



Fat and HIV persistence, role of fat metabolism and inflammation

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GENE EXPRESSION ANALYSIS

- **Healthy controls**
- **Naïve:** HIV-1-infected, untreated patients
- **Non-LD:** HIV-1-infected patients, on cART without clinical signs of lipoatrophy
- **LD:** HIV-1-infected patients, on cART with clinical lipoatrophy

Applied Biosystems Human Genome Survey Arrays

Gene markers of
mitochondrial biogenesis
adipogenesis
adipokines
inflammation



Biopsies of
subcutaneous
adipose tissue



RNA
DNA
protein
extraction

Transcriptomic analysis of subcutaneous adipose tissue from HIV-patients relative to healthy controls

	Naïve	Non-LD	LD
<u>IMMUNOLOGY-RELATED</u>	—	—	—
Complement-mediated immunity	*	*	*
T-cell activation	*	*	—
Interferon-mediated immunity	*	—	—
B-cell antibody-mediated immunity	*	*	—
Inflammation mediated by chemokines and cytokine signaling pathways	*	*	*
MHC-II mediated immunity	—	—	*
Macrophage-mediated immunity	—	*	** *

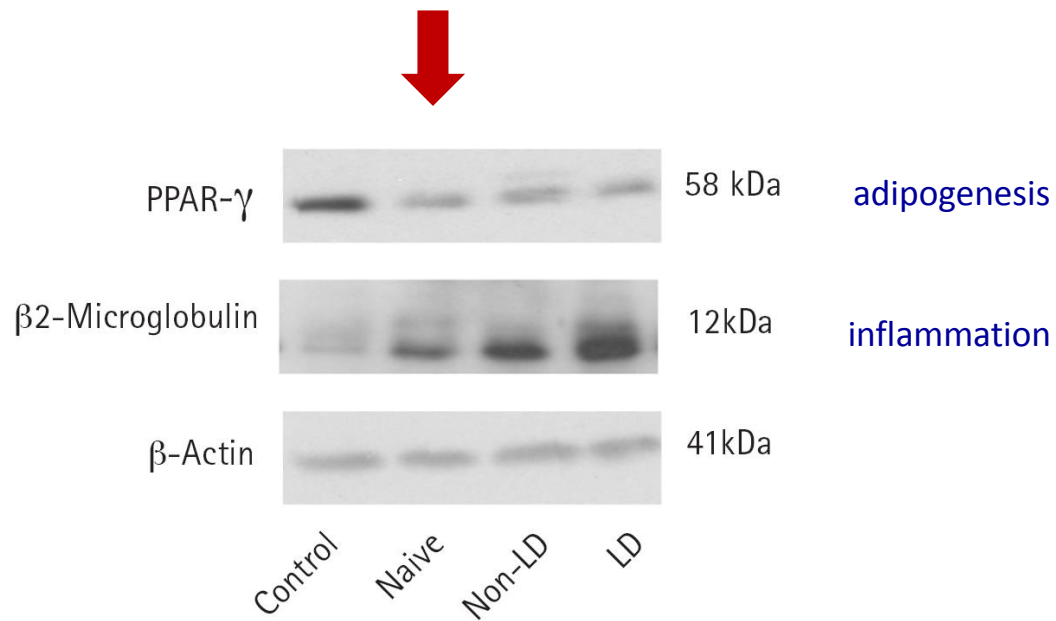
* significant change (either up- or down-regulation)

Transcriptomic analysis of subcutaneous adipose tissue from HIV-patients relative to healthy controls

	Naïve	Non-LD	LD
<u>METABOLISM</u>			
Lipid, fatty acid & steroid metabolism	*	* *	* *
Fatty acid metabolism	*	* *	* *
Fatty acid beta-oxidation	*	*	*
Lipid and fatty acid transport	-	-	*
Lipid and fatty acid binding	-	-	*
TCA Cycle	*	*	*
Oxidative phosphorylation	-	*	* *
Electron transport	-	*	* *
Detoxification	-	*	-
<u>OTHER</u>			
Apoptosis	-	-	*
Angiogenesis	*	*	*

* significant change (either up- or down-regulation)

Reciprocal repression of marker proteins of adipogenesis and induction of marker proteins of inflammation in subcutaneous adipose tissue

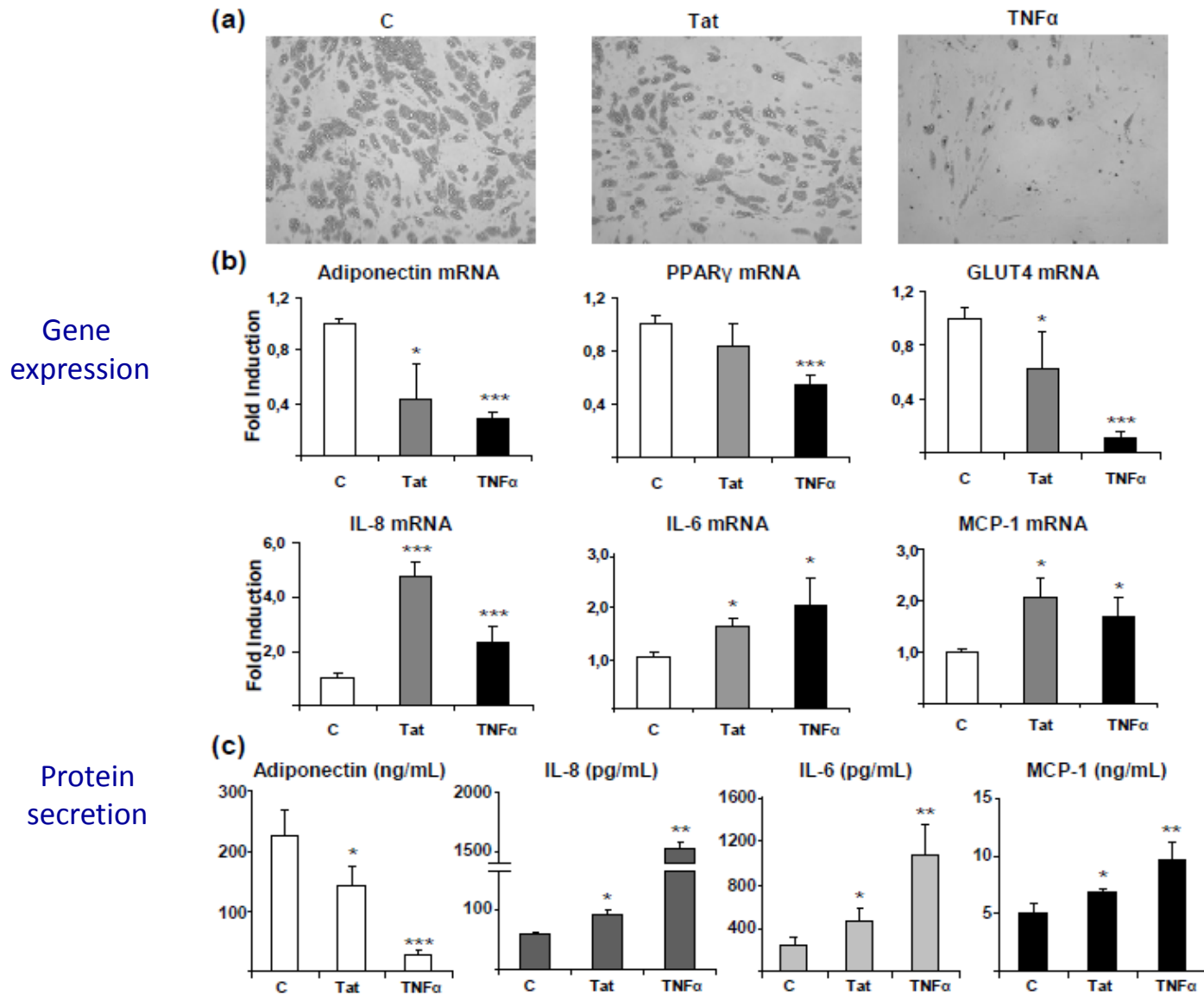


**Viral infection, in the absence of cART, impacts on
adipose tissue biology:**

associated pro-inflammatory environment?

direct action of viral proteins?

