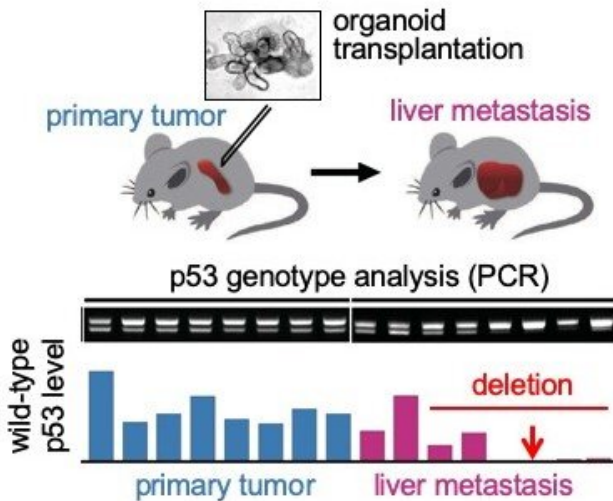


Notorious cancer protein mutations cooperate to proliferate disease

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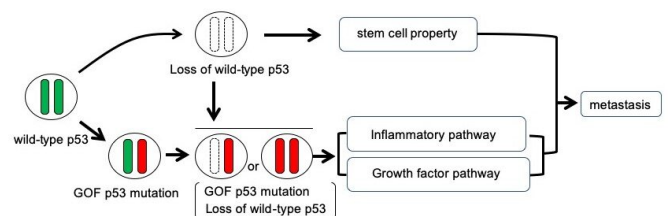
Tumor-derived organoids were transplanted to mouse spleen (top), and p53 status of the liver metastasized tumor cells were analyzed by PCR (middle). Notably, most metastasized tumor cells lost wild-type p53, suggesting that loss of p53 with GOF mutant p53 accelerates liver metastasis. Credit: Kanazawa University

Understanding the genetic mutations and protein changes that take place in the progression of cancer is key to its treatment. Mutations in the gene *TP53* and concomitant mutant p53 proteins in cancer cells have become notorious over the course of multiple studies of several different types of cancers. Yet so far, the mechanisms behind the role of these protein changes in propagating the disease has been poorly understood. Masanobu Oshima at Kanazawa University and his collaborators now report results that suggest just how these protein changes may lead to accelerated tumorigenesis and metastasis.

Previous studies of cancers in the lungs, intestines and pancreas, as well as colorectal [cancer](#), had suggested that changes in p53 lead to invasive

abilities and stem cell characteristics crucial for late-stage tumorigenesis. In addition to this 'gain of function (GOF)' mutation, wild-type protein p53 is reduced in [cancer cells](#) through a 'loss of heterozygosity' process, where instead of having two alleles for each gene corresponding to each parent, the gene has just one.

Oshima collaborated with researchers at Kanazawa University in Japan, Theragen Etx Bio Institute, and Seoul National University in South Korea. They found that [tumor cells](#) that carry GOF mutant p53 and lost wild-type p53 are enriched in liver metastasis lesions, suggesting combination of p53 LOH and GOF mutations promotes metastasis. In studies of single cells dissociated from each other, the researchers found that loss of heterozygosity was key to the cell's survival and proliferation, which has implications for metastasis. In addition, the structures formed were more complex and glandular.



Schematic illustration of the p53 mutation and metastasis. Combination of loss of wild-type p53 and GOF p53 mutation promotes metastasis through acquisition of stem cell property and activation of inflammatory and growth factor pathways. Credit: Kanazawa University

Further studies suggested that the combination of gain of function mutations and loss of heterozygosity provided some protection against cell death. When injected in live mouse models the

combination also accelerated tumorigenesis as well. The tumours formed had a fibrotic microenvironment, a signature of a poor prognosis for colorectal cancer.

Fibrotic structures are also linked with inflammation pathways. The researchers analysed the mutant proteins with the Ingenuity Pathway Analysis software package, which provides integrated analysis of gene expression and other cell traits. The analysis indicated [growth factor](#)/MAPK and inflammatory pathways were indeed upregulated in cells with gain of function mutations and loss of heterozygosity. What is more, trametinib treatment, (which inhibits a protein in the MAPK pathway) suppressed the cloning efficiency of these mutant cells. Ingenuity Pathway Analysis also revealed that stem cell pathways were up regulated in cells with gain of function mutations and loss of heterozygosity, as well as [cells](#) where both Trp53 alleles were disrupted. In their report, the researchers suggest their findings "will contribute to the future development of therapeutic strategies against CRC [colorectal cancer] metastasis."

More information: Mizuho Nakayama et al, Loss of wild-type p53 promotes mutant p53-driven metastasis through acquisition of survival and tumor-initiating properties, *Nature Communications* (2020). [DOI: 10.1038/s41467-020-16245-1](https://doi.org/10.1038/s41467-020-16245-1)

Provided by Kanazawa University

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