



This work has been fully supported by Croatian Science Foundation under the project 5699.

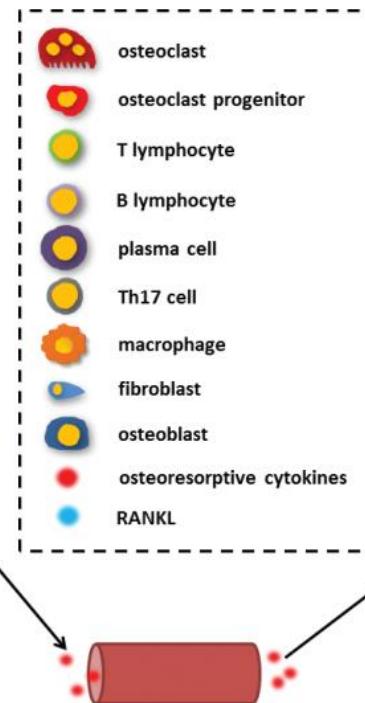
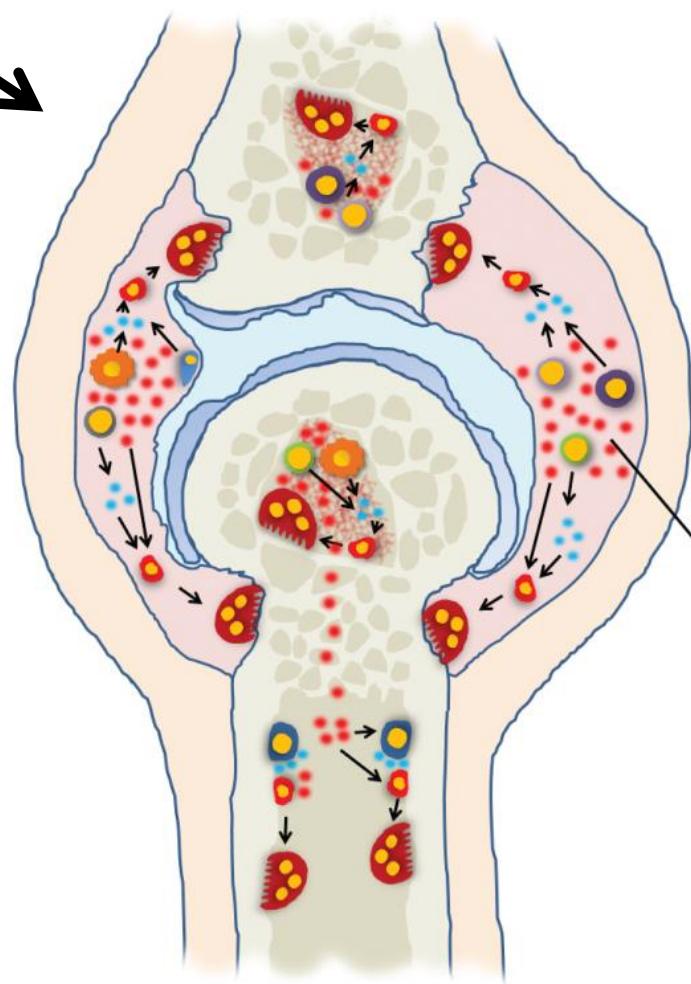
# **Lymphocytic subpopulations associated with disease activity and levels of proinflammatory cytokines in rheumatoid and psoriatic arthritis**