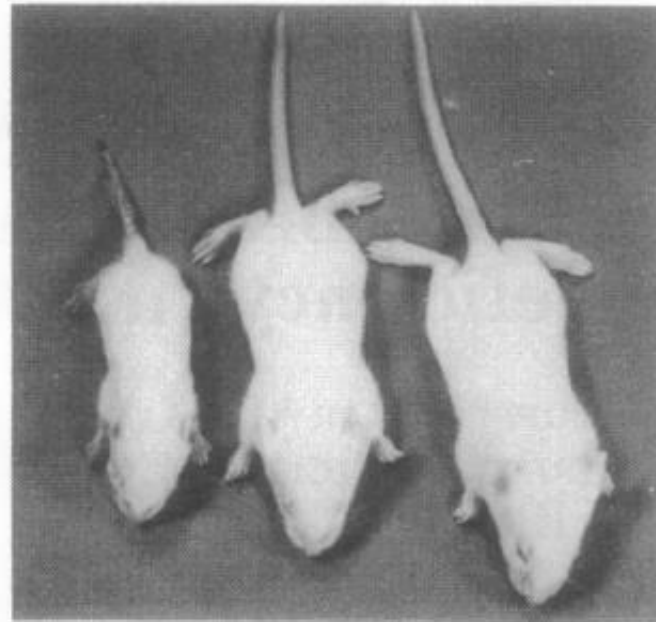
HIV-Tat inhibits human adipocyte differentiation and induces expression and release of pro-inflammatory cytokines



HIV-1 secreted proteins (e.g., Tat) may play a role in the adipose tissue alterations that ultimately lead to adipose tissue and systemic metabolic disturbances observed in HIV-1-infected patients



Transgenic mice expressing the HIV-1 genome devoid of the *gag-pol* genes (Tg26 mice)



Homozygous mice
seldom survive to
weaning

Heterozygous mice exhibit
premature death (around
6 month)

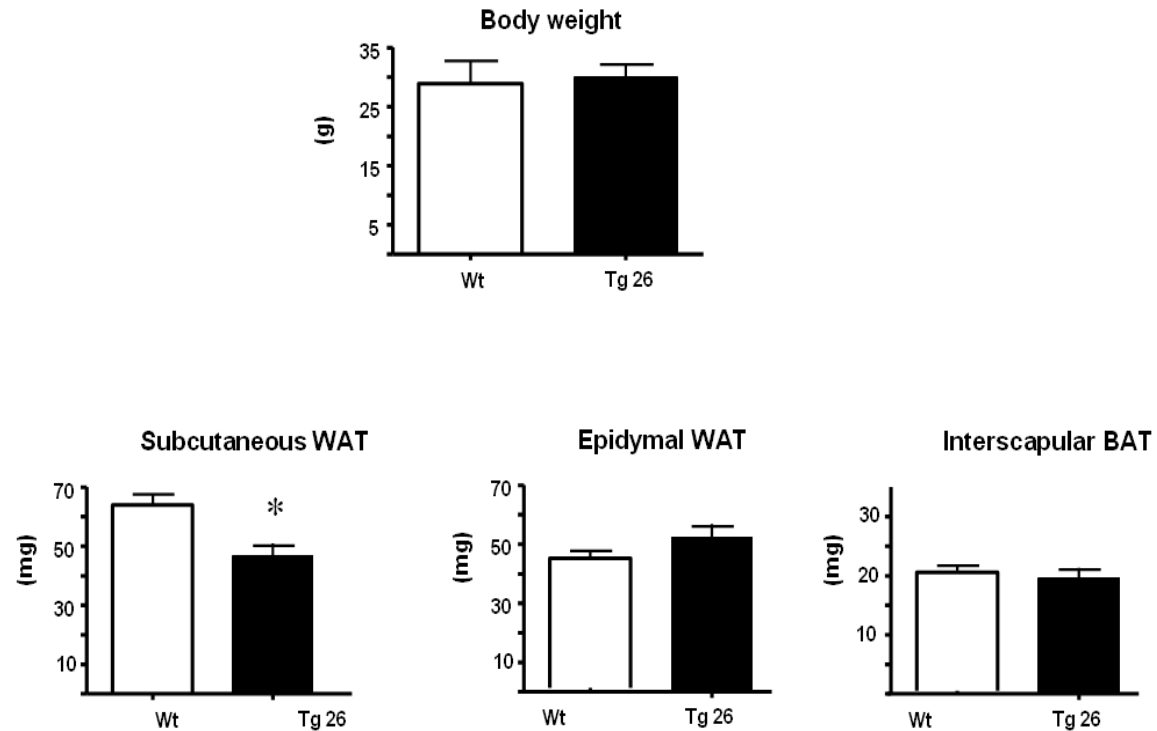
Tg26 +/+ **Tg26 +/-** WT

Transgenic mice containing 10-20 of a 7.4-kb proviral HIV-1 DNA lacking pol and gag genes to avoid spontaneous infection

Collaboration with PE Klotman, Mount Sinai Hospital, New York (now at Baylor College of Medicine, Houston) USA



