

Figure S6. Structuration of the 15 motifs identified in the 52 full-length *OeuAQP* protein sequences (**Figure 2**). Motif analysis was performed by the MEME web server. Details in amino acid constituting these motifs are listed in the legend, and residue occurrences are provided on page 2 of the file.

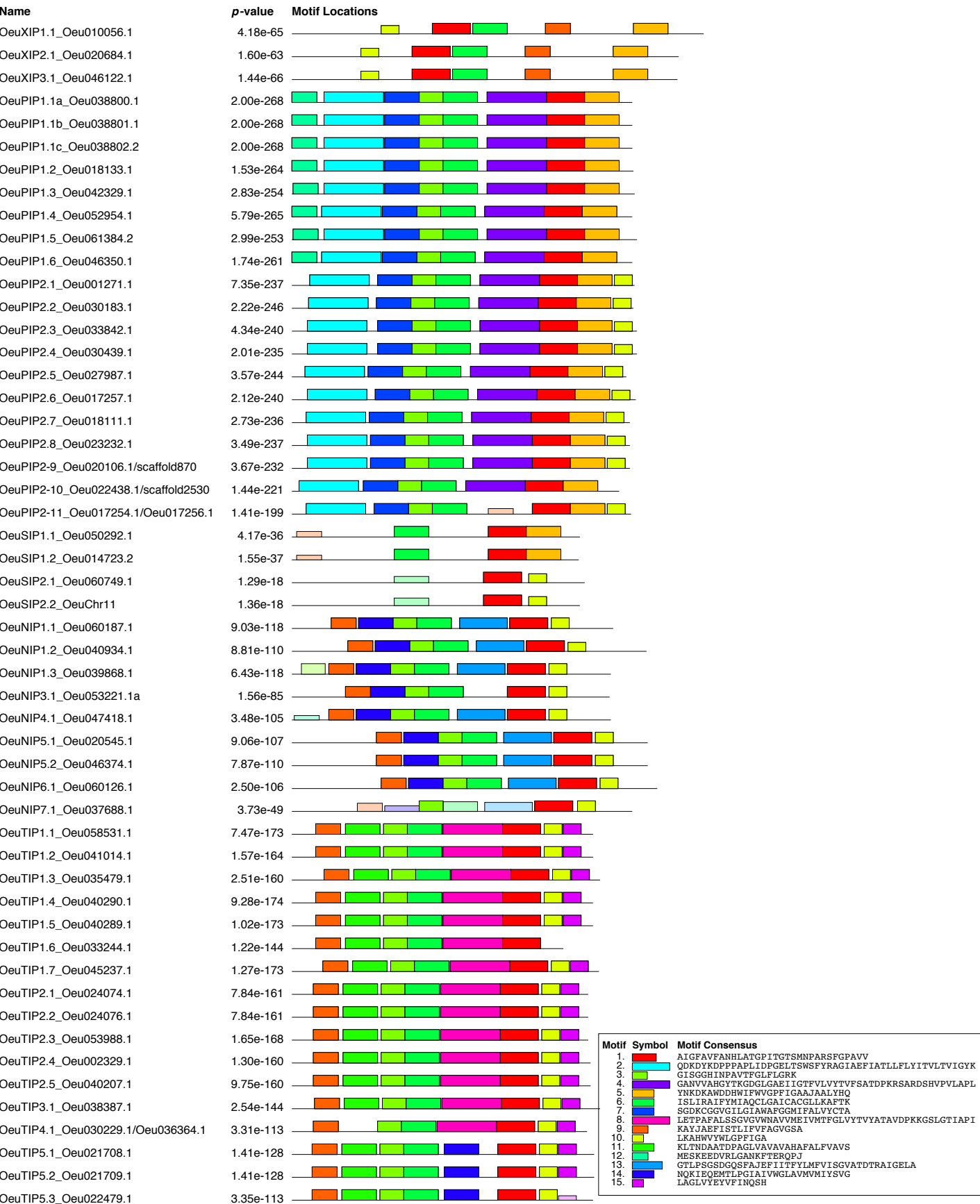


Figure S6, suite. Amino acids constituting the fifteen motifs of the 50 full-length *Oeu*AQP protein sequences (Figure 2). Motif analysis was performed by the MEME web server.

