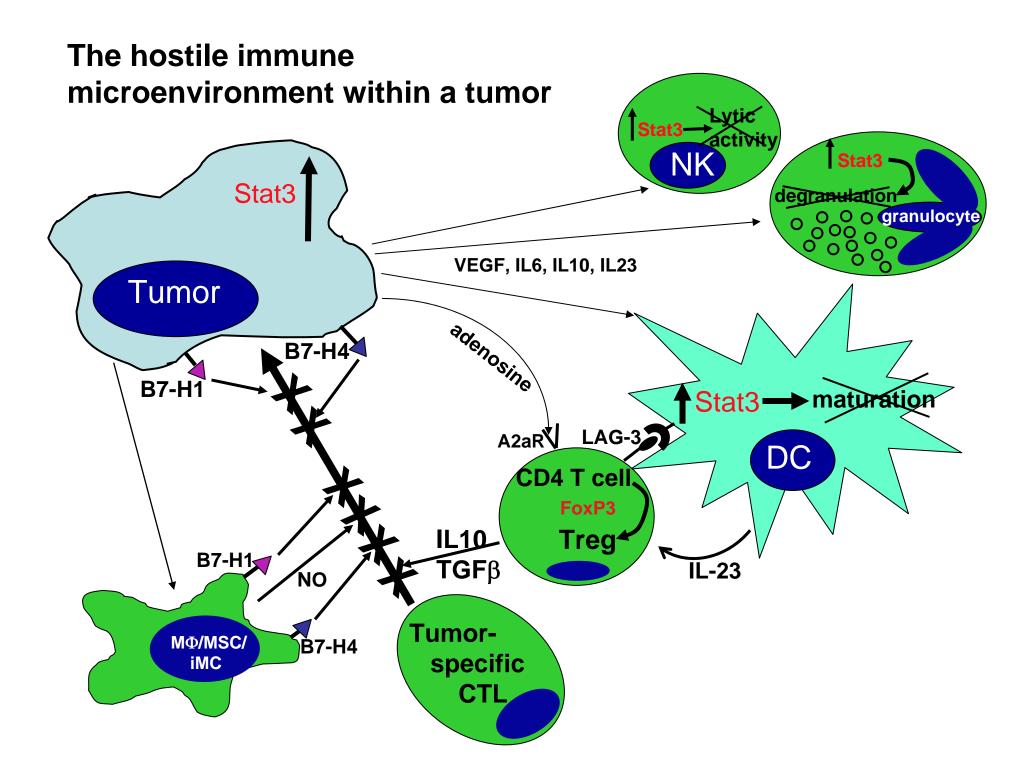
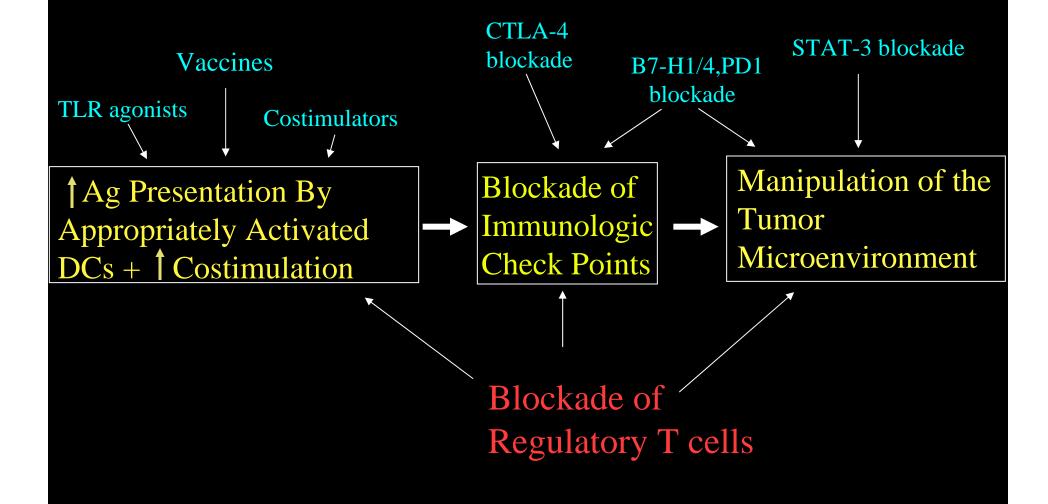
Regulating the Regulators for Cancer Immunotherapy: LAG-3 Finally Catches Up

