

Supplementary table 1

Digital PCR (dPCR) data are presented as absolute mRNA copy numbers per microliter (cp/ μ L) for each individual sample, along with the relative contribution of each muscarinic receptor subtype (M1, M2, M3) to the total M receptor transcript pool (expressed as a percentage of 100%) in 15 different patients (HAP#).

<i>Sample</i>	<i>M1 receptor</i>		<i>M2 receptor</i>		<i>M3 receptor</i>	
	<i>cp/ μl</i>	<i>cp%</i>	<i>cp/ μl</i>	<i>cp%</i>	<i>cp/ μl</i>	<i>cp%</i>
HAP1	0.92	0.18	491.36	97.76	10.35	2.06
HAP2	0.3	0.06	448.32	97.1	13.08	2.83
HAP3	0.1	0.03	306.8	94.8	16.71	5.16
HAP4	0.52	0.11	465.7	97.31	12.36	2.58
HAP5	0.41	0.09	431.9	99.15	3.3	0.76
HAP6	0.3	0.08	379.3	96.74	12.47	3.18
HAP7	0.33	0.13	247.17	93.46	16.96	6.41
HAP8	0.22	0.06	367.02	97.79	8.07	2.15
HAP9	0.46	0.21	204.61	93.62	13.48	6.17
HAP10	0.26	0.09	276.29	97.46	6.93	2.44
HAP11	0.26	0.05	533.9	98.72	6.65	1.23
HAP12	0.39	0.18	191.3	89.94	21.02	9.88
HAP13	0.26	0.04	623.8	96.32	23.55	3.64
HAP14	0.3	0.1	294.37	98.07	5.51	1.84
HAP15	0.03	0.01	382.06	96.84	12.44	3.15
average		0.09		96.34		3.57