

## Supplementary material

## *In silico* analysis and *in vitro* characterisation of the bioactive profile of three novel peptides identified from 19 kDa $\alpha$ -zein sequences of maize,



spipodszizzea\_malzej:-24 is7 a AF LQQQLL PF YPQVGN INA FLQQQLL PF YPQVAN NA FLQQQLL PF YQLALT NPA FLQQQLL PF YQLALT NPA ASYQH I IGGAL F spipodszizzea\_malzej:-24 is7 a AF LQQQL IL FF SQLAAN RAS FLTQQL IL PF YQLAN FLQQQLL PF YQLALT DPA ASYQH I IGGAL F spipodszizzea\_malzej:-24 is7 a AF LQQQL IL FF SQLAAN RAS FLTQQL IL PF YQLAN PT LLQLQQLL PF YQLALT DPA ASYQH I IGGAL F spipodszizzea\_malzej:-24 is7 a AF LQQQL IL FF SQLAAN RAS FLTQQL IL FY YQLAN PT LLQLQQLL PF YQLALT DPA ASYQH I IGGAL F spipodszizzea\_malzej:-24 is7 a AF LQQQL IL FF SQLAAN RAS FLTQQL IL FY YQLAN PT LLQLQQLL PF YQLALT DPA ASYQH I IGGAL F spipodszizzea\_malzej:-24 is7 a AF LQQQL IL FF SQLAAN SFA AF LTQQL IL FY YL MT AP NYGTLLQLQQLL FF YQLALT DPA ASYQH I IGGAL F spipodszizzea\_malzej:-24 is7 a AF LQQQL IL FF SQLAAN SFA AF LTQQL IF FY LMT AP NYGTLLQLQQLL FF YQLALT DPA ASYQH I IGGAL F spipodszizzea\_malzej:-24 is6 a AF LQQQL IFF SQLAAN SFA AF LTQQL IF FY LMT AP NYGTLLQLQQL IFF NQLALT NPA AF YQQF I IGGAL F spipodszizzea\_malzej:-24 is6 a AF LQQQL IFF SQLAG YS FA FL TQQL IF FY LMAY PNAGTLLQLQQL IFF NQLALT NPA AF YQQF I IGGAL F spipodszizzea\_malzej:-24 is6 i AF VLQQQL IFF SQLAG YS FA FL TQPQL IFF YQH YA PNAGTLLQLQQL IFF NQLALT NPA YF YQP I IGGAL F spipodszizzea\_malzej:-24 is6 i AF VLQQQL IFF SQLAG YS FA FL TQPQL IFF YQH YA PNAGTLLQLQQL IFF NQLALT NPA YF YQP I IGGAL F spipodszizzea\_malzej:-25 is6 i AF VLQQQL IFF SQLAG YS FA FL TQPQL IFF YQH YA PNAGTLLQLQQL IFF NQLAT NNA YF YQP I IGGAL F spipodszizzea\_malzej:-25 is6 i AF VLQQQL IFF SQLAG YS FA FL TQQL IFF YLMA PNAGTLLQLQQL IFF NQLAT NNA YF YQP I IGGAL F spipodszizzea\_malzej:-25 is6 AF VLQQQL IFF SQLAG YS FA FL TQQL IFF YLMA PNAGTLLQLQQL IFF NQLAT NNA YF YQP I IGGAL F spipodszizzea\_malzej:-25 is6 AF VLQQU IFF SQLAG YS FA FL TQQL IFF YLMA PNAGTLLQLQQL IFF NQLAT NNA YF YQP I IGGAL F spipodszizzea\_malzej:-25 is6 AF VLQQU IFF SQLAG YS FA FL TQQU IFF YLMA PNAGTLLQLQU IFF NQLAT NTAFY YQP I IGGAL F spipodszizzea\_malzej:-25 is6 AF VLQQU IFF SQLAG YS FA FL TQQU IFF YLMA PNAGTLLQLQU IFF NQLAT NT FYY YQP I IGGAL F spipodsziZZE



233

230

## Molecules 2020, 25, 5405

А



**Figure S2.** LC-ESI MS analytical data of 19ZP1 showing in (**A**) the ion current (upper), UV chromatogram at 220nm (middle) and full scan mass spectrum (lower), and (**B**) the UV chromatogram at 254 nm. The mobile phase consisted of a linear gradient of acetonitrile (20% to 70%) in 0.1% aqueous formic acid. \*: Peak was further isolated by reverse phase high pressure liquid chromatography (RP-HPLC) with buffers 35%–50% B in A (A = 0.1% TFA in water; B = 0.1% TFA in acetonitrile).





**Figure S3.** LC-ESI MS analytical data of CF19ZP1 showing in (**A**) the ion current (upper), UV chromatogram at 220nm (middle) and full scan mass spectrum (lower), and (**B**) the UV chromatogram at 254 nm. Chromatographic data of peptide samples used. Plot of analytical HPLC (left column) and ESI–MS spectrum (right column) of 19ZP1 (a), 19ZP2 (b) and 19ZP3 (c) peptides. The mobile phase consisted of a linear gradient of acetonitrile (20% to 70%) in 0.1% aqueous formic acid.



**Figure S4.** LC-ESI MS analytical data of 19ZP2 showing in (**A**) the ion current (upper), UV chromatogram at 220nm (middle) and full scan mass spectrum (lower). Purity was calculated from the ratio of the two abundant peaks, where the peak at RT 12.13 is the product peak. UV chromatograms at 220 (**B**) and 254 nm (**C**). (**D**) ESI mass spectrum of the peak at RT 12.21, which corresponds to the product peak. (**E**) ESI mass spectrum of the peak at RT 13.54, which corresponds to a side product in which deamidation of the aspartate residue occurred. The mobile phase consisted of a linear gradient of acetonitrile (20% to 70%) in 0.1% aqueous formic acid. Peak at 12.3 min was further purified by reverse phase high pressure liquid chromatography (RP-HPLC) with buffers 35%–50% B in A (A = 0.1% TFA in water; B = 0.1% TFA in acetonitrile)

А



**Figure S5.** LC-ESI MS analytical data of CF19ZP2 showing in (**A**) the ion current (upper), UV chromatogram at 220nm (middle) and full scan mass spectrum (lower), and (**B**) the UV chromatogram at 254 nm. Beside the product, a side product (MW 3128 g/mol) is visible, which presumably represents a piperidide adduct of the product. The mobile phase consisted of a linear gradient of acetonitrile (20% to 70%) in 0.1% aqueous formic acid.





**Figure S6.** LC-ESI MS analytical data of 19ZP3 showing in (**A**) the ion current (upper), UV chromatogram at 220nm (middle) and full scan mass spectrum (lower), and (**B**) the UV chromatogram at 254 nm. The mobile phase consisted of a linear gradient of acetonitrile (20% to 70%) in 0.1% aqueous formic acid.

А



**Figure S7.** LC-ESI MS analytical data of CF19ZP3 showing in (**A**) the ion current (upper), UV chromatogram at 220nm (middle) and full scan mass spectrum (lower), and (**B**) the UV chromatogram at 254 nm. The mobile phase consisted of a linear gradient of acetonitrile (20% to 70%) in 0.1% aqueous formic acid.