HIV-1 transgene expression in mice alters the amount of scWAT

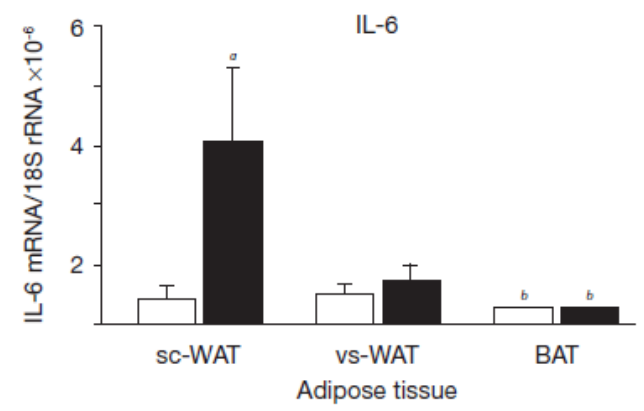
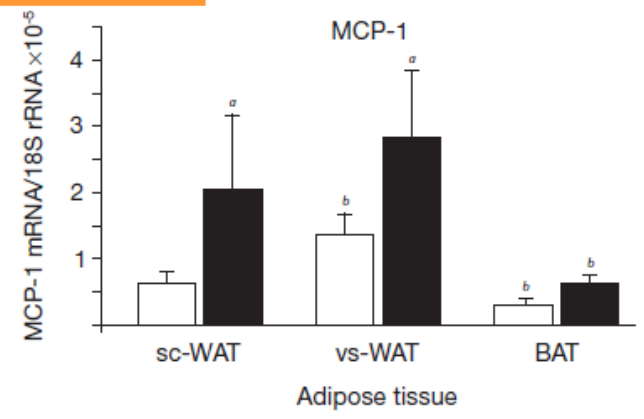
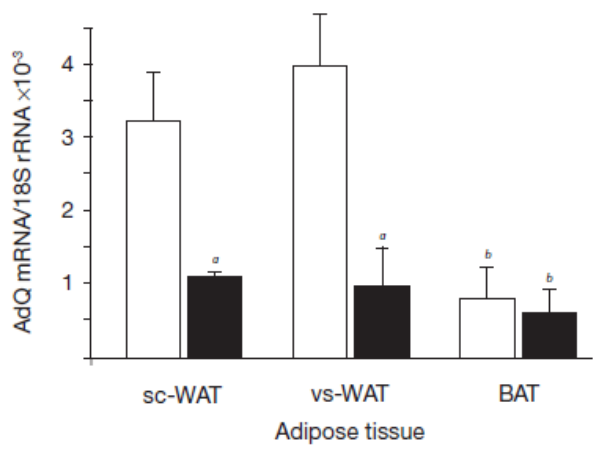
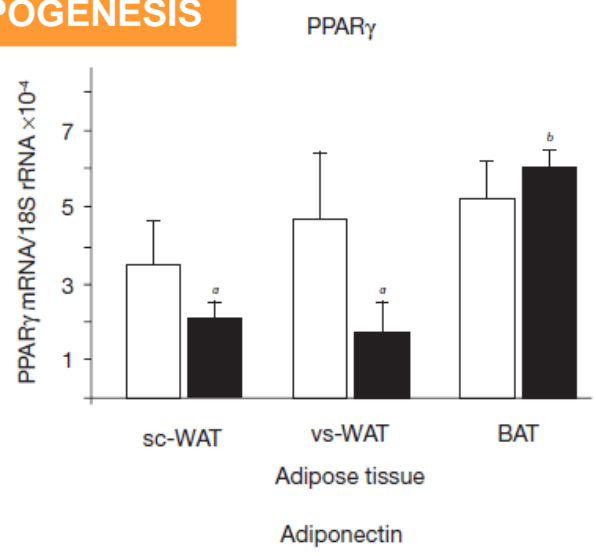


□ Wild-type ■ Tg26+/-

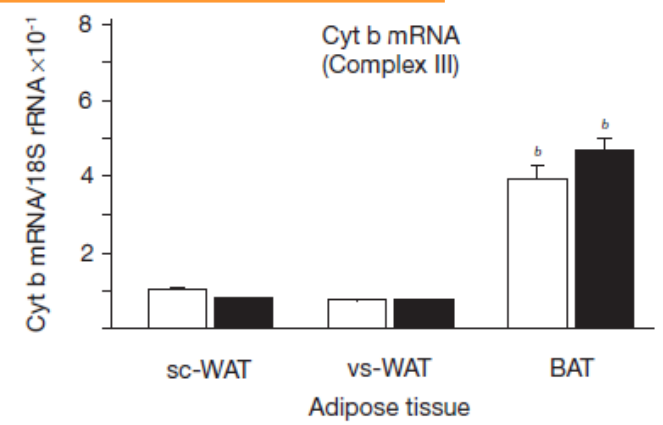


HIV-1 transgene expression in mice alters gene expression in WATs

ADIPOGENESIS



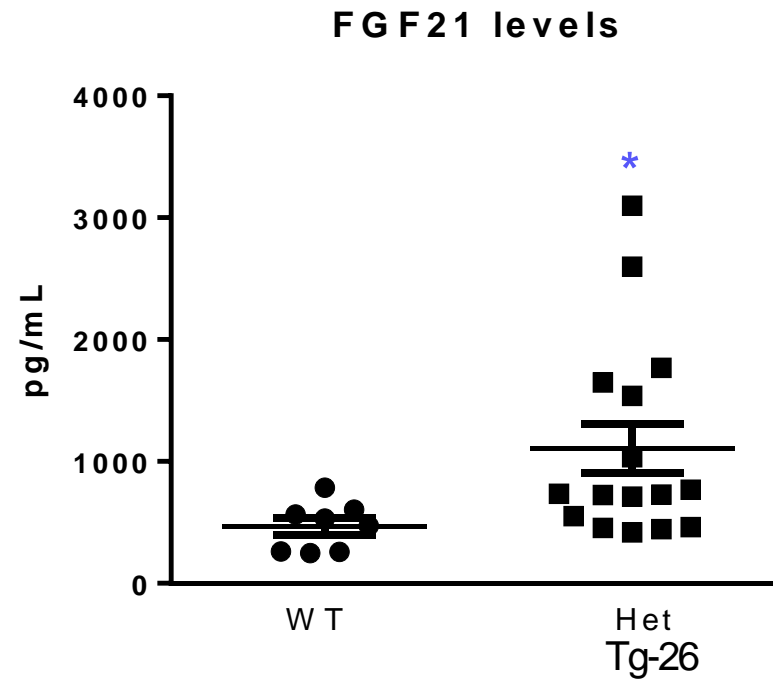
MITOCHONDRIAL FUNCTION



Transgenic HIV expression in mice recapitulates partially several features of the adipose tissue and adipokine alterations in HIV-1-infected patients associated with lipodystrophy, specially mild alterations not attributable to antiretroviral treatment



HIV-1 transgene expression in mice increases serum FGF21 levels





FGF-21 as a novel metabolic regulator

Alexei Kharitononkov,¹ Tatiyana L. Shiyanova,¹ Anja Koester,¹
Amy M. Ford,¹ Radmila Micanovic,¹ Elizabeth J. Galbreath,¹ George E. Sandusky,¹
Lisa J. Hammond,¹ Julie S. Moyers,¹ Rebecca A. Owens,¹ Jesper Gromada,²
Joseph T. Brozinick,¹ Eric D. Hawkins,¹ Victor J. Wroblewski,¹ De-Shan Li,¹
Farrokh Mehrbod,¹ S. Richard Jaskunas,¹ and Armen B. Shanafelt¹

¹Lilly Research Laboratories, Division of Eli Lilly and Co., Indianapolis, Indiana, USA. ²Lilly Research Laboratories, Hamburg, Germany.

