"Cellular heterogeneity as predictor for the differentiation and regenerative potential of bone marrow stromal cells"

The Danish 3R Symposium 2021

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Molecular Endocrinology Unit (KMEB),

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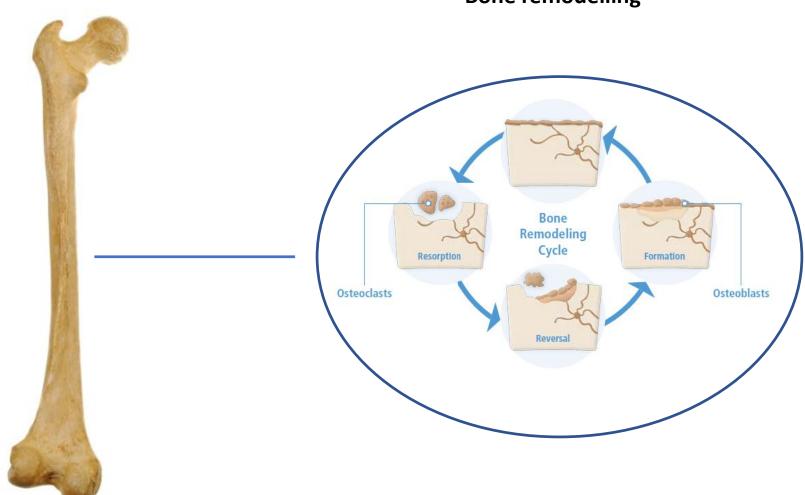








