Supplementary Information

Association between Oncostatin M expression and inflammatory phenotype in experimental arthritis models and osteoarthritis patients

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Citation: Lastname, F.; Lastname, F.; Lastname, F. Title. *Cells* **2021**, *10*, x. https://doi.org/10.3390/xxxxx

Academic Editor: Alessandra Colombini

Received: 18 January 2021 Accepted: 23 February 2021 Published: date

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Table 1. Reliability assessment between PGPS samples stained with and without antigen retrieval.

Structure	Intra-class correlation —	Confidence Interval (95 %)		
		Lower limit	Upper limit	
Cartilage	0.870	0.671	0.945	
Periosteum	0.919	0.826	0.962	
Synovium	0.922	0.536	0.987	
Meniscus	0.893	0.584	0.973	
Ligaments	0.975	0.805	0.996	



Figure 1. Effect of antigen retrieval on OSM scoring in the PGPS model samples. OSM score per anatomical structure: cartilage (**a**), periosteum (**b**), meniscus (**c**), synovium, and ligaments (**d**). Control (C), Induced (I). 0 = no staining, 1 = slight staining, 2 = moderate staining, 3 = extensive staining. Data are presented as median and 95 % CI.

Due to limited availability of samples, seven samples from the PGPS model were stained for OSM with and without antigen retrieval. OSM was scored as described before (0-3) for cartilage, periosteum, synovium, meniscus and ligaments. Scoring was done in a random order by two independent observers blinded for treatment and staining method, and scores were averaged.

The intra-class correlation coefficient (ICC) was used to calculate the agreement in OSM score between the two staining methods. A two-way mixed-effect model based on a mean-rating (k=2) and absolute agreement was used. ICC estimates and the 95% confidence intervals (CI) were reported. The ICC values are above 0.870 for all the joint structures, indicating that antigen retrieval does not affect the scoring for the PGPS model. Thus, the samples without antigen retrieval were used.



Figure 2. Negative control with mouse IgG isotype. Representative pictures of different joint structures: Femur condyle (FC) cartilage, meniscus, periosteum and synovium. Scale bar: 50 μ m.

Table 2. Joint	compartments.
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Structure	Compartment	
	Lateral femoral condyle	
Cartilago, short drogstop	Medial femoral condyle	
Cartilage - chondrocytes	Lateral tibia plateau	
	Medial tibia plateau	
	Lateral femur	
Dariastour	Medial femur	
renosteum	Lateral tibia	
	Medial tibia	
Moniague	Lateral	
Weniscus	Medial	
Synovium	Femorotibial synovium	
Ligaments	Collateral ligaments	



Figure 3. Hematoxylin & eosin staining of a rat joint. Histological overview of the rat joint with higher magnification pictures of the (medial side) compartments. For scoring purposes both medial and lateral compartments were taken into account. Scale bar = $50 \mu m$ (figures on the left).



Molecular weights (kDa)	Clone B-6	Clone A-9
Recombinant rat OSM (40 ng)	26.17	25.38
Recombinant rat OSM (20 ng)	25.95	25.49
PC-12	33.95	32.10
Primary rat chondrocytes	34.52	32.87

Figure 4. Western blot analysis showing binding of two different anti-OSM antibodies to the OSM protein recombinant rat OSM, PC-12 cells, and primary rat chondrocytes.



Figure 5. Intracellular OSM expression in chondrocytes in the PGPS and ACLT models. Scale bar: 10 $\mu m.$



Figure 6. Subchondral bone sclerosis. Subchondral bone thickness measured by μ CT in the PGPS (a) and ACLT + pMMx (b) models. Subchondral bone thickness was measured in the medial and lateral compartments of both femur condyle and tibia plateau.



Figure 7. Trap staining. Representative pictures of different joint structures quantified for TRAP staining: periosteum, growth plate and subchondral bone.



Figure 8. Quantification of TRAP staining. TRAP staining was performed in the knee sections of the PGPS (a and b), and ACLT + pMMx (c and d) models. TRAP staining was scored in periosteum (a and c), growth plate (GP) and subchondral bone (SB) (b and d) for tibia and femur in both medial (M) and lateral (L) compartments. Scoring was done in a random order by two independent observers blinded for treatment (JPG, AK), and scores were averaged. Data are presented as median and 95% CI.

Table 3. Spearman's	s correlation between	n OSM and	remaining	cytokines.
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	OSM	
Cytokines	Spearman's rho	<i>p</i> value
IL-1α	0.461	0.016*
IL-1β	0.193	0.335
IL-4	0.289	0.144
IL-6	0.350	0.074
IL-8	0.035	0.863
IL-7	0.450	0.019*
IL-10	0.106	0.599
IL-13	0.359	0.066
TNF-α	0.481	0.011*
IFN-Y	0.573	0.002*
IL-1Ra	0.038	0.852

Catalata	Difference	Sig.	95% Confidence Interval	
Cytokine			Upper	Lower
IFN-γ	-94.90	0.001	-39.43	-217.65
IL-1a	-20.91	0.003	-8.93	-45.78
TNF-α	-4.35	0.017	-1.28	-13.30
IL-7	-4.00	0.050	-0.97	-14.03
IL-4	-1.64	0.052	-0.61	-3.66
IL-6	-1743.25	0.094	-150.13	-16669.00
IL-13	-18.74	0.097	-5.13	-52.89
IL-1β	-6.92	0.291	-0.62	-33.31
IL-1Ra	557.27	0.399	968.29	27.67
IL-8	47.28	0.399	27.90	7.05
IL-10	-4.05	0.595	0.59	-38.39

Table 4. Differences in cytokine concentration between patient groups (OSM- and OSM+).