

Supplementary Figure 1. Schematic representation of the peptides resulting from processing of ADAMTS3 in the cellular layer and released in the extracellular medium. Mutations are marked in dark blue. Position of the HA-tag (9 amino acids) is shown in magenta. The areas detected by the antibodies are indicated by green dashed lines. **Fu**, furine cleavage site; **Fu***, likely site based on homology with ADAMTS2 (ref 18); **X**, probable autocatalytic process (ref 18); **SP**, signal peptide; **Dis**, disintegrin domain; **TSP**, thrombospondin domains.