

YING WAI LAM

Vermont Biomedical Research Network (VBRN) Proteomics Facility
 Department of Biology
 The University of Vermont
 Room 337, Marsh Life Science Building,
 109 Carrigan Drive, Burlington, Vermont 05405
<https://www.uvm.edu/cas/biology/profiles/ying-wai-lam>

ylam@uvm.edu
 Tel: 802-656-4709

<https://vbrn.org/proteomics/>

EDUCATION

- Ph.D. in Biochemistry**, The Chinese University of Hong Kong **Dec. 1998 - Dec. 2001**
 Thesis: *Purification and Characterization of Defense Proteins from Various Biological Sources.*
 Chromatographically isolated (ion exchange, size exclusion, and affinity) and functionally characterized new defense proteins (lectins, antifungal proteins, and ribosome-inactivating proteins), which play important roles in host defense against exogenous attacks/infections in plants.
- B.Sc. in Biochemistry**, The Chinese University of Hong Kong **Sep. 1995 - Dec. 1998**

PROFESSIONAL EXPERIENCE

- Director, Vermont Biomedical Research Network (VBRN) Proteomics Facility** **Jun. 2020 - present**
Director, Vermont Genetics Network (VGN) Proteomics Facility **Jun. 2011 - May 2020**
Research Associate Professor, Department of Biology, University of Vermont **Sep. 2019 - present**
- Research Assistant Professor, Department of Biology, University of Vermont** **Jun. 2011 - Jul. 2019**
 Direct the VBRN (formally VGN) Proteomics Facility (supported by Vermont INBRE, NIH NIGMS P20GM103449) with the goals of providing expertise and the latest proteomics technology to researchers, as well as establishing an educational environment for sharing experience and knowledge in proteomics. Work with faculty, staff, postdocs, and graduate students, providing guidance and support for their proteomics projects from initial experimental design, sample preparation, and instrument operation, to data analysis and interpretation. Assist investigators in initiating proteomics projects, publishing their findings, and obtaining intramural and external funding. Together with my colleagues, maintain the operation of facility instrumentation (LTQ linear ion trap MS, LTQ-XL-ETD linear ion trap MS, LTQ-Orbitrap Discovery MS, Q-Exactive Plus MS, and Orbitrap Fusion Tribrid MS), evaluate new technologies and software, and develop methodologies to address emerging proteomics and mass spectrometry needs. Give seminars, guest lectures and facility tours to faculty, faculty candidates, undergraduate, and graduate students from UVM and Primarily Undergraduate Institutions (PUIs) in Vermont. Prepare annual facility progress reports to NIH, External Advisory Committee and AAAS, as well as the proteomics core portion of the INBRE grant renewals. Worked with VBRN since 2020 to transition the facility operations to an income-expense model on July 1st, 2021. Preparing to relocate the facility to the new Firestone Building as part of the Center for Biomedical Shared Resources serving investigators from the Northern New England Region (VT, NH, ME).
- Research Assistant Professor** **Oct. 2005 - May 2011**
Department of Environmental Health, University of Cincinnati, OH

Main research focuses were: 1) elucidation of the unknown mechanisms underlying prostate aging and cancer development in an animal model of aging by stable isotope-based quantitative proteomics (ICAT) (Lam et al., *Proteomics*, 2008), and 2) understanding the role of oxidative/nitrosative stress-induced protein modifications

(nitration/S-nitrosylation) in prostate inflammation and cancer development (Lam et al., PLOS One, 2010). The results yielded clues for designing and interpreting subsequent hypothesis-driven molecular biology experiments on the importance of S-nitrosylation in prostate cancer cell migration/invasion.

Established a proteomics laboratory equipped with the LTQ-Orbitrap and capital proteomics equipment. Developed a number of research collaborations that involved protein identification, phosphorylation analyses, and structural elucidation of hydroxymetabolites using accurate mass and MSⁿ.

Postdoctoral Fellow

Oct. 2001 - Sep. 2005

Department of Surgery, University of Massachusetts Medical School, MA

Developed an approach utilizing a profiling platform (solid-phase extraction in conjunction with MALDI-TOF-MS) to classify patients with prostate cancer according to their serum protein profiles. One of the most relevant biomarkers pinpointed for metastatic prostate disease was isolated and identified by a multi-dimensional separation scheme in conjunction with mass spectrometry and validated by biochemical methods (Lam et al., Proteomics 2005).

RESEARCH AND COLLABORATIONS

I have worked directly with UVM investigators (**66** faculty, **16** staff, **15** postdocs, and **37** graduate students) from **18** UVM departments and **29** investigators from other institutions in multiple states, providing expertise, training and guidance for their proteomics projects. The facility begins projects with a consultation, during which details of the projects are discussed with investigators. The facility personnel help design proteomics experiments tailored to the experimental goals. As the projects progress, we discuss results, assist in data interpretation, paper preparation, and grant submission. We also provide education and training to faculty, staff and students depending on their needs and skill levels. Two to three consultations (a total of ~ 2 hours) are arranged per week.

I have collaborated with a number of investigators and applied an array of state-of-the-art mass spectrometry-based proteomics strategies to their studies. These studies range from routine protein identification, characterization of protein interaction, and post-translational modifications (e.g., acetylation, methylation, phosphorylation, glutathionylation, S-Sulfenylation, etc.), to large-scale quantitative proteomic analyses using stable isotopes (stable isotope labeling by amino acids in cell culture (SILAC), dimethyl labeling, and tandem mass tags (TMT)). Since 2012, I have co-authored **30** publications with UVM investigators and external users on a variety of projects. These collaborative efforts involve the use of large-scale proteomics discovery platforms to yield data and generate new hypotheses to address important research questions that cannot be interrogated by other biochemical or genomics strategies. I co-designed and helped develop in-house workflows and guided graduate students to establish methods specific to the research projects, for example, dimethyl labeling and strong cation exchange separation workflow for elucidating *A. actinomycetemcomitans* membrane proteome (Smith et al., Proteomics, 2015), and enrichment workflows for stable isotope-based phosphoproteomics (Hasan et al., MBio, 2020). I established specific mass spectrometry experiments with investigators for their specialized projects, for example, mapping novel cleavage sites (Lee et al., Molecular Neurodegeneration, 2017), identification and quantification of phosphorylation and S-sulfenylation using stable isotopes and parallel reaction monitoring (PRM) (Krishnamurthy et al., MBio, 2016; Heppner et al, Nature Communications, 2018), elucidating exosome protein content under asbestos exposure using gel-based fractionation method and TMT labeling (Munson et al., FASEB J, 2018), and quantifying estradiols in vomeronasal organs using stable-isotope dilution LC/MS techniques and Selected Reaction Monitoring (SRM) (Cherian, Wai Lam et al., Neuroscience, 2014). The results were published in the journals of investigators' interest and in leading proteomics journals with a main focus on proteomic discovery (e.g. Journal of Proteomics, and Proteomics).

Current efforts involve establishing new proteomics techniques for structural proteomics, including hydrogen-deuterium exchange mass spectrometry, isotope based cross-linking, and disulfide bond elucidation. These method developments target areas that are carefully assessed as the future needs of our investigators.

