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TITLE: Superoxide Dismutase and Transcription Factor SOX9 as Mediators of Tumor Suppression by MAC25 (IGFBP-RP1) in Prostate Cancer Cells

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| <b>13. ABSTRACT (Maximum 200 Words)</b><br>The hypothesis of this proposal is that: The mac25/IGFBP-rp1 gene functions as a tumor suppressor for prostate epithelium by induction of additional genes regulating response to oxidative stress, senescence, and differentiation. One of these, SOD2, impairs tumor cell growth by detoxifying superoxides which may initiate transformation, or by interaction with growth promoting signal pathways via modulation of protein phosphorylation. Another, SOX9, induces a subset of genes required for differentiation of normal cells with reduced rates of proliferation and higher susceptibility to apoptosis. We propose to adress this hypothesis by the following four Specific aims: 1) Determine if either SOD2 or SOX9 is necessary and/or sufficient for effective tumor suppression by mac25; 2) Analyze the effects of both SOD2 and SOX9 on specific proteins and activities of cell cycle and apoptotic pathways known to be regulated by mac25; 3) Analyze the interaction between mac25 and androgen-responsive pathways. Production of specific proteins associated with differentiation, including cytokeratins, cadherins, CD44 and CD57, and PSA, will also be determined; 4) Assess the extent to which either SOD2 or SOX9 can account for the gene expression pattern associated with tumor suppression by mac25. Herein, we report on studies that address specific aims 2 and 3. |   |  |  |
| <b>14. SUBJECT TERMS</b><br>SOX9; SOD-2; Cell Signaling; Apoptosis; Cell Cycle Proteins; Androgens; Androgen Receptor; Transcription Factor; Antisense RNA   |   |  | <b>15. NUMBER OF PAGES</b><br>8                |
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**Introduction:** We have previously determined that the senescence-associated gene mac25 is a tumor suppressor for prostate cancer. We have also determined, using cDNA microarrays that manganese superoxide dismutase (SOD-2) and the transcription factor SOX-9, are two genes that have their expression increased by mac35 in prostate epithelial cells. We have shown that the expression of SOD-2 decreases in prostate cancer and re-expression of SOD-2 results in cell senescence and decreased tumor growth. Recent data have shown that a polymorphism in the SOD-2 gene that results in a decrease in function is associated with higher rates of prostate cancer. The purpose of this study was to determine the effects of SOX9 on development of prostate cancer and the interaction with SOD-2. In the first year of this study we reported that :

- Sox-9 and SOD2 expression is decreased in prostate cancer and SOX9 decreases in models of prostate cancer metastasis.
- SOD-2 and SOX9 both contribute to the tumor suppressing action of mac25 but each by its unique mechanism.
- Antisense and sense adenoviral constructs have been developed for SOD-2 and SOX9.
- Confocal microscopy demonstrates nuclear localization of mac25.

In the second year of this study we have concentrated our efforts on Task 2 of the proposal in which we said that we would construct adenoviral vectors for SOD-2 and SOX9 and determine the effects of expression of these genes on downstream proteins and signaling pathways.

**Body:** Approved work statement for year 2

Task 2: Analyze the effects of both SOD2 and SOX9 on specific proteins and activities of cell cycle and apoptotic pathways known to be regulated by mac25.

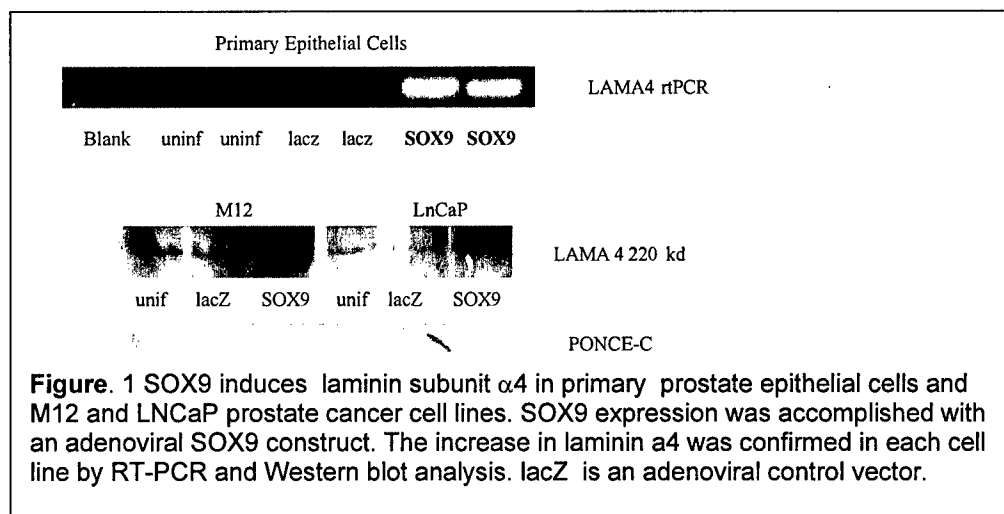
- Determine expression of specified cell cycle proteins, and perform flow cytometry, in M12/SOD2 and M12/SOX9. Repeat in presence of antisense oligos in cases where both SOD2 and SOX9 are upregulated. (Months 12-18)