Bone remodelling



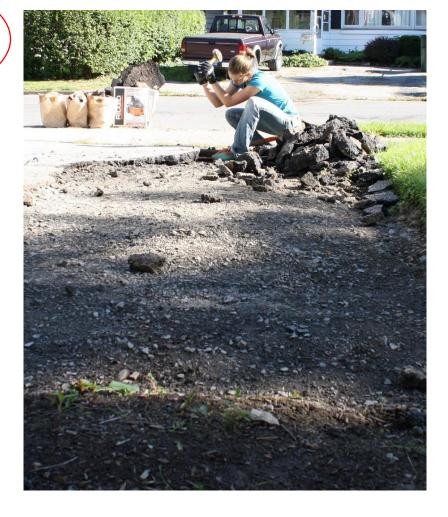








Step 1: Remove







Step 1: Remove

Step 2: Pour new







Step 1: Remove



Step 2: Pour new











Step 1: Remove



Osteo<u>clast</u> cells

Step 2: Pour new



Osteo**blast** cells

New bone





Old bone



Step 1: Remove



Osteo<u>clast</u> cells

p 1: Step 2: nove Pour new



Osteo**blast** cells

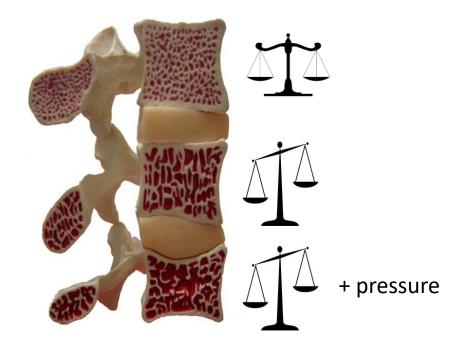


New bone



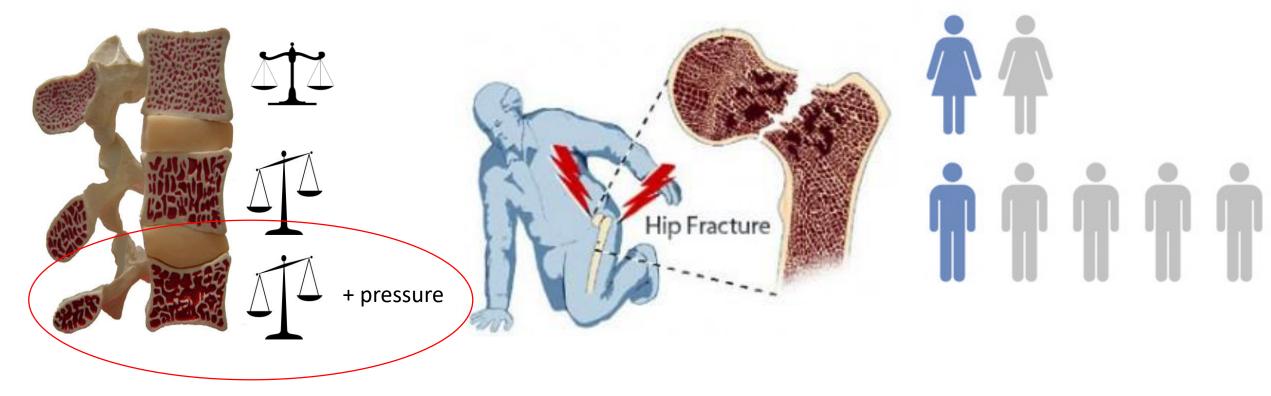


Osteoporosis



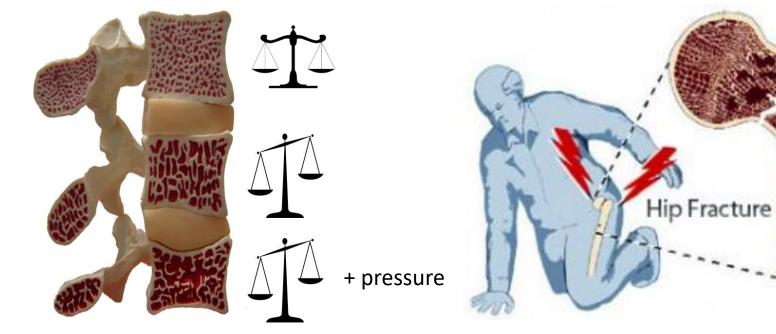


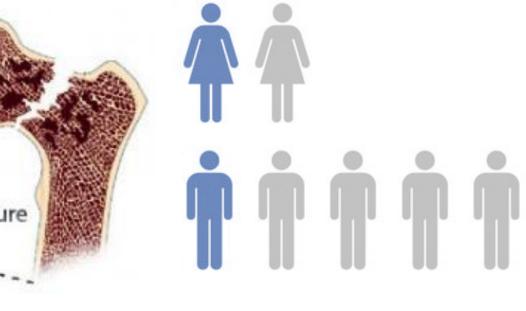
Osteoporosis



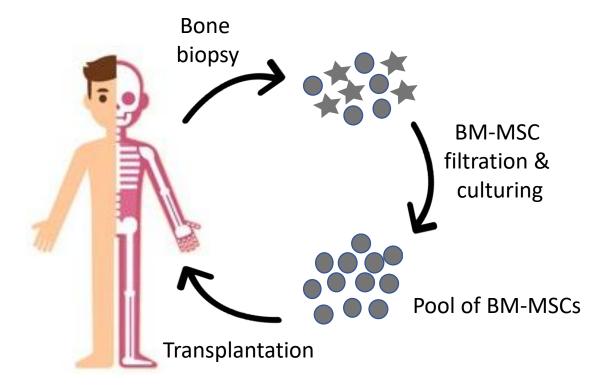


Osteoporosis

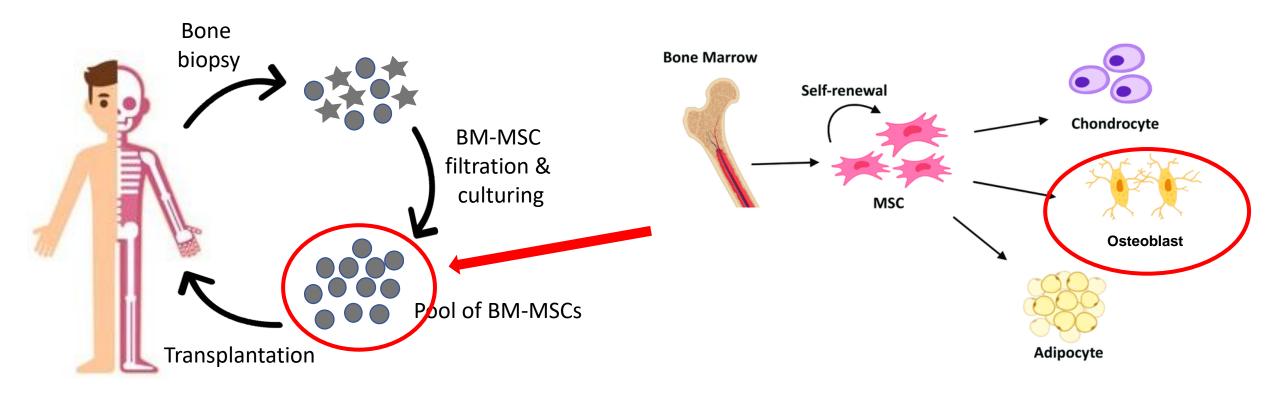




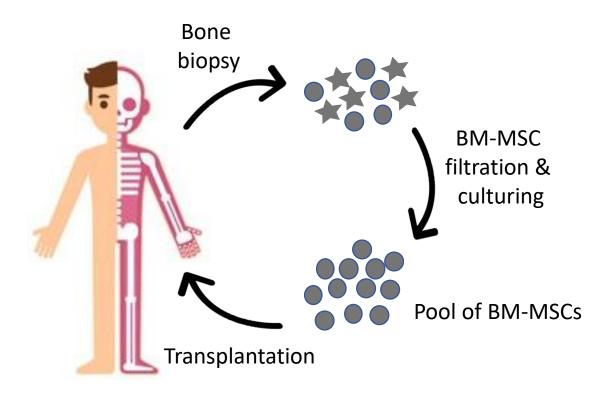


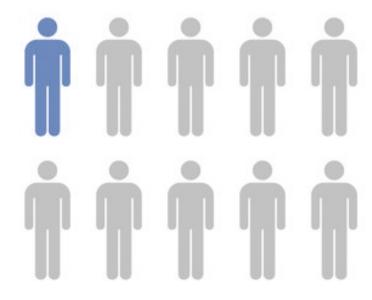




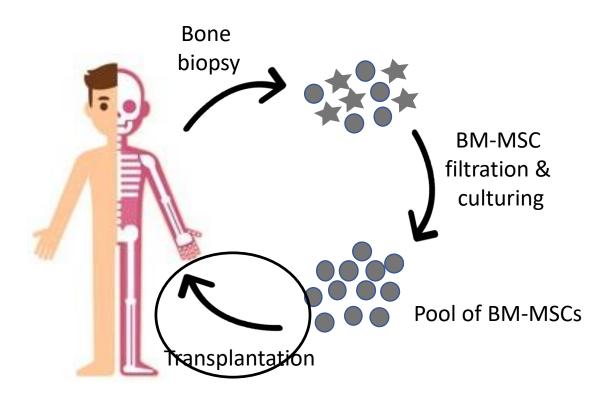


