

Research from value added analytical matter

Benjamin Katz

Pet Food Innovation Workshop

September 14th 2016



Acknowledgements

PI : Dr. John Tomich

Dr. Takeo Iwamoto

Nozomi M. Caton

Deane Lehman

Susan Whitaker

Past Students

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Kyle Steuber (Nutrisciences)

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Yanting Shen

Dr. Ying Fang

Rui Guo

Thomas Gallagher

External

Mike Molitor (U.Wisc. Madison)

Dan Abitz (GBA Engineering)

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Kansas State University

Kansas State University Biotech Core

- Non-Profit initiative in the Vice President for Research Office.
- Work with >70 KSU investigators in a wide range of disciplines.
- Facilitate collaboration between academy and industry.
- Goal: Help promote new product design and entrepreneurship in the Midwest.

Kansas State University Biotech Core

- **Equipment**

- Analytical and Preparative HPLC
- MALDI-TOF and ESI-MS/MS
- Proteomics and Metabolomics
- Wet Lab, Roto-vap Distillation, Lyophilization
- Microbiology culture, Fermentation Scale up
- Super Critical Fluid Extraction and Spray Drying

- **Types of Projects**

- Animal Nutrition and Disease
- Food and Human Health
- Agricultural Crops: Pest, Disease and Tolerance
- Biosecurity BSL3ag support: Animal and Zoonosis
- Fundamental Biology and Ecology
- Classical Chemistry and Peptide Synthesis

Analyzing “Enriched” Protein Products

- Fingerprinting hydrolysates for peptides signature
 - Using MALDI-TOF
 - Like Bruker Biotyper for microbes
- Link nutrition information to product name
 - Enriched with...
(Glutamine, BCAA, Sulfur AA...etc)
 - Blending example: Enriched Sunflower Meal



Outline of Talk

- By-product protein streams
- Negative image from consumers
- Rational design of hydrolysates
- Analytical service offered by KSU biotech core
- Controlling waste stream consistency
- Fingerprinting hydrolysates in blends
- Examples from industry

Grain Storage Proteins (prolamins)

- Great source of metabolic energy
- Essential source of nitrogen and sulfur rich amino acids
- Inherent poor digestibility
- Due to low/missed cleavage site by digestive enzymes
- Presences of trypsin inhibitors
- Negative image as cheap filler

Negative image from consumers

Consumer have been taught to AVOID:

- SOY FLOUR, WHOLE WHEAT FLOUR, CORN, CORN GLUTEN MEAL
- WHEAT BRAN, BREWER'S RICE, POTATO PRODUCT, MIDDLINGS/MIDS or MILL RUN of any kind.
- Unspecified grain sources like CEREAL FOOD FINES, CORN BRAN, OAT HULLS, RICE HULLS, PEANUT HULLS, DISTILLERS GRAIN FERMENTATION SOLUBLES, and CELLULOSE
- MEAT AND BONE MEAL, ANIMAL DIGEST from "4D" animals

Examples of protein by-products

- Corn Gluten Meal
 - Roquette America \$520.00/ton FOB IOWA
 - 60% Protein @ 43c/lb protein
 - Zein prolamin proteins

- Wheat Midds, dried
 - ADM \$88.00/ton FOB Kansas
 - 16% Protein @ 28c/lb protein
 - Gliadin prolamin proteins

[Waste Streams updated list](#)

Prolamin processing

- Soluble in ~70% ethanol
- Difficult processing due to aggregation
- Insoluble aggregate floats and sink in water
 - Heavy and light fractions
- Difficult for enzymes to access cleavage sites
- Processing can cause bad flavors
- Denatured proteins cleave differently

Rational Prolamin Processing

- Utilize fungal/plant/bacterial enzymes
 - Able to cleave proteins at unique sites
 - Proteases
 - Help solubilized bulk protein
 - Endo & exo peptidases
 - Tailor flavor by eliminate low threshold taste modulators
 - Improve nutrition

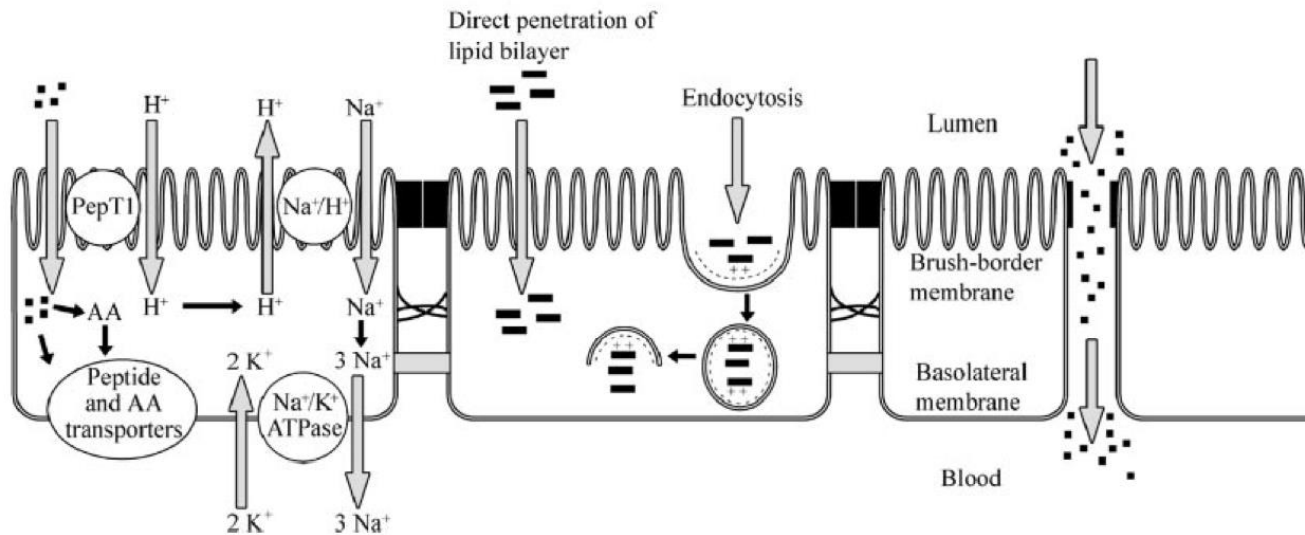
Amino Acid Nutrition

Gilbert et al.

A. Active transport via PepT1 transporter

B. Transcellular movement of CPP with peptides as cargo

C. Paracellular movement



Group I: Hydrolysate > amino acid mix. > protein

Essential amino acids

Nonessential amino acid

Thr, Lys

Tyr, His, Ser, Asx

Group II: Hydrolysate = amino acid mix. > protein

Essential amino acids

Nonessential amino acid

Ile, Leu, Val, Met, Phe

Arg, Ala, Gly, Glx, Cys

Group III: Amino acid mix. > hydrolysate > protein

Essential amino acids

Nonessential amino acid

—

Pro

Rational Prolamin Processing

- Use analytical data to create multiple enzyme processes.
 - Eliminate known bad flavors and indigestible fragments
 - Completely digest into di/tri peptide supplements
 - Use PepT1 transporter for animal specific nutrition
 - Tailor peptide enrichment to specific markets
 - High glutamine & branch chain amino acids. (muscle energy)
 - High methionine & sulfur rich amino acids. (hair, skin and milk)
- Create peptide fingerprint of final product for QC
 - Check starting materials and in-line processes

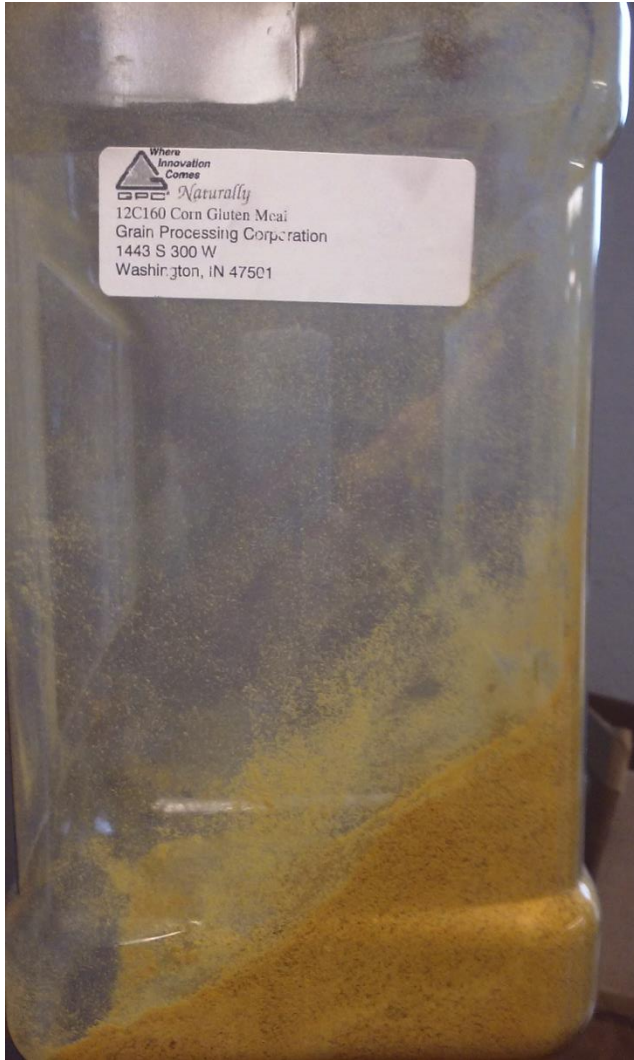
Analytical Service Provided

- ID proteins in by-products by MS1 mass
- Evaluate enzyme choices for desired product
- Fingerprint peptide signature of hydrolysates
- Analyze for AA length and content
- Link peptides to flavor and nutrition
- Track peptide fingerprints after blending

MALDI-TOF mass spectrometry

- Provides exact mass of proteins and peptides.
- Higher accuracy of intact large protein than 1D gels
- Better resolution of similar protein subunits
- Faster data acquisition with ability to automate

Corn Gluten Meal Example



- Corn Gluten Meal from corn syrup manufacturing
- Food grade 60% zein protein waste
- High in glutamine and proline
- Alpha and gamma Zein
 - High in BCAA
 - Leucine, Isoleucine & Valine
- Beta and delta Zein
 - High in sulfur containing AA
 - Methionine

Corn Gluten Meal Example



- Remove excess oil, color and other hydrophobic impurities with 95% Ethanol.
- Can further purify with hexane/ethanol blends depending on quality.
- Ethanol can be distilled and reused

[Biotech Core Instagram](#)

Corn Gluten Meal Example



- Repetitively extract prolamins in 70% ethanol.
- Can tailor ethanol % for desired protein composition.
- Purification and extraction may not be required for some products.

