

# Properties of Protein Targets of Reactive (Lipid) Species

Dennis Petersen, PhD

Department of Pharmaceutical Sciences

School of Pharmacy

University of Colorado Denver

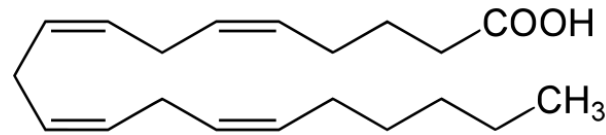
Anschutz Medical Campus

[dennis.petersen@ucdenver.edu](mailto:dennis.petersen@ucdenver.edu)

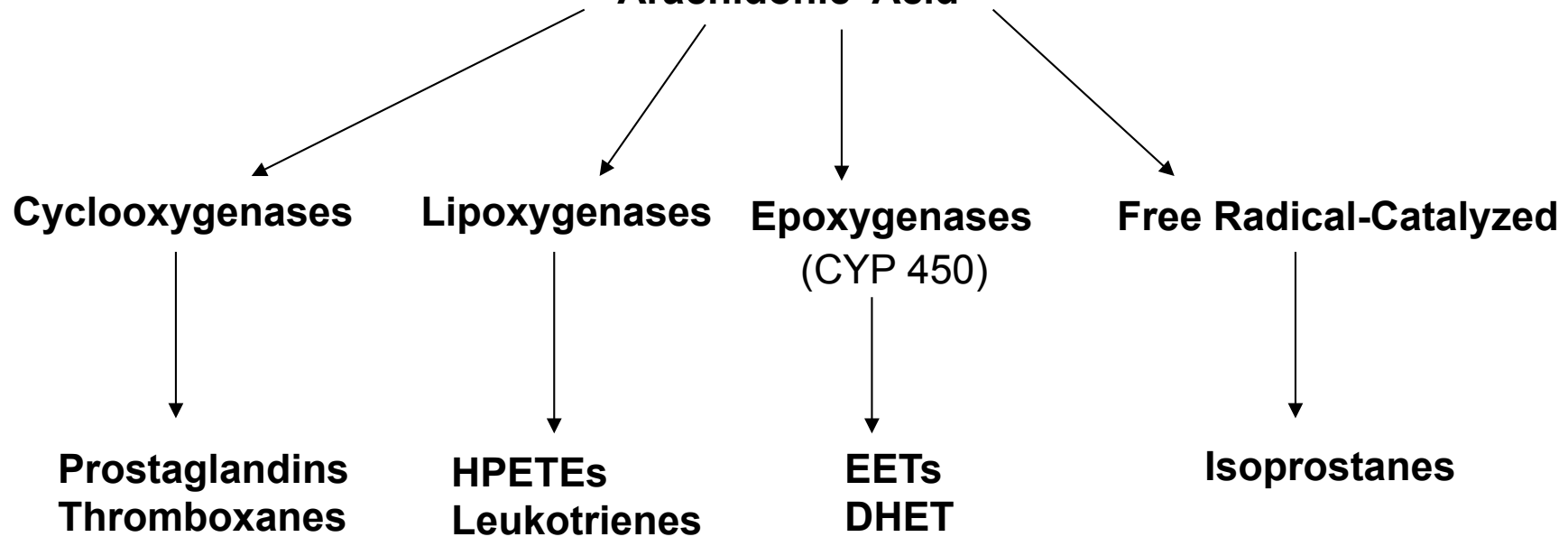
# Overview of Presentation

- Sources and characteristics of reactive lipid species (electrophiles)
- Characteristics of protein target residues (nucleophiles)
- Survey of relative rates of electrophile-nucleophile reactions
- Detection and Identification of post-translational protein modifications by lipid electrophiles
- Electrophile-protein modification: functional and structural characterization
- Limitations and challenges

