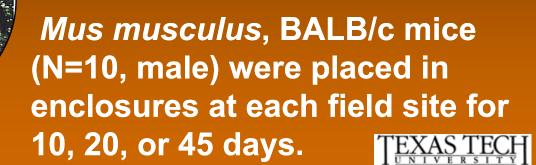
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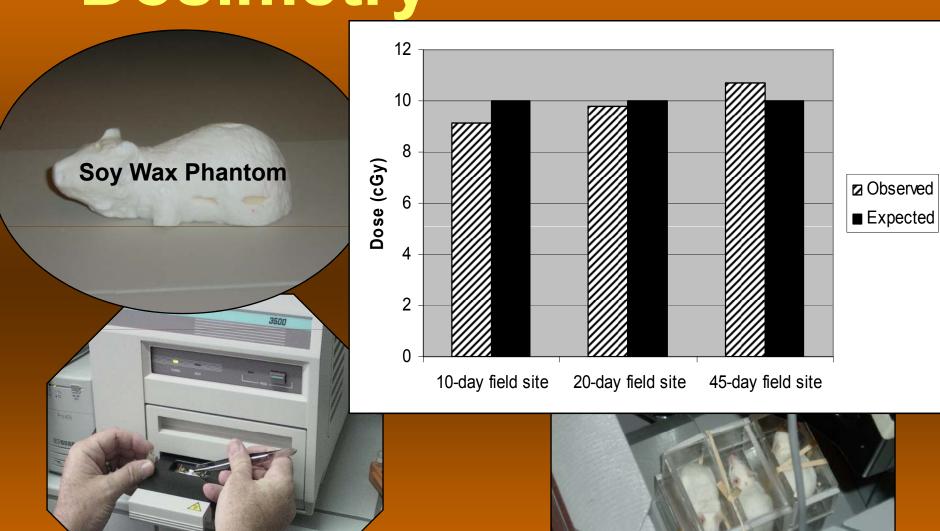


# **Experiments & Environmentally Relevant**Radiation Exposure Groups

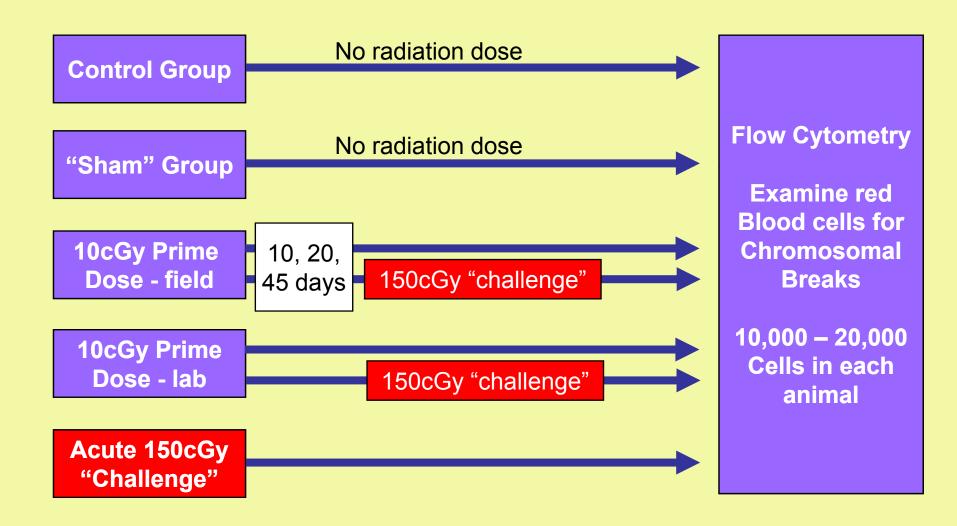
Areas in the Red Forest were identified in which a total absorbed dose of 10 cGy could be achieved.



