Abstract

No. 1661

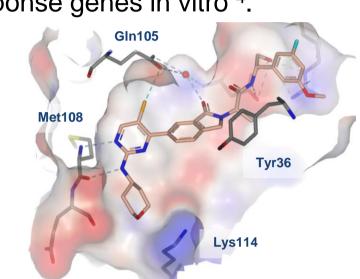
Immune modulation by the dual-mechanism ERK inhibitor, ASTX029, in MAPK-activated tumor models

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INTRODUCTION

- Activation of the MAPK pathway has been associated with a more immunosuppressive tumor microenvironment (TME) ¹
- Preclinical studies have demonstrated that targeting of BRAFV600mut or KRASG12C with agents such as dabrafenib or AMG510 leads to a more pro-inflammatory TME ^{2,3}.
- As a downstream node in the MAPK pathway, the inhibition of ERK1/2 (ERK) is an attractive therapeutic option for potentially overcoming acquired resistance and bypass signaling, with several ERK inhibitors under investigation in the clinic.
- Inhibition of ERK nuclear translocation by dual-mechanism ERK inhibitors has been reported to lead to an increase in expression of interferon response genes in vitro 4.
- We have recently described the discovery of ASTX029, a novel, dual-mechanism ERK inhibitor, through fragment screening and subsequent optimisation by structure-based drug design (SBDD) 5.
- ASTX029 is currently undergoing clinical development in a Phase 1 trial in advanced solid tumors (NCT03520075).
- Here we describe the immunomodulatory effects of ASTX029 in BRAF- and KRAS-mutant models.



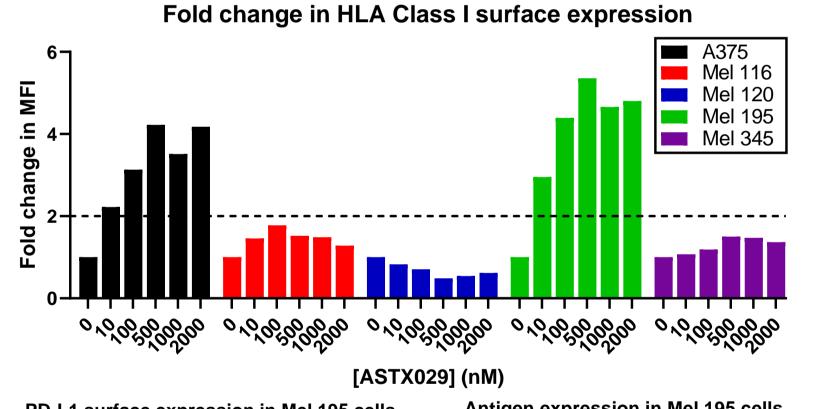
Surface expression of HLA

Class I was measured in

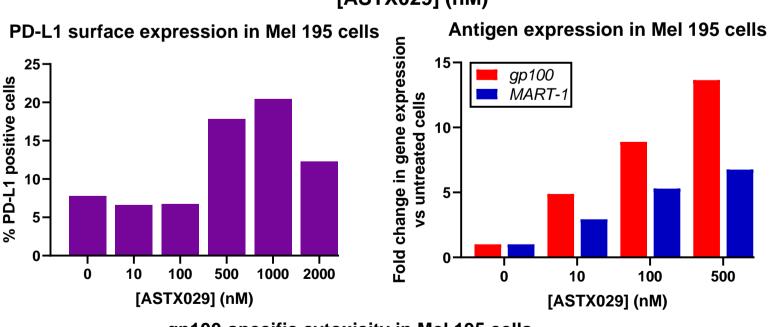
following

BRAF-mutant

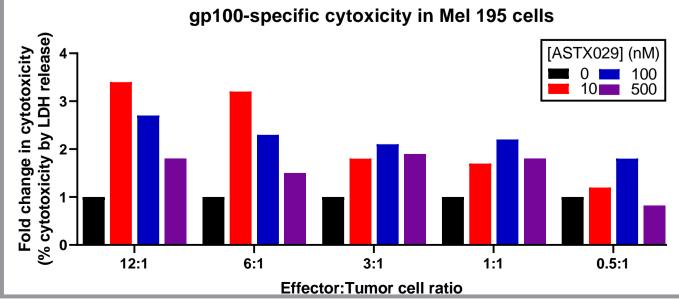
ASTX029 increases antigen presentation in human BRAF-mutant melanoma cell lines



treatment with ASTX029 2 out of 5 cell lines (A375 and Mel 195) showed an increase of >2-fold in HLA Class I expression upon



