

S10 Table

	Term	# Genes	% Genes	Adjusted p-value
CpA (3,412 genes)	1. Topological domain: extracellular	587	17.2	1.9×10^{-5}
	2. G-protein coupled receptor signaling pathway	213	6.2	1.8×10^{-4}
	3. Olfactory receptor activity	112	3.3	7.5×10^{-5}
	4. GPCR, rhodopsin-like, 7TM	174	5.1	1.5×10^{-4}
	5. G-protein coupled receptor, rhodopsin-like	170	5.0	1.0×10^{-4}
	6. Olfactory receptor	112	3.3	7.4×10^{-5}
	7. Detection of chemical stimulus involved in sensory perception of smell	111	3.3	2.5×10^{-4}
	8. Olfaction	114	3.3	5.4×10^{-5}
	9. Topological domain: cytoplasmic	701	20.5	3.7×10^{-4}
	10. Transmembrane region	981	28.8	2.5×10^{-4}
	11. Transducer	207	6.1	5.3×10^{-5}
	12. G-protein coupled receptor	194	5.7	5.4×10^{-5}
	13. Glycosylation site: N-linked (GlcNAc...)	844	24.7	8.1×10^{-4}
	14. Olfactory transduction	112	3.3	2.5×10^{-4}
	15. Integral component of membrane	989	29.0	1.0×10^{-3}
	16. Glycoprotein	906	26.6	1.9×10^{-4}
	17. Sensory transduction	146	4.3	1.9×10^{-4}
	18. G-protein coupled receptor activity	163	4.8	1.7×10^{-3}
	19. Transmembrane	1076	31.5	2.5×10^{-4}
	20. Transmembrane helix	1072	31.4	2.4×10^{-4}
	21. Cell membrane	635	18.6	4.1×10^{-4}
	22. Receptor	350	10.3	6.8×10^{-4}
	23. Nucleosome core	31	0.9	1.6×10^{-3}
	24. Sensory perception of smell	44	1.3	8.6×10^{-2}
	25. Histone-fold	35	1.0	4.0×10^{-2}
	26. Keratin-associated matrix	11	0.3	3.6×10^{-2}
	27. Disulfide bond	670	19.6	3.6×10^{-3}
	28. TAF	10	0.3	4.1×10^{-2}
	29. H4	10	0.3	4.1×10^{-2}
	30. TATA box binding protein associated factor (TAF)	10	0.3	3.9×10^{-2}
	31. Histone H4	10	0.3	3.9×10^{-2}
CpC (1,289 genes)	1. Mitochondrion	119	9.2	3.6×10^{-1}
	2. Neuron projection	30	2.3	5.0×10^{-1}
CpG (2,136 genes)	see Table 2 in the main manuscript			
CpT (625 genes)	1. Signal	153	24.5	7.1×10^{-1}
	2. Secreted	84	13.4	4.8×10^{-1}