Major collaborations (since 2011) for which I worked closely with the PIs, their postdocs and graduate students are listed below:

Institution Department	PI	Research Areas and Proteomics Approaches	Representative Publications
UVM MMG	Ward	Phosphoproteome SILAC PRM and Skyline analysis	Krishnamurthy S (2016) MBio 7, pii: e00754-16.
UVM MMG	Mintz	Membrane proteome SCX fractionation Stable isotope dimethyl labeling	Smith et al (2015) Proteomics 15:1859-67. Smith KP et al (2015) Mol Oral Microbiol 30:97-110.
UVM MMG	Huston	Phosphoproteomics (IMAC, TiO ₂) Stable isotope dimethyl labeling	Hasan et al (2020) MBio. 11(4):e00660-20.
UVM Pathology	Van der Vliet	Redox biology/S-sulfenylation Dimedone/dimedone-d6 labeling PRM and Skyline analysis Iodo-TMT	Hristova et al (2014) Redox Biol 2: 436-446. Heppner et al (2018) Nat Commun, 9:4522.
UVM Pathology	Shukla	Exosome proteomics TMT-based profiling	Munson et al (2018) J Cell Biochem 119:6266-6273. Munson et al (2018) FASEB J fj201701291RR.
UVM Pathology	Heininger	Redox biology/glutathionylation PRM and Skyline analysis TMT-based profiling	Corteselli et al (2021), <i>manuscript submitted</i>
UVM Pathology	Anathy	Protein interaction Spectral counting	Kumar et al (2021) Thorax, 77:669-678 Chandrasekaran et al (2021), <i>submitted</i>
UVM Pulmonary Medicine	Weiss	ECM protein profiling Spectral/peptide counting	Platz et al (2016) Tissue Eng Part C Methods 22: 725-739. Wagner et al (2014) Biomaterials 35: 2664-2679. Wagner et al (2014) Biomaterials 35: 3281-3297. Sokocevic et al (2013) Biomaterials 34: 3256-3269. Bonenfant et al (2013) Biomaterials 34: 3231-3245.
UVM Biology	Delay	Small molecule quantification SRM absolute quantification	Cherian et al (2014) Neuroscience 269: 43-58.
UVM Animal Sci.	Greenwood	Milk Proteome Spectral counting TMT-based profiling	Tacoma et al. (2016) J Proteomics 130: 200-210. Tacoma et al. (2016) Data Brief 6: 843-846.
Mayo Clinic	Lee	Cleavage site mapping TMT Pro based profiling	Lee et al. (2017) Mol Neurodegener 12: 55.
Univ. of Connecticut Chemistry	Yao	SILAC analysis	Li et al (2017) Anal Chem 89: 6295-6299.
Univ. of Cincinnati Environmental Health	Yadav	Small molecule characterization Accurate mass/MS ⁿ	Syed et al (2013) Appl Environ Microbiol 79: 2692-2702.
Mississippi. Sate Univ. Biochemistry	Li	Protein identification from 2D gels	Liu et al (2017) Acta Physiol Plant 39:215.

Besides research collaborations, I had also provided routine proteomics services and expertise to many investigators at UVM and nationwide (e.g., University of Connecticut, Albany College of Pharmacy, and Dartmouth College). From Aug. 2012 to 2014, recognizing the need for accurate mass measurements to fulfill publication requirements of chemistry journals, I had helped **6** UVM Chemistry faculty members authenticating their compounds of interest (a total of **11** publications acknowledging the facility).

SERVICES

Departmental

Thesis committee membership

UVM

Emily Price (PhD candidate)	Thesis committee member (Biology)	Oct. 2021 -
Lynda Menard (PhD candidate)	Thesis committee member (Biology)	Jan. 2017 - Dec. 2020
Wyatt Chia (PhD candidate)	Thesis committee member (CMB)	Jul. 2016 - Apr. 2020
Ravi Nagori (PhD candidate)	Thesis committee member (Biology)	Dec. 2014 -
Robert Bauer (PhD candidate)	Outside committee member, CMB Qualifying Exam (Phase II)	May 2014 - May 2016

Wei Du

Thesis Committee member (Biology)

October 2022
Jan. 2014 - Dec. 2014

ad-hoc RPT guidelines review committee

Department of Biology, UVM

Nov.21 2020 –
Jan. 31, 2021

Prior to 2011

University of Cincinnati

Jared Isaac (PhD, graduated 06/12) Co-mentor/thesis committee member Jul. 2009 - May 2011

Thesis title: Nitrosylation of Integrins in Prostate Cancer Progression (Dept. of Cancer and Cell Biology)

I served as mentor on the Gene-Environment Interactions Training Program (GEITP), NIEHS, T32ES016646

Regional / National

Reviewer for the NSF

2022

Session co-host “Mass spectrometry 101”

Northeast Regional Laboratory Staff and Core Directors (NERLSCD) Meeting
(Portsmouth, NH)

Nov. 4, 2021

MENTORING / TEACHING / OUTREACH

I) Facility technicians trained

Since 2012, I have trained five full-time facility technicians. They were involved in many aspects of the facility operation and were equipped with solid proteomics knowledge and technology, general laboratory techniques, and scientific writing/communications skills for the next stage of their careers.

Sydney Guthrie		Jul. 2022 -
Clarissa Gold	Coauthored 4 manuscripts	Sep. 2019 - May 2022
Catrina Hood	Coauthored manuscripts in preparation <i>Currently enrolled in UVM medical school</i>	May 2017- May 2019
Bethany Ahlers	Coauthored 2 publications	Jan. 2016 - Feb. 2017
Julia Fields	Coauthored 6 publications (as second author on 5 papers; 1 in <i>Proteomics</i> and 1 in <i>Journal of Proteomics</i>) <i>Enrolled in Master of Science in Medical Laboratory Science</i> <i>Currently a medical laboratory scientist</i>	Aug. 2012 - Dec. 2015

II) Provide support for UVM and Vermont PUI classes (sample analysis/lectures/facility tours)

Worked with the VGN Outreach Core, UVM and PUI faculty members to help incorporate a proteomics experimental component into their courses or laboratory curricula, aimed at equipping future scientists with state-of-the-art proteomics knowledge. Gave guest customized proteomics lectures (with in-class exercise and video demonstration), which are usually followed by a 15 to 45 min facility tour.

UVM Course	Course Director	Year	No. of Students	# of Sample Analysis	Facility Tour (Date)	Lecture (Date)
UVM/BIOL 205	Van Houten	2018, Spring	11	10	Apr. 17	Mar. 22/Apr. 19
	Van Houten	2017, Spring	10	15	Apr. 18	Mar. 23/Apr. 18
	Van Houten	2016, Spring	12	24	Apr. 7	Mar. 15/Apr. 14
	Van Houten	2015, Spring	9	--	Mar. 31	Mar. 10/Apr. 9
	Van Houten	2014, Spring	11	--	Apr. 2	Mar. 12/Apr. 10

	Van Houten	2013, Spring	13	--	Apr. 2	Mar. 12/Apr. 9
UVM/BIOC 207	Everse/Silveira	2022, Spring	21	21	No tour	By course instructor
	Everse/Silveira	2019, Spring	35	35	No tour	By course instructor
	Everse/Silveira	2018, Spring	30	30	No tour	By course instructor
	Everse/Silveira	2017, Spring	24	24	No tour	By course instructor
	Everse/Silveira	2016, Spring	26	31	No tour	By course instructor
	Everse/Silveira/Bouchard	2015, Spring	24	33	No tour	By course instructor
	Everse/Silveira/Bouchard	2014, Spring	16	35	No tour	By course instructor
UVM/MLS 221	Ray	2016, Spring	24	No	Mar. 4	Mar. 4
UVM/NSCI 306	Nishi	2016, Summer	5	No	Aug. 6	Aug. 6
	Nishi	2014, Summer	6	No	Aug. 19	Aug. 19
UVM/PBIO 187	Almstead	2015, Summer	11	18	No tour	By course instructor
	Almstead	2014, Summer	8	14	No tour	By course instructor
	Almstead	2013, Fall	109	No Analysis	Oct. 14-18 (4)	By course instructor
	Almstead	2013, Summer	13	No Analysis	No tour	By course instructor