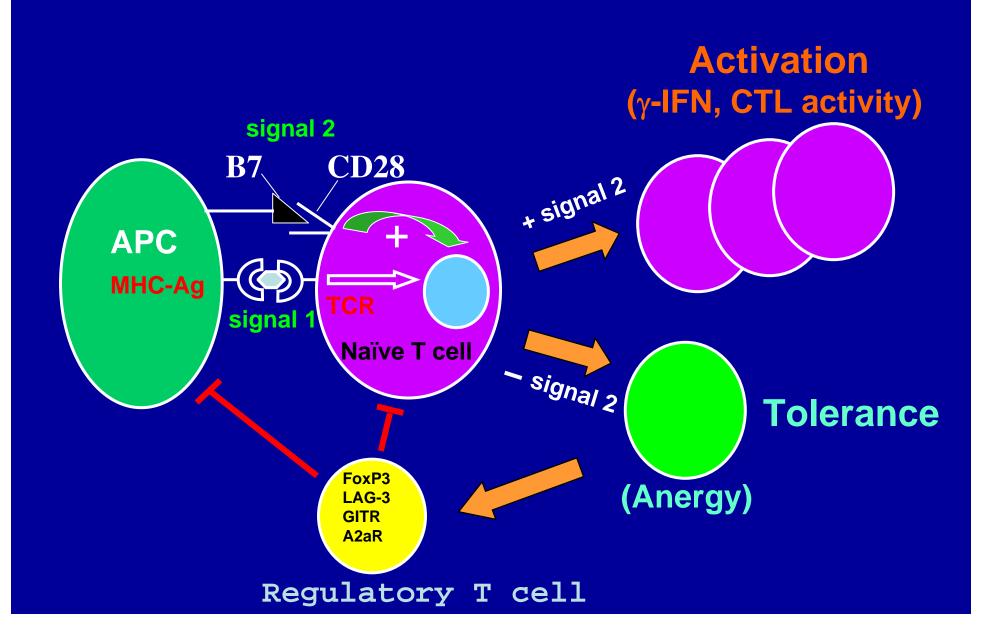
> Drew Pardoll Sidney Kimmel Cancer Center Johns Hopkins



Combinatorial immune therapies to hit distinct steps in the evolution of antitumor immunity



Peripheral T cell tolerance



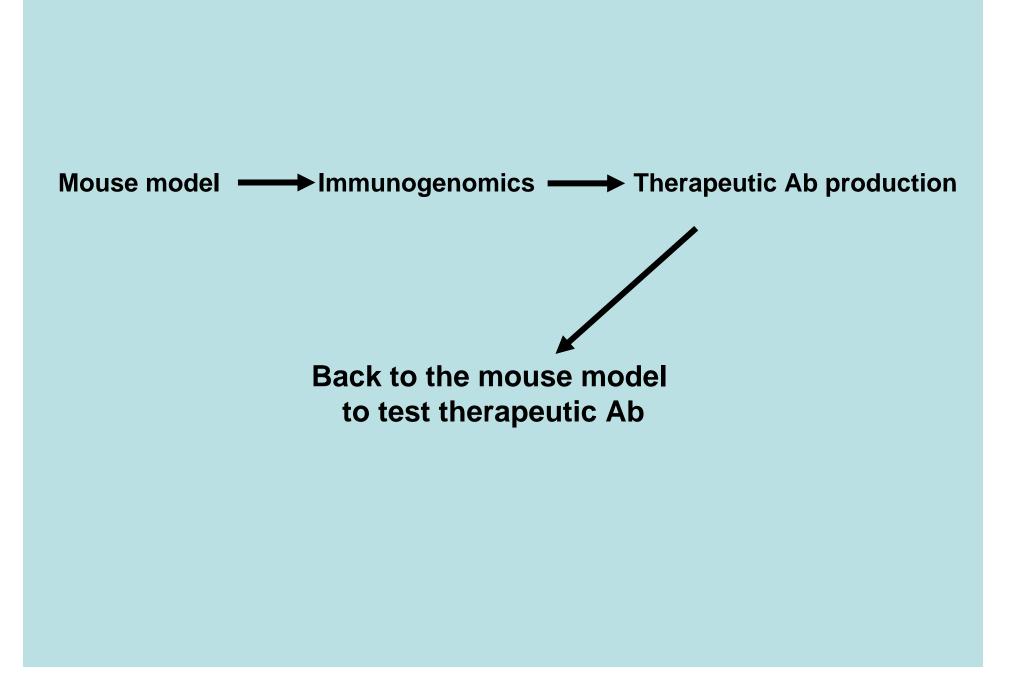
The Probasin – Hemagluttinin (ProHA) Transgenic Mouse