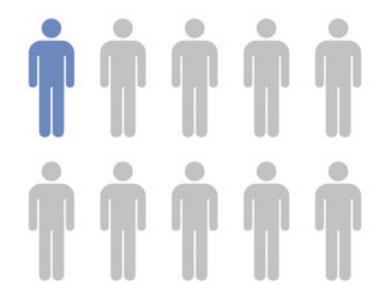














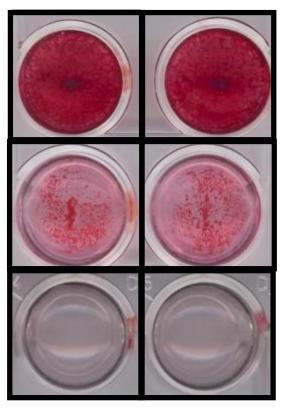
Cohort size = 56

Donor examples:

Donor 1

Donor 2

Donor 3















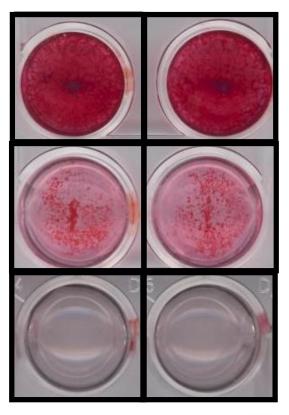
Cohort size = 56

Donor examples:

Donor 1

Donor 2

Donor 3





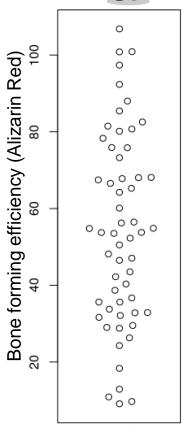
















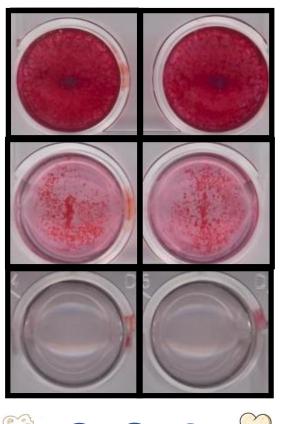
Cohort size = 56

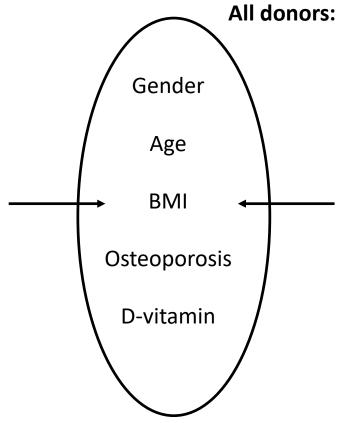
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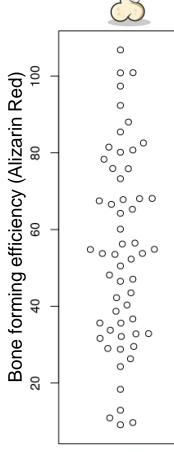
Donor 1

Donor 2

Donor 3





















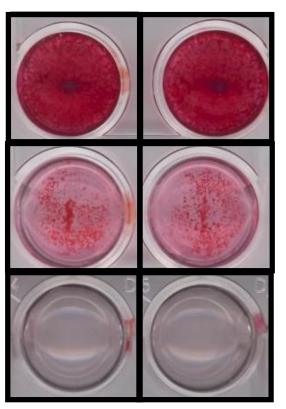
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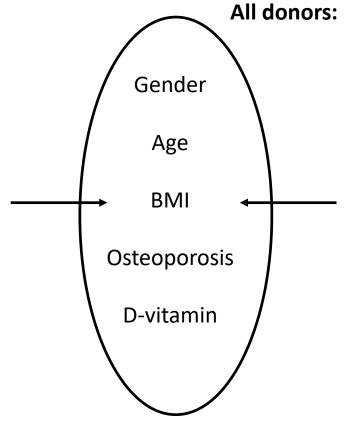
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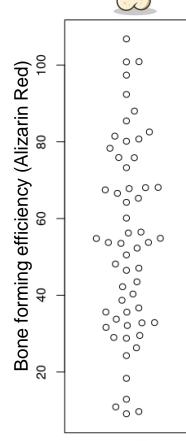
Donor 1

