

Wilms' Tumor

“The most common renal tumor in children”



Overview

- What is Wilms' Tumor?
- History.
- Cause and proposed mechanisms of tumor development.
- Identification of Wilms' tumor.
- Staging and Cellular classification.
- Treatments.
- Future outlook.

What is Wilms' tumor?

- It is the most common kidney tumor of children.
- Originates within the kidney during early childhood development.
- Occurs with a frequency of 1 in 10,000 live births.
- Average age of diagnosis—3 yrs.
- Average size—0.5 lbs.
- Two types:
 1. Favorable histology.
 2. Unfavorable histology
 - anaplastic and diffuse anaplastic

History



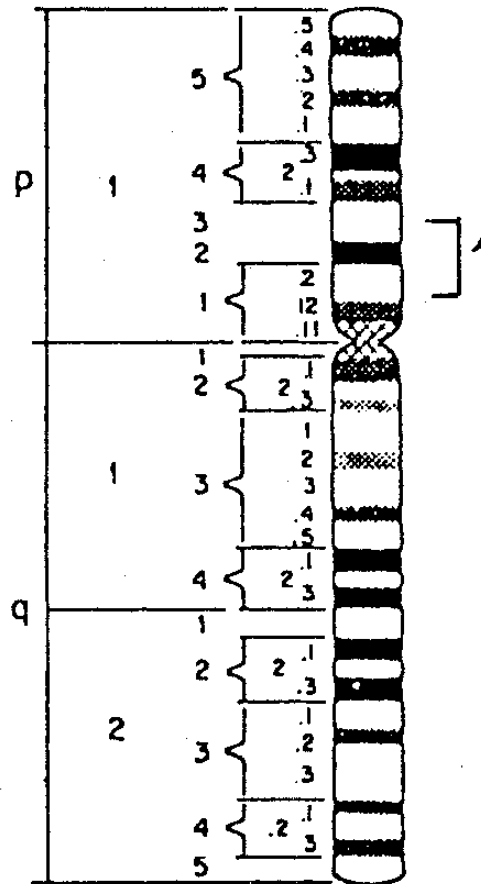
MAX WILMS (1867 - 1918)

- Discovered in 1899 by Max Wilms.
- Establishment of the National Wilms Tumor Study Group (NWTSG).
 - *Goals: --increase the survival rate of children.
 - identify adverse effects of treatments.
 - study long-term effects of treatments.
 - study epidemiology and biology of the Wilms tumor.

Cause and proposed mechanisms of tumor development.

- Caused by one or more changes in several genes.
- Two mutations recognized are on chromosome 11p; 11p13 & 11p15.
- Other loci possibly affected: 1p, 7p, 16p, 17p(the p53 suppressor gene), and 19p.
- Possible causes I will look at are:
 1. Mutations in gene WT1
 2. Mutations in gene WT2
 3. Mutations in p53
 4. Mutations in the beta-catenin pathway