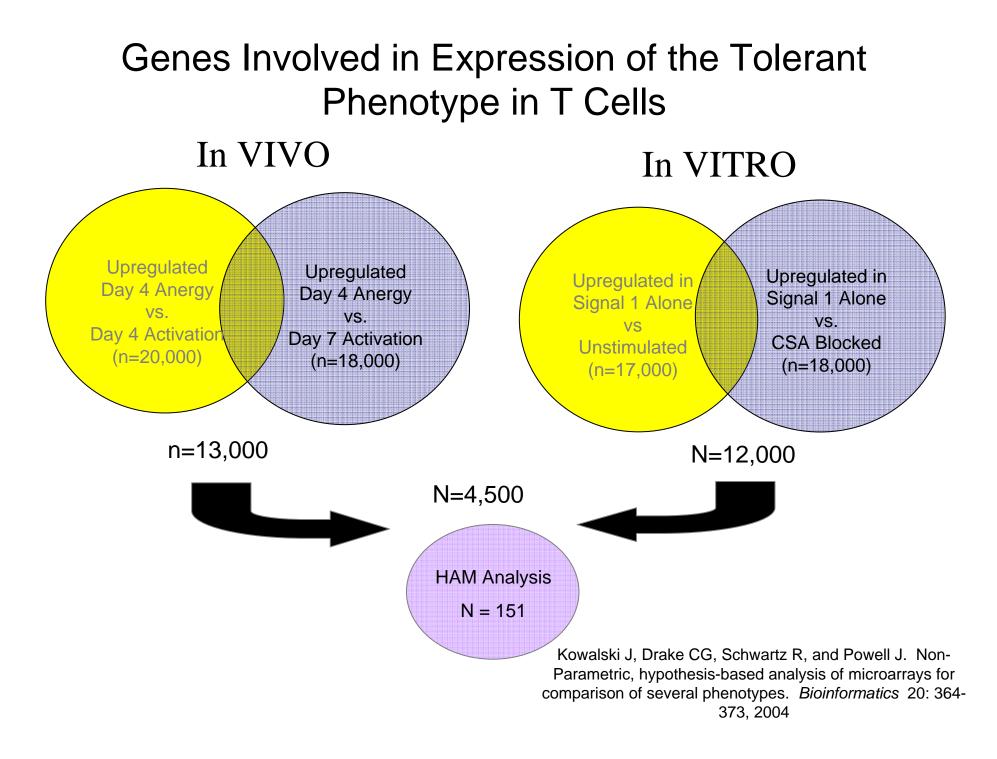


The ProHA x TRAMP Mouse(ProTRAMP) A tumor tolerance model

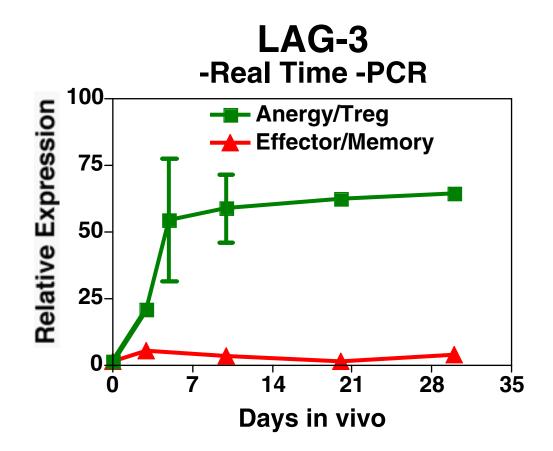
- ProSV40 Oncogenic
- ProHA A Tumor / and Tissue Specific Antigen
- Disease grossly identical to TRAMP
- 12'th Generation Intercross onto B10.D2
- HA-specific CD4 and CD8 TCR transgenic T cells transferred into ProTRAMP are TOLERIZED. HA-specific CD4s become Treg



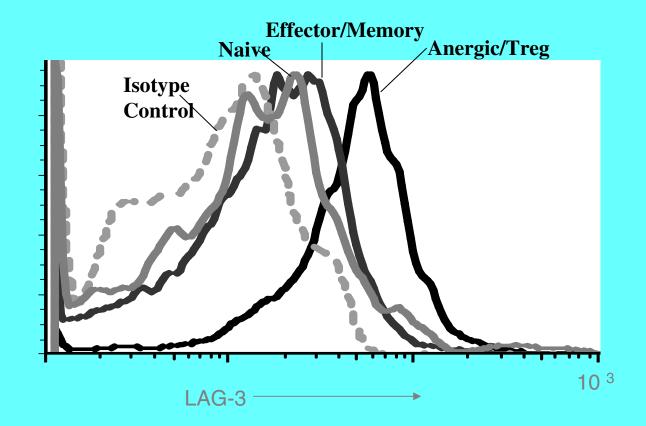




| Symbol | IN VIVO Ratio Anergy / Memory | IN VITRO Ratio Anergy / Resting | NAME - FROM GO Ontogeny Search | |
|----------|----------------------------------|------------------------------------|---|--|
| Lag3 | 3.6 | 8.7 | Lymphocyte-activation gene 3 | |
| Bcl3 | 3.5 | 14.6 | B-cell leukemia/lymphoma 3 | |
| Tnfrsf4 | 2.9 | 5.7 | tumor necrosis factor receptor superfamily member 4 | |
| Nfatc1 | 2.7 | 14.3 | NF-ATc isoform a (NF-ATca) | |
| 115 | 2.7 | 10.7 | Interleukin 5 | |
| Bcat1 | 2.6 | 34.0 | branched chain amino acid transferase 1, cytosolic | |
| Ptprs | 2.5 | 4.5 | Protein tyrosine phosphatase, receptor type, S | |
| Mapkapk2 | 2.4 | 10.5 | MAP kinase-activated protein kinase 2 | |
| Tubb5 | 1.8 | 4.5 | beta-tubulin (isotype Mbeta 5) | |
