# From 1p3 to PI3K

Studies of Neuroblastoma

#### AKADEMISK AVHANDLING

som för avläggande av medicine Doktorsexamen vid Sahlgrenska akademin vid Göteborgs Universitet kommer att offentligen försvaras i hörsal Arvid Carlsson, Medicinaregatan 3, Göteborg Fredagen den 4 februari 2011 kl 9.00

av

### Susanne Fransson

Fakultetsopponent: Professor Rogier Versteeg Department of Human Genetics, Academic Medical Center, University of Amsterdam, Amsterdam, Netherlands

Avhandlingen baseras på följande arbeten:

- I. Fransson S, Martinsson T & Ejeskär K. Neuroblastoma tumors with Favorable and Unfavorable outcome: Significant differences in mRNA expression of Genes Mapped at 1p36.2. *Genes, Chromosomes and Cancer (2007) 46:45-52*
- II. Carén H, Fransson S, Ejeskär K, Sjöberg R-M, Kogner P & Martinsson T. Genetic and epigenetic changes in the common 1p36 deletion in neuroblastoma tumors. Br j Cancer (2007) 19;97(10):1416-24
- III. Fransson S, Abel, F, Eriksson H, Kogner P, Martinsson T & Ejeskär K. Analysis of the PI3K/Akt signaling pathway in Neuroblastoma – Stage dependent expression of PI3K p110 isoforms. (2011) Submitted
- IV. Fransson S, Uv A, Eriksson H, Andersson M K, Wettergren Y, Bergö M, & Ejeskär K. p37delta, a new isoform of PI3K p110delta that increases cell proliferation, is over expressed in human tumors. (2011) Submitted



UNIVERSITY OF GOTHENBURG

## From 1p3 to PI3K

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### Susanne Fransson

#### Department of Medical and Clinical Genetics, Institute of Biomedicine, Sahlgrenska Academy at University of Gothenburg, Sweden

Neuroblastoma (NB) is a tumor of the sympathetic nervous system and is the most common extra-cranial tumor of childhood, accounting for 7% of all pediatric malignancies. Despite recent advances in therapeutics, outcome is still fatal for patients with aggressive NB and side-effects of treatment are severe. These are important reasons to gain further knowledge of the biology behind NB.

*Aims:* The objective of this thesis was to explore genes and gene products that might contribute to initiation and progression of NB and possibly also other malignancies. Main focus has been on the chromosomal region 1p36.2-3 and participants of the PI3K/Akt signaling pathway.

**Results:** Real-time expression analysis of 30 genes at 1p36.2-3 showed that *TNFRSF9* and *PIK3CD* were down regulated in 1p-deleted compared to non-deleted NB tumors. Studies of the same region showed four genes (*ERRFI, CASZ1, RBP7* and *PIK3CD*) possibly regulated by epigenetically means. Bisulphite sequencing of these four genes in NB cell lines and primary tumors showed that methylation probably is not involved but that histone deacetylation could be implicated in their regulation. Some rare sequence variants were also identified in *ERRFI* and *PIK3CD*. *PIK3CD* encodes a catalytic subunit of the phosphatidylinositol 3-kinase (PI3K) that is involved in activation of Akt. Analysis of mRNA levels in a set of 88 genes associated to PI3K/Akt signaling showed that *PDGFRA*, *PIK3R1*, *PIK3CD*, *PRKCBI*, *PRKCZ* and *EIF4EBP1* were differentially expressed comparing stage 1-2 to stage 4 NB. At the protein level a stage-dependent expression of the different catalytic isoforms were detected, where levels of p110a were higher in stage 4 tumors compared to stage 1-2, while the opposite was seen for p1108. Stage 4 NB also had higher levels of phosphorylated Akt (T308 and S473) compared to low stage NB. Furthermore, levels of phosphorylated Akt T308 showed inverse correlation to protein levels of Pten.

We have also identified a novel splice variant p378, encoded by *PIK3CD*. Usage of an alternative donor site leads to truncation in the RAS-binding domain and loss of the catalytic domain. Despite the truncation, p378 interacts with RAS and there is a strong correlation between protein levels of p378 and RAS in primary cells. Expression of p378 is increased in human cancers of the ovaries and colon and ubiquitous expression of the human p378 in *Drosophila* increased the body size of the fly. Furthermore, over-expression of p378 in HEK-293 and mouse embryonic fibroblasts increased proliferation and invasive properties compared to controls, indicating a role in tumorgenicity.

**Conclusion:** Analysis of expression levels of genes and proteins could be used for pinpointing important genes and pathways. This thesis has added more knowledge about the genes at 1p36.2-3, a region commonly deleted in NB, as well as the PI3K/Akt signaling in NB. We have also described a new splice variant of p1108 that is expressed in human cancer and increases proliferation *in vitro* and *in vivo*.

Keywords; cancer, tumor, neural crest, mucosa, premalignant, neuroblastoma, tumor suppressor gene, oncogene, gene expression, epigenetics, splicing, signaling, 1p36, PI3K, Akt, Western blot, TaqMan, *PIK3CD*, *PIK3R1*, p1108, p378, p85, RAS

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