

Molecular Chaperones & the Heat Shock Response

Session 1 CHAPERONE BIOCHEMISTRY AND PROTEIN FOLDING

WEDNESDAY 5/3/2006, 7:30 PM

E. Craig

| <u>#</u> | <u>Iname</u> | <u>Title</u> | <u>Talk Length</u> |
|----------|------------------|---|--------------------|
| 1 | Craig | Following the evolution of generalist, specialist and "degenerate" Hsp70s | 25 |
| 2 | Deuerling | Ribosome-associated chaperones and their role in protein folding | 25 |
| 3 | Ullers | Trigger factor antagonizes both SecB and DnaK/DnaJ chaperone functions during cold adaptation | 12 |
| 4 | Martinez-Hackert | Structure of trigger factor in complex with a substrate—Insights into the action of a chaperone | 12 |
| 5 | Hammarstrom | An unfolding machine in action—Stretching by the chaperonin GroEL of the substrate protein core | 12 |
| 6 | Rye | Unfolding compaction and release of substrate proteins by GroEL | 12 |
| 7 | Stirling | Phosducin-like proteins as modulators of CCT | 12 |
| 8 | Regini | α -crystallin as a chaperone, structural studies using X-ray and neutron scattering | 12 |

Session 2 QUALITY CONTROL AND PROTEIN TRAFFICKING

THURSDAY 5/4/2006, 9:00 AM

W. Neupert

| <u>#</u> | <u>Iname</u> | <u>Title</u> | <u>Talk Length</u> |
|----------|--------------|---|--------------------|
| 9 | Neupert | Protein trafficking to mitochondria and the roles of chaperones | 25 |
| 10 | Rutkowski | Selective posttranscriptional and posttranslational attenuation of gene expression produces adaptation to ER stress | 12 |
| 11 | Hollien | A new role for Ire1 in the metazoan unfolded protein response | 12 |
| 12 | Williams | Potent in vitro chaperone functions of calnexin under physiological conditions of the ER lumen | 12 |
| 13 | Fass | Reductive activation required for catalysis of disulfide bond formation by Ero1 | 12 |
| 14 | Sevier | Cellular regulation of disulfide bond formation | 12 |
| 15 | Mkrtchian | ERp29, a broad specificity escort chaperone/secretion factor is required for tumor | 12 |

Session 3 POSTER SESSION I

THURSDAY 5/4/2006, 2:00 PM

| <u>#</u> | <u>Iname</u> | <u>Title</u> | <u>Talk Length</u> |
|----------|--------------|---|--------------------|
| 16 | Ahner | A genomic yeast screen identifies small heat shock proteins as regulators of CFTR degradation | |

