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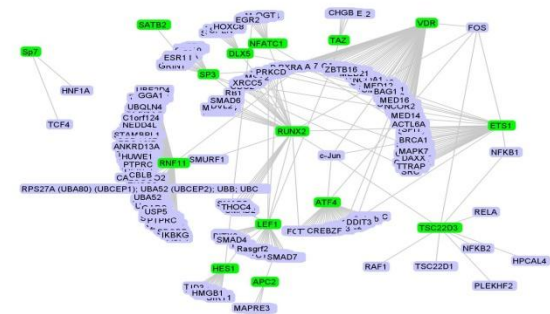
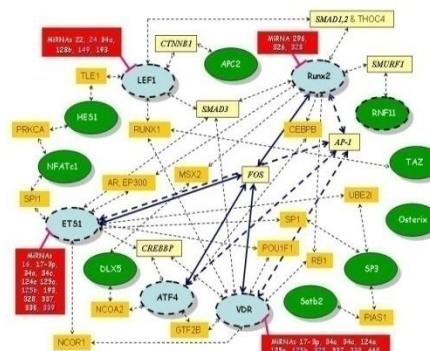
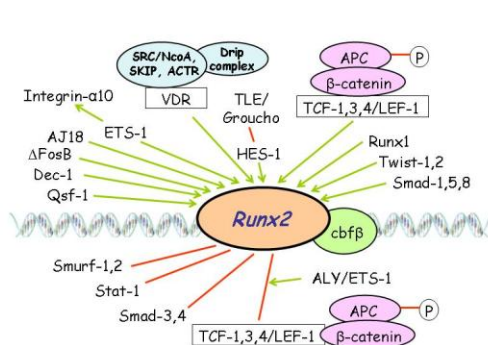


Key regulatory junctions stabilizing the osteoblast phenotype

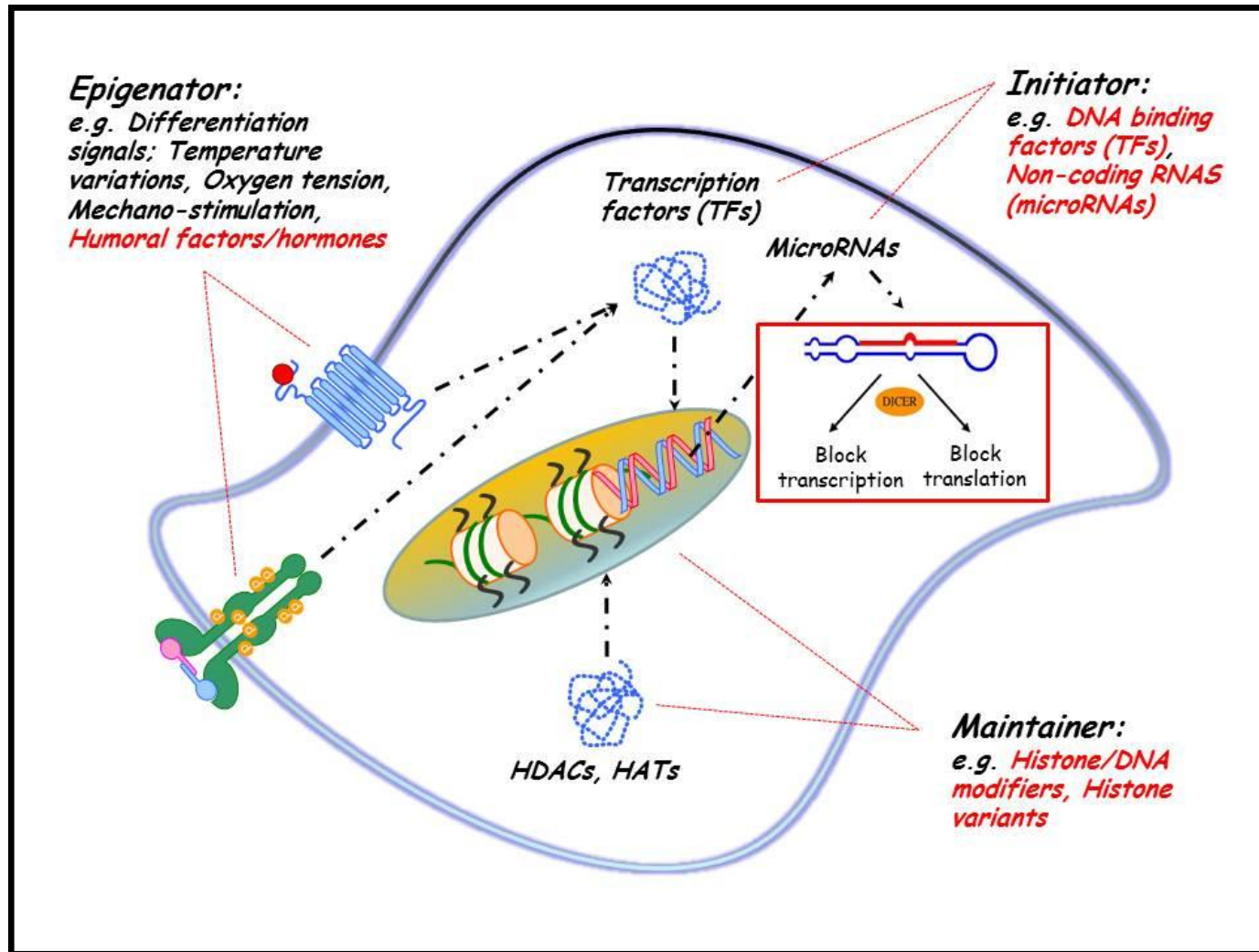
Implications for cell and tissue engineering