| Bcap37 | 1.8 | 4.1 | B-cell receptor-associated protein 37 | |
| Fhl2 | 1.7 | 8.3 | Four and a half LIM domains | |
| ll1r2 | 1.5 | 3.9 | Interleukin 1 receptor, type II | |
| Cish | 1.4 | 5.9 | Cytokine inducible SH2-containing protein | |
| Ndrg1 | 1.4 | 8.0 | N-myc downstream regulated gene 1 | |
| Etf1 | 1.4 | 4.6 | Eucaryotic tranlation termination factor 1 | |
| Prkch | 1.4 | 6.0 | Protein kinase C, eta | |
| Cnil | 1.3 | 24.7 | cornichon-like protein | |
| Tnfsf11 | 1.2 | 26.0 | Tumor necrosis factor (ligand) superfamily, member 11 | |
| II13 | 1.2 | 95.7 | Interleukin 13 | |
| Kcnn4 | 1.2 | 5.2 | calcium -activated potassium channel, small conductance | |
| Ccl1 | 1.1 | 159.9 | chemokine (C-C) motif ligand 1 | |
| Egr2 | 1.1 | 75.9 | Early growth response 2 | |
| ler3 | 1.1 | 8.7 | immediate early response 3 | |
| Gch | 1.1 | 30.5 | GTP cyclohydrolase 1 | |
| Rgs16 | 1.1 | 13.4 | regulator of G protein signalling 16 | |
| Csf1 | 1.1 | 3.7 | colony-stimulating factor 1 (macrophage) | |
| Fkbp8 | 1.0 | 8.2 | FK506 binding protein 8 | |
| Nr4a1 | 1.0 | 46.9 | nuclear receptor subfamily 4, group A member 1 | |
| Lcp2 | 1.0 | 4.4 | lymphocyte cytosolic protein 2 | |
| Dnajc5 | 1.0 | 16.1 | DnaJ (Hsp40) homolog | |



