

Identifying the function of VSIG4 as seen in the tumor microenvironment of Glioblastoma

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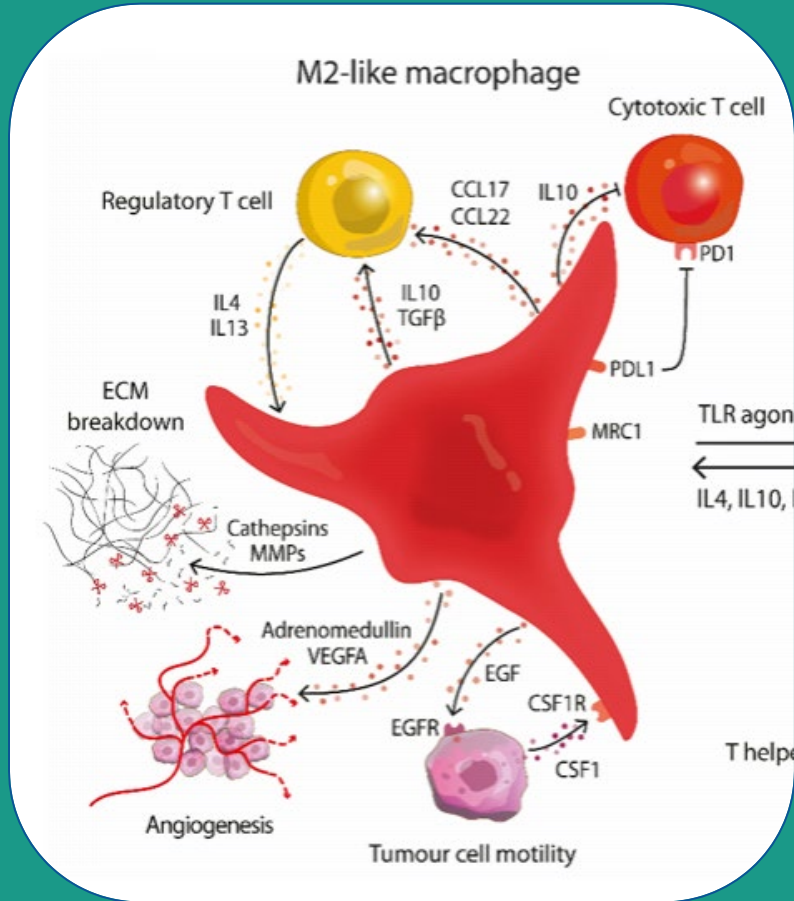


Glioblastoma (GBM)

- Most aggressive primary brain tumor
- 5 year survival rate less than a 10%
- Invasive nature makes it impossible to completely remove surgically
- Immunotherapies have failed to provide a survival benefit due to
 - Blood Brain Barrier
 - Tumor Microenvironment
 - Immune privileged organ



Tumor -associated macrophages (TAMs)



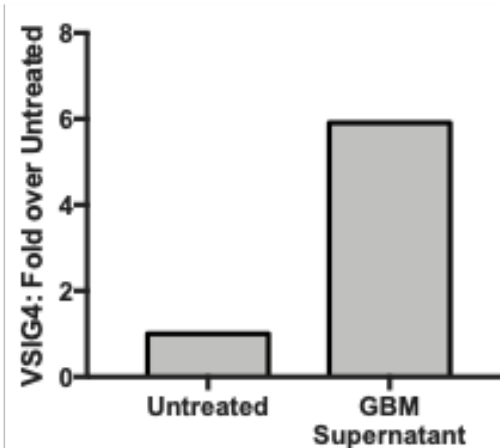
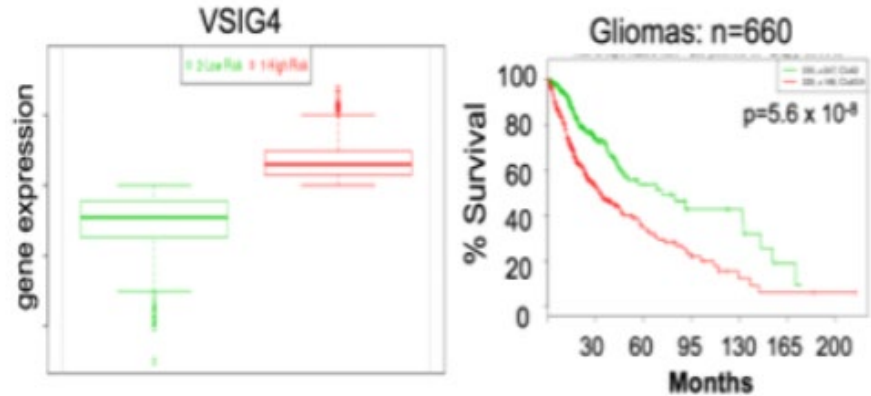
- The most abundant immune cell in GBM
- Initiate development of pro-tumor microenvironment
- Key in communication between the innate and adaptive immune system




