

INSTRUCTOR  
SCHOOL  
COURSE  
TERM

Name \_\_\_\_\_

Course Section/Period \_\_\_\_\_

## Exploring Genetic Characteristics of Diabetes Type 1

### Objective/s:

Learn about Diabetes Type 1 (insulin-dependant) at the genetic level and practice accessing bioinformatics on the Internet. At the conclusion of this exercise, you should be able to do the following:

- Search for and locate genetic information about a specific human disease
- Use bioinformatics tools to discover detailed information about the structure and function of particular genes
- Use the *BLAST* database to find information about nucleotide sequence homology
- Explore genetic characteristics of Diabetes Type I

### Background:

Diabetes results in an increase in glucose (sugar) in the blood. There are multiple genetic factors contributing to the progression of diabetes in an individual. Sugar provides the body with energy and is essential for humans to survive. Sugar is processed by a hormone (insulin) produced by the pancreas. In order for the body to function effectively, the body must have an appropriate balance between sugar and insulin; diabetes results when there is an imbalance between these two molecules in the bloodstream.

There are two types of diabetes that affect people's ability to process sugar – Type I and II. In Diabetes Type I, an individual may have to take insulin as a medication (insulin-dependant) because their pancreas is not producing enough insulin on its own. Without insulin, sugar may be present in the bloodstream but may not be available for cells to use as energy.

### Directions to Complete Web Activity:

1. Using your Internet browser (mozilla, explorer, netscape, etc.) and your favorite search engine (google, yahoo, etc.), search "OMIM" to find the gene map website. What does OMIM stand for?
2. **Follow the link to the OMIM site.** The resulting page should look like this:



3. Search “OMIM” (select in the drop-down menu); **type in “Diabetes Type 1” in the ‘for’ box and click Go.** You should see something like the following displayed at the top of your search results:

[1: +222100](#)  
DIABETES MELLITUS, INSULIN-DEPENDENT; IDDM  
DIABETES MELLITUS, INSULIN-DEPENDENT, 1, INCLUDED; IDDM1, INCLUDED  
Gene map locus [Xp11.23-q13.3, 12q24.2, 12q24.2, 1p13, 6p21.3](#)

- a) How many articles come up with this search?  
b) What is the name of the first article?  
c) What is the reference number of the first article (6-digit number)?
4. **Link to the first article**, read over it and fill in the following blanks based on the information on the web page:

Description:

Phenotype/Symptoms:

Clinical Features:

5. Look at the “**Gene map locus**” at the top of the page. Which chromosomes play a role in Diabetes Type 1? (\*Review chromosome notation)

\_\_\_\_\_

For the rest of the exercises, we will concentrate on one of the chromosomes in the list above. Specifically, we will explore chromosome 12q.24.2 (the one that appears twice on the list). **Click on the 12q24.2 link and select the fourth location in the list below (TCF1).**

**Map Locations available for 222100**

[PTPN22, PEP, PTPN8, LYP Protein tyrosine phosphatase, nonreceptor-type 22](#)

[IDDM1 Insulin-dependent diabetes mellitus-1](#)

[OAS1, OIAS 2',5'-oligoadenylate synthetase-1](#)

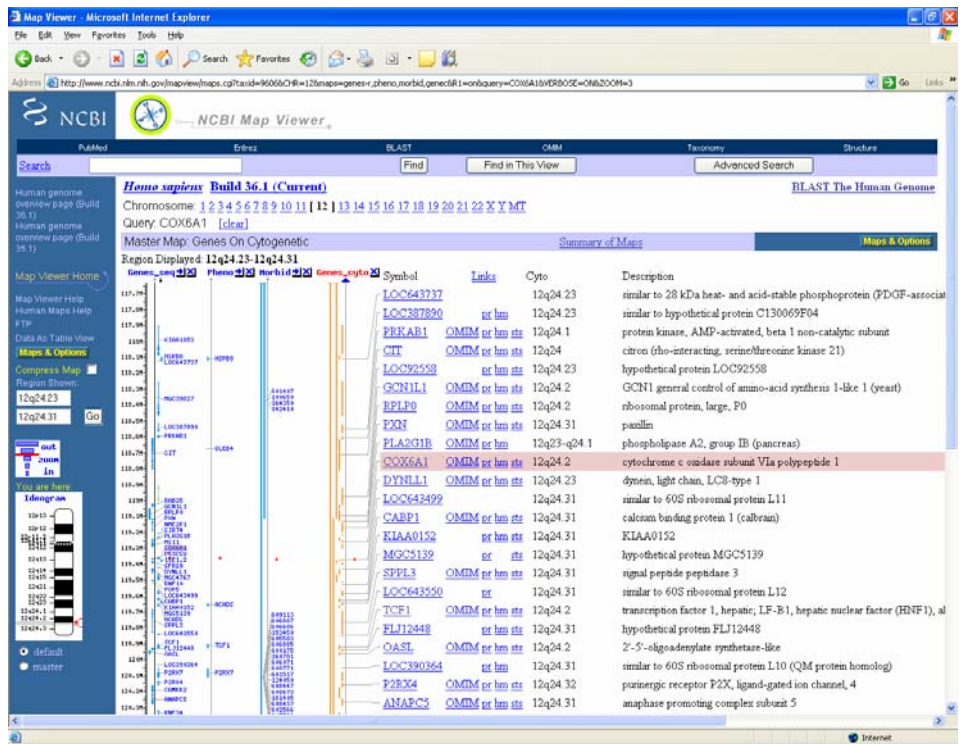
[TCF1, HNF1A, MODY3 Interferon production regulator factor \(HNF1\), albumin proximal factor](#) ←

