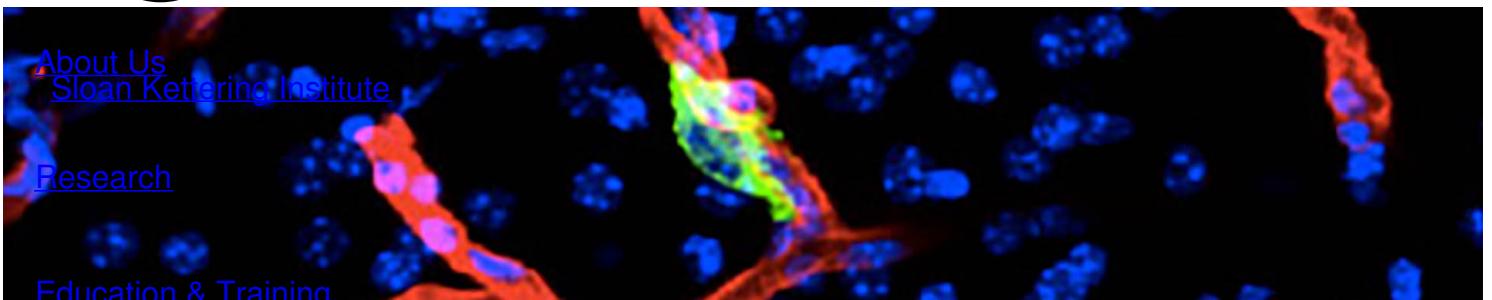


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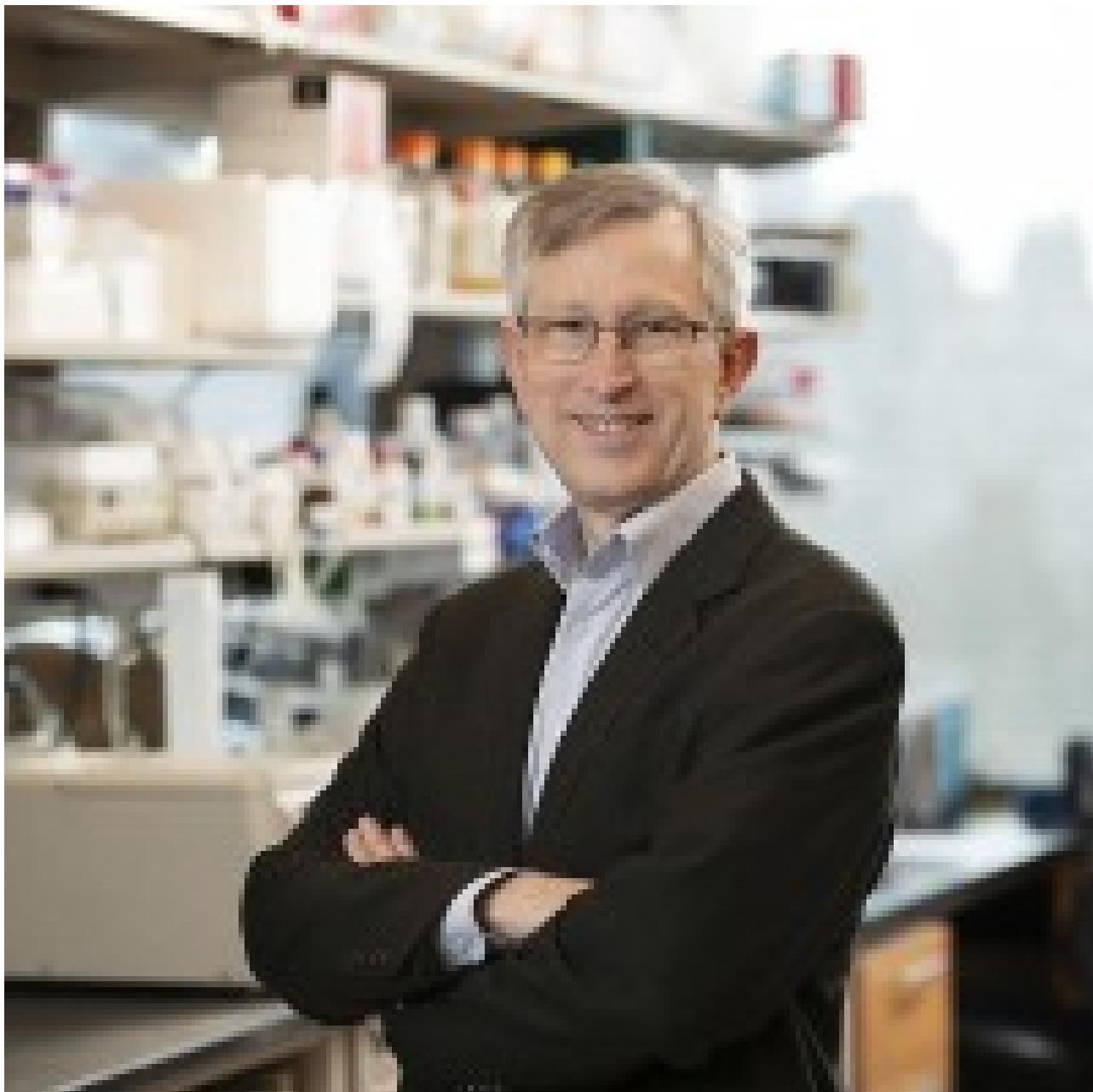
CANCER BIOLOGY & GENETICS PROGRAM

