Supplemental Table 5A. FPKMs by Gland

Gene ID	PG	SMG	SLG	
Ano1	*27.12	11.88	9.46	
Slc12a2	93.64	62.87	43.82	
Slc26a6	1.18	4.41	*38.57	
Slc9a1	1.89	*5.42	2.75	
Cftr	0.58	* 2.76	0.56	
Lrrc8a	2.85	2.59	2.00	
Clcn2	0.85	*2.30	0.39	
Kcnn4	* <i>75.65</i>	31.25	22.86	
Kcnma1	5.18	4.47	*1.21	
Scnn1a	1.15	*2.27	1.06	
Scnn1b	5.40	*36.28	4.59	
Scnn1g	2.09	*9.81	1.25	
Aqp5	87.56	69.60	80.03	
Chrm1	0.32	0.34	*1.82	
Chrm3	1.32	0.74	1.14	
Adrb1	4.21	5.15	*0.41	
Adrb2	1.12	*3.61	1.30	
Bsnd	1.85	*6.83	1.20	
Lrrc26	174.93	121.86	*202.24	
Slc26a2	*2.48	4.49	5.03	
SIc4a2	1.49	1.32	1.36	
SIc4a4	0.62	0.46	0.77	
SIc4a7	0.52	0.83	1.04	
Slc5a5	0.24	0.31	0.05	
SIc5a8	9.45	9.10	*17.32	

Supplemental Table 5B. FPKMs by Gland and Sex

Gene ID	PG M	PG F	SMG M	SMG F	SLG M	SLG F
Ano1	27.66	26.57	12.44	11.32	9.50	9.42
Slc12a2	85.93	101.36	50.53	75.20	40.68	46.96
Slc26a6	1.14	1.22	2.37	*6.44	39.17	37.96
Slc9a1	1.99	1.79	*8.05	2.79	3.30	2.20
Cftr	0.50	0.66	*4.36	1.16	*0.86	0.26
Lrrc8a	3.20	2.51	2.52	2.66	2.04	1.97
Clcn2	0.89	0.82	*3.59	1.02	0.49	0.30
Kcnn4	77.58	73.73	24.33	38.18	20.48	25.24
Kcnma1	4.24	6.12	3.47	5.48	1.19	1.23
Scnn1a	0.99	1.32	2.86	1.69	1.22	0.90
Scnn1b	5.27	5.52	*57.96	14.60	*7.91	1.27
Scnn1g	1.89	2.28	13.15	6.48	*1.96	0.55
Aqp5	82.39	92.74	47.28	91.91	69.96	90.09
Chrm1	0.24	0.39	0.33	0.34	1.83	1.80
Chrm3	1.42	1.22	0.68	0.80	1.02	1.26
Adrb1	3.54	4.88	3.41	6.89	0.51	0.30
Adrb2	1.02	1.22	4.21	3.00	1.85	0.74
Bsnd	2.18	1.51	8.50	5.15	*1.82	0.57
Lrrc26	186.92	162.93	109.07	134.66	206.72	197.75
Slc26a2	3.04	1.93	4.96	4.02	5.18	4.87
SIc4a2	1.49	1.49	1.19	1.45	1.34	1.39
SIc4a4	0.52	0.71	0.62	0.29	0.77	0.78
Slc4a7	0.51	0.54	0.69	0.97	0.99	1.10
Slc5a5	0.23	0.26	0.49	0.13	0.09	0.02
Slc5a8	9.77	9.13	7.40	10.80	16.29	18.35

Supplemental Table 5C. qPCR values of Cftr and Scnn1 by Gland and Sex

Gene ID	PG M	PG F	SMG M	SMG F	SLG M	SLG F
Cftr	0.45	0.18	5.69	2.12	0.29	*0.60
Scnn1a	3.05	2.34	6.73	5.22	3.14	2.86
Scnn1b	2.10	1.39	*24.26	10.38	0.53	*0.71
Scnn1g	0.58	0.39	*6.26	2.46	0.17	*0.27