## From Yeast to Humans—Essential Genes on the Evolutionary Continium.

## A. CBS protein in yeast and humans

Recall that earlier in the term we considered the human gene CBS and its yeast analog cys4. Recall that these genes each encode the protein cystathionine  $\beta$ -synthase that is responsible for converting homocysteine into cystathionine in the cellular pathway of creating cysteine.

- 1. What is cysteine? Is it important for organism's survival?
- 2. What would you expect to be the result of complete absence of the protein product of the yeast *CYS4* gene to be? What about the same question for the human CBS protein?
- 3. Would you expect cells that contain no functional copy of CBS enzyme to accumulate some kind of a compound? If no, why not? If yes, what kind of a compound would you expect that compound to be?
- 4. In the experiments we discussed earlier in the term, what was the phenotype of the *cys4* mutants on complete media?
- 5. As we told you a number of sessions ago, the deficiency in the human analog of *CYS4* gene, CBS, lead to a disease called cysteineurea, resulting in variety of serious conditions, including mental retardation, heart attack, and stroke. What accounts for such a big difference in phenotype between CBS protein deficient yeast and humans?

## **B.** Phylogenetic analysis

Below is a figure from a research paper showing alignment of the amino acid sequences of human, rat, yeast, and *E. coli* CBS proteins.

Figure removed due to copyright reasons.

Please see:

Kruger, W. D., and D. R. Cox. "A yeast system for expression of human cystathionine beta-synthase: structuraland functional conservation of the human and yeast genes." *Proc Natl Acad Sci USA* 91, no. 14 (July 5, 1994): 6614-8.

- 1. What do the dashes in the sequence represent?
- 2. Are the DNA sequences encoding amino acids that are conserved across species above necessarily the same? Why or why not?
- 3. What properties of the particular amino acids allow them to be grouped into the conservative groupings as described in the figure legend above?

4. Look at the genetic code table. Is there a relationship between the codons encoding amino acids in conservative groupings?

- 5. Is CBS a good candidate for creating a phylogenetic tree on the basis of its sequence? Why or why not?
- 6. Are human disease alleles of CBS likely to help with phylogenetic tree construction? Why or why not?
- 7. If constructing a phylogenetic tree on the basis of CBS alignment, would it be more useful to work with the protein or cDNA sequences? cDNA sequences or DNA sequences? Why?

8. Do you expect the human wild type gene to complement yeast CYS4 deficiency? Why or why not?

9. Is there a way to explore the relationship between human CBS protein and yeast CYS4 protein?

**The Genetic Code** 

	U	С	A	G	
U	UUU phe (F)	UCU ser (S)	UAU tyr (Y)	UGU cys (C)	U
	UUC phe	UCC ser	UAC tyr	UGC cys	C
	UUA leu (L)	UCA ser	UAA STOP	UGA STOP	A
	UUG leu	UCG ser	UAG STOP	UGG trp (W)	G
C	CUU leu (L)	CCU pro (P)	CAU his (H)	CGU arg (R)	U
	CUC leu	CCC pro	CAC his	CGC arg	C
	CUA leu	CCA pro	CAA gln (Q)	CGA arg	A
	CUG leu	CCG pro	CAG gln	CGG arg	G
Α	AUU ile (I)	ACU thr (T)	AAU asn (N)	AGU ser (S)	U
	AUC ile	ACC thr	AAC asn	AGC ser	C
	AUA ile	ACA thr	AAA lys (K)	AGA arg (R)	A
	AUG met (M)	ACG thr	AAG lys	AGG arg	G
G	GUU val (V)	GCU ala (A)	GAU asp (D)	GGU gly (G)	U
	GUC val	GCC ala	GAC asp	GGC gly	C
	GUA val	GCA ala	GAA glu (E)	GGA gly	A
	GUG val	GCG ala	GAG glu	GGG gly	G

## STRUCTURES OF AMINO ACIDS at pH 7.0