Colleges	Course Director	Year	No. of Students	# of Sample Analysis	Facility Tour	Lecture
<u>Vermont</u>						
Saint Michael's College	Lamos	2021, Fall	6	No Analysis	Nov. 1	By course instructor
	Lamos	2019, Fall	9	No Analysis	Dec. 2	By course instructor
	Schroll	2019, Spring	9	No Analysis	Apr. 25	Apr. 25
	Lamos	2018, Fall	6	No Analysis	Dec. 3	By course instructor
	Schroll	2018, Spring	10	18	Apr. 26	Apr. 26
	Schroll	2017, Spring	4	4	May 1	May 1
	Lamos	2017, Fall	8	No Analysis	Nov. 27	By course instructor
	Schroll	2016, Spring	6	6	Apr. 25	Apr. 25
	Schroll	2015, Spring	--	20	Apr. 27	By course instructor
	Schroll	2014, Spring	--	12	Apr. 14	By course instructor
	Lamos	2014, Fall	--	No Analysis	Oct. 24	By course instructor
	Schroll	2013, Spring	9	--	Apr. 18	By course instructor
	Lamos	2013, Fall	10	No Analysis	Nov. 21	Oct. 24
	Schroll	2012, Spring	--	--	Apr. 19	By course instructor
Schroll	2011, Spring	--	--	Apr.	By course instructor	
Northern Vermont University	Landesman	2021, Fall	4	4	Nov. 30	Nov. 30
Green Mountain College	Landesman	2018, Spring	7	12	No tour	--
	Coe	2016, Spring	13	No Analysis	Apr. 13	Apr. 13
	Coe	2013, Spring	8	No Analysis	Dec. 2	Dec. 2
Community College of Vermont	Joy	2017, Fall	20	No Analysis	Nov. 20	Nov. 20
Castleton University	Palmer	2017, Fall	9	No Analysis	Dec. 6	Dec. 6
	Palmer	2015, Fall	10	No Analysis	Oct. 27	Oct. 27
Marlboro College	Smith	2015, Spring	5	No Analysis	Apr. 10	Feb. 20/Mar. 6
Norwich University	Guth	2014, Spring	--	44	Feb. 24/28	Feb. 24/28
	Guth	2013, Spring	7	--	Nov. 11	Nov. 11
	Guth	2012, Spring	--	--	Apr. 16	Apr. 16
<u>Others</u>						
SUNY Plattsburgh	Valentine	2018, Fall	7	27	Nov. 1	By course instructor

(-- : information not collected)

III) Research collaboration with PUI faculty on proteomics and mass spectrometry

Since 2012

Meet with investigators from Vermont Colleges regularly to discuss their research projects. Provide research support, help design, run and interpret proteomics and mass spectrometry experiments. Discuss ideas for grant applications and facilitate their research connection with UVM investigators.

- Provided expertise for 8 PUI faculty (3 from St. Michael's College and 4 from Middlebury, 1 from NVU)
- Supported 3 VGN awards ('12-'13, '14-'16, '17-'18), 1 NSF, and 1 R01
- Served as faculty mentor on proteomics research for 2 BPI faculty (from St. Michael College ('12-'13) and Norwich Univ., '14-'15)
- Hosted a sabbatical stay for a St. Michael College PI in the facility (summer '14)

IV) Other services/outreach activities**Local (via VGN / VBRN)****Proteomics webinar speaker**

(with VGN Professional Development and Education Core)

- for PUIs (<https://www.youtube.com/watch?v=MaVbYhjingXQ>) 10/12/17
(131 views, Feb, 2020)
- for Community College of Vermont 11/30/18

Round table discussion guest, VGN Career Day

04/12/17, 04/11/18

Poster judge, VGN Career Day

04/16/14, 04/13/16, 04/12/17

Poster judge, New England Regional Shared Resources Conference

11/12/15

Facility tours, VGN Career Day (for undergraduates from Vermont PUIs)

04/11/18, 04/3/19, '04/13/22

Article contributor, VGN newsletters/magazines/handbooks

'12, '13, '15, '18, '19

Reviewer, grant proposal/pitch paper for VGN/VBRN grant writing workshop

'18, '20, '21, '22

GRANTS SUPPORTED

Since 2011, I have provided **95** letters of support, preliminary data, and write-ups (as a collaborator, key personnel, significant contributor, or consultant; no percent effort prior to 2021, as I was fully supported by VGN INBRE) for grant applications to federal and non-federal agencies. Percent effort on grant since 2021 is listed in "Research Support".

Twenty-two have been funded, **8** are currently active (underlined). (**12 from NIH**: F31HL142221 (Dustin), F32HL129706 (Heppner), R01DE014711 (Spatafora), R15GM123393 (Hass/Cen), R01HL122383 (Anathy), R01NS045940 (Cipolla), R01AI105191 (Ward), R01HL085646 (Van der Vliet), R01HL138708 (Van der Vliet), R01HL137268 (van der Vliet/Dixon), R01GM054899 (Francklyn), R01GM117155 (Jordan, Johns Hopkins Univ.);**1 from DOD**: IDeA W81XWH-14-1-0199 (Shukla); **2 from Foundation**: Hearing Health Foundation Res. Grant (Bond), Preeclampsia Foundation (Ko); **1 from NSF**: MCB 1817793 (Garcia, Castleton Univ.); **1 Hatch Award**, USDA-NIFA HATCH VT-H02009 (Greenwood), **3 from UVM**: CVRI ECAC award (Ko), FAHC Res. Award (Krag), COBRE PIP (Dixon); **3 VGN Pilot/Project Award**: (Lamos, Wuorinen, Garcia)).

AWARDS**Partial Travel Award, American Urological Association****Aug. 2005**

Society for Basic Urologic Research Summer Research Conference

*Oxidative and Nitrosative Stresses and Inflammation as Early Events of Hormone-induced Prostatic Carcinogenesis in Noble Rats***Department of Defense-Prostate Cancer Program Postdoctoral Research Award Apr. 2003 - Mar. 005***Proteomics Approach to Evaluate the Impact of Diet and Stress Reduction on Prostate Cancer Progression.*