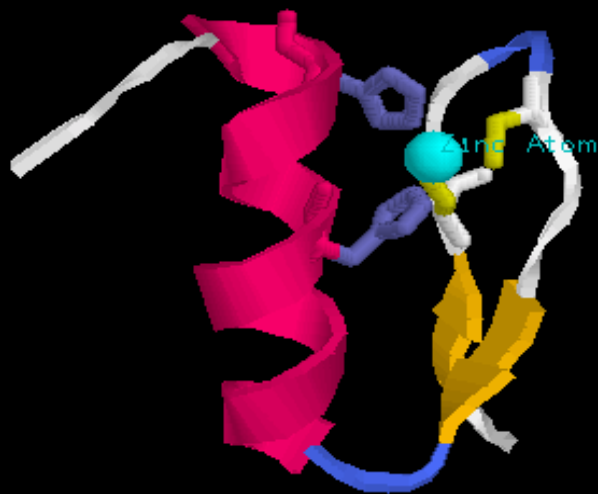
Chromosome 11



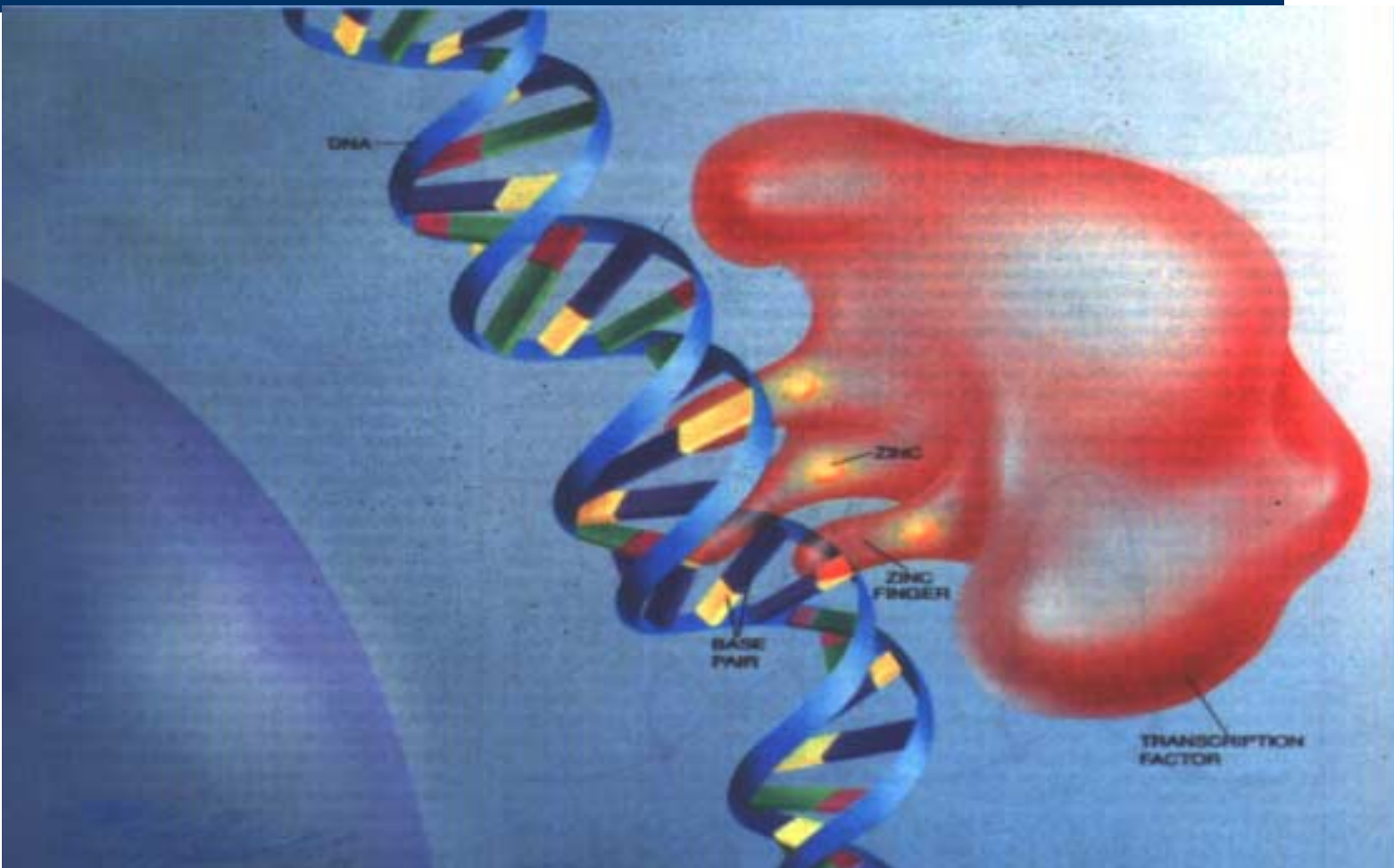
WT1 mutation

- WT1 functions as a zinc finger transcription factor.
 - Amino terminus is rich in proline and glutamine
 - Carboxyl terminus contains four C2H2 zinc finger DNA-binding motifs and a NLS sequence.
- In vivo there are 4 major isoforms that are generated by alternative splicing at two sites. These 4 isoforms are at a constant ratio, which suggest that there functions are independent of each other.
- DNA binding is isoform-dependent.
- WT1 has been shown to be a transcriptional activator or repressor.

Zinc Finger



Zinc Finger



Zinc Finger



Alternative Splicing of WT1



Figure 1 . The four patterns of alternative splicing of the *WT1* gene. The details are explained in the text.

Other mutations found:

- Mutations at 11p15 has now been identified and designated as WT2.
- *B*-catenin mutations were also found associated with Wilms' tumor.

*p*53 identified as a cyclin-dependent kinase (CDK) inhibitor.

