

Carbohydrate Engineering for Generating Sialylated Glycoproteins in Insect Cells

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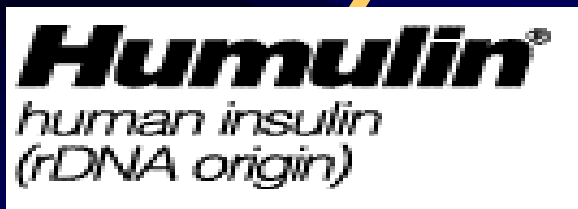
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+Nakano Vinegar, Co

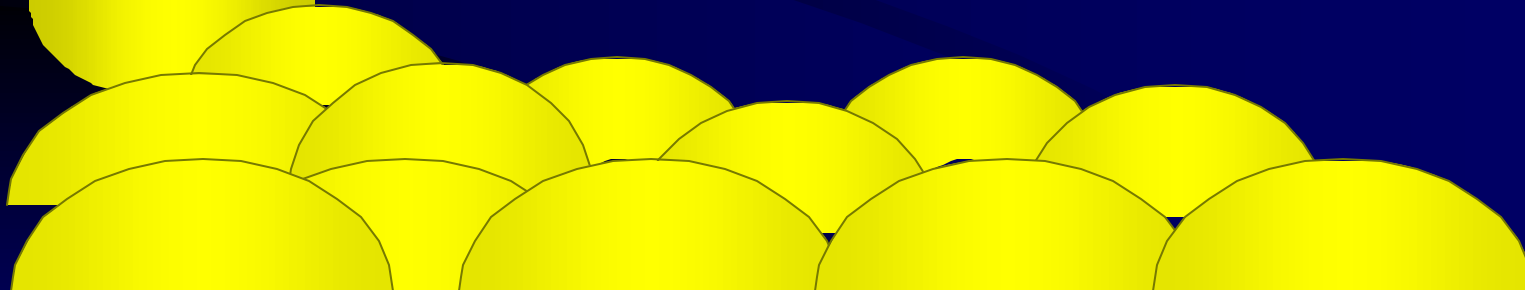
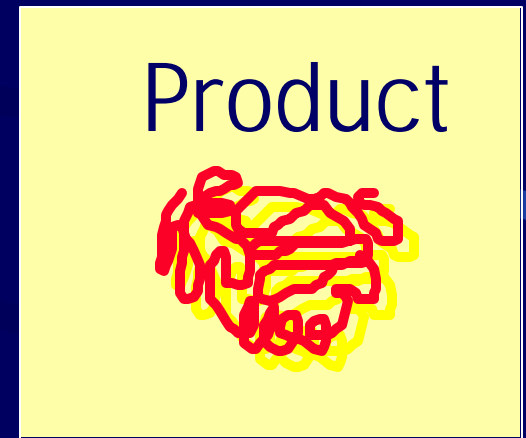
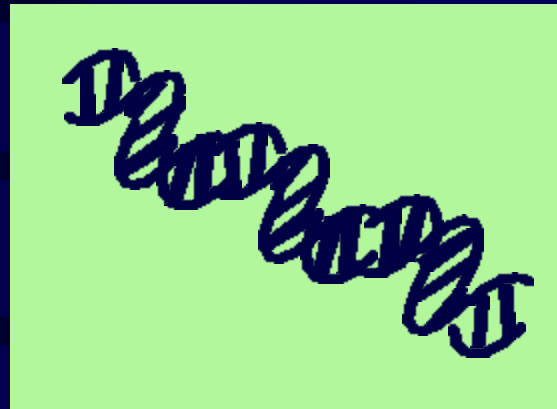
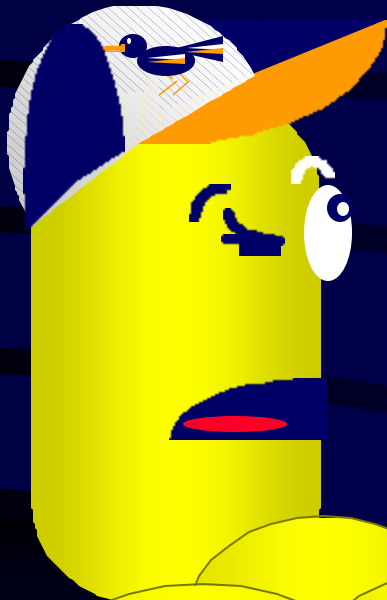
Supported by NSF Metabolic Engineering Program

Biotechnology & Pharmaceuticals

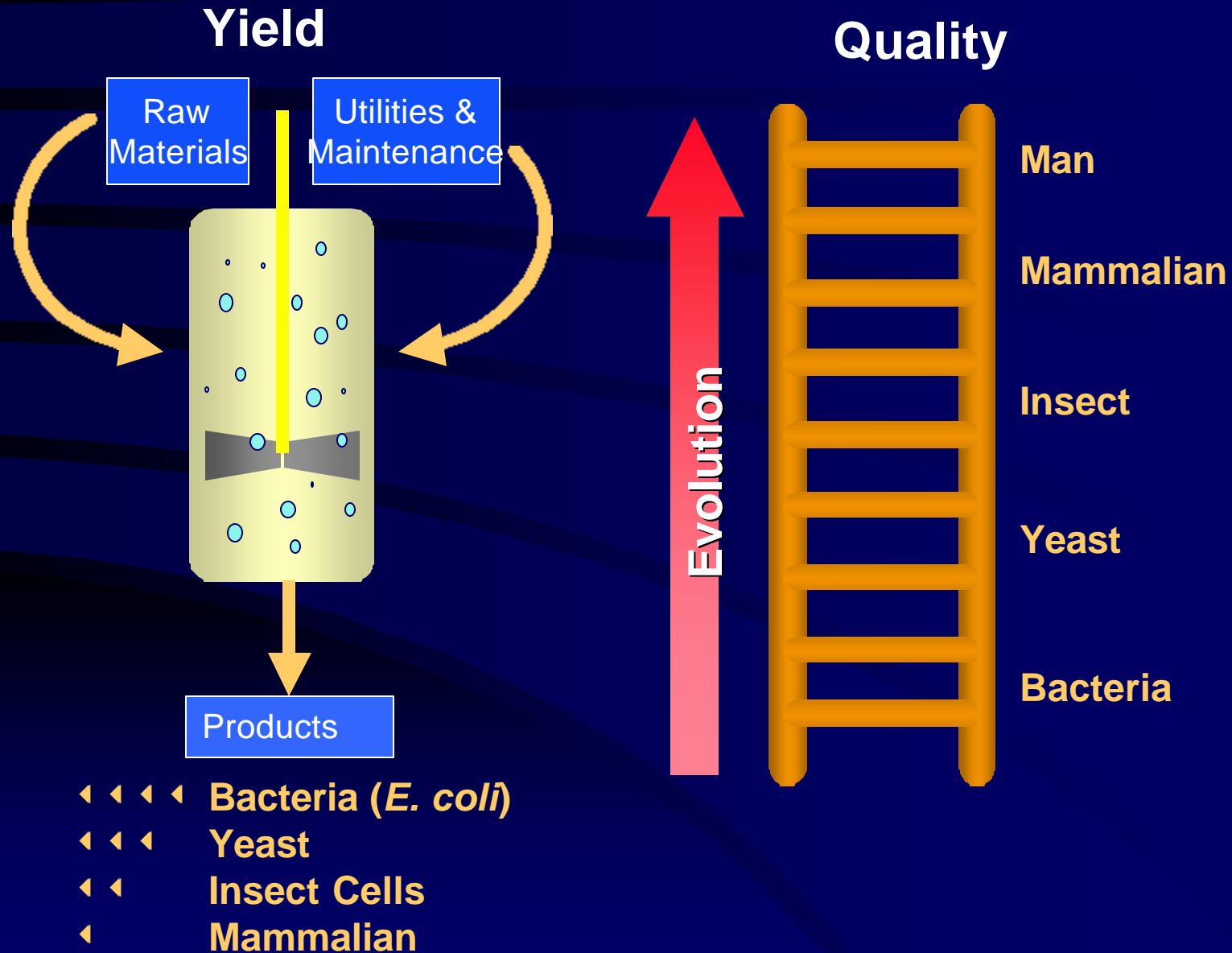


The Living Factory

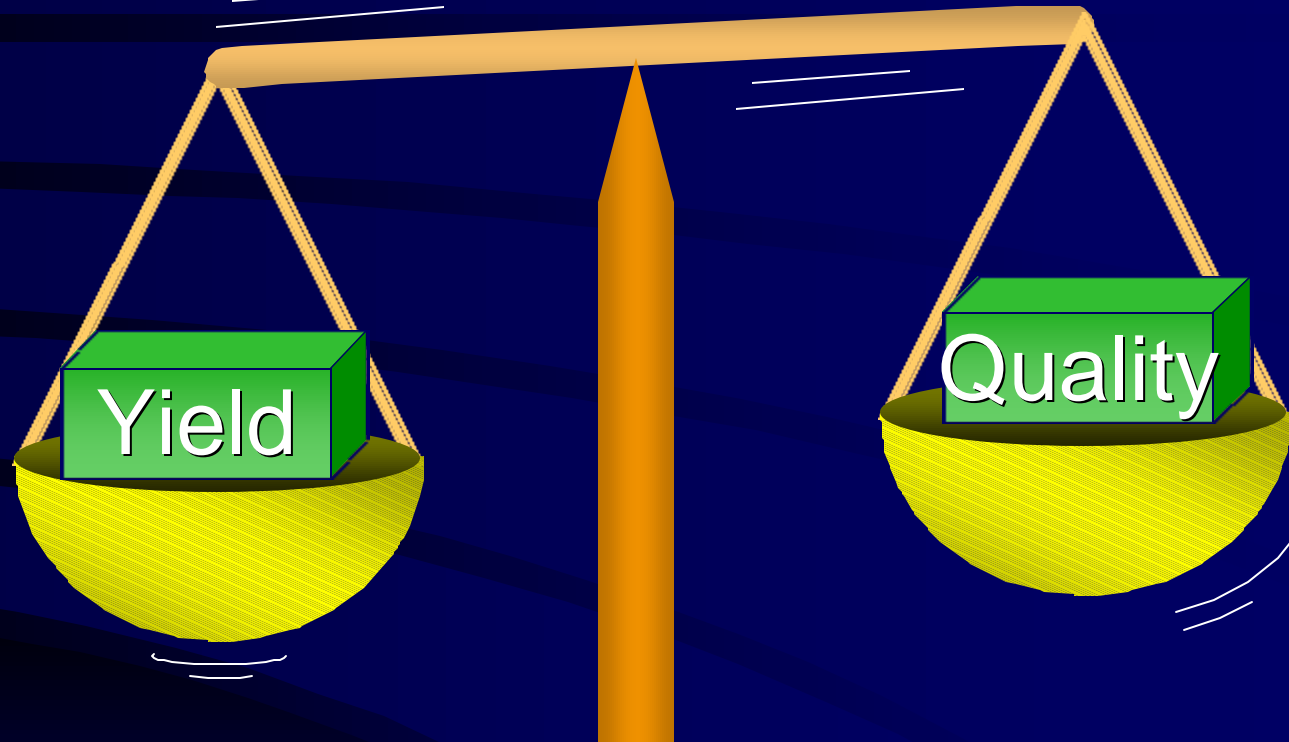
Okay, cells, here's what they're tellin' us to produce!



