





### Understanding Cancer Lecture 12 **Types of signalling** pathway: normal and dysregulated Notch

DR HAFSA WASEELA ABBAS www.hafsaabbas.com



# **RECAP:**

### What you hopefully should understand so far from Lecture 11



The human Philadelphia (Ph) chromosome arises from a translocation between chromosomes 9 and 22 of the t(9;22). This affects the ABL and BCR genes.



There are two main cancers associated with ABL-BCR translocation: chronic myeloid leukaemia (CML) that mainly affect middle-aged patients and B-cell acute lymphoblastic leukaemia (B-ALL) that affect paediatric patients under 15 years old

Other Abl fusion can occur that can cause T-cell acute lymphoblastic leukemia (T-ALL).





# What will we learn today?

- The structure of Notch receptors
- The types of Notch receptors



Receptor activation: Normal Notch signalling pathway

Signal transduction: Normal Notch signalling pathway

Cellular response: Normal Notch signalling pathway



Examples of cancers where dysregulated Notch signalling pathway

### GENTLE REMINDER An ideal way of learning:

Monday Tuesday Wednesday Thursday Friday Saturday Sunday

### Mini-lectures.

### Approximate total time: 1 hour Divide over 7 days at your <u>own pace</u>. Challenge yourself with a quiz!



www.hafsaabbas.com

### **RECAP: How to support your learning?**



Glossary to help understand what key words mean.



Summary doodle revision posters by HN designs.



Quizzes to test your knowledge and reflect.



**Reference list for further reading.** 

Acknowledgements: Special thanks to my parents, family, friends and colleagues for their support and the respected teachers and health professions who taught me and installed the passion of cancer/oncology.

# The structure of Notch receptors

# The structure of Notch receptors

They are transmembrane proteins expressed on the surface of cell membrane.

They consist of **Extracellular region**  *N-terminal EGF repeats between 29-36 Juxtamembrane Negative regulatory region (NRR):* It consists of two divisions: Three Lin-12/Notch repeat (LNR) Heterodimerization domain (HD)



#### Transmembrane domain

#### Intracellular region

- Protein-binding RAM region
- Seven ankyrin (ANK) repeat domains
- □ Transcriptional activation domain (TAD)
- *C-terminal degron domain with high levels of the amino acids proline, glutamate, serine and threonine (PEST)* (Aster et al. 2017; Yuan et al. 2015)

### The types of Notch receptors



Notch1 and Notch2 are expressed in tissue development and adult mammals.

Notch3 are expressed in vascular smooth muscle and pericytes.

Notch4 are expressed in endothelium.



DLL1 and DLL4 are transmembrane proteins and members of the Delta family of ligands.

DLL4 cannot activate Notch receptors and encodes a decoy receptor.

JAG1 and JAG2 are members of Serrate family of ligands.

### Receptor activation

### Receptor activation

### The ligand binds to the Notch receptor.

#### Depends on cleavage by three enzymes

• Furin-like protease.

2

3

- Location: Cleaves at site S1 in the heterodimerization domain
- Function: Creates a heterodimer made of Notch extracellular subunit and a transmembrane Notch subunit.
- It is held off state by NRR.
- ADAM disintegrin and metalloproteases family predominantly ADAM17 or ADAM10
- Location: Cleaves at site S2 outside the transmembrane domain.
- Following the binding of Notch to its ligand, autoinhibitory effect of NRR takes place, ADAM cleaves.

- Two E3 ligases: Neuralized and Mind Bomb
- Function: ubiquitination of ligand cytoplasm.

(Aster et al. 2017)

### Signal transduction

## Signal transduction

An enzyme called presenilindependent gamma secretase releases the Notch intracellular domain (NICD) of the activated receptor and translocates it into the nucleus.



In the nucleus, the NICD binds to the ubiquitous transcription factor that can repress transcription called CSL. CSL is an abbreviation for CBF-1/suppressor of hairless/Lag1. Together, a NICD-CSL complex.

Other ways of activating transcription for is binding with the DNA sequence and the SMART complex in absence of NICD.

The NICD-CSL complex is converted to an activator for the transcription of Notch target genes.

(Yuan et al. 2015)

## Cellular response

### Target genes

#### **NICD-CSL** complex in the nucleus

#### p27cip1/waf1

- Inhibited cyclin dependent kinase 2 (cdk2) activity and halted cell cycle
- G1 phase of the cell cycle by p27(Kip1)
- G2 phase by p21(Waf1)
- □ Cyclin D1 cell cycle regulation
- □ c-Myc growth and cell cycle regulation

#### □ p21

- Inhibitor of cyclin E/cdk2.
- Its expression regulated by tumour suppressor p53 and oncogene K-ras.

(Schmidt et al. 2001; Hayat, 2005)

### Target genes

#### **NICD-CSL** complex in the nucleus

#### **Given** Survivin

• A member of Inhibitors of apoptosis (IAPS)

#### Slug

• Regulates Epithelial-mesenchymal transition (EMT) and is involved in invasion and evading apoptosis.