VSIG4

The Crane lab performed a single cell RNA sequencing of TAMs and found that the protein VSIG4 is significantly upregulated in GBM TAMs (in vivo)

- Increased VSIG4 is also correlated with poor patient prognosis
- The Crane lab found that upon the treatment of macrophages with GBM supernatant, VSIG4 expression increased in vitro via qPCR



The experiment: mRNA analysis (by Nanostring)

Question

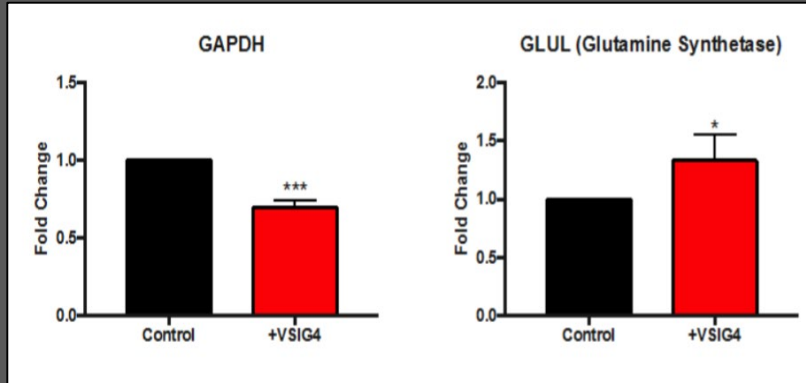
Does the overexpression of VSIG4 lead to changes in global gene expression?

Groups:

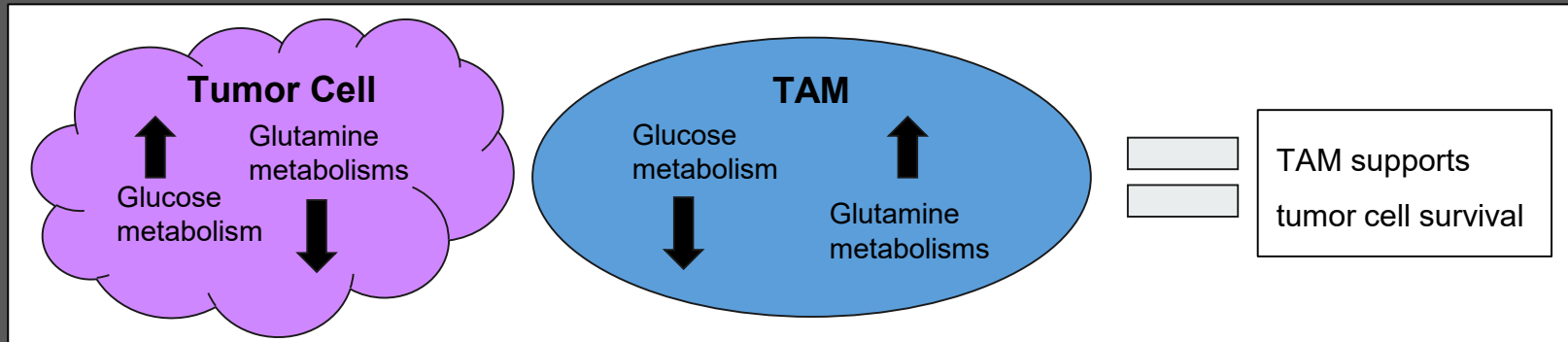
1. Control human macrophages (n=3)
2. VSIG4-overexpressing human macrophages (n=3)



Changes in glucose and glutamine metabolism



- Upregulation of GLUL (glutamine Synthetase)
- Downregulation of GAPDH (glyceraldehyde-3-phosphate dehydrogenase)





Conclusions and future directions

VSIG4 plays a crucial role in TAM survival and due to VSIG4's role in TAM survival, it is a strong candidate for further study

Using this information, we can:

- Run further experiments on VSIG4 examining the molecular factors causing its overexpression
- Examine other genes that may have a role in shifting TAMs to a fatty acid metabolism
- Engineer immunotherapies targeting VSIG4
- Block the upregulation of VSIG4 in TAMs



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