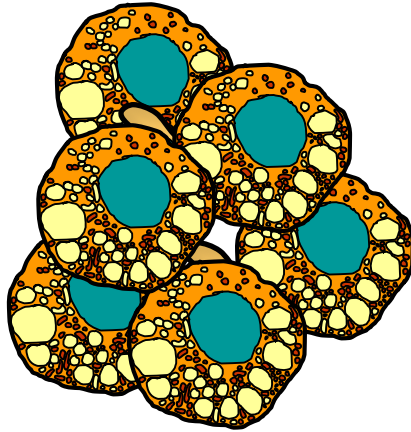
Diabetes mellitus is a major health concern, affecting more than 5% of the population. Here we describe a potential novel therapeutic agent for this disease, FGF-21, which was discovered to be a potent regulator of glucose uptake in mouse 3T3-L1 and primary human adipocytes. FGF-21-transgenic mice were viable and resistant to diet-induced obesity. Therapeutic administration of FGF-21 reduced plasma glucose and triglycerides to near normal levels in both *ob/ob* and *db/db* mice. These effects persisted for at least 24 hours following the cessation of FGF-21 administration. Importantly, FGF-21 did not induce mitogenicity, hypoglycemia, or weight gain at any dose tested in diabetic or healthy animals or when overexpressed in transgenic mice. Thus, we conclude that FGF-21, which we have identified as a novel metabolic factor, exhibits the therapeutic characteristics necessary for an effective treatment of diabetes.

In mouse models of diabetes and obesity, FGF21 administration lowers glucose, improves insulin sensitivity and decreases body weight

JCI, 2005

CONTROL OF METABOLIC CIRCADIAN BEHAVIOR AND FEMALE REPRODUCTIVE HORMONES

Boockout et al., & Owen et al., Nature Med. 2013

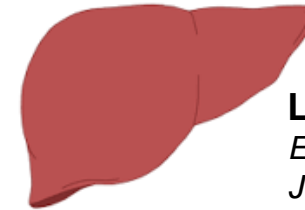
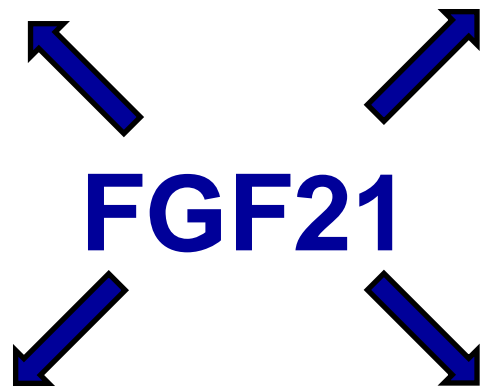


SWEET AND ALCOHOL PREFERENCE

*Talukdar et al., von Holstein
Rathlou et al., and Soberg
et al. Cell Metab 2016, 2017*

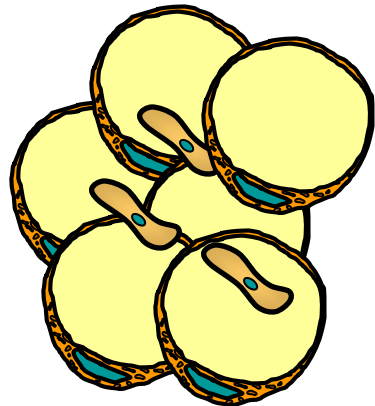
↑ THERMOGENIC ACTIVATION

*Hondares et al,
Cell Metab 2010*



LIPID METABOLISM

*Emanuelli et al.,
J Clin Invest 2014*



↑ GLUCOSE UPTAKE AND OXIDATION

Kharitononkov et al. JCI 2008

↑ «BROWNING»

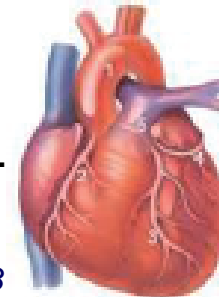
Fisher et al, Genes Dev. 2012

↑ ADIPONECTIN

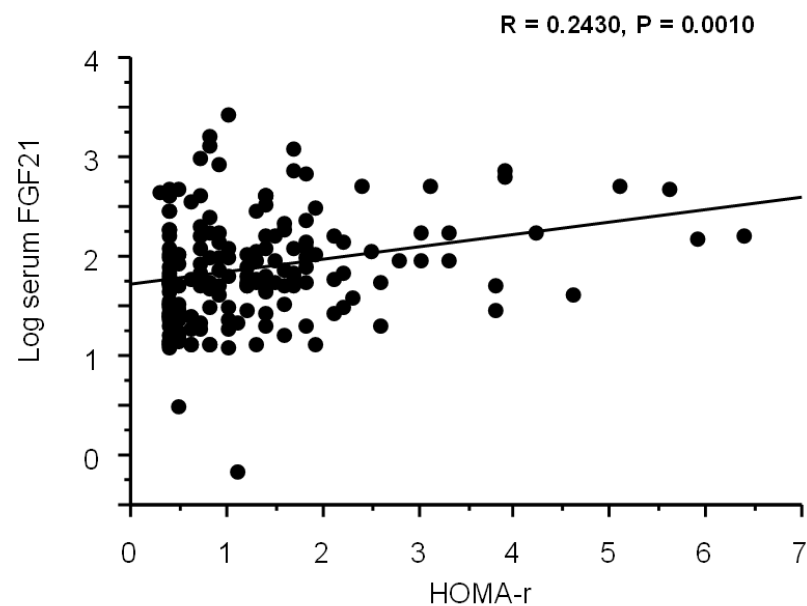
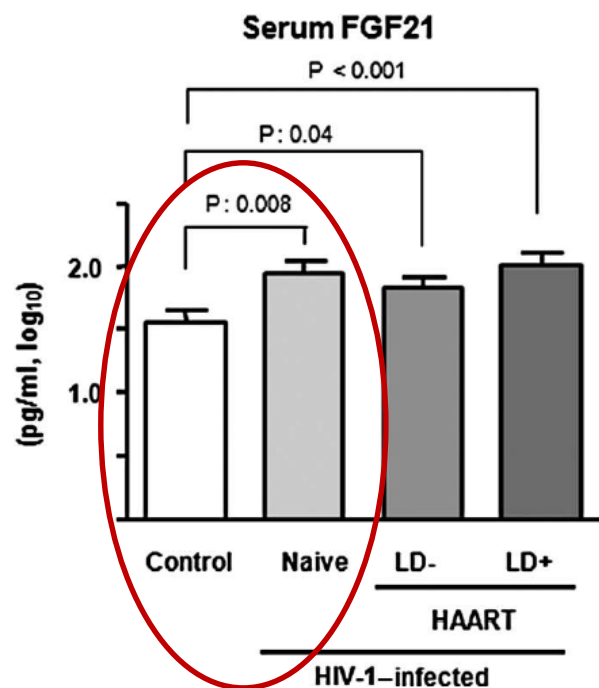
Holland et al.; Li et al. Cell Metab, 2013