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The Joan Massagué Lab

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Research



Joan Massagué, PhD

Director, Sloan Kettering Institute; Member, Cancer Biology & Genetics Program; Marie-Josée and Henry R. Kravis Chair

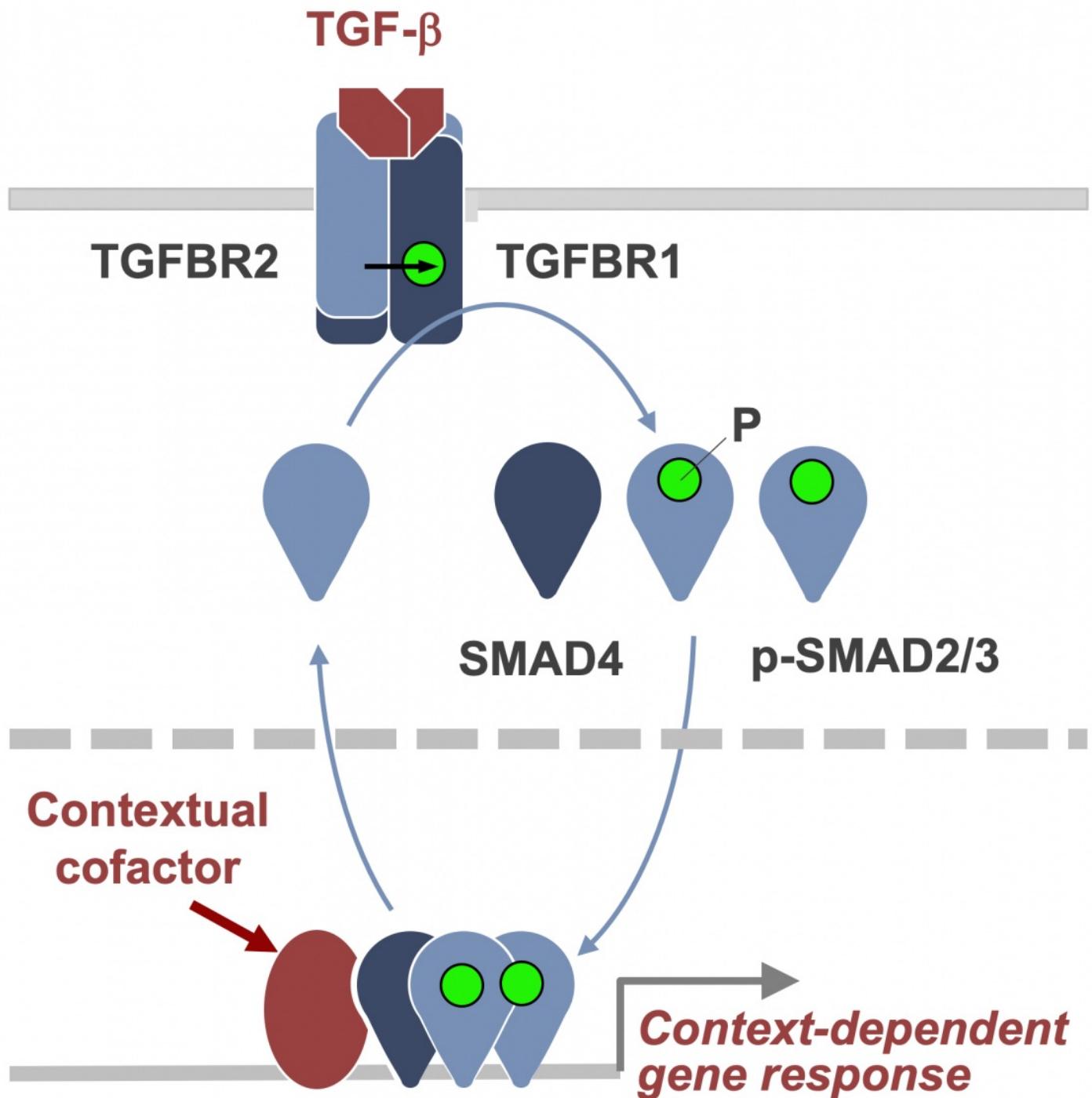
The Massagué lab investigates metastasis stem cells and their stromal niches throughout the metastatic cascade, with a growing interest in the dormant phase of metastasis. We are defining the phenotypic plasticity and evolution of metastatic cell populations from dormancy to outbreak, identifying drug targets, and enabling clinical trials to treat metastasis. We entered this area on realizing the central role of TGF- β and its interplay with oncogenic signals in metastasis. We collaborate with biologists, data scientists, and clinicians to leverage these findings for clinical

benefit.

Stem Cell Signaling, Growth, Control, and Cancer Metastasis

TGF- β signaling in development and disease

The TGF- β regulatory system plays crucial roles in the preservation of organismal integrity by simultaneously regulating different cell types and cellular functions. TGF- β signaling controls embryo development, tissue homeostasis, and injury repair through coordinated effects on cell proliferation, phenotypic plasticity, migration, metabolic adaptation, and immune surveillance of multiple cell types in shared ecosystems. Defects of TGF- β signaling disrupt immune tolerance, promote inflammation, underlie the pathogenesis of fibrosis and cancer, and contribute to the resistance of these diseases to treatment. Having elucidated the TGF- β signaling pathway, we are investigating how cells interpret TGF- β signals depending on the presence of context-dependent SMAD cofactors and chromatin configurations. Recent work uncovered a major interface between the TGF- β and RAS-MAPK pathways. TGF- β and RAS, signaling through SMAD and RAS-responsive element-binding protein 1 (RREB1) respectively, jointly activate a multi-arm regulatory program in epithelial progenitors and carcinoma cells (Su et al *Nature* 2020; Lee et al *Cell* 2024). RREB1 localizes to acetylated histone H4 marks in histone H2A.Z-loaded nucleosomes in cell plasticity genes, priming these enhancers for activation by a TGF- β activated SMAD4-INO80 nucleosome remodeling complex. These findings illuminate the operation of a bifunctional program that promotes metastatic outgrowth.