Donor 2

Donor 3











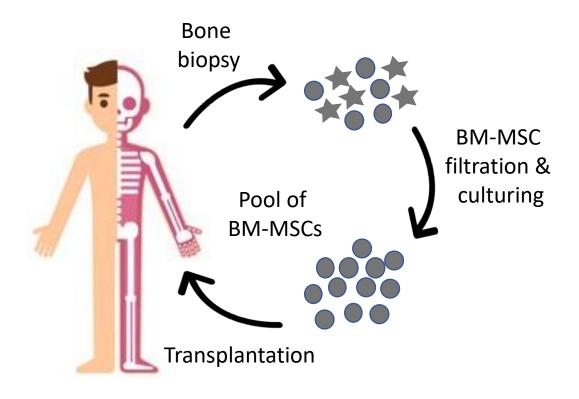




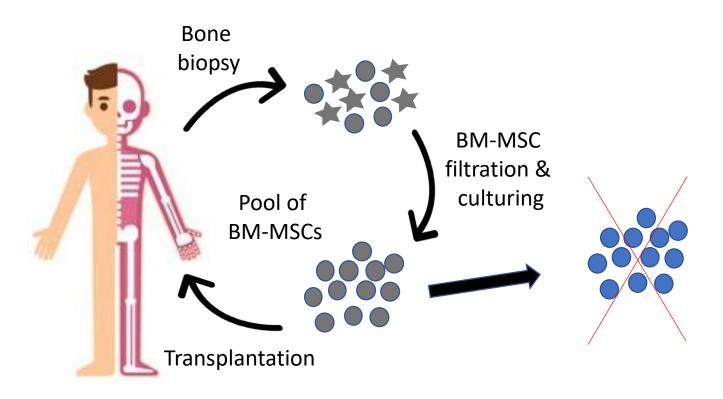


50% potency explained

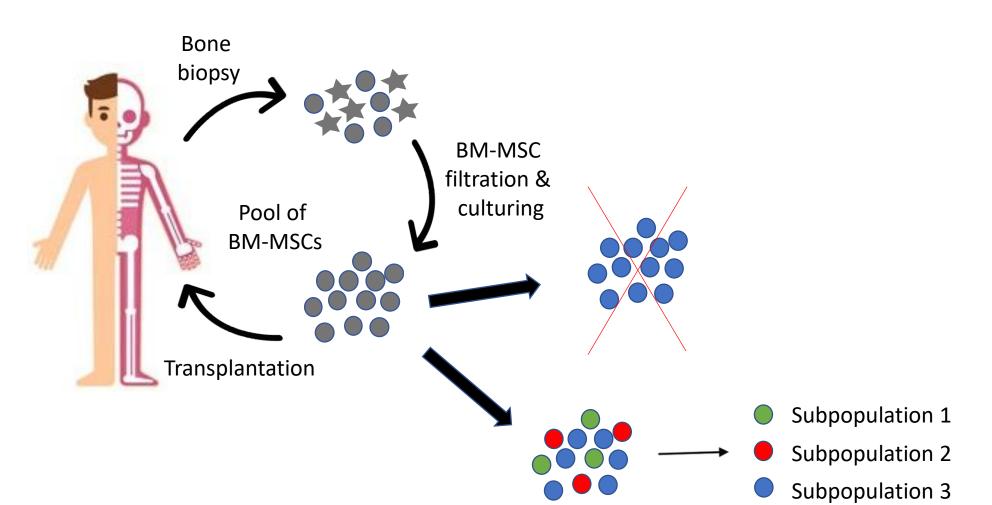














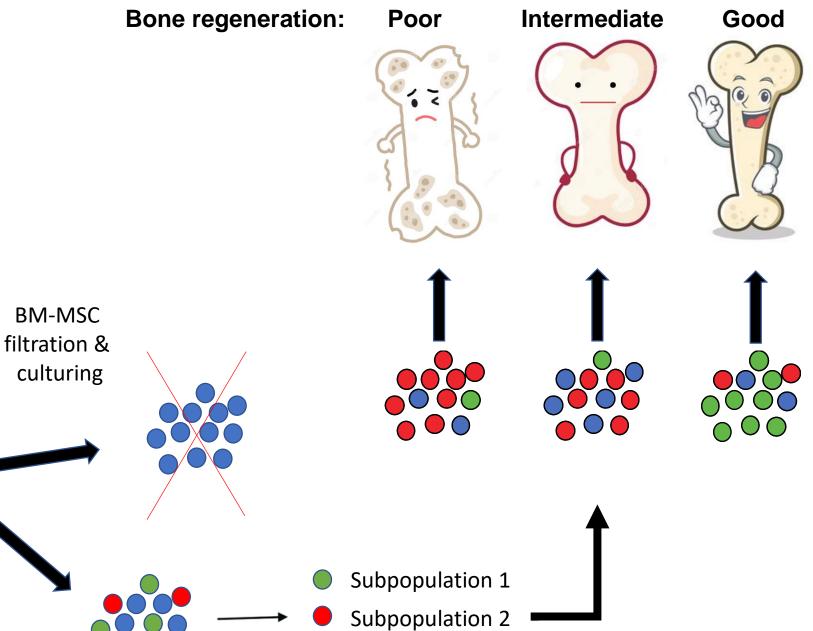
Bone

biopsy

Pool of

BM-MSCs

Transplantation

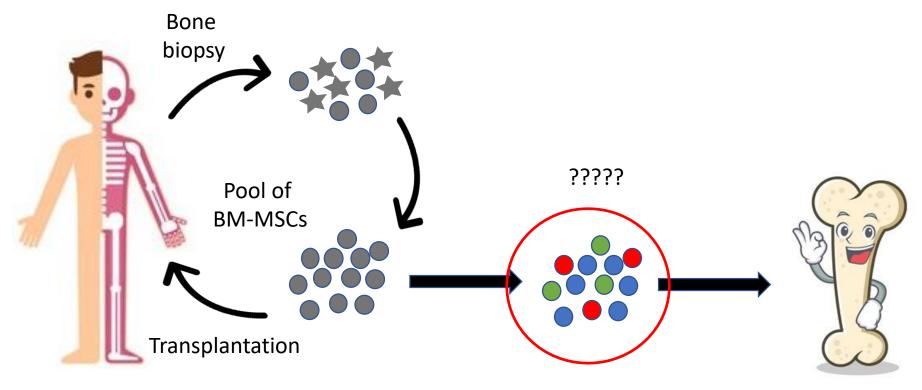


Subpopulation 3



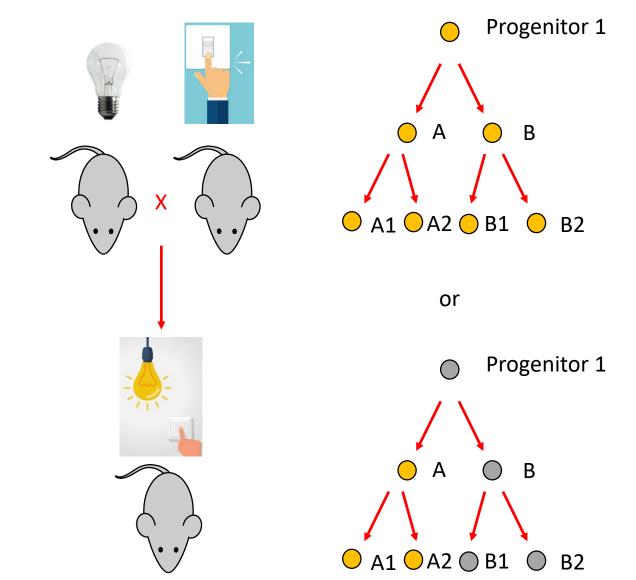
Overall aim

To investigate how cellular heterogeneity within patient-derived BM-MSCs affect osteogenic differentiation and bone tissue regeneration potential.