↑ PROTECTION AGAINST PATHOLOGICAL HYPERTROPHY

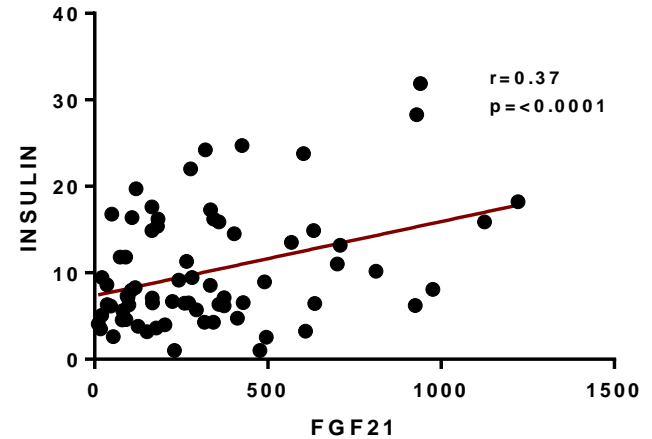
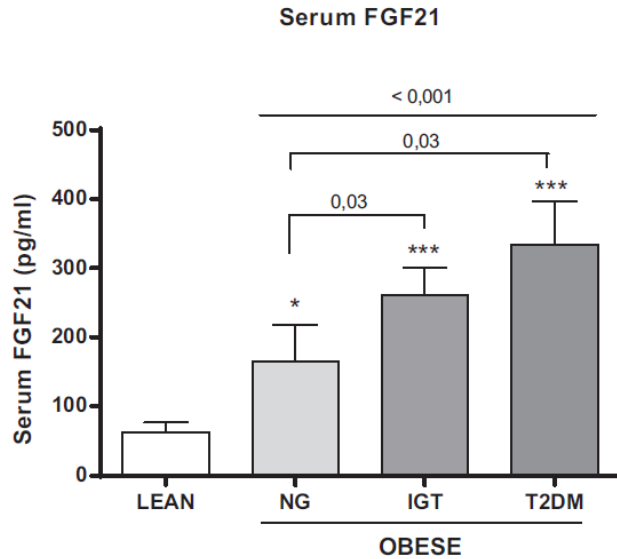
*Planavila et al.
Nature Comm. 2013*



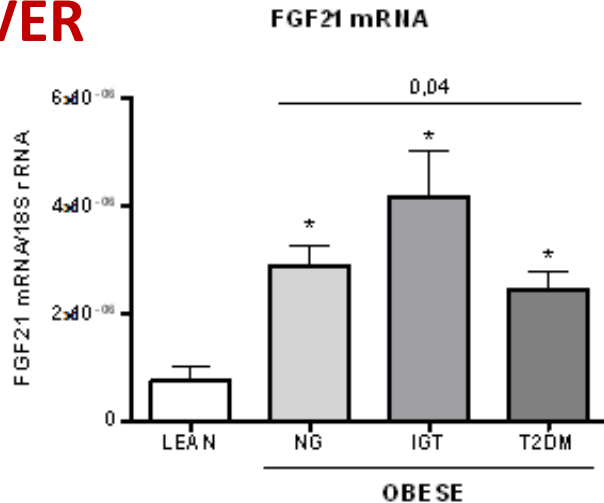
HIV-1-infected patients show increased levels of serum FGF21 associated with insulin resistance, even in the absence of cART



Serum FGF21 levels and FGF21 expression in liver are increased in obese and T2DM patients



LIVER



WAT

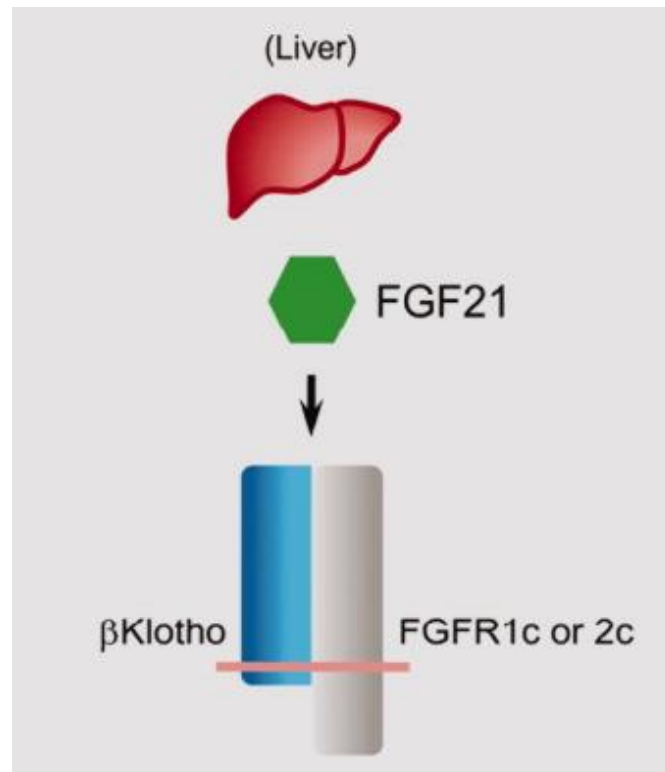
FGF21 mRNA was undetectable in scAT and vAT from lean and obese individuals

In collaboration with Gema Frühbeck, Clínica Universidad de Navarra (Pamplona, Spain)

In a Mexican population with a high prevalence of obese HIV-infected patients, no additive effects on abnormally high FGF21 levels are observed

**Increased serum FGF21 levels in obese
and HIV-patients despite altered
metabolism, ... a paradox?**

FGF21-resistance in these patients?

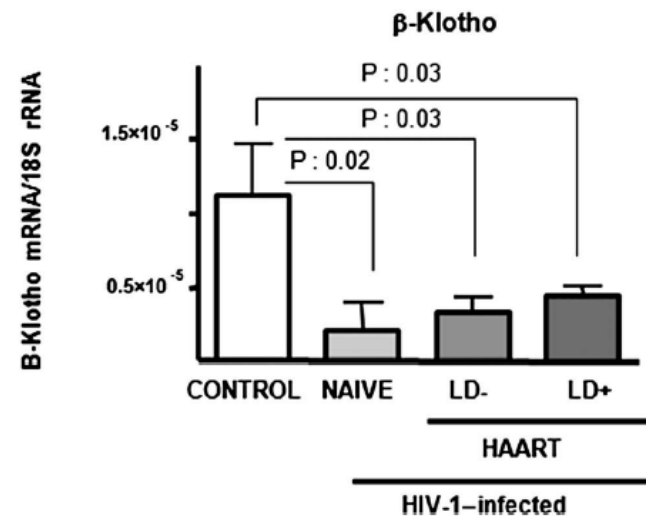
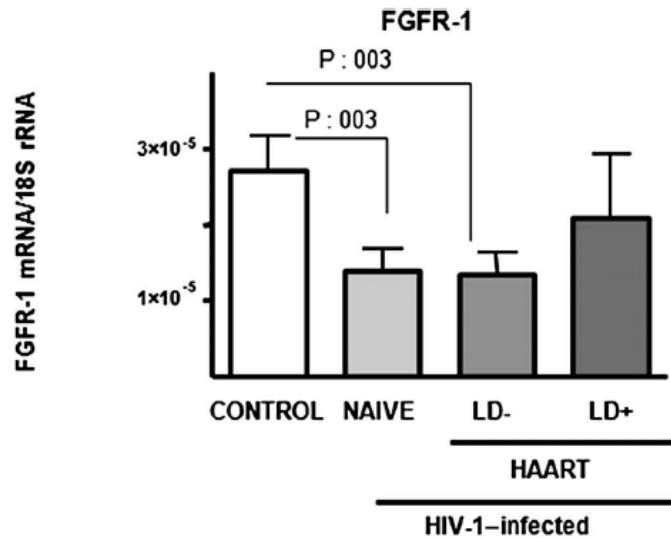