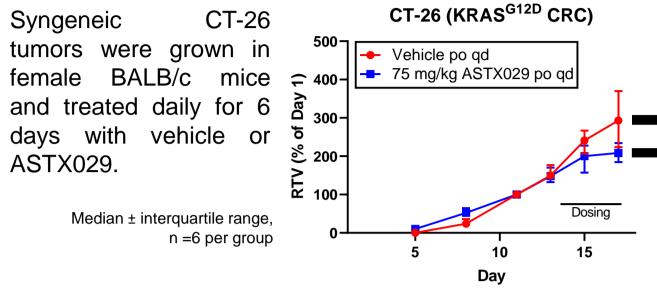
The Mel 195 cell line was selected for further analysis. Surface profiling revealed an upregulation of PD-L1 at higher concentrations of ASTX029. There was also a dose-dependent increase in expression melanoma antigens gp100 and MART-1.



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Treatment with ASTX029 caused an increase in antigen-specific T cell cytotoxicity following co-culture of Mel 195 cells with gp100-specific T cells at several ratios of effector to tumor cells.

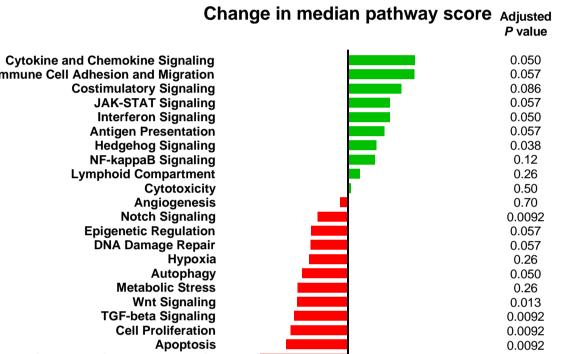
days with ASTX029.



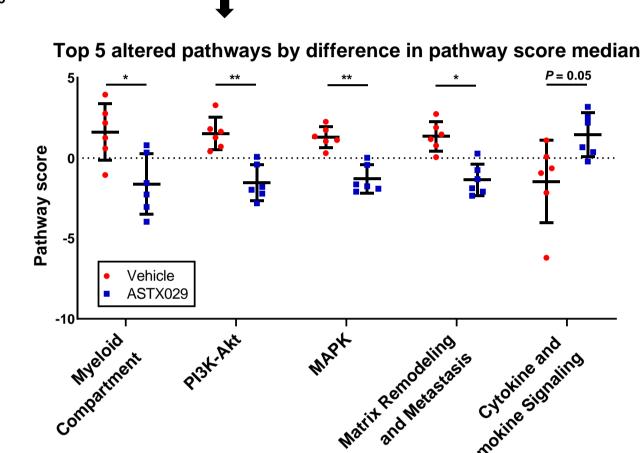
murine KRAS-mutant colorectal cancer model umors were collected on day 6 of dosing and divided, if large enough, f

gene expression analysis and for FFPE preparation.

Vehicle samples: 6 for IO 360 NanoString® panel, 6 FFPE samples for GeoMx® analysis ASTX029 samples: 6 for IO 360 NanoString® panel, 5 FFPE samples for GeoMx® analysis



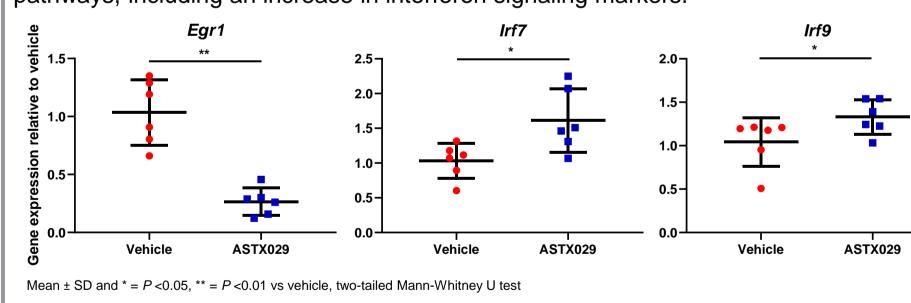
Difference in pathway score median ASTX029 vs vehicle



