

Immunosuppression underlies resistance to anti-angiogenic therapy

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A Massachusetts General Hospital (MGH) research team has identified a novel mechanism behind resistance to angiogenesis inhibitors - drugs that fight cancer by suppressing the formation of new blood vessels. In their report published in the *Journal of Clinical Investigation*, the team based in the Edwin L. Steele Laboratories for Tumor Biology in the MGH Department of Radiation Oncology describes finding in mouse models how anti-angiogenesis treatment induces a microenvironment that suppresses immune systems actions that would otherwise help to eliminate a tumor. They also developed a potential strategy for getting around this resistance mechanism.

"Deciphering and targeting mechanisms involved in <u>resistance</u> to antiangiogenic therapy is critical to realizing the full potential of this promising cancer therapy," says Dai Fukumura, MD, PhD, deputy director of the Steele Labs, co-senior author of the paper. "Not only is this the first report investigating the role in anti-angiogenic cancer therapy of a subset of innate immune cells - Ly6Clow or non-classical monocytes - it is also the first to find an immunosuppressive function for these cells and to identify that as the key mechanism conferring resistance to anti-angiogenic therapy."

Angiogenesis inhibitors - often given in combination with traditional therapies - can improve treatment for several types of cancer, both by restricting the growth of new blood vessels and by "normalizing" the abnormal vessels in and around a <u>tumor</u> that can interfere with both chemotherapy and radiation therapy. But resistance to anti-angiogenic



therapy limits the drugs' survival benefits. Several studies have suggested a role in the development of resistance for the immune system particularly innate immune cells that suppress immune activity. But how specific subsets of these cells contribute to resistance has not been defined.

A series of experiments in mouse models of colorectal cancer first revealed that treatments blocking the vascular endothelial growth factor (VEGF) pathway - the target of approved anti-angiogenic drugs - induce the accumulation of monocytes and neutrophils. It soon became apparent that the buildup of non-classical monocytes - a subset previously identified as patrolling healthy blood vessels and possibly having an antitumor effect in lung cancer - was responsible for development of an immunosuppressive tumor microenvironment in colorectal cancer.

The research team identified the signaling pathway by which VEGF blockade induces expression of a molecule called CX3CL1 on tumor cells, <u>attracting non-classical monocytes</u> that carry the matching receptor. Those cells, in turn, attract neutrophils with another molecule called CXCL5; and both immune cells express factors that inhibit the proliferation of T cells, reducing the overall immune response to the tumor. Examination of biopsy samples taken from human patients before and after anti-angiogenic therapy revealed that expression of these chemokines - molecules that attract immune <u>cells</u> - increased in response to anti-VEGF treatment.

Since several methods of experimentally blocking the pathway improved the effects of anti-VEGF therapy in the mouse models, the team collaborated with Massachusetts Institute of Technology investigators to develop a potential gene therapy approach. Utilizing nanoparticledelivered RNA interference against the interaction between CX3CL1 and its receptor, the approach significantly reduced the infiltration of non-classical monocytes into treated tumors and increased the beneficial



effects of anti-VEGF therapy in a mouse model.

"Targeting resistance mechanisms can improve the efficacy of antiangiogenic therapy drugs and help fulfill their promise against cancer," says Fukumura, an associate professor of Radiation Oncology at Harvard Medical School (HMS). "Our study's unveiling of a novel resistance mechanism to anti-VEGF <u>therapy</u> and the molecular mechanism underlying that resistance offers a basis for the development of novel and efficient immunotherapeutic strategies to treat solid tumors."

Rakesh K. Jain, PhD, director of the Steele Labs and co-senior author of the *Journal of Clinical Investigation* report, adds, "Tumors also escape the immune system through immune checkpoint molecules like PD-1 and CTLA-4, which are the targets of recently approved drugs. But that strategy has been effective only in some tumor types and in only a fraction of patients. Therapeutic strategies based on our findings regarding the immunosuppressive action of non-classical monocytes may prove to have synergistic effect with those approved immune checkpoint inhibitors." Jain is the Cook Professor of Radiation Oncology (Tumor Biology) at HMS.

More information: Keehoon Jung et al, Ly6Clo monocytes drive immunosuppression and confer resistance to anti-VEGFR2 cancer therapy, *Journal of Clinical Investigation* (2017). <u>DOI:</u> <u>10.1172/JCI93182</u>

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