FGF21

Requires the presence of a specific transmembrane protein of the klotho family, **β -klotho (KLB)** in order to bind the receptor effectively and to activate intracellular pathways

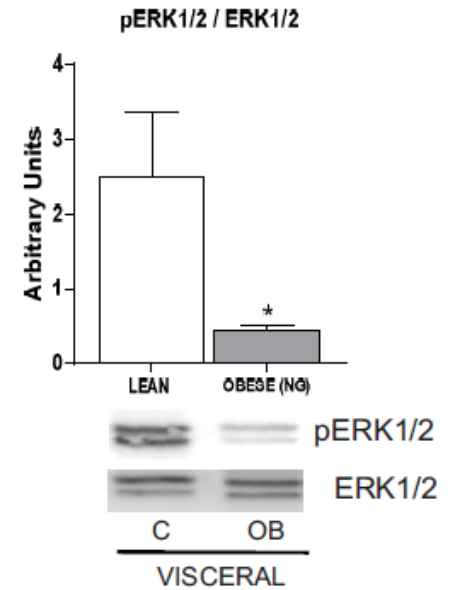
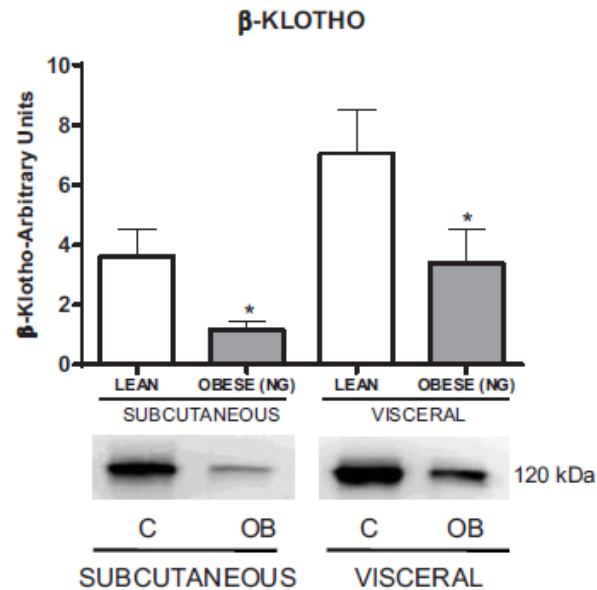
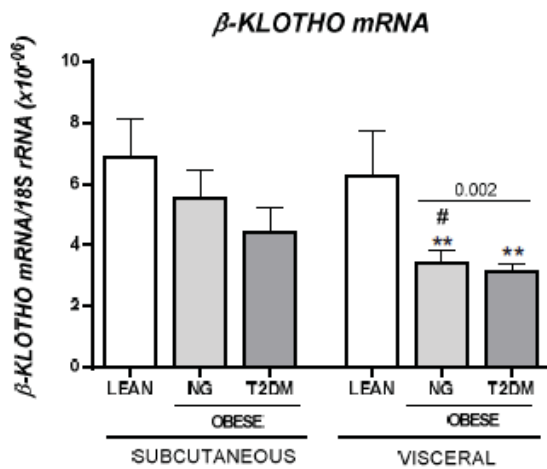
HIV-1-infected patients show impaired expression of β -Klotho and FGFR1 in adipose tissue, consistent with impaired FGF21 action, even in the absence of cART

SC ADIPOSE TISSUE



Obese individuals show impaired expression of β -Klotho in adipose tissue and impaired FGF21 action

ADIPOSE TISSUES



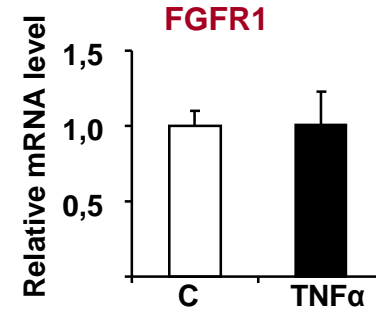
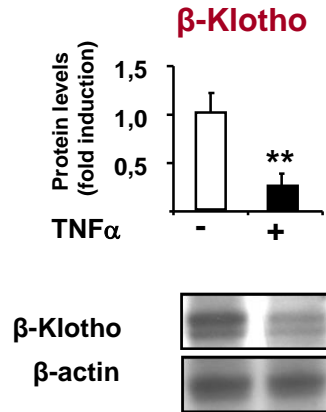
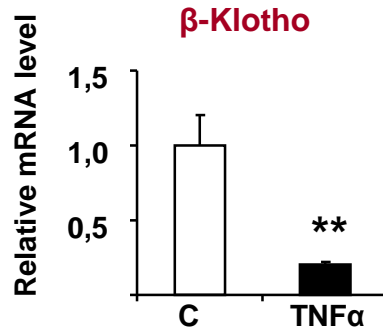
HIV-1-infected and obese patients show similar alterations in the endocrine FGF21 pathway

Abnormalities in the FGF21 pathway appear in HIV-1-patients, even without cART

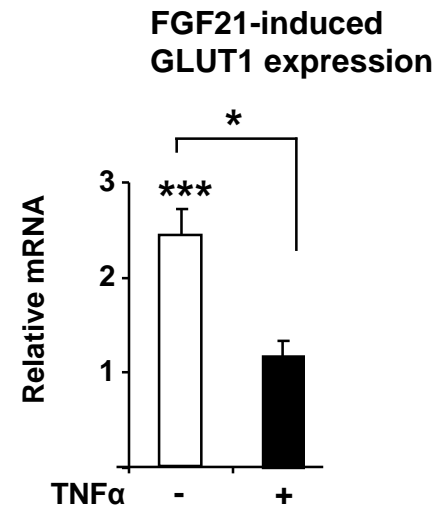
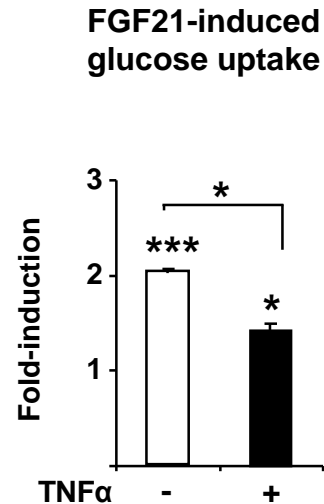
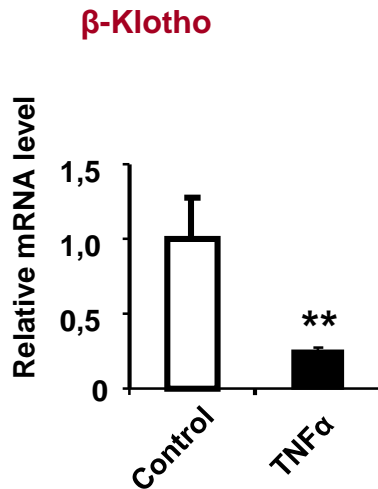