Jan O. Gordeladze^{1,2,3}, Janne E. Reseland⁴
Unni syversen⁵ and Mauro Valtieri⁶

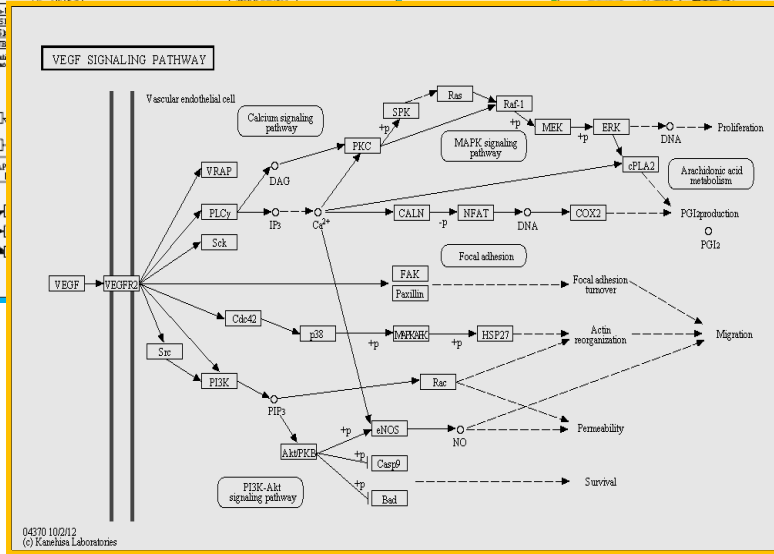
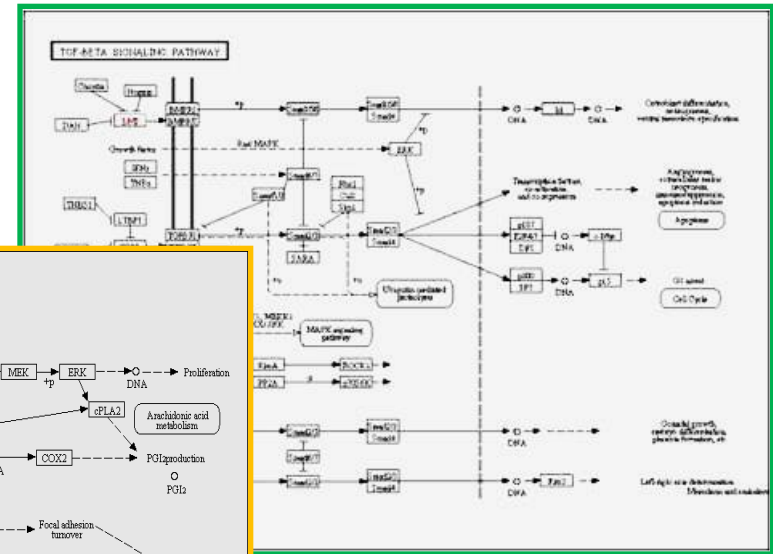
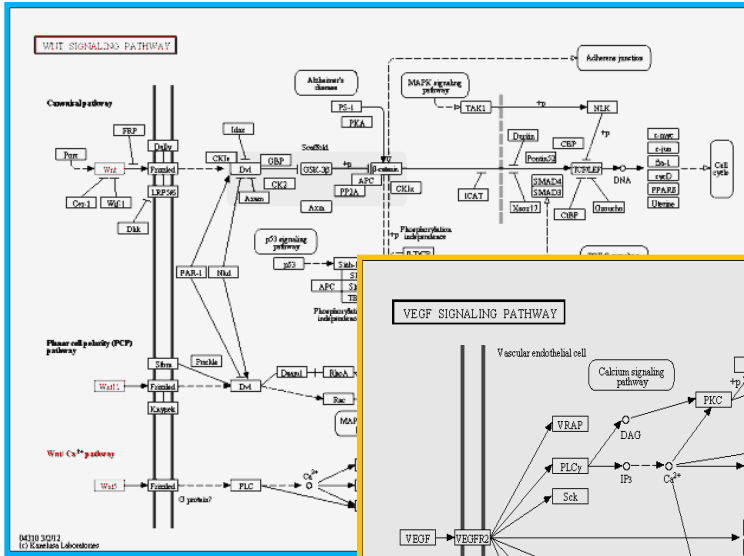
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The «epigenator, initiator, and maintainer» principle of epigenetics on phenotype acquisition and maintenance