# Dosimetry

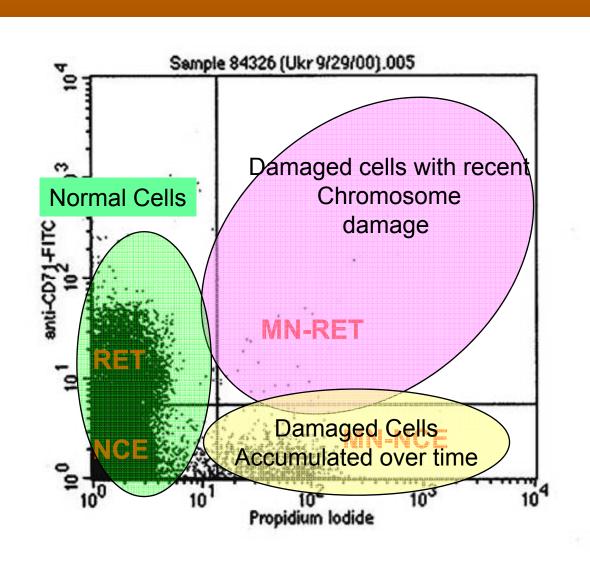


#### BALB/c Experimental Design





#### MicroFlow® micronucleus assay



Enumerates MN in peripheral blood reticulocytes

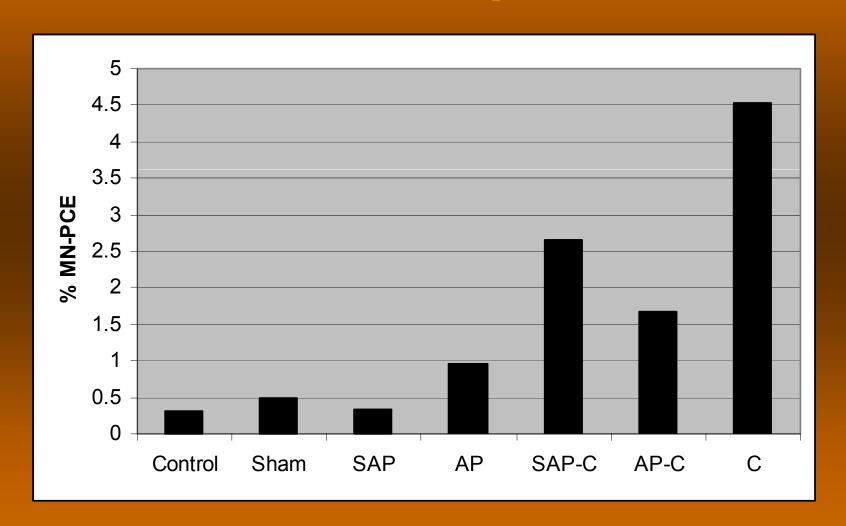
 10-20,000 reticulocytes analyzed per blood sample



#### **Overview**

- Define Radioadaptive Response
- Units of Radioactivity
- Objectives of Radioadaptive Experiments
- Experimental Design
- Results
- New Directions

### **MN-RET** frequencies

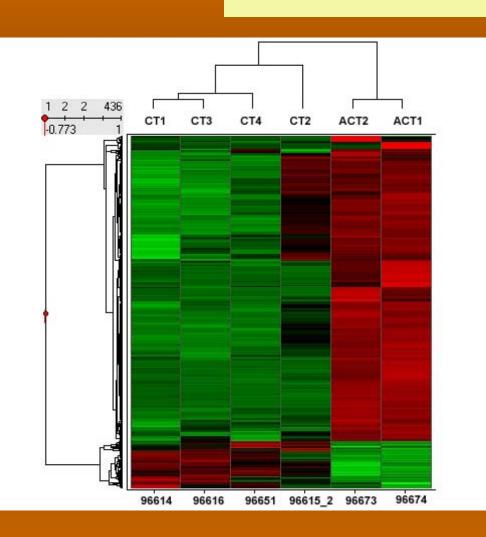




# Expected and Observed Mean MN-RET Frequencies

Group	Expected % MN-RET	Observed % MN-RET	Reduction
SAP + 1.5 Gy	4.86	2.65	45%
AP + 1.5 Gy	5.49	1.68	69%

