Supplementary Table 1. Identity and features of differentially expressed genes in immature and differentiated cells of the ultimobranchial lineage

A. Calca_pos vs Calca_neg Ubb cells

Gene	Protein	Development	Endocrine	Function(s)
Nnat	Neuronatin	neural fate, brain	pituitary, pancreatic beta-cells	calcium regulation
Mest	Mesoderm-specific transcript homolog	mesoderm	adrenal	EMT
Foxa2	Forkhead box A	endoderm, lung, liver, pancreas	thyroid C cells, pancreas	pioneer transcription factor
Btg2	B-cell translocation gene 2	neurogenesis	pancreatic beta-cells	cell cycle
Clpx2	Complexin 2 (synaphin)	-	neuroendocrine	exocytosis
Ascl1	Achaete-scute homolog 1 (MASH1)	neurogenesis	adrenal, thyroid C cells, MTC	pioneer transcription factor
Cdc25b	Cdc25 isoform	early embryogenesis	MTC	cell cycle
Socs2	Suppressor of cytokine signaling 2	neuronal differentiation	pancreatic beta-cells	JAK/STAT signaling
Hair1	Hoxa adjacent long noncoding RNA 1	ESC lineage differentiation	-	RA regulation
Sst	Somatostatin	neuroendocrine differentiation	neuroendocrine, MTC	peptide hormone
Krt7	Keratin 7	epithelial differentiation	neuroendocrine	cytoskeleton
Prox1	Prospero homeobox 1	cell fate determination	neuroendocrine, thyroid C cells, MTC	EMT, secretory pathway

B. Calca neg vs Calca pos Ubb cells

2.			
Gene	Protein	Function(s)	
Hmcn1	Hemicentin 1 (HMCN1)	ECM, basement membrane organization, cell anchorage	
Plagl1	PLAGL1 (ZAC-1)	transcription factor, ECM regulation, anti-proliferative	
Hs6st2	Heparan sulfate 6 sulfotransferase 2	ECM regulation, morphogen/growth factor activation	
Rbm12b2	RNA binding motif protein 12	post-transcriptional regulation	
Smtnl2	Smoothelin-like 2	apical actin turnover	
Dscc1	DNA replication and sister chromatid cohesion 1	DNA replication	
Hs6st2 Rbm12b2 Smtnl2	Heparan sulfate 6 sulfotransferase 2 RNA binding motif protein 12 Smoothelin-like 2	ECM regulation, morphogen/growth factor activation post-transcriptional regulation apical actin turnover	

List of genes in A and B relates to data presented in Supplementary figures 10a and b, respectively.

Based on scRNAseq analysis of E12.5 cells confined to Ubb clusters 4, 6 and 8

Abbr: Ubb ultimobranchial body, MTC medullary thyroid carcinoma, EMT epithelial-mesenchymal transition, ECM extracellular matrix