Role: Principal Investigator. \$98,000

PROFESSIONAL SOCIETY MEMBERSHIPS

Member, American Society of Mass Spectrometry

Since 2002

Member, American Chemical Society

Since 2002

Member, Association of Biomolecular Resource Facilities

Since 2019

Faculty, Graduate College, University of Vermont

Since 2014

PEER-REVIEWED PUBLICATIONS

1. **2021** Kumar A, Elko E, Bruno SR, Mark ZF, Chamberlain N, Korwin-Mihavics B, Chandrasekaran R, Walzer J, Ruban M, Gold C, Lam YW, Ghandikotad S, Jeggad AG, Gomez JL, Janssen-Heininger YMW and Anathy V. Inhibition of PDIA3 in Club Cells Attenuates Osteopontin Production and Lung Fibrosis. **Thorax** 77:669-678. PMC8847543
2. **2021** Aboushousha R, Elko E, Chia SB, Manuel AM, van de Wetering C, van der Velden J, MacPherson M, Erikson C, Reisz JA, D'Alessandro A, Wouters EFM, Reynaert NL, Lam YW, Anathy V, van der Vliet A, Seward DJ, Janssen-Heininger Y. Glutathionylation chemistry promotes interleukin 1 beta-mediated glycolytic reprogramming and pro-inflammatory signaling in lung epithelial cells. **FASEB J**, 35: e21525 PMID: PMC8073242
3. **2021** Yano J, Wells R, Lam YW, Van Houten JL. Two Ca²⁺ Pumps that Regulate Intraciliary Ca²⁺ Following the Action Potential Are Found in the Same Ciliary Membrane Fractions as Ciliary Voltage-Gated Ca²⁺ Channels. **J Exp Biol**, 224: jeb232074
4. **2020** Hasan M, Teixeira JE, Lam YW, Huston CD. Coactosin phosphorylation controls *Entamoeba histolytica* 1 cell membrane protrusions and cell motility **MBio** 11: e00660-20. PMID: PMC7407079
5. **2020** Chia SB, Elko EA, Aboushousha R, Manuel AM, van de Wetering C, Druso JE, van der Velden J, Seward DJ, Anathy V, Irvin CG, Lam YW, van der Vliet A, Janssen-Heininger Y. Dysregulation of the glutaredoxin/S-glutathionylation redox axis in lung diseases. **Am J Physiol Cell Physiol** 318:C304-C327. PMID: PMC7052607
6. **2018** Heppner DE, Dustin CM, Liao C, Hristova M, Veith C, Little AC, Ahlers BA, White SL, Deng B, Lam YW, Li J, van der Vliet A Direct cysteine sulfenylation drives Src kinase activation. **Nature Commun** 9:4522. PMID: 6207713
7. **2018** Wrenn SM, Griswold ED, Uhl FE, Uriarte JJ, Park HE, Coffey AL, Dearborn JS, Ahlers BA, Deng B, Lam YW, Huston DR, Lee PC, Wagner DE, Weiss DJ Avian lungs: A novel scaffold for lung bioengineering. **PLoS One** 13: e0198956. PMID: 6021073
8. **2018** van der Velden JL, Wagner DE, Lahue KG, Abdalla ST, Lam YW, Weiss DJ, Janssen-Heininger YMW TGF-beta1-induced deposition of provisional extracellular matrix by tracheal basal cells promotes epithelial-to-mesenchymal transition in a JNK1-dependent manner. **Am J Physiol Lung Cell Mol Physiol** 314: L984-L997. PMID: PMC6032072
9. **2018** Munson P, Lam YW, MacPherson M, Beuschel S, Shukla A Mouse serum exosomal proteomic signature in response to asbestos exposure. **J Cell Biochem** 119: 6266-6273. PMID: PMC6335961
10. **2018** Munson P, Lam YW, Dragon J, MacPherson M, Shukla A Exosomes from asbestos-exposed cells modulate gene expression in mesothelial cells. **FASEB J**: fj201701291RR. PMID: PMC6044058
11. **2018** Burgess EJ, Hoyt LR, Randall MJ, Mank MM, Bivona JJ, 3rd, Eisenhauer PL, Botten JW, Ballif BA, Lam YW, Wargo MJ, Boyson JE, Ather JL, Poynter ME Bacterial lipoproteins constitute the TLR2-stimulating activity of serum amyloid A. **J Immunol** 201: 2377-2384. PMID: 6179936
12. **2017** Tacoma R, Gelsinger SL, Lam YW, Scuderi RA, Ebenstein DB, Heinrichs AJ, Greenwood SL Exploration of the bovine colostrum proteome and effects of heat treatment time on colostrum protein profile. **J Dairy Sci** 100: 9392-9401. PMID: PMC6350923
13. **2017** Tacoma R, Fields J, Ebenstein DB, Lam YW, Greenwood SL Ratio of dietary rumen degradable protein to rumen undegradable protein affects nitrogen partitioning but does not affect the bovine milk proteome produced by mid-lactation Holstein dairy cows. **J Dairy Sci** 100: 7246-7261. PMID: PMC6350925
14. **2017** Liu Y, Wang BX, Li J, Song ZQ, Lu BG, Chi M, Yang B, Liu JB, Lam YW, Li JX, Xu DY Salt-response analysis in two rice cultivars at seedling stage. **Acta Physiologiae Plantarum** 39: 215. PMID: PMC6858053
15. **2017** Li S, Diego-Limpin PA, Bajrami B, Keshipeddy S, Lam YW, Deng B, Farrokhi V, McShane AJ, Nemati R, Howell AR, Yao X Scaling proteome-wide reactions of activity-based probes. **Anal Chem** 89: 6295-6299. PMID: PMC6368408

16. **2017** Lee CW, Stankowski JN, Chew J, Cook CN, Lam YW, Almeida S, Carlomagno Y, Lau KF, Prudencio M, Gao FB, Bogyo M, Dickson DW, Petrucelli L The lysosomal protein cathepsin L is a progranulin protease. **Mol Neurodegener** 12: 55. PMID: 5526245
17. **2016** Tacoma R, Fields J, Ebenstein DB, Lam YW, Greenwood SL Characterization of the bovine milk proteome in early-lactation Holstein and Jersey breeds of dairy cows. **J Proteomics** 130: 200-210. PMID: 4859431
18. **2016** Tacoma R, Fields J, Ebenstein DB, Lam YW, Greenwood SL Comparative proteomics dataset of skimmed milk samples from Holstein and Jersey dairy cattle. **Data Brief** 6: 843-846. PMID: 4749939
19. **2016** Platz J, Bonenfant NR, Uhl FE, Coffey AL, McKnight T, Parsons C, Sokocevic D, Borg ZD, Lam YW, Deng B, Fields JG, DeSarno M, Loi R, Hoffman AM, Bianchi J, Dacken B, Petersen T, Wagner DE, Weiss DJ Comparative decellularization and recellularization of wild-type and alpha 1,3 galactosyltransferase knockout pig lungs: a model for ex vivo xenogeneic lung bioengineering and transplantation. **Tissue Eng Part C Methods** 22: 725-739. PMID: 4991572
20. **2016** Krishnamurthy S, Deng B, Del Rio R, Buchholz KR, Treeck M, Urban S, Boothroyd J, Lam YW, Ward GE Not a simple tether: binding of *Toxoplasma gondii* AMA1 to RON2 during invasion protects AMA1 from rhomboid-mediated cleavage and leads to dephosphorylation of its cytosolic tail. **MBio** 7. PMID: 5021801
21. **2016** Smith KP, Fields JG, Voogt RD, Deng B, Lam YW, Mintz KP Alteration in abundance of specific membrane proteins of *Aggregatibacter actinomycetemcomitans* is attributed to deletion of the inner membrane protein MorC. **Proteomics** 15: 1859-67. PMID: 4456248
22. **2015** Smith KP, Fields JG, Voogt RD, Deng B, Lam YW, Mintz KP The cell envelope proteome of *Aggregatibacter actinomycetemcomitans*. **Mol Oral Microbiol** 30, 97-110 PMID: 4305030
23. **2014** Wagner DE, Bonenfant NR, Sokocevic D, DeSarno MJ, Borg ZD, Parsons CS, Brooks EM, Platz JJ, Khalpey ZI, Hoganson DM, Deng B, Lam YW, Oldinski RA, Ashikaga T, Weiss DJ Three-dimensional scaffolds of acellular human and porcine lungs for high throughput studies of lung disease and regeneration. **Biomaterials** 35: 2664-2679.
24. **2014** Wagner DE, Bonenfant NR, Parsons CS, Sokocevic D, Brooks EM, Borg ZD, Lathrop MJ, Wallis JD, Daly AB, Lam YW, Deng B, DeSarno MJ, Ashikaga T, Loi R, Weiss DJ Comparative decellularization and recellularization of normal versus emphysematous human lungs. **Biomaterials** 35: 3281-3297.
25. **2014** Hristova M, Veith C, Habibovic A, Lam YW, Deng B, Geiszt M, Janssen-Heininger YM, van der Vliet A Identification of DUOX1-dependent redox signaling through protein S-glutathionylation in airway epithelial cells. **Redox Biol** 2: 436-446. PMID: 3949091
26. **2014** Cherian S, Wai Lam Y, McDaniels I, Struziak M, Delay RJ Estradiol rapidly modulates odor responses in mouse vomeronasal sensory neurons. **Neuroscience** 269: 43-58. PMID: PMC4270699
27. **2013** Syed K, Porollo A, Lam YW, Grimmett PE, Yadav JS (2013) CYP63A2, a catalytically versatile fungal P450 monooxygenase capable of oxidizing higher-molecular-weight polycyclic aromatic hydrocarbons, alkylphenols, and alkanes. **Appl Environ Microbiol** 79: 2692-2702. PMID: 3623170
28. **2013** Sokocevic D, Bonenfant NR, Wagner DE, Borg ZD, Lathrop MJ, Lam YW, Deng B, Desarno MJ, Ashikaga T, Loi R, Hoffman AM, Weiss DJ (2013) The effect of age and emphysematous and fibrotic injury on the re-cellularization of de-cellularized lungs. **Biomaterials** 34: 3256-3269.
29. **2013** Purmessur D, Guterl CC, Cho SK, Cornejo MC, Lam YW, Ballif BA, Laudier JC, Iatridis JC Dynamic pressurization induces transition of notochordal cells to a mature phenotype while retaining production of important patterning ligands from development. **Arthritis Res Ther** 15: R122. PMID: 3978427
30. **2013** Bonenfant NR, Sokocevic D, Wagner DE, Borg ZD, Lathrop MJ, Lam YW, Deng B, Desarno MJ, Ashikaga T, Loi R, Weiss DJ (2013) The effects of storage and sterilization on de-cellularized and re-cellularized whole lung. **Biomaterials** 34: 3231-3245.

