



Supplementary Figure S1. Chromatograms showing Alamandine standard at different concentrations.



Supplementary Figure S2. Amino acid sequence alignment of teleost ace2 and ace2-like protein isoforms by NCBI BLAST. The phylogenetic tree (built with Neighbor Joining method) comprises 95 sequences from the result of processing (by BLASTP) the ace2 reference protein sequence of *C. auratus* (Acc. N. XP_026131313.1) for matching the *refseq_protein* database, subset of Teleostei (taxid: 32443).