Choosing a Production System



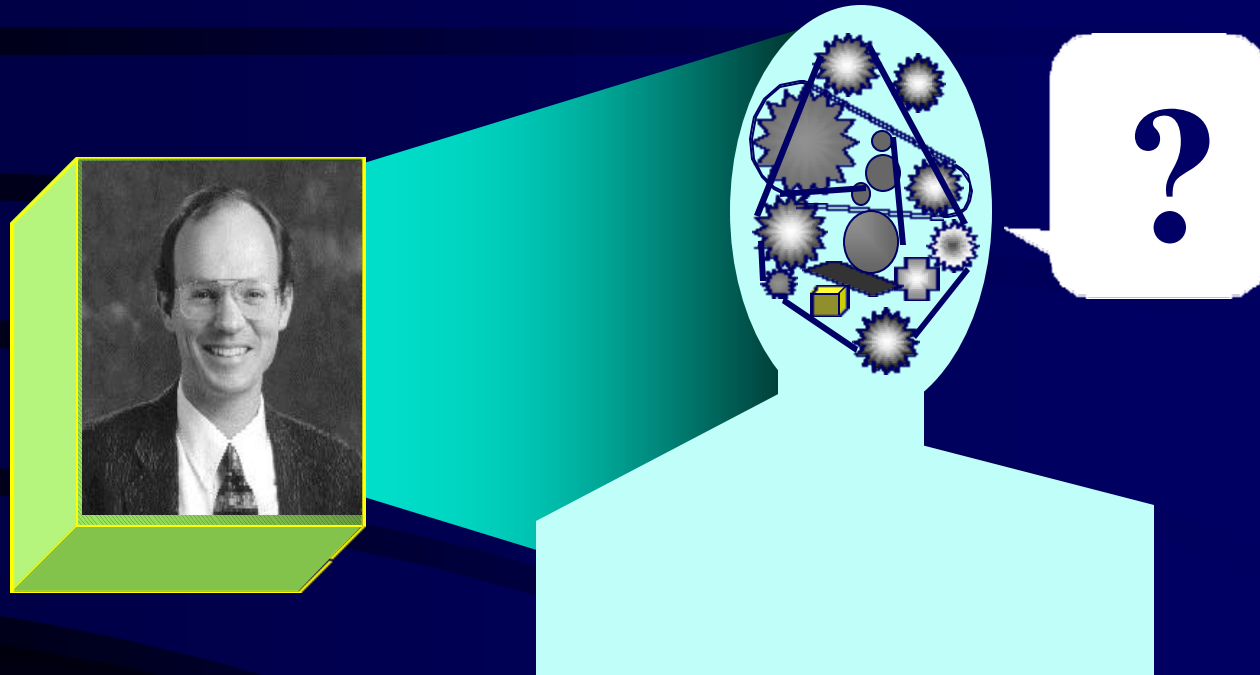
The Engineering Problem



Efficient and inexpensive
production scheme

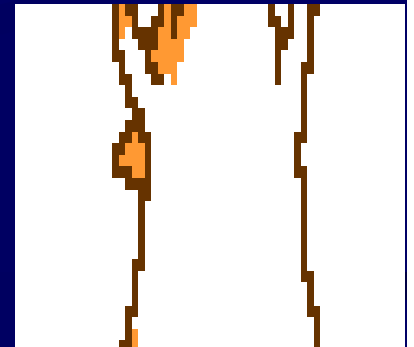
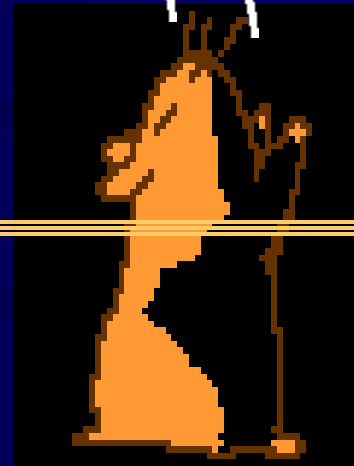
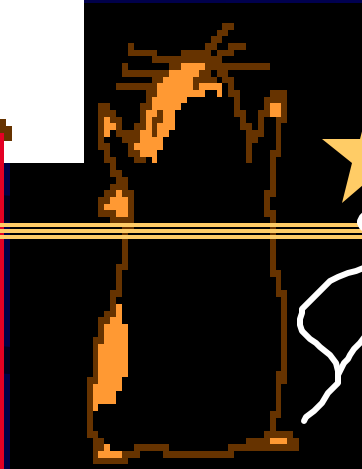
Essential pharmacological
properties

Once upon a time...



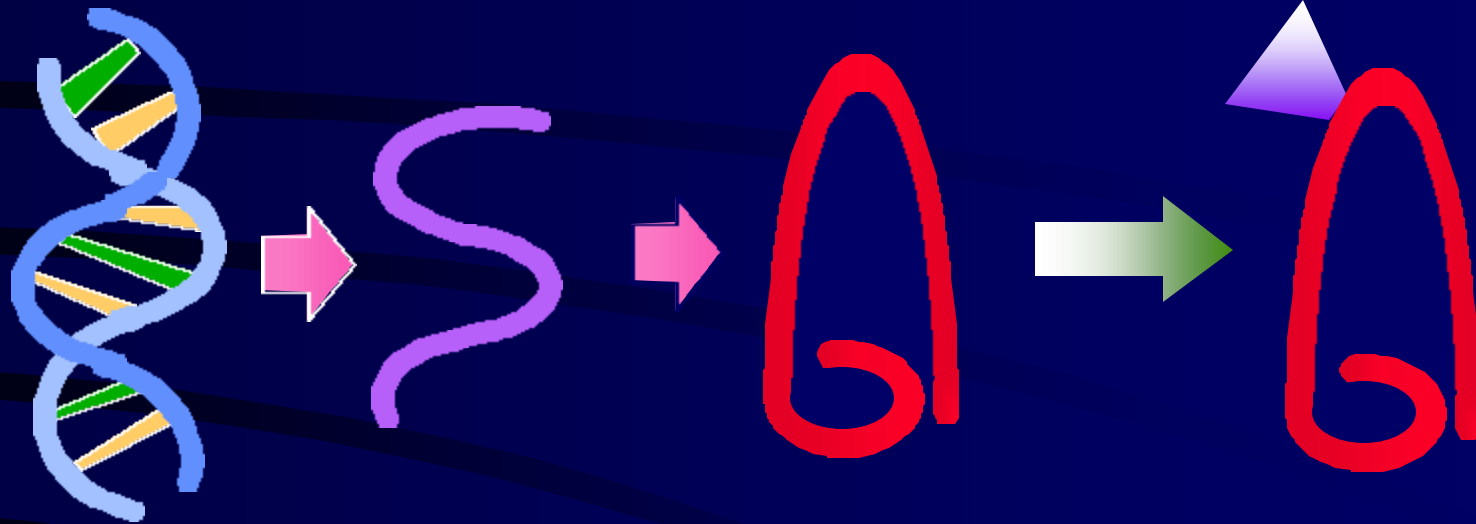
Goal: High yields and high quality

Insect vs. Hamster!



Masters of Production Match

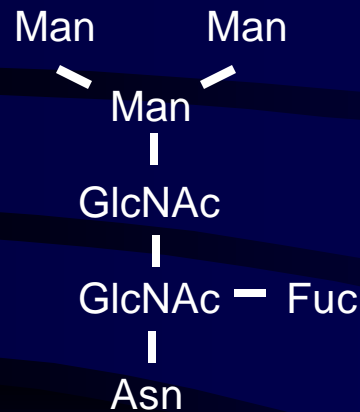
Where do insect cells fall short?



