A Highly Selective and Potent HPK1 Inhibitor Enhances Immune Cell Activation and Induces Robust Tumor Growth Inhibition in a Syngeneic Tumor Model

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BACKGROUND

HPK1, a member of the MAP4K family of protein serine/threonine kinases, is involved in regulating signal transduction cascades in cells of hematopoietic origin. Recent data from HPK1 knockout animals and kinase-inactive knock-in animals underscores the role of HPK1 in negatively regulating lymphocyte activation. This negative-feedback role of HPK1 downstream of lymphocyte activation and function, combined with its restricted expression in cells of hematopoietic origin, make it a compelling drug target for enhancing anti-tumor immunity.

METHODS

A structure-based drug design approach was used to identify potent and selective inhibitors of HPK1. Biochemical assays, as well as primary human and mouse immune cellbased activation assays, were utilized for multiple iterations of structure-activity relationship (SAR) studies. In vivo efficacy, target engagement and pharmacodynamic data were generated using murine syngeneic tumor models.







FIGURE 5. Increased Serum Cytokines and Circulating Antibodies (Pharmacodynamic Responses) in NMBS-2 Treated CT-26 Animals











FIGURE 6. NMBS-2 Induces Robust Tumor Growth Inhibition as a Single Agent in the B16-F10 Syngeneic Model



CONCLUSIONS AND FUTURE STEPS

An oral, potent, highly selective small molecule HPK1 inhibitor demonstrates in vivo tumor growth inhibition as a single agent and in combination with anti-PD1 in syngeneic models. Observed pharmacodynamic responses in syngeneic models, including increased serum cytokines and circulating plasma antibody levels, are consistent with comprehensive immune cell activation. Further evaluation of this selective inhibitor in additional immune cell-based assays will continue to build upon the mechanistic understanding of HPK1 inhibition as a novel immunomodulatory approach for anti-tumor immunity.

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