Supporting Information

Protein/protein nanocomposite based on whey protein nanofibrils in a whey protein

matrix

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Secondary structure		Relative area of Gaussian components (size of each resolved peak in %)										
Position (cm ⁻¹)	Assignment	WPI-0	WPI-N-2.5	WPI-N-5	WPI-N-10	WPI-N-15	WPI-N-100	WPI-D-2.5	WPI-D-5	WPI-D-10	WPI-D-15	WPI-D-100
1618	β-Sheets, strongly hydrogen-bonded peptide group	13.0 ± 0.5	12.5 ± 1.0	12.8 ± 1.0	12.4 ± 0.7	13.1 ± 1.1	17.7 ± 1.9	12.6 ± 0.7	11.2 ± 1.3	11.0 ± 1.5	9.6 ± 1.7	28.7 ± 1.4
1625	β-Sheets, strongly hydrogen-bonded peptide group	16.8 ± 0.5	18.8 ± 1.9	18.0 ± 2.3	20.4 ± 1.3	18.8 ± 2.2	30.7 ± 2.7	18.7 ± 0.8	22.0 ± 3.0	26.1 ± 3.7	31.5 ± 3.6	27.3 ± 1.7
1634	β-Sheets, weakly hydrogen-bonded peptide group	20.4 ± 0.5	19.1 ± 0.5	19.1 ± 1.9	17.8 ± 0.7	18.9 ± 1.0	12.7 ± 3.5	19.2 ± 1.3	17.7 ± 1.5	14.1 ± 2.1	11.9 ± 2.9	3.4 ± 1.5
1644	Unordered	8.3 ± 0.5	8.4 ± 0.6	8.9 ± 1.3	8.9 ± 1.0	8.5 ± 1.1	5.8 ± 0.9	8.9 ± 0.9	8.1 ± 0.3	10.5 ± 1.9	7.3 ± 2.9	14.8 ± 2.6
1651	α -Helices and random coils	7.5 ± 1.2	8.2 ± 1.4	7.8 ± 1.5	7.4 ± 1.5	7.3 ± 0.8	4.2 ± 1.3	7.7 ± 0.5	11.5 ± 1.0	10.0 ± 0.5	13.2 ± 1.3	2.3 ± 1.0
1658	α-Helices	15.2 ± 2.0	14.9 ± 2.3	15.1 ± 1.4	15.5 ± 2.2	14.7 ± 3.1	10.4 ± 2.4	15.8 ± 1.1	12.2 ± 1.8	10.8 ± 2.8	9.5 ± 1.2	4.4 ± 0.9
1667	β-Turns	9.1 ± 1.3	8.4 ± 2.0	8.4 ± 1.6	7.8 ± 2.9	8.0 ± 2.3	8.3 ± 2.0	7.1 ± 1.6	8.1 ± 1.2	8.4 ± 1.2	7.9 ± 1.1	12.7 ± 1.6
1680	β-Sheets, weakly hydrogen-bonded peptide group	6.6±0.5	6.9 ± 1.4	7.0 ± 1.1	7.3 ± 1.5	8.0 ± 0.8	6.4 ± 1.9	7.4 ± 1.3	6.7±0.6	6.7±0.7	6.9 ± 0.5	2.4 ± 0.8
1691	β-Turns	3.1 ± 0.2	2.7 ± 0.5	2.8 ± 0.3	2.6 ± 0.3	2.7 ± 0.2	3.9 ± 0.8	2.6 ± 0.5	2.5 ± 0.3	2.3 ± 0.1	2.1 ± 0.2	3.9 ± 0.5

Table S1. Relative content of secondary structure in the amide I region (N: non-dialysed. D: dialysed)

±-values are standard deviations



Figure S1. Modulus (a), stress (b) and extensibility (c) as a function of PNF content. Filled and unfilled symbols correspond to dialysed and non-dialysed films.