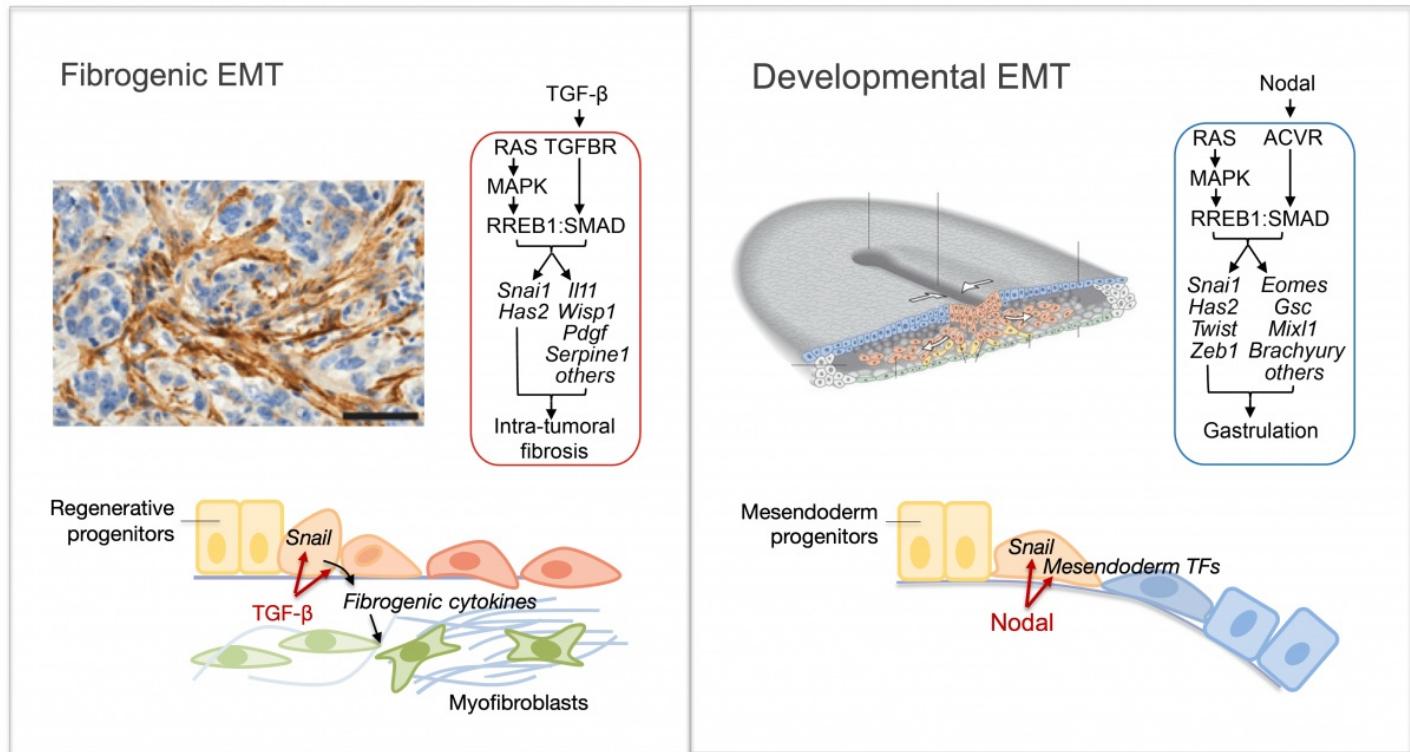


TGF- β signaling from membrane to nuclear target genes and basis for context-dependent effects (from David & Massagué Nature Revs. 2018).