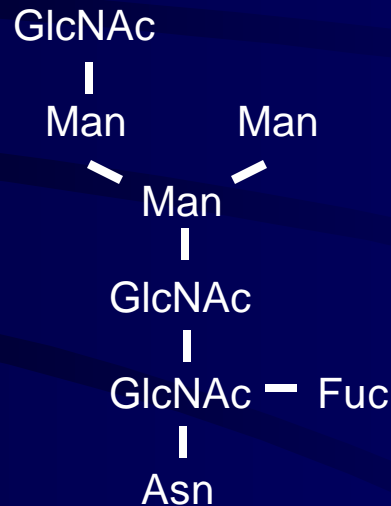
DNA → RNA → Protein → Active Sugar-modified Protein

N-glycan Processing: Insect versus Mammalian

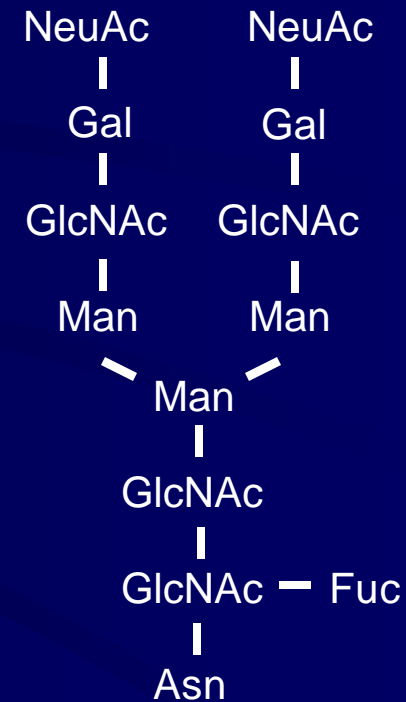
*Paucimannosidic
(Low Mannose)*



Hybrid



Complex



Insect Cells

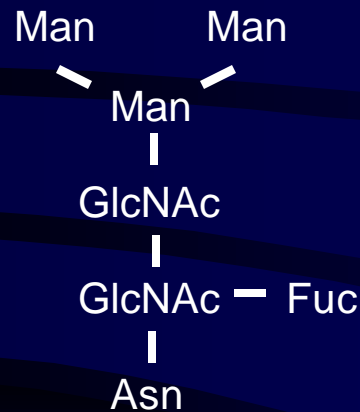
Mammalian Cells

Importance of Sialylation (NeuAc)

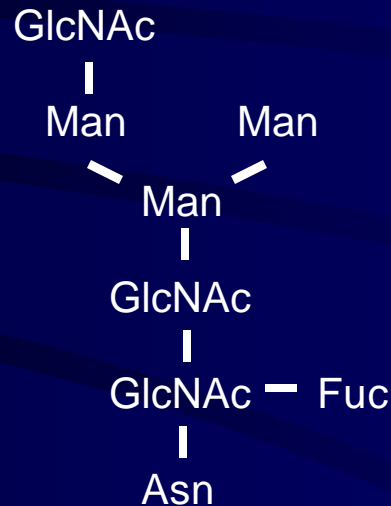
- Protein Structure
- Protein Stability
- Biological Activity
- *In Vivo* Circulatory Half-Life

