

Mesenchymal stem cells conditioning by multiple myeloma cells: translation initiation as the playing field

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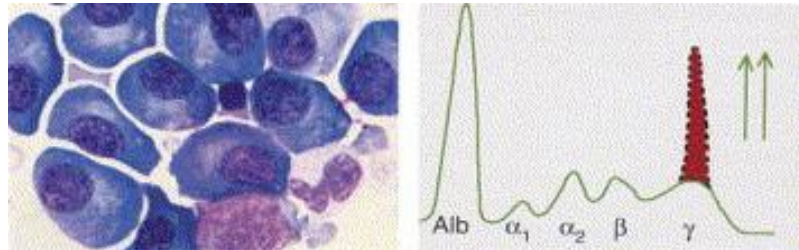
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Multiple myeloma characteristics

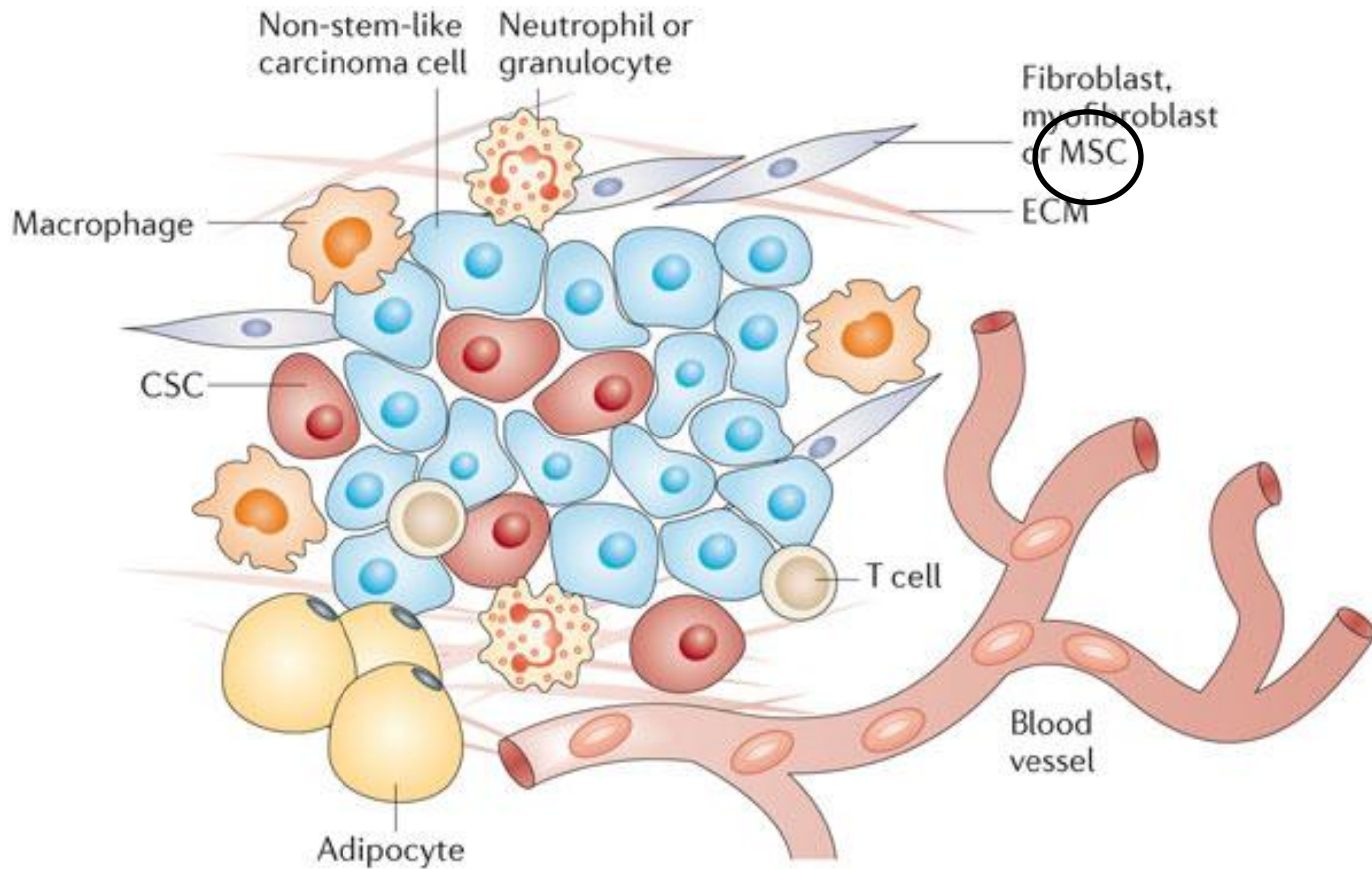
- A neoplastic disorder of plasma cells (10% of hematological malignancies)

MM remains an incurable disease

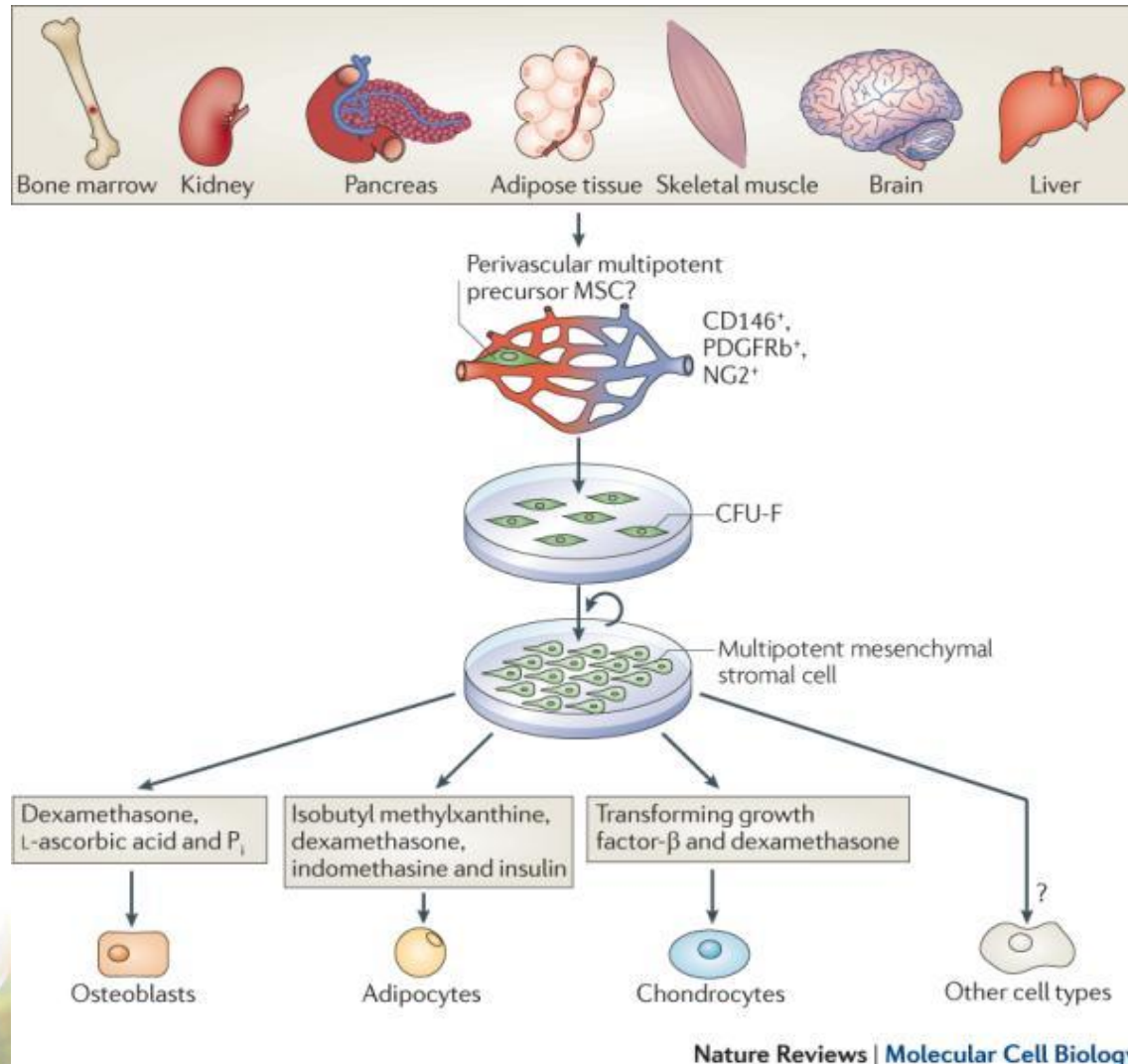


MM characterized by extensive protein synthesis.

