## Supplementary materials

Table S1: Protein content of potato protein hydrolysate fractions and amount of protein in emulsions.

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| --- | --- | --- | --- |
| Sample | Protein content (%) | Protein in emulsion (mg) | Protein in emulsion (wt%) |
| AFPPH | 71.8 ± 0.5 | 78.9 | 0.04 |
| AF35 | 75.3 ± 1.2 | 82.9 | 0.04 |
| AF13 | 73.4 ± 0.1 | 80.7 | 0.04 |
| AF1 | 62.9 ± 0.2 | 69.1 | 0.03 |
| TFPPH | 68.2 ± 0.0 | 75.0 | 0.03 |
| TF35 | 73.0 ± 0.3 | 80.3 | 0.04 |
| TF13 | 66.8 ± 0.6 | 73.4 | 0.03 |

Table S2: Peptidomics data from PPHs and their fractions. Table includes all peptide-level information from the MaxQuant output data (peptides.txt) in addition to sample-wise relative signal intensity (I\_rel) and predicted antioxidant scores from AnOxPePred (FRS and CHE). Table also includes peptide-wise calculations of molar AA distribution as well as intensity-weighted mean length and charge. Table is appended as separate file (Table S2.xlsx).

Table S3: Zeta potential (mV) and droplet size D[3,2] and D[4,3] in emulsions. Values are given as mean ± standard deviation. Letters (a-e) in the same column indicate a statistically significant difference between samples. \* Show a significant difference in droplet size between day 1 and day 9

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
| Emulsion | D[3,2]  (µm) | | D[4,3]  (µm) | | Zeta potential (mV) |
|  | Day 1 | Day 9 | Day 1 | Day 9 | Day 1 |
| Control | 0.120±0.001bc | 0.120±0.000b | 0.185±0.001cd | 0.184±0.001bc | (-)18.1±3.8a |
| EDTA | 0.121±0.001de | 0.122±0.000d,\* | 0.190±0.001e | 0.192±0.000de | (-)19.6±2.8a |
| AFPPH | 0.118±0.000a | 0.119±0.000a,\* | 0.180±0.000a | 0.181±0.002a | (-)15.4±3.2a |
| AF35 | 0.119±0.000b | 0.120±0.001ab | 0.182±0.000ab | 0.182±0.001ab | (-)15.1±1.8a |
| AF13 | 0.121±0.000e | 0.123±0.001d | 0.191±0.001e | 0.194±0.001e | (-)14.1±1.0a |
| AF1 | 0.120±0.000cd | 0.120±0.000bc,\* | 0.187±0.001d | 0.187±0.000c | (-)14.3±1.6a |
| TFPPH | 0.120±0.000b | 0.122±0.001cd | 0.189±0.000e | 0.190±0.001d | (-)14.1±0.8a |
| TF35 | 0.120±0.001bc | 0.120±0.001ab | 0.184±0.000bc | 0.183±0.000ab | (-)13.3±1.3a |
| TF13 | 0.119±0.001d | 0.120±0.000b,\* | 0.183±0.001bc | 0.184±0.001abc | (-)12.4±1.0a |

Table S4: alpha-tocopherol consumption in % during 9 days of storage

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| --- | --- | --- | --- |
|  | Day 3 | Day 6 | Day 9 |
| Control | 96.2 | 100 | 100 |
| EDTA | 19.5 | 23.4 | 21.7 |
| AFPPH | 22 | 78 | 77.7 |
| AF35 | 26.2 | 50.8 | 71 |
| AF13 | 21.9 | 50 | 57.8 |
| Af1 | 14.5 | 40.3 | 54.3 |
| TFPPH | 35.5 | 74.2 | 81.5 |
| TF35 | 32.3 | 54.1 | 78.9 |
| TF13 | 26.9 | 41.2 | 63.7 |