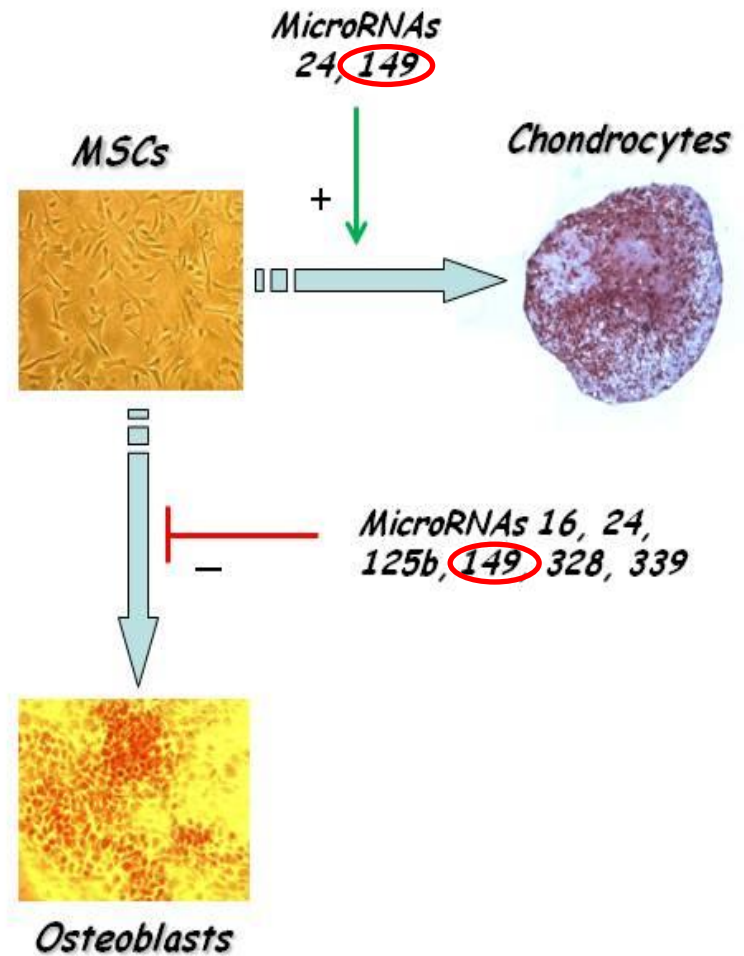
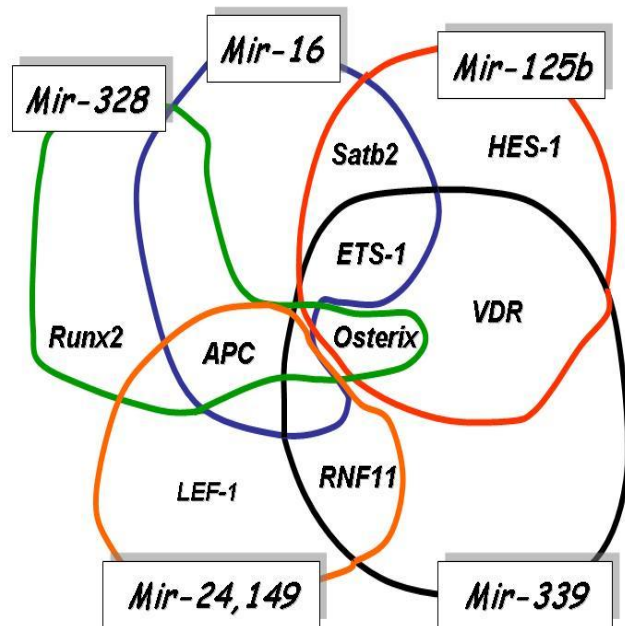