| UNIGENE   | NAME  | LNcap_SOX <sup>9</sup> | M12_SOX9 | PEC_SOX9 |
|-----------|---|------------------------|----------|----------|
| Hs.283722 | START_DOMAIN_CONTAINING_7_(STARD7)  | 1.89                   | 3.2      | 2.08     |
|           | UNKNOWN_162   | 2.04                   | 1.82     | 1.51     |
|           | HUMAN_DNA_SEQUENCE_FROM_CLONE_RP11-503C24_ON_CHROMOSOME_6_COMPLETE_SEQUENCE | 1.37                   | 1.38     | 1.35     |
| Hs.150477 | WERNER_SYNDROME_(WRN)   | 1.4                    | 1.58     | 1.65     |
| Hs.78672  | LAMININ_ALPHA_4_(LAMA4)   | 1.46                   | 1.57     | 1.83     |
| Hs.183805 | ANKYRIN_1_ERYTHROCYTIC_(ANK1)   | 1.43                   | 1.74     | 1.48     |
| Hs.75659  | MPV17_TRANSGENE_MURINE_HOMOLOG_GLOMERULOSCLEROSIS_(MPV17)                   | NA                     | 1.52     | 1.29     |
|           |   | 1.28                   | 1.42     | 1.37     |

|           |  |      |      |      |
|-----------|--|------|------|------|
| Hs.380843 | RIBOSOMAL_PROTEIN_S6_(RPS6)  | 1.85 | 2.11 | 1.36 |
| Hs.124854 | CHROMOSOME_7_OPEN_READING_FRAME_13_(C7ORF13)                         | 1.42 | 1.91 | 1.47 |
| Hs.5920   | UDP-N-ACETYLGLUCOSAMINE-2-EPIMERASE/N-ACETYLMANNOSAMINE_KINASE_(GNE) | 1.67 | 1.68 | 1.3  |
| Hs.132071 | OVARIAN_CARCINOMA_IMMUNOREACTIVE_ANTIGEN_(OCIA)                      | 1.62 | 1.47 | 1.3  |
| Hs.21411  | HOMO_SAPIENS_CDNA_FLJ30370_FIS_CLONE_BRACE2007832                    | 1.35 | 1.43 | 1.27 |
| Hs.9071   | PROGESTERONE_RECEPTOR_MEMBRANE_COMPONENT_2_(PGRMC2)                  | 1.45 | 1.62 | 1.29 |
| Hs.93868  | KIAA0662_GENE_PRODUCT_(KIAA0662)                                     | 1.37 | 1.35 | 1.24 |
| Hs.87372  | ESTS   | 1.3  | 1.53 | 1.33 |
| Hs.301132 | KIAA0195_GENE_PRODUCT_(KIAA0195)                                     | 1.42 | NA   | 1.5  |
| Hs.349961 | RIBOSOMAL_PROTEIN_L6_(RPL6)  | 1.45 | 1.61 | 1.27 |
| Hs.2007   | TUMOR_NECROSIS_FACTOR_(LIGAND)_SUPERFAMILY_MEMBER_6_(TNFSF6)_MRNA    | 1.39 | 1.26 | 1.53 |

Table 1. cDNA microarray of genes significantly upregulated after infection of the respective cell lines with an adenoviral construct containing the construct for SOX9 or lacZ control. , >0.67 = 2 =fold increase.