N-glycan Processing: Insect versus Mammalian

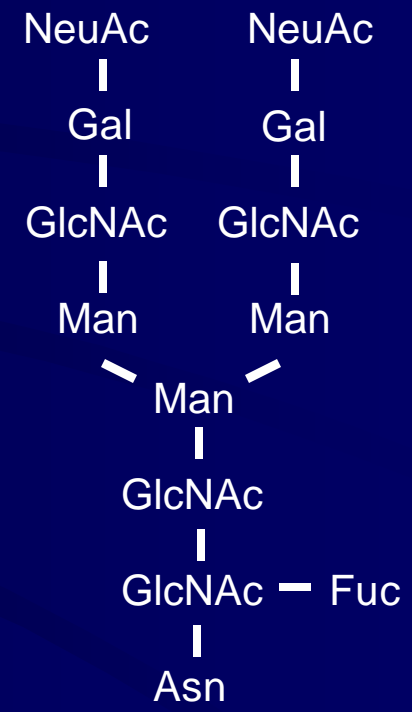
*Paucimannosidic
(Low Mannose)*



Hybrid



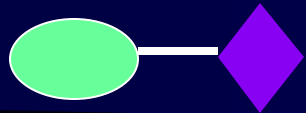
Complex



Insect Cells

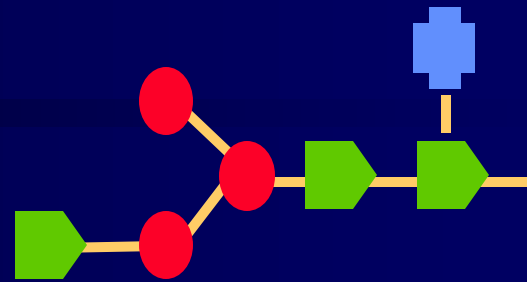
Mammalian Cells

Mammalian Galactosylation Reaction

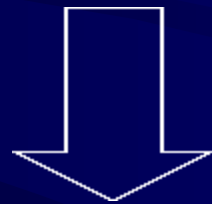


UDP-Galactose

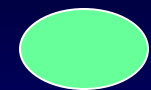
+



N-acetylglucosamine Structure

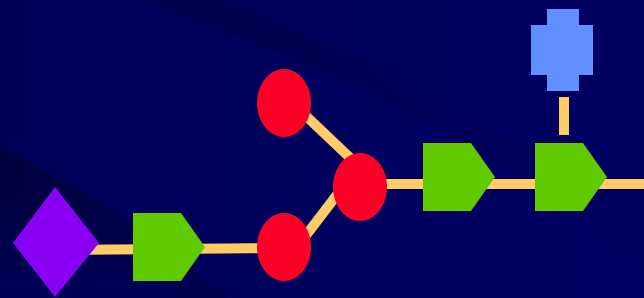


*Mammalian
Galactose transferase*



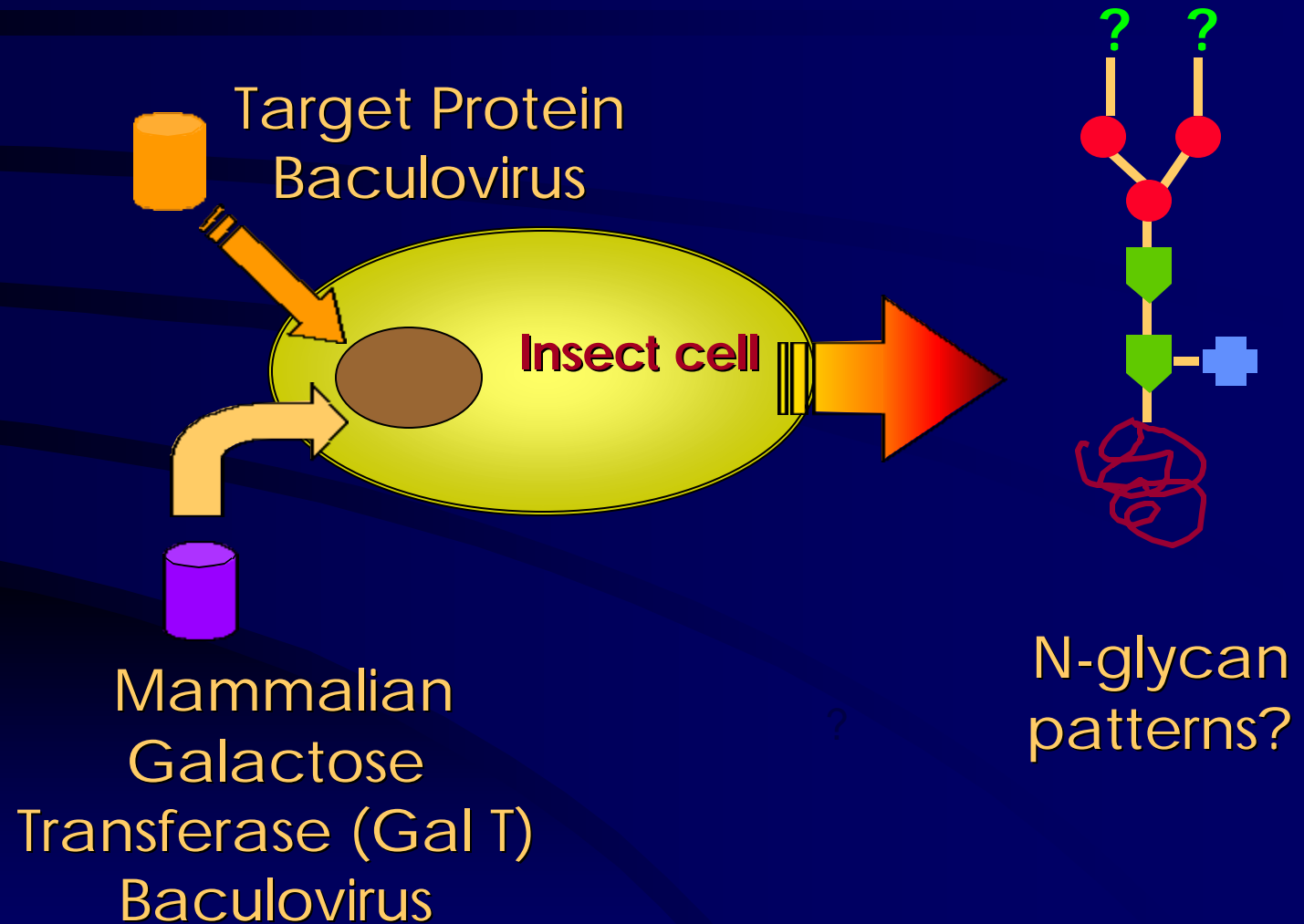
UMP

+

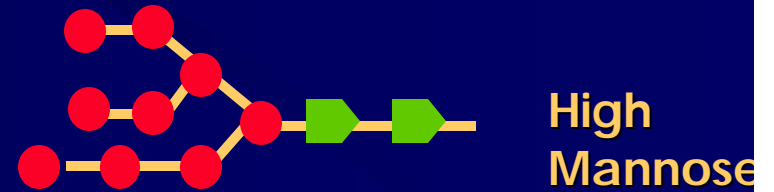
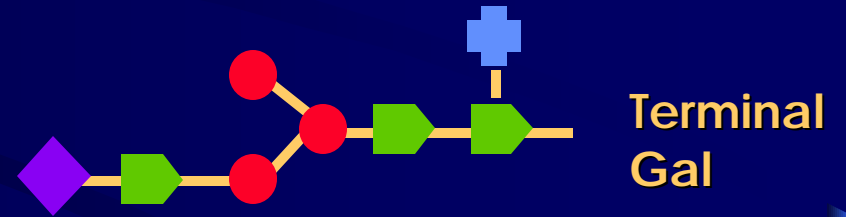
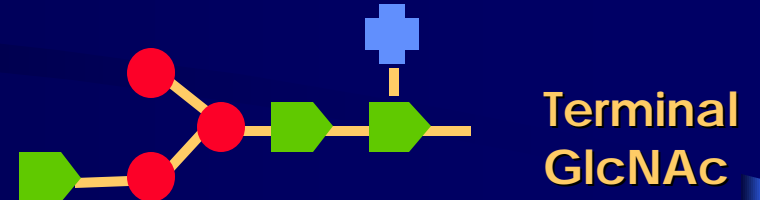
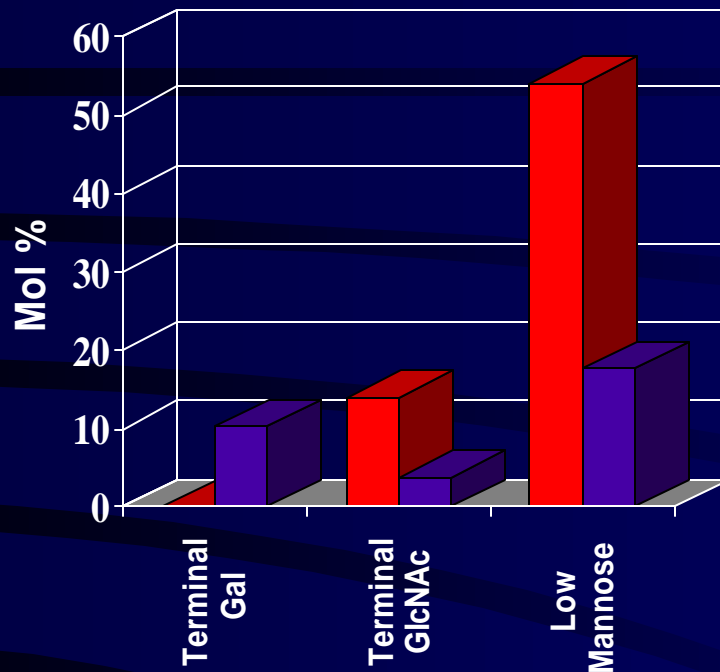


Galactose-terminated Structure ?

Engineering Galactosylation in Insect Cells

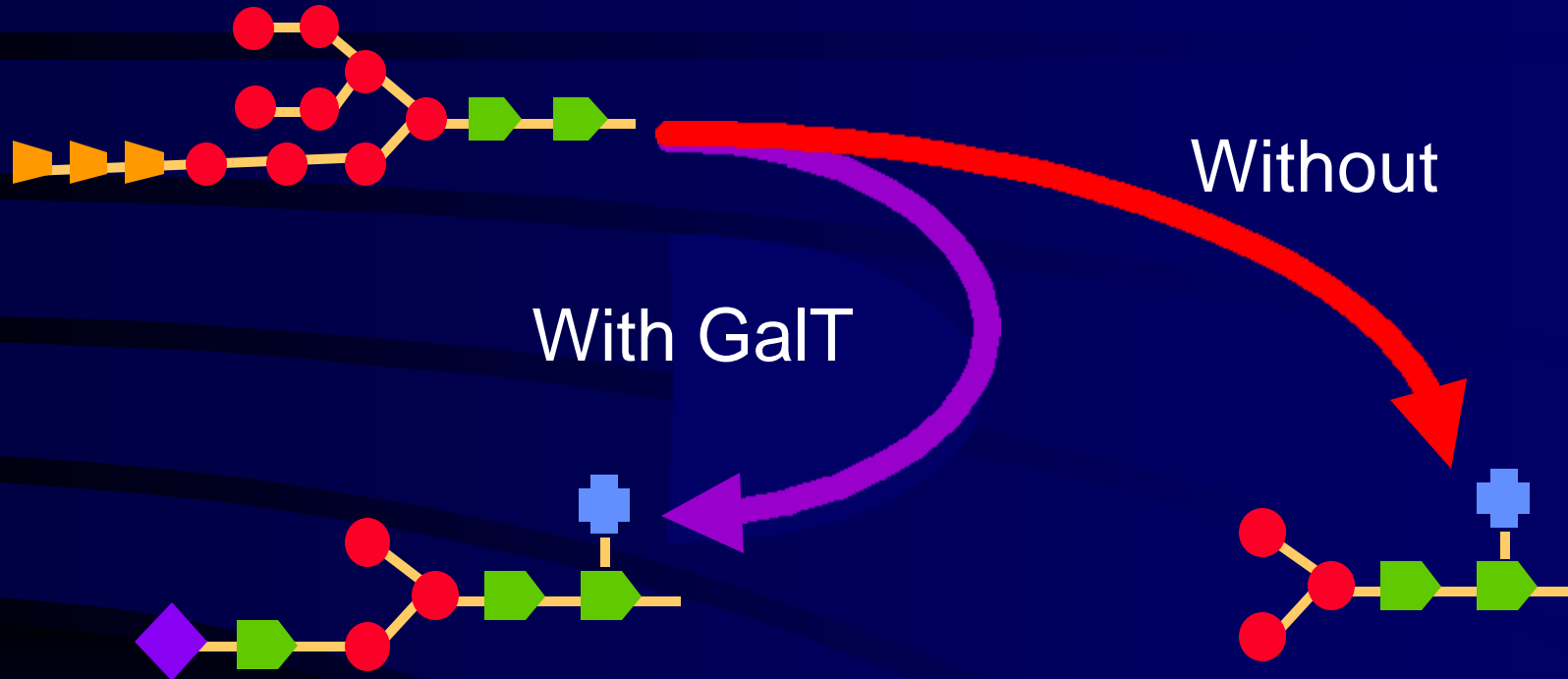


N-Glycan patterns from secreted transferrin



- Transferrin from *T. ni*
- Transferrin (+ GalT) from *T. ni*

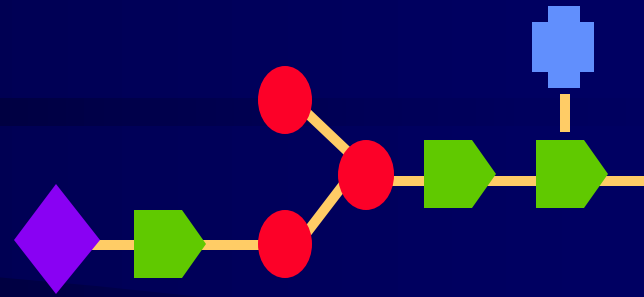
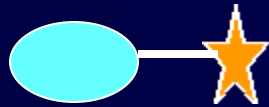
Gal Transferase make N-glycans more “mammalian-like”



1. Adds Terminal Galactose
2. Blocks N-acetylglucosamine Removal

+ = Fucose; ▲ = Glucose; ◆ = Galactose; ▶ =

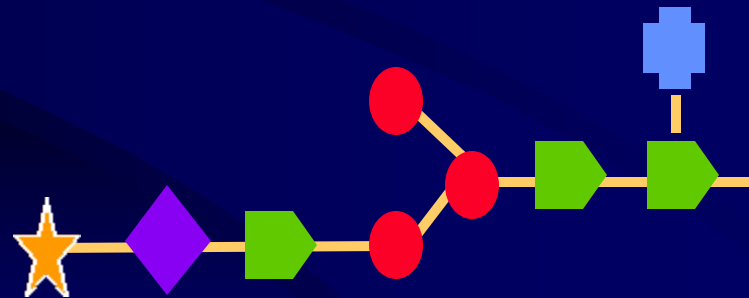
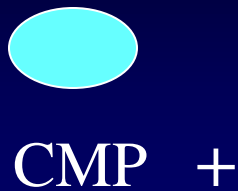
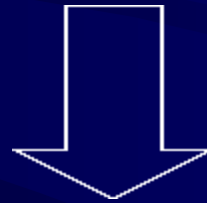
Sialylation Reaction



CMP-NeuAc +
(Cytidine Monophospho-
N-acetylneuraminic acid/Sialic Acid)