Cancer Microenvironment

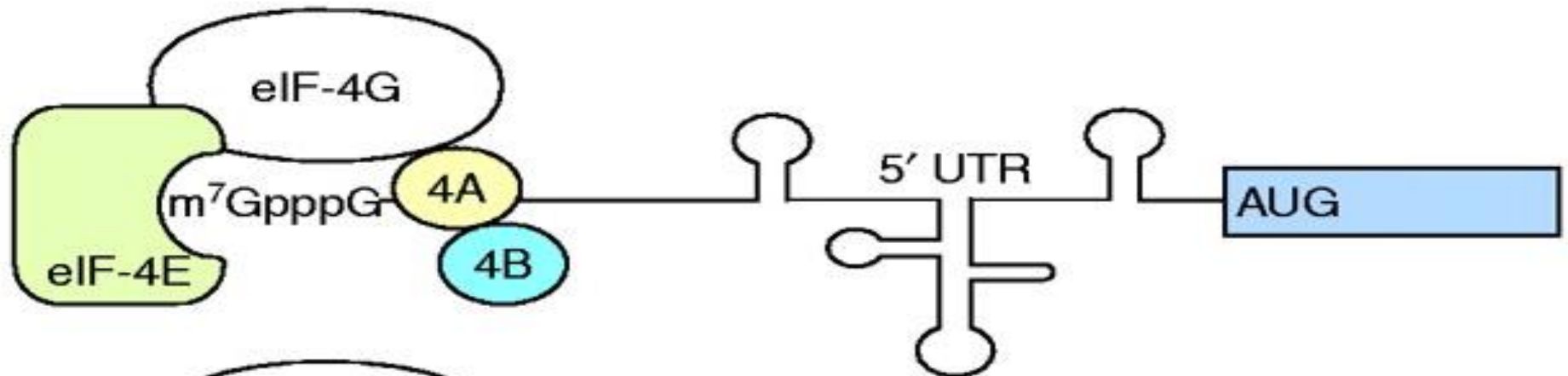


Mesenchymal Stem Cells



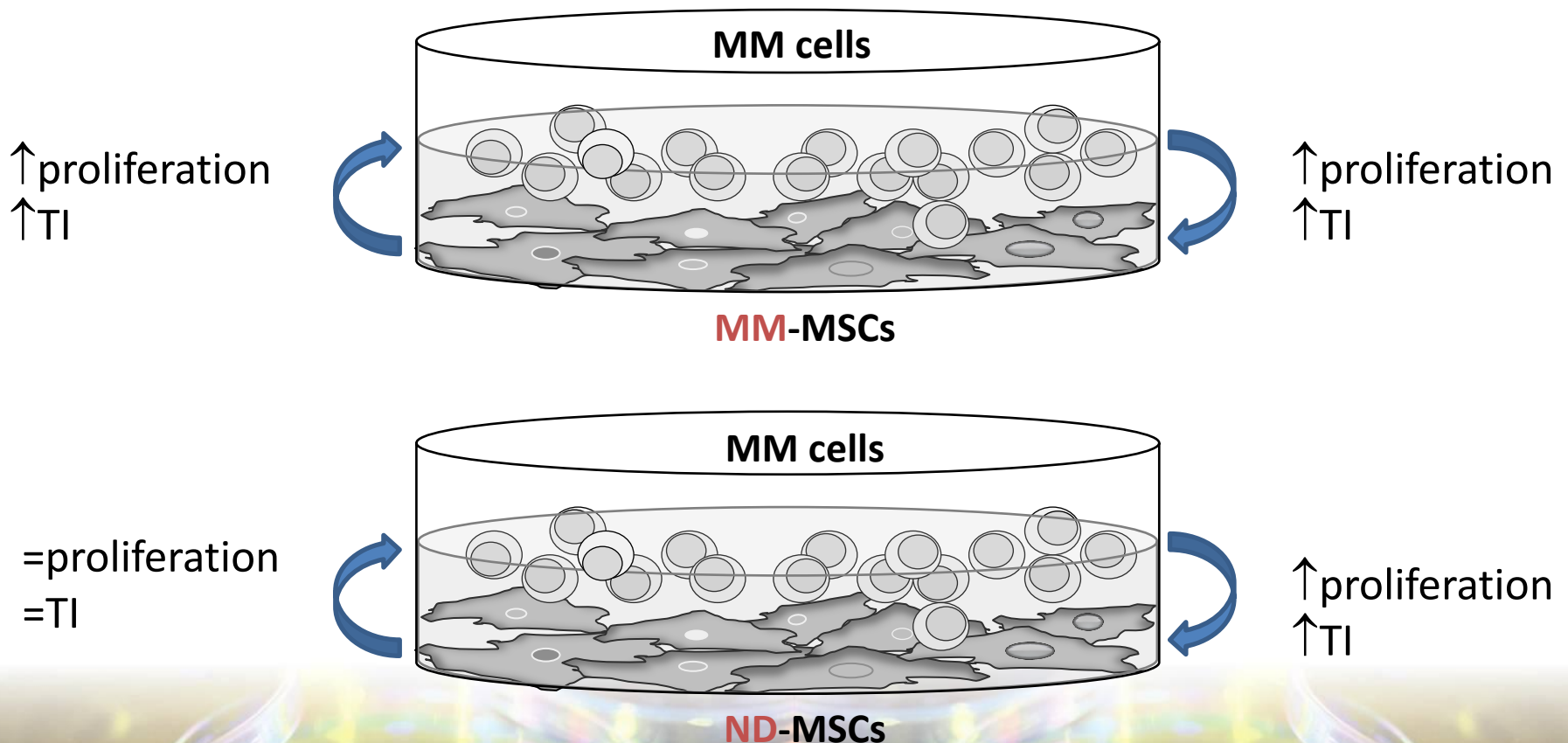
Translation initiation factors (eIF4E, eIF4G): Rate limiting stage of translation process

- Major component of the dynamic dialogue between BM-MSCs and MM cells.
- TI Regulators (MNK, mTOR, 4EBP).
- TI Targets (Smad5, NFkB, HIF1- α , CyclinD, cMyc).



