

Characterization of an oncogenic isoform of *TP53*: $\Delta 160p53$

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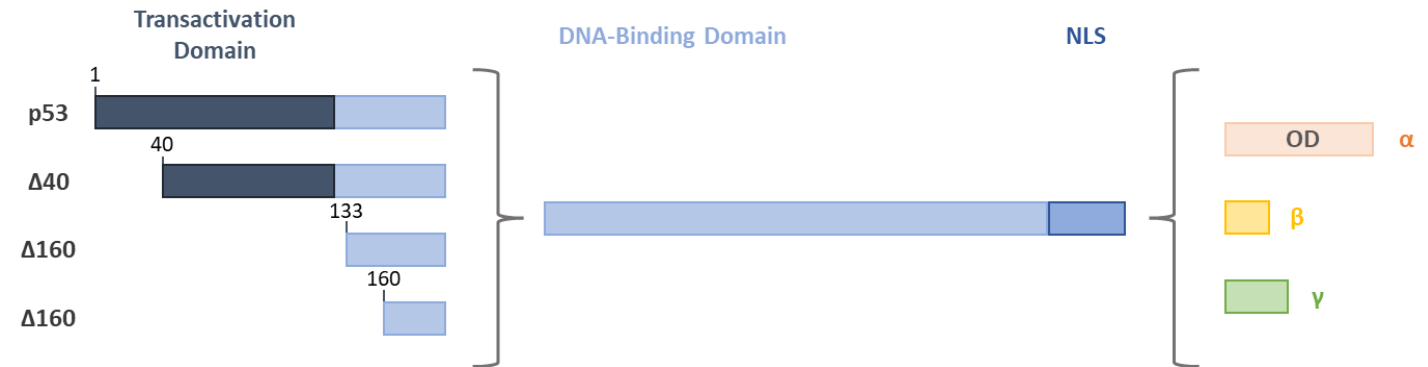
Introduction

The *TP53* gene

- Tumor suppressor gene.
- Encodes for **12 protein isoforms**.

- p53 α is a transcription factor kept at low levels in normal conditions.
- In response to stress, such as DNA damage, p53 α is activated to promote cell cycle arrest, apoptosis or senescence.

Guardian of the genome



- The *TP53* gene is mutated in over half of all human cancers (mostly missense mutations), leading to:

- Loss of wild-type tumor suppressor functions;
- Gain of new functions that actively drive tumorigenesis, invasion and metastasis.

Proto-oncogene

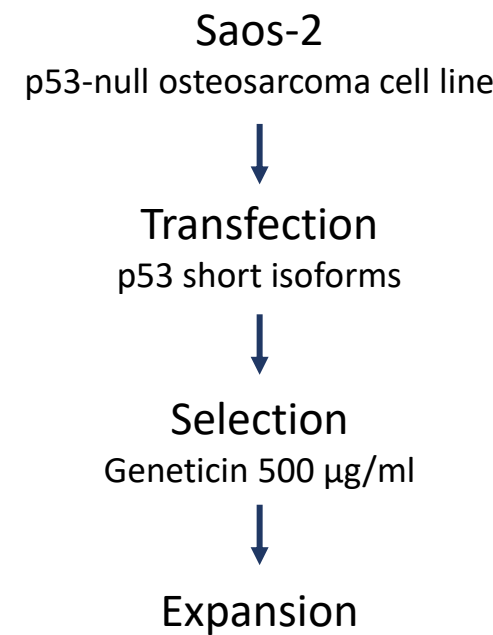
$\Delta 160$ p53 α

- This isoform is overexpressed in cancer cells and presents oncogenic functions.
- It is proposed that $\Delta 160$ p53 α is a proto-oncogene and the missense mutations commonly found in tumor cells promote oncogenesis by favoring the translation of this isoform.