# Reactive Lipid Species: The Eicosinoids



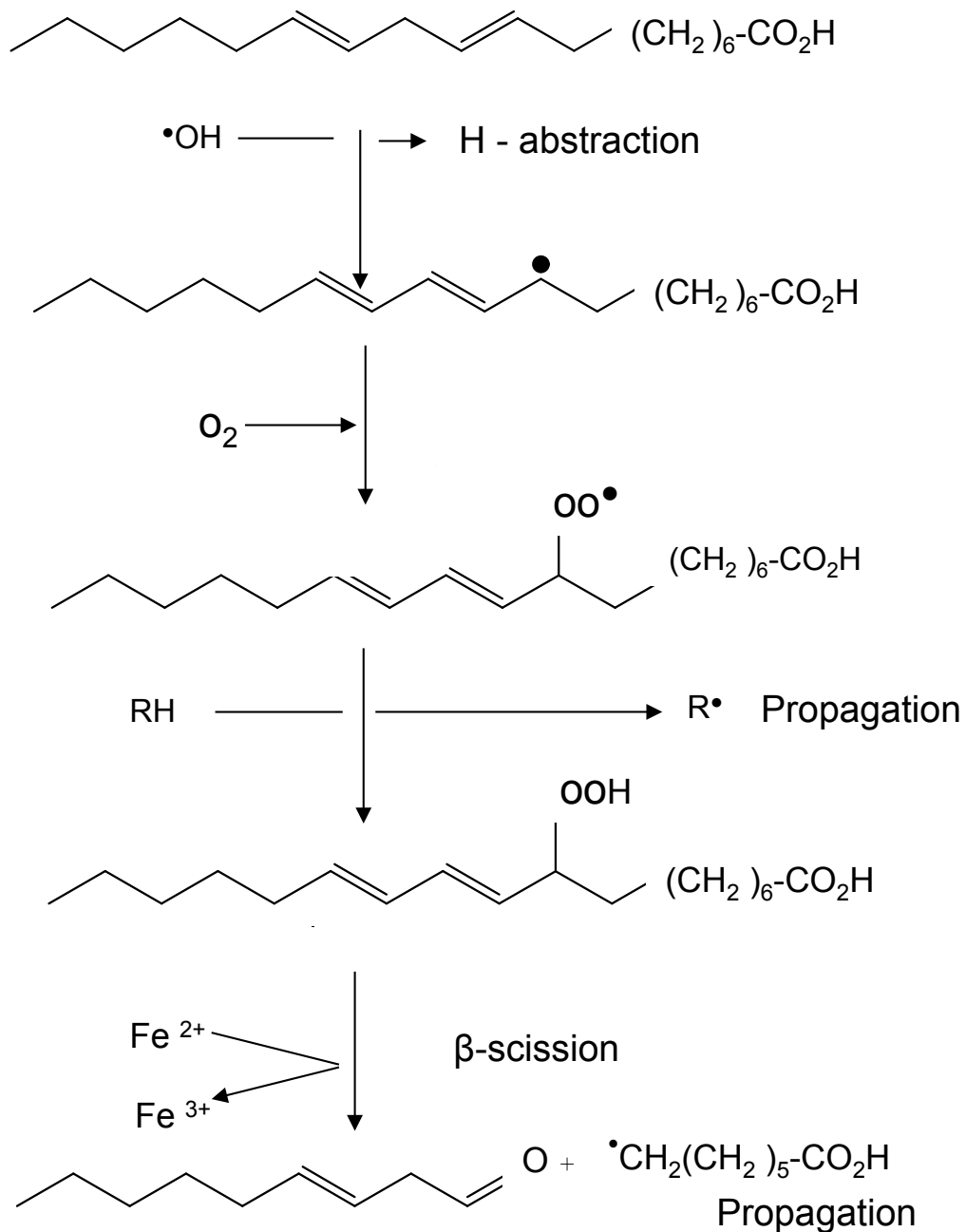
**Arachidonic Acid**



**Characteristics:** Biosynthetic pathways are in response to specific stimuli and are well-regulated

Bioactivities are diverse, pronounced and mediated *via* specific receptors

# Endogenous Generation of Lipid Peroxidation Products

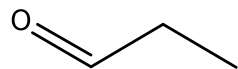




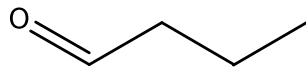
# Endogenous Lipid Peroxidation Products

## Saturated Alkanals

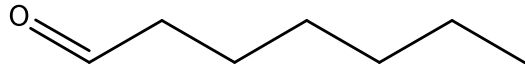
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Propanal



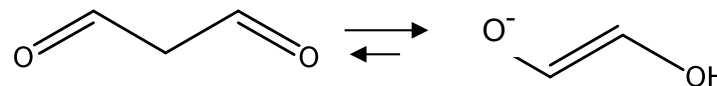
Butanal



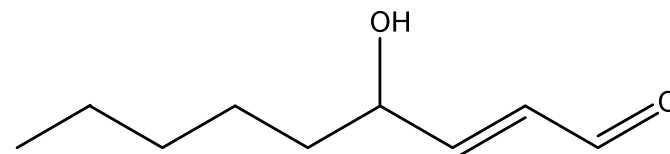
Heptanal

## Alkenals and Hydroxyalkenals

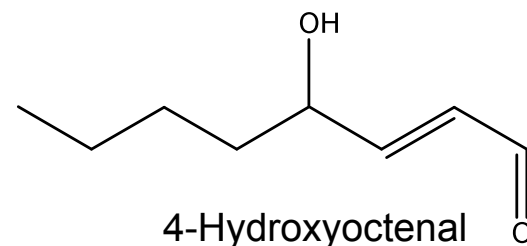
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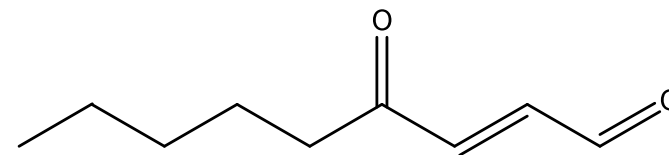
Malondialdehyde



4-Hydroxynonenal



4-Hydroxyoctenal



4-Oxononenal

Esterbauer et al. *Biochem. J.* 208:129, 1982

Poli, et. al. *Biochem. J.* 227:629, 1985

Lee & Blair, *Science* 292:2083, 2001

## **Endogenous Lipid Peroxidation Products (cont.)**

Characteristics: Generated endogenously and stimulated by oxidative stress

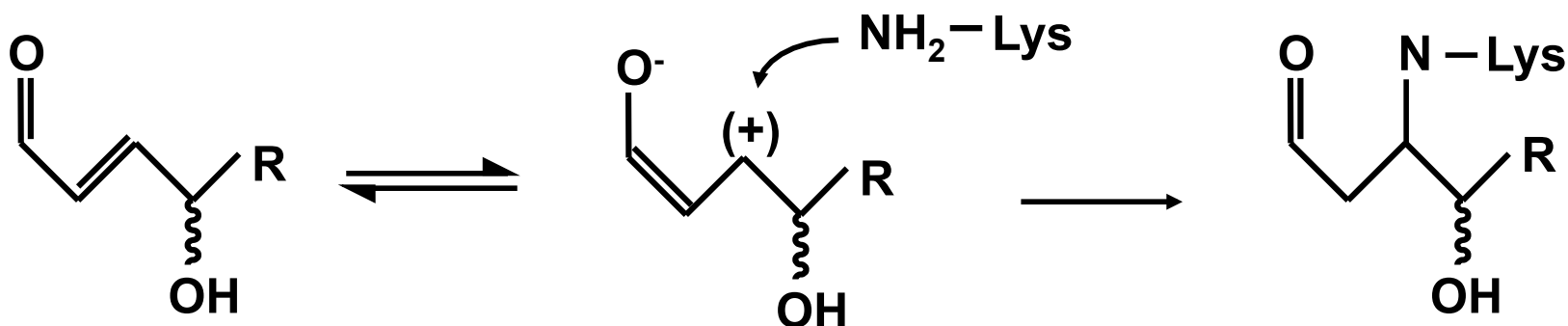
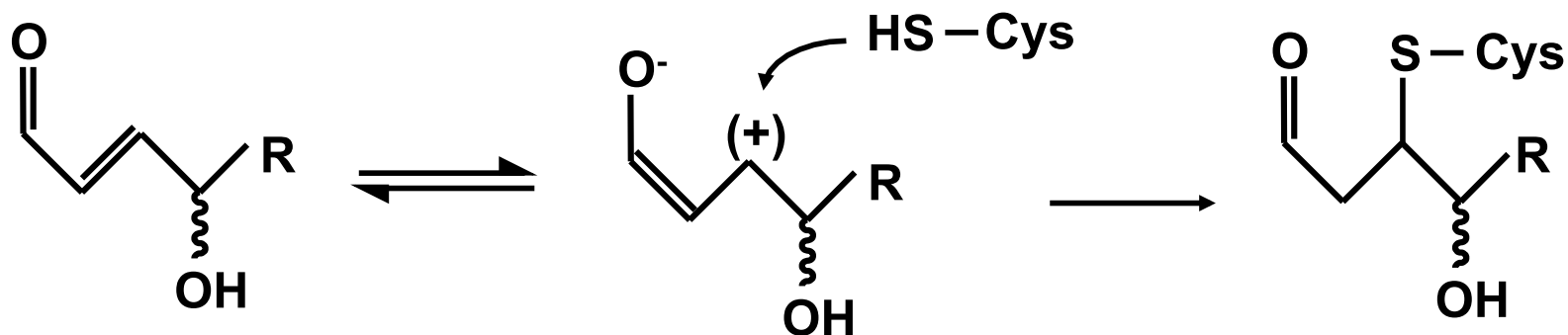
Cascades of generation are non-enzymatic and autocatalytic