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| 17 | Akai | Analysis of XBP1 splicing mechanism with improved ER stress-indicators |
| 18 | Ali | Crystal structure of the Hsp90-nucleotide-Sba1 closed chaperone complex |
| 19 | Anckar | Specificity determinants in SUMO-mediated negative regulation of heat shock |
| 20 | Andersson | A novel Clp proteolytic core in the cyanobacterium <i>Synechococcus</i> |
| 21 | Lederkremer | PERK-dependent recruitment of ER-associated degradation and unfolded protein response machineries to the ER quality control compartment |
| 22 | Lukacs | Quality control mechanism of CFTR at post-Golgi compartments |
| 23 | Beatrix | Structure of the human NAC domain—Implications for NACs mode of function |
| 24 | Bernardini | Stress response activation by 50hz sinusoidal magnetic fields in porcine aortic endothelial cells |
| 25 | Birch | Two analogs of cobalamin protect endothelial cells against homocysteine induced oxidative stress |
| 26 | Breiman | The <i>Arabidopsis</i> heat stress induced co-chaperones Rof1 and Rof2 (FKBPs) are developmentally regulated and interact with HSP90 and novel proteins |
| 27 | Bross | The c.297G>C (Val98Ile) variation in the human <i>HPS60</i> gene is associated with hereditary spastic paraplegia SPG13, and affects folding of chaperonin substrates |
| 28 | Bublik | hGTSE-1 protein can modulate the cellular response to stress by regulating the stability of the cyclin-dependent kinase inhibitor p21 ^{CIP1/WAF1} in an Hsp90- |
| 29 | Burch | Modification and reorganization of the cytoprotective chaperone Hsp27 during HSV-1 infection |
| 30 | Catlett | Sgt1 is an Hsp90 adaptor co-chaperone |
| 31 | Chadli | GCUNC-45 is a novel regulator for the progesterone receptor/Hsp90 chaperoning |
| 32 | Vierling | Quaternary structure and substrate interactions of small heat shock proteins probed by mass spectrometry |
| 33 | Chien | Tripeptide motifs in ClpXP mediated protein degradation |
| 34 | Chien | Inhibition of amylin fibril formation by chaperones |
| 35 | Christensen | Genetic models of chronic neuronal degeneration causing hereditary spastic paraplegia—Production and characterization of an animal model for Hsp60 |
| 36 | Shapira | Revisiting the chaperone mediated assembly of Rubisco—Exposure of a hidden RRM blocks LSU translation |
| 37 | Coulstock | Mechanism of human Hsp90 ATPase activity and regulation by co-chaperones |
| 38 | Cox | FKBP52 phosphorylation—A potential mechanism for regulating steroid hormone receptor signaling pathways |
| 39 | Cullinan | The role of Hsp90 in the evolution of chronic myelogenous leukemia |
| 40 | Daugaard | Heat shock protein (Hsp) 70-2 cooperate lens epithelium derived growth factor (LEDGF/p75) in cancer cell survival |

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| 41 | Dickey | The impact of chaperones on non-aggregated tau clearance |
| 42 | Naylor | Chaperonin 10 regulates the dynamics of the response to TLR ligation |
| 43 | Gewirth | Structure of the GRP94 N- and middle domains |
| 44 | Endo | The ER stress pathway involving CHOP, is activated in the lung of LPS-treated mice |
| 45 | Reem | DAP1—A new ER stress response partner of BiP |
| 46 | Eriksen | Tau clearance through protein degradation pathways |
| 47 | Sorensen | The Hsc70 E3 ligase is a stress-related tumor suppressor that reduces ER stress through elimination of misfolded proteins |
| 48 | Fang | Mechanism of glucocorticoid receptor—Hsp90 interactions |
| 49 | Felts | Different cochaperones are required to enable Hsp90-dependent folding of steroid receptors and protein kinases in vitro |
| 50 | Urano | WFS1 is a novel negative regulator of ER stress signaling and regulates apoptosis through WIND |
| 51 | Galadari | Akt degradation is an important determinant of L929 cell death following heat stress |
| 52 | Ganea | Protection of α -crystalline chaperone activity by SOS molecules against modifying |
| 53 | Garg | Hsp90 is a peptide translocator in the MHC class I antigen presentation pathway |
| 54 | Ben-Zvi | Progressive disruption of cellular protein folding in models of polyglutamine |
| 55 | Goeckeler | The ATPase activity of the yeast Hsp110 molecular chaperone, Sse1p, is not peptide-stimulated |
| 56 | Gotoh | Molecular mechanisms and pathological roles of the ER stress-CHOP pathway |
| 57 | Graner | Characterization of the Hsp70 co-chaperone HspBP1 in brain tumors |
| 58 | Grimminger | Regulation of Hsp104p by its co-chaperone the cyclophilin Cpr6p |
| 59 | Knowlton | Extracellular Hsp60—Role of lipid rafts and exosomes in Hsp60 secretion |
| 60 | Hansen | Decreased transcript levels of the Lon and ClpP protease genes in cultured cells from patients with hereditary spastic paraplegia (SPG13) caused by a mutation in the <i>HSP60</i> gene |
| 61 | Haslbeck | Hsp20.2, a small heat shock protein from <i>A. fulgidus</i> is a highly dynamic oligomer |
| 62 | Nakai | Decision of death or life by heat shock transcription factor 1 |
| 63 | Heikkila | Analysis of the expression and function of Hsp27 and Hsp30 in <i>X. laevis</i> embryos |
| 64 | Hoffmann | Trigger Factor forms a protective shield for nascent polypeptides at the ribosome |
| 65 | Elguindi | Membrane protein Hti1 is a transcription factor activated by endoplasmic reticulum stress and a novel transducer of the UPR |
| 66 | Jin | Insights into the mechanisms underlying the neurodegenerative disease observed in <i>hsf-1</i> -deficient mice |