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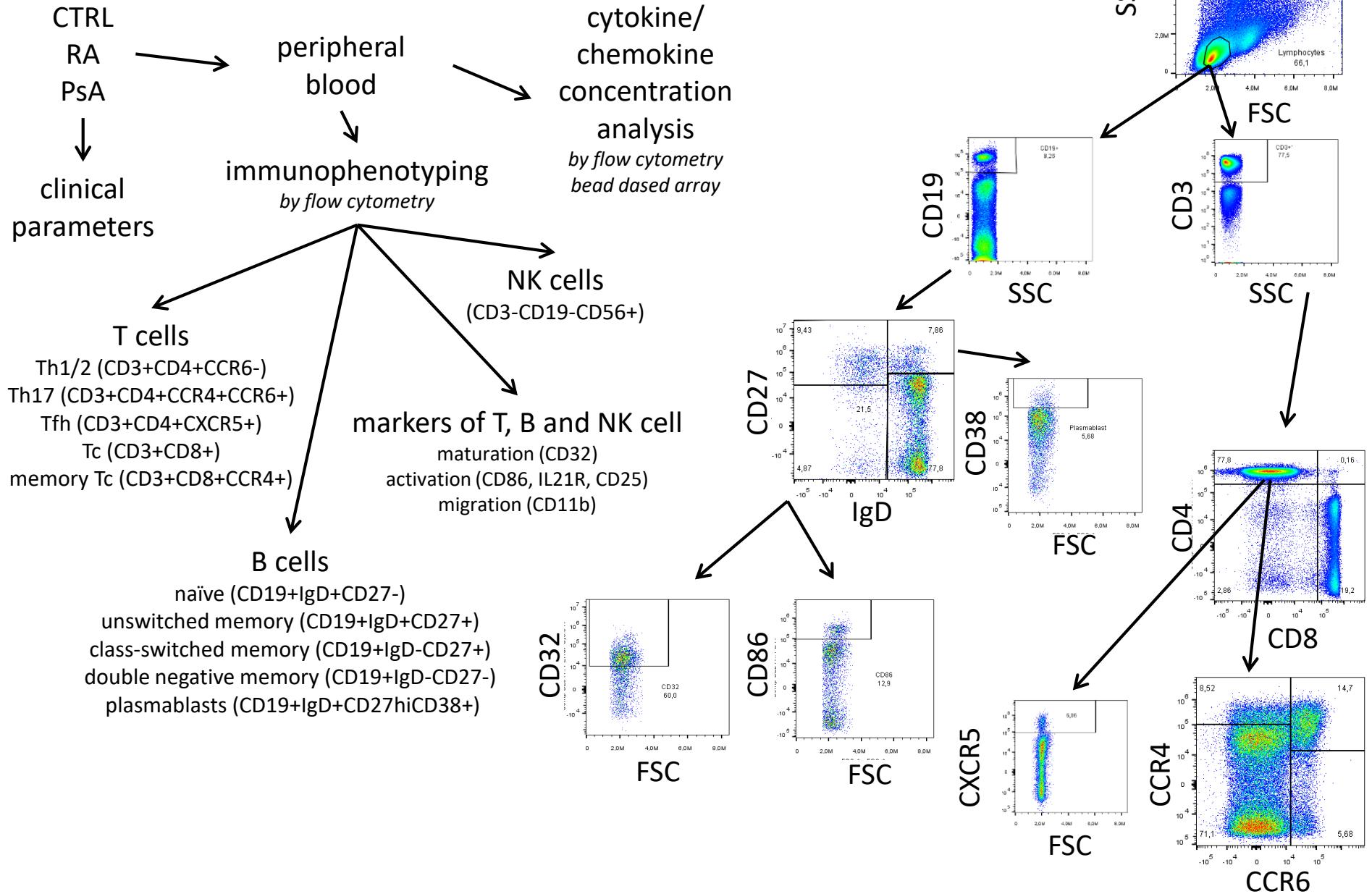
# Rheumatoid arthritis and peripheral psoriatic arthritis



