

S9 Table

	Term	# Genes	% Genes	Adjusted p-value
ApA (530 genes)	1. Detection of chemical stimulus involved in sensory perception of smell	97	18.3	1.4×10^{-61}
	2. Olfactory receptor	97	18.3	1.1×10^{-61}
	3. Olfactory receptor activity	97	18.3	3.5×10^{-61}
	4. Olfaction	97	18.3	7.1×10^{-61}
	5. Olfactory transduction	97	18.3	3.4×10^{-53}
	6. Sensory transduction	103	19.4	5.8×10^{-52}
	7. G-protein coupled receptor activity	108	20.4	1.6×10^{-49}
	8. GPCR, rhodopsin-like, 7TM	105	19.8	1.5×10^{-46}
	9. G-protein coupled receptor	112	21.1	1.5×10^{-46}
	10. G-protein coupled receptor, rhodopsin-like	103	19.4	6.2×10^{-46}
	11. Transducer	113	21.3	2.4×10^{-44}
	12. G-protein coupled receptor signaling pathway	112	21.1	7.4×10^{-42}
	13. Receptor	132	24.9	4.6×10^{-29}
	14. Topological domain: extracellular	161	30.4	9.4×10^{-20}
	15. Disulfide bond	162	30.6	2.8×10^{-18}
	16. Topological domain: cytoplasmic	179	33.8	1.3×10^{-17}
	17. Glycosylation site: N-linked	202	38.1	3.9×10^{-16}
	18. Odorant binding	26	4.9	3.3×10^{-16}
	19. Disulfide bond	173	32.6	2.8×10^{-16}
	20. Glycoprotein	209	39.4	1.1×10^{-14}
	21. Cell membrane	156	29.4	3.5×10^{-13}
	22. Detection of chemical stimulus involved in sensory perception	22	4.2	1.2×10^{-11}
	23. Plasma membrane	184	34.7	2.7×10^{-11}
	24. Sensory perception of smell	24	4.5	1.4×10^{-9}
	25. Transmembrane region	202	38.1	1.0×10^{-8}
	26. Transmembrane helix	216	40.8	1.8×10^{-8}
	27. Integral component of membrane	199	37.5	9.1×10^{-8}
	28. Transmembrane	216	40.8	2.2×10^{-8}
	29. Transmembrane signaling receptor activity	22	4.2	6.1×10^{-5}
	30. Membrane	255	48.1	4.3×10^{-5}
ApC (392 genes)	1. Extracellular space	45	11.5	3.6×10^{-1}
	2. Biosynthesis of antibiotics	12	3.1	2.8×10^{-1}
ApG (1,011 genes)	1. Domain: Leucine-zipper	17	1.7	5.7×10^{-1}
	2. DNA-binding region: basic motif	20	2.0	9.5×10^{-1}
ApT (223 genes)	1. Commissural neuron axon guidance	3	1.3	9.8×10^{-1}
	2. Disulfide bond	55	24.7	7.6×10^{-1}