Maize prolamins

γ (27k)

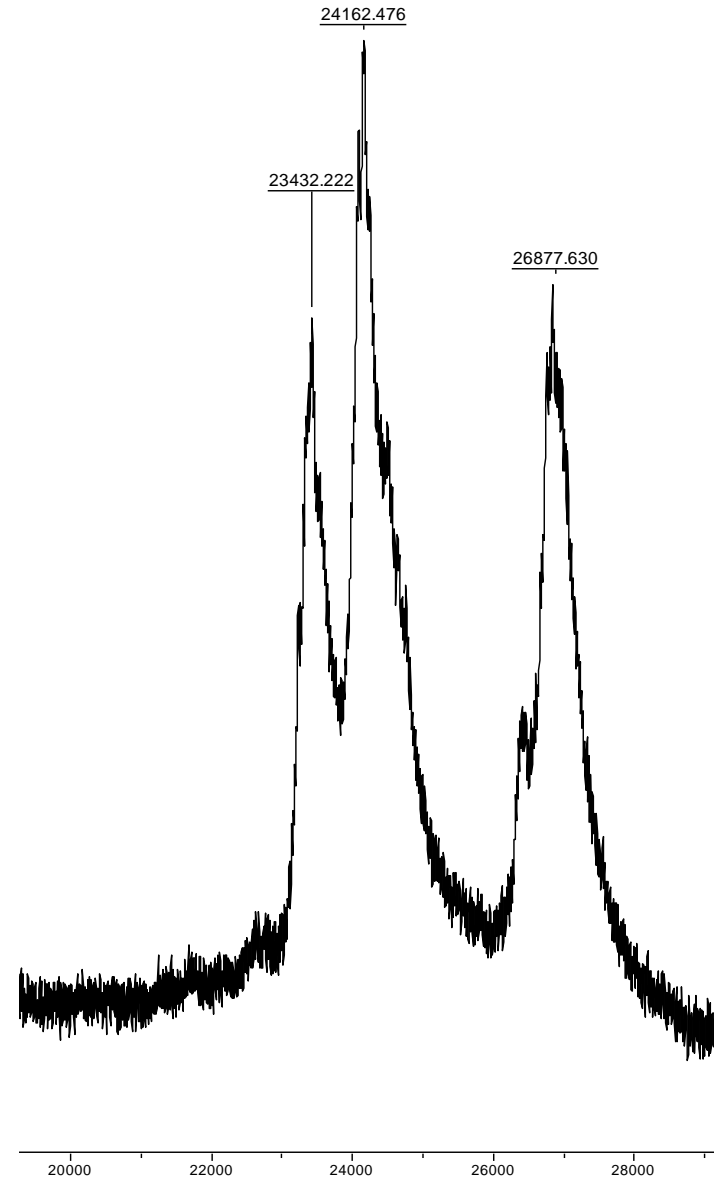
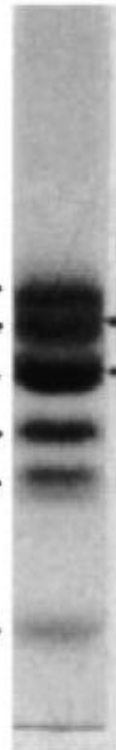
α (22k)

α (19k)

γ (16k)

β (14k)

δ (10k)



Corn Gluten Meal Example

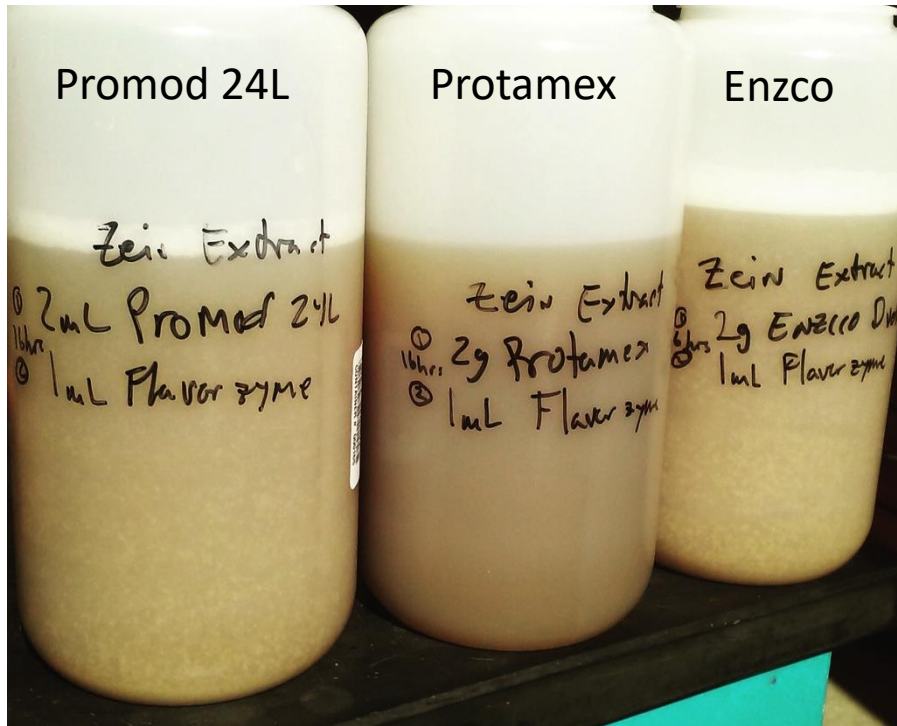


- Soluble Zein proteins aggregate and precipitate in water.
- Add to pH adjusted water with 0.1-1% enzyme protease.
- Keep final ethanol concentration at ~20%
 - Engineered food enzymes are still active.
 - Prevents microbial growth.
 - Helps solubilize peptides to improve accessibility.

Food Grade Enzyme Examples

- Promod™ 24L - Biocatalysts
 - Bacterial broad spectrum protease
- ENZECO® Dual - Enzyme Development Corp.
 - Bromelain and Papain from meat tenderizing
- Protamex® - Novozyme
 - Bacillus sp. bacterial protease from fish processing
- Flavourzyme – Novozyme
 - Protease and peptidase preparation from *Aspergillus oryzae* culture.
 - Complex mix of at least 8 enzymes

CGM Double Digest Example



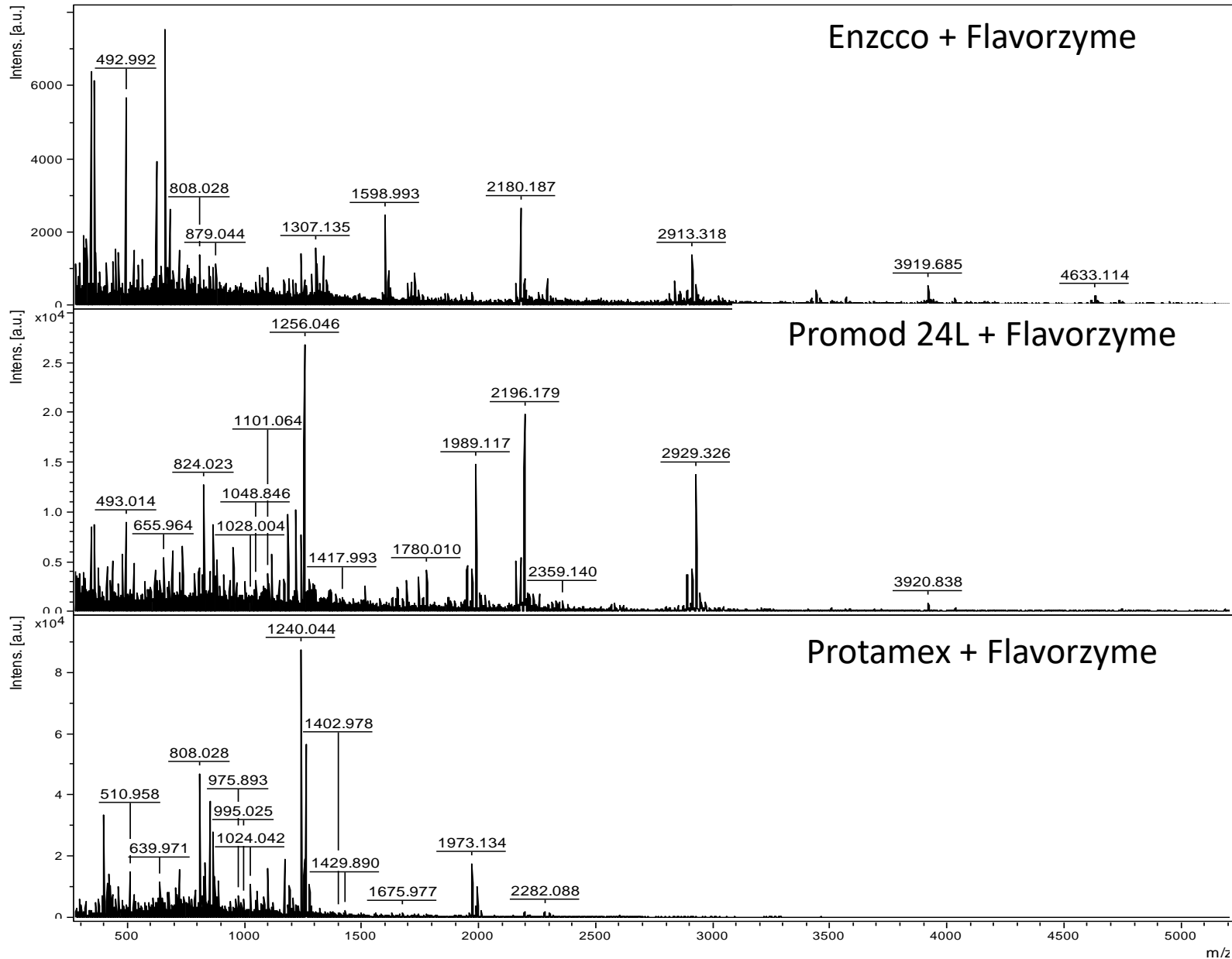
- Digest Zein with first protease until aggregate is broken.
- Digest with secondary enzyme, Flavourzyme, until entire solution is liquefied.
- Monitor hydrolysates process by MALDI-TOF fingerprinting.

