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Mass spectrometry based proteomics as foodomics tool in research and assurance of food quality and safety

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2	assurance of food quality and safety
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18	ABSTRACT
19	Background: As a comprehensive discipline that studies food and nutrition, foodomics requires
20	reliable qualitative and quantitative information about the food proteome component in order to
21	extract new, integrative information from the complex multivariable space of omics. This
22	information is necessary to achieve a higher level of understanding of processes in food science
23	and technology, consequently new functions of food and improved markers of food quality and
24	safety and transform the concept of food safety.
25	Scope and Approach: We are presenting mass spectrometry (MS) based proteomic approaches
26	that are being utilized in different proteomic studies, not necessarily only in the field of
27	foodomics. Current analytical capabilities of MS-based proteomics together with sample
28	preparation procedures and quantification strategies, and recent technical developments were
29	presented.
30	Key Findings and Conclusions: MS-based proteomics enables the analysis of different aspects of
31	proteins and provides a variety of approaches for reliable quantification of individual proteins
32	and/or food proteome. This is a complex field and its successful implementation requires a

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