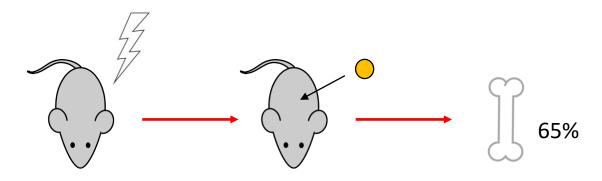
Lineage tracing

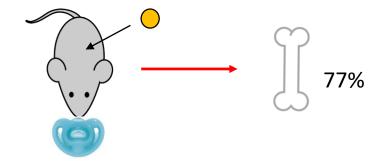




Lineage tracing - verification

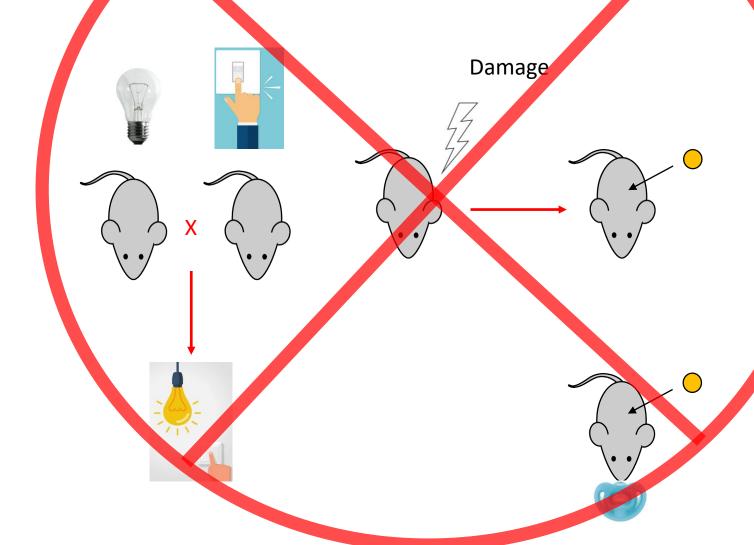
Damage







Lineage tracing verification





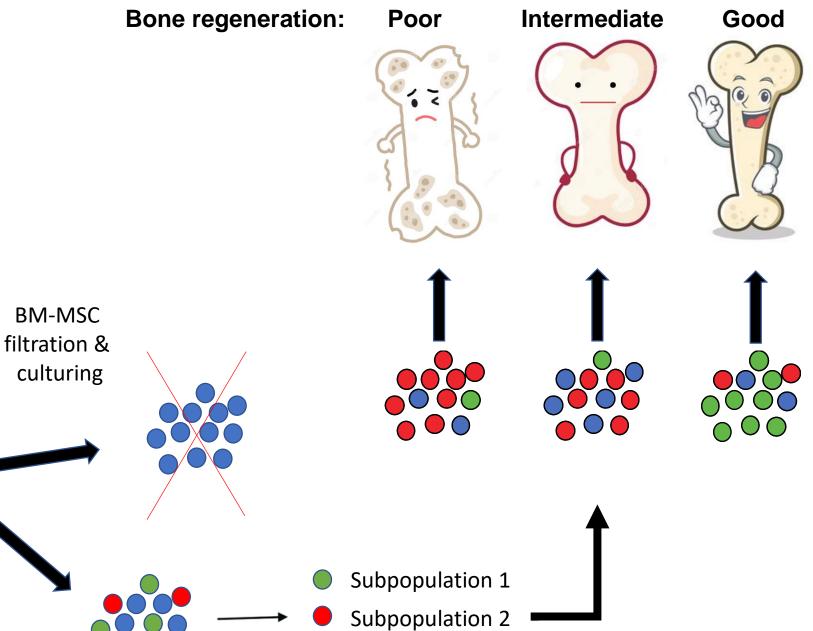
Bone

biopsy