Analytical Procedure

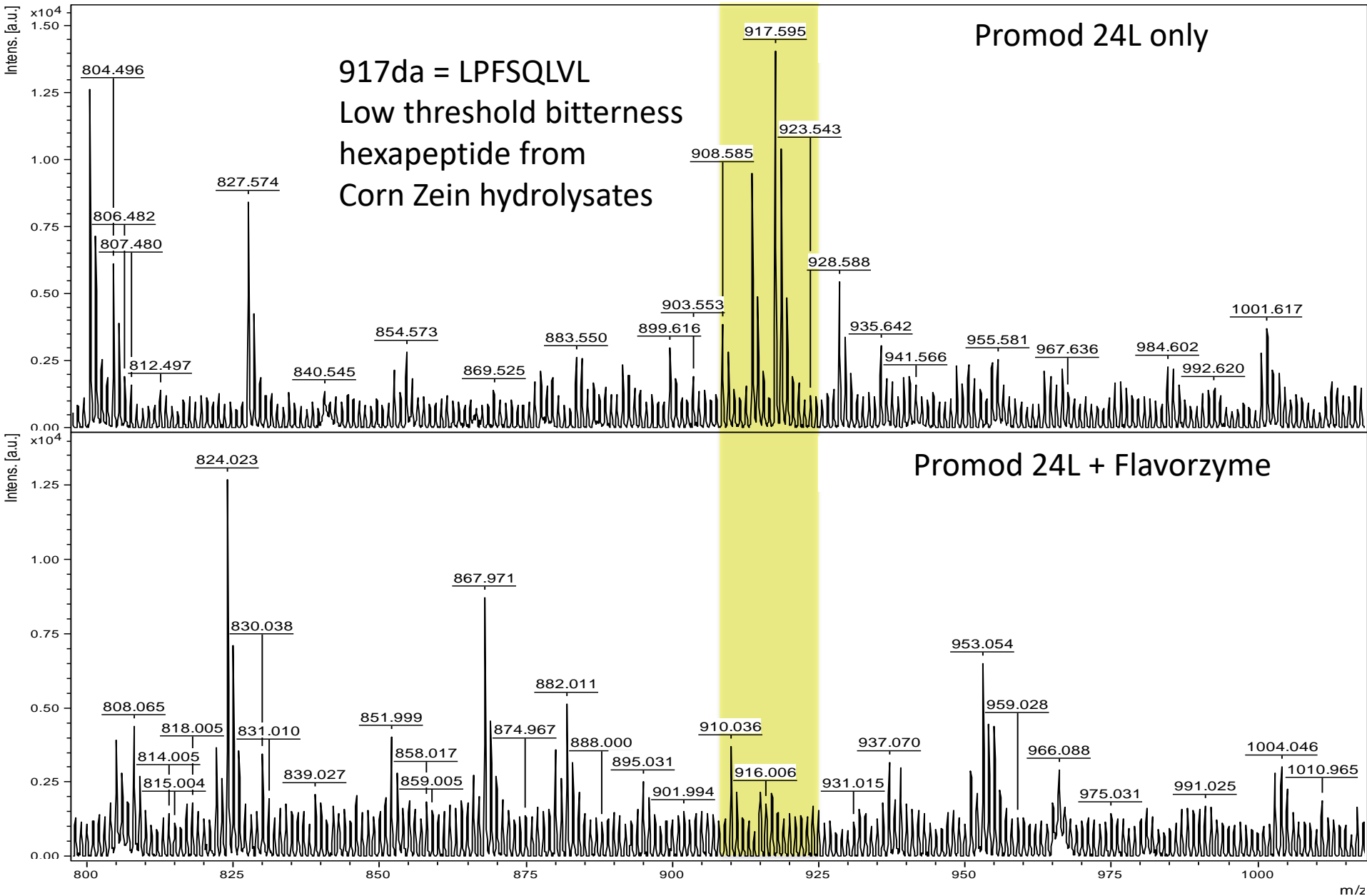
- Dilute 10uL of hydrolysate into:
 - 100uL of Acetonitrile and 90uL 1% TFA in water.
- Spot 1uL extract to MALDI-TOF
 - Use DHB matrix for peptide fingerprinting.



CGM Double Digests

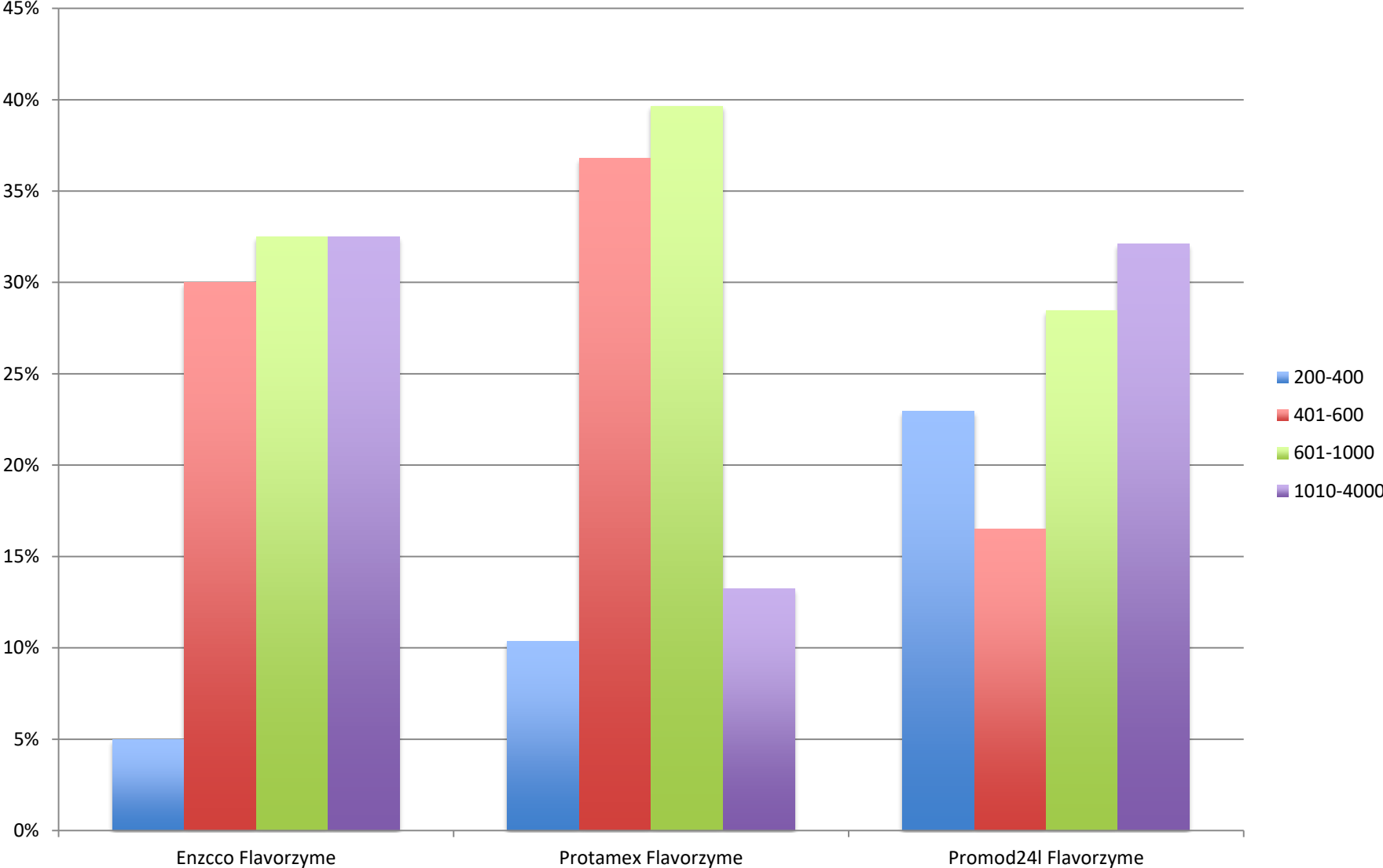


Why double digests?

