



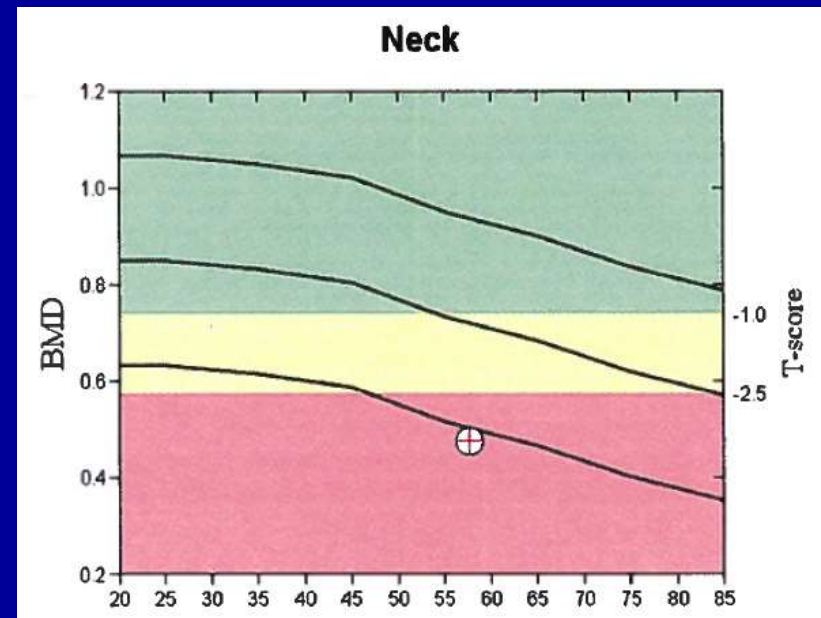
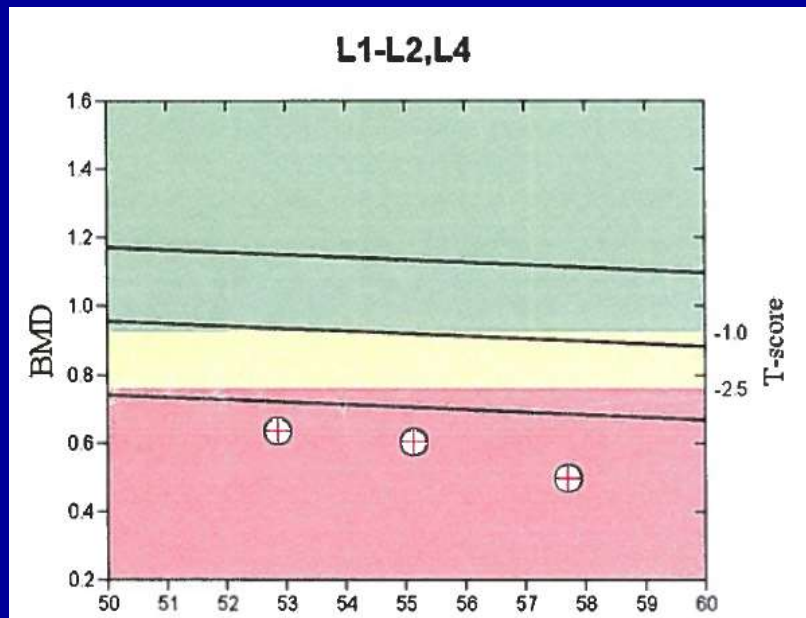
# Conduite devant une DMO très basse

Serge Ferrari

Service des maladies osseuses

Hôpitaux universitaires et Faculté de médecine de Genève

Mme H., fracture vertébrale à 40 ans en jouant au badminton, perte de taille 10 cm depuis lors...



IBN → TPT

### DXA Results Summary: L1-L2,L4

| Messungsdatum | Alter | BMD (g/cm <sup>2</sup> ) | T - Score | BMD-Änderung   |               |
|---------------|-------|--------------------------|-----------|----------------|---------------|
|               |       |                          |           | Vgl. mit Basis | Vgl. Vorherig |
| 26.09.2017    | 57    | 0.498                    | -4.9      | -21.4%#        | -17.6%#       |
| 02.03.2015    | 55    | 0.604                    | -3.9      | -4.6%*         | -4.6%*        |
| 15.11.2012    | 52    | 0.633                    | -3.6      |                |               |

# Mme H, suite

- Parodontolyse juvénile, rétro-maxillaire opéré
- Cervicalgies chron. s/ ostéochondrose C5/C6
- Acro-ostéolyse phalanges distales
- Hyperlaxité
- 154 cm (était 164), 58kg
- Ménopause 52 ans non-substituée, pas d'autres FR OP
- Bilan sg sp: 0 Sy. Inflamm, 25OHD 105, PTH et Ca N, creat N, CTx 0.19, PAL 54