Factors to play with: *Signaling molecules*

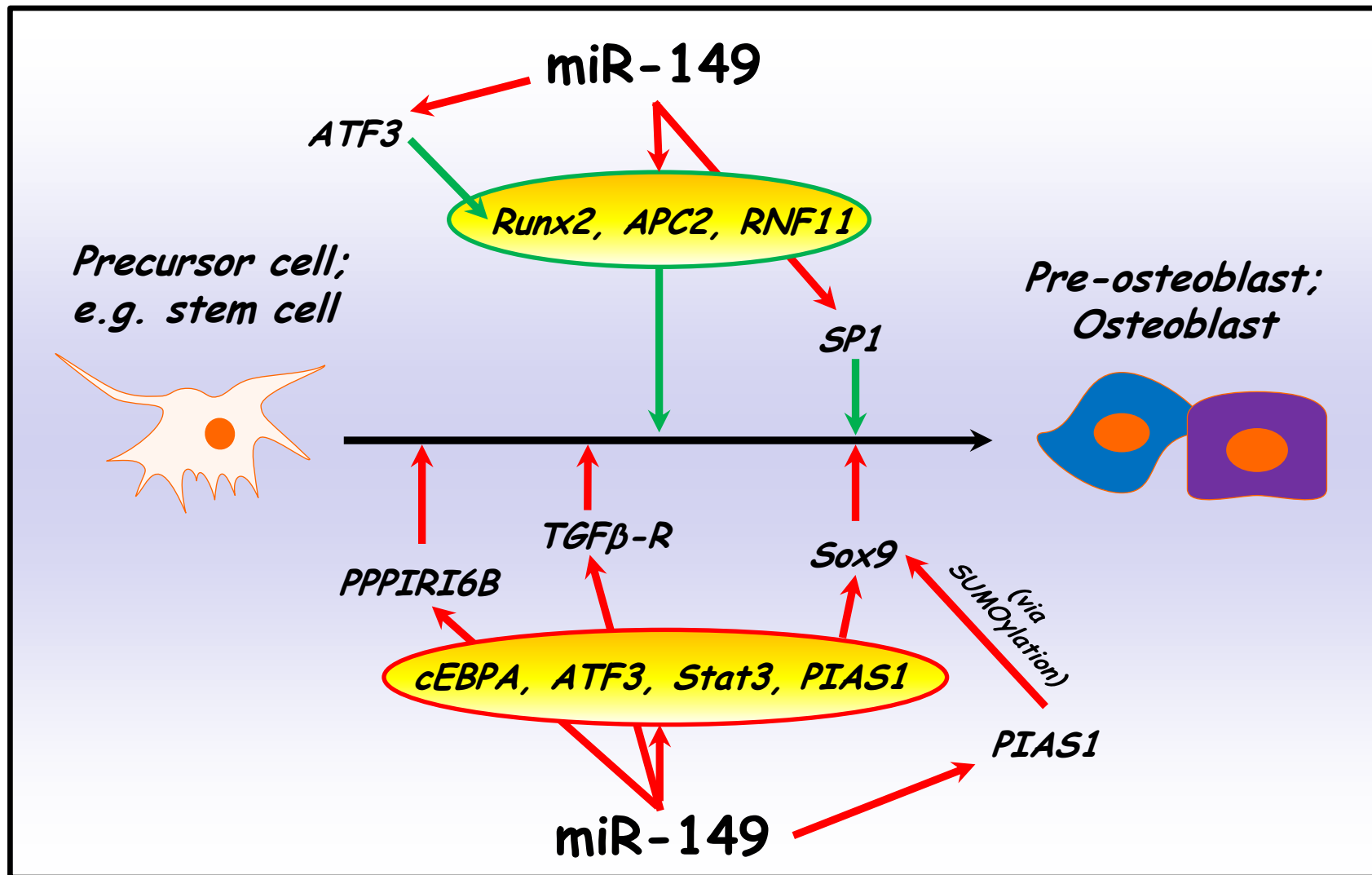


How the microRNA signature affects differentiation of osteoblasts and chondrocytes from hMSCs

6 microRNA species specifically block osteoblastogenesis, thereby promoting chondrogenesis, targeting at least 9 transcriptional modulators:



MiR-149 may serve as a switch-MiR yielding either osteoblasts or chondrocytes from stem cells





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Experimental approaches:

- 1) Identify and challenge **regulatory loops** including **microRNA species** of an osteoblast signature (**16, 24, 125b, 149, 328, and 339**), as well as **204 and 211**, and **transcription factors (TFs)** instrumental in «guiding» stem cells to differentiate into osteoblasts, and
- 2) Test «stabilized» osteoblasts for **resilience towards exposure to cytokines** produced by Th-cells.



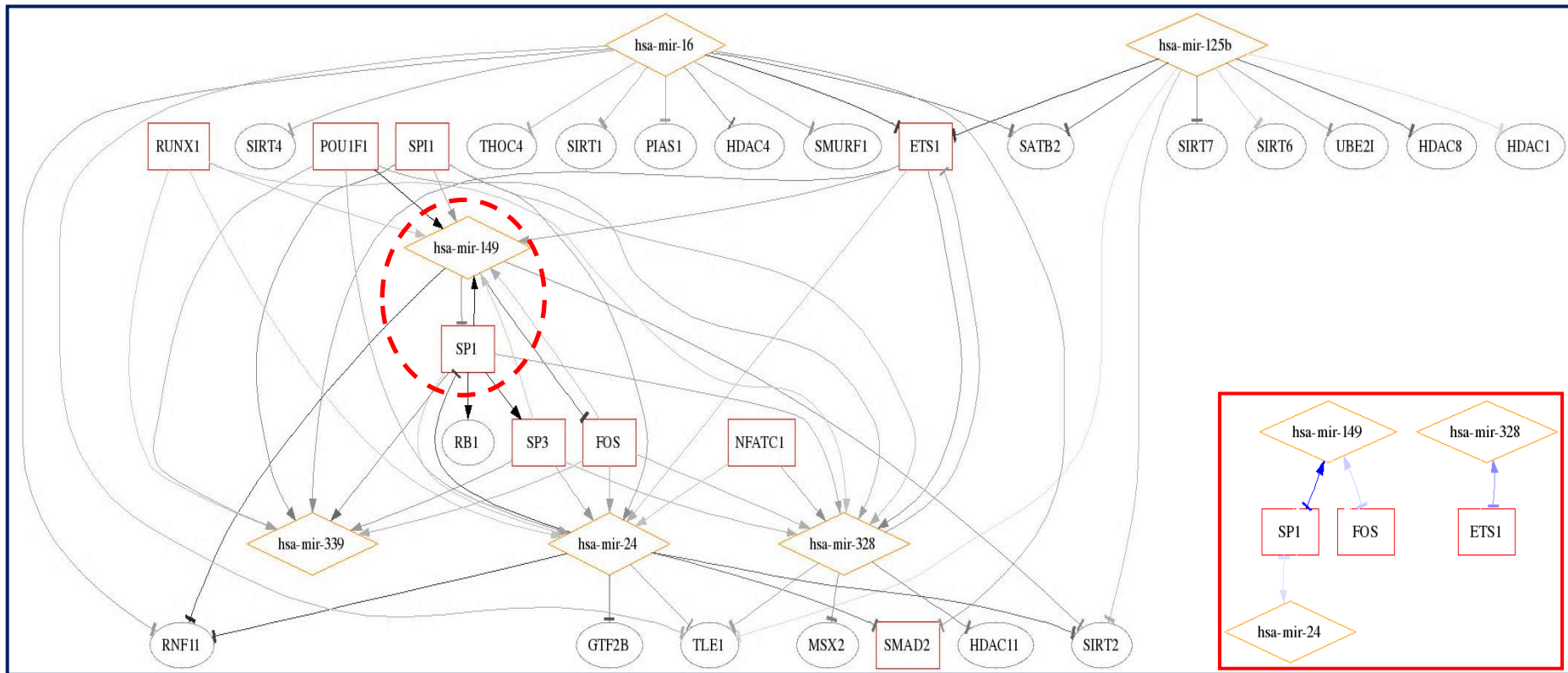
The use of the Mir@nt@n algorithm predicting interactions between transcription factors (TFs) and microRNAs