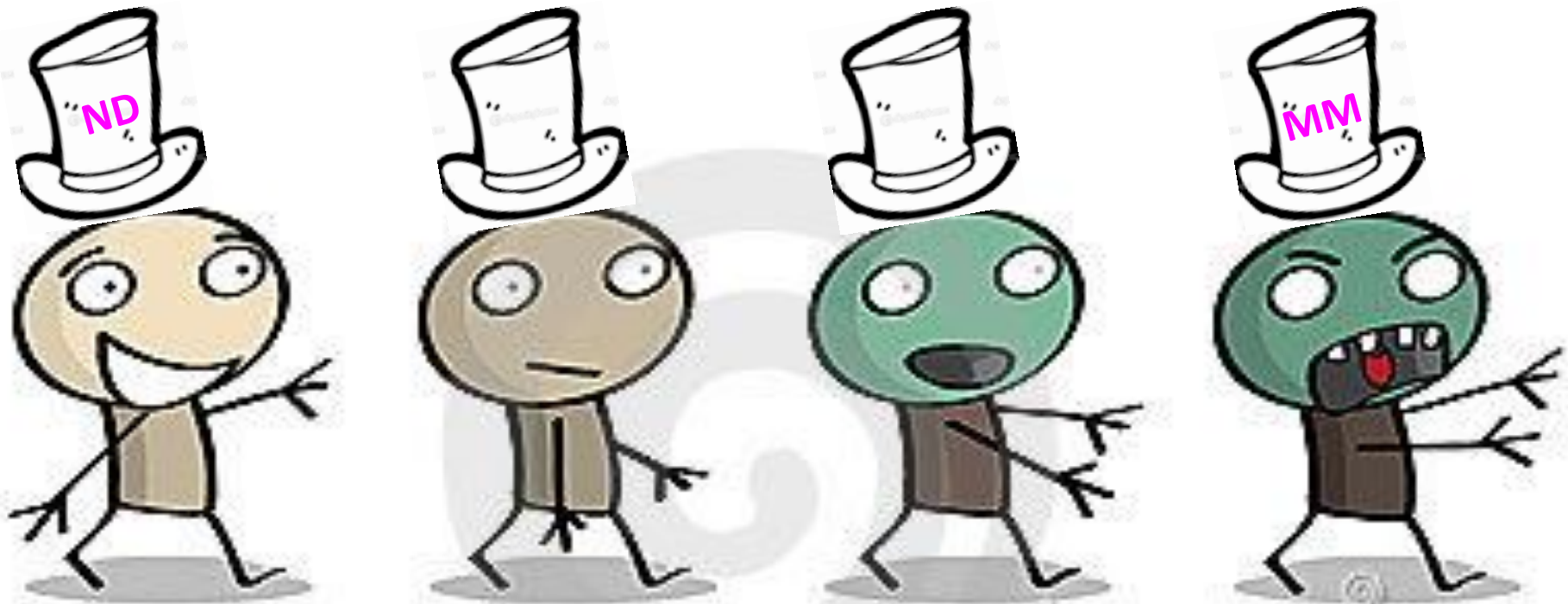
MAJOR FINDINGS

Dynamic dialogue between BM-MSCs and MM cell lines



ND-MSCs transformation

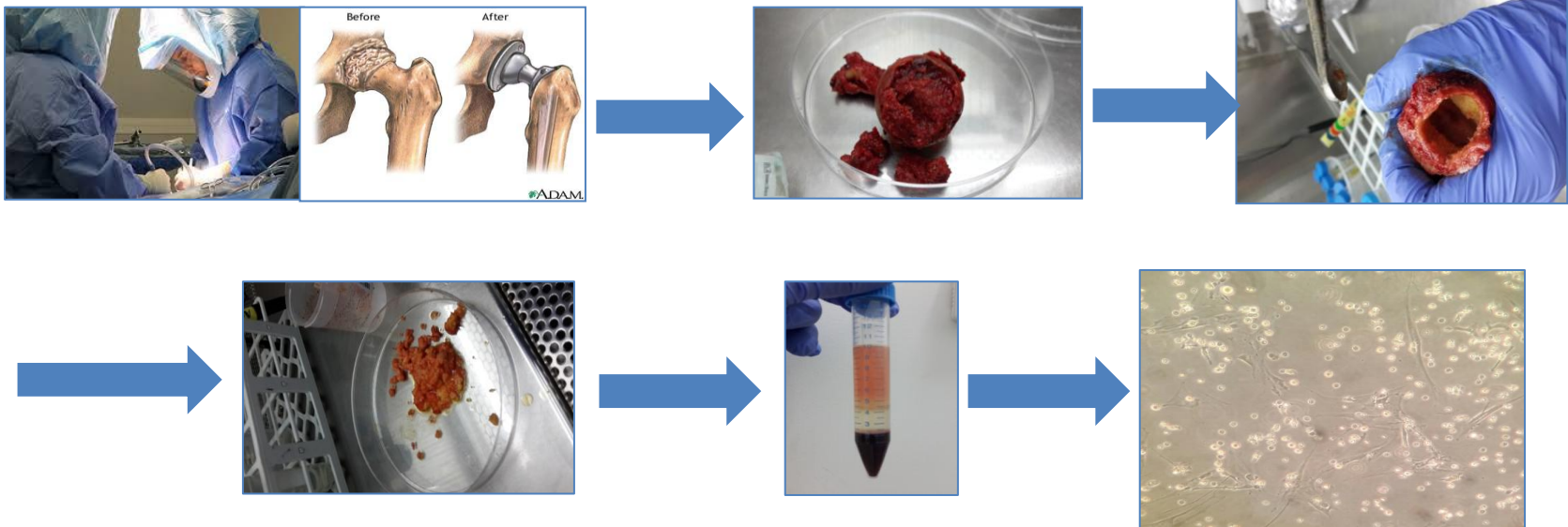
- My aim was to explore the changes that ND-MSCs undergo in MM proximity



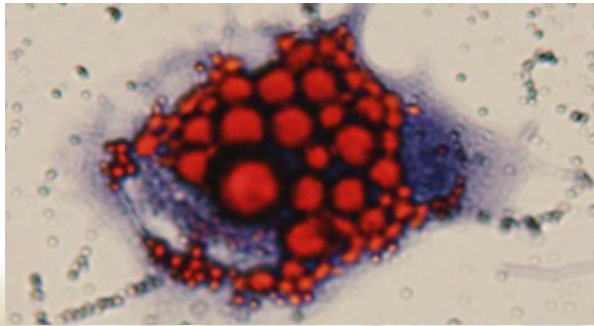
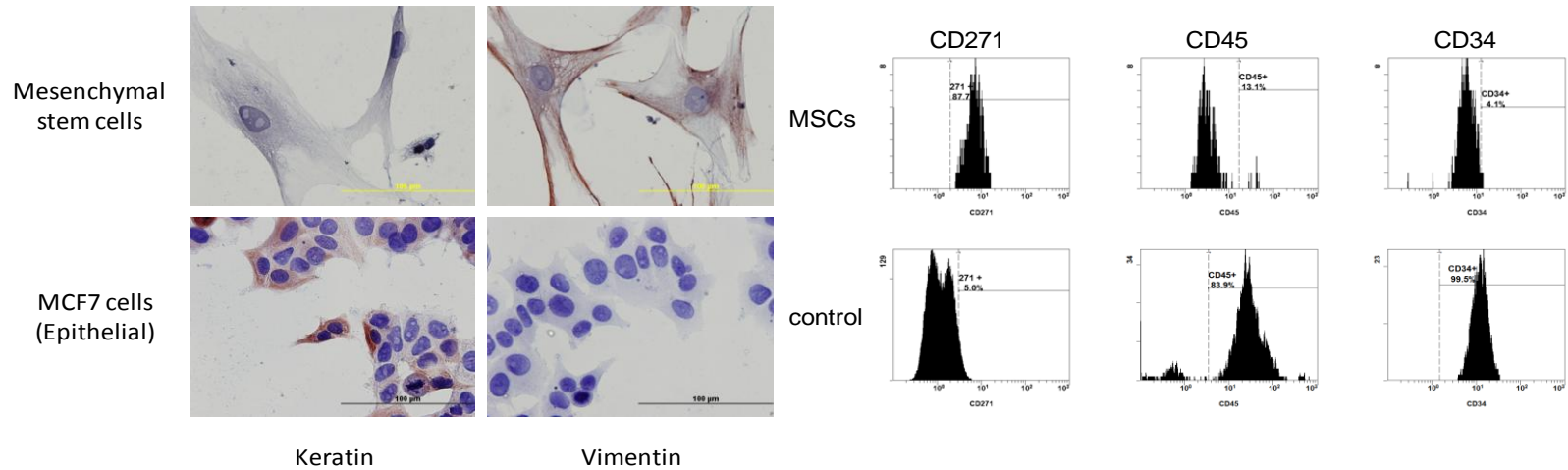
Results..