REVIEW

Open Access

# Hajdu-Cheney syndrome: a review

Ernesto Canalis\* and Stefano Zanotti

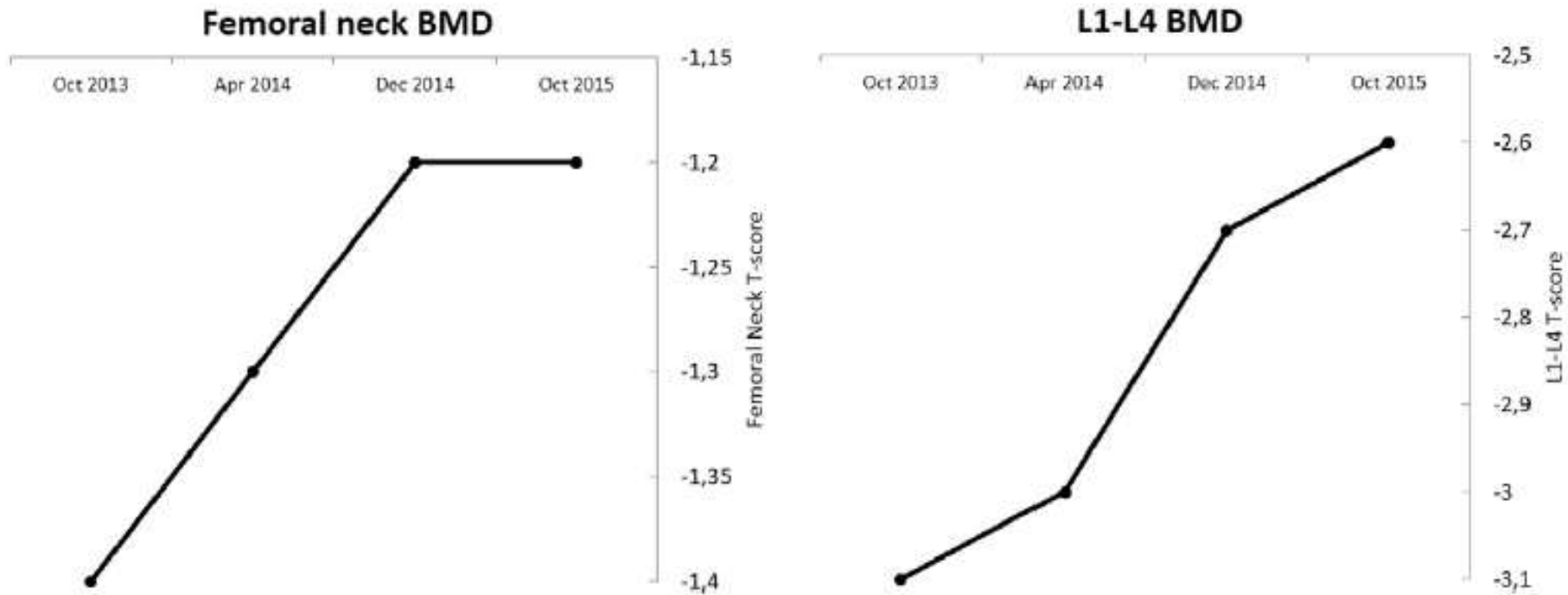
## Abstract

Hajdu Cheney Syndrome (HCS), Orpha 955, is a rare disease characterized by acroosteolysis, severe osteoporosis, short stature, specific craniofacial features, wormian bones, neurological symptoms, cardiovascular defects and polycystic kidneys. HCS is rare and is inherited as autosomal dominant although many sporadic cases have been reported. HCS is associated with mutations in exon 34 of *NOTCH2* upstream the PEST domain that lead to the creation of a truncated and stable *NOTCH2* protein with enhanced *NOTCH2* signaling activity. Although the number of cases with *NOTCH2* mutations reported are limited, it would seem that the diagnosis of HCS can be established by sequence analysis of exon 34 of *NOTCH2*. Notch receptors are single-pass transmembrane proteins that determine cell fate, and play a critical role in skeletal development and homeostasis. Dysregulation of Notch signaling is associated with skeletal developmental disorders. There is limited information about the mechanisms of the bone loss and acroosteolysis in HCS making decisions regarding therapeutic intervention difficult. Bone antiresorptive and anabolic agents have been tried to treat the osteoporosis, but their benefit has not been established. In conclusion, Notch regulates skeletal development and bone remodeling, and gain-of-function mutations of *NOTCH2* are associated with HCS.

**Keywords:** Notch, Skeleton, Bone remodeling, Hajdu-Cheney syndrome, Fractures, Polycystic kidneys, B cell lymphoma

# Hajdu-Cheney: Response to denosumab (1 case report)

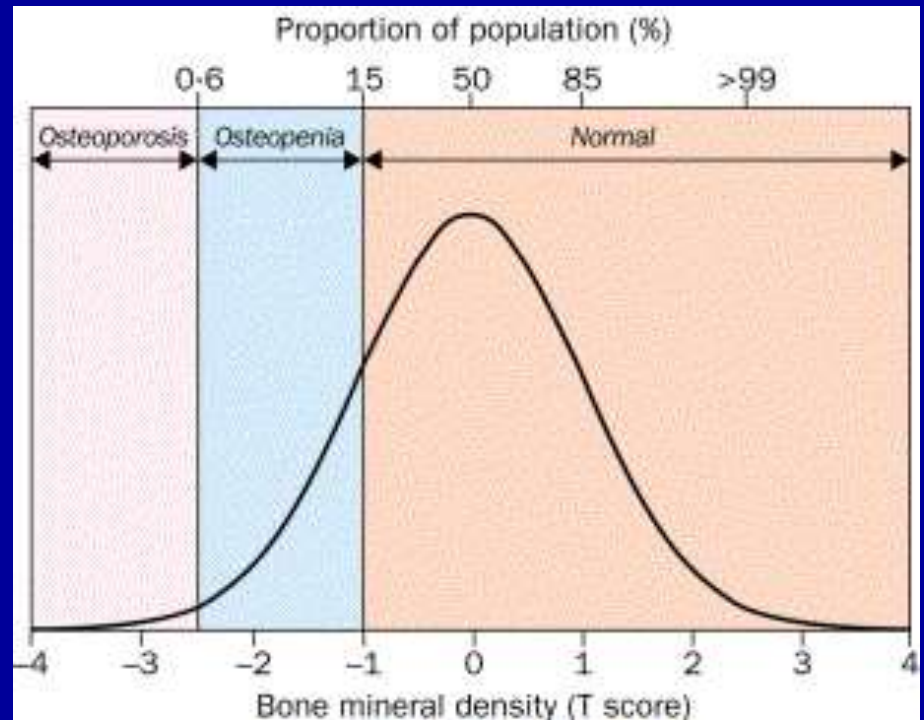
*G. Adami et al. / Bone 92 (2016) 150–156*





# BMD distribution

- Prevalence of osteoporosis in the general population:
  - Osteoporosis:  $\leq 0.5\%$
  - Osteopenia: 10-15%



(Kanis, OI 1997; Liu, JBMM 2008;  
Diaz Curiel, Med Clin 2001)

- Osteoporosis in subjects with chronic diseases:
  - 15-50%

# Osteoporosis in the young: Definition (IOF)

- T-score  $\leq -2.5$  (spine or hip) in men and women  $> 20$  yrs (when growth is completed)

And/or

- Low trauma and/or multiple fractures (vertebrae !)
- In absence of low trauma and/or multiple fractures
  - If
  - A chronic disease (secondary osteoporosis)
- Excluding vitamin D deficiency (osteomalacia)