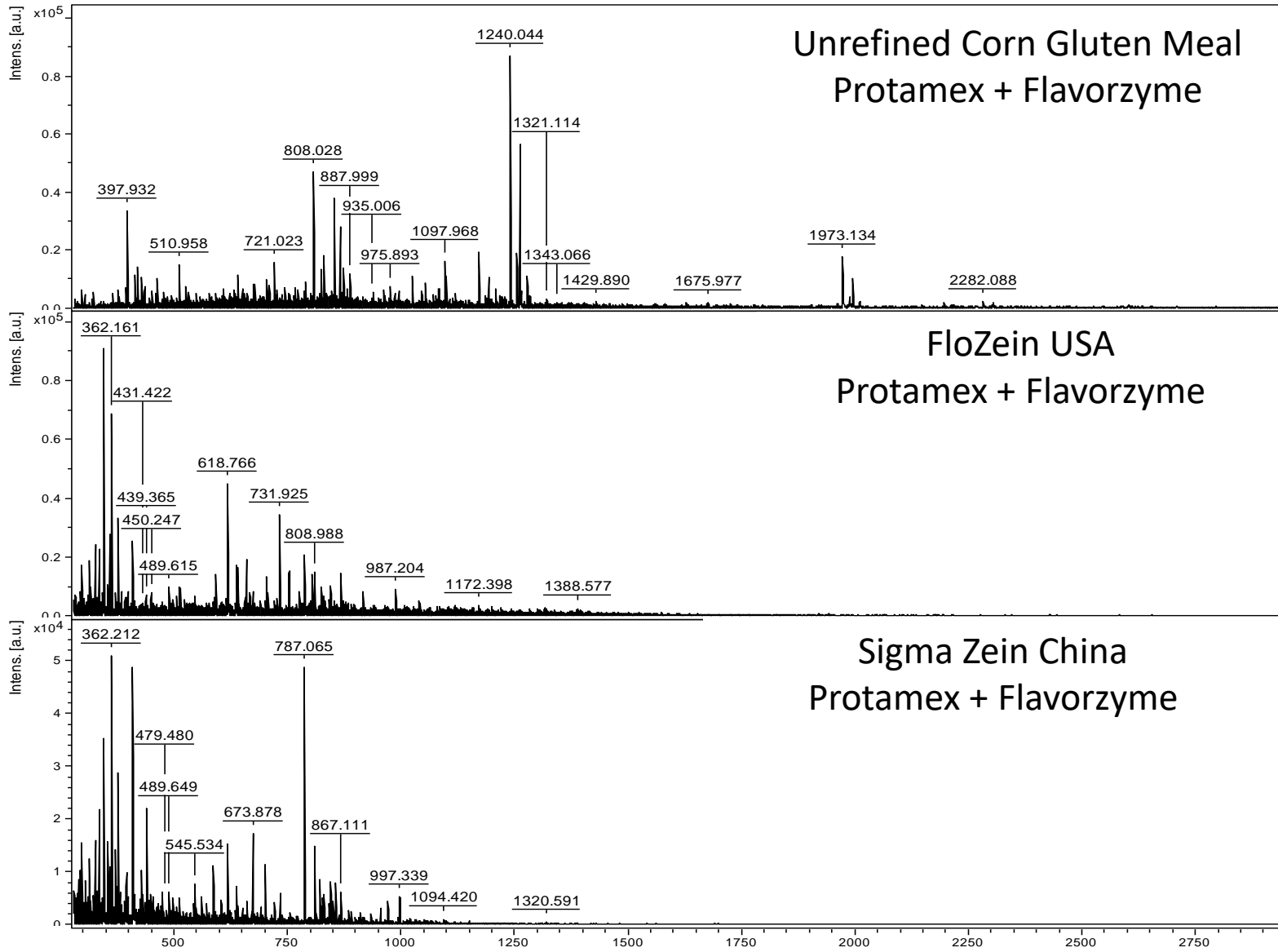


Measuring improved digestibility

Double digests of CGM Zein



Comparing different Zein sources



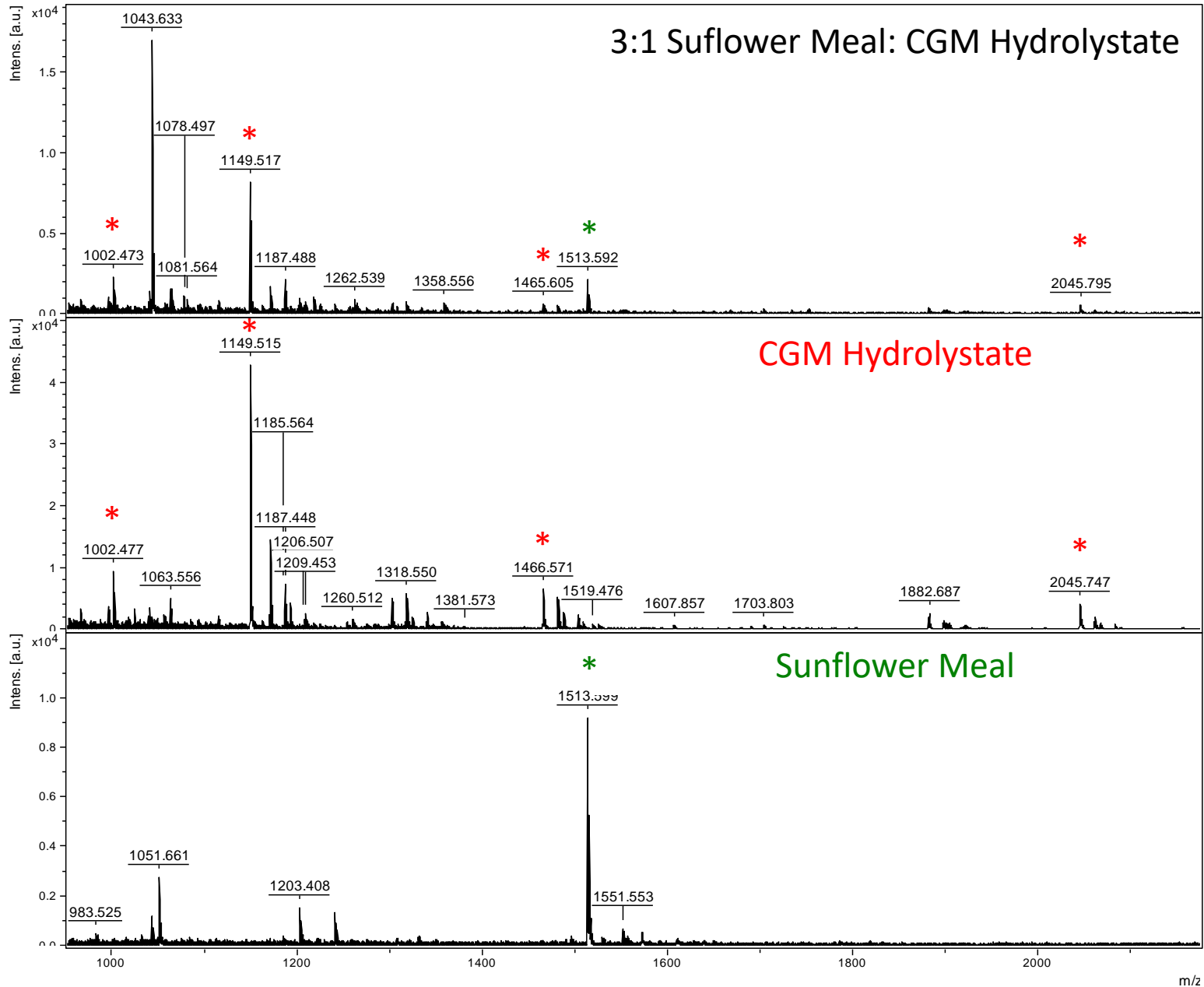
Unique bulk filler sources

- Canola meal
 - \$191/ton FOB from ADM
 - 36% protein
 - Accepted as a grain-free nutrient source.
- Sunflower meal
 - \$280.00/ton FOB from CPE Feeds Inc – Texas
 - 30% protein
 - High in methionine, adds great flavor.

Blending with Filler

- New naming of product
 - Hydrolysate enrich by-products
 - “Protein enriched”
- Example product:
 - 3 parts Sunflower Meal : 1 part CGM hydrolysate
- Mix and heat extrude to kill enzymes
- Rename as “Enriched sunflower meal”
- GRAS due to all source ingredients being GRAS

Tracking peptides thru blending



CGM Enriched Products

- Turns high protein by-products into digestible nutrition without changing GRAS status.
- Masks the source of protein hydrolysates by blending into alternative sources.
- Analytical data used to normalize by-product streams.
- QC data can be used by plant manager to assign CAS#'s for new enrich products.

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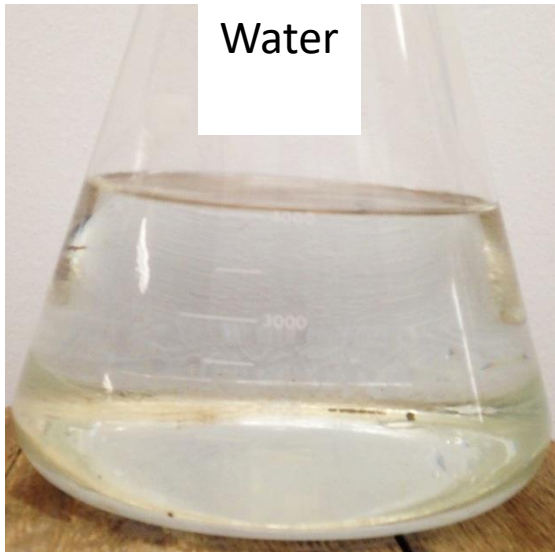
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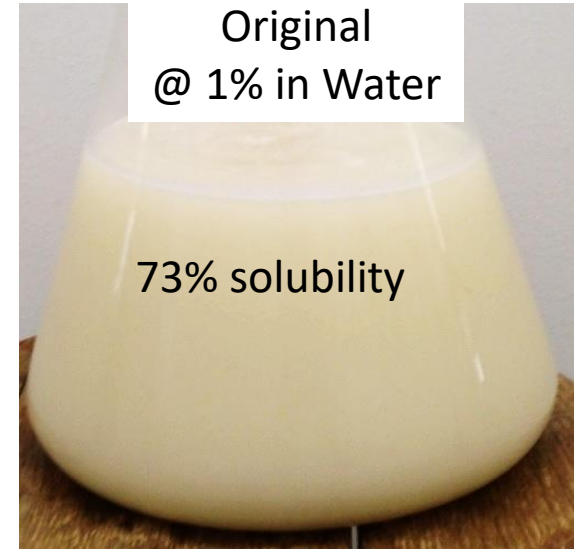
Example 2: Gluten hydrolysate project



New Product

?

A white rectangular box with a black border. Inside the box, the words "New Product" are written in a large, bold, black font at the top. In the center of the box, a large black question mark is displayed.

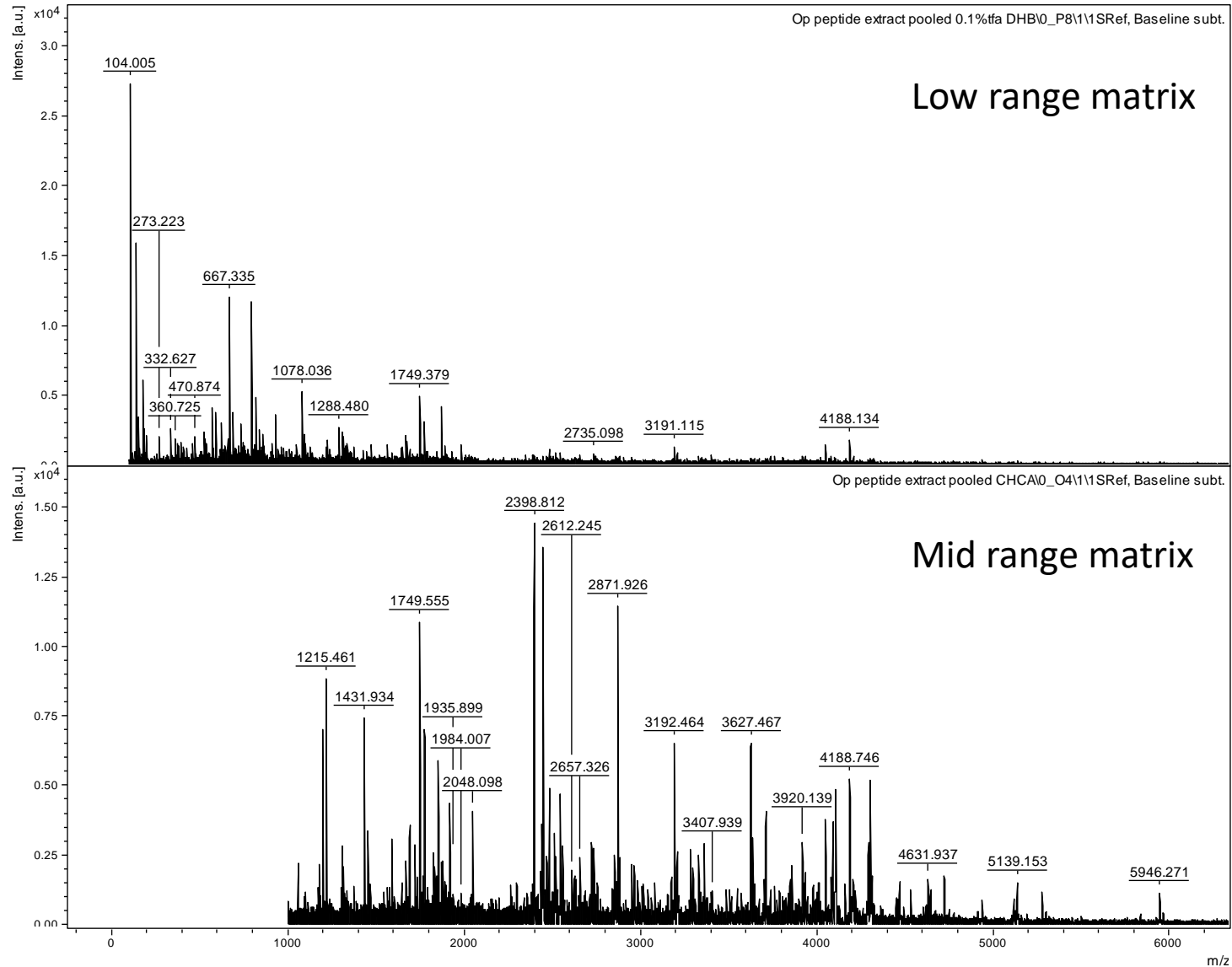


- Goal: Increase solubility in water
- Shorten peptide length for absorption
- Cleave known peptide epitopes and reduce bitterness

Full Analytical Extraction Protocol

- 50mg of spray dried hydrolysate was defatted in CHCl_3 /MeOH/Water and freeze dried.
- Freeze dried samples were split into 4 fractions and resolubilized.
 - Fraction 1: w/ 0.1%TFA
 - Fraction 2: w/ 25% Acetonitrile : 75% 0.1% TFA
 - Fraction 3: w/ 50% Acetonitrile : 50% 0.1%TFA
 - Fraction 4: w/ 75% Acetonitrile : 25% 0.1%TFA
- Fractions were centrifuged and supernatant was collected and pooled.