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**Table S5**

**Control**

**EDTA**

**AFPPH**

**AF35**

**AF13**

**AF1**

**TFPPH**

**TF35**

**TF13**

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**Table S6**

**Control**

**EDTA**

**AFPPH**

**AF35**

**AF13**

**AF1**

**TFPPH**

**TF35**

**TF13**

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**Table S7**

**Control**

**EDTA**

**AFPPH**

**AF35**

**AF13**

**AF1**

**TFPPH**

**TF35**

**TF13**

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**Table S8**

**Control**

**EDTA**

**AFPPH**

**AF35**

**AF13**

**AF1**

**TFPPH**

**TF35**

**TF13**

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**Table S9**

**Control**

**EDTA**

**AFPPH**

**AF35**

**AF13**

**AF1**

**TFPPH**

**TF35**

**TF13**

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**Table S10**

**Control**

**EDTA**

**AFPPH**

**AF35**

**AF13**

**AF1**

**TFPPH**

**TF35**

**TF13**

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**Table S11**

**Control**

**EDTA**

**AFPPH**

**AF35**

**AF13**

**AF1**

**TFPPH**

**TF35**

**TF13**

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**Table S12**

**Control**

**EDTA**

**AFPPH**

**AF35**

**AF13**

**AF1**

**TFPPH**

**TF35**

**TF13**

A table with numbers and letters

Description automatically generated

**Table S13**

**Control**

**EDTA**

**AFPPH**

**AF35**

**AF13**

**AF1**

**TFPPH**

**TF35**

**TF13**

A table with numbers and letters

Description automatically generated

**Table S14**

**Control**

**EDTA**

**AFPPH**

**AF35**

**AF13**

**AF1**

**TFPPH**

**TF35**

**TF13**

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**Table S16**

**Table S15**

**Control**

**EDTA**

**AFPPH**

**AF35**

**AF13**

**AF1**

**TFPPH**

**TF35**

**TF13**

**Control**

**EDTA**

**AFPPH**

**AF35**

**AF13**

**AF1**

**TFPPH**

**TF35**

**TF13**



Figure S1: SDS-PAGE analysis of crude supernatant and fractionated PPHs following sequential hydrolysis with Alcalase and Flavourzyme or Trypsin and Flavourzyme. The substrate (Protafy 130) is included as reference.

Figure S2: Relative, intensity-weighted length distribution (n = 3-30) of peptides identified in fraction AF1 from peptidomics analysis by unspecific analysis of LC-MS/MS data in MaxQuant.

Figure S3: Relative, intensity-weighted length distribution (n = 3-30) of peptides identified in fraction AF35 from peptidomics analysis by unspecific analysis of LC-MS/MS data in MaxQuant.

Figure S4: Relative, intensity-weighted length distribution (n = 3-30) of peptides identified in fraction AF35 from peptidomics analysis by unspecific analysis of LC-MS/MS data in MaxQuant.

Figure S5: Relative, intensity-weighted length distribution (n = 3-30) of peptides identified in unfractionated AFPPH from peptidomics analysis by unspecific analysis of LC-MS/MS data in MaxQuant.

Figure S6: Relative, intensity-weighted length distribution (n = 3-30) of peptides identified in fraction TF13 from peptidomics analysis by unspecific analysis of LC-MS/MS data in MaxQuant.

Figure S7: Relative, intensity-weighted length distribution (n = 3-30) of peptides identified in fraction TF35 from peptidomics analysis by unspecific analysis of LC-MS/MS data in MaxQuant.

Figure S8: Relative, intensity-weighted length distribution (n = 3-30) of peptides identified in unfractionated TFPPH from peptidomics analysis by unspecific analysis of LC-MS/MS data in MaxQuant.

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Figure S9: Depiction of one-way ANOVA and Tukey test for FRS (A) and CHE (B) score distribution by pairwise comparison. Significance level is indicated by adjusted p value as: not significant (ns), < 0.0332 (\*), < 0.0021 (\*\*), < 0.0002 (\*\*\*), and < 0.0001 (\*\*\*\*).

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Figure S10: Full UpSet plot showing overlap of identified peptides between PPH fractions and indicating the arithmetic magnitude of the intersects (i.e., number of shared peptides) as well as the total number of peptide IDs within each fraction.

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| Figure S11: Venn diagrams showing shared peptides between fractions of the AF PPH (A) and the TF PPH (B). | |

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| Figure S12: Surface plasmon resonance sorption isotherms. From top to bottom: AF1, AF13, AF35, AFPPH, TF13, TF35, TFPPH |

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| Figure S13: Evolution of SVOPs in 5% fish oil-in-water emulsions during 9 days of storage. | |