S Ferrari, ML Bianchi, JA Eisman, AJ Foldes, S Adami, J Stepan, MC de Vernejoul, JM Kaufm, for the Osteoporosis Pathophysiology Working group  
*Osteoporosis Int 2012*

# Masse osseuse très basse chez la femme jeune

JCEM 2008

**TABLE 1.** Body weight composition, BMD, nutritional and food intake parameters, and several hormones (mean levels  $\pm$  SEM) in AN, CT, and controls

|                                       | AN (n = 44)                   | CT (n = 25)                   | Controls (n = 28) |
|---------------------------------------|-------------------------------|-------------------------------|-------------------|
| Anthropometry and body composition    |                               |                               |                   |
| Age (yr)                              | 23.4 $\pm$ 1.2                | 23.1 $\pm$ 1.2                | 23.9 $\pm$ 1.4    |
| Height (m)                            | 1.62 $\pm$ 0.1                | 1.63 $\pm$ 0.02               | 1.63 $\pm$ 0.01   |
| BMI (kg/m <sup>2</sup> )              | 15.5 $\pm$ 0.1 <sup>a</sup>   | 15.8 $\pm$ 0.1 <sup>a</sup>   | 20.7 $\pm$ 0.4    |
| FM %                                  | 9.8 $\pm$ 1.1 <sup>a</sup>    | 18.6 $\pm$ 0.7 <sup>a,b</sup> | 26.3 $\pm$ 1.2    |
| BMD                                   |                               |                               |                   |
| Femoral neck BMD (g/cm <sup>2</sup> ) | 0.795 $\pm$ 0.03 <sup>a</sup> | 0.809 $\pm$ 0.02 <sup>a</sup> | 0.951 $\pm$ 0.02  |
| Lumbar spine BMD (g/cm <sup>2</sup> ) | 0.849 $\pm$ 0.03 <sup>a</sup> | 0.873 $\pm$ 0.02 <sup>a</sup> | 0.986 $\pm$ 0.03  |
| Hormonal parameters                   |                               |                               |                   |
| Leptin ( $\mu$ g/liter)               | 2.4 $\pm$ 0.5 <sup>a</sup>    | 6.0 $\pm$ 0.8 <sup>a,b</sup>  | 11.2 $\pm$ 1.9    |
| GH (mIU/liter)                        | 8.5 $\pm$ 0.7 <sup>a</sup>    | 4.8 $\pm$ 0.6 <sup>b</sup>    | 4.7 $\pm$ 0.6     |
| IGF-I ( $\mu$ g/liter)                | 163 $\pm$ 16 <sup>a</sup>     | 295 $\pm$ 34 <sup>b</sup>     | 283 $\pm$ 20      |
| Cortisol (ng/liter)                   | 364 $\pm$ 31 <sup>a</sup>     | 216 $\pm$ 12 <sup>b</sup>     | 266 $\pm$ 17      |
| 17 $\beta$ -estradiol (ng/liter)      | 14.3 $\pm$ 1.4 <sup>a</sup>   | 73.1 $\pm$ 8.6 <sup>b</sup>   | 51.6 $\pm$ 11.4   |





# Clinical Approach:

Med Hx, Phys exam

DXA (VFA)

Labs (1st line)



|                                      |   |
|--------------------------------------|---|
| Mineral metabolism                   | Serum calcium (corrected for albumin)<br>Serum phosphate<br>Creatinine<br>25(OH)D<br>iPTH<br>ALP (bone specific)<br>BTMs (for instance, s-CTX, s-PINP)* |
| Inflammation, hematopoietic disorder | Blood cell count<br>ESR or CRP  |
| Hepatic disease                      | GOT, GPT, $\gamma$ -GT  |
| Diabetes (primary or secondary)      | Fasting glucose, Hba1C  |
| Thyroid dysfunction                  | TSH   |
| Hypogonadism (men)                   | Total testosterone  |
| Malabsorption, Celiac disease        | 24-h urinary calcium<br>Anti-endomysial , anti -transglutaminase  |