Pool of

BM-MSCs

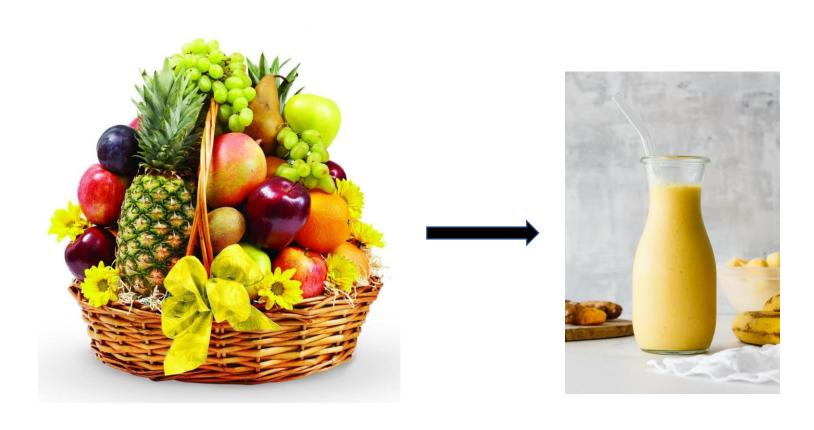
Transplantation



Subpopulation 3

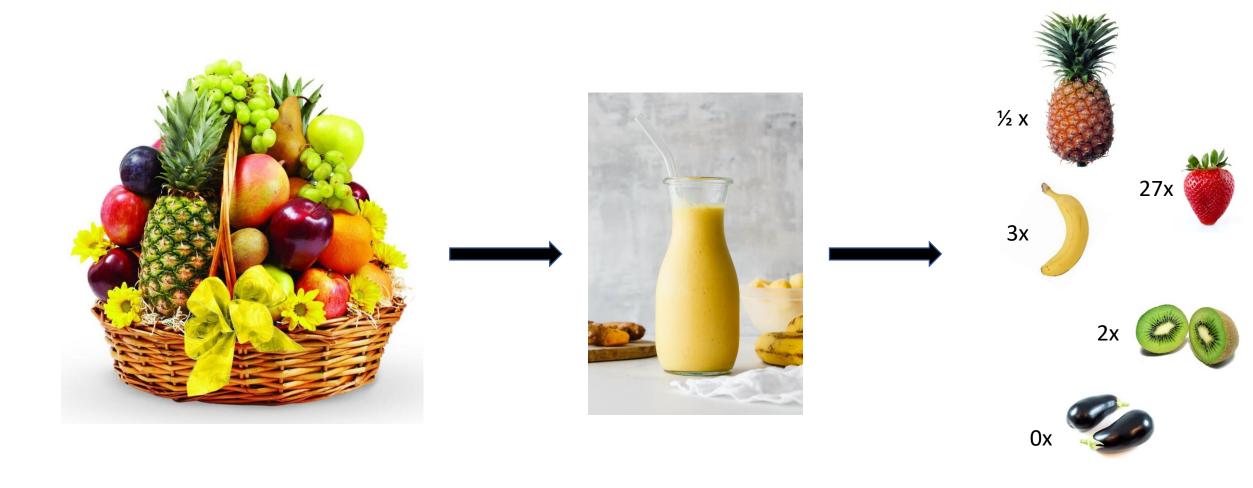


scRNA-sequencing

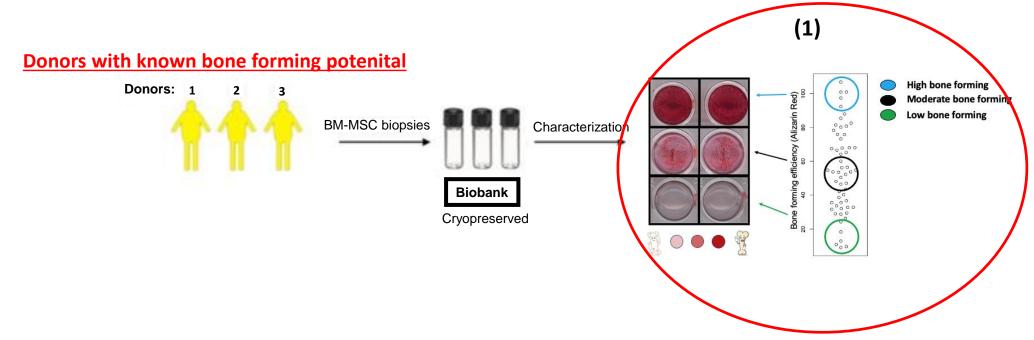




scRNA-sequencing



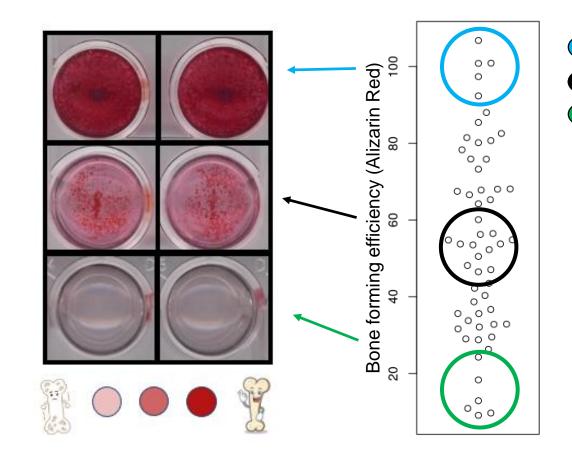




Practicalities:

1) Sample selection





High bone formingModerate bone forming

Low bone forming

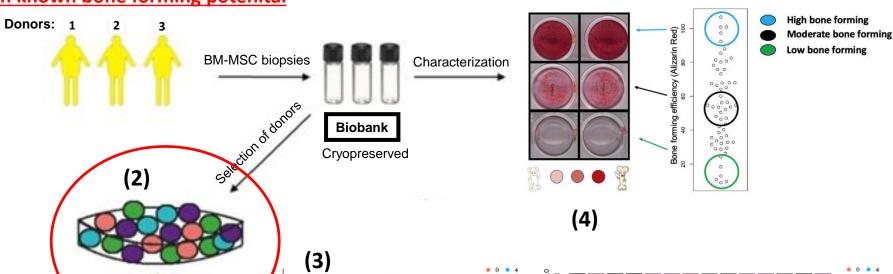
Practicalities:

1) Sample selection



Donors with known bone forming potenital

scRNA-seq



UMAP 1

(1)

Donor samples

Practicalities:

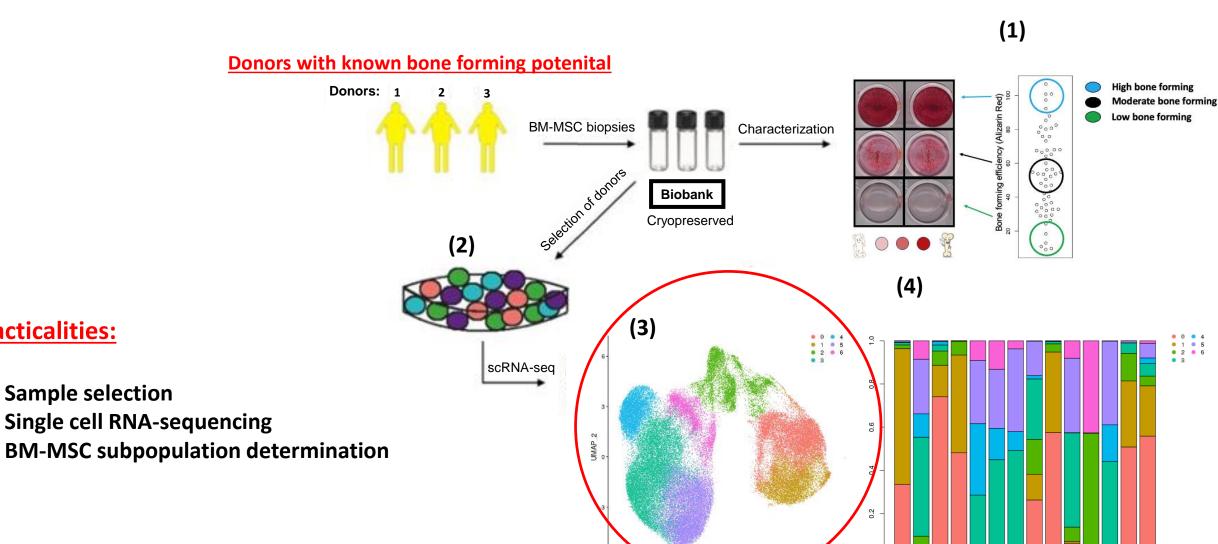
- 1) Sample selection
- 2) Single cell RNA-sequencing



Practicalities:

Sample selection

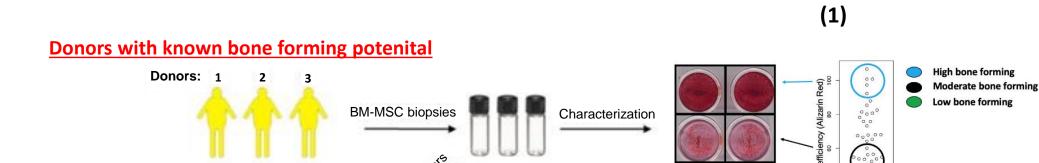
Study design - identification



Donor samples

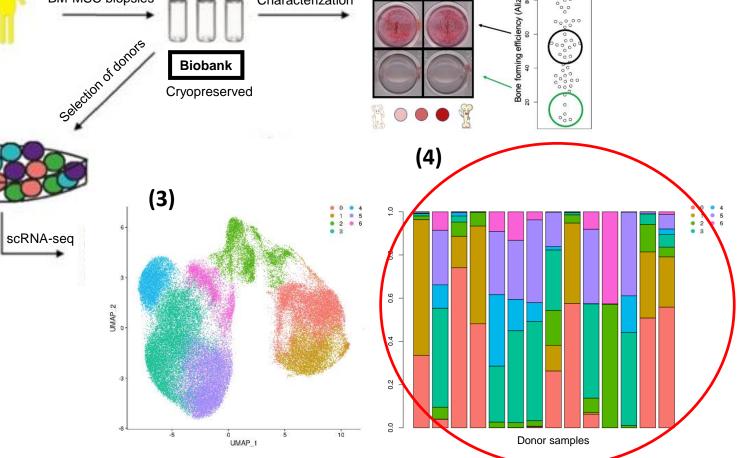


(2)