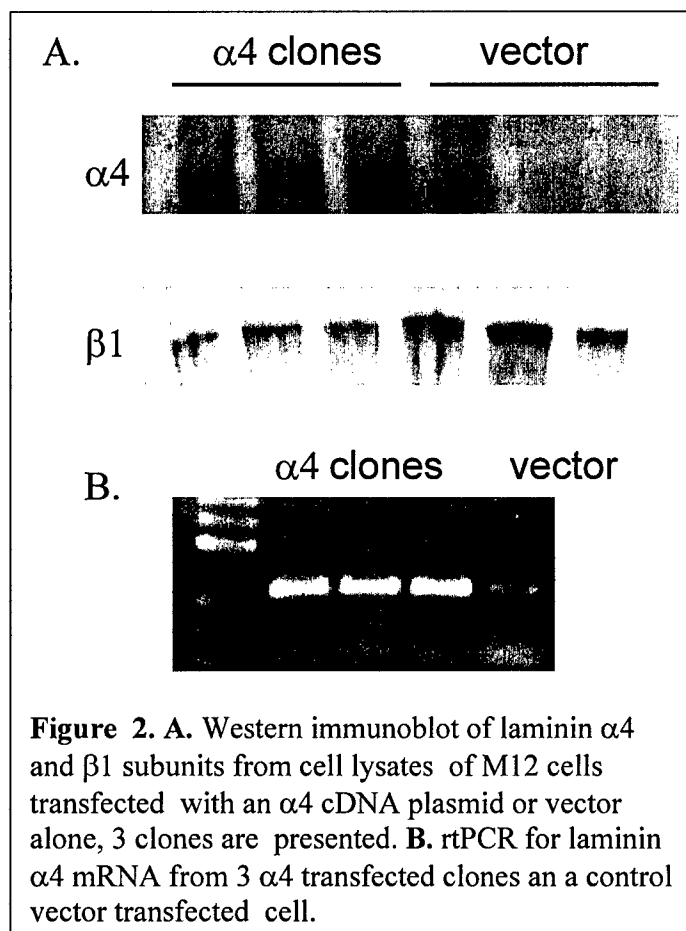
b) Clone SOX2 and SOX9 in adenoviral vector, optimize conditions, and test production. (Months 12-18) Figure 1

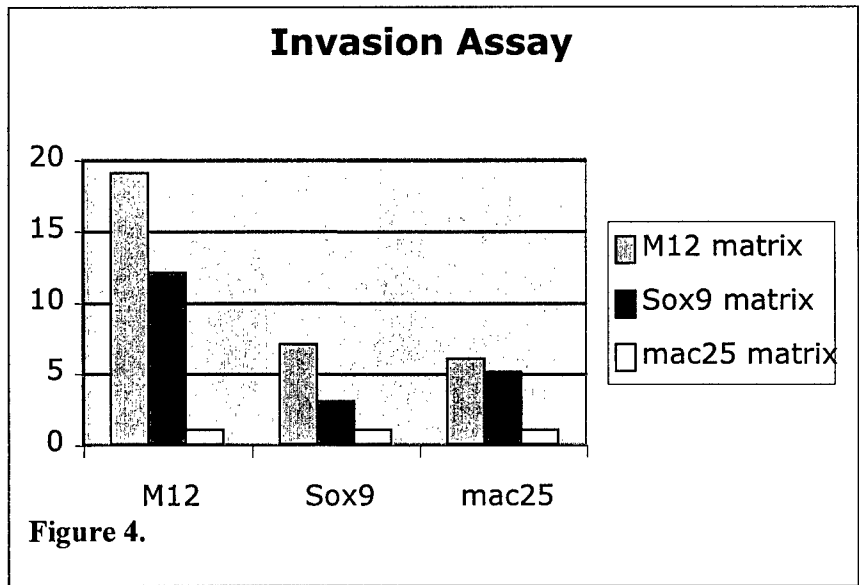
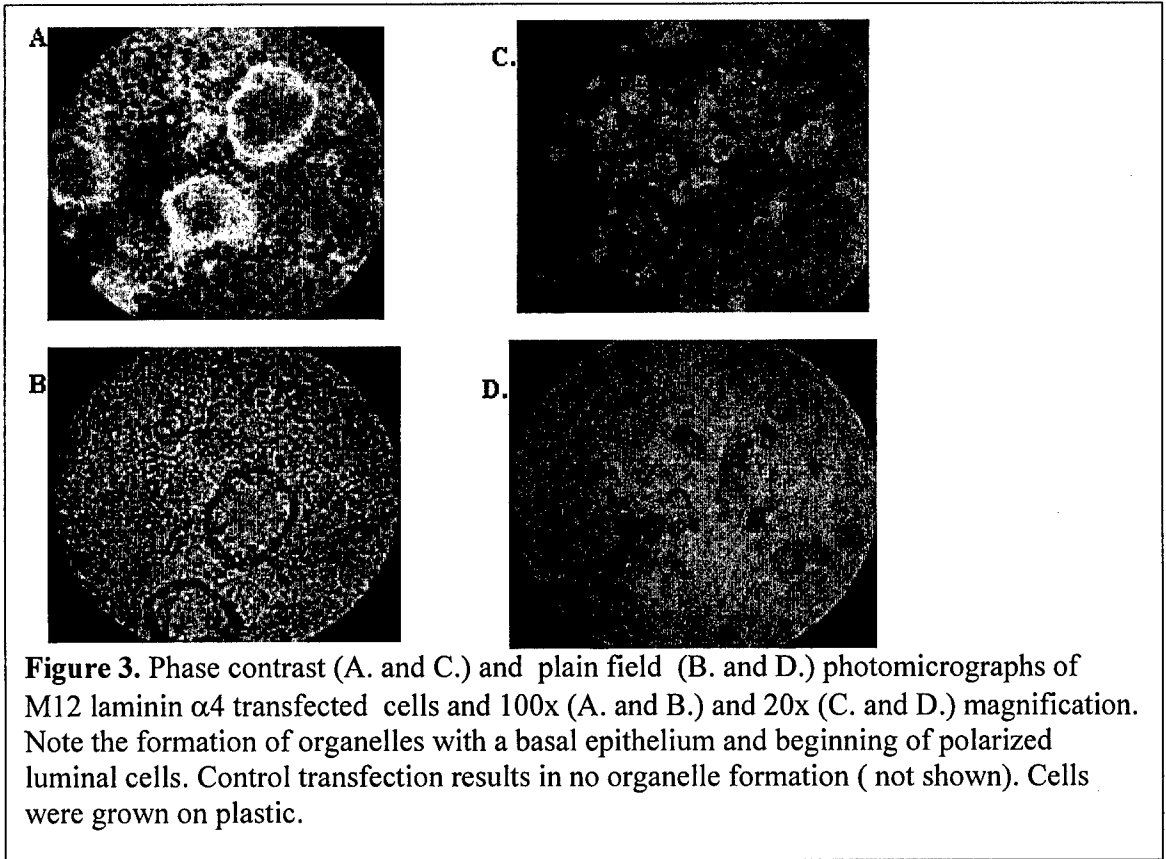