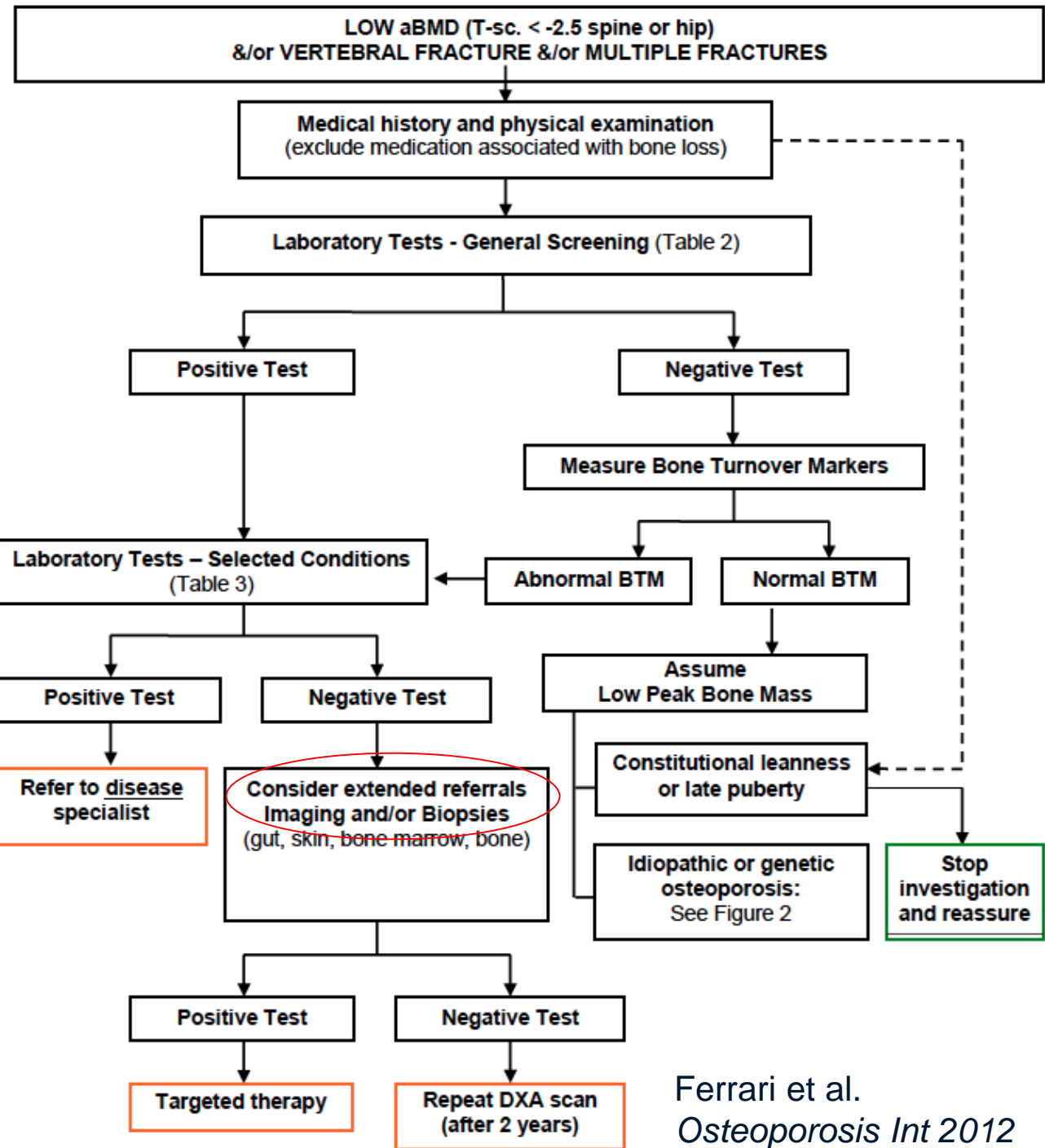
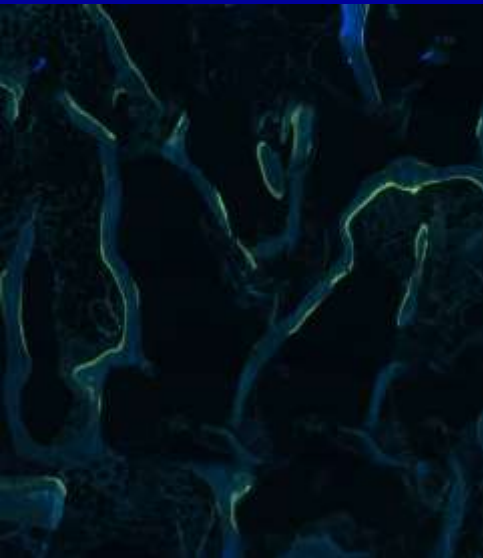
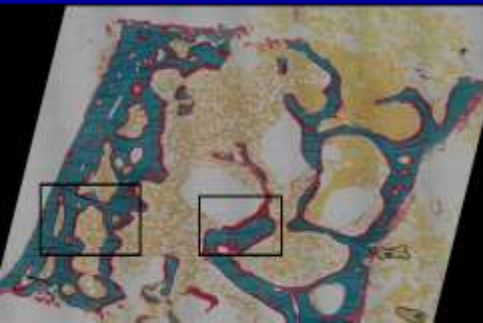


# Clinical Approach:

Labs (selected conditions)



| Background                       | Test                                 |
|----------------------------------|--------------------------------------|
| TSH alterations                  | Free T4                              |
| Altered glucose, Cushing's       | 24-hour free urinary cortisol        |
| Altered testosterone (men)       | LH /SHBG (free testosterone)         |
| Amenorrhea, hypogonadism (women) | FSH/estradiol                        |
| Altered renal function (CRF)     | 1,25(OH) <sub>2</sub> D <sub>3</sub> |
| Hemochromatosis                  | Ferritin                             |
| Hypophosphatasia                 | ALP, BALP                            |
| Mastocytosis                     | Tryptase, IgE                        |
| Gaucher's disease                | Glucocerebrosidase                   |

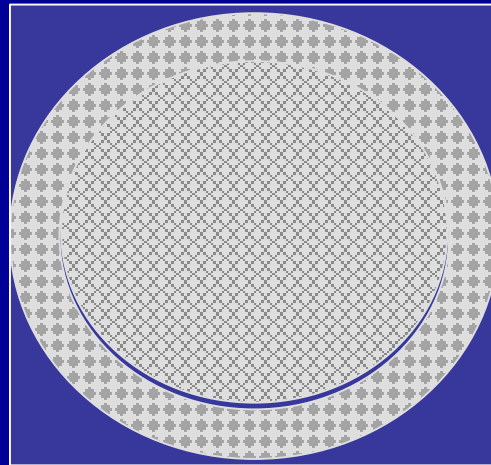


# Mme B., 70 ans

- Minéralo de “routine” à 65 ans: T-sc -4.6 rachis, -2.7 hanche
- Pas de fractures
- Ménopause non-substituée à 52 ans
- Pas de FR OP , histoire familiale ou Sy. Inflamm
- 150 cm / 47kg
- 25OHD 65, CTx 0.76, P1NP 55

# What is bone mass (aBMD) ?

- Degree of mineralization (max 1200mg/cm<sup>3</sup>)
- Trabecular bone volume
- Cortical thickness and porosity
- Bone size



aBMD = how much  
Bone mineral there is  
in the square

Hence Bone quantity and quality are inter-related !

