Growth Differentiation Factor-15 slows the growth of murine prostate cancer by stimulating tumor immunity

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Supplementary methods

Mouse genotyping

Transgenic mice were identified by PCR on DNA extracted from tail samples, using the transgene specific primers identified in the identified below.

1. For PB-SV40 T antigen sequence in TRAMP mice:

Pb-forward: 5'-CCGGTCGACCGGAAGCTTCCACAAGTGCATTTA-3' and SV40Tag-reverse: 5'-CTCCTTTCAAGACCTAGAAGGTCCA-3'.

2. For GDF15 transgene in TRAMP^{fmsmic-1} mice:

171MIC1 Flag-forward: 5'-GACTACAAGGACGACGATGACAAG-3' and 172MIC1MS8-reverse: 5'-CTCGGTGCACGCGGTAGGCTTCG-3'.

3. For RAG-KO mice:

1179_RAG1-KO-U1: TGGATGTGGAATGTGTGCGAG 1188_RAG1-WT-U1: GAGGTTCCGCTACGACTCTG 1189_RAG1-WT-L1: CCGGACAAGTTTTTCATCGT

Detection of TRAMP tumor infiltrating lymphocytes

Lymphocytes were identified following the gating strategy shown in S1 Fig. After identifying leukocytes using anti-CD45⁺, the antibodies used in Supplementary Table S1 identified T cells (CD3⁺), CD4 T cells (CD3⁺CD4⁺) CD8 T cells (CD3⁺CD8⁺), CD11c expressing CD8 cells (CD3⁺CD8⁺CD11c⁺), T cells bearing both CD4 and

CD8 (CD3⁺CD4⁺CD8⁺), T cells with no CD4 or CD8 which may be gamma delta T cells (CD3⁺CD4⁻CD8⁻), NK cells (CD3⁻CD11c⁻CD11b⁻B220⁻NK1.1⁺) and B cells (CD3⁻CD11c⁻CD11b⁻B220⁺NK1.1⁻).

In vivo CD8 T cell depletion

To validate CD8 T depletion procedure, we depleted CD8 T cells from WT mice by twice weekly intraperitoneal injection of 400μg of rat anti-CD8-α monoclonal antibody (clone 2.43, Bio X Cell). A control was similarly injected with 400μg of isotype matched control rat anti KLH antibody (Clone LTF-2, Bio X Cell). After one week spleen and axillary and brachial lymph nodes were collected, dispersed to single cell suspension and 1 million cells were stained with CD45-PerCP, CD3-BV21 and CD8B-BV786 (Clone H35-17.2, BD Biosciences) and subjected to flow cytometry. Flow cytometry results indicated that there was essentially depletion of CD8 T cells in both spleen and lymph nodes (S2 Table).