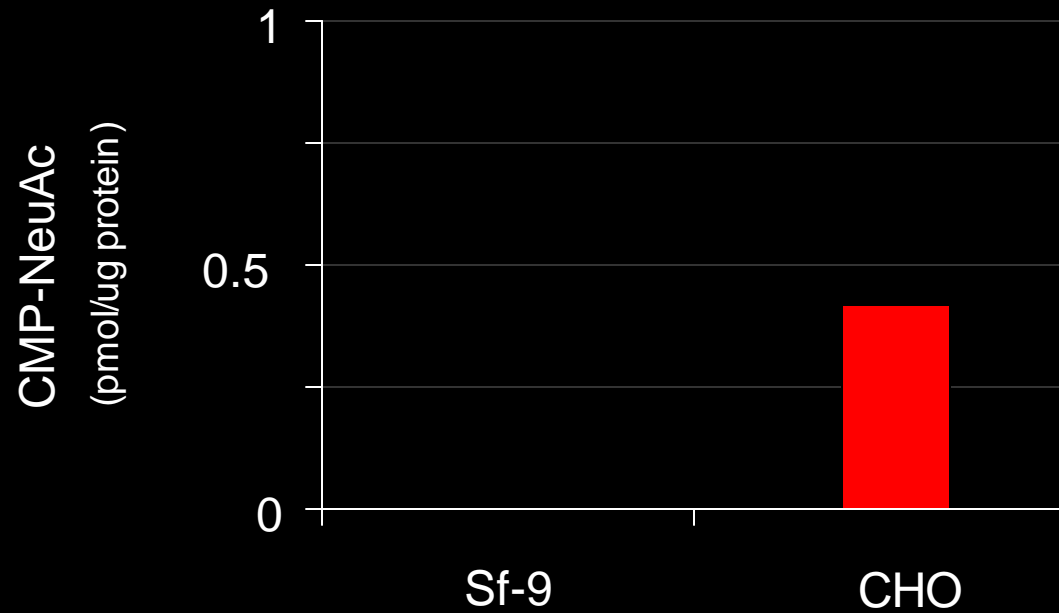
Galactose-terminated Structure

Sialyltransferase

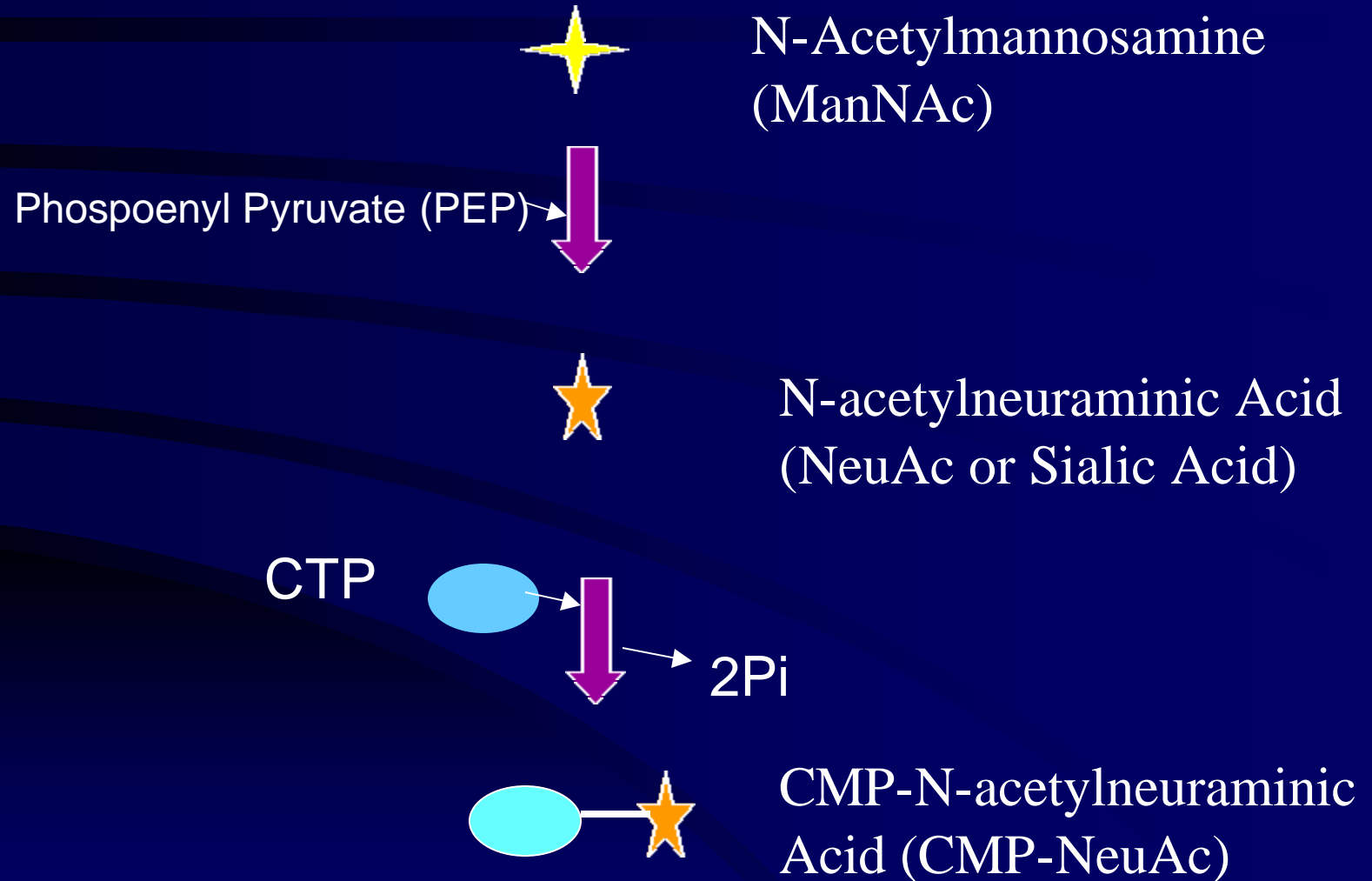


Sialylated Structure
(terminal NeuAc)