Bioactivities/toxicities are diverse and mediated via interactions with proteins containing highly reactive nucleophilic amino acid residues

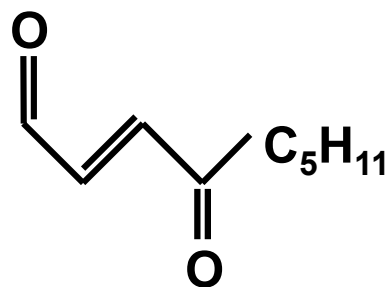
# The Lipid Peroxidation Product 4-Hydroxynonenal

## 4-HNE as a Michael Acceptor

R: C<sub>5</sub>H<sub>11</sub>

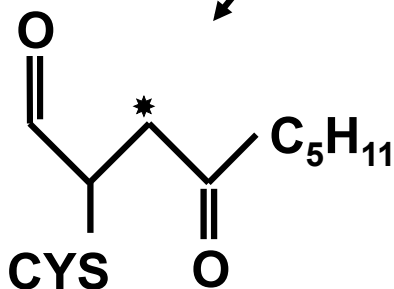


# Proposed Interactions of 4-ONE with Proteins



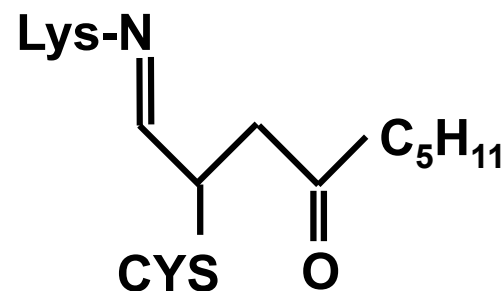
4-ONE

*Protein Cys-adduct formation*



4-ONE-Cys Michael Adduct

*Intra – or interpeptide  
cross-linking*



4-ONE-Cys Michael Adduct followed  
by formation of Lys-Schiff base

## Rate Constants of 4-ONE and 4-HNE with Peptides

Compound	$k(\text{M}^{-1}\text{s}^{-1})$		Ratio 4ONE/4HNE
	4-ONE	4-HNE	
NAC	$186 \pm 29$	$1.2 \pm 0.03$	153
GSH	$145 \pm 10$	$1.3 \pm 0.05$	109
Ac-Lys-amide	$7 \pm 0.6 \times 10^{-3}$	$1.3 \pm 0.08 \times 10^{-3}$	6
Ac-His-amide	$2 \pm 0.17 \times 10^{-2}$	$2.1 \pm 0.30 \times 10^{-3}$	10

Overall Reactivity  $\longrightarrow$  Cys  $\gg$  His  $>$  Lys

**4-HNE hepatocellular  $t_{1/2} = \sim 2$  min**

**4-ONE Hepatocellular  $t_{1/2} = < 1$  sec**

Doorn and Petersen *Chem Res Tox* 15:1445, 2002  
LoPachin et al. *Chem Res Tox* 22:1499, 2009

# Hepatocellular Biotransformation of 4-Hydroxynonenal

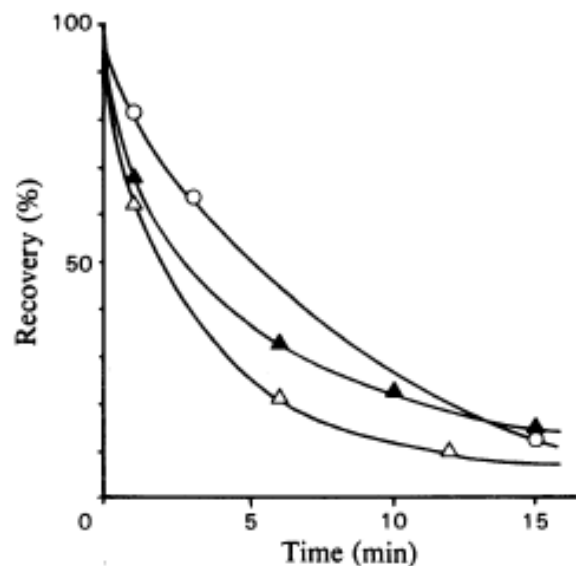


Fig. 4. Kinetics of the metabolism of 4-hydroxyalkenals by isolated rat hepatocytes