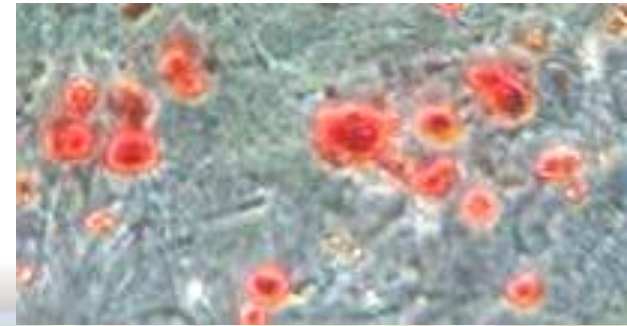
ND-MSCs extraction model



ND-MSCs characterization

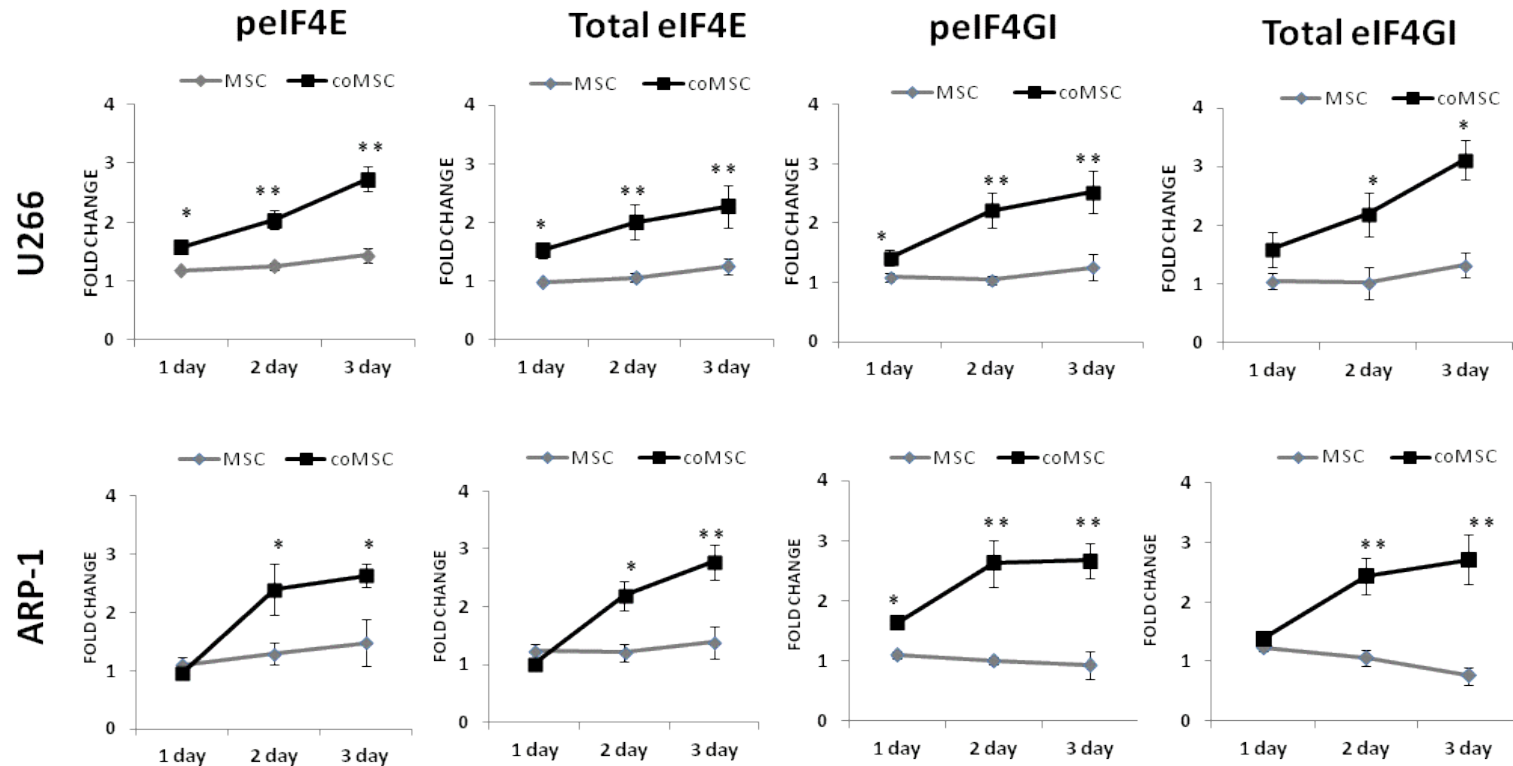


Adipocytes



Osteoblasts

Translation initiation factors are increased in MMcond-MSCs: Timing

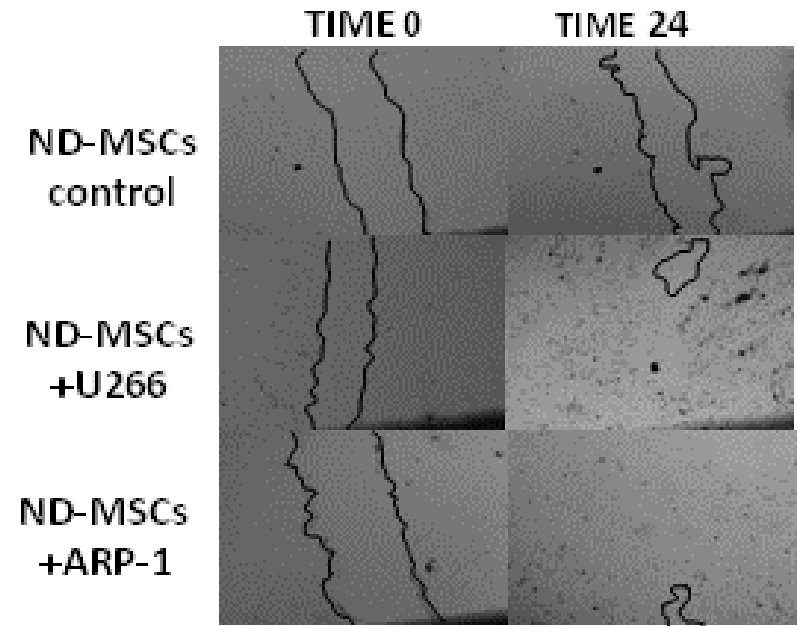
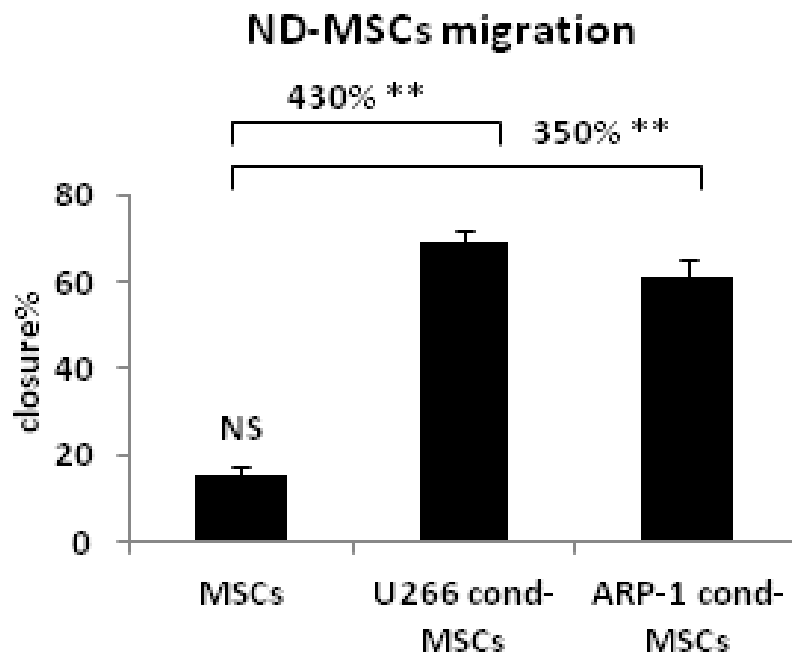


Plato at 4 and 5 days.