Phenotypic plasticity regulation

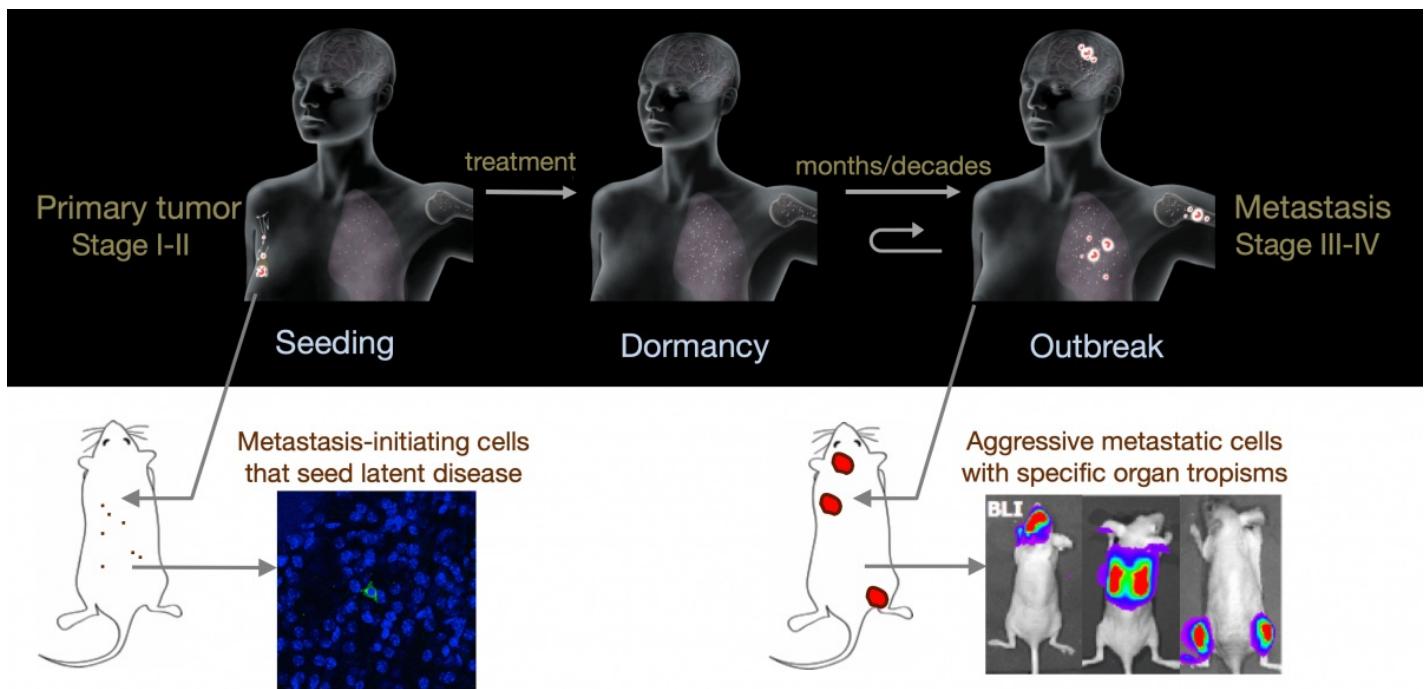
Epithelial-to-mesenchymal transitions (EMTs) are phenotypic plasticity processes that confer migratory and invasive properties to epithelial cells during development, wound-healing, and cancer. TGF- β is a potent inducer of EMTs implicated in liver disease, pulmonary fibrosis, and carcinoma.

metastasis. Lung adenocarcinoma (LUAD) cells undergo a TGF- β -dependent EMT associated with fibroblast activation and extracellular matrix remodeling. RAS-activated RREB1 primes enhancers of EMT and fibrogenic genes for activation by chromatin remodeling complexes that the TGF- β /SMAD pathway recruits to these enhancers. Both the EMT arm and the fibrogenic arm of this response program are essential for pulmonary metastasis in LUAD models. Inhibiting RREB1 disables this pro-metastatic process.



TGF- β orchestrates fibrogenic and developmental EMTs via the RAS effector RREB1 (from Su et al. *Nature* 2020).

In primitive, stem-like SOX2+ LUAD progenitors, TGF- β induces growth arrest accompanied by a full EMT response that subsequently transitions into an atypical mesenchymal state of round morphology and lacking actin stress fibers. TGF- β drives this long-term transition by inducing the expression of gelsolin, which converts a stress fiber-rich mesenchymal phenotype into a cortical actin-rich spheroidal state of low biomechanical stiffness to protect metastatic stem cells from killing by CD8 T cells and NK cells. Thus, LUAD primitive progenitors undergo an atypical EMT as part of a strategy to evade immune-mediated elimination (Wang et al *Nature Cancer* 2026). We are building on these insights to gain a better understanding of epithelial plasticity regulation by TGF- β in development, fibrosis and metastasis.



Stages of metastasis and mouse models for their analysis.

Metastasis initiating cells

Tissue homeostasis is maintained by stem cells, whereas damaged tissues are repaired by facultative progenitors that are activated upon injury. Developmental processes underlying normal tissue regeneration also operate in metastasis. Metastasis remains the main cause of death from cancer. The persistence and lethal relapse of disseminated cancer is driven by stem-like metastasis initiating cells. We are interested in understanding how the metastasis initiating phenotype emerges during tumor progression. Using patient tumor tissues, organoids, mouse models, lineage tracing, advanced imaging, and single cell analytics, the lab is investigating the nature of metastasis initiating cells and their stromal niches and evaluating L1CAM and related molecules as therapeutic targets.

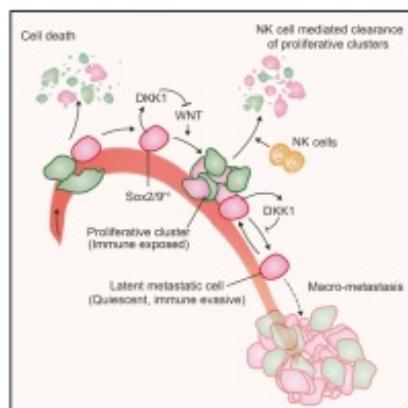
From metastatic dormancy to outbreak

Metastasis is a complex, multiorgan, and often fatal process (accounting for 90% of cancer-related deaths). Cancer cells that disseminate from a tumor to distant sites enter a period of dormancy that may persist from months to decades before giving rise to detectable metastasis. Adjuvant therapy treatments seek to prevent overt metastasis by eliminating residual malignant cells during this dormancy period. Efforts to improve adjuvant therapy are hindered by an insufficient understanding of the molecular mechanisms that preserve the long-term viability of dormant metastatic cells. Identifying these mechanisms is needed to improve treatments and prevent relapse. Using mouse