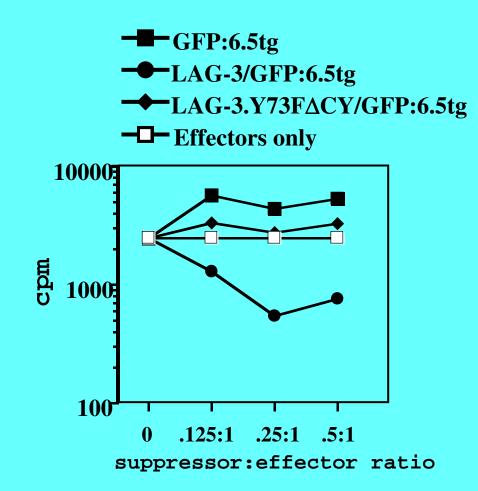
LAG-3 is highly expressed on the surface of induced Treg



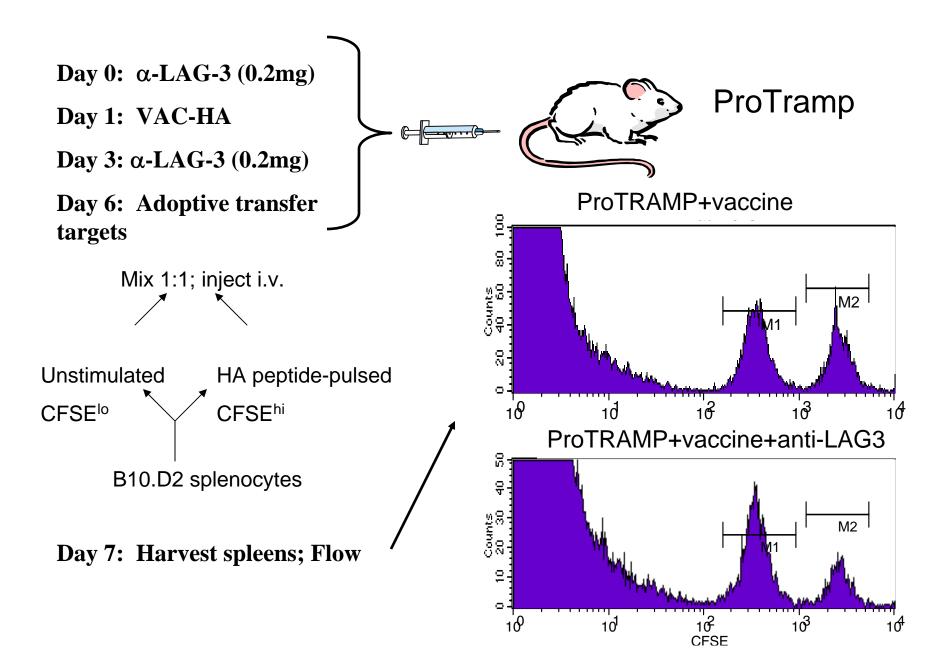
LAG-3

- Cloned in early 1990s
- CD4 homologue
- Does not substitute for CD4 in T cell development or helper T cell function
- Binds MHC II with higher affinity than CD4 but at a distinct site from CD4
- Function unclear reported to play a role in modulating NK function, T cell function, APC function but no clear conclusions
- Cytoplasmic tail completely different from CD4. Signaling pathways unclear