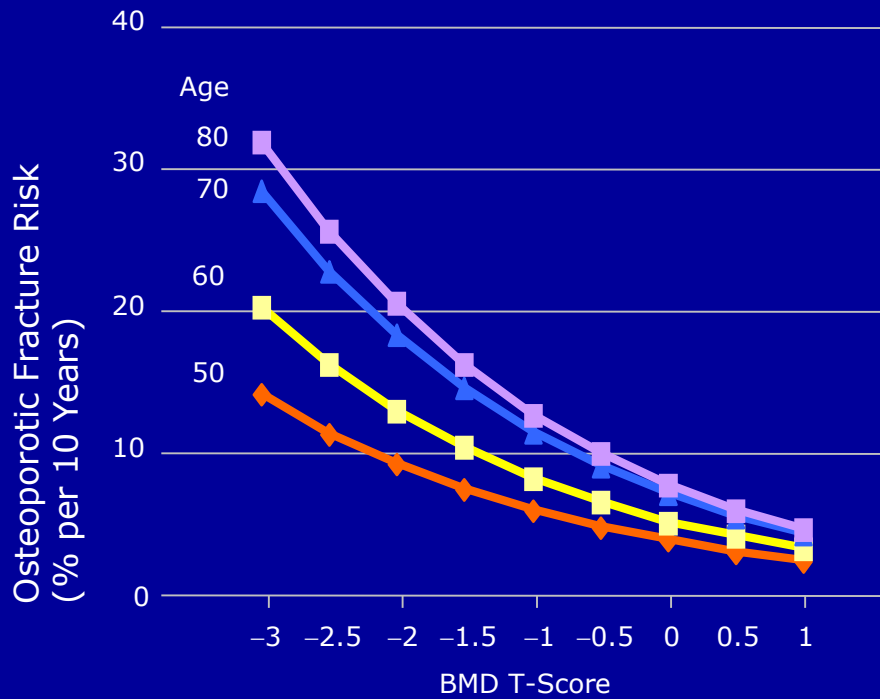


# Correlations between spine BMD and microstructure (ex vivo)

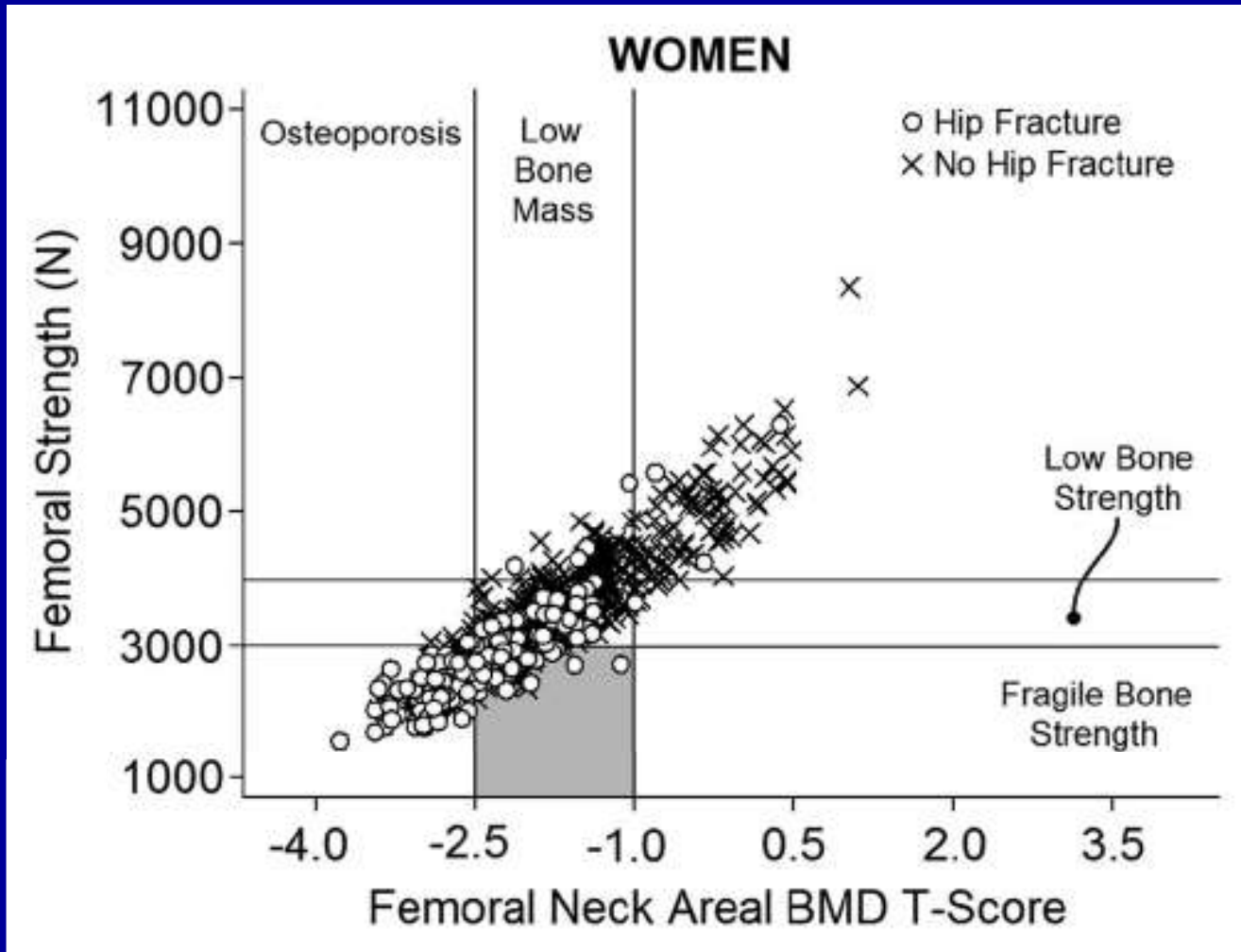
**Table 2** Pearson correlation coefficients between DXA, microarchitecture parameters, mechanical behavior, and TBS

