

receptor	conservation between orthologs (% aa identity)
Rhodopsin	94.2±1.3
ADRB1	92.2±2.8
ADRB2	87.3±2.8
MC4R	94.0±2.0
V2R	85.4±3.1
TAAR1	78.9±5.9
TAAR2	88.9±2.3
TAAR3	87.2±2.7
TAAR4	83.5±3.2
TAAR5	85.6±2.6
TAAR6	84.1±3.3