

## BOTTOM-UP APPROACH BASED ON S-LAYERS AS TEMPLATES IN THE FORMATION OF ORDERED ARRAYS OF METALLIC NANOPARTICLES

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Crystalline bacterial cell surface layer (S-layer) proteins have the intrinsic tendency to self-assemble into two-dimensional arrays in suspension and at various interfaces. S-layer lattices exhibit either oblique, square or hexagonal lattice symmetry with unit cell dimensions in the range of 3.5 to 35 nm. S-layers are generally 5 to 10 nm thick. In addition to the unique properties of native S-layer proteins as well defined binding matrices, genetically engineered S-layer fusion proteins opened a new horizon for the tuning of their structural and functional features.

Metal binding or precipitating peptides as functional domains were fused to S-layer proteins. In detail, a silver or cobalt binding sequence was fused to a C-terminally truncated form of the S-layer protein SbpA of *Bacillus sphaericus* CCM 2177. Amino acid position 1068 is known to be located on the outer surface of the S-layer lattice and therefore the fused functional domains are accessible to metal binding after oriented recrystallization of the S-layer fusion proteins.

Such S-layer fusion proteins can be expected to recrystallize on technologically important substrates (e.g., silicon wafers, gold chips) and shall be exploited as patterning element for the bottom-up fabrication of regularly arranged nanoparticles or carbon nanotubes.

### References:

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