

Surfactant protein G

The results indicate [Surfactant protein G](#) (SP-G) as a new [surfactant protein](#) which represents an until now unknown surfactant protein class.

Rausch et al. found mRNA for SP-G to be present in kidneys, heart, testis, umbilical cord and trophoblast, ¹⁾.

[Surfactant protein G](#) was identified by means of [bioinformatics](#) and named surfactant protein G (SP-G) or surfactant-associated protein 2 (SFTA 2) ²⁾.

The protein (SP-G) is encoded on the human chromosome 6, its primary theoretical translation product consist of 78 amino acid residues resulting in a molecular weight of approximately 8 kDa. This putative surfactant protein shows no sequential or structural similarities to surfactant proteins or other known proteins in general and therefore seems to represent a new group of proteins. Furthermore, there is no hard evidence or information neither on the organ or tissue distribution nor on the function of the protein. It is carrying an N-terminal signal peptide of 19 amino acid residues which is essential for protein secretion ³⁾.

Its presence and possible functions in the central nervous system are unknown. Therefore, a study of Krause et al., from the Department for Neurosurgery, University Hospital [Leipzig, Germany](#), aimed to elucidate the presence of SP-G in the [brain](#) and its concentration in normal and pathologic samples of [cerebrospinal fluid](#) in order to gain first insight into its regulation and possible functions. A total of 121 samples of human cerebrospinal fluid (30 controls, 60 [hydrocephalus](#) patients, 7 [central nervous system infections](#), and 24 [brain hemorrhage](#) patients) and 21 [rat](#) brains were included in the study.

CSF samples were quantified using a commercially available [ELISA](#) system. Results were analyzed statistically using [SPSS 22](#), performing Spearman Rho correlation and [ANOVA](#) with Dunnett's post hoc analysis. Rat brains were investigated via [immunofluorescence](#) to determine SP-G presence and colocalization with common markers like [aquaporin-4](#), [glial fibrillary acidic protein](#), [platelet and endothelial cell adhesion molecule 1](#), and [neuronal nuclear antigen](#). SP-G occurs associated with brain [vessels](#), comparable to other conventional SPs, and is present in a set of cortical [neurons](#). SP-G is furthermore actively produced by ependymal and [choroid plexus epithelium](#) and secreted into the [cerebrospinal fluid](#). Its concentrations are low in control subjects and patients suffering from [aqueductal stenosis](#), higher in [normal pressure hydrocephalus](#) ($p < 0.01$), and highest in [infections](#) of the central nervous system and brain hemorrhage ($p < 0.001$). Interestingly, SP-G did correlate with total CSF protein in patients with CNS infections and hemorrhage, but not with cell count. Based on the changes in CSF levels of SP-G in hydrocephalus, brain hemorrhage, and CNS infections as well as its abundance at CSF flow-related anatomical structures closely associated with immunological barrier systems, importance for CSF rheology, brain waste clearance, and host defense is assumable.

Thus, SP-G is a potential new [cerebrospinal fluid biomarker](#), possibly not only reflecting aspects of CNS innate immune responses, but also rheo-dynamically relevant changes of CSF composition, associated with CSF malabsorbtion. However, further studies are warranted to validate this findings and increase insight into the physiological importance of SP-G in the CNS ⁴⁾.

References

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