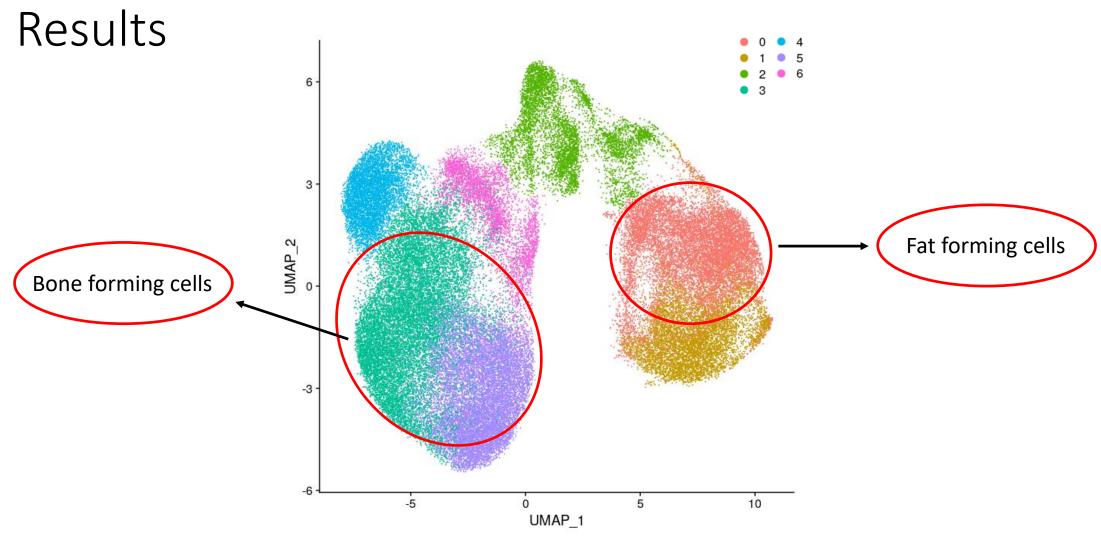


Practicalities:

- 1) Sample selection
- 2) Single cell RNA-sequencing
- B) BM-MSC subpopulation determination
- 4) BM-MSC subpopulation abundancies quantification

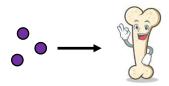








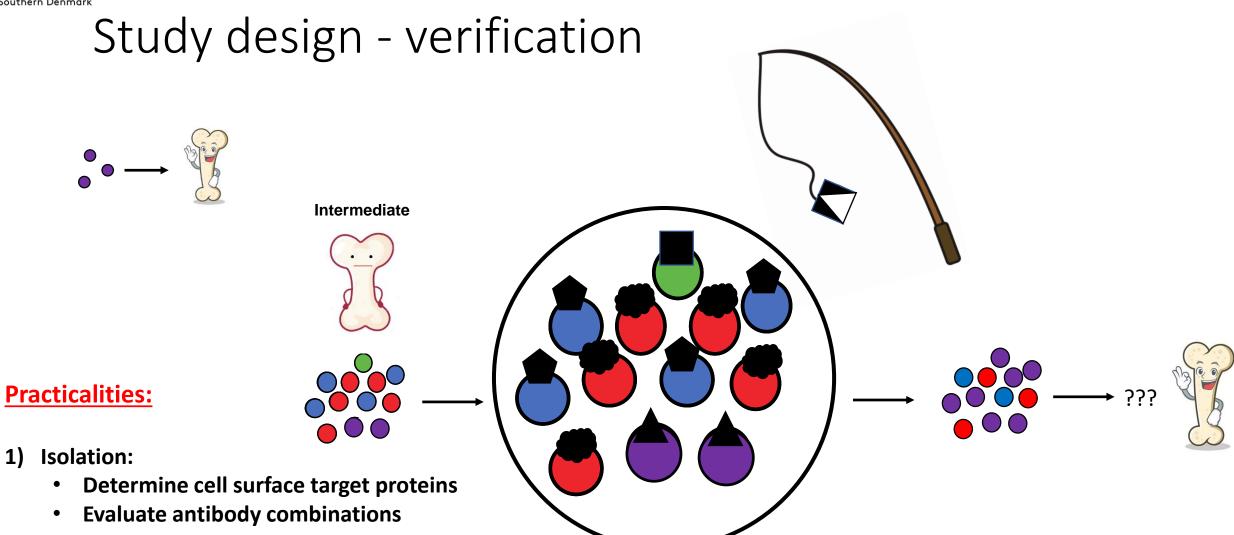
Study design - verification



Practicalities:

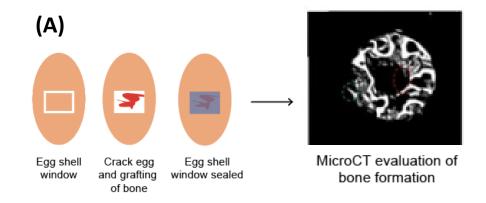
- 1) Isolation:
 - Determine cell surface target proteins
 - Evaluate antibody combinations (A)







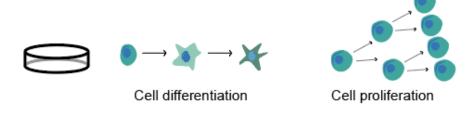
Study design - verification



Practicalities:

- 1) Isolation:
 - Determine cell surface target proteins
 - Evaluate antibody combinations
- 2) Intrinsic osteogenic potential determination:
 - In vivo differentiation (A)
 - In vitro differentiation (B)

(B)



Functional analysis in vivo and in vitro



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