Transduction of CD4+CD25- T cells with wild type LAG-3 confers regulatory capacity

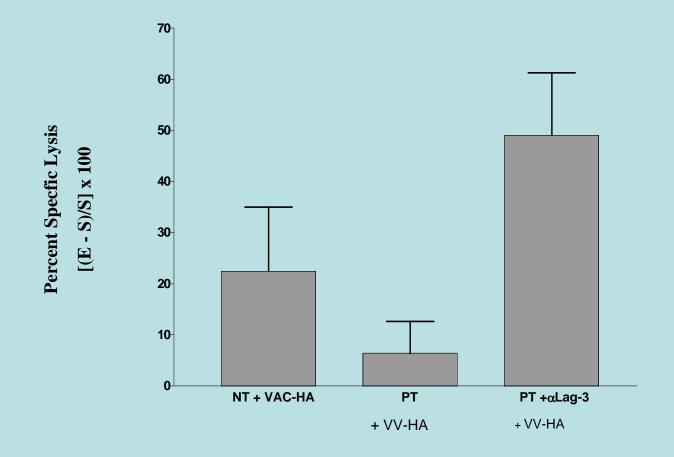


Can LAG-3 blockade alter endogenous T cell function?

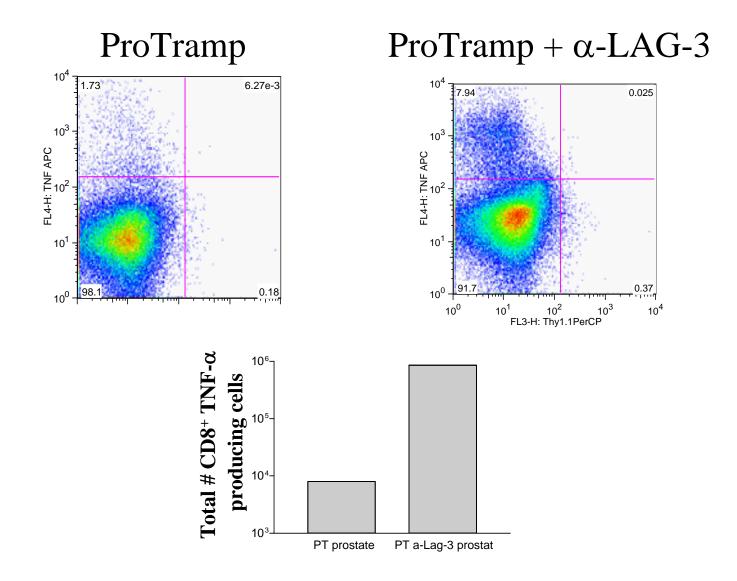


Tumor-tolerized endogenous CTL regain effector function in vivo after LAG-3 blockade