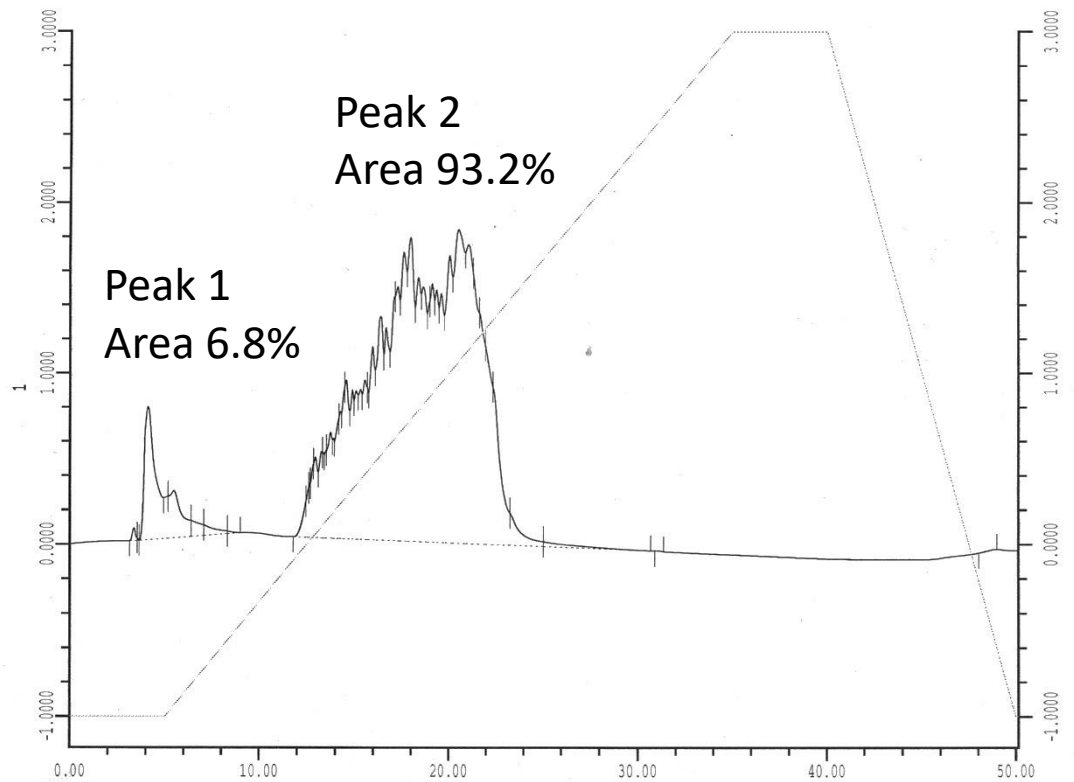
Raw MALDI-MS data of Pooled Sample



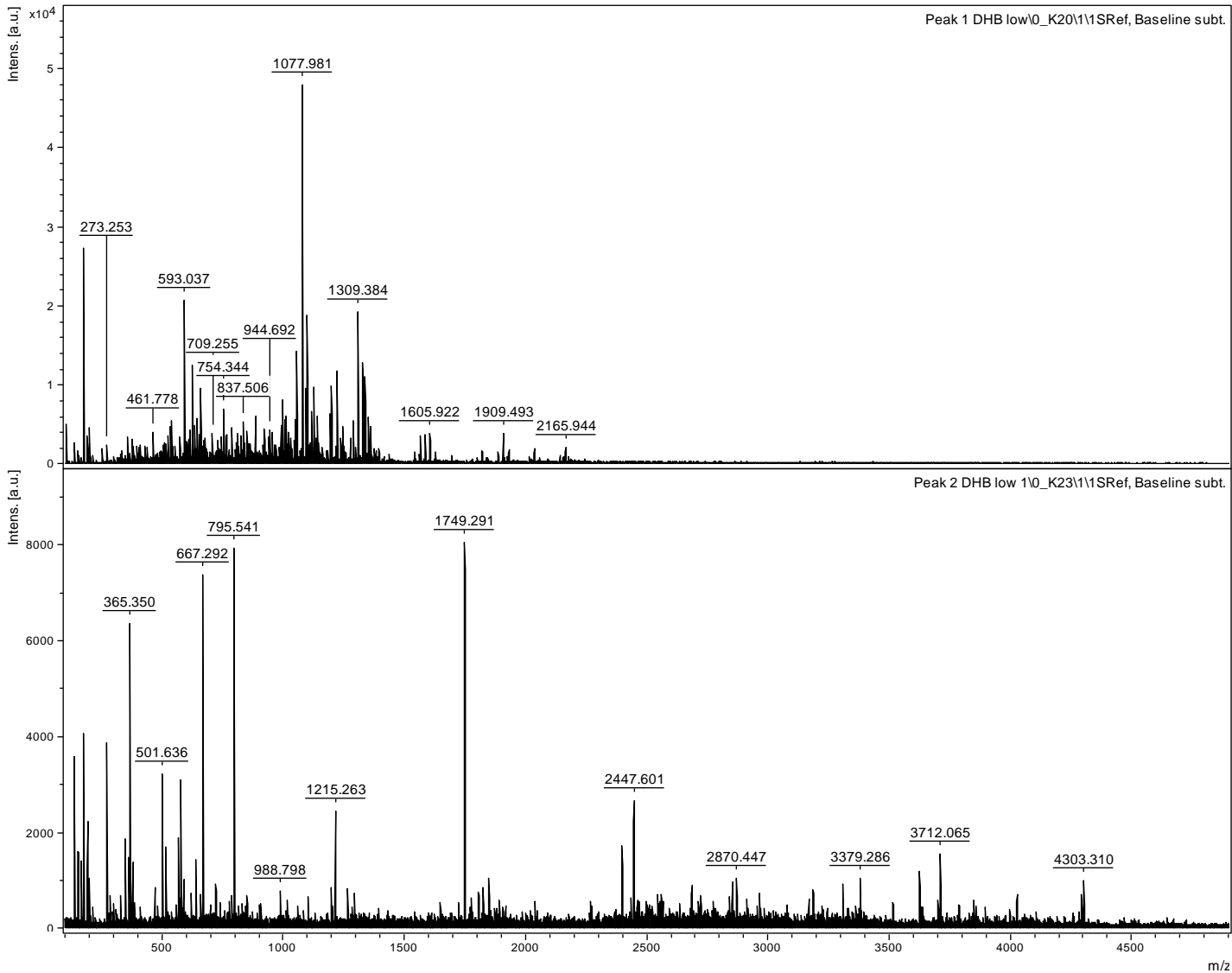
Peptide fractionation and distribution

- Pooled extracts were freeze dried and solubilized in 0.1% TFA for HPLC and Mass Spec (MS) analysis.
- HPLC was run on a 250 mm x 4.6 mm C4 column with Acetonitrile as the mobile phase.
- Two distinct broad peaks were collected and analyzed by MALDI-TOF MS.