c) Determine cell cycle protein expression and cell kinetics in cells infected with adenoviral constructs. (Months 18-24).

As we have reported ( Drivdahl, Oncogene ) , SOX9 induced the expression of the androgen receptor(AR) in M12 cells. During this past year we have further demonstrated that the induction of AR is cell cycle dependent with expression beginning in late S-phase and greatest in G2M.

Of the genes that were regulated by SOX9, we were particularly interested in laminin subunit  $\alpha$  4. This subunit has not been previously described to be expressed by the prostate epithelial cell. It is of particular interest because it is associate with the production of laminins 8 and 9. The laminins may significantly alter the invasive properties of tumors but little is known of their function in prostate cancer. We have transfected the  $\alpha$ 4 laminin subunit into our M12 cells, figure 2 and have confirmed that the cell now expresses laminin 8 in its extracellular matrix. When the cells are grown in culture *in vitro*, as seen in figure 3, they began to form prostate gland appearing organelles. Boyden chamber assays demonstrated that matrix prepared from M12mac25 and M12SOX9 expressing cells in which the matrix is enriched for laminins containing the  $\alpha$ 4 subunit markedly suppressed invasion, figure 4.





### **C. Key Research Accomplishments**

- SOX9 expression in M12 cells results in the increased expression of 18 genes by 2 fold or more.
- Of the genes that are upregulated by SOX9, the  $\alpha 4$  laminin subunit was of particular interest because of significant changes that occur to the ECM.
- Extracellular matrix enriched in  $\alpha 4$  laminin subunit results in epithelial differentiation and significantly impairs *in vitro* assays of prostate cancer cell invasion.

### **D. Reportable outcomes:**

Drivdahl R, Tennant MK, Sprenger CT, Nelson PS, Plymate SR 2004 Transcription Factor SOX9 Regulation of Prostate Cancer Growth. *Oncogene*. 3;23(26):4584-93

Plymate SR, Roberts CTJr, Tennant MK, Haugk K, Woodke L, Marcelli M, Ware JL, 2004 IGF-IR Regulation of Androgen Receptor Signaling in progression to Metastatic Prostate Cancer. *Prostate*. 61:276-284

York TP, Plymate SR, Nelson PS, Eaves LJ, Webb HM, Ware JL. 2005  
cDNA Microarray Analysis Identifies Genes Induced in Common by Peptide Growth Factors and Androgen in Human Prostate Epithelial Cells. Revision Submitted to *Mol Cancer Res*.

**E. Conclusions:** The data generated during the past funding period (2004-2005) have demonstrated that SOX9 causes a significant decrease in prostate cancer progression, in part, by altering the structure of the extracellular matrix to promote differentiation and become a barrier to invasion.

### **F. References:**

Drivdahl R, Tennant MK, Sprenger CT, Nelson PS, Plymate SR 2004 Transcription Factor SOX9 Regulation of Prostate Cancer Growth. *Oncogene*. 3;23(26):4584-93

Plymate SR, Roberts CTJr, Tennant MK, Haugk K, Woodke L, Marcelli M, Ware JL, 2004 IGF-IR Regulation of Androgen Receptor Signaling in progression to Metastatic Prostate Cancer. *Prostate*. 61:276-284

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### **G. Appendices: None**