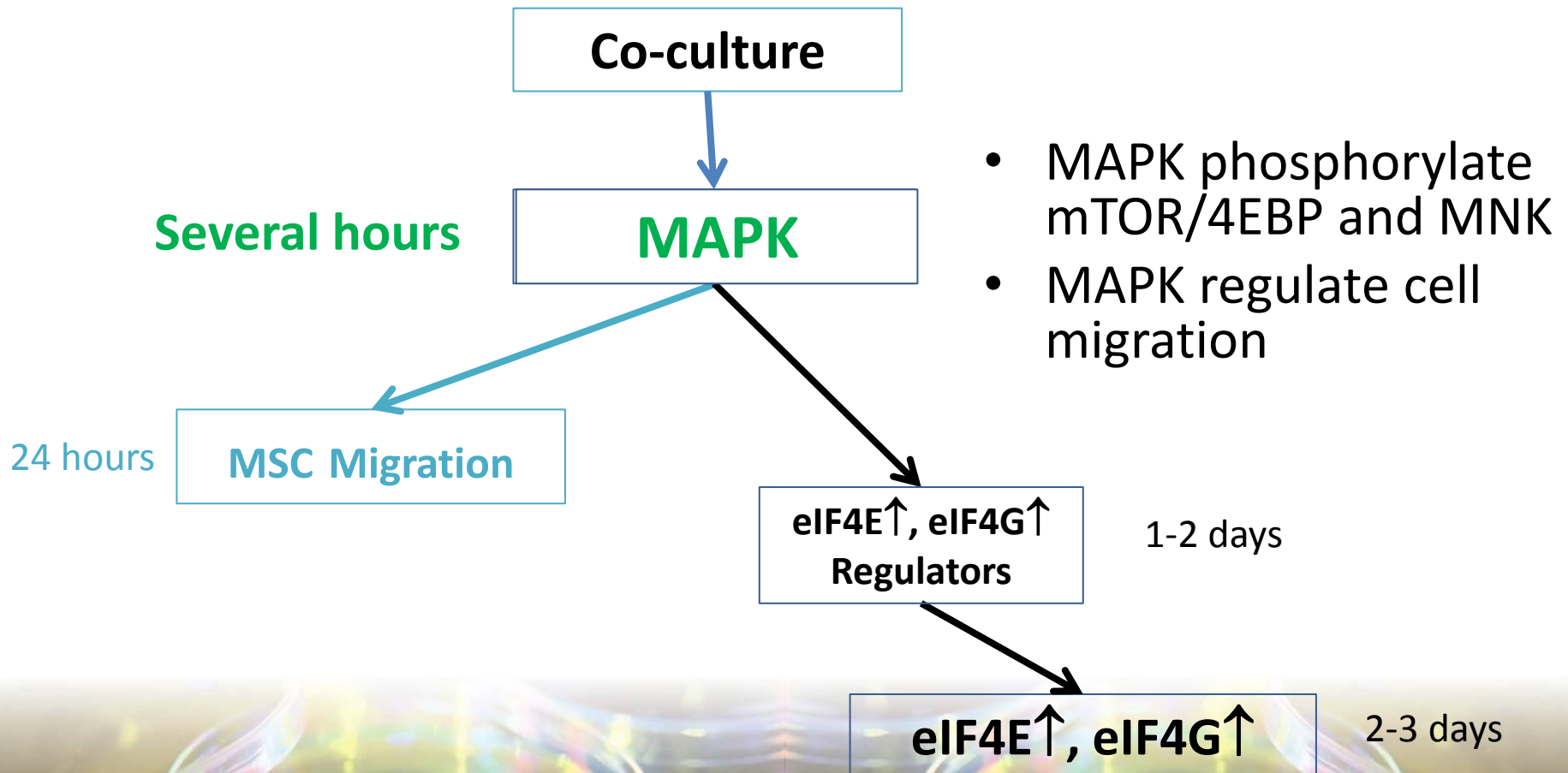
eIF4E, eIF4G regulators were elevated at the 1st and the 3rd days.

eIF4E, eIF4G targets were elevated at the 3rd day.

MMcond-MSCs display elevated migration (24h)

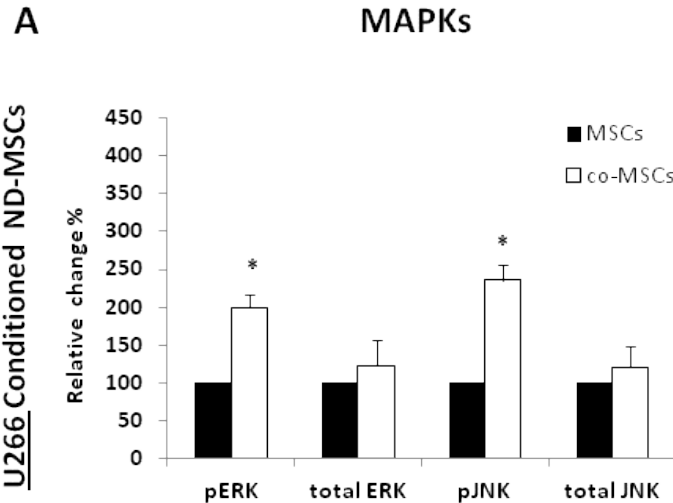


MMcond-MSCs possible signals

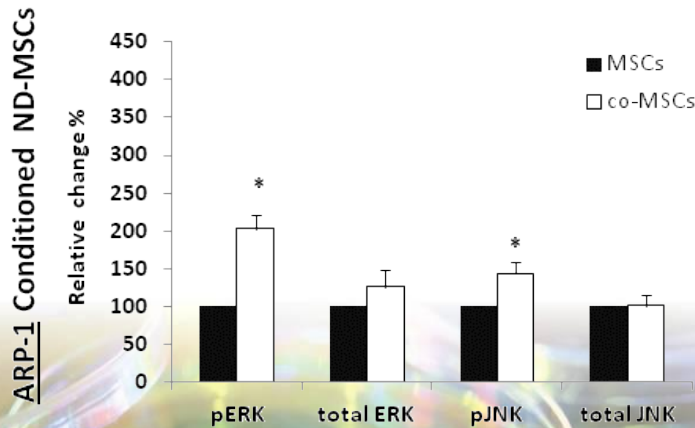


MMcond-MSCs show early elevated MAPKs signals and TI factors phosphorylation (1.5h)

U266



ARP-1

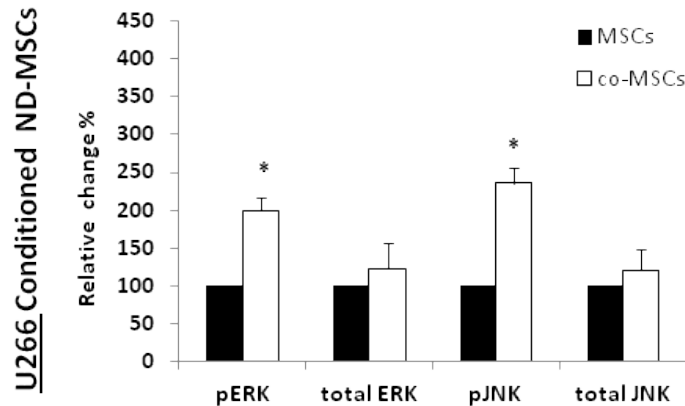


MMcond-MSCs show early elevated MAPKs signals and TI factors phosphorylation (1.5h)