Regulation the cell cycle

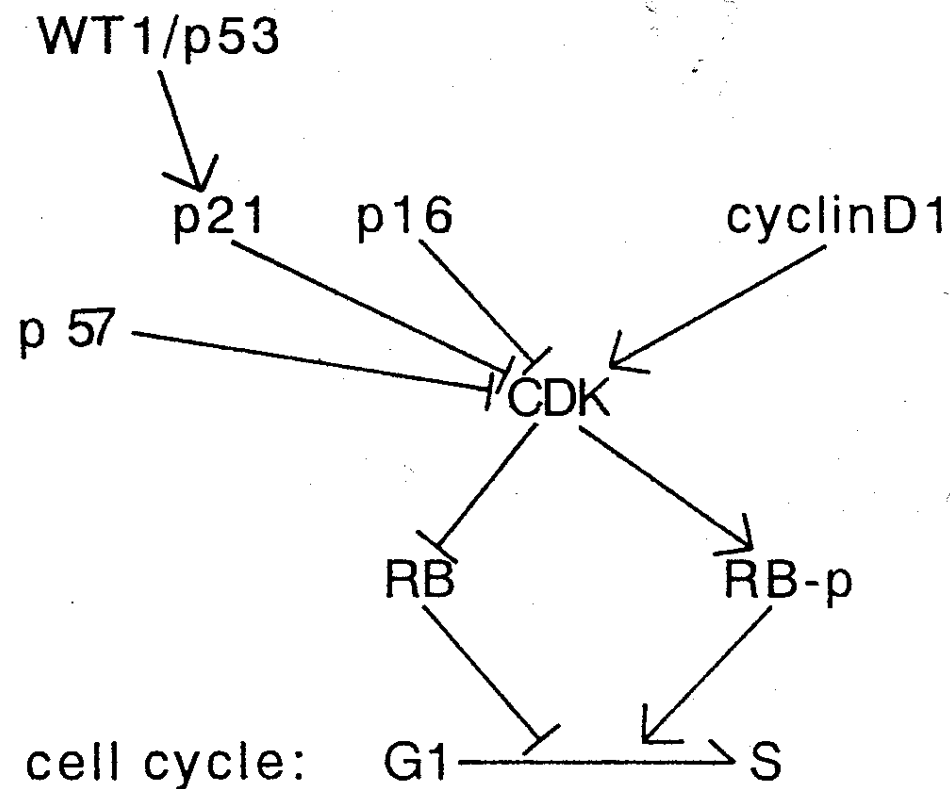


Figure 2. The relation between the proteins that regulate cell cycle progression.

Identification of Wilms' tumor.

- 1st sign is a large lump or swelling in the abdomen.
- 25% also have other symptoms:
 - stomach pain
 - fever
 - blood in the urine
 - high blood pressure
- Medical history, examination, and imaging testing.
- Surgical removal of the tumor is performed in 95% of the cases.

Staging and Cellular classification.

- Staging of the tumor.
 - Stages 1, 2, 3, 4, & 5.
- 2 prognostic groups on the basis of histology:
 - Favorable histology
 - Unfavorable histology
 - Anaplastic
 - Diffuse anaplastic

Treatments.

- Surgery—complete removal of the tumor and surrounding tissue without tumor rupture.
- Treatments range from:
 - 18 weeks of chemotherapy /no radiation
 - 24 weeks of chemotherapy plus abdominal and whole lung radiation
- Relapse treatment depends on these factors:
 - Site of recurrence
 - Tumor histology
 - Length of remission
 - Initial chemotherapy regimen

Future outlook.

- Amount of chemotherapy and radiation used has decreased over the years.
- Further studies associated with WT1, WT2, b-catenins, etc...
- Discoveries of new mutations that lead to Wilms' tumor.
- Therapies that might use anti-telomerase.
- Study #5 will be finished in 2003.



References

- P. Stanhope-Baker, B. Williams, "Identification of Connective Tissue Growth Factor as a Target of WT1 Transcriptional Regulation." *JBC online*. <http://www.jbc.org/cgi/content/full/275/49/38139>.
- J. Laity, J. Chung, J. Dyson, and P. Wright, "Alternative splicing of Wilms' Tumor Suppressor Protein Modulates DNA Binding Activity through Isoform-Specific DNA-Induced Conformational Changes." *Biochemistry* (2000), 39: 5341-5348.
- "Wilms' Tumor." CancerNet. <http://www.cancernet.nci.nih.gov/cgi-bin/srchcgi>.
- M. Hirose, "The role of Wilms' tumor genes." *The Journal of Medical Investigation* (1999), Vol. 46:130-140.
- S. Oh, Y. Song, J. Yim, T. K. Kim, "The Wilms' Tumor Suppressor Gene Represses Transcription of the Human Telomerase Reverse Transcriptase Gene." *The Journal of Biological Chemistry* (1999), Vol. 274, Dec. 24: 37473-37478.

Cont.

- M. J. Coppes, R.M. Egeler, “Genetics of Wilms Tumor.” *Semin. Urol. Oncol.* (1999), Feb. 17(1):2-10.
- A. Menke, L. McInnes, N. D. Hastie, A. Schedl, “The Wilms’ tumor suppressor WT1: approaches to gene function.” *Kidney Int.* (1998), June;53(6): 1512-8.
- D. Haber, “Characterization of the Wilms tumor suppressor gene WT1.” 4/23/2001, <http://www.mgh.harvard.edu/depts/CancerCenter/haber.html>
- “National Wilms Tumor Study Group Mission Statement.” <http://www.nwtsg.org/public/mission.html>
- “Wilms Tumor.” ACS::Wilms’ Tumor Resource Center. <http://www.3.cancer.org/cancerinfo/load -cont.asp>