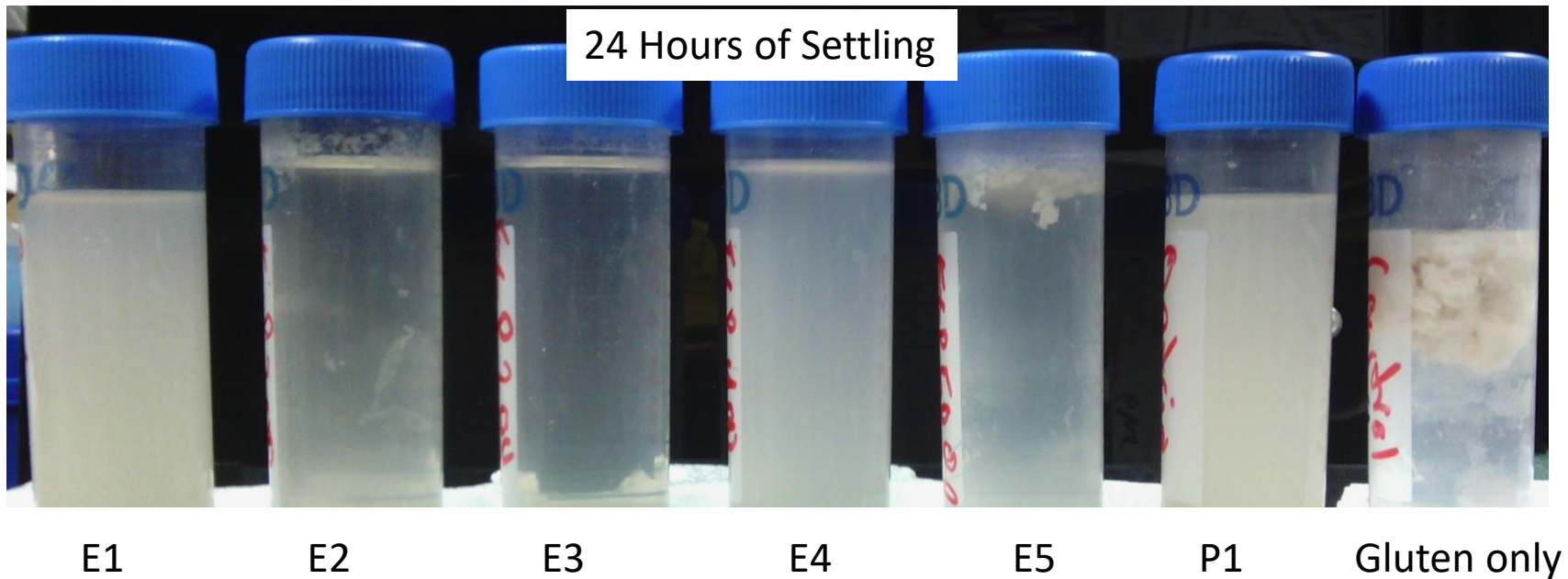
C4 HPLC of pooled extract



HPLC MALDI-MS data of Peak 1 & 2

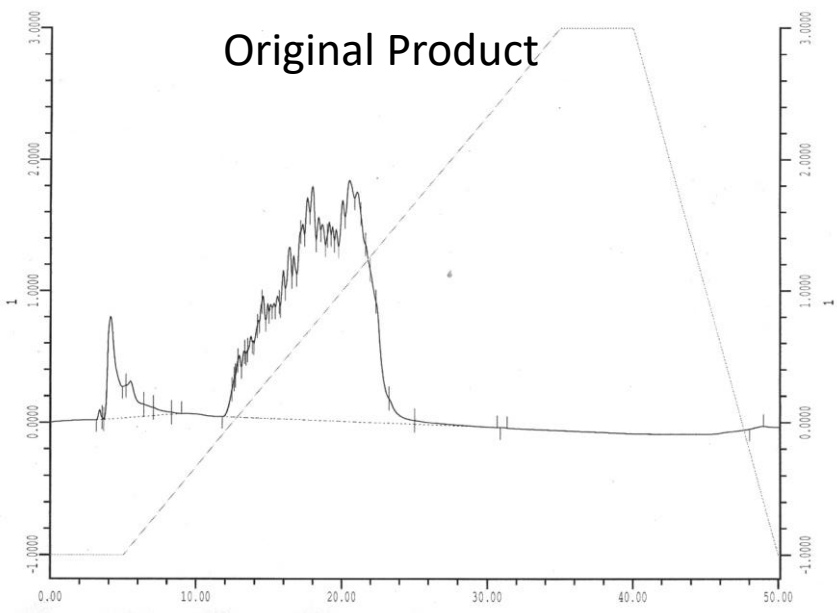
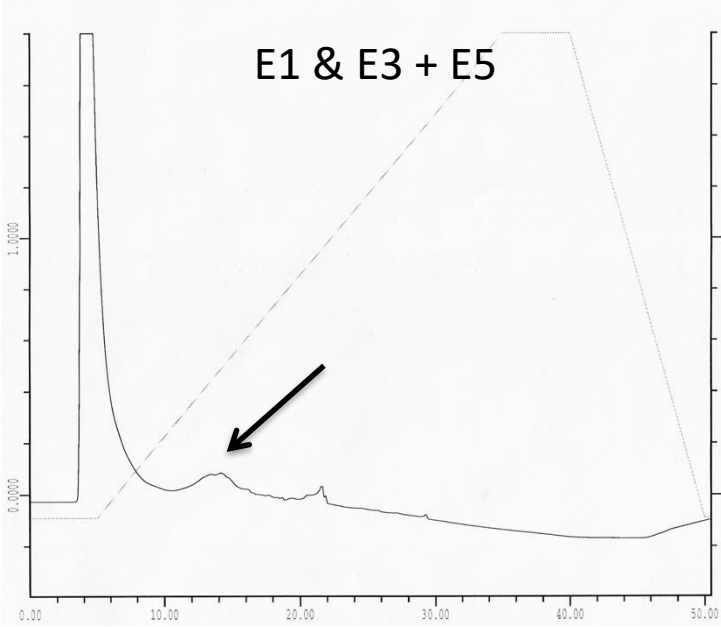
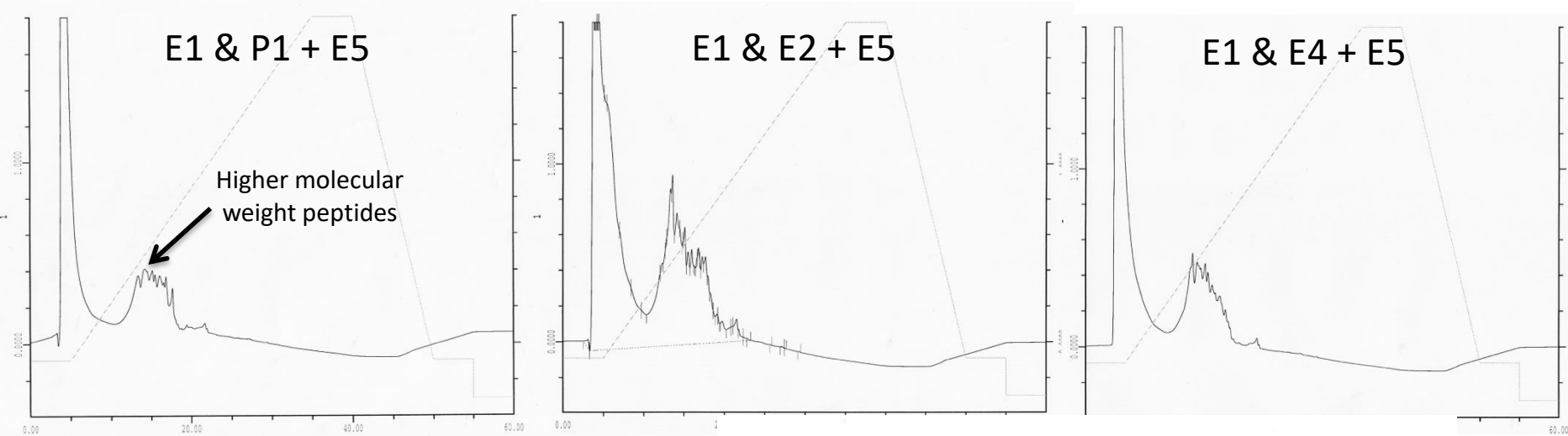


Single enzyme Solubilization and Dispersion of Gluten

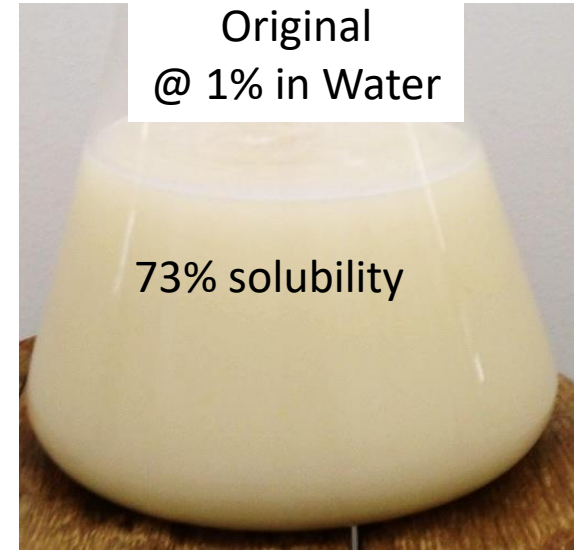
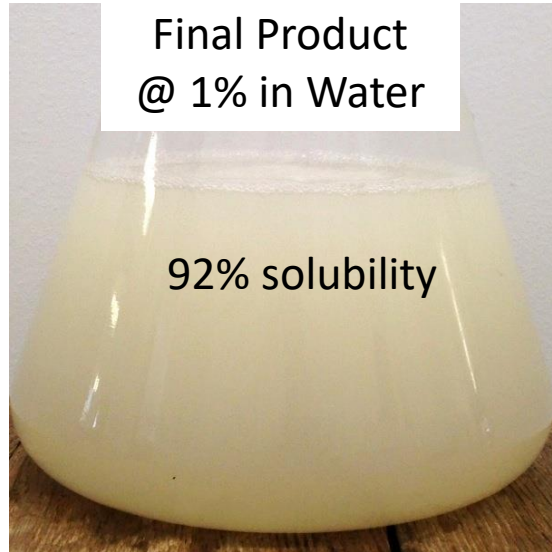
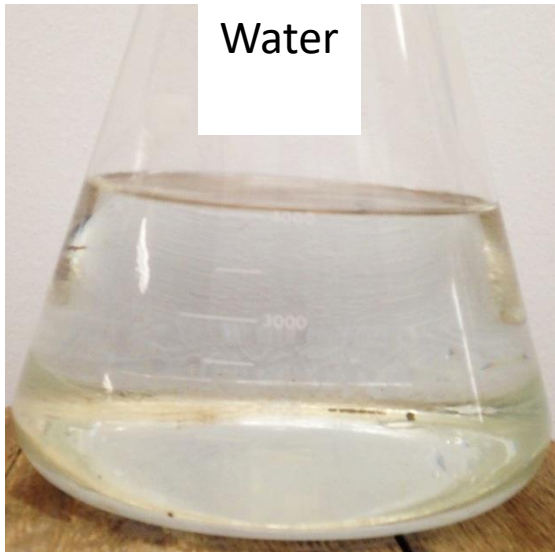


- All enzymes are at 1% w/w of gluten.
- E1, E4 and P1 were best at dispersing the gluten peptides after 24hrs of settling.
- E2 & E3 produced peptide rich soluble fractions but were unable to disperse large amount of gluten peptides.
- An enzyme combination strategy is necessary to both disperse and solubilize gluten peptides.

HPLC comparison of Enzyme Combos



Example 2: Gluten hydrolysate project



- Decreased insoluble precipitates by >300%
 - 27% w/v to 8% w/v
- Shorten peptide length for absorption
- Cleaved known peptide epitopes
- Reduced Bitterness

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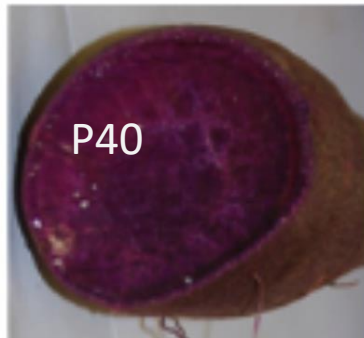
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Other projects and services

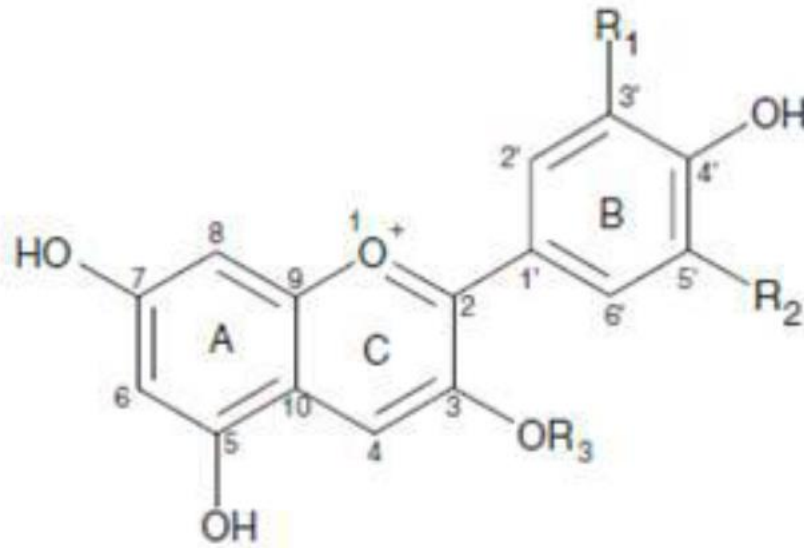
- Purple tomato and sweet potato anthocyanins
 - Discovering new small molecule with ESI-MS/MS
- PRRS virus using nanotubes for intercellular spread
 - Protein ID and Mapping

Anthocyanin Rich Sweet Potatoes and Tomatoes



Good dietary antioxidants for cancer prevention specifically colorectal cancer.

Structures of common anthocyanidins and anthocyanins



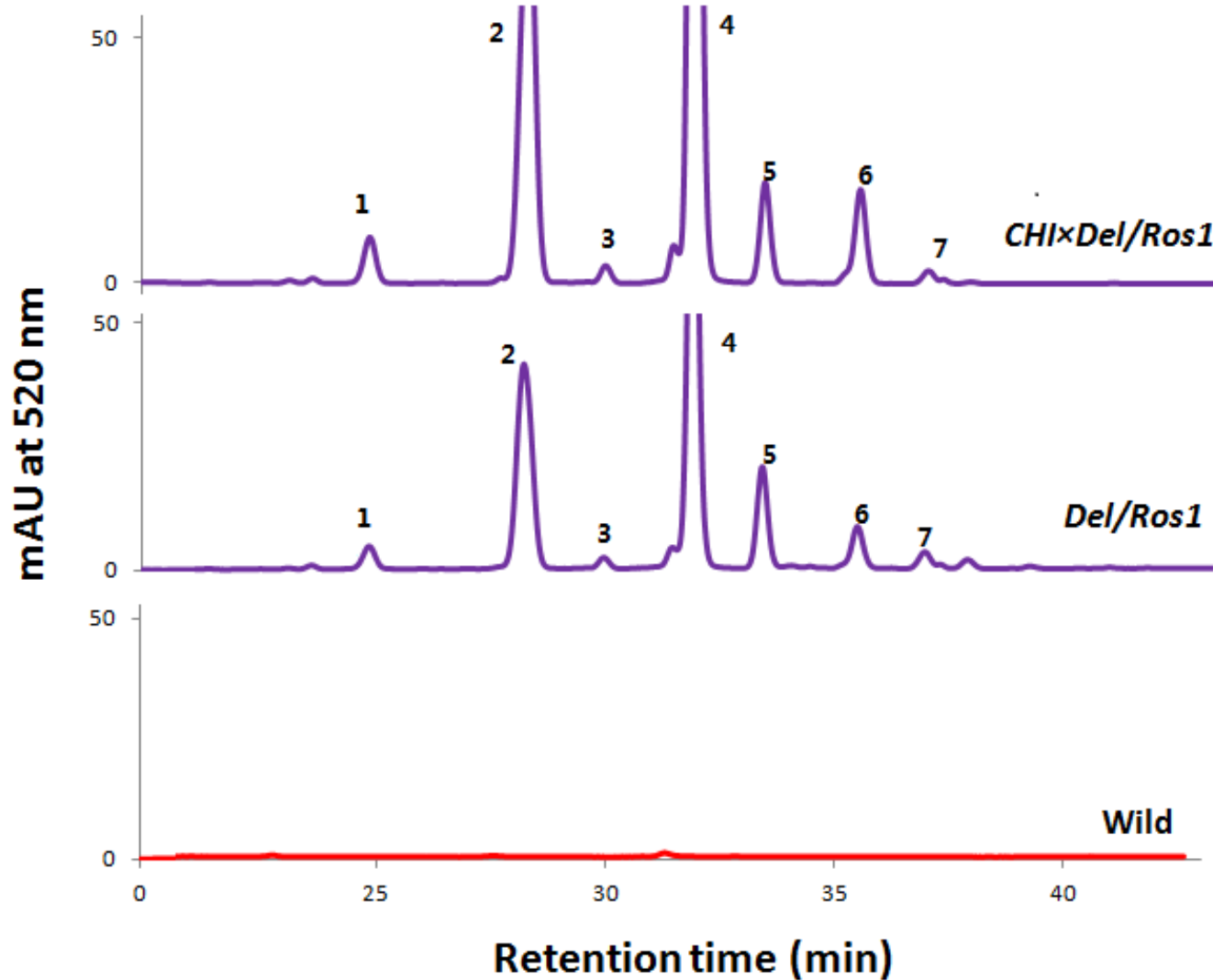
	<u>R₁</u>	<u>R₂</u>
Cyanidin	OH	H
Pelargonidin	H	H
Delphinidin	OH	OH
Petunidin	OCH ₃	OH
Peonidin	OCH ₃	H
Malvidin	OCH ₃	OCH ₃