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Osteoblast





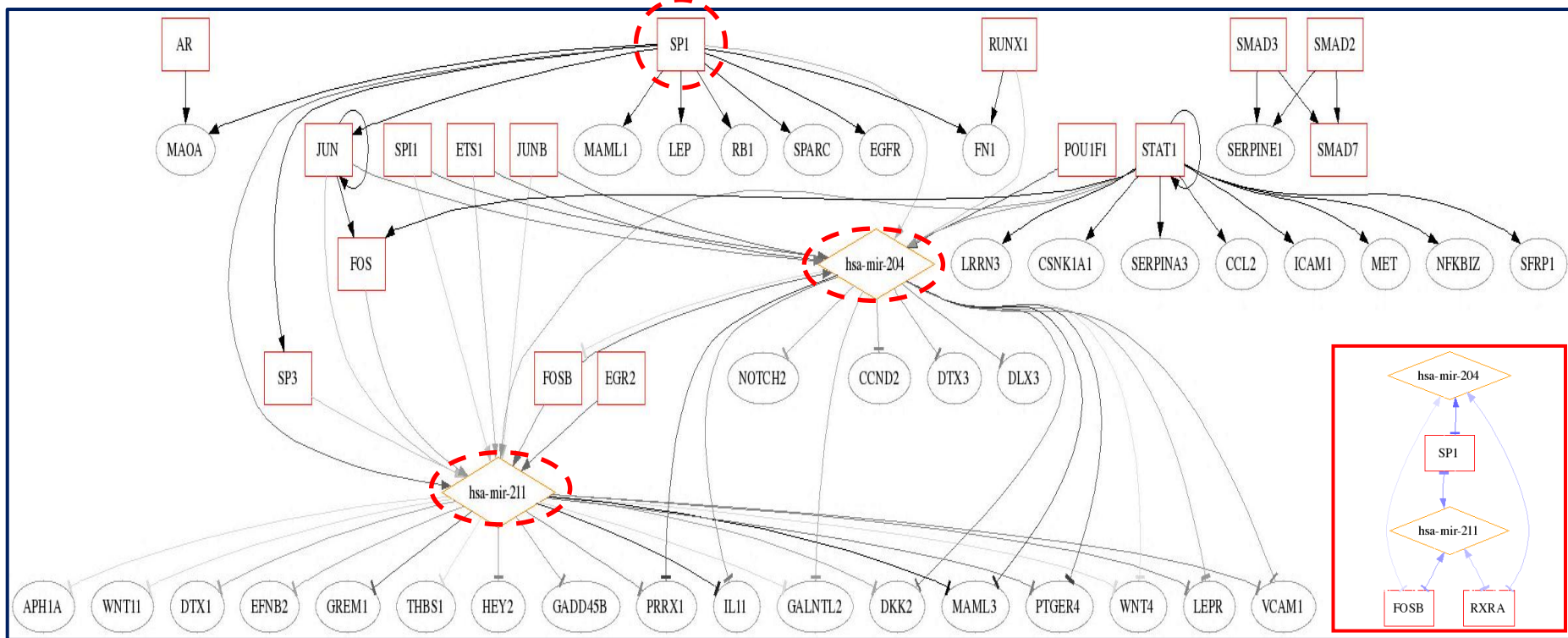
The use of the Mir@nt@n algorithm predicting interactions between transcription factors (TFs) and microRNAs