31. **2012** Lam HM, Suresh Babu CV, Wang J, Yuan Y, Lam YW, Ho SM, Leung YK Phosphorylation of human estrogen receptor-beta at serine 105 inhibits breast cancer cell migration and invasion. **Mol Cell Endocrinol** 358: 27-35. PMID: 3348253
32. **2012** Isaac J, Tarapore P, Zhang X, Lam YW, Ho SM Site-specific S-nitrosylation of integrin alpha6 increases the extent of prostate cancer cell migration by enhancing integrin beta1 association and weakening adherence to laminin-1. **Biochemistry** 51: 9689-9697. PMID: 3567210
33. **2011** Syed K, Porollo A, Lam YW, Yadav JS A fungal P450 (CYP5136A3) capable of oxidizing polycyclic aromatic hydrocarbons and endocrine disrupting alkylphenols: role of Trp(129) and Leu(324). **PLoS One** 6: e28286. PMID: 3229547

prior to 2011

34. **2010** Syed K, Doddapaneni H, Subramanian V, Lam YW, Yadav JS Genome-to-function characterization of novel fungal P450 monooxygenases oxidizing polycyclic aromatic hydrocarbons (PAHs). **Biochem Biophys Res Commun** 399: 492-497. PMID: 2943217
35. **2010** Lam YW, Yuan Y, Isaac J, Babu CV, Meller J, Ho SM Comprehensive identification and modified-site mapping of S-nitrosylated targets in prostate epithelial cells. **PLoS One** 5: e9075. PMID: 2816712
36. **2009** Xiong SD, Yu K, Liu XH, Yin LH, Kirschenbaum A, Yao S, Narla G, DiFeo A, Wu JB, Yuan Y, Ho SM, Lam YW, Levine AC Ribosome-inactivating proteins isolated from dietary bitter melon induce apoptosis and inhibit histone deacetylase-1 selectively in premalignant and malignant prostate cancer cells. **Int J Cancer** 125: 774-782. PMID: 3778503
37. **2008** Lam YW, Tam NN, Evans JE, Green KM, Zhang X, Ho SM Differential proteomics in the aging Noble rat ventral prostate. **Proteomics** 8: 2750-2763.
38. **2005** Lam YW, Mobley JA, Evans JE, Carmody JF, Ho SM Mass profiling-directed isolation and identification of a stage-specific serologic protein biomarker of advanced prostate cancer. **Proteomics** 5: 2927-2938.
39. **2004** Mobley JA, Lam YW, Lau KM, Pais VM, L'Esperance JO, Steadman B, Fuster LM, Blute RD, Taplin ME, Ho SM Monitoring the serological proteome: the latest modality in prostate cancer detection. **J Urol** 172: 331-337
40. **2003** Mobley JA, Leav I, Zielie P, Wotkowitz C, Evans J, Lam YW, L'Esperance BS, Jiang Z, Ho SM Branched fatty acids in dairy and beef products markedly enhance alpha-methylacyl-CoA racemase expression in prostate cancer cells in vitro. **Cancer Epidemiol Biomarkers Prev** 12: 775-783.
41. **2002** Lam YW, Ng TB Purification and characterization of a rhamnose-binding lectin with immunoenhancing activity from grass carp (*Ctenopharyngodon idellus*) ovaries. **Protein Expr Purif** 26: 378-385.
42. **2001** Lam YW, Ng TB, Wang HX Antiproliferative and antimitogenic activities in a peptide from puffball mushroom *Calvatia caelata*. **Biochem Biophys Res Commun** 289: 744-749.
43. **2001** Lam YW, Ng TB A monomeric mannose-binding lectin from inner shoots of the edible chive (*Allium tuberosum*). **J Protein Chem** 20: 361-366.
44. **2000** Lam YW, Wang HX, Ng TB A robust cysteine-deficient chitinase-like antifungal protein from inner shoots of the edible chive *Allium tuberosum*. **Biochem Biophys Res Commun** 279: 74-80.

MANUSCRIPTS SUBMITTED / IN PREPARATION

1. **2022** Chandrasekaran R, Bruno S, Mark Z, Walzer J, Caffry S, Gold C, Kumar A, Chamberlain N, Lam YW, van der Vliet A, Janssen-Heininger Y, Poynter M, Dixon AE, Anathy V. MitoQ alleviates airway inflammation, remodelling, hyperresponsiveness and tissue eosinophilia in obese allergic asthma, **submitted**

2. **2022** Corteselli EM, Sharafi M, MacPherson M, White S, Lam YW, Gold C, Manuel A, van der Vliet A, Schneebeli ST, Anathy V, Li J, and Janssen-Heininger Y. Structural and functional fine mapping of cysteines in mammalian glutaredoxin reveal a hierarchy of susceptibility to oxidative inactivation, **submitted**
3. **2022** Chandra H, Gupta ML, Lam YW, and Yadav JS. A Predominantly Orphan Secretome in Mycobacterium abscessus as Revealed by a Multi-pronged Strategy. **In preparation**

TALKS

Gave talks/seminars covering proteomics technologies and applications since 2011.

- 2022 Vermont Biomedical Research Network Retreat**
“Overview of the VBRN Proteomics Facility” (Jul. 21)
- 2021 Northeast Regional Laboratory Staff and Core Directors (NERLSCD) Meeting (Portsmouth, NH) Nov. 3 to Nov. 5, 2021.**
“Proteomics 101” (Nov. 4)
- 2020 Symposium for Proteomics Core Directors and Staff (IDeA National Resource for Proteomics)**
“Crosslinked peptide analysis” (Jan. 29)
- 2019 Northeast Regional IDeA Conference (NERIC)**
Section: Core presentations of resources available to NERIC investigators
“Overview of the VGN Proteomics Facility” (Aug. 15)
- 2019 Symposium for Proteomics Core Directors and Staff (IDeA National Resource for Proteomics)**
“Overview of the VGN Proteomics Facility” (Feb. 22)
- 2018 Webinar (for Community College of Vermont)**
“Proteomics” (Nov. 30)
- 2018 University of Vermont, Department of Biology**
“VGN Proteomics Facility: How can we help you?” (Sept. 10)
- 2018 Vermont Genetics Network Retreat**
“Overview of the VGN Proteomics Facility” (Aug. 15)
- 2017 Webinar (for primarily undergraduate institutions in Vermont)**
“From Basic to Advanced Proteomic Applications and How They Might Benefit Your Research: UVM Proteomics” (Oct. 12)
- 2015 New England Regional Shared Resources Conference (NERSRC) at Dartmouth**
“Brief Overview of the VGN Proteomics” (Nov. 12)
- 2015 Vermont Genetics Network Retreat**
“Proteomics” (Aug. 12)
- 2014 University of Vermont / Department of Pathology / Redox Biology Meeting**
“How Can Proteomics Help Your Research? Expression, PTMs, Metabolites, and More” (Oct. 27)
- 2014 Saint Michael’s College / Department of Chemistry**
“Proteomics at UVM” (Apr. 11)
- 2012 Vermont Cancer Center Basic Science/Translational Strategic Planning Meeting**
“UVM/VGN Proteomics” (Sep. 29)
- 2012 Vermont Cancer Center Population Health Sciences Strategic Planning Meeting**
“UVM/VGN Proteomics” (Sep. 8)
- 2012 University of Vermont / Department of Pathology / Environmental Pathology and Carcinogenesis Seminar Series**
“How Can Proteomics Help Your Research? Expression, PTMs, Metabolites, and More” (Feb. 27)
- 2011 University of Vermont / Behavior and Health Focus Group Meeting**
“How Can Proteomics Help Your Research? Expression, PTMs, Metabolites, and More” (Dec. 13)
- 2011 University of Vermont / VGN Proteomics Facility Open House**
“How Can Proteomics Help Your Research? Expression, PTMs, Metabolites, and More” (Nov. 15)