# **Gene Chip Analysis**

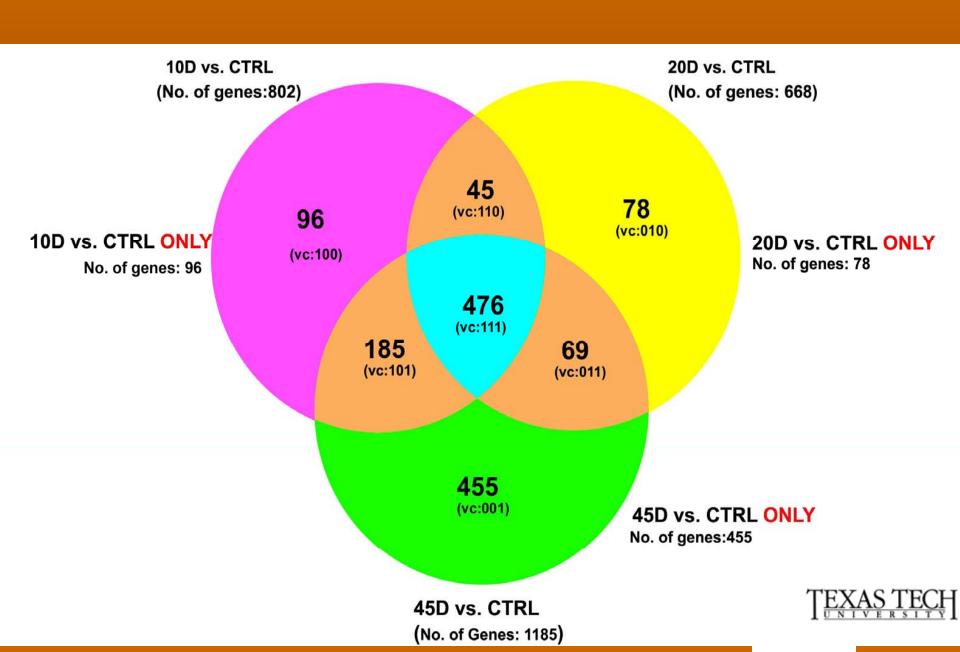


Heat Map comparing controls to 45d exposed

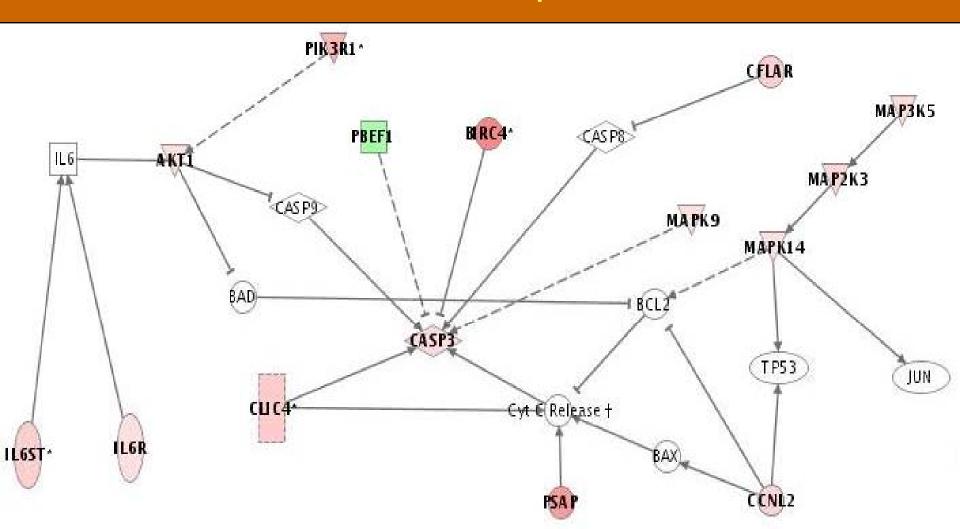
**Total RNA wa** isolated from liver, RNA from each mouse was then hybridized to its own **Affymetrix GeneChip** Mouse Genome 430 2.0 Array®



#### Differentially Expressed Genes



Biological network suggested by IPA ver. 4.1 in which several genes products involved in apoptosis are present. Many of these are differentially transcribed in all exposed animals in comparison to unexposed animals thus possibly representing a signature of low-dose rate exposure.



# Future Directions for Chernobyl Research

- Principal Components Analysis (PCA) on genomics datasets.
- Transcriptomic analysis from skeletal muscle tissue (a subset of the same mice used in the liver analyses).
- Real-time PCR array (n = 41 transcribed genes) Molecular Genomics Core facility (UTMB) to validate a subset of genes in the genechip arrays.



# Future Directions for Chernobyl Research

- Genetic Susceptibility
- In utero exposures
- MN origin
- SEM for Metals Analysis
- Evaluate trans-generational responses (mutational load)



# Acknowledgments

- William Osorio, Kristen Holmes, Adam Brown, Heather Meeks
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- International Radioecology Laboratory, UA



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