|              | AP. BMC | AP. BMD | AP. area | Lat. BMC | Lat. area | Lat. BMD | Tb.BV/TV | DA       | SMI     | Tb.Th | Failure load | Stiffness |
|--------------|---------|---------|----------|----------|-----------|----------|----------|----------|---------|-------|--------------|-----------|
| AP. BMD      | 0.72**  |         |          |          |           |          |          |          |         |       |              |           |
| AP. area     | 0.66**  | -0.03   |          |          |           |          |          |          |         |       |              |           |
| Lat. BMC     | 0.86*** | 0.50*   | 0.73***  |          |           |          |          |          |         |       |              |           |
| Lat. BMD     | 0.91*** | 0.70**  | 0.56*    | 0.86***  |           |          |          |          |         |       |              |           |
| Lat. area    | 0.42    | 0.00    | 0.64**   | 0.73***  |           |          |          |          |         |       |              |           |
| Tb.BV/TV     | 0.49    | 0.68**  | -0.05    | 0.22     | -0.17     | 0.44     |          |          |         |       |              |           |
| DA           | -0.32   | -0.49   | 0.11     | -0.23    | -0.06     | -0.24    | -0.69**  |          |         |       |              |           |
| SMI          | -0.32   | -0.68** | 0.29     | 0.03     | 0.48      | -0.33    | -0.85*** | 0.40     |         |       |              |           |
| Tb.Th        | 0.16    | 0.40    | -0.22    | 0.16     | 0.02      | 0.19     | 0.49*    | -0.87*** | -0.28   |       |              |           |
| Failure load | 0.41    | 0.34    | 0.26     | 0.27     | -0.13     | 0.49*    | 0.33     | 0.25     | -0.56*  | -0.37 |              |           |
| Stiffness    | 0.49*   | 0.24    | 0.44     | 0.32     | -0.07     | 0.52*    | 0.32     | 0.22     | -0.36   | -0.46 | 0.72**       |           |
| TBS          | 0.25    | 0.36    | -0.04    | -0.01    | -0.32     | 0.24     | 0.58*    | -0.09    | -0.62** | -0.03 | 0.46         | 0.64**    |