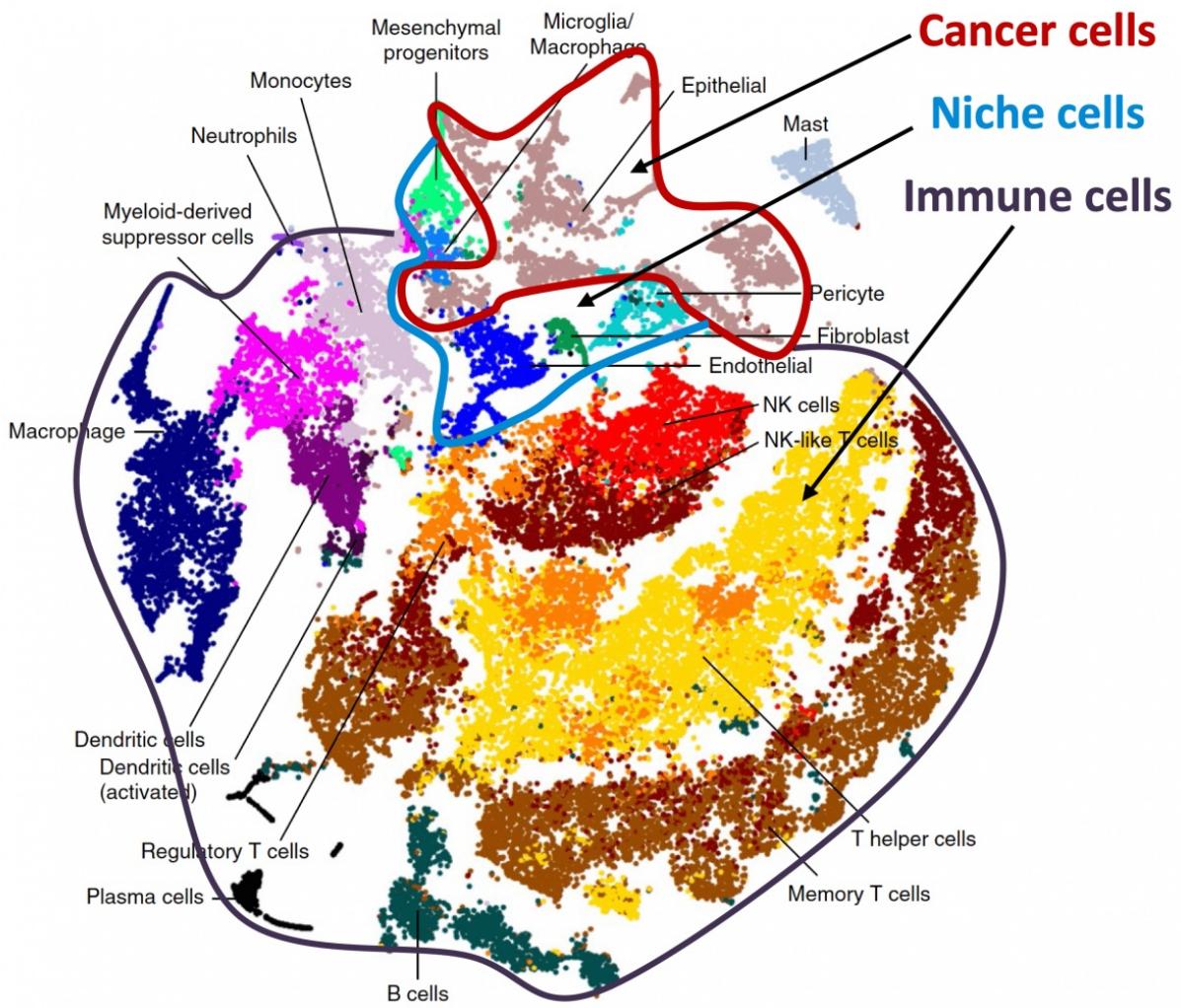
models that the lab developed, we are investigating the evolution of SOX2+ metastasis stem cells that enter a dormant phase after infiltrating target organs, remain viable under immune surveillance, and eventually reinitiate tumor growth. This work revealed roles of TGF- β as a driver of immune evasive metastatic dormancy and reveal vulnerabilities of metastatic cells. We uncovered key roles of STING (Hu et al *Nature* 2023) and L1CAM in this process (Park et al *bioRxiv* 2025) that can be exploited for the eradication of residual disease.