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| 67 | Inoue | Fragmentation of Sup35 amyloid fibers |
| 68 | Inouye | Modulation of Il-6 gene expression by heat shock transcription factor 1 |
| 69 | Takayama | BAG3 regulates cell motility |
| 70 | Johnson | In vivo interaction of Hsp90 with co-chaperones and potential client proteins |

Session 4 CELLULAR RESPONSE TO STRESS

THURSDAY 5/4/2006, 7:30 PM

A. Horwich

| <u>#</u> | <u>Iname</u> | <u>Title</u> | <u>Talk Length</u> |
|----------|--------------|--|--------------------|
| 71 | Horwich | GroEL-GroES-mediated protein folding | 25 |
| 72 | Jakob | The chaperone Hsp33—Activation by stress-induced domain unfolding | 25 |
| 73 | Barral | Real-time observation of trigger factor function on translating ribosomes | 12 |
| 74 | Zuiderweg | The structural basis of interdomain communication of the Hsp70 chaperones in | 12 |
| 75 | Morano | The Sse1/Hsp110 molecular chaperone—An essential modulator of Hsp70 function | 12 |
| 76 | Nicchitta | A stress sensor function for GRP94—Adenine nucleotides as repressors of GRP94 molecular chaperone function | 12 |
| 77 | Tsai | Visualizing the ATP-driven conformational changes in ClpB | 12 |
| 78 | Mogk | M-domain mediated activity control of the AAA+ chaperone ClpB | 12 |

Session 5 CHAPERONE FUNCTION IN DISEASE AND DEVELOPMENT

FRIDAY 5/5/2006, 9:00 AM

F.U. Hartl

| <u>#</u> | <u>Iname</u> | <u>Title</u> | <u>Talk Length</u> |
|----------|--------------|---|--------------------|
| 79 | Hartl | Cytotoxicity of protein misfolding and its modulation by molecular chaperones | 25 |
| 80 | Lee | Oxidative stress-dependent functional switching of yeast 2-Cys Prxs from a peroxidase to a molecular chaperone function | 25 |
| 81 | Houry | MoxR AAA+ ATPase—A novel family of molecular chaperones | 12 |
| 82 | Hager | Molecular chaperones function as steroid receptor mobility factors | 12 |
| 83 | Smith | FKBP co-chaperones and steroid receptor function | 12 |
| 84 | Vaughan | Structure and assembly of Hsp90-Cdc37-protein kinase complexes | 12 |
| 85 | Freeman | The p23 and Hsp90 molecular chaperones mediate telomerase dynamics | 12 |
| 86 | Dillin | Insulin-like signaling couples the aging process and toxic protein aggregation by regulating opposite detoxification activities | 12 |