[FOXP3, IPEX, AIID, XPID, PIDX Forkhead box P3 \(scurlin\)](#)

Now link on the first 'location' (12q.24.2) to see the chromosome ideogram.

6. Take a look at the ideogram on the left side of the page. On which arm (long or short) of the chromosome does the highlighted gene appear?

7. Scan the column labeled (Cyto) for genes mapped on 12q24.2. Write down the names (from the Symbol column) of the genes mapped to this locus.



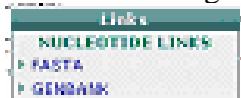
8. Click on the link labeled TCF1. What does TCF1 stand for?

The following page should now be displayed on your screen. Use the information on this page to answer questions 9 and 10:

The screenshot shows the Entrez Gene page for TCF1. The 'Genomic regions, transcripts, and products' section displays the RefSeq for NC\_000012.10. The 'Genomic context' section shows the gene's location on chromosome 12, between LOC398363 and LOC645559, with TCF1 and ORSL genes nearby.

9. Look in the section labeled “Genomic regions, transcripts, and products.” The gene’s reference sequence (RefSeq) is illustrated there. The chromosome number contains several digits, starting with the letters NC. What is the gene’s chromosome number?

10. Click on the number you just wrote down; you will see a dropdown menu like the one shown below. Select **FASTA** from this list, and you will go to a page showing the nucleotide structure of the gene. At the top of this page is the NC number you just clicked on, and next to it a number in square brackets that begins with the letters gi. Write down this number (the gene index?) below:



11. Examine the nucleotide string shown on the page. In the space below, write down the first 21 nucleotide bases in the string.

12. For the nucleotide sequence you wrote above, list the complementary nucleotides in the space below:

13. For the nucleotide string in number 10, give the mRNA transcript below:

14. List the 7 amino acids (in order) described by the mRNA in number 12, using the chart below:

		Second letter				
		U	C	A	G	
First letter	U	UUU } Phe UUC } UUA } Leu UUG }	UCU } UCC } Ser UCA } UCG }	UAU } Tyr UAC } UAA Stop UAG Stop	UGU } Cys UGC } UGA Stop UGG Trp	U C A G
	C	CUU } CUC } Leu CUA } CUG }	CCU } CCC } Pro CCA } CCG }	CAU } His CAC } CAA } Gln CAG }	CGU } CGC } Arg CGA } CGG }	U C A G
	A	AUU } AUC } Ile AUA } AUG Met	ACU } ACC } Thr ACA } ACG }	AAU } Asn AAC } AAA } Lys AAG }	AGU } Ser AGC } AGA } Arg AGG }	U C A G
	G	GUU } GUC } Val GUA } GUG }	GCU } GCC } Ala GCA } GCG }	GAU } Asp GAC } GAA } Glu GAG }	GGU } GGC } Gly GGA } GGG }	U C A G
						Third letter

15. **Click Back** to return to the TCF1 page, the top of which is shown below:

All Databases PubMed Nucleotide Protein Genome Structure PMC Taxonom

Search Gene for [ ] Go Clear

Limits Preview/Index History Clipboard Details

Display Full Report Show 5 Send to

All: 1 Current Only: 1 Genes Genomes: 1 SNP GeneView: 1

1: **TCF1 transcription factor 1, hepatic; LF-B1, hepatic nuclear factor (HNF1), albumin proximal factor** [*Homo sapiens*]  
GeneID: 6927 Primary source: [HGNC:11621](#) updated 28-Jul-2006

Summary ? ↑

16. **Scroll down to the Genomic regions, transcripts, and products section** and find the Nucleotide Messenger RNA number (begins with NM). Write this number below:

17. **Click on this link and choose FASTA.** You will see a nucleotide sequence display similar to the one you used for steps 10-14.