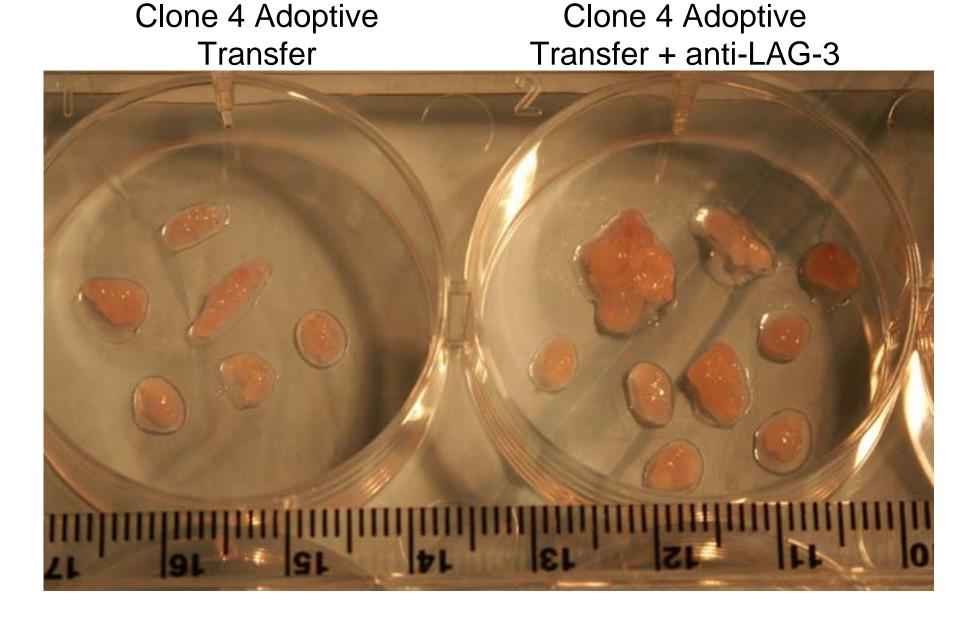
ProTramp (14-16 wk old)



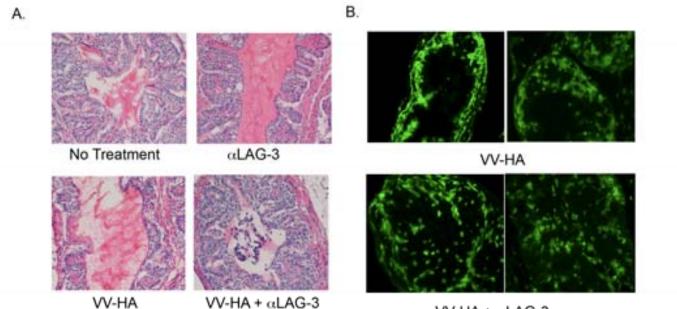
 α -LAG-3 leads to endogenous CD8 migration and TNF- α production within prostates of ProTRAMP mice



Prostate tissue from Pro-Tramp mice 7 days post-AT

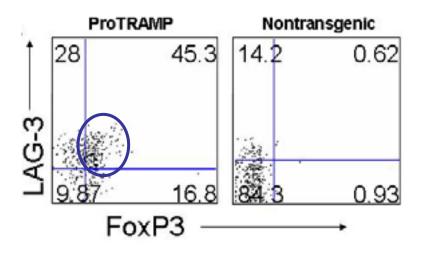


Histology of prostate cancers treated with vaccine + anti-LAG-3 antibodies

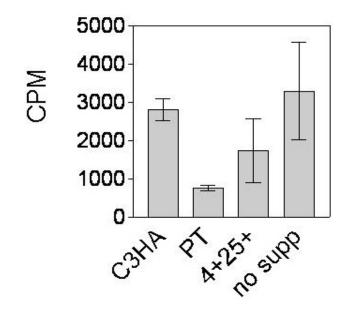


VV-HA + αLAG-3

LAG-3 and FoxP3 expression are concordant on antigen-specific CD4 T cells only when antigen is present as self or tumor



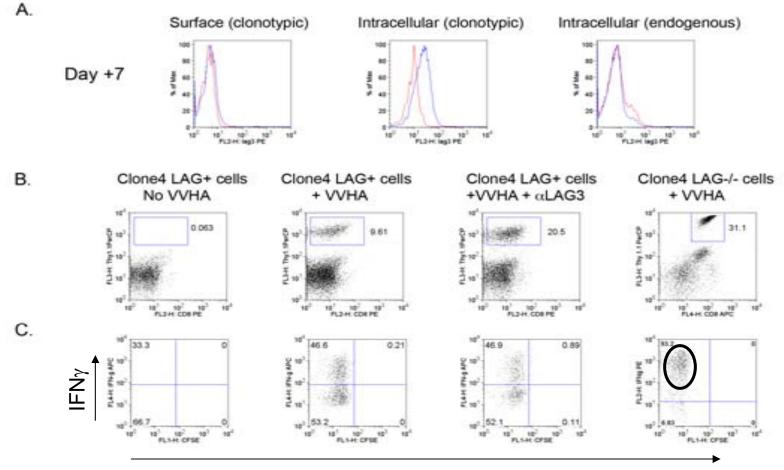
CD4+25+ Treg from tumor bearing mice are highly suppressive in *in vitro* assays