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|----------|--------------|---|--------------------|
| 87 | Kanzaki | Functional analysis of group II chaperonin using subunit-linkage mutants | |
| 88 | Kim | Molecular chaperones regulating cellular toxicity of α -synuclein in yeast | |
| 89 | Kirstein | Adaptor protein controlled oligomerization activates the AAA+ protein ClpC | |
| 90 | deHaseth | Mutational study of σ 32, the <i>E. coli</i> heat shock σ factor—The role of a conserved histidine in region 2.3 | |
| 91 | Krantz | Single-particle dynamics of Hsc70 mediated clathrin uncoating | |
| 92 | Krojer | The protease state of the heat-shock protein DegP from <i>E. coli</i> | |
| 93 | Kumar | Directed evolution of functional GroELs from <i>M. tuberculosis</i> GroELs—Phenotypic and sequence analysis of the mutants | |
| 94 | Lachapelle | The ability of Hsp70 to inhibit apoptosis is impaired in cells heat-shocked under acidic conditions | |
| 95 | Lee | Structural and functional switching of a human 2-Cys peroxiredoxin by oxidative stress enhances HeLa cell resistance to H ₂ O ₂ -induced cell death | |
| 96 | Vierling | S-nitrosoglutathione reductase (GSNOR) is required for the development of thermotolerance in <i>Arabidopsis</i> | |
| 97 | Lev | Generation of novel TCR-like antibodies reveals the presence of an antigenic peptide pool depended on molecular chaperones | |
| 98 | Lin | Hsp60 localizes to lipid rafts in a rat model of heart failure | |
| 99 | Liu | VIPP1 forms rod-shaped supercomplexes that are disassembled by the chloroplast Hsp70B-CDJ2-CGE1 chaperone machinery | |
| 100 | Lum | Peptide binding to the molecular chaperone Hsp104 | |
| 101 | Stirling | Prefoldin—A flexible molecular clamp with essential functions in a multicellular | |
| 102 | Madan | Triggering of productive protein folding by GroEL | |
| 103 | Voos | Proteomic analysis of mitochondrial protein turnover—Identification of novel substrate proteins of the matrix protease Pim1 | |
| 104 | Mandal | Quality control of protein kinases by Cdc37 is independent of the ubiquitin/proteasome pathway | |
| 105 | Martin | Rebuilding the ClpX unfoldase reveals operating principles for other AAA+ motor | |
| 106 | McClellan | Genome-wide screens uncover the essential cellular functions and substrates of | |
| 107 | McFall | The expansion of the Hsp70 network function in <i>C. elegans</i> | |
| 108 | McNally | Functional divergence between the multiple GroEL proteins of chlamydiae | |

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| 109 | Merz | The C-terminal domain of <i>E. coli</i> trigger factor represents the central module of its chaperone activity |
| 110 | Christians | Search for Hsf1 oocyte-specific target genes |
| 111 | Meyer | Domain analysis of Jjj1—A cytosolic j-protein in <i>S. cerevisiae</i> |
| 112 | Miller | Phenotypes of mice and cells that are null for the p23 co-chaperone |
| 113 | Miller | A role for the unfolded protein response in rescuing defects associated with an aberrant ABC transporter |
| 114 | Millson | MAP kinase clients of the Hsp90 chaperone system |
| 115 | Mivechi | Genetic interplay between heat shock transcription factor Hsf1 and p53 tumor suppressor gene during tumorigenesis |
| 116 | Miot | Toward the in vitro reconstitution of the Cpx system, an extracytoplasmic stress response pathway of <i>E. coli</i> |
| 117 | Cashikar | Genes essential for thermotolerance in yeast |
| 118 | Mok | Tumoricidal activity in the partially folded-protein-lipid complex HAMLET—Structural characterization and its molecular basis of transport into cells |
| 119 | Mokranjac | Import motor of the mitochondrial Tim23 complex |
| 120 | Morris | Structure/function studies on the mammalian small heat shock protein Hsp25—The role of the C-terminal extension |
| 121 | Nair | Coordinating protein folding with Hsp90 ATPase cycle |
| 122 | Nakatsukasa | Reconstitution and chaperone requirements for the ubiquitination of misfolded multispanning ER-membrane proteins |
| 123 | Naylor | Chaperonin 10 has an immunomodulatory activity independent of its essential chaperone function |
| 124 | Neuwald | Statistical evidence for specific DNA clamp loading mechanisms in replication factor C chaperone-like AAA+ ATPase |
| 125 | Noma | Transmission of phenotypic trait of yeast prion in single living cells |
| 126 | Ohtaki | Role of N-terminal residues of <i>Pyrococcus</i> prefoldin in the interaction with substrate and chaperonin |
| 127 | orton | Altered regulation of HSF1 and the neuronal heat shock response |
| 128 | Osherovich | DAF-2 and TOR modulate misfolded protein toxicity |
| 129 | Argon | The essential functions of GRP94 in muscle differentiation and protection from apoptosis are due to its regulation of insulin-like growth factor secretion |
| 130 | Ron | Co-translational degradation protects the stressed endoplasmic reticulum from misfolded client protein overload |
| 131 | Bedard | Menin is a regulator of the stress response in <i>D. melanogaster</i> |