local bone loss

generalized bone loss

# Immunophenotyping



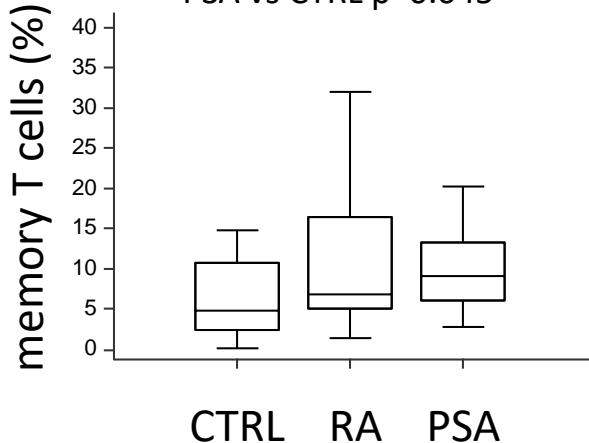
# Patient data

	Rheumatoid arthritis	Psoriatic arthritis	Controls
Age (years)	65 [51-74]	56 [44-74]	65 [48-68]
Sex (male/female)	7/73	21/34	21/89
DAS28	5.97 [4.99-6.71]	5.5 [4.05-6.78]	
DAS28 <5.1 / DAS28 >5.1 (no. of patients)	20/60	23/32	
CDAI	37 [24.03-44.10]	33.6 [14.0-52.08]	
Disease duration (years)	18.5 [9-29]	14 [8-30]	
SE (mm/h)	27.5 [19.0-39.5]	14.5 [4.0-46.5]	
CRP (mg/L)	11.9 [4.75-24.5]	2.7 [0.66-29.13]	
RF (IU/L) (n=57)	79.7 [15.0-239.0]	-	
aCCP (EU/L) (n=33)	68.7 [2.23-270.75]	-	
Seropositive (no. of patients)	44	-	
Tender joint count (out of 28)	15 [8-22]	17 [3-26]	
Swollen joint count (out of 28)	7 [2-12]	6 [0-14]	
Disease activity – physician VAS	7 [5.45-8.78]	6.4 [4.7-9.0]	
Disease activity – patient VAS	7 [5.0-8.8]	6.85 [4.9-8.23]	
NSAID	51	40	
DMARD	51	18	
Corticosteroids	33	6	

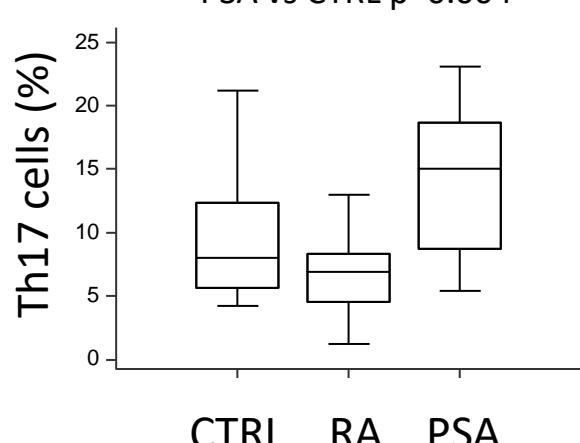
pg/mL	Control		Rheumatoid arthritis	
	Median	IQR	Median	IQR
<b>TNF</b>	70,32	[38,0-93,6]	<b>90,45</b>	[39,0-131,0]
<b>IL2</b>	11,65	[ND-27,3]	<b>30,5</b>	[ND-54,0]
<b>IL4</b>	25,94	[16,0-34,6]	<b>34,15</b>	[23,6-40,9]
<b>IL10</b>	122,8	[90,3-163,6]	<b>175,99</b>	[119,1-211,3]
<b>IL6</b>	330,22	[210,3-428,4]	<b>360,28</b>	[237,5-535,5]
<b>IL17</b>	21,99	[ND-78,4]	<b>81,66</b>	[46,125-121,500]
<b>CCL2</b>	115,9	[17,2-141,3]	<b>142,22</b>	[20,5-239,2]
<b>CCL3</b>	13,19	[7,5-17,0]	<b>21,2</b>	[15,0-23,9]
<b>CCL4</b>	39,86	[30,4-52,840]	<b>60,99</b>	[45,7-69,8]
<b>CCL5</b>	4436,35	[2182,8-7192,2]	<b>12094,74</b>	[5077,3-17134,7]
<b>CXCL9</b>	48,74	[31,4-102,6]	<b>251,57</b>	[150,8-480,8]
<b>CXCL10</b>	81,995	[40,0-141,0]	<b>230,4</b>	[133,6-315,1]