Organ-specific metastasis

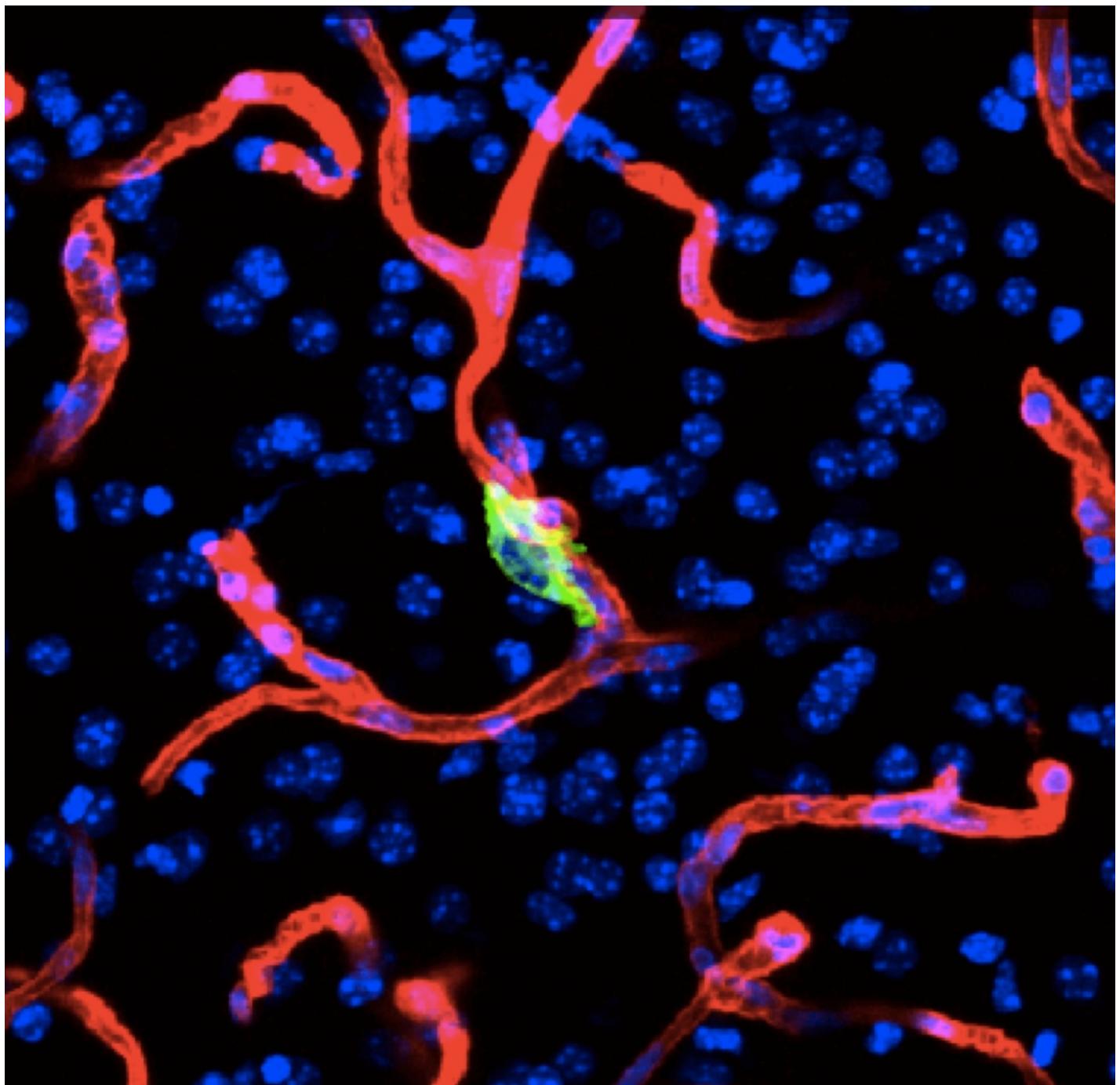
During metastasis, disseminated cancer cells undergo profound, tissue-specific reprogramming that reshapes their phenotype, metabolism, and therapeutic responses, giving rise to organ adaptations and organ tropisms characteristic of each type of cancer. The lab has identified a series of mediators of organ-specific metastasis and is currently focusing on the specific case of brain metastasis. Brain metastasis is highly lethal. Its incidence is ten-fold higher than that of all other brain tumor types combined. We created mouse models of brain metastasis from lung cancer and breast cancer and are using these models to identify relevant mediators of brain metastasis. In recent work we demonstrated that distinct tumor architectures and microenvironments for the initiation of metastasis in the brain (Gan et al *Cancer Cell* 2024). This work argues for an precision oncology framework that integrates these organ site-specific programs into treatment design.



Metastatic latency and immune evasion through self-imposed growth arrest in metastasis initiating cells (from Malladi et al *Cell* 2016).



Regenerative lineages and immune-mediated pruning of cancer cells in lung cancer metastasis as revealed by single-cell analysis (from Laughney et al Nature Medicine 2020).



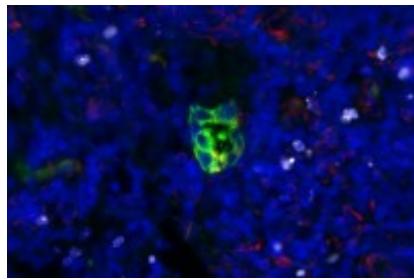
A metastasis initiating cell (green) has emerged from the circulation in the brain and is spreading on a capillary (red) to restart tumor growth. (from Valiente et al Cell 2014)

[View Lab Overview](https://www.mskcc.org/research/ski/labs/joan-massague/overview) (<https://www.mskcc.org/research/ski/labs/joan-massague/overview>)

VIDEO | 02:08

Go inside the lab of Joan Massagué, Chief Scientific Officer and Director, Sloan Kettering Institute.

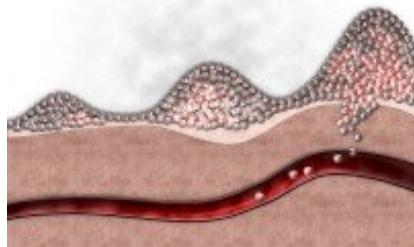
Featured News



[MSK Scientists Identify Potential New Strategy Against Metastasis](#)

A team of scientists at the Sloan Kettering Institute has identified the STING cellular signaling pathway as a key player in keeping dormant cancer cells from progressing into aggressive tumors months, or even years, after they've escaped from a primary tumor.