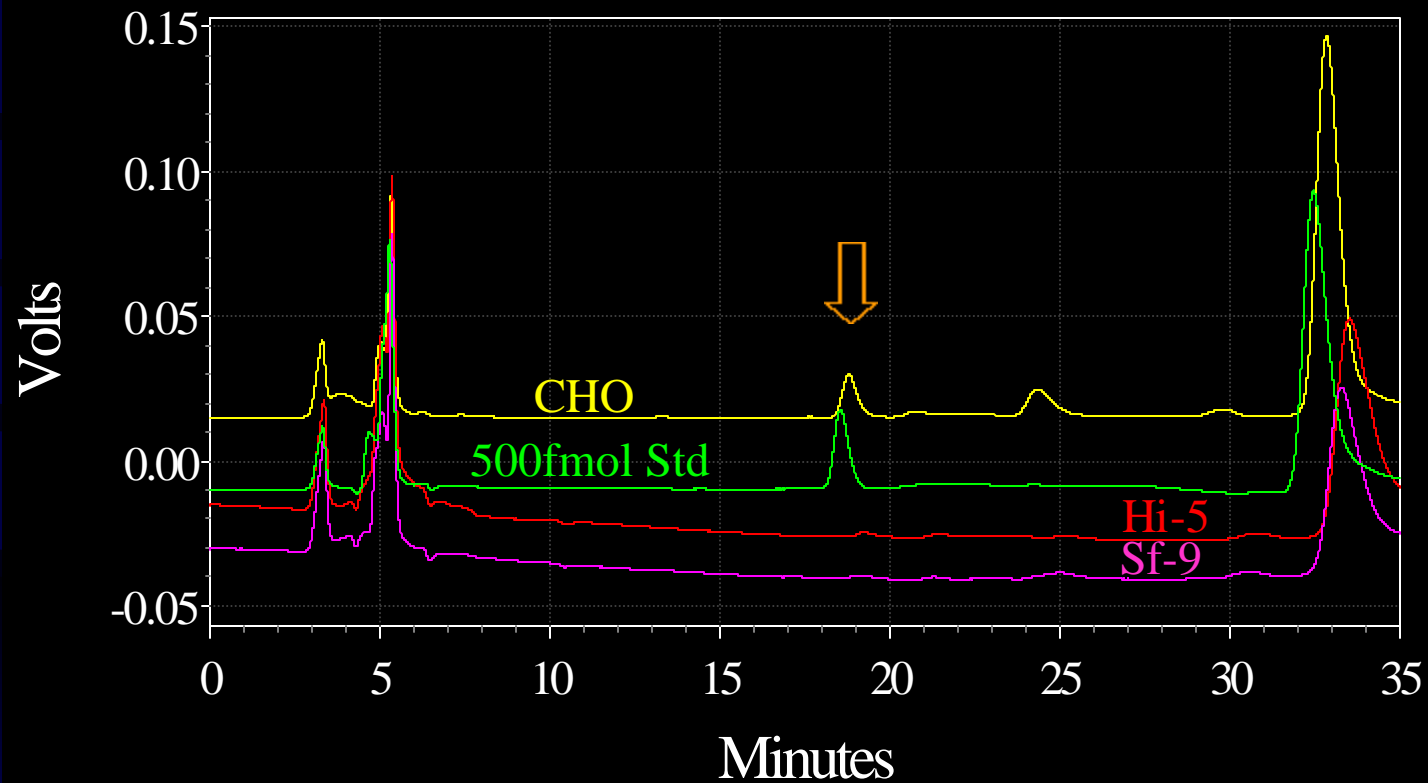
CMP-NeuAc Levels



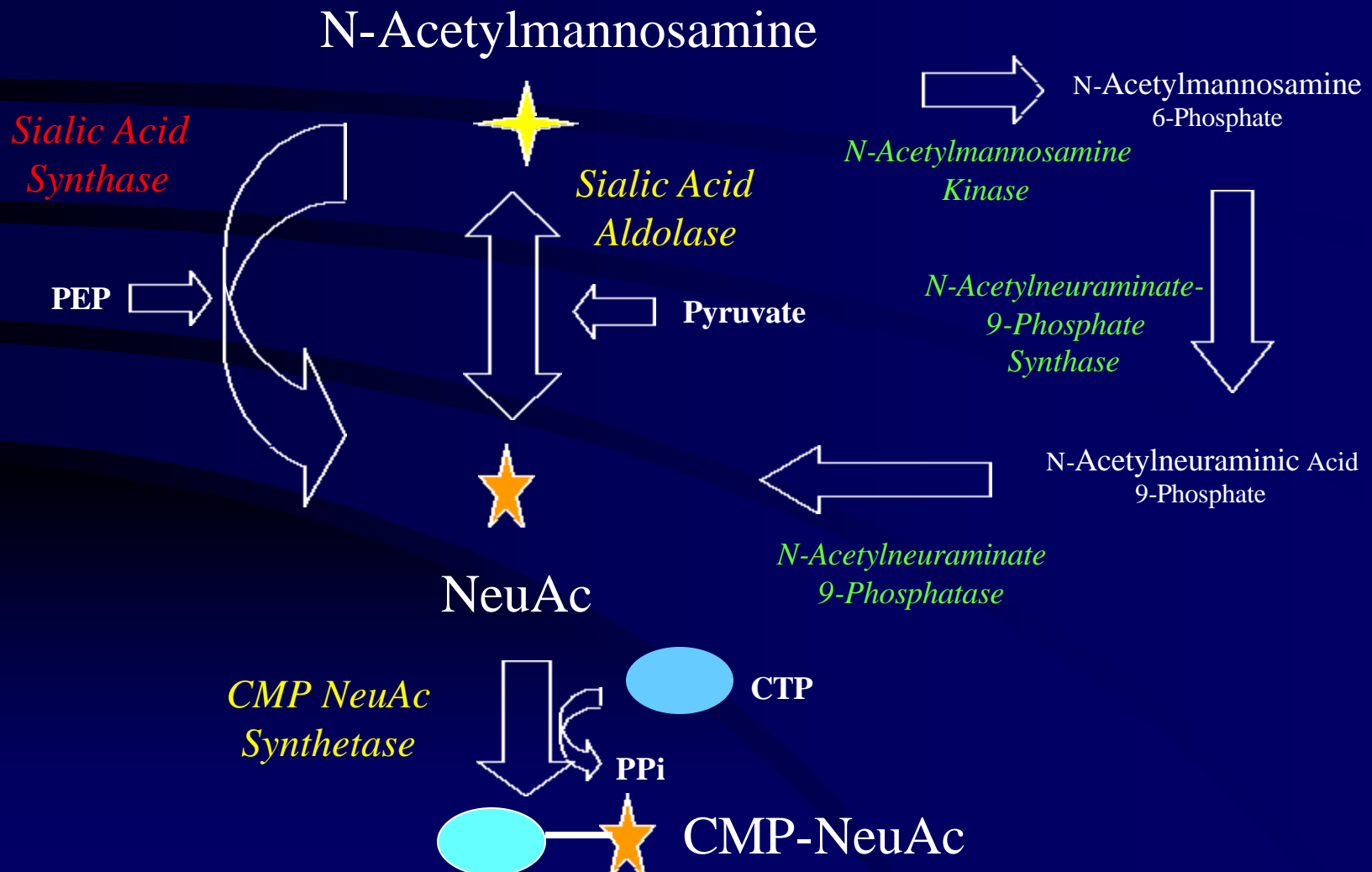
Sialic Acid Pathway



Free NeuAc (Sialic Acid) Levels

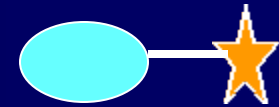


Sialic Acid Pathway



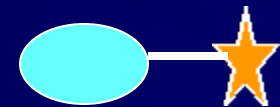
Engineer Sialic Acid Pathway

Insects



DNA

Humans

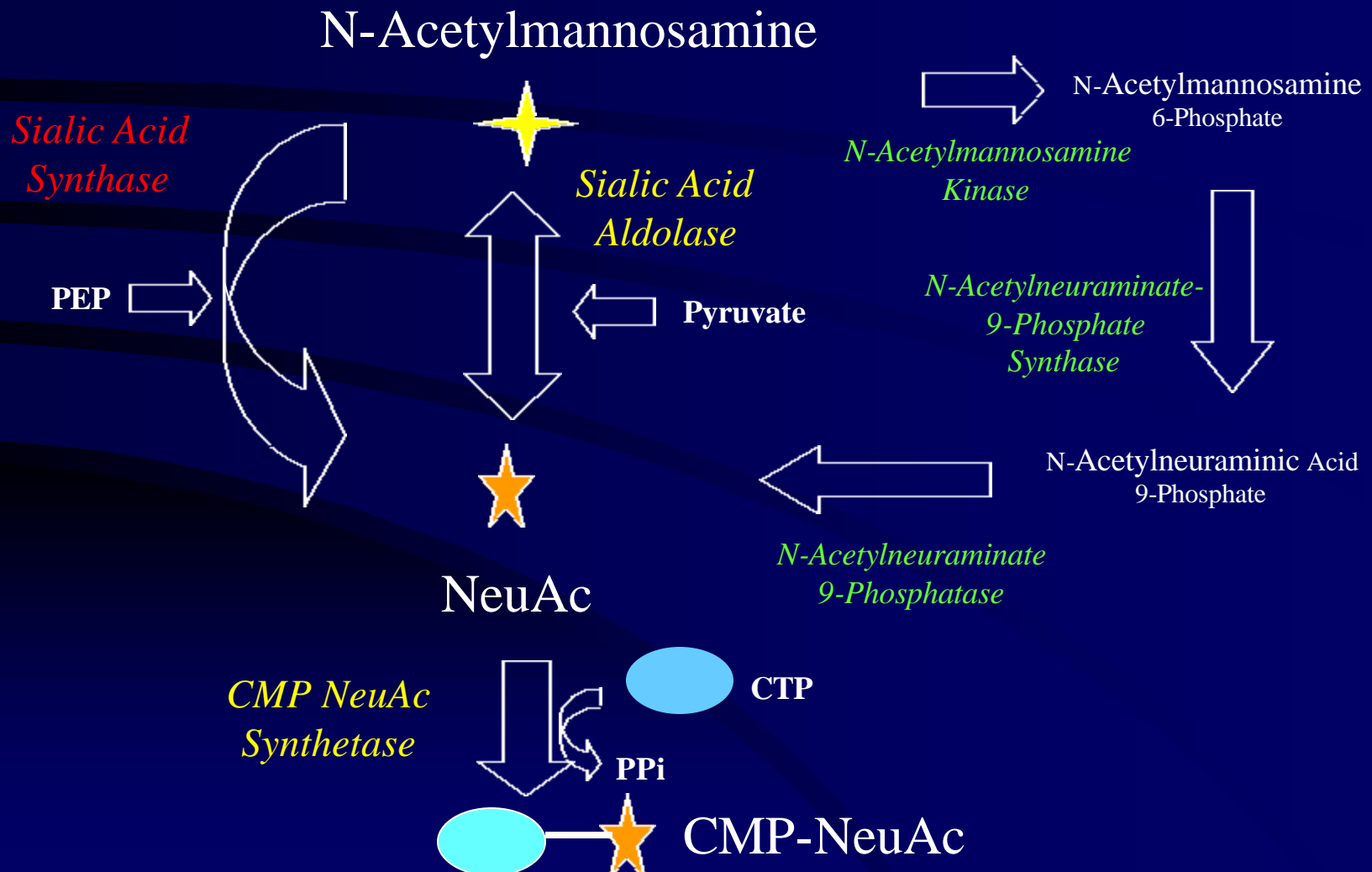


CMP-NeuAc

Bioinformatics Strategy

- Human Gene Identification: Human Genome Sciences, Inc.
 - mRNA → cDNA → EST
 - EST's collected into database
 - EST's assembled into contigs based on homology with themselves
 - Perform homology search using bacterial gene
 - Probe contigs for putative ORF of human gene