Membrane protein encoding genes upregulated among CD4+ cells infiltrating human prostate cancer

| Probe ID | Fold Increase | Gene Definition | Gene Symbol | Cellular Component |
|-------------|---------------|--|-------------|-----------------------|
| TNFSF9 | 122 | Tumor necrosis receptor superfamily, member 9 (41BB) | TNFRSF9 | Membrane |
| 234895_at | 95 | cytotoxic T-lymphocyte-associated protein 4 | CTLA4 | Membrane |
| 206486_at | 86 | lymphocyte-activation gene 3 | LAG3 | Membrane |
| 211269_s_at | 58 | interleukin 2 receptor, alpha (CD25) | IL2RA | Membrane |
| 223851_s_at | 31 | tumor necrosis factor receptor superfamily, member 18 (GITR) | TNFRSF18 | Membrane |
| 224211_at | 17 | forkhead box P3 | FOXP3 | Nucleus |

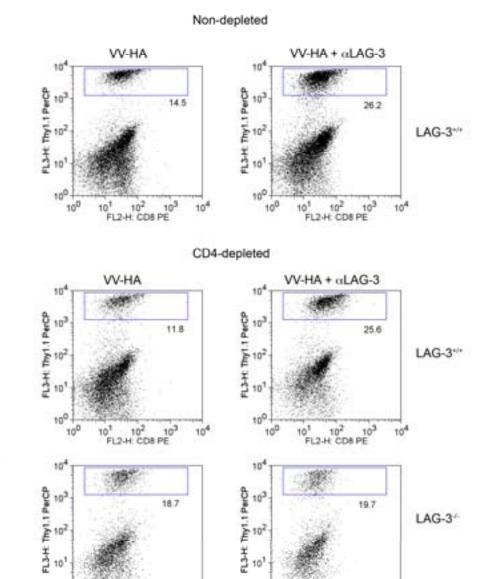
LAG3 expression on tumor specific CD8 T cells restrains their accumulation and function of in prostate cancers



CFSE

CD4-independent role of LAG-3 for tumor-specific CD8 T cells

Α.



100

100

10¹

0¹ 10² 10³ FL2-H: CD8 PE

104

C.

102

100

100

10¹

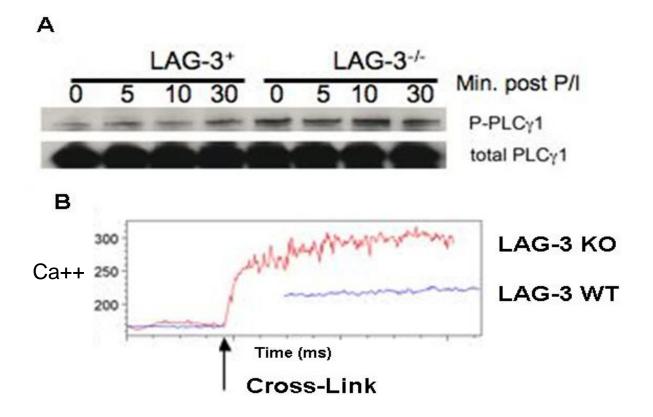
0¹ 10² 10 FL2-H: CD8 PE

103

104

B.

LAG-3 downregulates TCR dependent signaling



ProTRAMP model Charles Drake Adam Adler

LAG-3 Charles Drake Ching-Tai Huang Joe Grosso Tulia Bruno Ed Hipkiss Christin Kelleher Dario Vignale Craig Workman