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Osteo-chondro-adipocyte





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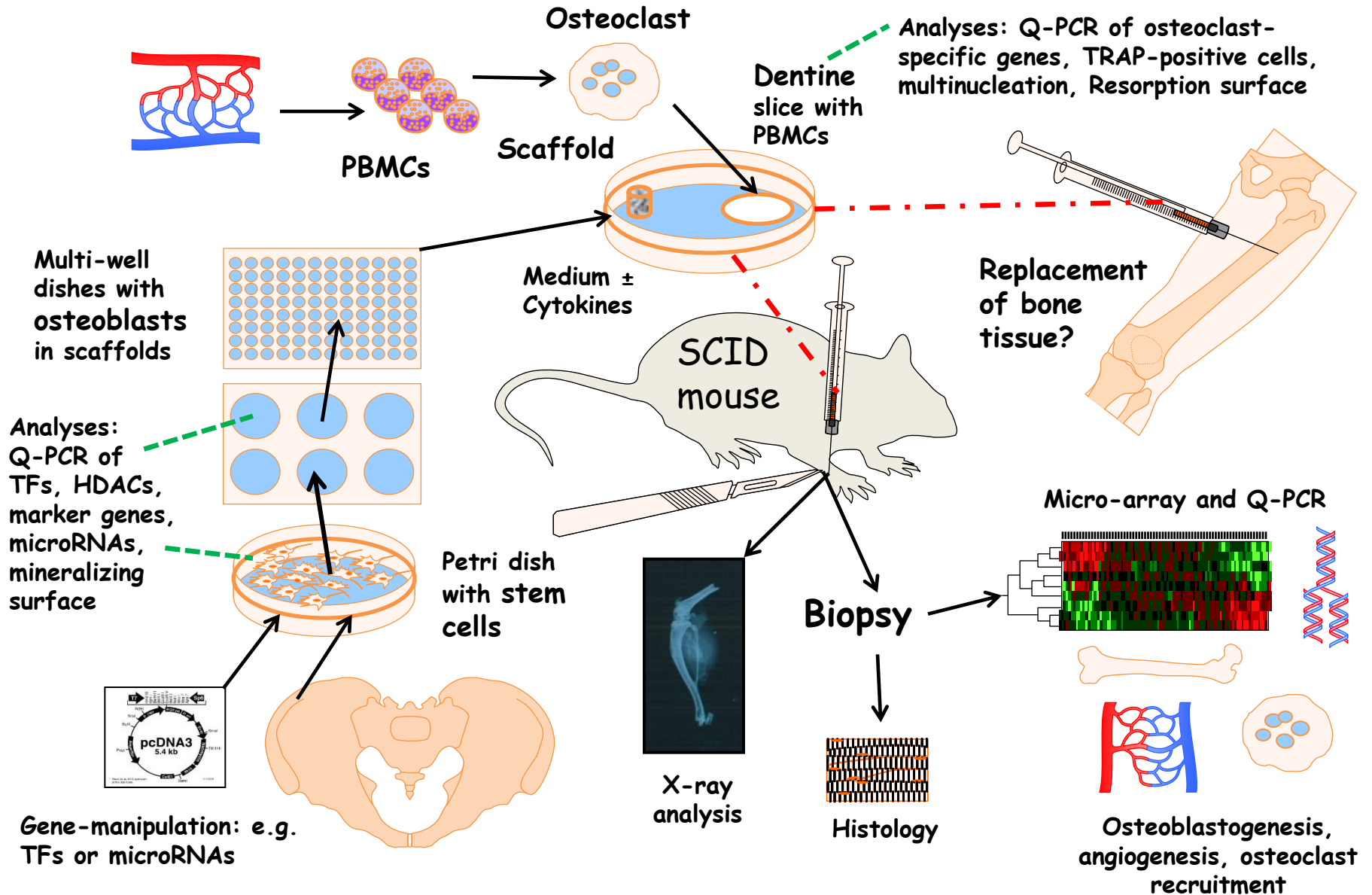
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*Is **SP1** important for Osteoblastogenesis?*