#### Nanog

- Stem cell differentiation where cells are pluripotent.
- Pluripotency they can become any cell type.

**□** Nuclear factor-kappa B (NF-κB) pathway – later discussed in the series.

(Shih and Yang, 2011; DeBerardinis and Thompson, 2008; Baker, 2009)

### Target genes

**CSL-DNA sequence-SMART complex in the nucleus** 

□ Hes (Hairy Enhance of Split)

Cell fate decisions during tissue development.

□ Hey (Hairy/Enhancer of Spit related with YRPW motif)

Hey2 decreases skeletal mass and regulates bone remodeling.

**Example of tissue: heart** 

Development and maturation of heart cells in the chambers of the heart and blood vessels (arterial and ventricular cardiomyocytes) and the inner layer of the heart (endocardium)

(Piscione et al. 2004; Zanotti and Canalis, 2013; Yuan et al. 2015)

### Overview of cellular responses



Embryo development Apoptosis

(Yuan et al. 2015)

Examples of cancers where dysregulated Notch signalling pathway

### Examples of cancers

Liver	Small cell lung cancer	T-cell acute lymphoblastic leukemia (T- ALL)
Breast	Prostate	Colorectal
Central Nervous System	Throat cancer	Ewing sarcoma (bone cancer)

# Notch

Promotes tumour progression and metastasis

### Suppresses tumour growth

### Promotes tumour progression

Mutations in the PEST domains of Notch1, Notch2 and Notch3 dysregulated the Notch signaling.

Amplification of wild-type Notch receptors and ligands.

Notch4 and Notch ligands genes were mutated at low levels.

This increased the expression of Notch target genes which increased the progression of cancer.

(Aster et al. 2017; Yuan et al. 2015)

### Promotes tumour progression

**DLL4-Notch regulates NF-κB signalling in liver metastasis of human small cell lung cancer (SCLC).** 

- Over 50% of T-ALL tumours have Notch 1 mutations. It stimulated cell proliferation, invasion and chemoresistance.
- □ Notch1 and Notch3 involved in tumour progression and prognosis in Non-small cell lung cancer.
- □ DLL3 and Hes1 overexpression poor survival.

(Aster et al. 2017; Yuan et al. 2015)

### Suppresses tumour progression

□ Notch suppresses SIRT1 and activates the tumour suppressor p53 to induce growth arrest

in B-cell tumours and Ewing sarcoma.

□ Low expression of DLL4-Notch decreased angiogenesis and tumour growth.

(Aster et al. 2017; Yuan et al. 2015)

# By the end of this lecture, you should understand

Notch are transmembrane protein receptors.



Four receptors found in mammals: Notch1, Notch2, Notch3 and Notch4.



The ligand-receptor complex can be cleaved by three enzymes: Furin-like protease, ADAM family and E3 ligases.



The intracellular domain of the activated receptor is released and can bind to CSL to mediate transcription in the nucleus. CSL can also bind without the intracellular domain to induce transcription.



Notch can promote tumour progression and can also suppress.

# Reference list for further reading

Aster, J., Pear, W. and Blacklow, S. (2017) The Varied Roles of Notch in Cancer. Annual Reviews of Pathology. 12:245-275.

Baker, M. (2009). What does Nanog do? *Nature Reports Stem Cells*.

DeBerardinis, R.J. and Thompson, C.B. (2008). Metabolism of Cell Growth and Proliferation. *The Molecular Basis of Cancer*, pp.189–203.

Hayat, M.A. (2005). Pancreatic Carcinoma: An Introduction. *Handbook of immunohistochemistry and in situ hybridization of human carcinomas*, pp.279–304.

Piscione, T.D., Wu, M.Y.J. and Quaggin, S.E. (2004). Expression of Hairy/Enhancer of Split genes, Hes1 and Hes5, during murine nephron morphogenesis. *Gene Expression Patterns*, 4(6), pp.707–711.

Schmidt, M., Lu, Y., Parant, J.M., Lozano, G., Bacher, G., Beckers, T. and Fan, Z. (2001). Differential Roles of p21<sup>Waf1</sup> and p27<sup>Kip1</sup>in Modulating Chemosensitivity and Their Possible Application in Drug Discovery Studies. *Molecular Pharmacology*, 60(5), pp.900–906.

Shih, J.-Y. and Yang, P.-C. (2011). The EMT regulator slug and lung carcinogenesis. *Carcinogenesis*, [online] 32(9), pp.1299–1304.

Yuan, X., Wu, H., Xu, H., Xiong, H., Chu, Q., Yu, S., Wu, G. and Wu, K. (2015) Notch signaling: An emerging therapeutic target for cancer treatment. *Cancer Letters*, 369 (1).

Zanotti, S. and Canalis, E. (2013). Hairy and Enhancer of Split-related with YRPW Motif (HEY)2 Regulates Bone Remodeling in Mice. *Journal of Biological Chemistry*, 288(30), pp.21547–21557.







### Understanding Cancer Lecture 13 **Types of signalling** pathway: normal and dysregulated Wnt DR HAFSA WASEELA ABBAS www.hafsaabbas.com