# Altered T lymphocyte subpopulations

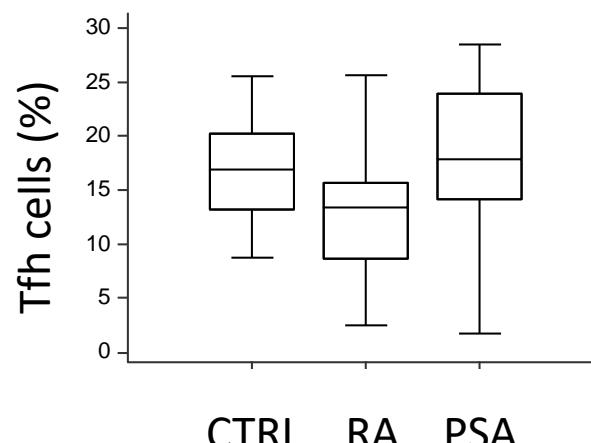
RA vs CTRL p=0.020  
PSA vs CTRL p=0.045



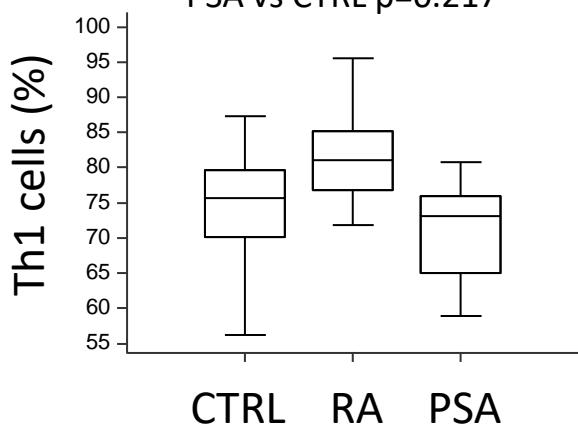
RA vs CTRL p=0.038  
PSA vs CTRL p=0.004