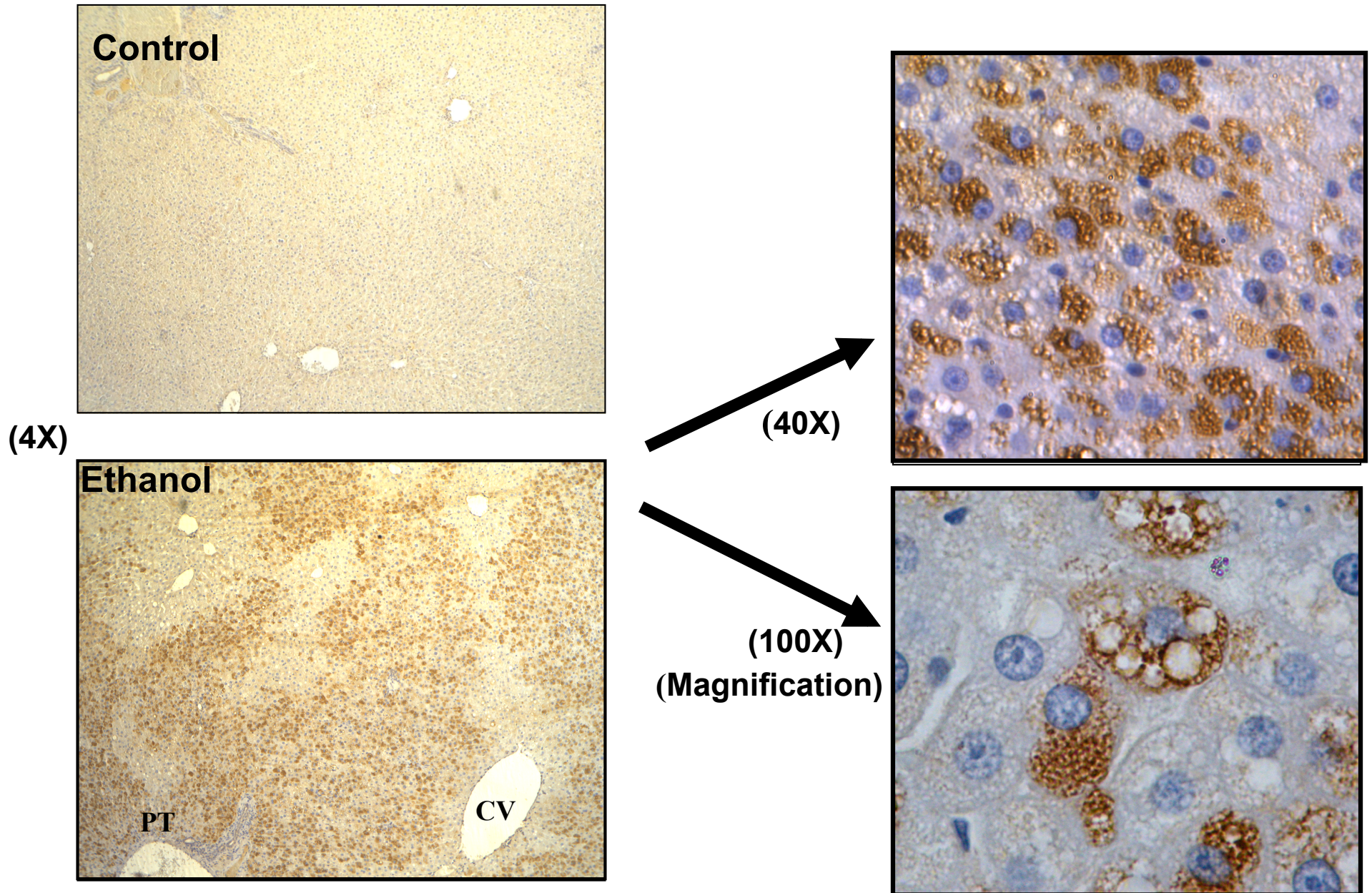
○, 4-Hydroxypent-2-enal; ▲, 4-hydroxyoct-2-enal; △, 4-hydroxynon-2-enal. Hepatocytes ( $0.2 \times 10^6$  cells/ml) were incubated in the presence of each aldehyde (0.1 mM), and the disappearance of the aldehyde was monitored by h.p.l.c.

Poli, et. al. *Biochem. J.* 227:629,1985

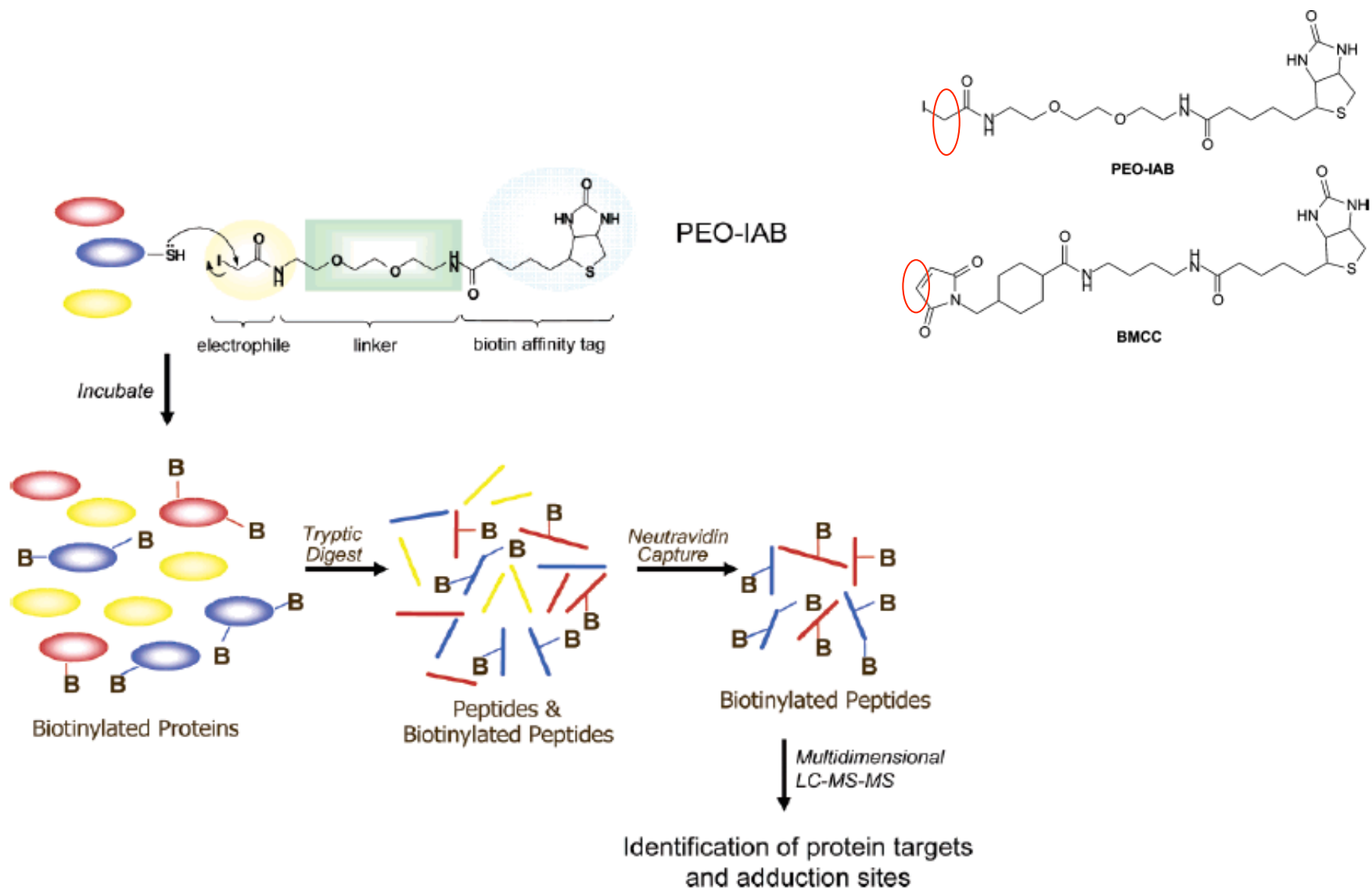
Bioconversion Pathways	Overall Contribution
Alcohol/aldehyde dehydrogenase	10%
GSH/glutathione transferases	70%
Unidentified	20%

