

Supplemental Table 4.
Up-regulated genes in the proliferative area of partial thyroidectomized thyroid vs. intact thyroid¹

Genbank	Gene Symbol	Description	Fold Change	p-value²
NM_029662	Mfsd2	major facilitator superfamily domain containing 2	2.547466933	0.0001
NM_144860	Mib1	mindbomb homolog 1	2.4223825	0.0131
NM_010779	Mcpt4	mast cell protease 4	2.35909455	0.0005
NM_007529	Bcan	brevican	2.258720667	0.0112
NM_145927	Fntb	farnesyltransferase, CAAX box, beta	2.213885667	0.0231
NM_016974	Dbp	D site albumin promoter binding protein	2.1370478	0.0274
NM_010149	Epor	erythropoietin receptor	2.079813667	0.0018
NM_001001182	Baz2b	bromodomain adjacent to zinc finger domain, 2B	1.724448267	0.0135
AK032347	Frmd7	FERM domain containing 7	1.669752533	0.0041
AF260925	Nmnat1	nicotinamide nucleotide adenyltransferase 1	1.6465104	0.0139
NM_001025572	Ankrd12	ankyrin repeat domain 12	1.623278733	0.0263
NM_025566	Tnfaip8l1	tumor necrosis factor, alpha-induced protein 8-like 1	1.587914917	0.0289
NM_001024468	Bcat1	branched chain aminotransferase 1, cytosolic	1.58205225	0.0193
NM_130859	Card10	caspase recruitment domain family, member 10	1.5804634	0.0134
NM_016969	Myadm	myeloid-associated differentiation marker	1.561795083	0.034
NM_027769	Cpne3	copine III	1.557383667	0.0153
NM_010942	Nsg1	neuron specific gene family member 1	1.5470056	0.0172
AK036956	Mpp7	membrane protein, palmitoylated 7 (MAGUK p55 subfamily member 7)	1.54085345	0.0409
NM_144945	Lgi2	leucine-rich repeat LGI family, member 2	1.502684533	0.0323

¹Only genes with known gene symbols that were up-regulated more than 1.5-fold in log2 scale are listed.