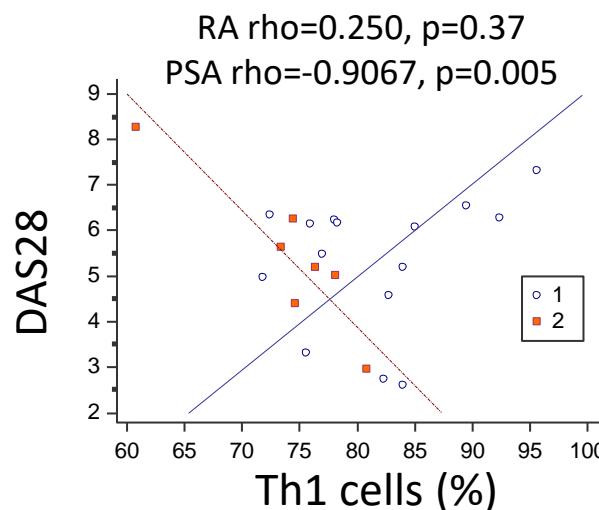
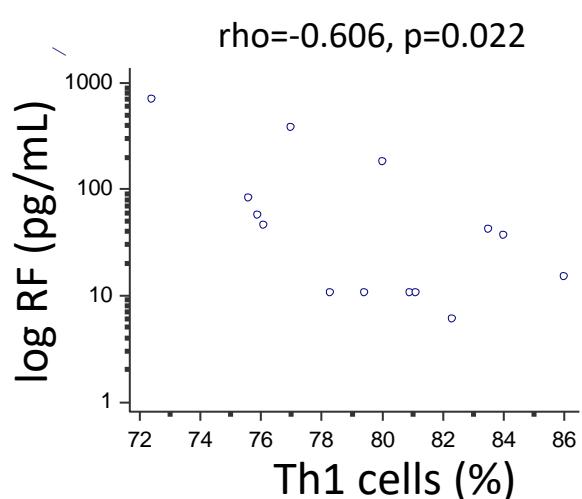
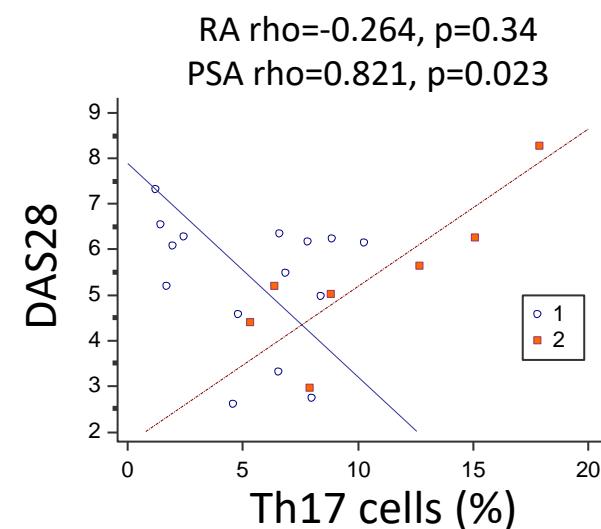
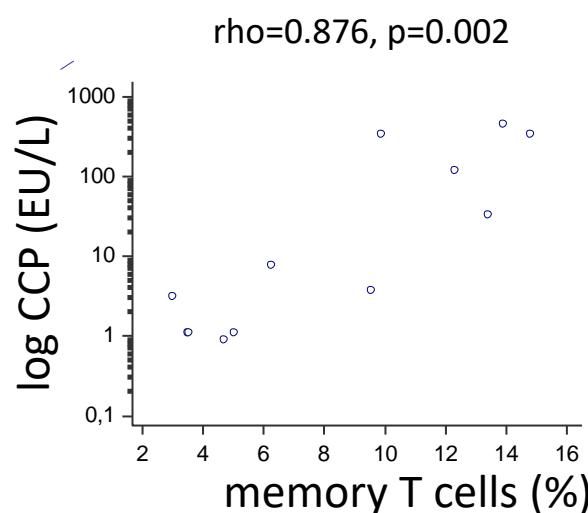
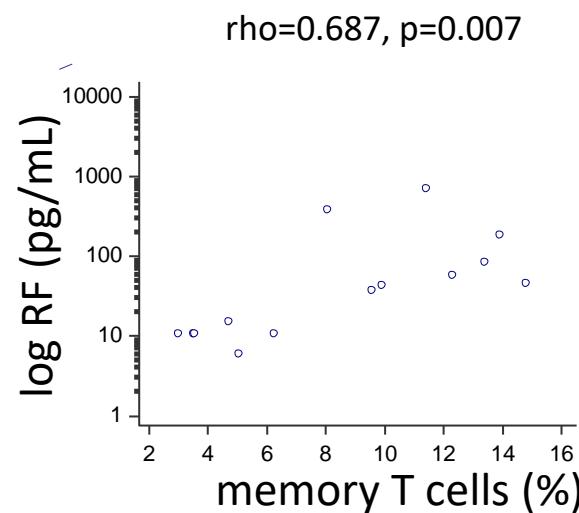
RA vs CTRL p=0.011  
PSA vs CTRL p=0.286