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| 132 | Wolf | The cytoplasmic Hsp70-chaperone machinery subjects misfolded and EF import incompetent proteins to degradation via the ubiquitin-proteasome system |
| 133 | Tatu | Heat shock protein 90 controls development of <i>P. falciparum</i> during intra-erythrocytic cycle |
| 134 | Bar-Nun | ATPases in endoplasmic reticulum-associated protein degradation |
| 135 | Raden | A systems biology approach to understanding the early secretory pathway |
| 136 | Ran | Analysis of defects in Hsp90 chaperone complex interaction with mutant <i>MAL</i> -activator proteins |
| 137 | Rasouly | Induction of σ^{32} at low temperatures is sufficient to induce thermotolerance in <i>E. coli</i> |
| 138 | Retzlaff | Segmental and structural contribution of the N-terminal domain of Hsp90 to its intrinsic ATPase activity |
| 139 | Ricketson | Regulation of nuclear and cytoplasmic localization of apo-glucocorticoid receptors by Hsp90 |
| 140 | Riggs | Genetic analysis of FKBP52 function in steroid hormone signaling |
| 141 | Roos-Mattjus | Regulation of HSF2 stability and DNA-binding activity by HSF1 |
| 142 | Van Wijk | Reduction of the non-catalytic ClpR2 protease subunit of <i>A. thaliana</i> disrupts plastid biogenesis and up-regulates soluble chaperones and membrane proteases |
| 143 | Ryoo | <i>Xbp1</i> splicing and the unfolded protein response is activated as a protective mechanism in the <i>Drosophila</i> model for autosomal dominant retinitis pigmentosa |

Session 7 REGULATION OF THE STRESS RESPONSE

FRIDAY 5/5/2006, 7:30 PM

| R. Morimoto | | | |
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| # | <u>Iname</u> | <u>Title</u> | <u>Talk Length</u> |
| 144 | Morimoto | <i>C. elegans</i> systems analysis of misfolded protein stress | 25 |
| 145 | Nudler | RNA-mediated response to heat shock in mammalian cells | 25 |
| 146 | Sarge | Bookmarking of the Hsp90, Hsp27, and c-fos promoters by Hsf2 during mitosis and its role in modulating constitutive expression of these genes | 12 |
| 147 | Frydman | The substrate spectrum of the chaperonin TRiC/CCT revealed by genomic | 12 |
| 148 | Matts | Structural features that modulate the interaction of Hsp90 and Cdc37 with protein | 12 |
| 149 | Gleiter | Directed evolution to eliminate rate limiting steps in folding pathways in vivo | 12 |
| 150 | Kohnno | The ER stress-sensory mechanism by transmembrane protein Ire1—Dimerization of Ire1 is not sufficient for its activation | 12 |
| 151 | Haynes | Genetic analysis of regulated mitochondrial chaperone gene expression | 12 |

Session 8 CHAPERONES AND PROTEOLYSIS

SATURDAY 5/6/2006, 9:00 AM

B. Sauer

| <u>#</u> | <u>Iname</u> | <u>Title</u> | <u>Talk Length</u> |
|----------|--------------|---|--------------------|
| 152 | Sauer | ClpXP and hslUV—Old dogs, new tricks? | 25 |
| 153 | Langer | Substrate recognition by AAA proteases within mitochondria | 25 |
| 154 | Hill | Activating the proteasome | 25 |
| 155 | Smith | Regulation of the 20S proteasome by the ATPase ring complex | 12 |
| 156 | Tian | A conserved proteasomal processing mechanism regulates the activity of transcription factors Cubitus interruptus and NF- κ B | 12 |
| 157 | Christianson | A complex containing the ubiquitin ligase Hrd1 mediates ERAD of the AMPA receptor subunit GluR1 | 12 |
| 158 | Waksmonski | Sse1p, an Hsp110, contributes to ApoB stabilization | 12 |
| 159 | Epstein | The UNC-45 chaperone regulates myosin assembly and degradation—(A model for muscle wasting) | 12 |