Mean \pm SD and $^* = P < 0.05$, $^{**} = P < 0.01$ vs vehicle, two-tailed Mann-Whitney U test adjusted by the

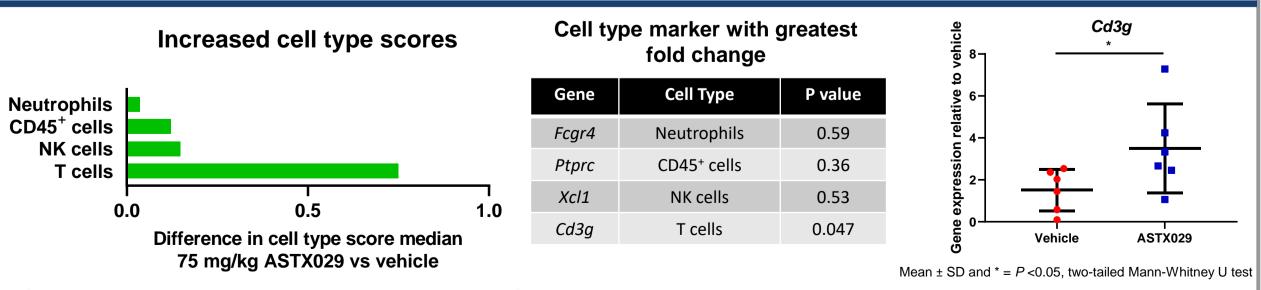
Gene expression changes in tumors were determined using the mouse PanCancer IO 360™ Gene Expression Panel of 770 genes. Pathway analysis of gene expression changes by nSolver™ software demonstrated changes in several pathways, including an increase in interferon signaling markers.

2. ASTX029 promotes pro-inflammatory TME gene expression changes in the CT-26



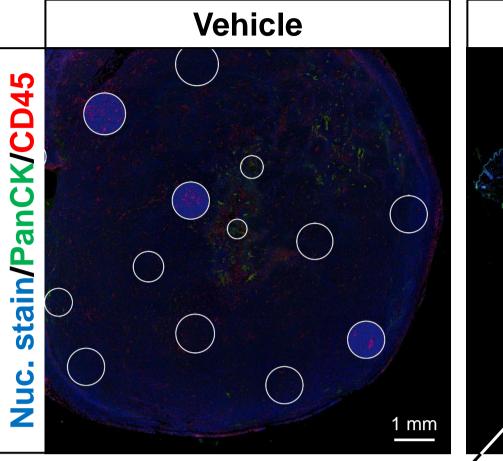
qRT-PCR analysis significant confirmed decrease in expression of the negative regulator Egr1 and an increase in expression interferon response genes and Irf9 following treatment with ASTX029.

3. ASTX029 increases T cell marker expression in CT-26 tumors

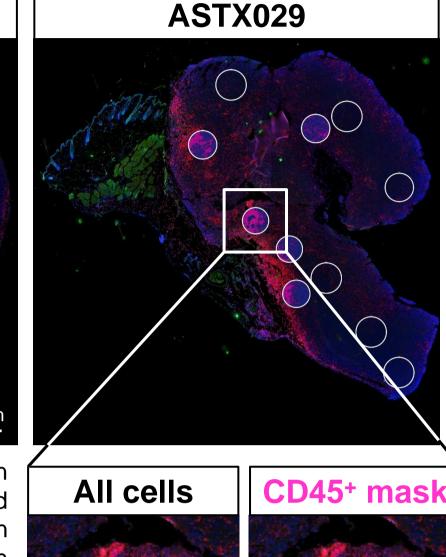


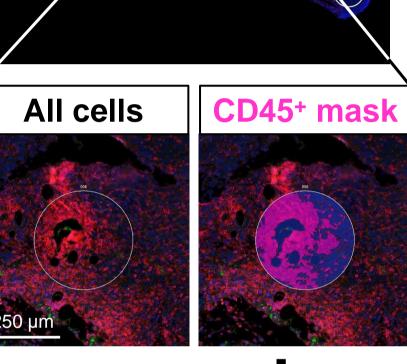
ASTX029 treatment resulted in increases in nSolver™ cell type score for several cell types, the largest increase being that for T cells. Treatment with ASTX029 induced a significant increase in the T cell marker gene Cd3g, which encodes a signaling component of the T cell receptor complex, as assessed by qRT-PCR.