TNF α -induced repression of β -Klotho expression is associated with impaired responsiveness to FGF21 action

Murine white adipocytes



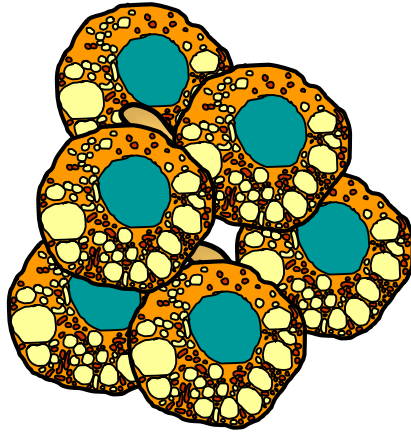
Human adipocytes



**Inflammatory signaling contributes to altered
FGF21 action in adipose tissue**

CONTROL OF METABOLIC CIRCADIAN BEHAVIOR AND FEMALE REPRODUCTIVE HORMONES

Boockout et al., & Owen et al., Nature Med. 2013



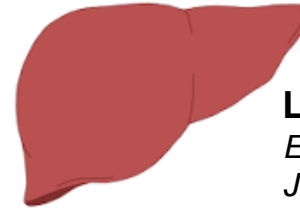
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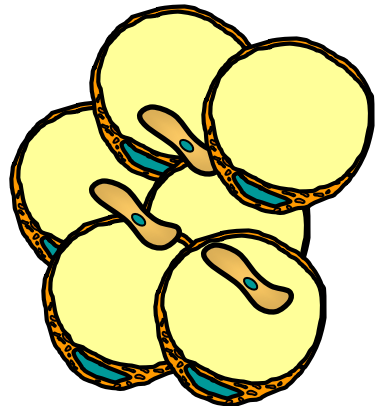
*Emanuelli et al.,
J Clin Invest 2014*

FGF21



BONE LOSS

*Wei et al. PNAS, 2012; Wang et al,
Cell Metab. 2015*



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Kharitononkov et al. JCI 2008

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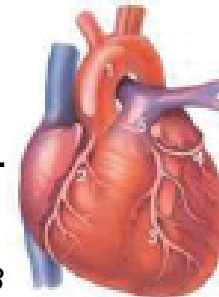
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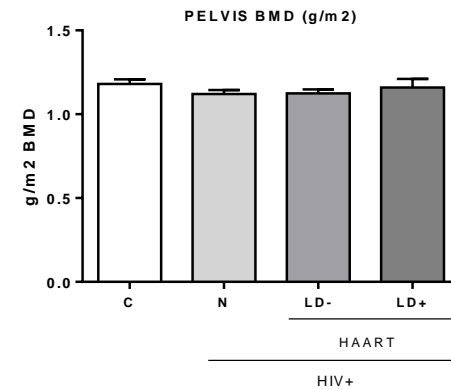
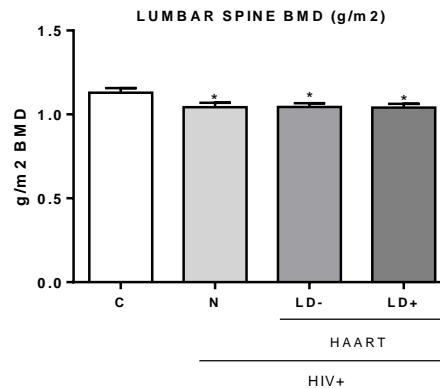
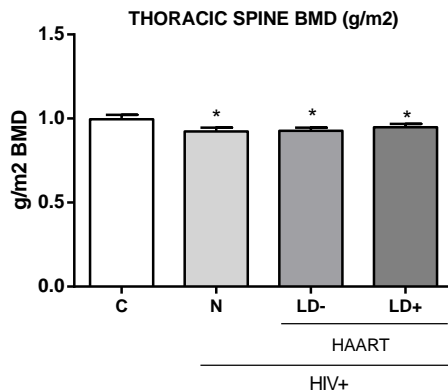
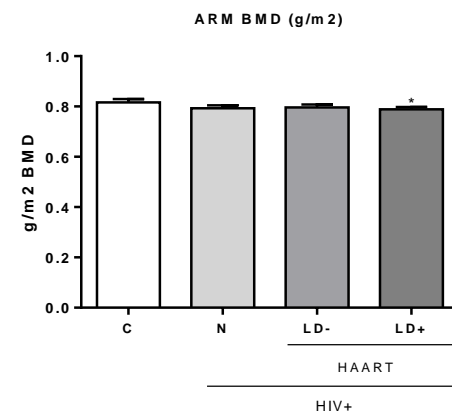
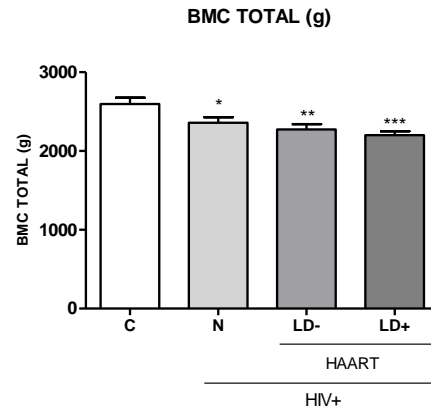
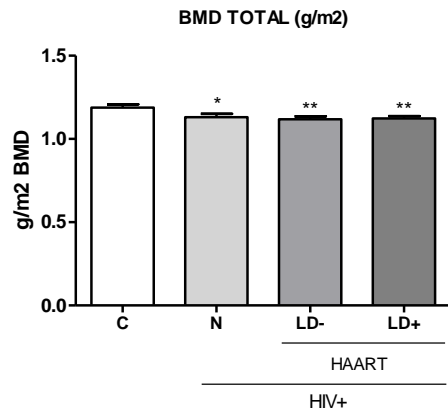
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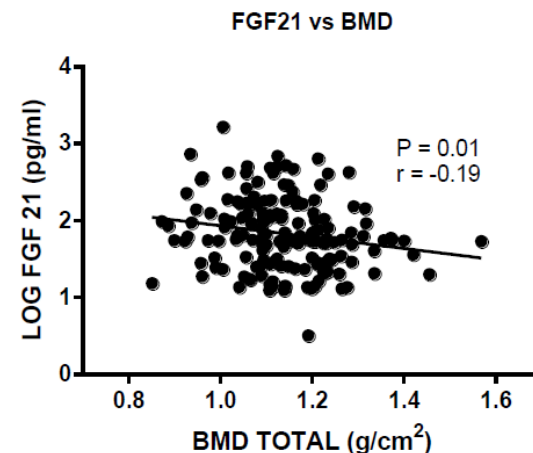
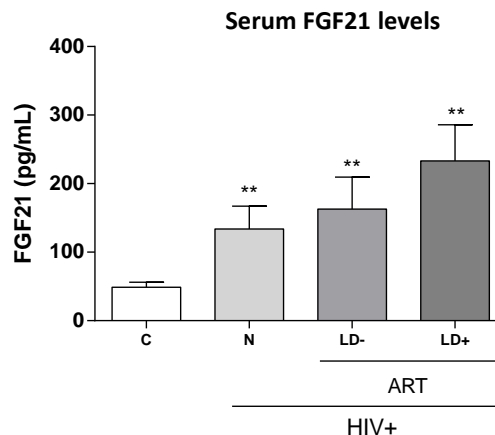
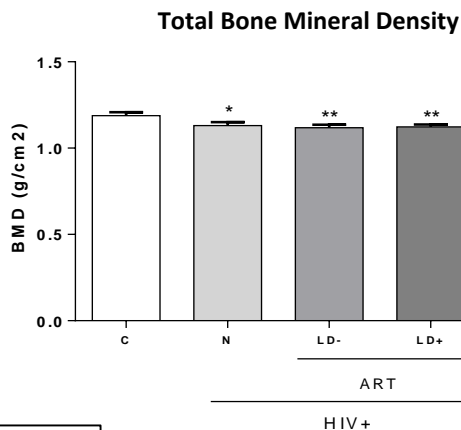
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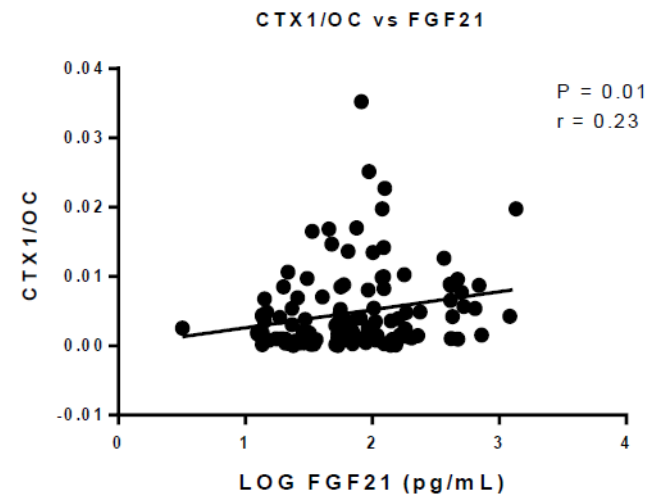
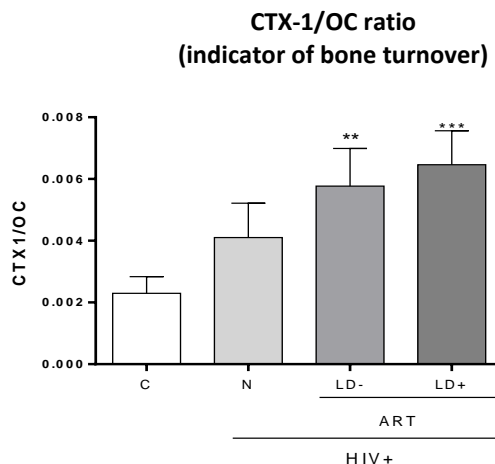
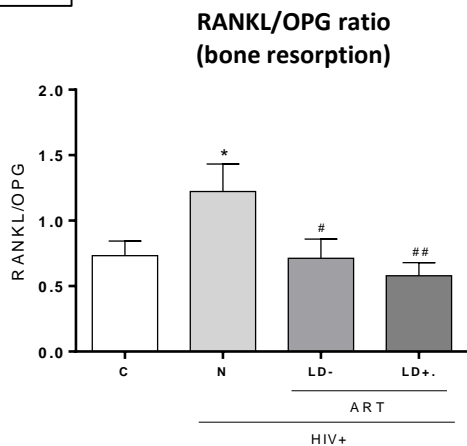
HIV-1-infected patients show altered bone parameters, even in the absence of cART