R₃ = Glucose, galactose, rhamnose, xylose, or arabinose

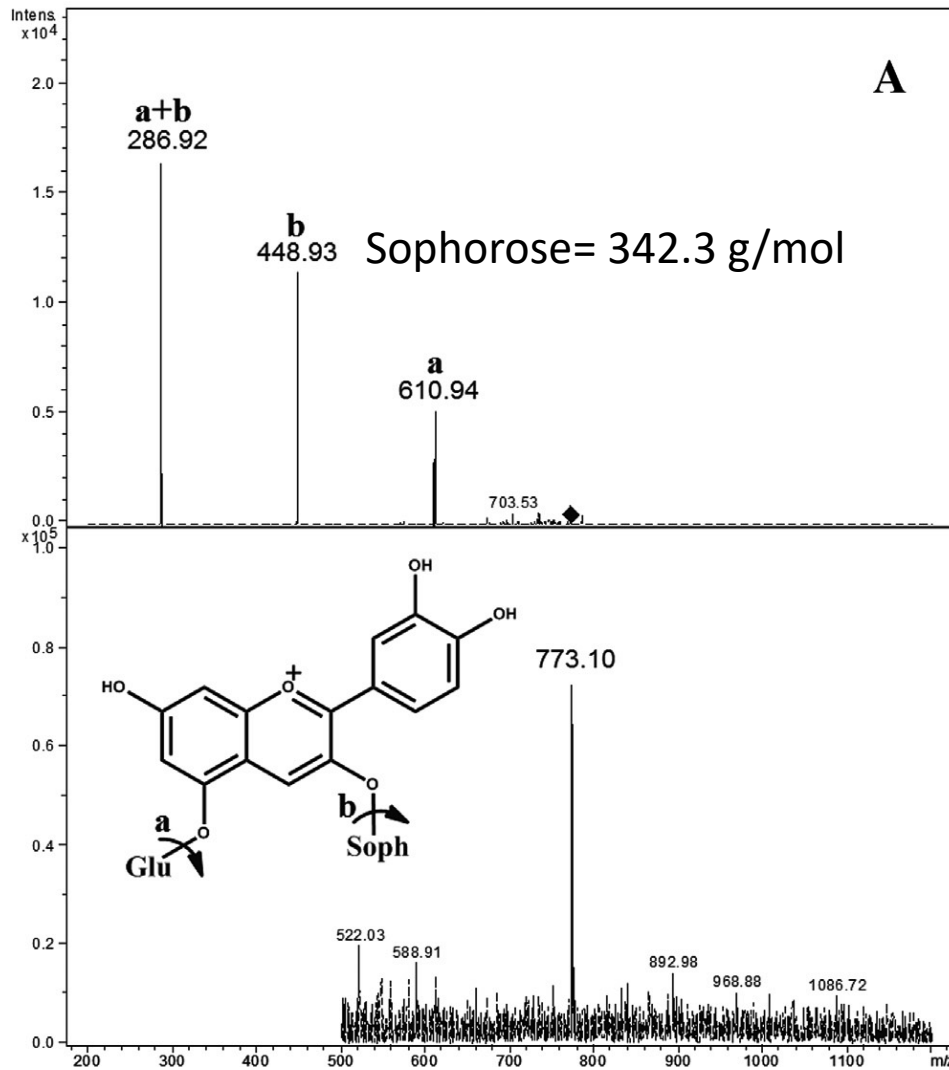
Cyanidin = 287.2 g/mol

Malvidin = 331.3 g/mol

C18 Reverse phase HPLC separation of a methanolic extract

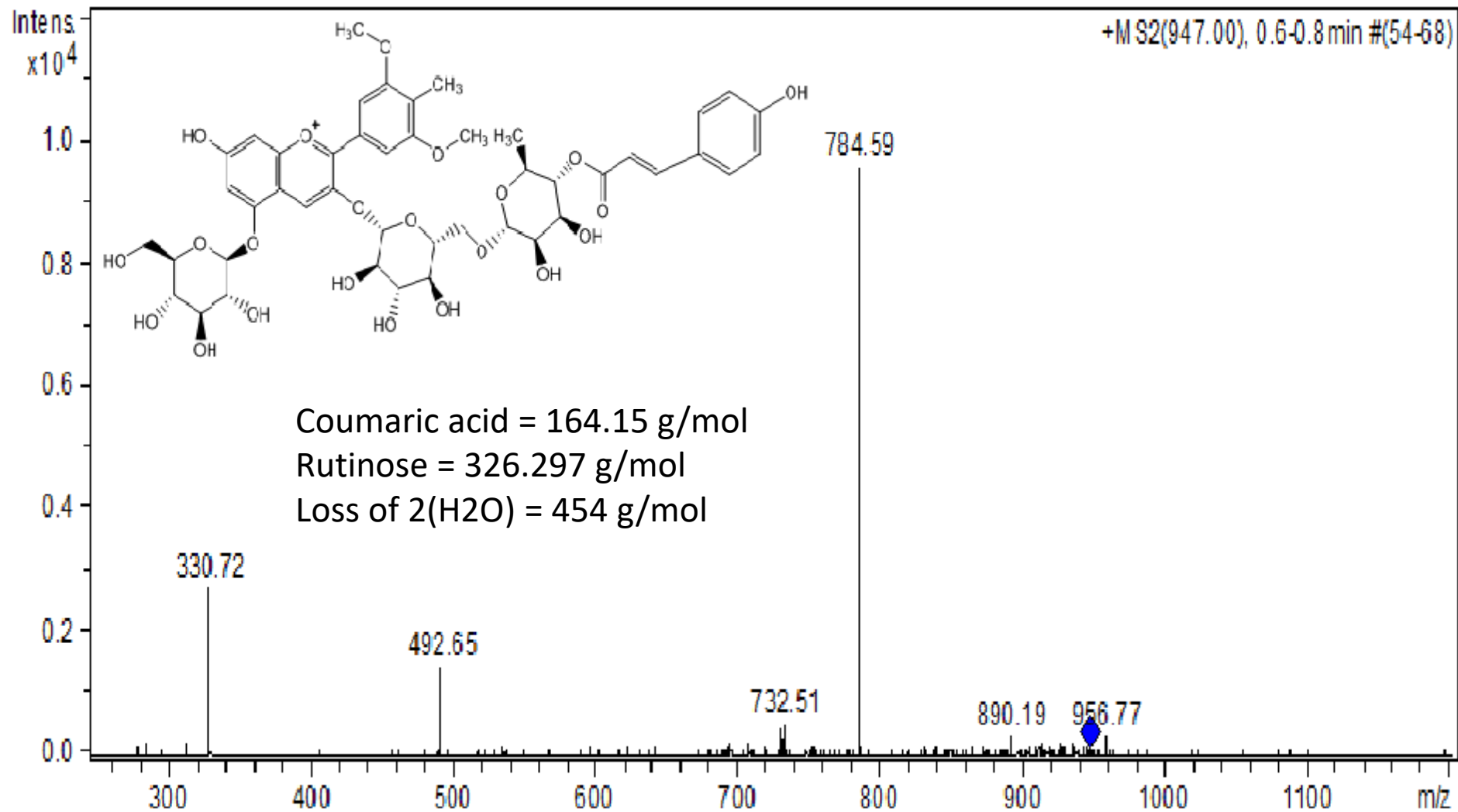


Dissecting the ESI-MS/MS daughter ions



Peak 2 : Cyanidin 3-sophoroside-5-glucoside

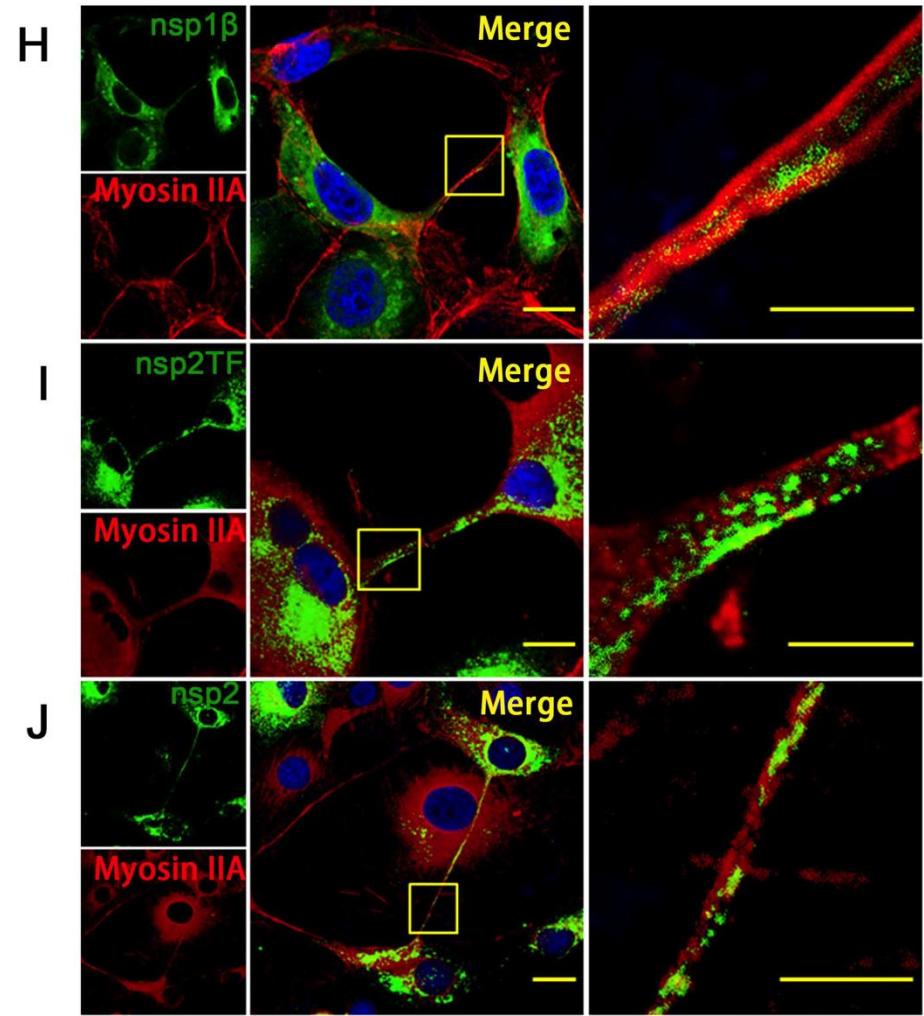
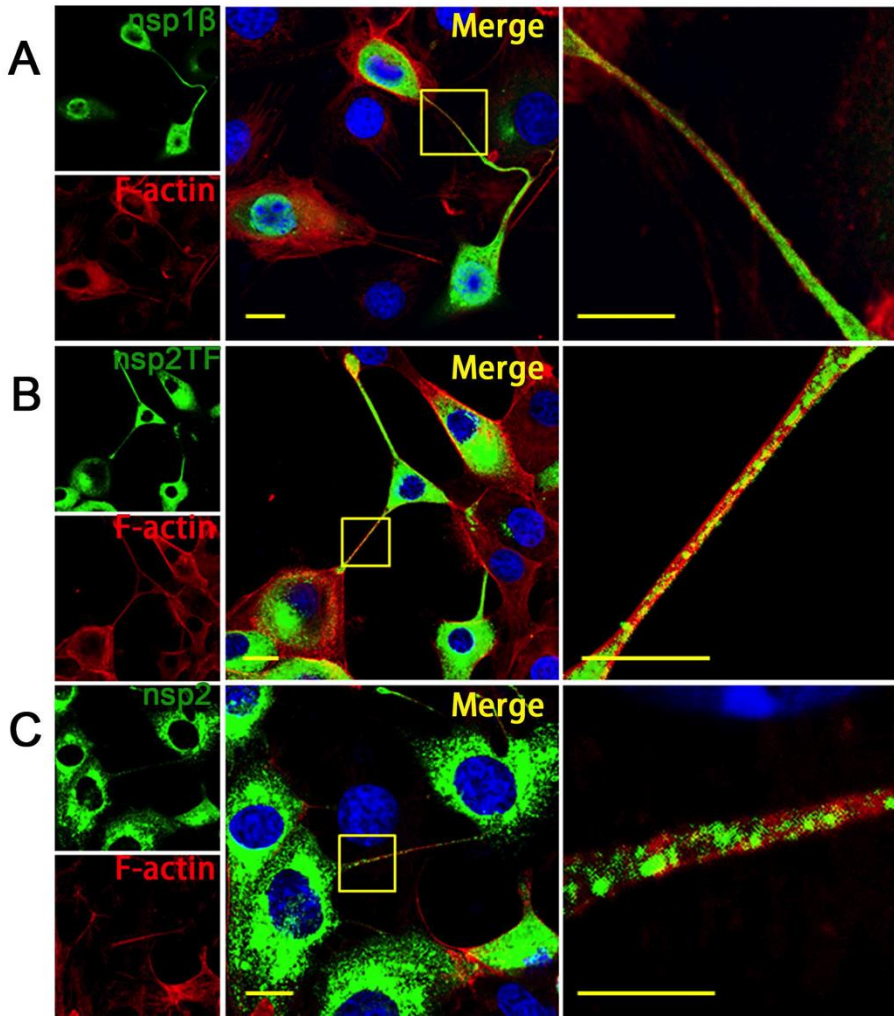
Discovering unknowns by fragmentation pattern



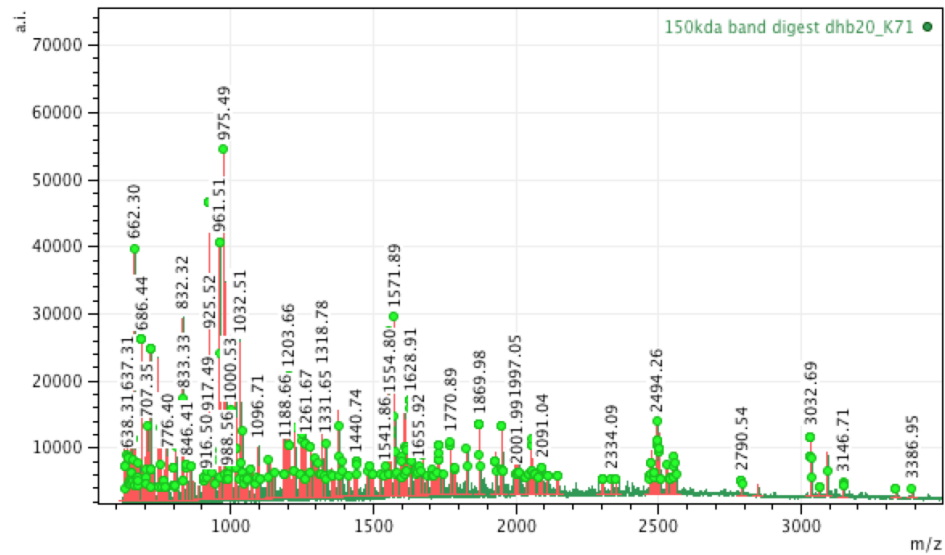
Peak 6: malvidin-3-(p-coumaroyl)-rutinoside-5-glucoside

Porcine reproductive and respiratory syndrome virus utilizes nanotubes for intercellular spread.

Rui Guo, Benjamin B. Katz, et al. In Press Journal of Virology. 2016



Myosin II Identification using In-gel Trypsin Digestion with MALDI-TOF



Sequence - >gi|635102017|ref|XP_007973837.1| PREDICTED: myosin-9 [Chlorocebus sabaues]

Accession	Length	Mo. Mass	Av. Mass	Coverage	Matched Int.							
	1960	228713.77	228853.49	88.2 %	59.3 %							
MAQQAADKYL	YVDKNFINNP	LAQADWAAKK	LVWVPSDKSG	FEPASLKEEV	GEEAIVELVE	NGKKVKVKNK	DIQKMNPPKF	SKVEDMAELT	CLNEASVLHN	LKERYSGLI	YTYSGLFCVV	INPYKNLPIY
SEEIIVEMYK	KKRHEMPPHI	YAITDTAYRS	MMQDREDQSI	LCTGSEGAGK	TENTKKVIQY	LAVASSHKS	KKDQGELERQ	LLQANPILEA	FGNAKTVKND	NSSRFGKIR	INFDVNGYIV	GANIETYLLE
KSRAIRQAKE	ERTFHIFYYL	LSGAGEHLKT	DLLLEPYNKY	RFLSNGHVTI	PGQQDKMFQ	ETMEAMRMIG	IPEEEQMGLL	RVISGVLQLG	NIVFKKERNT	DQASMPDNTA	AQKVSHLLGI	NVTDFTRGIL
TPRIKVGDRD	VQKAQTKQEA	DFAIEALAKA	TYERMFRLV	LRINKALDKT	KRQGASFIGI	LDIAGFEIFD	LNSFQQLCIN	YTNKQLQQLF	NHTMFILEQE	EYQREGIEN	FIDFGLDLQP	CIDLIEKPG