Session 9 POSTER SESSION III

SATURDAY 5/6/2006, 2:00 PM

| <u>#</u> | <u>Iname</u> | <u>Title</u> | <u>Talk Length</u> |
|----------|--------------|--|--------------------|
| 160 | Sahi | Global analysis of cytosolic J-protein function in <i>S. cerevisiae</i> | |
| 161 | SAKURAI | Immaturity of microtubule networks in α B-crystallin antisense C2C12 mouse myoblast cells | |
| 162 | Santos | Protein disulfide isomerase (PDI)/NAD(P)H oxidase interaction is a pathway bridging the unfolded protein reaction (UPR) to oxidative stress in vascular smooth muscle cells (VSMC) | |
| 163 | Schroder | The unfolded protein response represses differentiation through the <i>RPD3-SIN3</i> histone deacetylase | |
| 164 | Schroda | Are heat shock factors involved in counteracting transgene silencing in <i>Chlamydomonas</i> ? | |
| 165 | Scroggins | Regulatory acetylation site of Hsp90 and characterization of mutations | |
| 166 | Shenhar | Expression of heat shock proteins in <i>E. coli</i> is affected by the mRNA chaperone | |
| 167 | Sjögren | New insights into the chloroplast Clp-protease of higher plants—Oligomeric structure and substrate identification | |
| 168 | Smith | Effect of sulforaphane on heat shock protein mediated apoptosis in colon cancer | |
| 169 | Snyman | Salicylic acid potentiates heat-induced Hsp70 accumulation through the activation in Hsf-DNA binding in tomato seedlings | |

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| 170 | Spiess | Identification of substrate binding sites in eukaryotic chaperonin TriC/CCT reveals novel mode of binding |
| 171 | Stahl | Structural and functional role of chaperones in duck HBV polymerase activation |
| 172 | Stankowski | Interaction between Hsp70 and ChiP leads to protection against cell death |
| 173 | Steed | Small-molecule inhibitors of Hsp90 display selectivity between Hsp90 client |
| 174 | Strbuncelj | The prefoldin chaperone complex mediates the stable propagation of Sup35 forms |
| 175 | Sun | Chaperone displacement from mutant cystic fibrosis transmembrane conductance regulator (CFTR) restores its biogenesis and function in human airway epithelial |
| 176 | Takaki | Ciliary dyskinesia in mice deficient for heat shock transcription factor 1 |
| 177 | Niwa | Direct involvement of NF- κ B in transcription of the UPR target genes |
| 178 | Tanaka | From conformation to phenotype—The physical basis of prion strain diversity |
| 179 | Hayer-Hartl | Mechanism of accelerated protein folding by GroEL/GroES chaperonin |
| 180 | Tani | A novel membrane protein recognizing the C-terminal diversity among heat shock protein 70 |
| 181 | Terada | Differential roles of cytosolic type I DnaJ homologs in mammals |
| 182 | Tessarz | Conserved and novel functions of the AAA+ chaperone Hsp104 |
| 183 | Thibault | Specificity in substrate and cofactor recognition by the N-terminal domain of the chaperone ClpX |
| 184 | Tipton | Chaperone-mediated prion amplification |
| 185 | Travers | Coevolutionary analysis identifies putative functional amino acid residues within Hsp70, Hsp90 and hop |
| 186 | Needham | Primate chaperones Hsc70 (constitutive) and Hsp70 (induced) differ functionally in supporting growth and prion propagation |
| 187 | Tveten | Rescue of the low density lipoprotein receptor 2A mutant G544V by 4-phenylbutyrate |
| 188 | Uryu | Convergence of Hsp90 with ubiquitin in filamentous α -synuclein inclusions of α -synucleinopathies |
| 189 | Vabulas | Protein synthesis upon acute nutrient restriction relies on proteasome function |
| 190 | Vang | Prediction of aggregation propensity for actin variants in congenital myopathies |
| 191 | Vembar | Peptide affinity and Hsp40 J-domain interaction differentially affect BiP function |
| 192 | Virador | Differential BAG3 binding and behavior of Hsp70 as Hsp27 in breast cancer cells |
| 193 | Voisine | Identification of chaperone networks in <i>C. elegans</i> |
| 194 | Wendler | Comparison of hexameric structures of full-length and Δ N variants of Hsp104 by |
| 195 | Wilkinson | HSV-1 infection disrupts the ATR-dependent DNA damage response through sequestration of repair factors into virus-induced chaperone-enriched nuclear |