High FGF21 levels are associated with poor bone homeostasis in HIV-1-infected patients



*P≤0.05 vs C
#P≤0.05 vs N

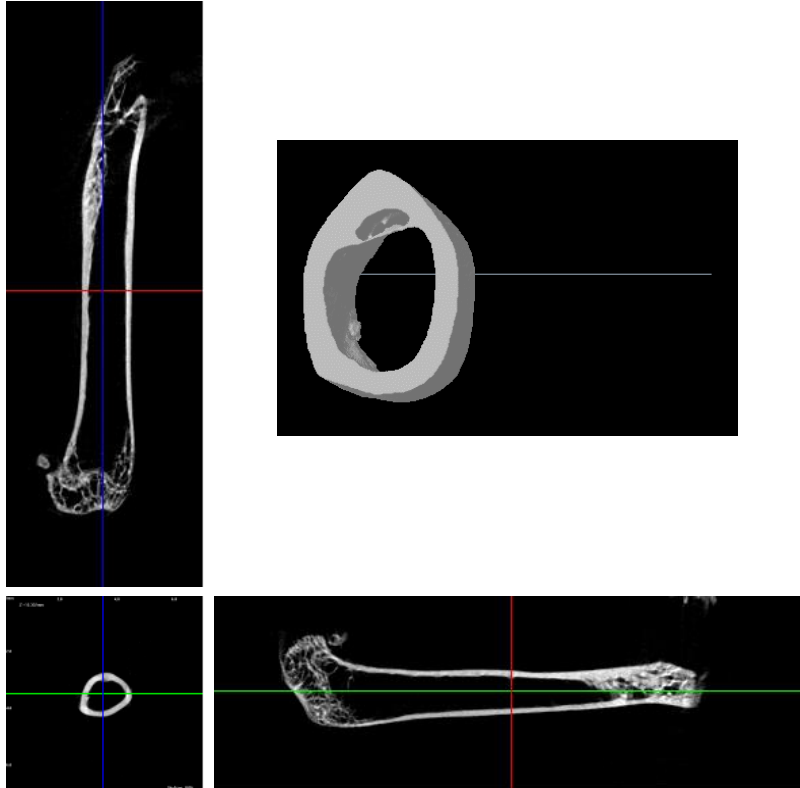


RANKL: receptor activator of NFκB ligand (osteoclastogenic regulator)
OPG: osteoprotegerin (bone turnover regulator)

CTX-1: c-terminal telopeptide of type-1 collagen (soluble marker of bone resorption)
OC: osteocalcin (soluble marker of bone formation)



Femur bone morphometric parameters



Micro-Computed Tomography Scans

- Bone volume (BV)
- Total cross-sectional area (B.Ar)
- Total cross-sectional perimeter (B.Pm)
- Cross-sectional thickness (Cs.Th)

- Trabecular thickness (Tb.Th)
- Trabecular separation (Tb.Sp)
- Trabecular number (Tb.N)

- Bone volume fraction (BV/TV)*



HIV-1 transgene expression in mice alters bone homeostasis

Normal bone development

Bone malformation

Bone loss

HIV-1 transgene expression in mice increases markers of bone turnover

Increased gene expression and circulating levels of osteoclastogenic markers

Wild-type
 Tg26 +/-

The HIV-Tg26 mouse model largely mimics bone alterations in HIV-1 patients

Alterations in HIV-patients without treatment (naïve) evidence the pathogenic capacity of infection-related events on adipose tissue, bone and metabolic homeostasis

***In vitro* studies reveal that adipocytes are targets of HIV-proteins and inflammatory signals, as occurring in HIV-patients**

The HIV-Tg26 mouse may be useful as a model to study alterations in patients non-attributable to cART