

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0 Q96R19	TAAR9_HUMAN	Homo sapiens	Trace amine-associated receptor 9	22.903221	T65;T67	NaN	38253038	MVNNFSQAEAVELCYKNVNESCIKT PYSFGPPRSILYAVLGFGAVLAAFGNL LVMIAILHFQQLHTPTNFILASLACA DFLVGVTVMPPFSTVRSVESCWYFGD SYCKFHTCFDTSFCFASLFHLCCISV DRYIAVTDPLTYPTKFTVSVSGICIVL SWFFSVTYSFIFYTGANEEGIEELV VALTCVGGCQAPLNQNWVLLCFLLF FIPNVAMVFIYSKIFLVAKHQARKIES TASQAQSSSESYKERVAKRERKAAK TLGIAMAAFLVSWLPYLVDVIDAYM NFITPPYVYEILVWCVYNSAMNPLI YAFFYQWFGKAIKLIVSGKVLRTDSS TTNLFSEEVETD	False	True	1.328	1.782	1.353	1.01	0.559	5.0	1.565