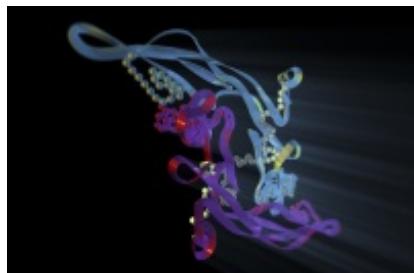
IN THE LAB



[What Does Cancer Metastasis Have to Do with Wound Healing? More than You Might Think](#)

Scientists are learning that — in a literal sense — metastasis is wound healing gone wrong.

FEATURE



[An Old Protein Gets a New Look: Researchers Target TGF-β to Make](#)

[Immunotherapy More Effective](#)

Drugs that block the action of TGF- β may make checkpoint inhibitors more effective.

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Publications Highlights

Wang Z, Elbanna Y, Godet I, Peters L, Lampe G, Chen Y, Xavier J, Huse M, Massagué J. TGF- β induces an atypical EMT to evade immune mechanosurveillance in lung adenocarcinoma dormant metastasis. *Nature Cancer*. (2026) Jan 5; DOI: 10.1038/s43018-025-01094-y [PMID:] [PMCID:]

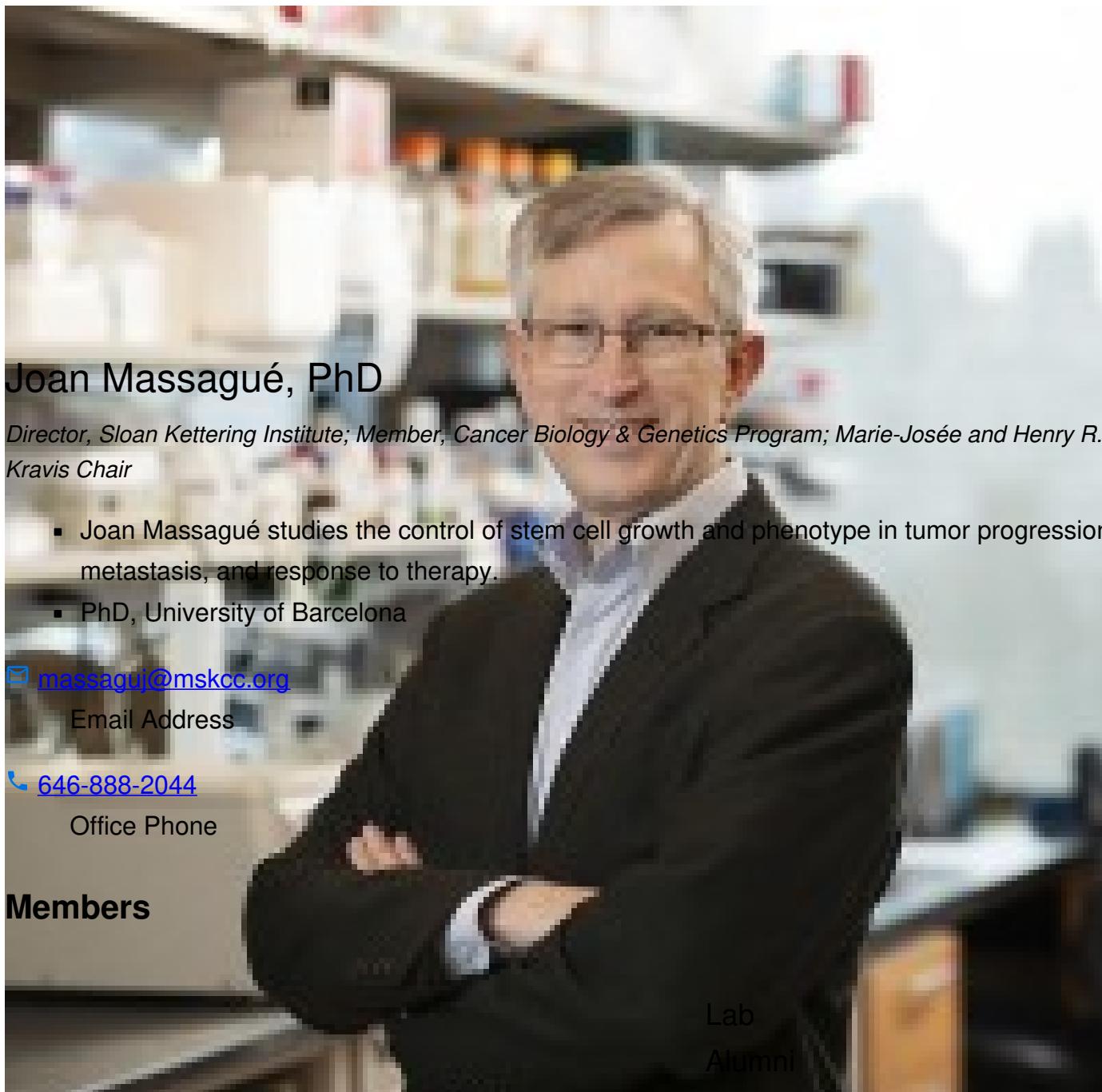
Kawasaki K, Salehi S, Zhan YA, Chen K, Lee JH, Salataj E, Zhong H, Manoj P, Kinyua D, Mello BP, Sridhar H, Tischfield SE, Linkov I, Ceglia N, Zatzman M, Havasov E, Shah NJ, Meng F, Loomis B, Bhanot UK, Redin E, de Stanchina E, Hamard PJ, Koche RP, McPherson A, Quintanal-Villalonga A, Shah SP, Massagué J* and Rudin CM* (*corresponding). FOXA2 promotes metastatic competence in small cell lung cancer. *Nature Commun.* 16:4865, (2025) [PMID: 40419484] [PMC1210783]
<https://doi.org/10.1038/s41467-025-60141-5>