Hartley et al. *Arch Biochem Biophys* 316:197, 1995

# 4-HNE Protein Adduct Formation In Liver of Rats Chronically Consuming Alcohol



# Cellular Protein Targets of Thiol-Reactive Electrophiles



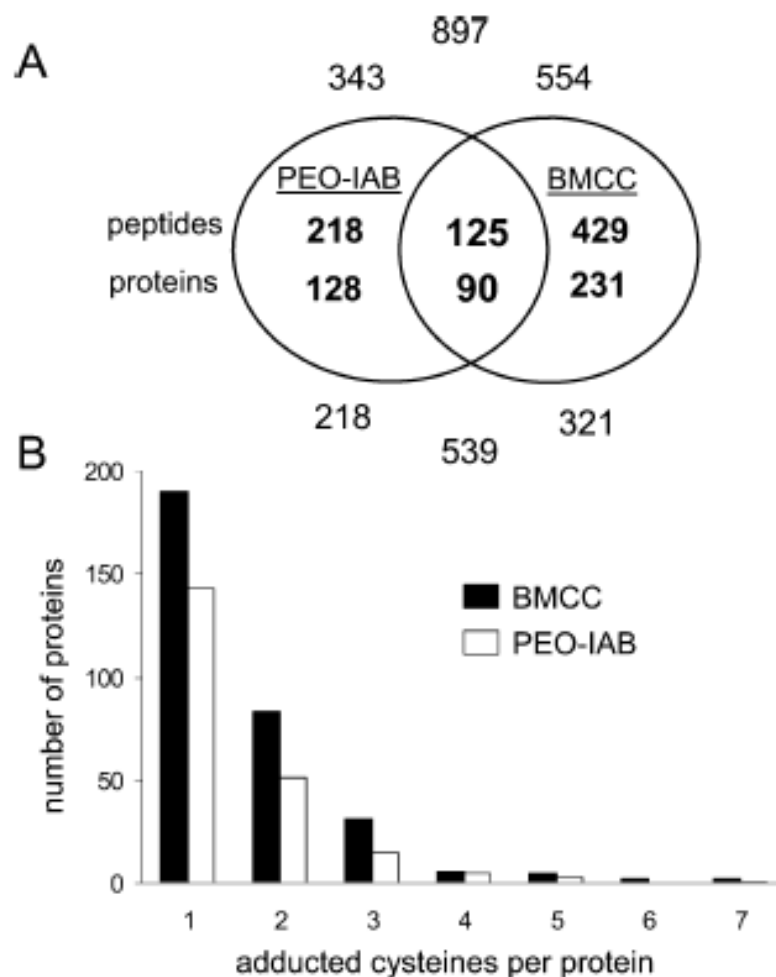


# Cellular Protein Targets of Thiol-Reactive Electrophiles

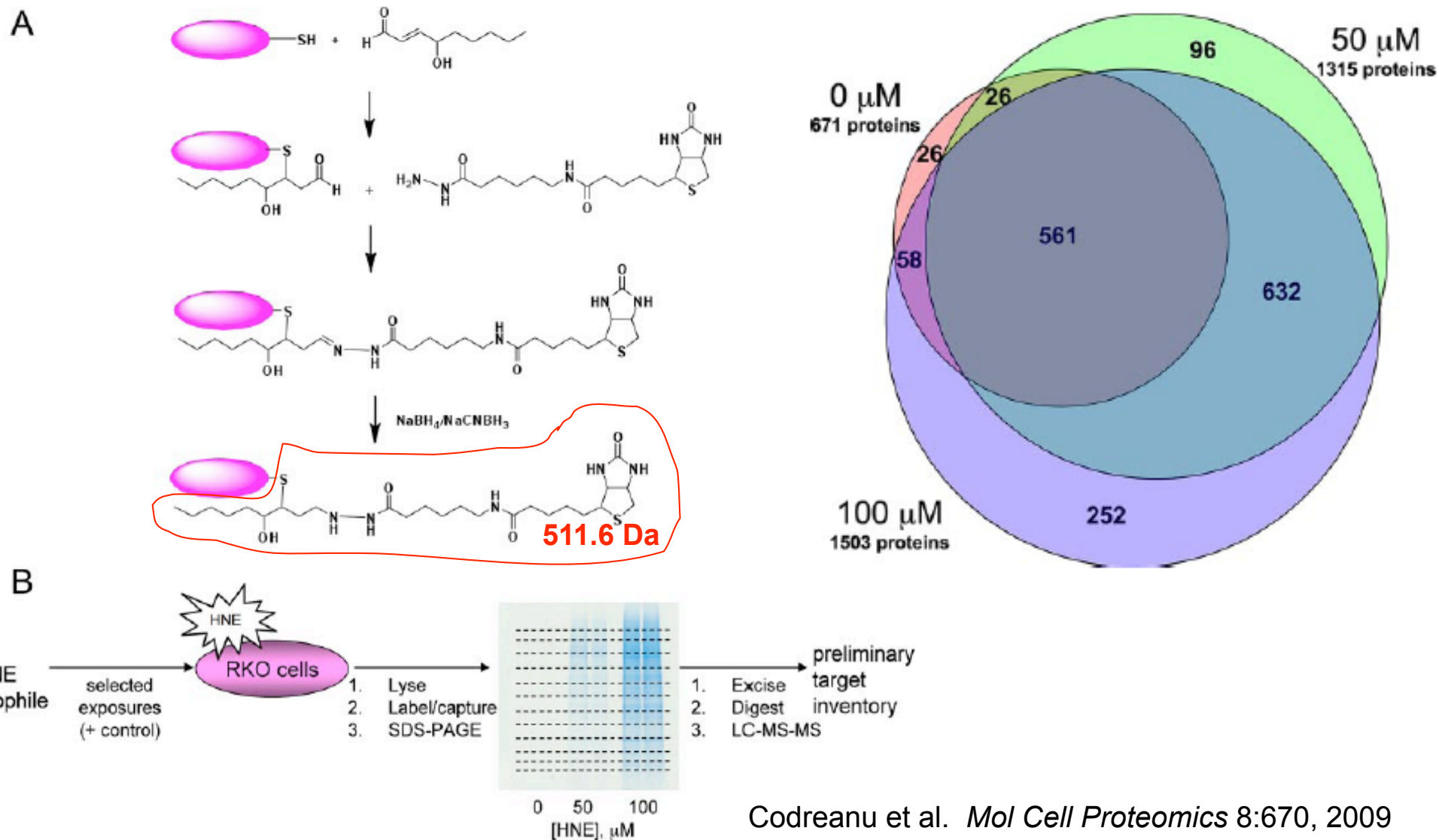
Only cysteine adducts were identified

Greater than 90% of adducted proteins were modified at only 1 or 2 cysteines