“From a literature search (PubMed) on “SP1 transcription factor and osteoblasts”, **SP1** is somehow interfering with the effect of **Runx2**, **SP7** (osterix), **FIAT** (inhibitor of **ATF4**), **ETS-like TFs**, **MZF1** (myeloid zinc finger), **JUNB**, and also directly affecting the transcription of marker genes like **Col1a1**, **Col5a1**, **Col5a3**, **Col11a2**, fibromodulin, osteocalcin, **MGP** (matrix-gla protein), **RANKL**, Pit phosphate transporter, **Integrin β 5**, and **TGF β -R1**.

Outline of interconnected experiments





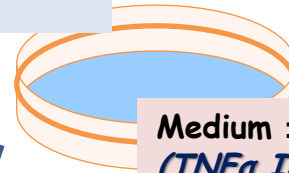
Outline of key experiments

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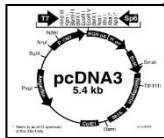
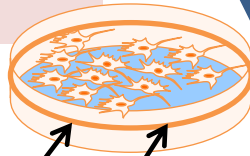


Analyses conducted:
Mineralized surface,
Immunohistochemistry,
Q-PCR of microRNAs and
Osteoblast marker mRNAs
and others

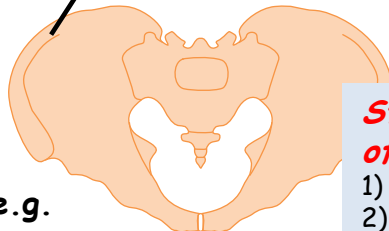


**Medium ± Cytokines
(TNF α , IL-1, IL-8, and
IL-17A)**

**Petri dish
with stem
cells**



**Gene-manipulation: e.g.
of SP1 or microRNAs**



Stem cells (MSCs or ASCs) are manipulated in terms of either:

- 1) SP1-expressing vector or Sh-RNA vs SP1
- 2) Polycistronic constructs with mir-204/211 or mir-149 and antago-mirs vs same microRNA species



The effect of SP1 overexpression in engineered osteoblasts

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Parameters	Control = 100%	+ Cytokines	+ SP1	+ SP1 + Cytokines
Runx2	100	27	432	324
Collagen1a1	100	47	145	133
Osteocalcin	100	23	345	288
Osterix	100	18	182	171
Mineralizing «surface»	100	22	234	198
OPG/Rank-L ratio	100	435	27	321
PPAR γ	100	534	28	43
HSL	100	385	35	58
Oil-Red-O «surface»	100	689	36	32
Mir-149	100	546	34	75
Mir-328	100	465	37	58



The effect of mir-204/211 suppression in engineered osteoblasts

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Parameters	Control = 100%	+ Cytokines	+ Antago-mirs	+ Antago-mirs + Cytokines
Runx2	100	25	389	319
Collagen1a1	100	36	319	272
Osteocalcin	100	27	321	251
Osterix	100	31	301	199
Mineralizing «surface»	100	47	247	167
OPG/Rank-L ratio	100	378	21	271
PPAR γ	100	444	19	39
HSL	100	417	27	47
Oil-Red-O «surface»	100	571	27	41
Mir-149	100	449	31	69
Mir-328	100	577	41	48



Summary and Conclusion

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- 1) *The use of bioinformatics (the Mir@nt@n algorithm) backed by PubMed searches yields interesting paradigm shifts as to which of many TFs are the better markers for cell phenotypes (SP1 instead of Runx2; Osterix = SP7) characterizing osteoblasts?*
- 2) *Manipulating members of regulatory loops encompassing TFs and microRNAs makes it easier to either disrupt or reinforce the stability of a certain phenotype (e.g. enhance osteoblast resilience against exposure to inflammation).*
- 3) *Identifying regulatory loops encompassing microRNAs and TFs may thus be important/mandatory for the success of cell engineering/replacement cell therapy.*



*To be commended for their scientific and technical support and never-ceasing enthusiasm:
INSERM U844, Montpellier, France
&
Université Montpellier 1, enabling me to work within the U844 as a guest professor for 3 years*

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Thank you for your attention!