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[Hu, J., Sánchez-Rivera, F.J., Johnson, G., Wang, Z., Ho, YJ., Ganesh, K., Gan, S.. Hampton, J., Zhao, H., Kottapalli, S., De Stanchina, E., Iacobuzio-Donahue, C.A., Pe'er, D., Lowe, S.W., Sun, J., and Massagué, J. STING suppresses the reactivation of dormant metastasis in lung adenocarcinoma. Nature 616\(7958\):806-813 \(2023\) PMC10569211](#)

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People



Joan Massagué, PhD

Director, Sloan Kettering Institute; Member, Cancer Biology & Genetics Program; Marie-Josée and Henry R. Kravis Chair

- Joan Massagué studies the control of stem cell growth and phenotype in tumor progression, metastasis, and response to therapy.
- PhD, University of Barcelona

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Zhenghan Wang

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Lab
Alumni

Lab Affiliations

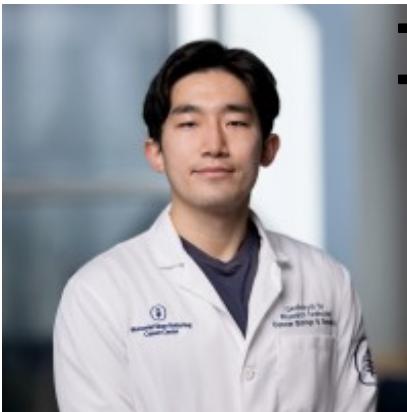
Achievements

- Chief Scientific Officer, MSKCC (2023–2025)

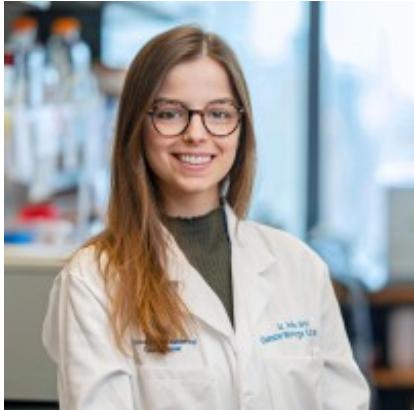
- Director, Sloan Kettering Institute (2014–)
- HHMI Scientific Review Board (2014–2025)



Derniele David
Sr. Administrative Assistant



Dechok De
Sr. Research Technician



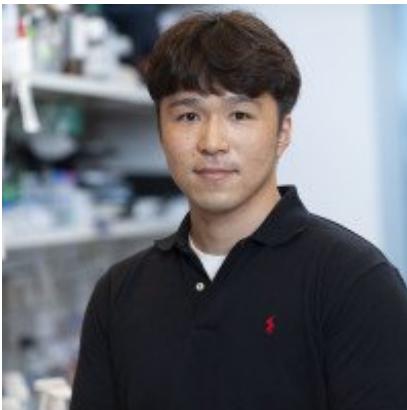
Ines Godet
Research Associate



Alhaji Janneh
Research Scholar



Carson Kenum
Research Assistant



Jun Ho Lee
Sr. Research Scientist

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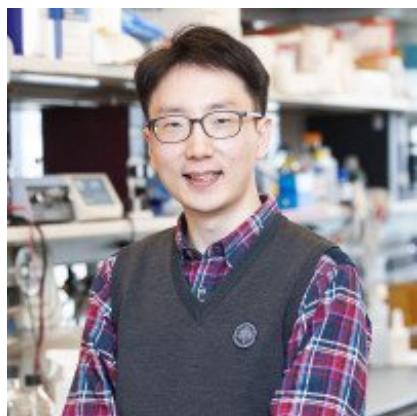
- HHMI Investigator (1990-2013)
- Fellow, American Association for Cancer Research (AACR) Academy (2016)

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Jin Suk Park
Research Associate

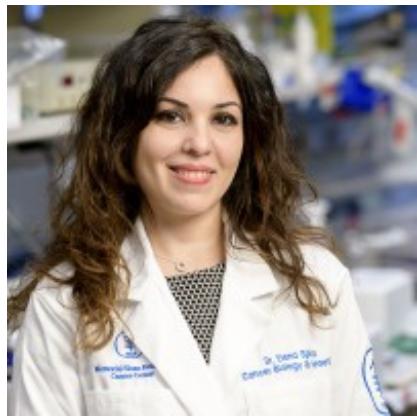


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Members of the MSK Community often work with pharmaceutical, device, biotechnology, and life

sciences companies, and other organizations outside of MSK, to find safe and effective cancer treatments, to improve patient care, and to educate the health care community. These activities outside of MSK further our mission, provide productive collaborations, and promote the practical application of scientific discoveries.

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Joan Massagué discloses the following relationships and financial interests:

- Institute for Research in Biomedicine Barcelona
 - Professional Services and Activities
- Scholar Rock
 - Equity

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