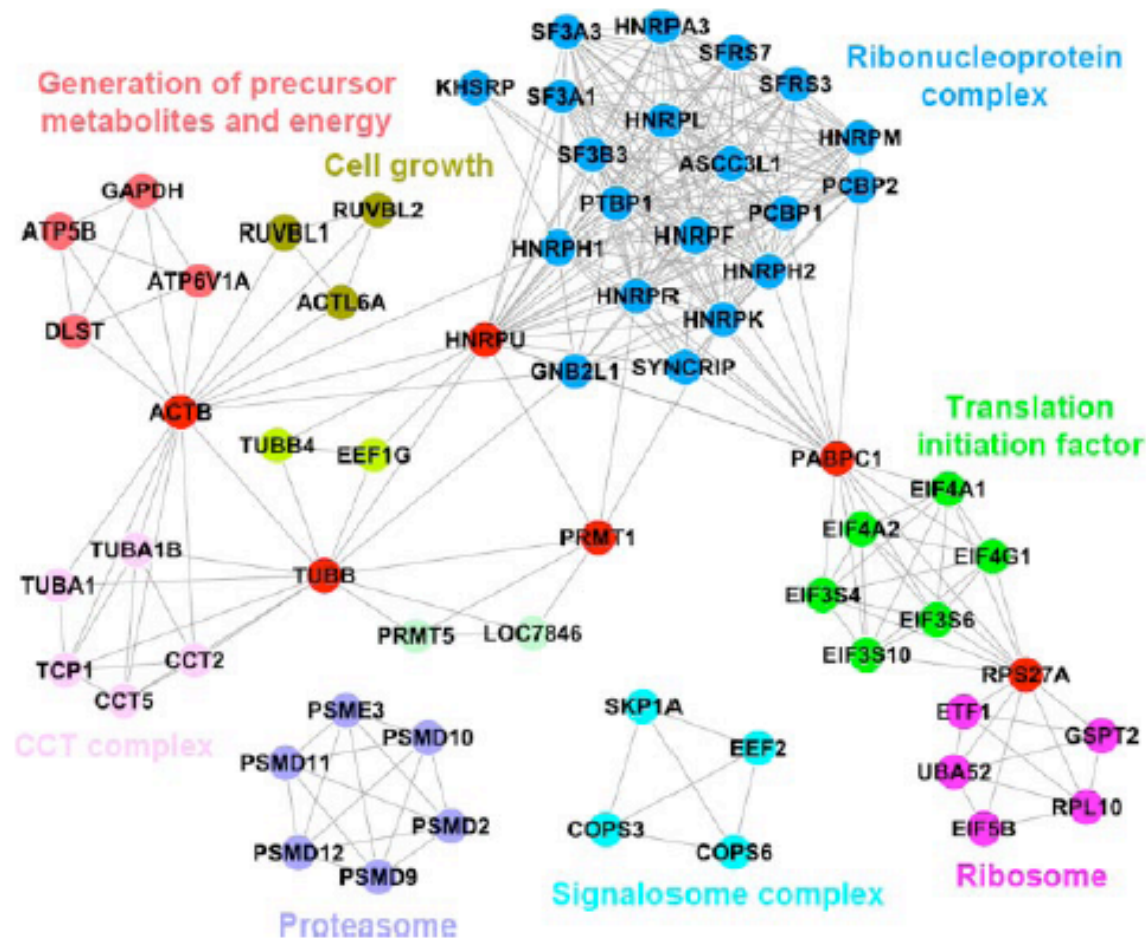
PEO-IAB & BMCC modified a core group of 125 cysteines



# Cellular Protein Targets of Thiol-Reactive Electrophiles



# Protein Communities of 4-HNE-Modified Proteins



# Alcoholic Liver Disease: A multifactorial Disorder Involving Oxidative Stress

## Ethanol Oxidation

↑ NADP(H) redox state

↑ CYP2E1 activity

Increased ROS/RNS Production

↓ Antioxidant status (GSH)