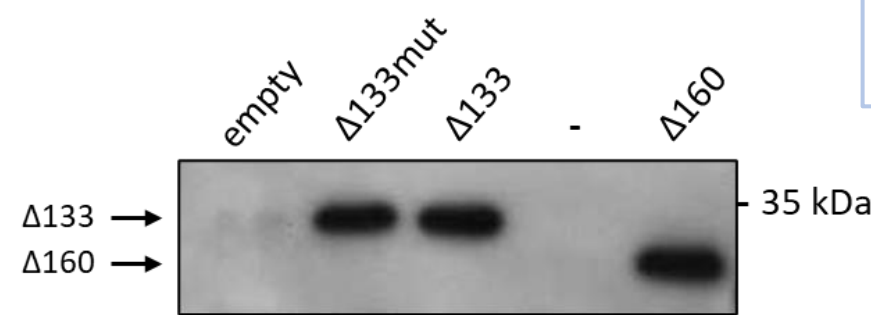
Anchorage-independent growth of cells expressing the p53 isoforms Δ133 and Δ160

Soft agar colony formation assay

Stable clones expressing the p53 isoforms:



Construct name	Description
Δ133	Δ133p53 with part of its 5'UTR sequence that corresponds to the beginning of exon 5
Δ133mut	Δ133p53 with kozak sequence; ATG160 and ATG169 are mutated
Δ160	Δ160 with kozak sequence
Empty	empty pcDNA3.1+ vector

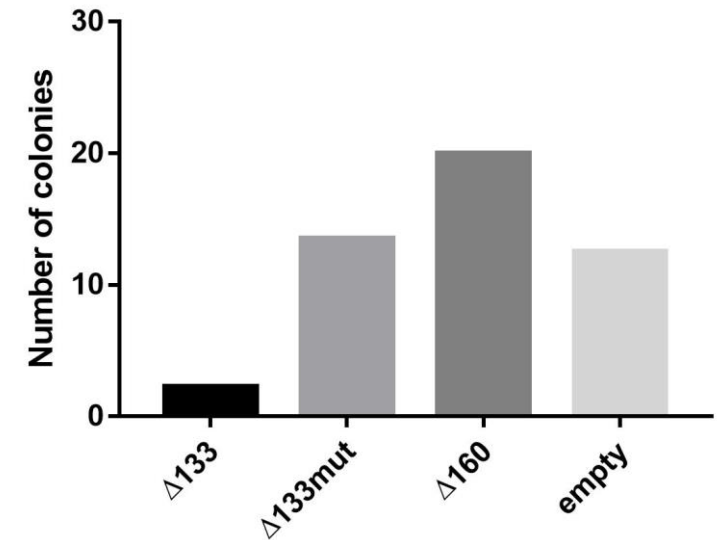
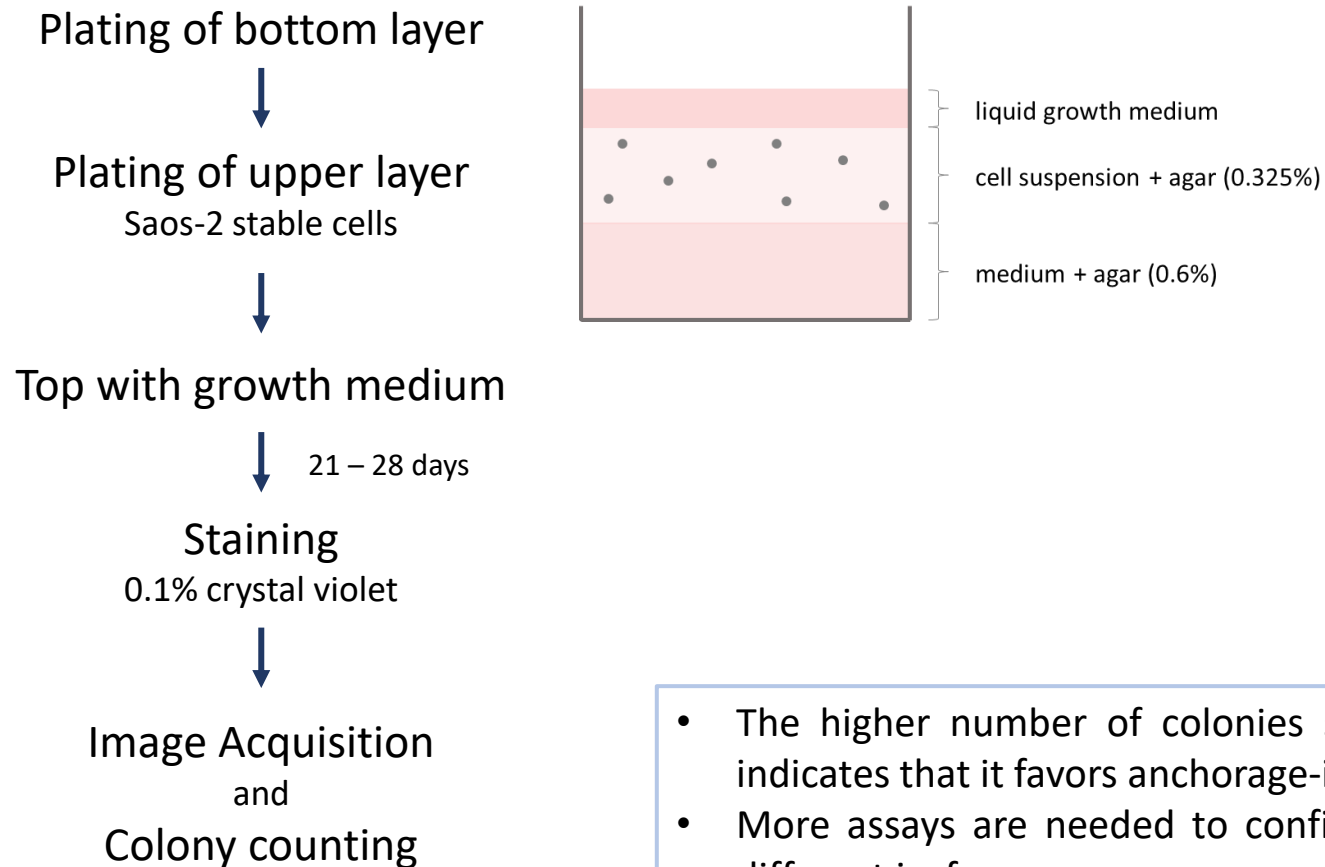


