

Multivariate diagnostic model for osteoarthritis highlights the importance of osteoclastogenesis and expression of interleukin 1 receptors

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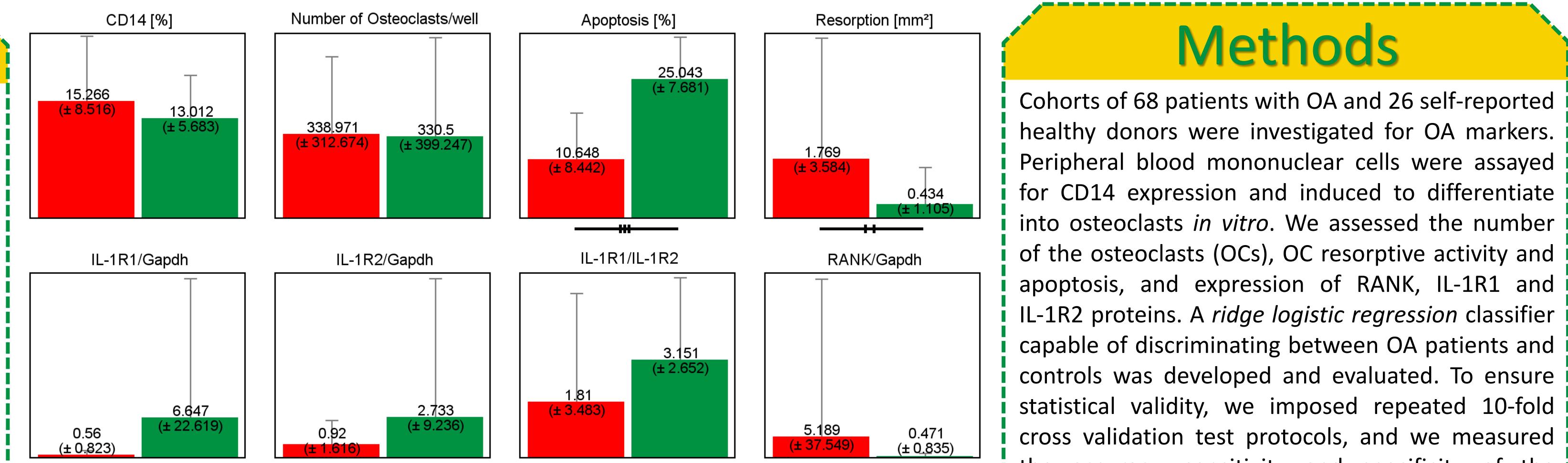


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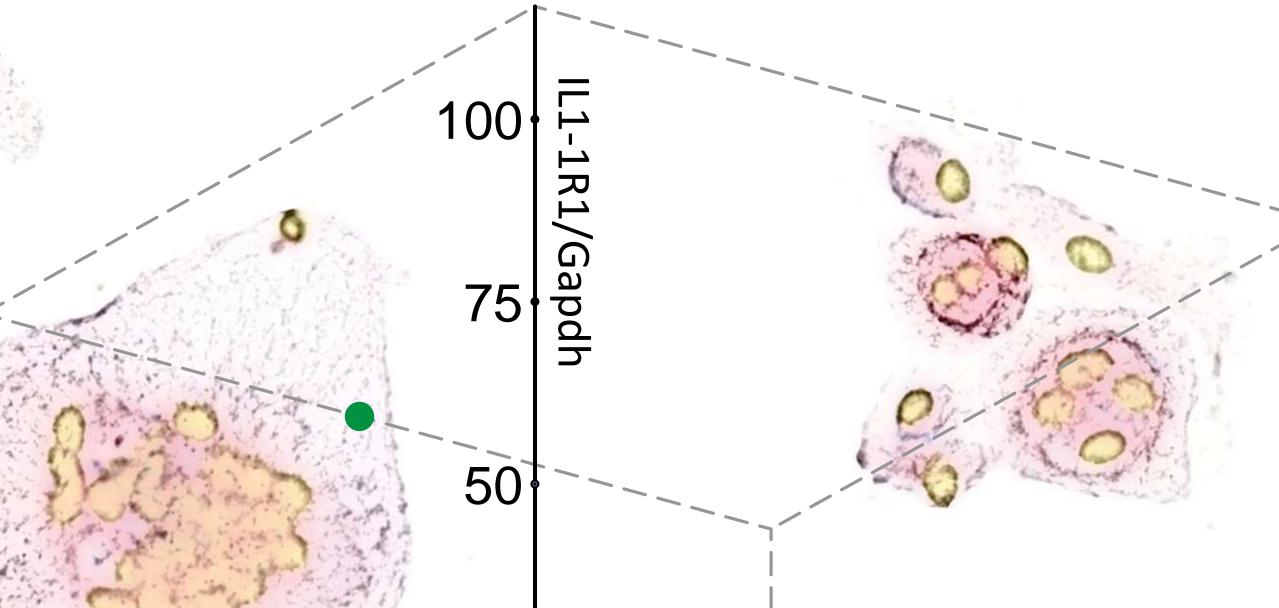
Introduction

Bone is a central element in the pathophysiology of degenerative and inflammatory arthropathies. Subchondral bone sclerosis, bone cysts and osteophytes are hallmarks of the osteoarthritis (OA). There is indirect evidence implicating



osteoclasts (OCs) in the pathogenesis of OA. The involvement of OCs in bone and joint destruction has been confirmed, yet their role in disease onset and progression has not been previously addressed. We have observed that the capacity for in vitro osteoclastogenesis varies widely in a normal human population, distinguishing two differentiators, subgroups, high and low regardless of age, gender, weight, or other demographic variables. We hypothesize that variations in osteoclastogenic capacity in OA population could underlie variation in predisposition for this disease and its severity.

Statistical analysis showing the difference in mean values between the OA (red) and control (green) groups for the statistically significant markers. * P value < 0.05, ** P value < 0.01, *** P value < 0.001; T-test was used for normal variables, otherwise the non-parametric Mann-Whitney test was used; Shapiro-Wilk test was used to verify normality.



of the osteoclasts (OCs), OC resorptive activity and apoptosis, and expression of RANK, IL-1R1 and IL-1R2 proteins. A ridge logistic regression classifier capable of discriminating between OA patients and controls was developed and evaluated. To ensure statistical validity, we imposed repeated 10-fold cross validation test protocols, and we measured the accuracy, sensitivity and specificity of the model. We also performed classical statistical analysis to validate the significance of the univariate markers.

Results

We found that OC parameters (in particular OC apoptosis) and expression of IL-1R1 and IL-1R2 can be used to build a well-performing multivariate diagnostic model for OA. Our regression model provides strong diagnostic input, with cross validation-based accuracy of about 87%. The multivariate model provides a significant improvement (p < 0.001) compared to the bestperforming univariate model, indicating that OC parameters and expression of interleukin 1 receptors provide complementary diagnostic information. 100 x 10 cv Self check Specificity Accuracy Sensitivity Specificity Sensitivity Inputs Accuracy Apoptosis 83.8% (±0.7) 53.8% 95.4% (±0.8) 53.4% (±1.6) 95.6% 84.0% + IL-1R1 85.5% (±0.9) 94.9% (±1.1) 61.0% (±1.8) 86.2% 95.6% 61.5%

osteoclastogenic phenotype and expression of the presence of OA. Our aim is to investigate, using modern data mining techniques, the possible multivariate relationships between osteoclastogenic characteristics and protein expression in OA patients compared to a cohort of self-reported normal individuals to develop a wellperforming diagnostic model.

