Table 1. Relative sst1, 2, 3 and 5 mRNA expression

	Patient	Median*
sst1	3.192	0.677
sst2	0.834	0.791
sst3	11.696	0.739
sst5	2.721	0.961

*median expression in 23 somatotropinomas
sst: somatostatin receptors.
Values are expressed as fold change, which indicates how many times the gene is over- or underexpressed in the tumour tissue compared to normal pituitary tissue.