U266

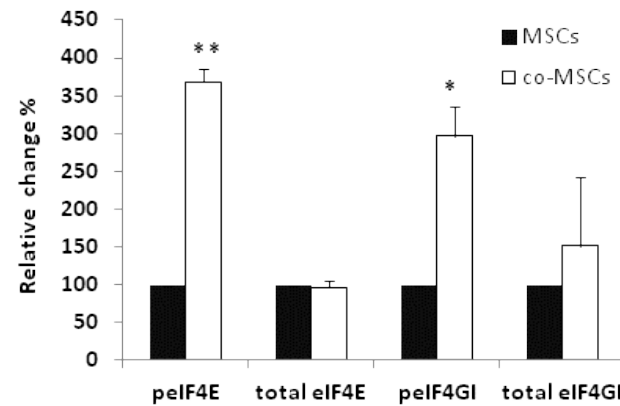
A

MAPKs



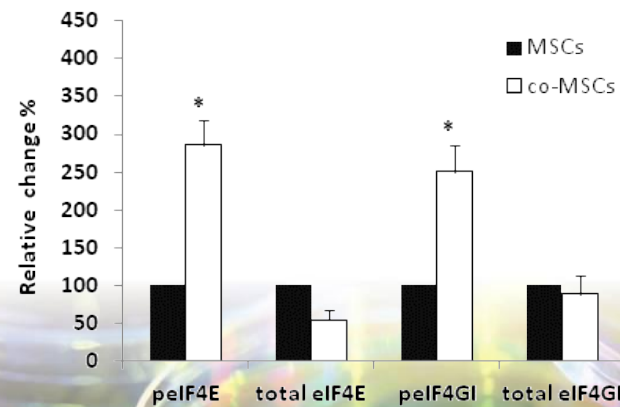
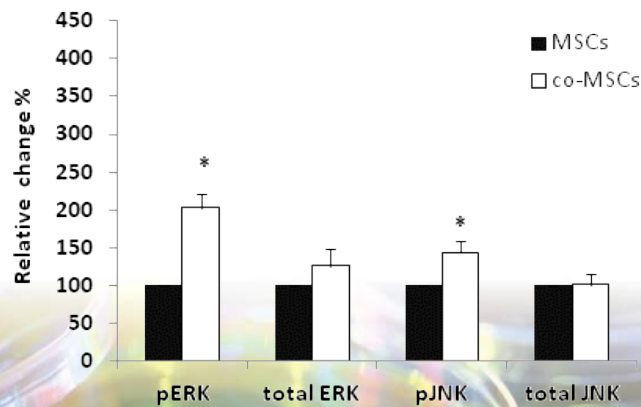
B

Translation initiation factors



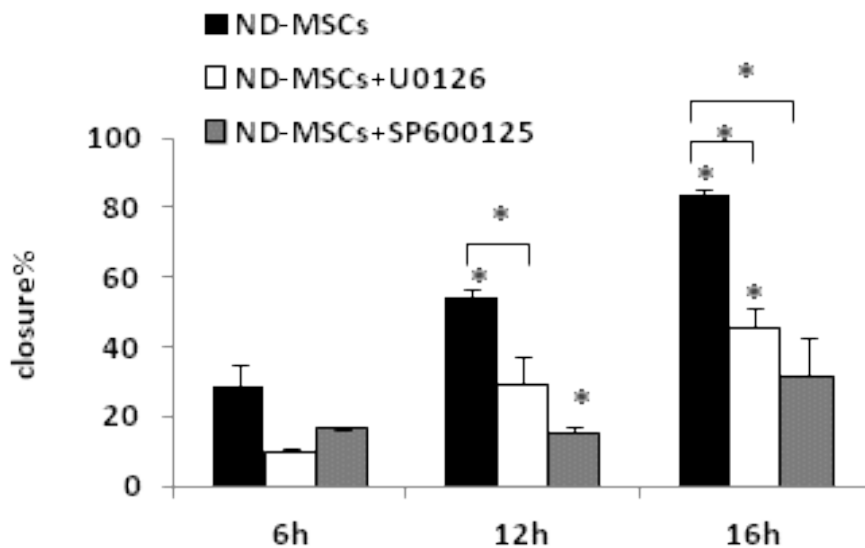
ARP-1

ARP-1 Conditioned ND-MSCs

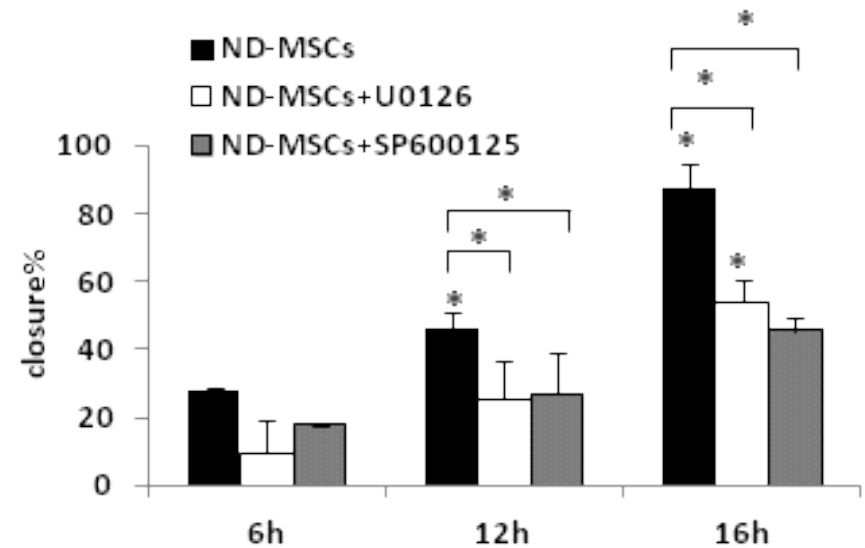


MAPKs inhibitors decrease MMcond- MSCs elevated migration (16h)