- 2011 University of Vermont / Department of Biochemistry**
 "How Can Proteomics Help Your Research? Expression, PTMs, Metabolites, and More" (Oct. 21)
- 2011 University of Vermont / Department of Molecular Physiology and Biophysics**
 "How Can Proteomics Help Your Research? Using Mass Spectrometry as a Tool" (Oct. 10)
- 2011 University of Vermont / Department of Biology / "Cell Lunch" Graduate Student Seminars**
 How Do I Apply Proteomics to My Research? "Stop by MLS 335" (Sep. 21)
- 2011 University of Vermont / Department of Biology**
 "How Can Proteomics Help Your Research? Using Mass Spectrometry as a Tool" (Sept. 19)
- 2011 Vermont Genetics Network Retreat**
 "How Can Proteomics Help Your Research? Interrogate the Proteome at Multiple Levels" (Aug. 17)
- 2011 University of Cincinnati / Department of Environmental Health**
 "How Can Proteomics Help Your Research? Take Prostate Cancer as an Example" (May 25)
- 2011 University of Vermont / Department of Biology**
 "How Can Proteomics Help Your Research? Take Prostate Cancer as an Example" (Mar. 07)
- 2011 Miami University / Department of Chemistry and Biochemistry**
 "Proteomics Applications in Prostate Research". (Feb. 25)

prior to 2011

- 2010 7th Annual Ohio Mass Spectrometry Symposium, Columbus, OH.**
 "Nitrosative Stress in the Prostate Epithelial Cells". (Apr. 19-20)
- 2009 6th Annual Ohio Mass Spectrometry Symposium, Columbus, OH.**
 "Comprehensive Identification and Modified Site Mapping of Nitrosylated Proteins in Normal Prostate Cells". (Apr. 20-21)
- 2007 39th Central Regional Meeting of the American Chemical Society (Mass Spectrometry II Session), Covington, KY.**
 "Differential Proteomics in the Aging Noble Rat Ventral Prostate: Profiling by Isotope-Coded Affinity Tags and Mass Spectrometry". (May 20-23)
- 2006 University of Cincinnati / Prostate Cancer Working Group Symposium**
 "Mass Spectrometry-Based Proteomics Technologies in Molecular Pathway Analysis". (June 22)
- 2006 3rd Annual Ohio Mass Spectrometry Symposium, Columbus, OH. Extended Poster (podium) Presentation**
 "Comparison of Relative Protein Levels in the Prostates of Young and Old Noble Rats by ICAT". (Mar. 20-21)
- 2005 University of Cincinnati / Prostate Cancer Working Group Symposium**
 "Proteomic Application in Prostate Cancer Research: Biomarker discovery in Serum". (Dec. 6)

POSTER PRESENTATIONS

Presented or contributed to >30 facility/research posters since June 2011.

FACILITY POSTERS

These facility posters were presented in 5 conferences to promote the facility (Northeast Regional IDeA Meeting, Biennial National IDeA Symposium of Biomedical Research Excellence, Northeast Regional Life Sciences Core Directors Meeting, Dartmouth Shared Resources Fair, and Annual Vermont Genetics Network Retreat).

National

Association of Biomolecular Resource Facilities (ABRF) Annual Meeting

- 2019** Lam YW, Deng B, Hood C. Vermont Genetics Network Proteomics Facility. San Antonio, TX (Mar. 23 – 26) J Biomol Tech. 2019 Dec;30 (Suppl):S41.

Biennial National IDEa Symposium of Biomedical Research Excellence (NISBRE), Washington, DC

- 2018 Lam YW, Deng B, Hood C. Vermont Genetics Network Proteomics Facility. (June 24-26)
iPoster: <https://nisbre2018-nisbre.ipostersessions.com/default.aspx?s=9A-EC-A3-B2-CF-7E-01-A3-8F-E0-C9-4D-EB-B8-1A-E1>
- 2016 Deng B, Ahlers B, Lam YW. Driving to a sustainable proteomics core facility. (June 26-28)

Regional**Northeast Regional IDEa Conference (NERIC)**

- 2019 Lam YW, Deng B, Hood C. Vermont Genetics Network Proteomics Facility. Bretton Woods, New Hampshire (Aug. 16-18)
- 2017 Lam YW, Deng B, Hood C. Vermont Genetics Network Proteomics Facility. Burlington, Vermont (Aug. 16-18)
- 2015 Fields J, Deng B, Lam YW. Growing Success - Vermont Genetics Network Proteomics Facility. Bar Harbor, ME (Sept. 24-26)
- 2015 Vincent J, Hunter T, Murray J, Driscoll H, Deng B, Lam YW, Van Houten J. Highlights of the Vermont Genetics Network. Bar Harbor, ME (Sept. 24-26)
- 2013 Fields J, Deng B, Lam YW. Growing success in the Vermont Genetics Network Proteomics Facility. Newark, Delaware (Aug. 21-23)
- 2011 Deng B, Lam YW. The establishment and growth of Vermont Genetics Network Proteomics Facility. New Port, Rode Island (Aug. 10-12)

Northeast Regional Life Sciences Core Directors (NRLSCD) Meeting

- 2015 Uhl F, Zvarova B, Fields J, Deng B, Lam YW, Weiss D, Wagner D. Enhanced mass spectrometry proteomics can be used to distinguish differences in protein compositions of acellular emphysematous versus normal lungs. Burlington, VT (Oct. 14-16)
- 2015 Fields J, Lam YW, Deng B, Smith K, Voogt R, Mintz K. Quantification of the membrane differential proteomes by stable isotope labeling. Burlington, VT (Oct.14-16)
- 2015 Lam YW, Fields J, Deng B. Growing Success - Vermont Genetics Network Proteomics Facility. Burlington, VT (Oct. 14-16)

Conferences at Dartmouth College

- 2015 **New England Regional Shared Resources Conference (NERSRC)** Lam YW, Fields J, Deng B. VGN Proteomics. (Nov. 12)
- 2013 **Dartmouth Shared Resources Fair.** Fields J, Deng B, Lam YW. Proteomics at UVM. (Oct. 9)

Local**Annual Vermont Genetics Network Retreat, South Burlington, VT**

- 2014 Fields J, Deng B, Lam YW. Growing success in the Vermont Genetics Network Proteomics Facility. (Aug. 13)
- 2013 Fields J, Deng B, Lam YW. Growing Success in the Vermont Genetics Network Proteomics Facility. (Aug. 7)
- 2012 Lam YW, Deng B. Growing success in the Vermont Genetics Network Proteomics Facility. (Aug. 15)

RESEARCH POSTERS

These posters describe proteomics applications or method development.