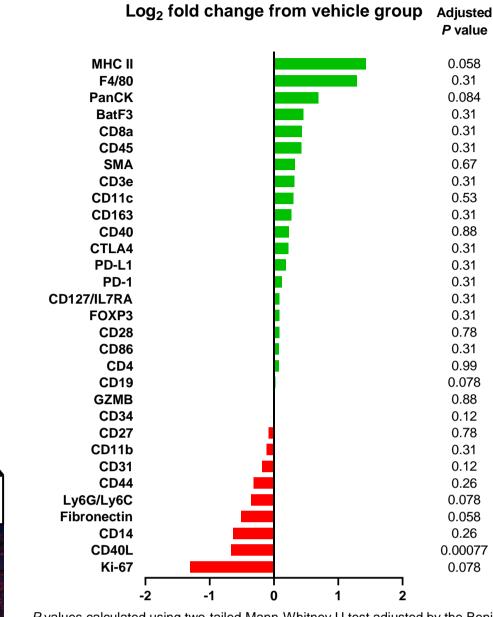
4. Treatment with ASTX029 induces changes in the myeloid compartment in CT-26 tumors



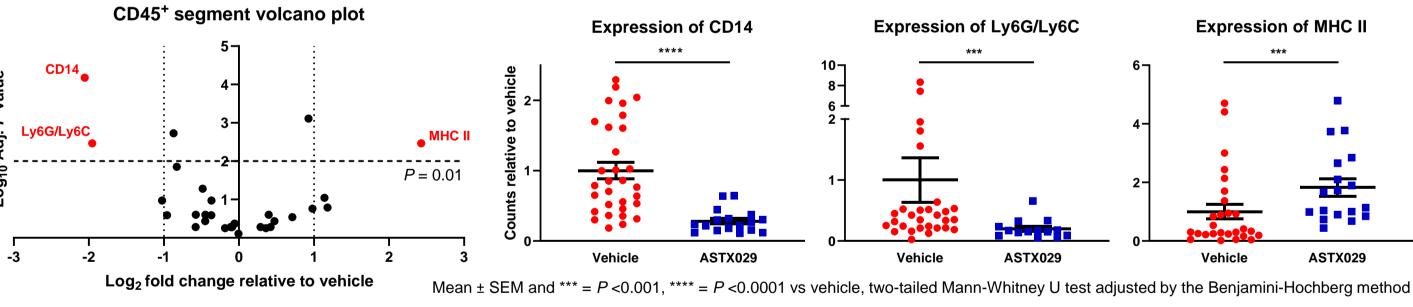
To further investigate the changes in gene expression observed in the myeloid lymphoid compartments, protein levels of 31 markers were assessed in FFPE sections of CT-26 tumors using NanoString® GeoMx® Digital Spatial Profiling. Sections were stained for nuclei, PanCK and CD45 before selection of regions of interest (ROIs) for analysis. Representative images and ROIs shown







Increased levels of CD45 following treatment NanoString® pathway downregulation of myeloid markers (CD14) and upregulation of antigen presentation (MHC II).



Regions were segmented for CD45

expression and selected for further analysis

Analysis of CD45+ regions demonstrated changes in myeloid cell phenotype on treatment with ASTX029, with a significant increase in MHC II levels, consistent with increased antigen presentation, and a significant decrease in Ly6G/Ly6C and CD14 levels.

SUMMARY AND CONCLUSIONS

- Treatment with ASTX029 leads to increased antigen presentation and antigen-specific T cell mediated killing in BRAFmutant melanoma cell lines.
- ASTX029 induces gene expression changes consistent with increased antigen presentation and a more proinflammatory TME with increased interferon signaling in the CT-26 KRAS-mutant colorectal cancer model.
- Changes in myeloid cell phenotype in CT-26 tumors following treatment with ASTX029 were detected by digital spatial profiling. Future studies will further define these phenotypic changes and expand the results beyond this model.
- These data will aid rational combination of ASTX029 with other tumor-directed or immunomodulatory agents as part of the ongoing clinical development of ASTX029.

Please see the related poster #CT108 LoRusso et al., A first-in-human, Phase 1 study of ASTX029, a dual-mechanism inhibitor of ERK1/2, in relapsed/refractory solid tumors.