

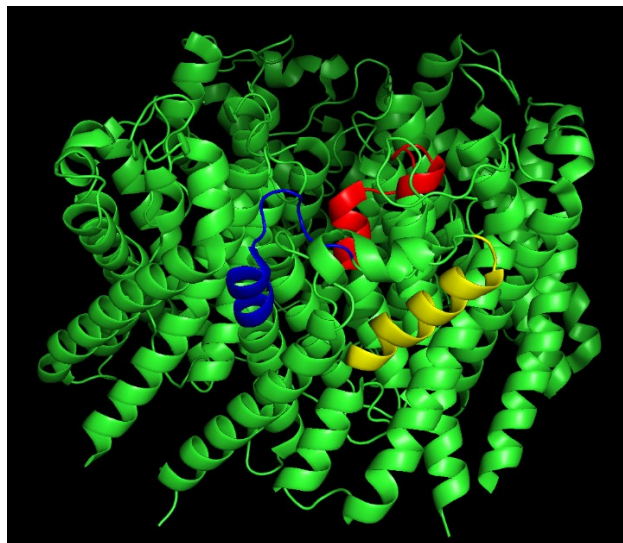


Figure S2 Localization of the three peptides in the predicted 3D models of the HqAQP1 tetramer. The homology models of HqAQPs were generated with the Swiss-Model Work-space (<https://swissmodel.expasy.org/>), using crystal structures of the human AQP10 (PDB ID: 6f7h) as templates. Peptide 1 is shown in red, peptide 2 is shown in blue and peptide 3 is shown in yellow.