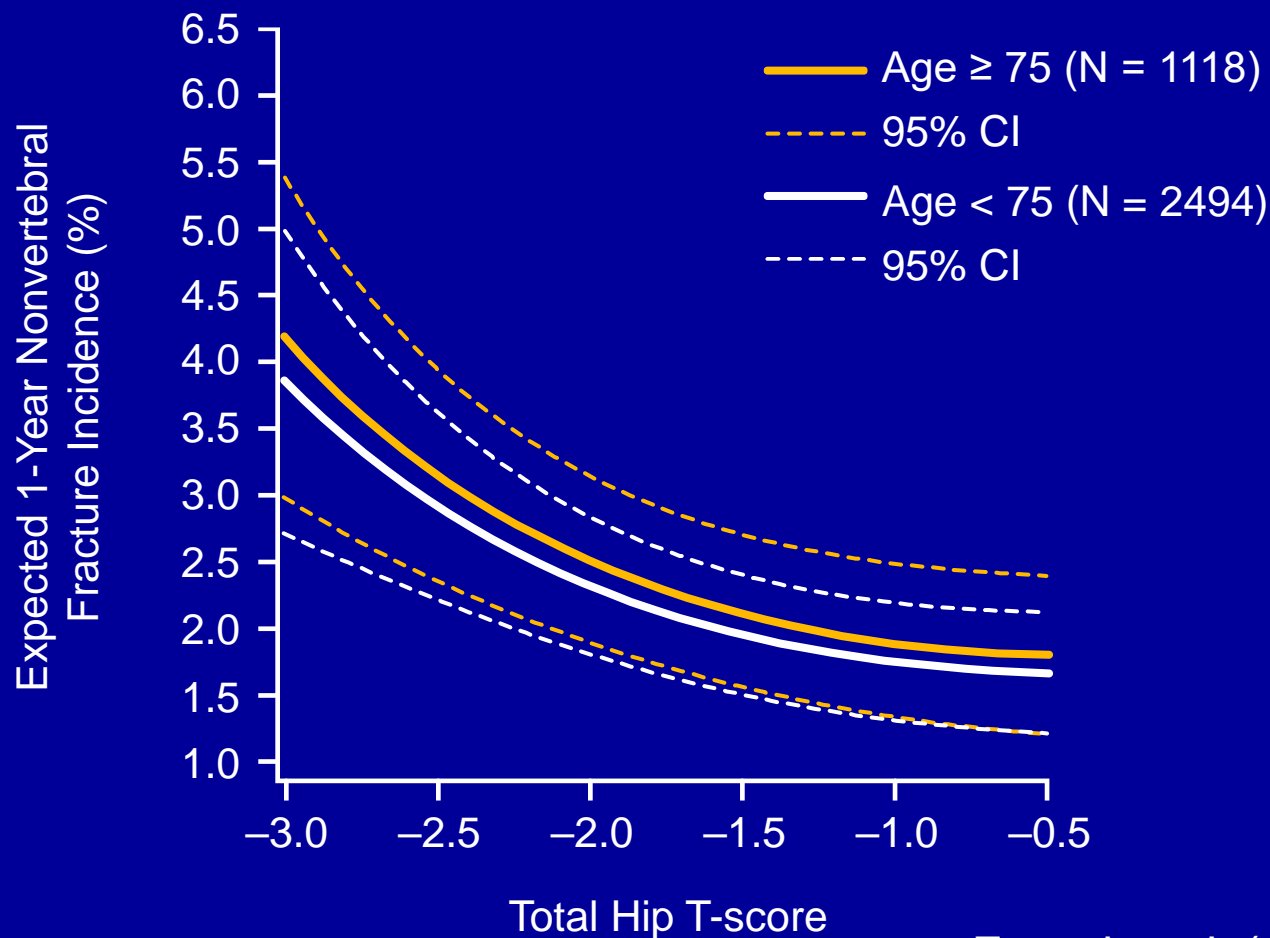
# DMO basse et risque fracturaire



# aBMD vs CT-derived hip strength (FEA) and incident hip fractures

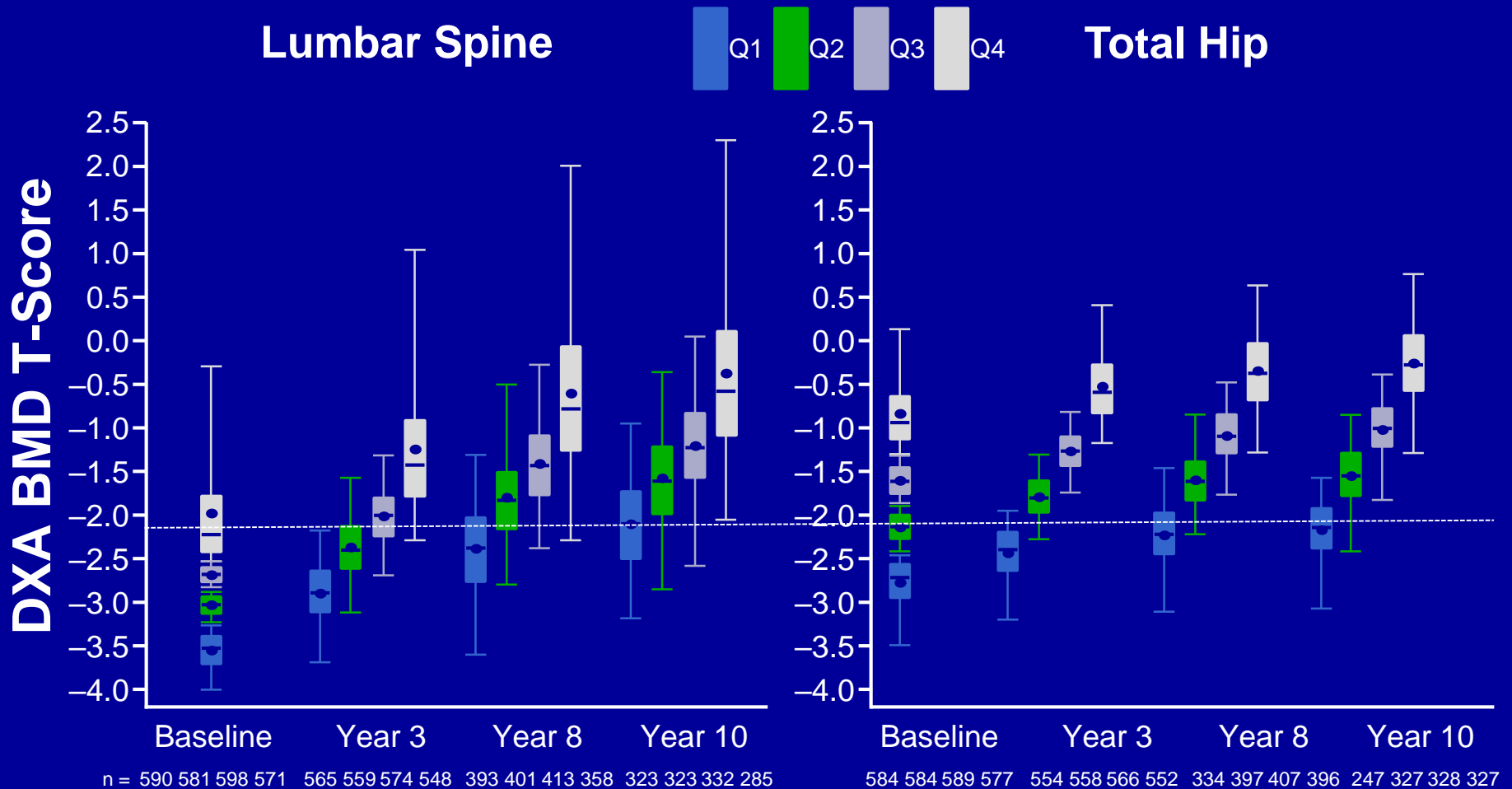


# Relationship Between Total Hip T-score and Nonvertebral Fracture by Age in denosumab-treated subjects: Freedom extension



Ferrari et al. (submitted)

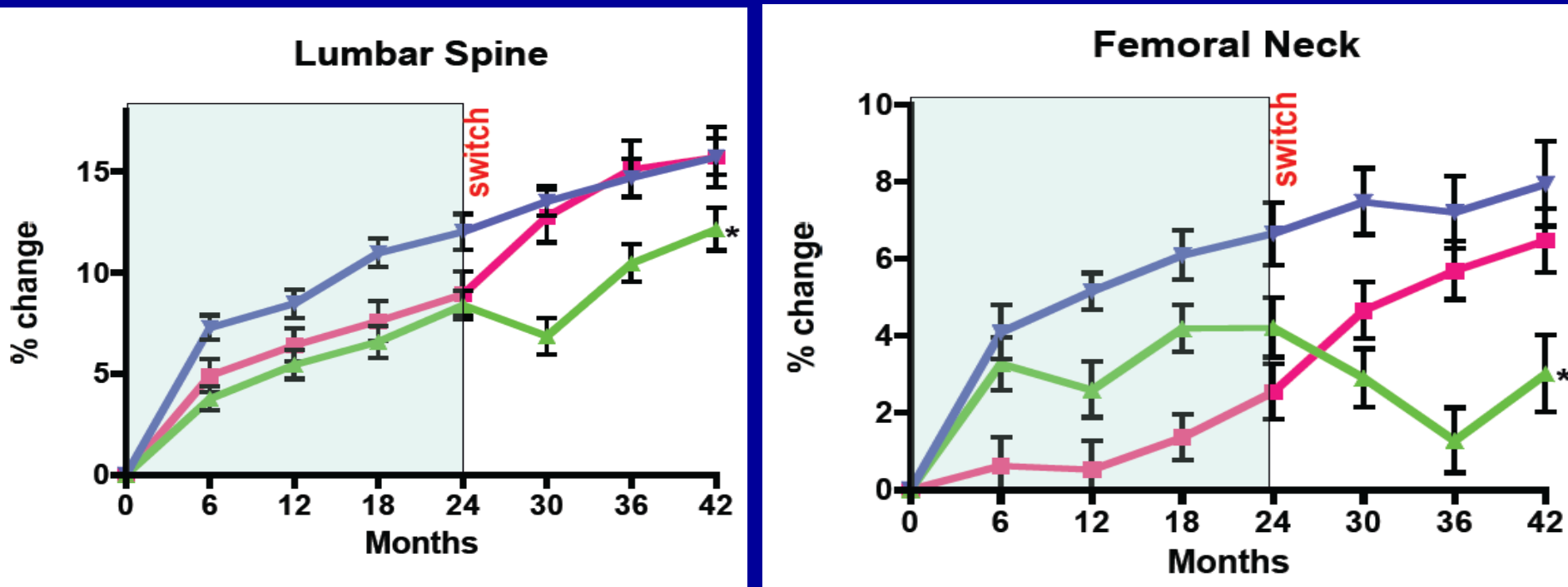
# Improvement in BMD T-scores Remained Largely Consistent With the Respective Baseline BMD T-score Quartile



- The T-scores showed a similar magnitude of improvement in BMD across subjects regardless of their initial BMD