18. **Copy and paste the entire sequence** (beginning with the character “>” – very important!) **into a Notepad file.**

19. **Save the Notepad file (named est1) to your desktop.**

20. **In your browser, scroll back to the top of the page and left-click on “NCBI”** to go to the NCBI (National Center for Biotechnology Information) home page (shown below).

NCBI National Center for Biotechnology Information  
National Library of Medicine National Institutes of Health

PubMed All Databases **BLAST** OMIM Books TaxBrowser Structure

Search All Databases for [ ] Go

21. **In the blue toolbar select BLAST.**

22. **Choose “Nucleotide-nucleotide BLAST (blastn)”** from the list of BLAST programs.

23. Copy and paste entire text of your Notepad file into the Search box of the BLAST page (shown below):

NCBI *nucleotide-nucleotide* **BLAST**  
Nucleotide Protein Translations Retrieve results for an RID

[Search](#)

[Set subsequence](#) From:  To:

[Choose database](#) nr

Now: **BLAST!** or **Reset query** **Reset all**

24. Click the **BLAST!** button. A new screen appears, a portion of which is shown below. Click the **FORMAT** button.

The request ID is

**Format!** or **Reset all**

25. Use the information on the BLAST results page to answer the following questions:

- How many BLAST hits did you get?
- How many of these were close matches (score of 200 or higher)?
- How many very close matches (e-value of 0.0) were found?

26. Click on the link labeled “Distance tree of results” to view a phylogenetic tree of our selected gene. A portion of the top of the resulting window is shown below. Click on the tab labeled “slanted” for a clearer view of the tree and answer the questions below.

Tree method ?      Sequence Label ?      Max Seq Difference ?

Fast Minimum Evolution      Sequence Title (if available)      0.75      Reset

rectangle    **slanted**    radial    force     Show distance

Mouse over an internal node for a subtree or alignment

- a) How many primate species are represented in the tree?
- b) How many human (*Homo sapiens*) chromosomes are represented?
- c) How many rodent chromosomes are represented?

#### Future plans for Curriculum:

- Have students this exercise using different genes/chromosomes related to Diabetes Type 1
- Have students explore genetic information on a disease of their choice using a similar procedure
- Have students run an ORF and explain what it means
- Exercises to look for ESTs
- BLAST exercises for nucleotides/proteins related to Diabetes Type 1 (or other disease)

#### PubMed references

(free, full-text articles in English using subject line “TCF1 and Diabetes Type I”)

1: Pearson ER, Badman MK, Lockwood CR, Clark PM, Ellard S, Bingham C, Hattersley AT.

Contrasting diabetes phenotypes associated with hepatocyte nuclear factor-1alpha and -1beta mutations.

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2: McKinney J, Cao H, Behme MT, Mahon JL, Hegele RA.

Maturity-onset diabetes of the young (MODY) mutation in type 2 diabetes and latent autoimmune diabetes of the adult.

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PMID: 14633836 [PubMed - indexed for MEDLINE]

3: Vilsboll T, Knop FK, Krarup T, Johansen A, Madsbad S, Larsen S, Hansen T, Pedersen O, Holst JJ.

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PMID: 12663611 [PubMed - indexed for MEDLINE]

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Hepatocyte nuclear factor-1 alpha gene mutations and diabetes in Norway.

J Clin Endocrinol Metab. 2003 Feb;88(2):920-31.

PMID: 12574234 [PubMed - indexed for MEDLINE]

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Identifying hepatic nuclear factor 1alpha mutations in children and young adults with a clinical diagnosis of type 1 diabetes.

Diabetes Care. 2003 Feb;26(2):333-7.

PMID: 12547858 [PubMed - indexed for MEDLINE]

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Diabetes Metab. 2002 Feb;28(1):39-44.

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The generalized aminoaciduria seen in patients with hepatocyte nuclear factor-1alpha mutations is a feature of all patients with diabetes and is associated with glucosuria.

Diabetes. 2001 Sep;50(9):2047-52.

PMID: 11522670 [PubMed - indexed for MEDLINE]

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EMBO Rep. 2000 Oct;1(4):359-65.

PMID: 11269503 [PubMed - indexed for MEDLINE]

- 10: Cha JY, Kim H, Kim KS, Hur MW, Ahn Y.  
Identification of transacting factors responsible for the tissue-specific expression of human glucose transporter type 2 isoform gene. Cooperative role of hepatocyte nuclear factors 1alpha and 3beta.  
J Biol Chem. 2000 Jun 16;275(24):18358-65.  
PMID: 10748140 [PubMed - indexed for MEDLINE]
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J Clin Endocrinol Metab. 2000 Jan;85(1):331-5.  
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A case of hepatocyte nuclear factor-1 alpha diabetes/MODY3 masquerading as type 1 diabetes in a Mexican-American adolescent and responsive to a low dose of sulfonylurea.  
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