Supplementary Figure 1. Phylogenetic tree of TRPA subfamily. *H. sapiens, M. musculus, D. melanogaster* and *C. elegans* members of these families are color coded, and the proposed names of the two *C. elegans* members is in parenthesis. ClustalW was used to align the transmembrane domains and the alignment is presented as a phylogram. The number of predicted ankyrin repeats is listed.