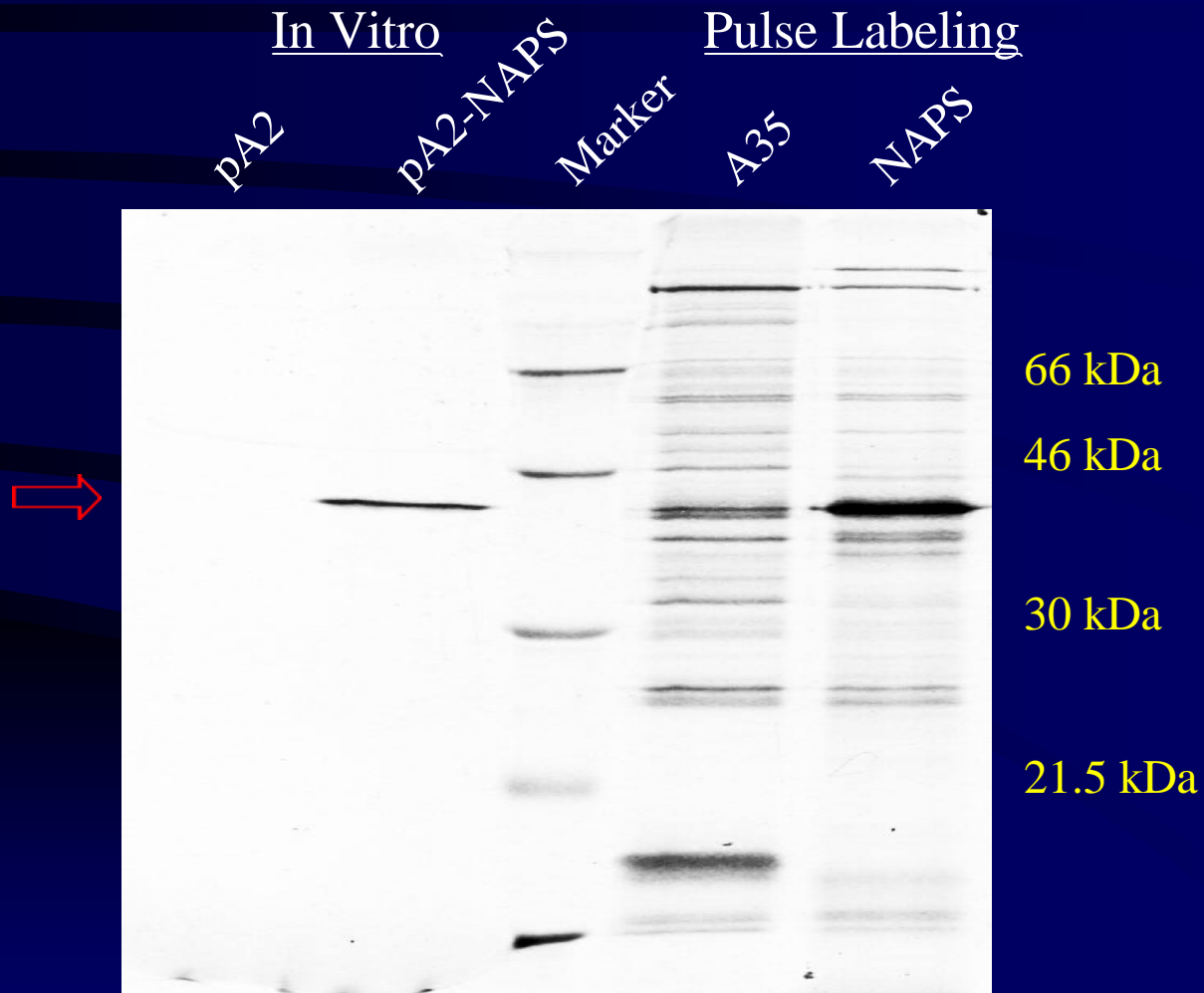
Sialic Acid Pathway



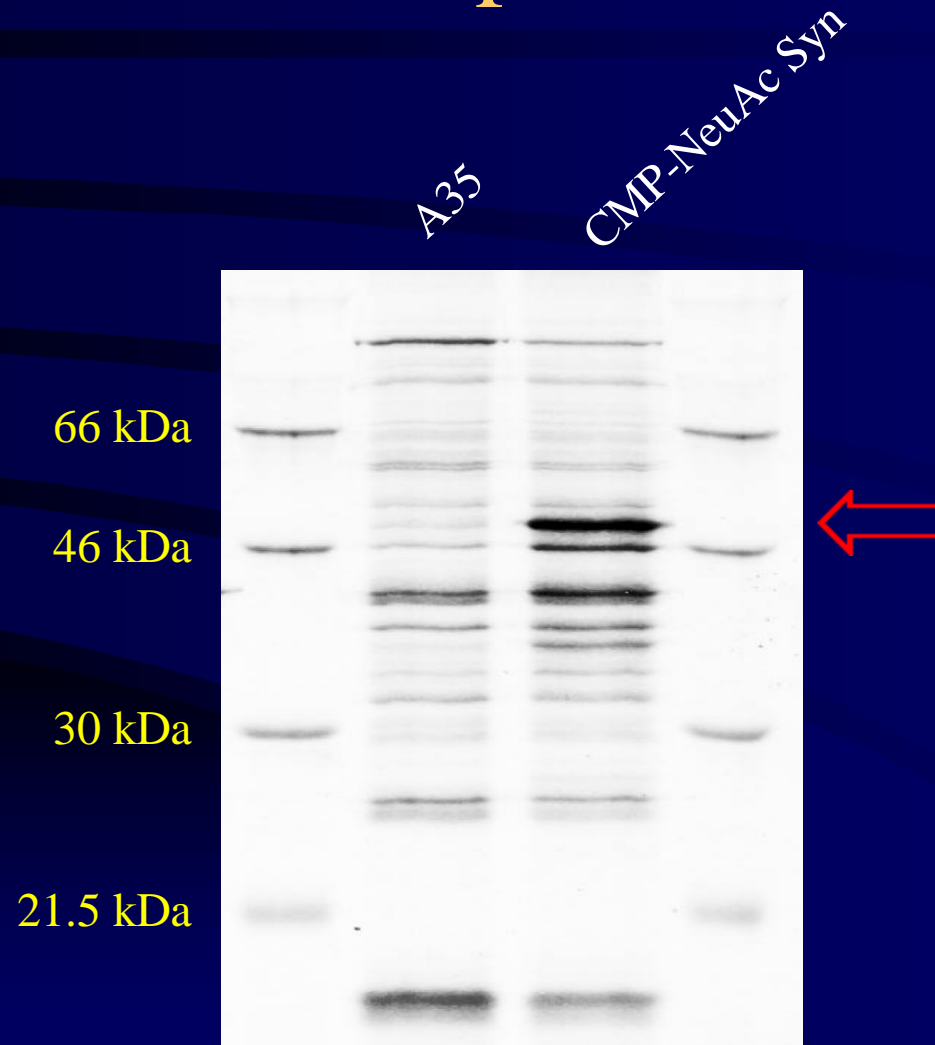
Putative Human Gene Homology

| | Similarity | Identity |
|--------------------------|------------|----------|
| CMP NeuAc Synthase | | |
| <i>E. coli</i> | 25% | 19% |
| <i>M. musculus</i> | 91.2% | 90.7% |
| NeuAc Phosphate Synthase | | |
| <i>E. coli</i> | 56% | 36% |