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| 196 | Willmund | Interaction between the chloroplast Hsp90 and Hsp70 chaperone systems |
| 197 | Schroda | The N-terminal domain of the chloroplast GrpE homolog CGE1 is required for dimerization and cochaperone function in vivo |
| 198 | Gestwicki | Small molecule agonists of heat shock protein (HSP) 70 |
| 199 | Wright | The yeast Hsp40 chaperone Ydj1p plays a role in the PKC pathway and cell wall |
| 200 | Wu | Curing of yeast prions by guanidine inactivation of Hsp104 does not require cell |
| 201 | Xiao | Swa2p/auxilin domain requirements for co-chaperoning Hsp70 clathrin uncoating activity in vivo |
| 202 | Xu | Strap—A heat shock-responsive transcriptional co-factor |
| 203 | Yaglom | Heat shock protein Hsp72 functions in cancer cells in avoiding p53-dependent and p53-independent senescence |
| 204 | Yamagishi | Importance of the phenolic hydroxyl group of sodium salicylate derivatives for the induction of heat shock response |
| 205 | Yang | Association with Hsp90 inhibits Cbl-mediated downregulation of mutant EGF |
| 206 | Yohda | Small heat shock protein protects denatured proteins from aggregation through transient interaction in the dissociated state |
| 207 | Yoshida | The modular structure of pXBP1(U), a negative feedback regulator of mammalian unfolded protein response, encoded in XBP1 pre-mRNA |
| 208 | Young | Systematic analysis of molecular chaperone interactions within the endoplasmic reticulum of <i>S. cerevisiae</i> |
| 209 | Younger | Recognition of folding defects in CFTR Δ F508 at sequential quality control checkpoints by the ER membrane-localized ubiquitin ligase RMA1 and cytosolic CHIP |
| 210 | Zako | Localization of the prefoldin interaction sites in the hyperthermophilic group II chaperonin and correlation between affinity and protein transfer rate between them |
| 211 | Zavialov | Resolving the energy paradox of chaperone/usher-mediated fiber assembly |
| 212 | Zhang | Chaperoned ubiquitylation—Structure of the CHIP E3 ubiquitin ligase and a CHIP-Uev1a complex |
| 213 | Zhang | The thiol oxidoreductase ERp57 promotes rapid disulfide formation in MHC class I molecules |
| 214 | Back | The structure of an unfolded protein response sensor reveals a novel dimerization interface required for IRE1 and PERK activation |

Session 10 DISEASES OF PROTEIN MISFOLDING

SUNDAY 5/7/2006, 9:00 AM

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| J. Kelly | | | |
| # | Iname | Title | Talk Length |
| 215 | Soto | Prion protein misfolding and the mechanism of neurodegeneration | 25 |
| 216 | Eisenberg | Structural studies of amyloid | 25 |

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| 217 | Kelly | Constitutive amyloidogenicity becomes toxic with aging | 25 |
| 218 | Masison | Hsp104 mutations that overcome impairment of yeast prions by Hsp70 uncover role for conserved domain in prion propagation | 12 |
| 219 | Serio | Hsp104 remodels existing prion complexes in vivo | 12 |
| 220 | Taguchi | Direct observation of amyloid fibril growth using a total internal reflection fluorescent microscopy | 12 |
| 221 | Viswanathan | Role of <i>C. elegans sir-2.1</i> and prion-like glutamine/asparagine-rich proteins in ER-stress and life span regulation | 12 |
| 222 | Zaarur | Targeting heat shock response to sensitize cancer cells to proteasome and Hsp90 inhibitors | 12 |