**Steatosis** - Increased Hepatocellular Lipid Accumulation

↑ Depot of peroxidizable substrate

↑ Lipid Peroxidation

↑ Production of 4-HNE

↑ Protein Modification

# Experimental Procedures

Harvest hepatic tissue & subcellular fractions



2-D Gel electrophoresis analysis for detection of 4-HNE modified proteins



Extract 4-HNE immunopositive spots for tryptic digestion

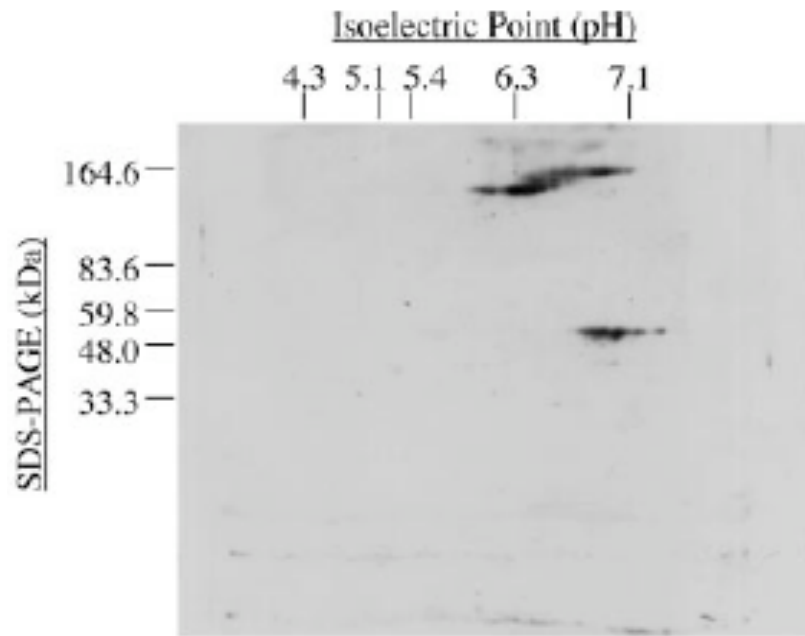


Identify 4-HNE modified proteins using MALDI-TOF/TOF or LC-MS/MS

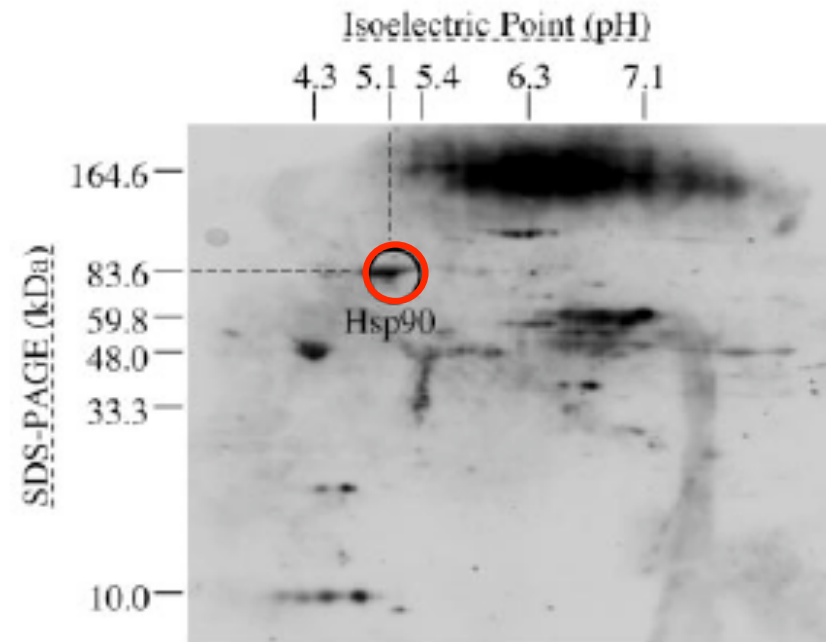


Characterize functional effects of 4-HNE modification using recombinant protein

# HSP90 Detection by Anti-4-HNE Immunoblots of 2-D Gels



**Control**

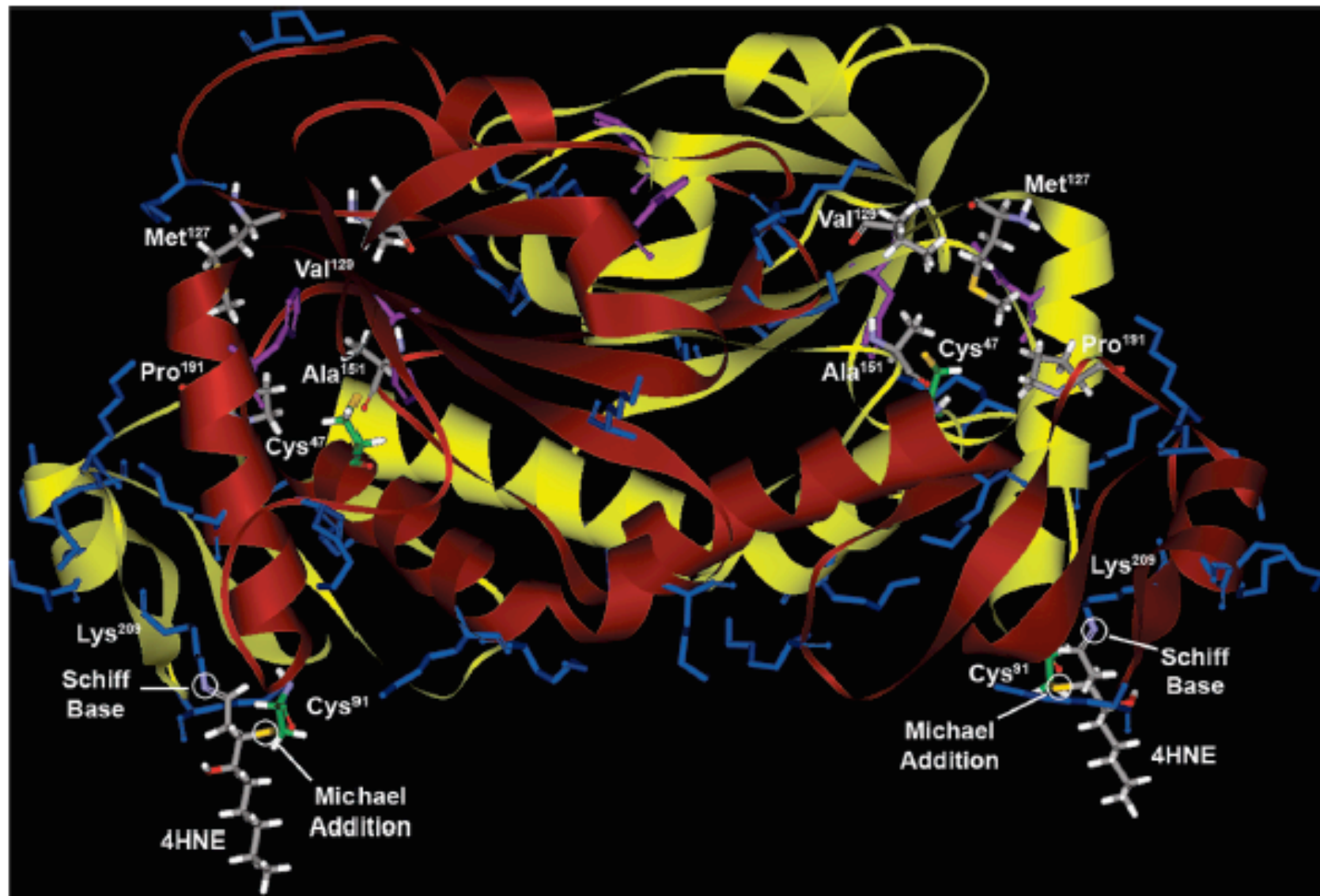


**Ethanol**

# Hepatic Proteins Modified by 4-HNE

Target Protein	Modified Residue	Functional Consequences
<b>HSP70</b> <i>Chem Res Toxicol.</i> 2004; 17:1459	Cys 267    TAC <b>ER</b>	Impaired ATPase activity & protein folding
<b>PDI</b> <i>Chem Res Toxicol.</i> 2005; 18:1324	Cys    57    W <b>CG</b> HC	Dysregulated restoration of mismatched intraprotein disulfide bonds
<b>ADH</b> <i>Free Radic Biol Med</i> 2004; 37:1430	Cys    46    NF <b>CL</b> K	Activity not impaired; enhanced proteosomal degradation
<b>HSP90</b> <i>JPET</i> 2005; 315:8	Cys 572    EN <b>LCK</b>	Impaired refolding of damaged proteins
<b>ERK1/2</b> <i>JBC</i> 2007; 282:1925 <i>Molec Pharmacol</i> 2007; 37:1430	His 178    PD <b>H</b> DH	Inhibition of ERK1/2 phosphorylation and impaired liver regeneration