National**ASMS Conference on Mass Spectrometry and Allied Topics (ASMS)**

- 2017 Krishnamurthy S, Deng B, del Rio R, Buchholz KR, Treeck M, Urban S, Boothroyd J, Lam YW, Ward GE. Dephosphorylation of AMA1 cytosolic tail during *Toxoplasma gondii* invasion. Indianapolis, IN (June 19-22)

- 2015** Lam YW, Deng B, Fields J, Smith K, Voogt R, Mintz K. Quantification of the membrane differential proteomes by stable isotope labeling and spectral counting strategies. St. Louis, MO (May 31-June 4)
- 2013** Li S, Diego P, Keshipeddy S, Bajrami B, Farrokhi V, McShane A, Lam YW, Deng B, Nemati R, Howell A, Yao X. High-throughput scheduled MRM for multiplexed quantitation of chemical probe labeled enzymes in human cells. Minneapolis, MN (June 9-13)
- 2012** Li S, Keshipeddy S, Diego P, Bajrami B, Mcshane A, Deng B, Lam YW, Howell AR, Yao X. Using chemical probes and multiple reaction monitoring mass spectrometry for monitoring multiple enzymes in human cells. Vancouver, Canada (May 20-24)

prior to 2011

- 2010** Lam YW, Isaac J, Yuan Y, Suresh Babu CV, Meller J, Ho SM. Comprehensive identification and modified-site mapping of S-nitrosylated targets in normal prostate cell Line. Salt Lake City, Utah (May 23-27)
- 2005** Lam YW, Sun P, Green KM, Evans JE, Ho SM. Protein profiling of serum from prostate cancer patients by isotope-coded affinity tag (ICAT) analysis. San Antonio, Texas (June 5-9)
- 2004** Lam YW, Tam NNC, Sun P, Green KM, Evans JE, Ho SM. Comparison of relative protein levels in prostate from young and old Noble rat by ICAT. Nashville, Tennessee (May 23-27)

Other conference presentations prior to 2011

- 2009** Xiong SD, Liu X-H, Yin HF, Kirschenbaum A, Yao S, Narla G, DiFeo A, Yu K, Wu JB, Yuan Y, Ho S-M, Lam YW, Levine AC. Ribosome-inactivating proteins isolated from dietary bitter melon induce apoptosis and inhibit histone deacetylase-1 selectively in premalignant and malignant prostate cancer cells. **Annual Meeting of the American Association for Cancer Research (AACR)**, Denver, CO (Apr. 2009)
- 2006** Lam YW, Tam NNC, Green KM Evans JE, Ho SM. Comparison of relative protein levels in the prostates of young and old Noble rats by ICAT. **3rd Annual Ohio Mass Spectrometry Symposium**, Columbus, Ohio (Mar. 20-21)
- 2005** Tam NN, Lam YW, Ho SM. Oxidative and nitrosative stresses and inflammation as early events of hormone-induced prostatic carcinogenesis in Noble rats. **American Urological Association/Society for Basic Urologic Research Summer Research Conference**, "Inflammation in Prostate Diseases", Baltimore, DC (Aug. 19-20)
- 2004** Lam YW, Mobley JA, Evans JE, Carmody JF, Ho SM. Mass profiling and multidimensional separation of serologic proteome of patients with advanced disease. **95th American Association for Cancer Research (AACR) Annual Meeting**, Orlando, Florida (Mar. 27-31)
- 2002** Lam YW, Mobley JA, Carmody JF, Ho SM. Proteomics approach to evaluate the impact of diet on prostate cancer progression. **9th Prouts Neck Meeting on Prostate Cancer**, Maine (Nov. 7-10)

SELECTED POSTERS FROM COLLABORATIONS

These posters cover a wide range of research areas of collaborators' interest, including lung physiology, neuroscience, parasitology, periodontal diseases, stem cell biology, redox biology, animal science, and mass spectrometry.

- 2021** Biomedical Research Institute of New Jersey
- Memon N, Eckman E, Lam YW, Herdt A, Lee C. Proteomic Alterations in the Plasma of Rat Pups Exposed to Phytosterols. **Pediatric Academic Societies 2021 Virtual Meeting** (Apr. 30 – Jun. 4)
- 2020** Department of OBGYN and Reproductive Sciences, UVM
- Ko NL, Mukhtarova N, Hood C, Lam YW, Osol G Restricted Uterine Venous Blood Flow Results in Attenuated Uterine Vascular Remodeling and Preeclampsia-Related Placental Secretomics During Pregnancy. **Society of Reproductive Investigation (SRI) Annual meeting**, Vancouver, BC, Canada (Mar. 10 – 14) (Canceled due to Covid-19)
- 2019** Department of Medicine: Immunobiology, UVM
- Teixeira JE, Hasan MM, Lam YW, Huston CD Coactosin phosphorylation controls *Entamoeba histolytica* cell membrane protrusions and cell motility. **Annual Woods Hole Molecular Parasitology Meeting**, Woods Hole, MA (Sept. 15-19)

2018 Department of Pathology, UVM

- Manuel A, Qian X, Chia SB, Aboushousha R, van de Wetering C, van der Velden J, Dixon AE, Lam YW, Irvin CG, Janssen-Heininger YMW. S-Glutathionylation of pyruvate kinase M2 is associated with metabolic reprogramming and cytokine production in the development of allergic obese airway disease. **25th Annual Conference of the Society for Redox Biology and Medicine**. Chicago, IL. (Nov. 14-17)

Department of Medicine: Pulmonary Disease & Critical Care Medicine, UVM

- Uhl FE, Zvarova B, Ahlers BA, Hood CM, Deng B, Lam YW, Beatman E, Schweitzer KS, Petrache I, Weiss DJ, Wagner DE. Characterization of decellularized COPD lung matrices using mass spectrometry proteomics. **16th Lung Science Conference**, Estoril, Portugal (Mar. 8-11)

2017 Department of Pharmacology, UVM

- Madasu S, Lam YW, Morielli A. Quantitative MS identifies potential targets for cerebellar disorders. **Discovery on Target**, Boston, MA (Sept. 25-29)

Department of Pathology, UVM

- Chia SB, Aboushousha R, Qian X, Lahue KG, Lam YW, Irvin CG, Anathy V, Poynter M, van der Vliet A, Heininger YMW. Glutaredoxin-1 regulates airway basal cell plasticity. **Gordon Research Conference** (Lung Development, Injury and Repair), New London, NH. (Aug. 20-25)

Department of Medicine: Pulmonary Disease & Critical Care Medicine, UVM

- Wrenn SM, Griswold ED, Uhl FE, Uriarte JJ, Park HE, Coffey AL, Dearborn JS, Hommel RJ, Lam YW, Deng B, Ahlers BA, Lee PC, Huston DR, Wagner DE, Weiss DJ. Lung bioengineering using avian tissue. **Stem Cell Conference**, Burlington, VT (July 24-27)
- Wrenn SM, Griswold ED, Uhl FE, Coffey AL, Dearborn JS, Hommel RJ, Lam YW, Deng B, Ahlers BA, Wagner DE, Weiss DJ. Avian lung de-and recellularization: A novel biomaterial for pulmonary therapeutics. **American Thoracic Conference**, Washington, DC (May 19-24)

Department of Animal Science, UVM

- Scuderi RA, Ebenstein DB, Lam YW, Kraft J, Greenwood SL. The effect of dietary grape marc on the bovine milk proteome. **ADSA Annual Meeting**, Pittsburgh, PA (June 25)

2016 Department of Pathology, UVM

- Chia SB, Nolin JD, Qian X, Aboushousha R, Lahue KG, van der Velden J, Schneider R, Lam YW, Irvin CG, Anathy V, van der Vliet A, Heininger YMW. Glutaredoxin-1 regulates allergic remodeling in house dust mite-induced allergic airways disease. **Gordon Research Conference** (Thiol-Based Redox Regulation and Signaling), Stowe, VT (Aug. 7-12)
- Heppner DE, Dustin CM, Liao C, Hristova M, Veith C, Little AC, Ahlers BA, White SL, Deng B, Lam YW, Li J, van der Vliet A. Molecular origin of thiol-based redox-regulation of the Src Kinase. **Gordon Research Conference** - Thiol-Based Redox Regulation and Signaling, Stowe, VT. (Aug. 7-12)
- Heppner DE, Dustin CM, Liao C, Hristova M, Veith C, Little AC, Ahlers BA, White SL, Deng B, Lam YW, Li J, van der Vliet A. Molecular origin of redox regulation of the Src kinase. **23rd National Meeting of the Society for Redox Biology and Medicine**, San Francisco, CA (Nov. 16 - 19)