Confirmation that the cells stably express the transfected isoforms.

Western blot of Saos-2 cells stably transfected with p53 isoforms.

Anchorage-independent growth of cells expressing the p53 isoforms $\Delta 133$ and $\Delta 160$

Soft agar colony formation assay

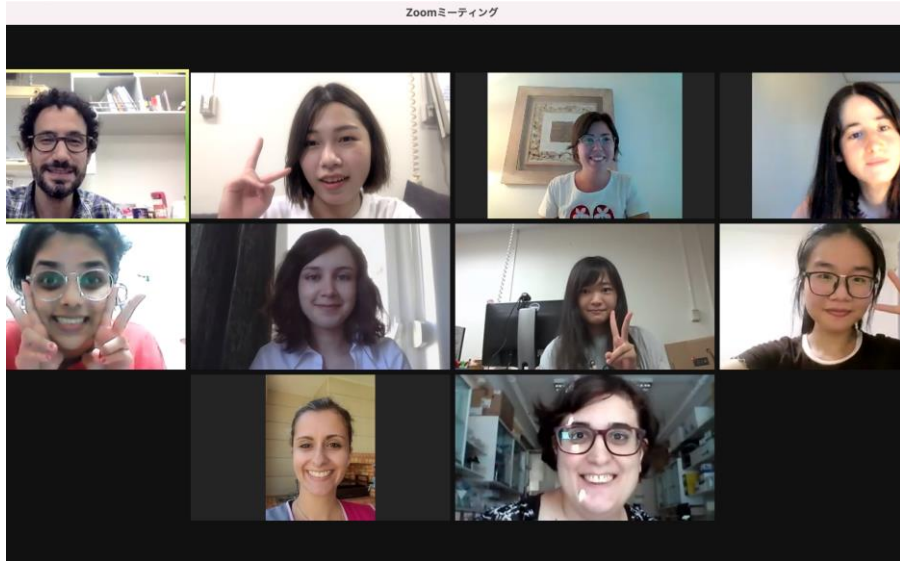


Soft agar colony formation assay of Saos-2 stable cells expressing p53 isoforms. Results from one experiment with 3000 cells (2 replicates).

- The higher number of colonies seen in $\Delta 160\text{p}53$ cells indicates that it favors anchorage-independent growth.
- More assays are needed to confirm the results for the different isoforms.

Acknowledgments

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