PPGILALLDE	ECWFPKATDK	SFVEKVMQEQ	GTHPKFQPKP	QLKDKADFCI	IHYAGKVVDYK	ADEWLMKNMD	PLNDNIATLL	HQSSDKFVSE	LWKDVDRIIG	LDQVAGMSET	ALPGAFKTRK	GMFRVTGQLY
KEQLAKLMAT	LRNTNPNFVR	CIIPNHEKKA	GKLDPHLVLD	QLRRCNGVLEG	IRICRQGFNP	RVVQEFQRQ	YEILTPNSIP	KGFMGKQAC	VLMIKALELD	SNLYRIGQSK	VFFRAGVLAH	LEEERDLKIT
DVIIGFQACC	RGYLARKAFA	KRQQQLTAMK	VLQRNCAAYL	KLRNWQWRL	FTKVKPLLQV	SRQEEEMAK	EEELVKVREK	QLAAENRLTE	METLQSQLMA	EKLQLEQLQ	AETELCAEAE	ELRARLTAKK
QELEEI CHDL	EARVEEEER	CQHLQTEKKK	MQONIQELEE	QLEEEESARQ	KLQLEKVTTT	AKLKKLEEEQ	IILEDNCKL	AKEKKLLEDR	IAEFTTNLTE	EEEEKSKLAK	LKNKHEAMIT	DLEERLRREE
KQRQLEKTR	RKLEGDSTD	SDQIAELQAO	IAELKMQLAK	KEEELQAALA	RVEEEAAQKN	MALKKIRELE	SQISELQEDL	ESERASRNKA	EKQKRDGEE	LEALKTELED	TLDTTAAQQE	LRSKREQEVN
ILKKTLEEEA	KTHEAQIQEM	RQKHSQAVEE	LAEQLEQTKR	VKANLEKAKQ	TLENERGELA	NEVKVLLQGK	GDSEHKRKKV	EAQLQELQVK	FNEGERVTE	LADKVTKLQV	ELDNVTGLLS	QSDSKSKL
KDFSALSQL	QDTQELQEE	NROKLSLSTK	LKQVEDEKNS	FREQLEEEEE	AKHNLEKQIA	TLHAQVADMK	KKMEDSVGCL	ETAEEVKRKL	QKDLEGLSOR	HEEKVAAYDK	LEKTKTRLQO	ELDDLLVDLD
HQRQSA CNLE	KKQKFPDQLL	AEEKTISAKY	AEERDRAEAS	AREKETKALS	LARALEEAME	QKAELERLTK	QFRTEMEDLM	SSKDDVGSV	HELEKSKRAL	EQQVEEMRTQ	LEEELELQ	TEDAKLRLEV
NLQAMKAQFE	RDLQKDEQS	EKKKQLVRQ	VREMEAELED	ERKQSRMAVA	ARKKLEM DLK	DLEAHIDSAN	KNREAIKQL	RKLQAQMKDC	MRELD DTRTS	REEILAQAKE	NEKKLKS MEA	EMIQLQEELA
AAERAKRQAO	QERDELADEI	ANSSGKALA	LEEKRRLEAR	IAQLEEELEE	EQNGTELIND	RLKKANLQID	QINTDLNLER	SHAQKNENAR	QQLERQNKEL	KVKLQEMEGT	VKSKYKASIT	ALBAKTAQLE
EQLDNETKER	QAACKQVVRT	EKKLKDVLLO	VDDERRNAEQ	YKQDQAKAST	RLKQLKROLE	EAEEEAQRAN	ASRRKLQREL	EDATETADAM	NREVSSLKKN	LRRGDLPLFVM	PRRMARKGAG	DGSDEVDGK
ADGAEAKPAE												

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