# Transition from Denosumab to TPTD or from TPTD to Denosumab in Postmenopausal Women with Osteoporosis: The DATA-Switch Study

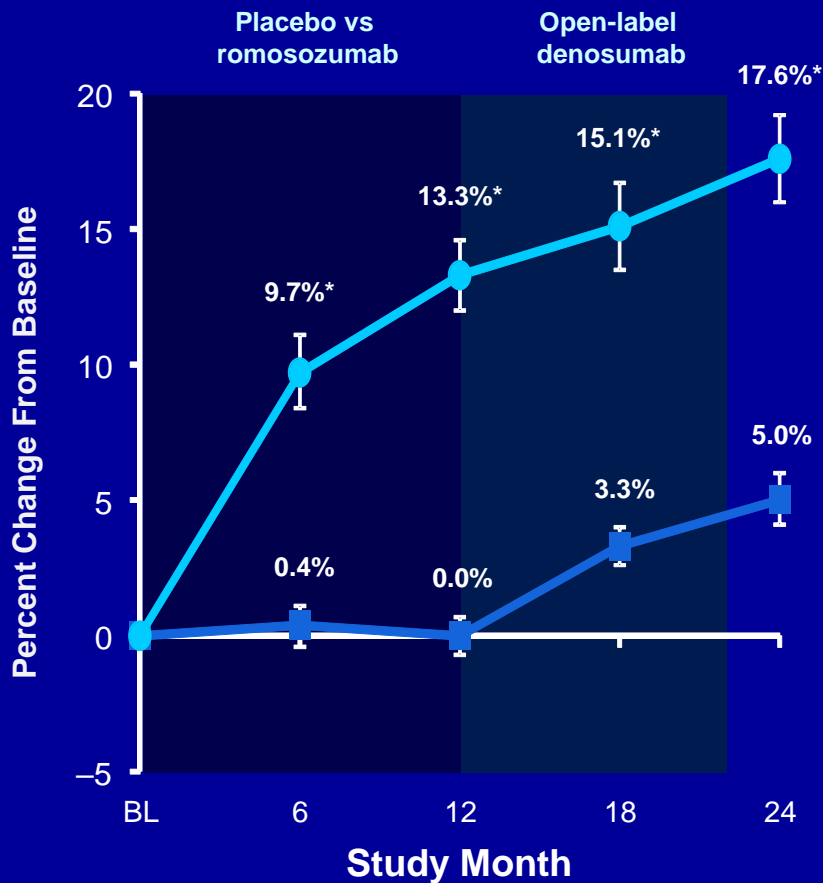


■ 24-mo TPTD, 18-mo DMAB  
▲ 24-mo DMAB, 18 mo TPTD  
◆ 24-mo Both, 18-mo DMAB

# FRAME: Lumbar Spine and Total Hip BMD Through Month 24

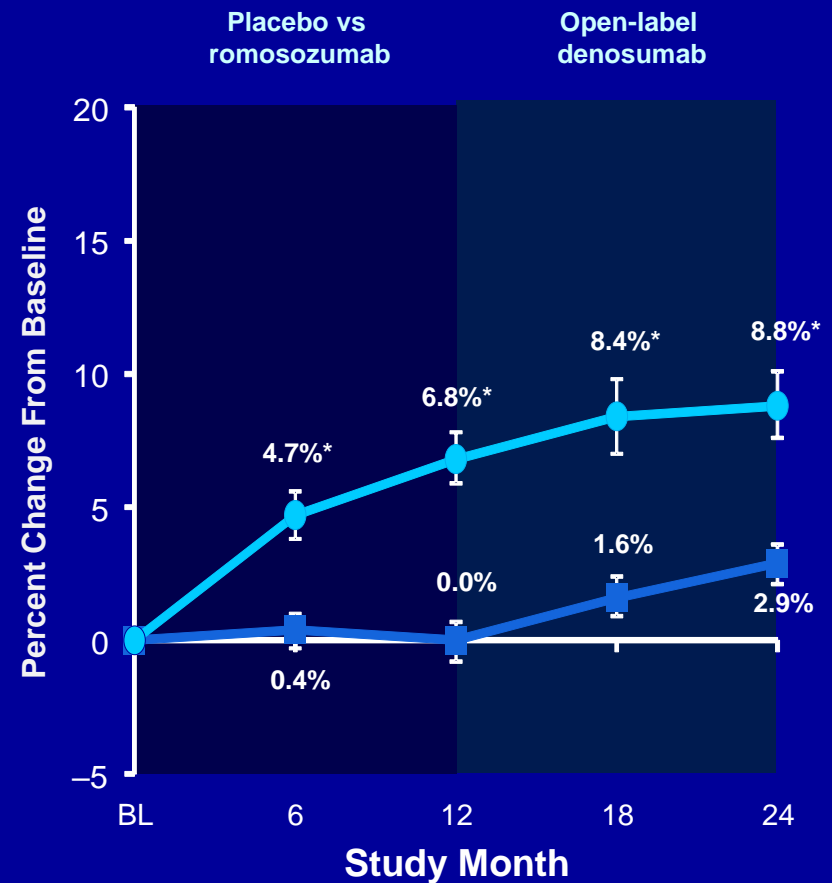
Placebo-to-denosumab (N = 61)  
 Romosozumab-to-denosumab (N = 65)

## Lumbar Spine



Placebo-to-denosumab (N = 62)  
 Romosozumab-to-denosumab (N = 66)

## Total Hip



\*p < 0.001 compared with placebo (M6 and M12) or placebo/denosumab (M18 and M24)  
 Data are least square mean (95% CI) adjusted for relevant baseline covariates

# Conclusions

- DMO très basse avant la ménopause:
  - faible pic de masse osseuse > hyperremodellage
  - cause 2<sup>re</sup> ou génétique > OP idiopathique ou constitution maigre
  - Dans la mesure du possible traiter la cause
  - Traiter l'OP...si évidence de remodelage très haut ou bas (role de la biopsie) ou fragilité osseuse avérée
- DMO très basse après la ménopause:
  - hyperremodellage presque toujours en cause
  - Altérations microstructurelles cohérentes avec la DMO (=vraie définition de l'OP)
  - Traitements séquentiels ou combinés AR + OF