<b>Prx6</b> <i>Free Rad Biol Med</i> 2008;45:1551 <i>Chem Res Toxicol.</i> 2008; 12:2289	Cys 91    YN <b>CEP</b>	Modification of solvent accessible thiol; activity only marginally impaired
<b>Tubulin</b> <i>Chem Res Toxicol</i> 2007; 20:1111 <i>Biochim. Biophys Acta</i> 2009;1791:772	Cys 347 $\alpha$ , CYS 376 $\alpha$ Cys 303 $\beta$	Inhibition of tubulin polymerization; Lipid accumulation and inhibition of lipid transport in HepG2 cells

## In Silico Modeling of 4-HNE: Peroxiredoxin 6

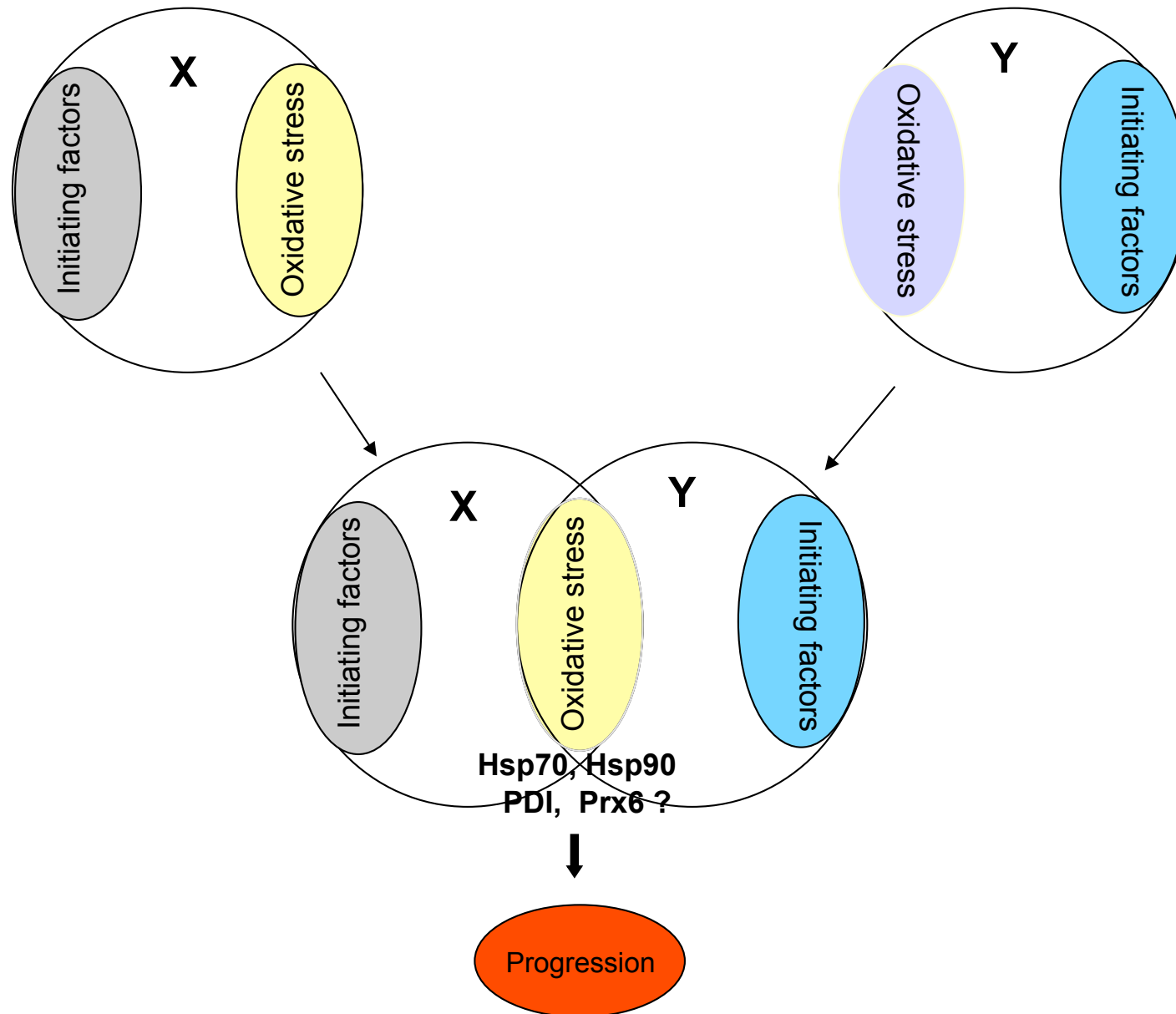




## 4-HNE Modification Does Not Impair the Function of All Proteins

Protein	Functional Consequence
Alcohol Dehydrogenase	Increased/decreased UPS degradation
Catalase	None
Glutamate Dehydrogenase	None

# Proposed Model for Identification of Common Target Proteins and Roles in Disease Processes



# Limitations and Challenges

- Large numbers of proteins are modified by electrophiles making it difficult to identify specific proteins involved in cellular dysregulation.
- There is a low stoichiometry of protein modification. This magnifies the analytical challenges suggesting the necessity of specific affinity enrichment procedures.

# Acknowledgements

## *Petersen Lab*

James Galligan

Rebecca Smathers

Colin Shearn, Ph.D.

Kristofer Fritz Ph.D

## *Funding Sources*

NIH R37 AA09300

NIH RO1 DK074487

NIH F31 AA018818

NIH F31 AA018606

NIH F32 AA018898

## *Former Lab Members*

Dr. Jon Doorn, University of Iowa

Dr. David Carbone, University of Arizona

Dr. James Roede (Emery University)

Dr. Ben Stewart (LLNL)