# *nif* genes functions and regulation of expression

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## nif genes

- The two main classes of nitrogen-fixing genes, the *nif* genes and *fix* genes, are present in rhizobia.
- The *nif* genes encode nitrogenase and show structural and functional resemblance with the nitrogen-fixing genes present in *Klebsiella pneumoniae* and other microbial groups.
- Although most of the *nif* genes are found on plasmids of rhizobia, it was also reported to be found on chromosomes of *Bradyrhizobium*.
- *Klebsiella pneumoniae* is the best studied nitrogen-fixing species because of its similarity to *E. coli*.
- *Klebsiella pneumoniae* is a free-living facultative anaerobic nitrogen-fixing bacterium. It contains a total of 20 *nif* genes located on the chromosome in a 24-Kb region. *nifH*, *nifD*, and *nifK* encode the nitrogenase subunits, while *nifE*, *nifN*, *nifU*, *nifS*, *nifV*, *nifW*, *nifX*, *nifB*, and *nifQ* encode proteins involved the assembly and incorporation of <u>iron</u> and <u>molybdenum</u> atoms into the nitrogenase subunits. *nifF* and *nifJ* encode proteins related to electron transfer taking place in the reduction process and *nifA* and *nifL* are regulatory proteins in charge of regulating the expression of the other *nif* genes.

#### • The nonsymbiotic *K. pneumoniae* carries at least 20 *nif* genes which are organized in about eight operons.

| Nif Genes | Role in Nitrogen Fixation  |
|-----------|--|
| nifH      | Dinitrogenase reductase  |
| nifD      | $\alpha$ -Subunit of dinitrogenase   |
| nifK      | B-subunits of dinitrogenase. B clusters are present at B subunit-interface                       |
| nifY      | In Klebsiella pneumoniae, aids in the insertion of FeMo-co into apodinitrogenase                 |
| nifE      | Forms a2 B2 tetramer with nifN. Required for FeMo-co synthesis                                   |
| nifN      | Required for FeMo-co synthesis   |
| nifx      | Involved in FeMo-co synthesis  |
| nifU      | Involved in mobilization of Fe-S cluster synthesis and repair                                    |
| nifS      | Involved in mobilization of S for Fe-S cluster synthesis and repair                              |
| nifV      | Homocitrate synthesis involved in FeMo-co synthesis  |
| nifW      | Involved in stability of dinitrogenase. Proposed to protect dinitrogenase from O2 inactivation   |
| nifM      | Required for the maturation of <i>nifH</i>   |
| nifF      | Flavodoxin, physiologic electron donor to nifH   |
| nifL      | Negative regulatory element  |
| nifA      | Positive regulatory element  |
| nifB      | Required FeMo-co synthesis. Metabolic product. NifB-co is the specific Fe and S donor to FeMo-co |
| fdxN      | Ferredoxin serves as electron donor to nitrogenase   |
| nifQ      | Involved in FeMo-co synthesis. Proposed to function in early MoO42 processing                    |
| nifJ      | Pyruvate flavodoxin (ferredoxin) oxidoreductase involved in electron transport to nitrogenase    |

### nif genes expression

- Gene expression in *nif* operons is dependent on the RNA polymerase σ54 factor and on the NifA transcriptional activator. The σ54 subunit (also known as RpoN, σN, or NtrA product of *ntrA* gene) recognizes a promoter-specific sequence located at positions -24 to -12.
- In nif operons, transcription initiation by the σ54dependent RNA polymerase requires that the enhancer-binding-protein (EBP) and NifA binds to DNA regions known as upstream activator sequences.

### Regulation of *nif* genes expression

- Regulation of *nif* gene expression has two elements:
  - An external system designated *ntr* (nitrogen regulatory)
  - An internal system mediated by *nif A* and *nif L* gene products
- In most bacteria, regulation of *nif* genes transcription is done by the nitrogen sensitive NifA protein.
- When there isn't enough fixed nitrogen available for the organism's use, NtrB the product of *ntrB* gene (function as protein kinase) phosphoylates NtrC. NtrC-P triggers NifA expression, and NifA activates the rest of the *nif* genes.
- If there is a sufficient amount of reduced nitrogen or oxygen is present, another protein NifL is activated.
- NifL inhibits NifA activity resulting in the inhibition of nitrogenase formation.
- NifL is regulated by the products of *glnD* and *glnK*.

#### **REGULATION OF NITROGEN FIXATION**



# Regulation of nif gene-

Where P, P1 and P2 = Promoter region of operon



#### Questions

- Write in detail about nif genes and regulation of their expression.
- Write a short note on nif genes functions.
- Write a short note on regulation of nif genes expression.