Department of Medicine: Pulmonary Disease & Critical Care Medicine, UVM

- Uhl F, Zvarova B, Fields J, Deng B, Lam YW, Weiss D, Wagner D. Enhanced mass spectrometry proteomics can be used to distinguish differences in protein compositions of acellular emphysematous versus normal lungs. **American Thoracic Conference**, San Francisco, CA (May 13-18, 2016)
- Uhl F, Costa R, Zvarova B, Fields JG, Deng B, Lam YW, Königshoff M, Weiss DJ, Wagner DE. Acellular emphysematous lungs retain pathophysiological attributes and can be used as a novel *in vitro* culture system for studying disease biology. **German Lung Center (DZL) conference**. (Jan. 30-31)

2015 Department of Animal Science, UVM

- Tacoma R, Lam YW, Fields JG, Greenwood S. Characterization of the bovine milk proteome produced by Holstein and Jersey breeds of dairy cows. **ADSA-ASAS Joint Annual Meeting**, Orlando, FL (July 12-16).

2014 Department of Biochemistry, UVM

- Robey-Bond S, Lam YW, Ebert A, Fields J, Lefebvre P, Francklyn C. Molecular basis of histidyl-tRNA synthetase-associated Usher Syndrome Type 3B in cochlear-derived mouse cells and zebrafish. **25th tRNA Conference**, Kyllini, Greece (Sept. 21-25)
- Robey-Bond S, Fields J, Lam YW, Francklyn C. Identification of protein networks disrupted by a mutation in HARS. **Neuroscience, Behavior and Health Forum**, UVM (Jan. 25)

2013 Department of Biochemistry, UVM

- Abbott J, Deng B, Lam YW, Francklyn CS, Robey-Bond S. The role of human histidyl-tRNA synthetase mutations in human diseases, Type IIIB Usher Syndrome and Peripheral Neuropathy. **RNA Metabolism in Neurological Disease Conference** (8th Brain Research Conference), San Diego, CA (Nov. 7-8)
- Robey-Bond S, Fields J, Lam YW, Francklyn C. Identification of protein networks disrupted by a mutation in HARS. **9th International Symposium on Aminoacyl-tRNA Synthetases Conference**, Hakone, Japan (Oct. 6 – 11)

Department of Microbiology and Molecular Genetics, UVM

- Smith K, Voogt R, Lam YW, Mintz K. Membrane proteome changes in *Aggregatibacter actinomycetemcomitans* dependent on one protein. **Penn Periodontal Conference**, Philadelphia, PA (June 23-28)

Department of Biochemistry, Middlebury College

- Borsinger T, Lam YW, Cluss B. Initial determination of the proteome of *Borrelia burgdorferi*, the Lyme disease spirochete. **Vermont Genetics Network Undergraduate Student Career Day**, South Burlington (Apr. 17)

PROFESSIONAL DEVELOPMENT

Courses, workshops, on-site trainings and other informational meetings attended.

	<u>Organizer</u>	<u>Topic</u>
2022	Thermo Fisher Scientific	3 - day on-site Eclipse and Exploris 240 training (Jul. 26-28 & Sep. 13 - 15)
2021	VT IDeA Retreat	StrategicDoing Workshop (Aug. 19)
2021	Arkansas/Oklahoma IDeA	Symposium for Proteomics Core Directors and Staff (Jan.28 - 31)
2020	Celdara Medical Inc./NIGMS/ DRIVEN	Core Business Virtual Meeting (Oct. 5)
2020	UVM	Online Teaching Boot Camp (Aug. 4)
2020	ASMS	2 - day virtual conference short course: LC-MS: Practical Maintenance and Troubleshooting
2020	Arkansas/Oklahoma IDeA	Symposium for Proteomics Core Directors and Staff (Jan.28 - 31)
2019	Maine INBRE	Bioinformatics (T3): Train the Trainer (Jun. 29 - July 6)
2019	ASMS	2 - day conference short course: Native Mass Spectrometry
2019	Arkansas/Oklahoma IDeA	Symposium for Proteomics Core Directors and Staff (Feb. 20 - 23)
2018	May Institute	Computation and Statistics for Mass Spectrometry and Proteomics (Apr. 30 – May 9)
2018	Arkansas/Oklahoma IDeA	Symposium for Proteomics Core Directors and Staff (Apr. 4 - 5)
2017	ASMS	2 - day conference short course: DMPK: Experimentation and Data Interpretation
2017	Arkansas/Oklahoma IDeA	Symposium for Proteomics Core Directors and Staff (Apr. 4 - 6)
2016	ASMS	2 - day conference short course: Protein Therapeutics: Practical Characterization and Quantitation by Mass Spectrometry
2015	Thermo Fisher Scientific	3 - day on-site Q-Exactive training (Sep. 8 - 10)
2015	ASMS	2 - day conference short course: Protein Structural Analysis by Mass Spectrometry: Hydrogen Exchange and Covalent Labeling
2014	ASMS	2 - day conference short course: Ion Mobility in Mass Spectrometry
2014	Waters Corporation	SYNAPT G2-Si Workshop in Salem, MA (Feb. 19)
2014	AB SCIEX	New Developments in Clinical Research and Forensic Toxicology Using LC/MS/MS (Apr. 22)

2013	ASMS	2 - day conference short course: Glycans and Glycoproteins in Mass Spectrometry
2013	Thermo Fisher Scientific	MS Thought Leader Summit (Feb. 26)
2012	ASMS	2 - day conference short course: Metabolomics
2010	ASMS	2 - day conference short course: High Resolution LC-MS for Structural Identification and Quantitation
2009	ASMS	2 - day conference short course: Protein Structural Analysis by Mass Spectrometry: Hydrogen Exchange and Covalent Labeling
2007	ASMS	2 - day conference short course: FTMS: Principles and Applications
2006	ASMS	2 - day conference short course: Mass Spectrometry of Peptides and Proteins
2006	Institute for Systems Biology	2 - day Proteomics Informatics Course (May 25 - 26)
2006	University of Louisville	Metabolomics Symposium
2005	ASMS	2 - day conference short course: Quadrupole Ion Trap Mass Spectrometry
2004	ASMS	2 - day conference short course: Practical LCMS: Fundamentals, Techniques and Applications

RESEARCH SUPPORT

The VBRN Proteomics Facility is supported by P20GM103449 and operates with a cost-recovery model prior July 1st, 2021. Consumables are provided by investigators via various grants that support the proteomics projects. The facility operations has transitioned into an income expense model since July 1st, 2021.

Ongoing Research Support

5P20GM103449-19 Rex Forehand (PI) 06/01/20 - 05/31/25
NIH/NIGMS
Vermont INBRE

The major goal of this grant is to increase the culture of research in the state's four-year institutions by facilitating faculty members' research/grant productivity and undergraduate education in the sciences.
Role: Core Facility Director

Completed Research Support

P20 GM103449 Rex Forehand (PI) 06/01/15 - 05/31/20
NIH/NIGMS
INBRE Vermont Genetics Network

The goal of this grant is to invest in the Vermont biomedical research infrastructure, including physical and human resources, in order to bring about sustainable changes in how we in Vermont carry out research and educate our next generation of scientists and doctors.
Role: Core Facility Director

P20 GM103449 formerly P20RR16462 Van Houten (PI) 06/01/11 - 05/31/15
NIH/NIGMS

The goal of this grant is to invest in the Vermont biomedical research infrastructure, including physical and human resources, in order to bring about sustainable changes in how we in Vermont carry out research and educate our next generation of scientists and doctors.
Role: Core Facility Director