Supplementary Figure S1: Gene ontology analysis. The graph was generated using Gorilla gene ontology analysis tools. It shows a hierarchical clustering by biological process, metabolic process, or biological regulation, of the genes upregulated in neonatal intermediate cells of the stria vascularis compared to the rest of the cochlear lateral wall. P value is a representation of the confidence that the genes analyzed are involved in a pathway. For genes involved in melanin biosynthesis and melanocyte differentiation, $p < 10^{-9}$. ipsum