ARP-1 Co-cultured ND-MSCs treated with MAPK inhibitors

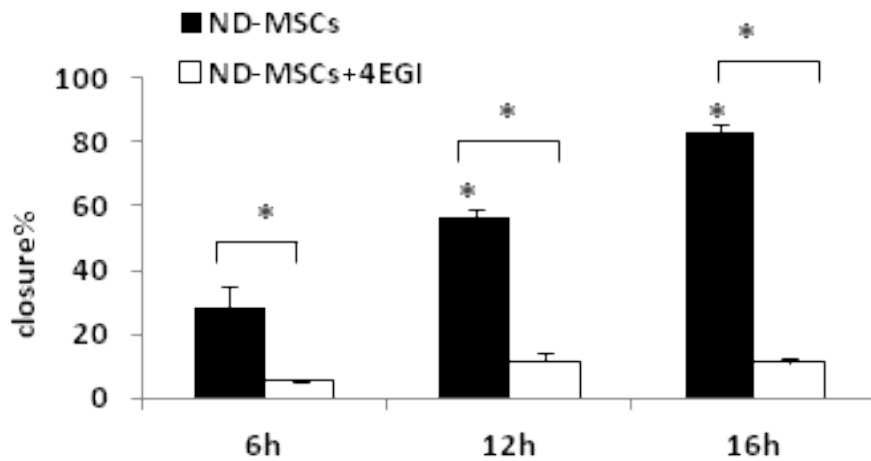


U266 Co-cultured ND-MSCs treated with MAPK inhibitors

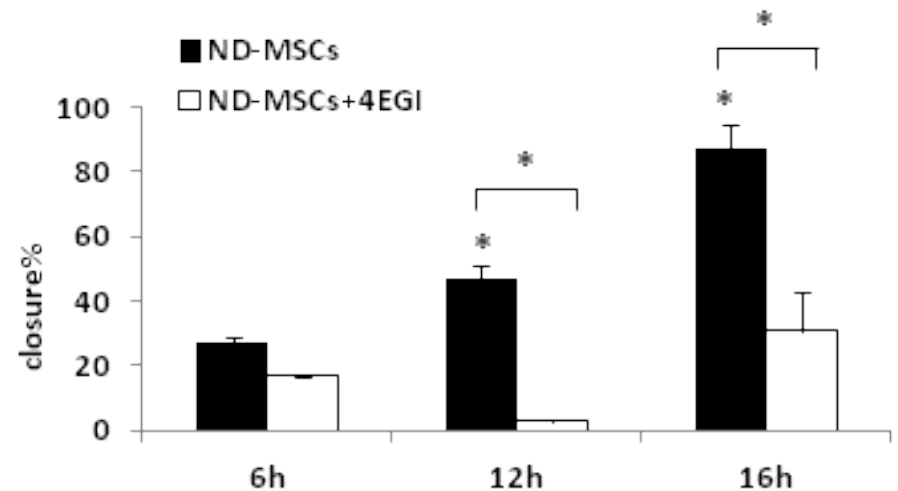


4EGI decreases MMcond-MSCs elevated migration (16h)