Putative NeuAc Phosphate Synthase (NAPS) Expression in Insect Cells



Putative CMP-NeuAc Synthase Expression



N-Terminal Protein Sequencing

NeuAc Phosphate Synthase

M**PLELE**LCP

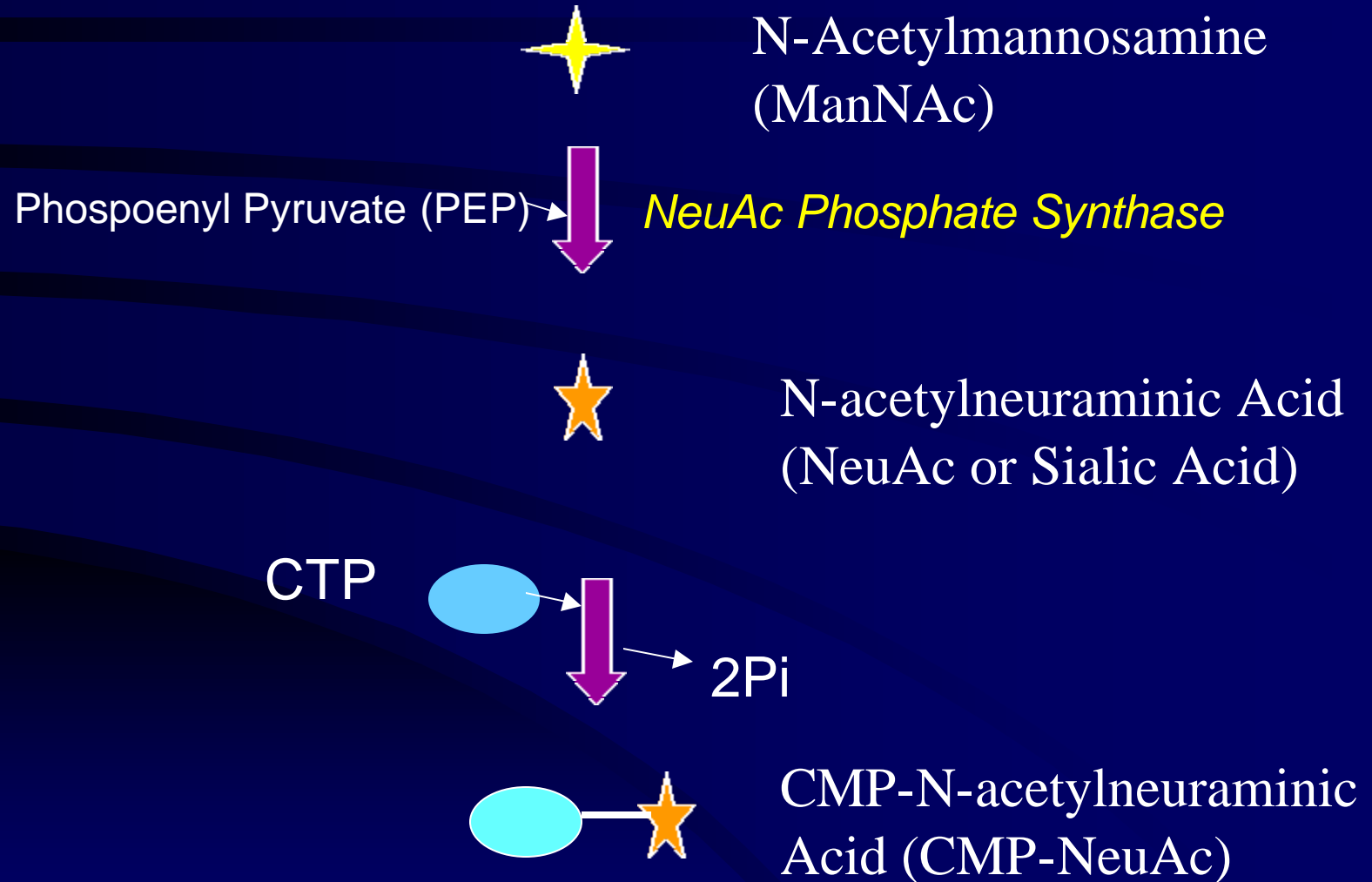
1 . . . 5

CMP-NeuAc Synthase

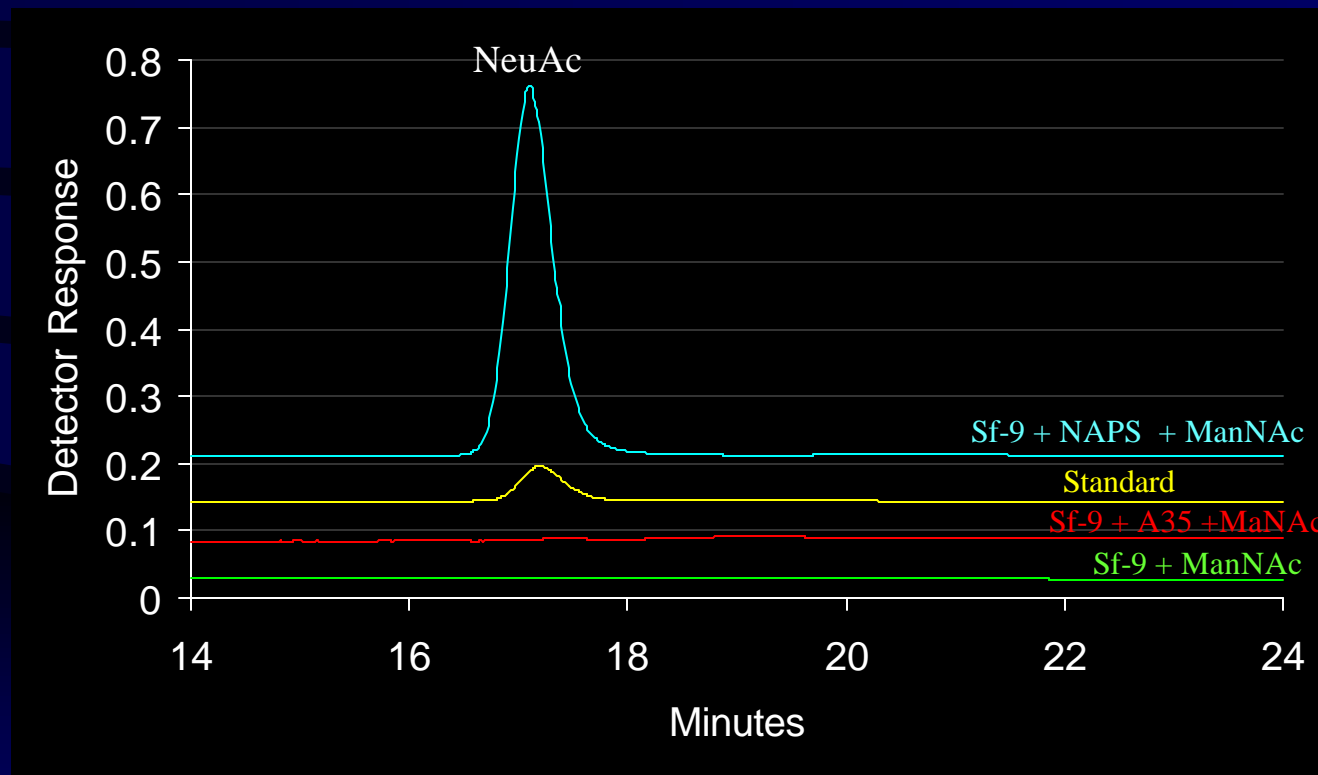
MDS . . . LQ**RNSRG**GQG

1 . . . 27 . . 30 35 .

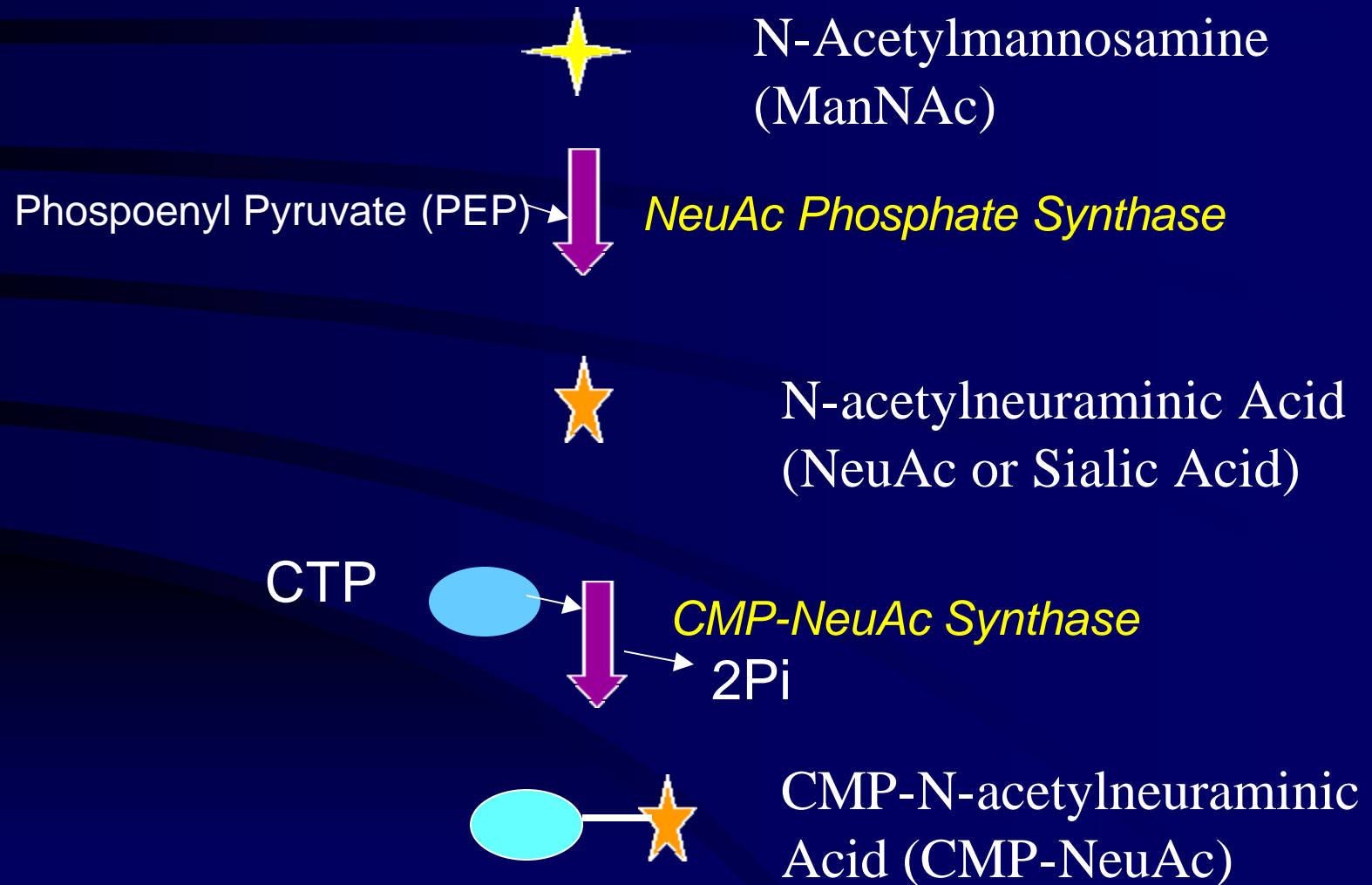
Sialic Acid Pathway



NeuAc Phosphate Synthase Infection with ManNAc Feeding



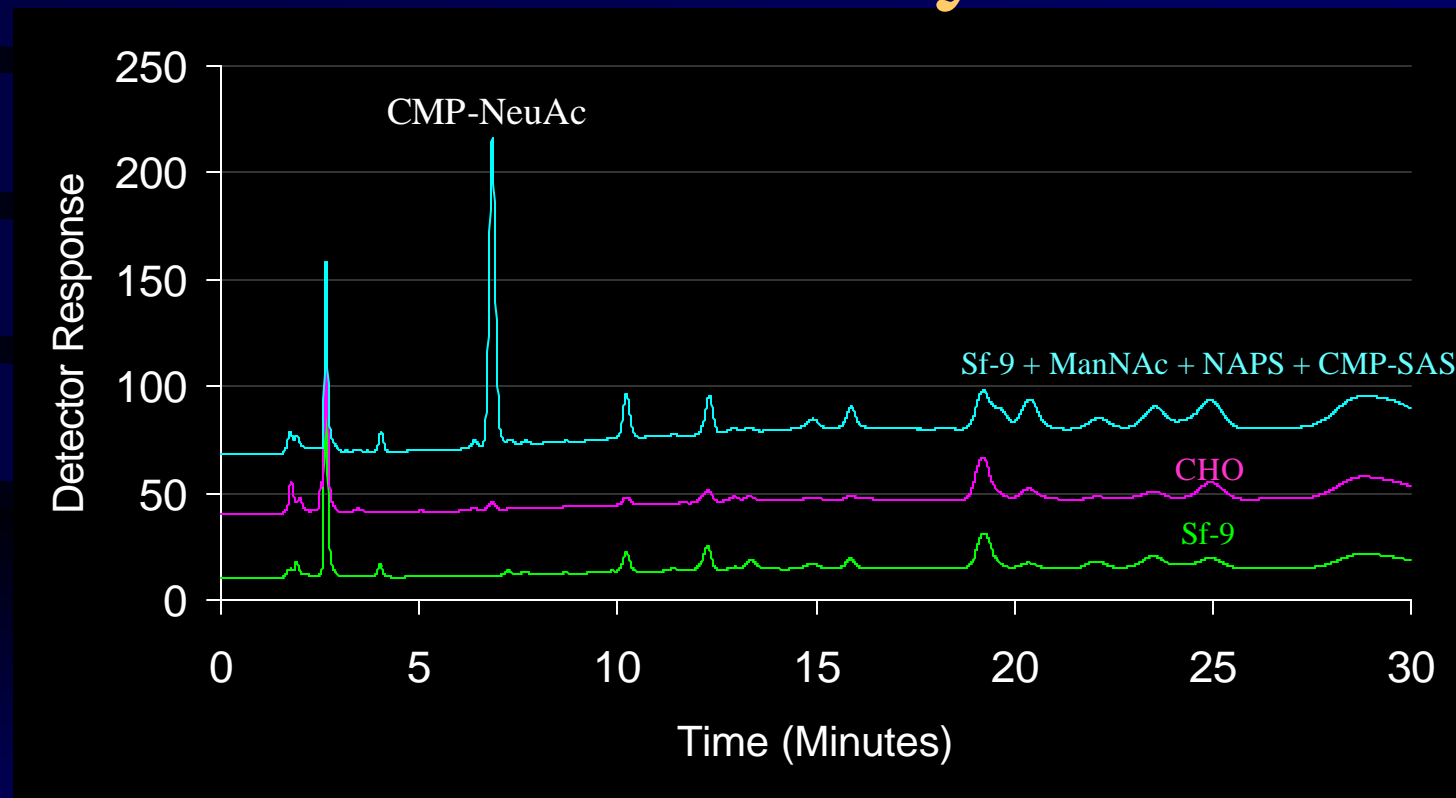
Sialic Acid Pathway



Coinfection: NeuAc-P-Synthase

+

CMP-NeuAc Synthase



CMP-Neu5Ac Levels

| Cell Line | Culture Conditions | CMP-Neu5Ac (pmol/ μ g protein) |
|-----------|--------------------------------|------------------------------------|
| Sf-9 | Uninfected | 0 |
| | Uninfected + ManNAc | 0 |
| | + A35 | 0 |
| | + NAPS + CMP-NeuAc S. | 0 |
| | + NAPS + CMP-NeuAc S. + ManNAc | 10.3 |
| CHO | Unsupplemented | 0.3 |
| | + ManNAc | 1.8 |

From Pest to Princess

