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MAML3 Overexpression Increases Tumorigenicity in Several Neuroendocrine Tumor Types



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BACKGROUND: The development of metastatic disease in neuroendocrine tumors (NETs) is not well-understood. MAML3 is a transcriptional co-activator. Recurrent MAML3 fusion genes are found in pheochromocytomas/parangliomas (PPGLs) associated with aggressive disease. Increased MAML3 protein expression is associated with metastatic disease in small bowel NETs, and high mRNA expression is seen in a subset of small cell lung cancers (SCLCs). Therefore, we aimed to investigate the role of MAML3 in NET tumorigenesis.

METHODS: Human PPGLs were used for immunohistochemistry and genetic analysis. Four NET cell lines, neuroblastoma (SK-N-SH), PNET (QGP1 and BON1) and SCLC (DMS114), were transiently transfected with MAML3 (FL) or exon 1 deleted MAML3 (dEx1; mimicking the fusion), and biologic effects of overexpression were examined in vitro. All assays performed in triplicate with >3 biological replicates. Statistical significance determined by ANOVA.

RESULTS: Through RT-PCR, 7% (4/55) of human PPGL have UBTF~MAML3 fusions, similar to TCGA, and all were sporadic cases with metastatic disease. Fusion-positive PPGL had intense MAML3 nuclear staining and increased β -catenin by IHC and showed increased WNT4 mRNA expression.

In vitro, overexpression of FL and dEx1 MAML3 increased invasion in SK-N-SH, QGP1, and BON1 (all $p < 0.05$), and increased soft agar colony formation in QGP1 and BON1 (all $p < 0.05$). In DMS114, only FL overexpression increased invasion ($p < 0.01$) whereas both FL and dEx1 increased colony formation (all $p < 0.01$). When investigating mechanism, we found increased TCF promoter activation by luciferase activity and co-immunoprecipitation between MAML3 and β -catenin in SK-N-SH, QGP1 and BON1 cells, suggesting WNT signaling pathway activation. Interestingly, this mechanism was not seen in DMS114 and transcriptome analysis suggested different pathway activation.

CONCLUSION: MAML3 overexpression is associated with increased tumorigenicity in several different NET cells. The mechanism of action may involve WNT signaling pathways in neuroblastoma and PNET cells, leading to metastatic disease.

ABSTRACT ID: 187