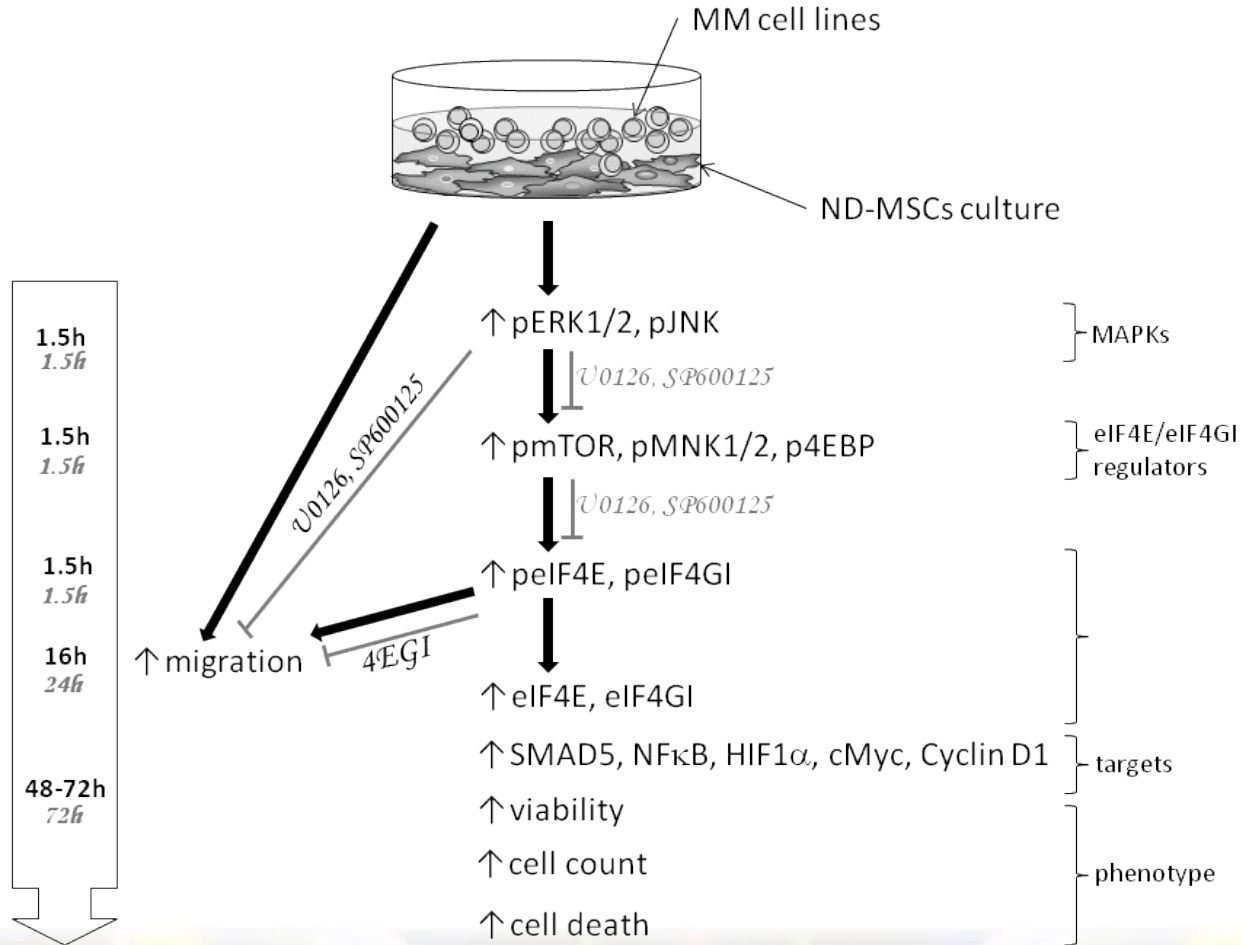
ARP-1 Co-cultured ND-MSCS with 4EGI



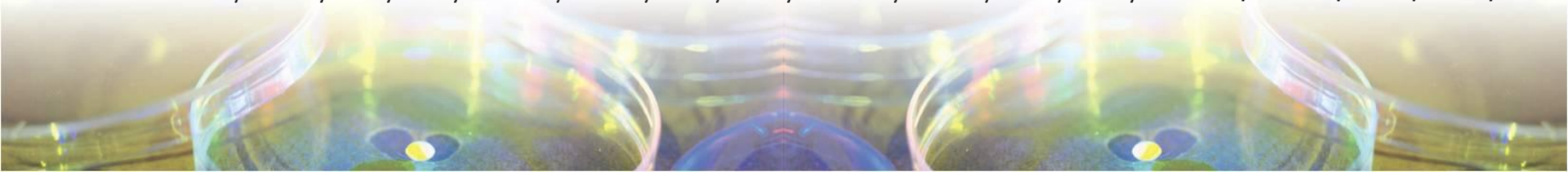
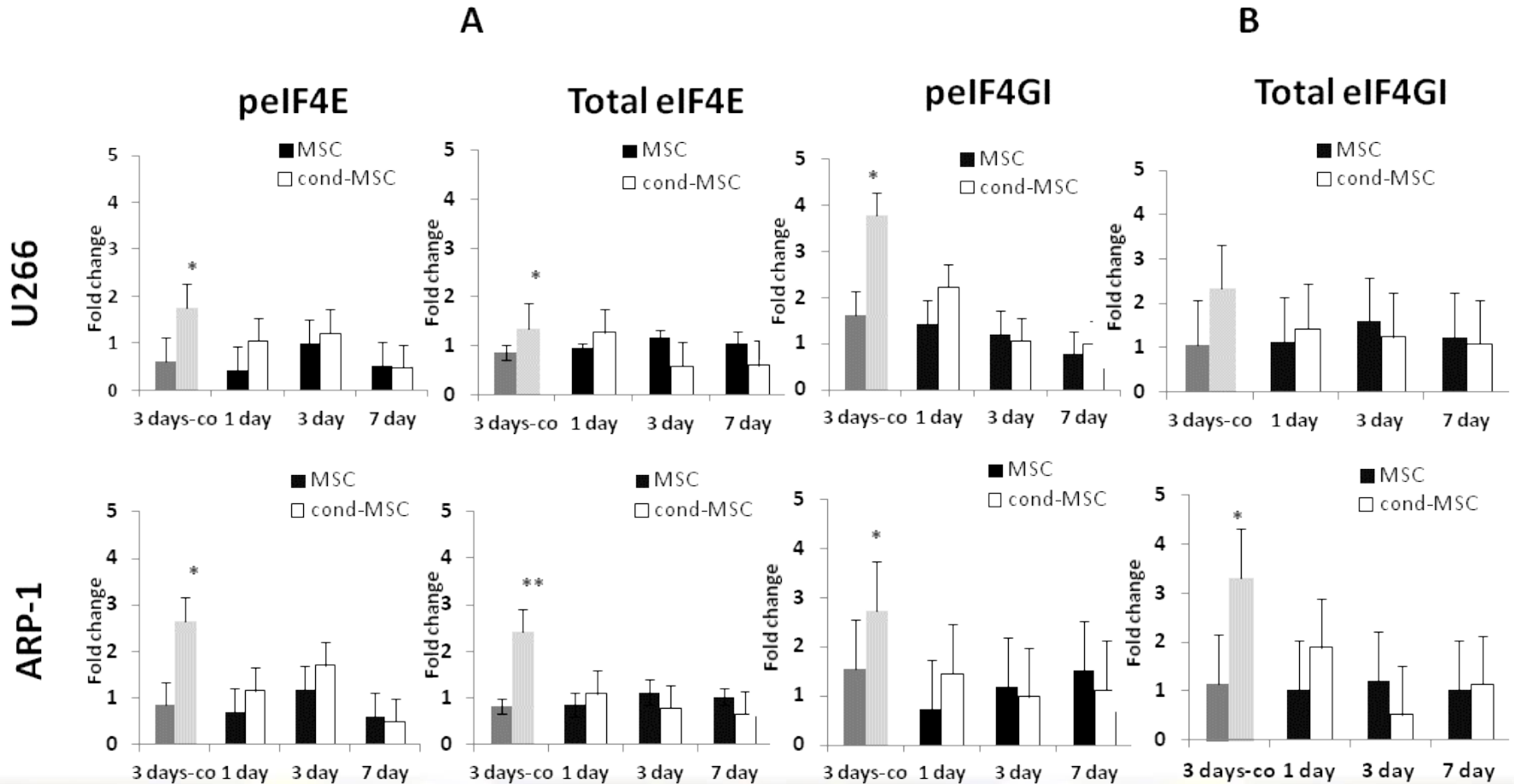
U266 Co-cultured ND-MSCS with 4EGI



Results summary

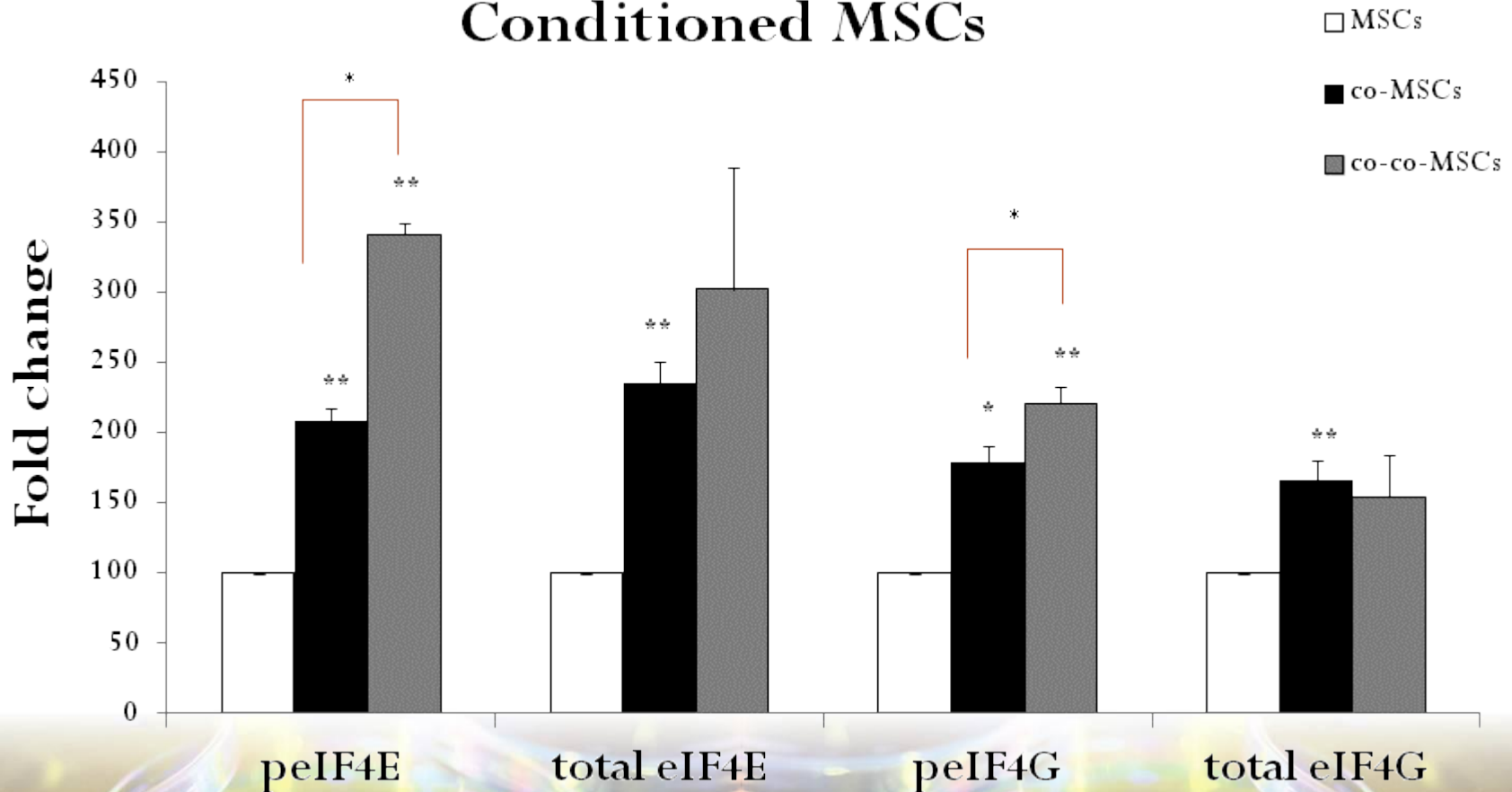


The increase in TI in MMcond-MSCs is reversible



Reconditioning of MMcond-MSCs display another elevation of TI factors

Conditioned MSCs



Results summary

- ND-MSCs exposed to MM cell lines undergo significant changes in protein synthesis and repertoire as well as migration.
- The changes are time dependent, reversible and can be exacerbated by reintroduction of “fresh MM”.
- Co-culture affected signaling cascade:
MAPK/eIF4E and eIF4G1/migration and targets.



Significance of my results

- *In vivo*: constant and dynamic refreshment of the niche with new MM cells: in actuality- non reversible and even enhanced ND-MSCs conditioning.
- Increased migration may facilitate the tropism of BM-MSCs to the tumor. Intervention may sabotage the MM niche.
- eIF4E and eIF4G1 are important for BM-MSCs protein repertoire and migration, underscoring their potential as therapeutic targets.



Submitted for publication..

Multiple myeloma cells reprogram bone marrow mesenchymal stem cells' translation initiation thereby promoting their migration

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Thank you

Professor Michael Lishner
Dr. Liat Drucker

Orthopedic department
Surgery rooms
Hematocytological laboratory

Dr. Shelly Tartakover Matalon
Dr. Oshrat Attar Schneider
Dr. Gali Epstein Shochet
Oded Komemi
Michal Bar