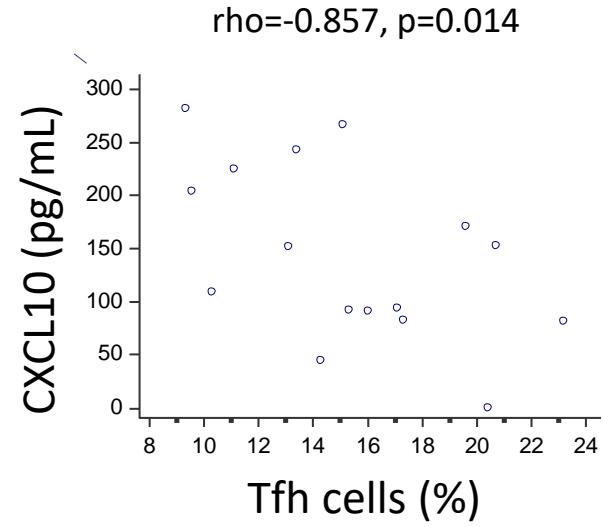
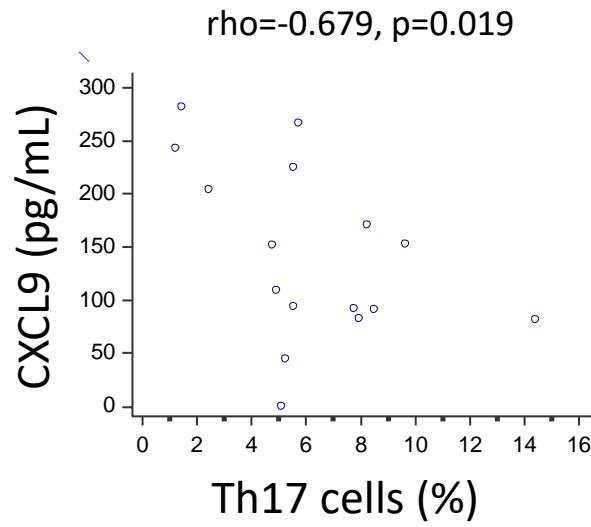
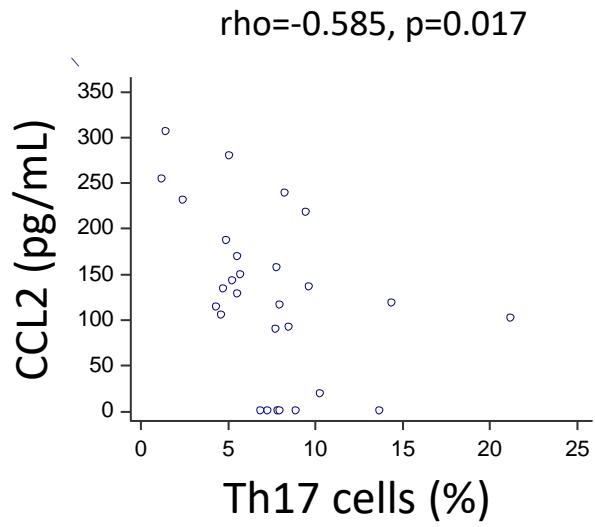
RA vs CTRL p=0.002  
PSA vs CTRL p=0.217



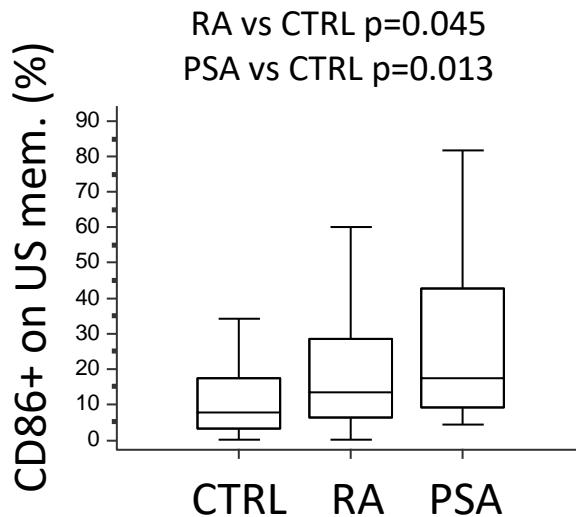
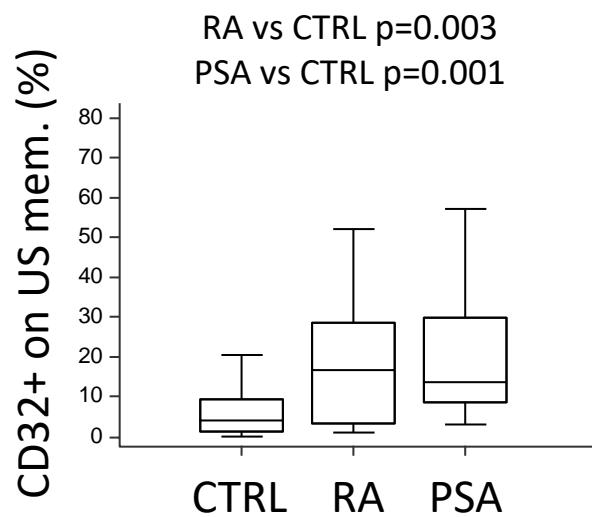
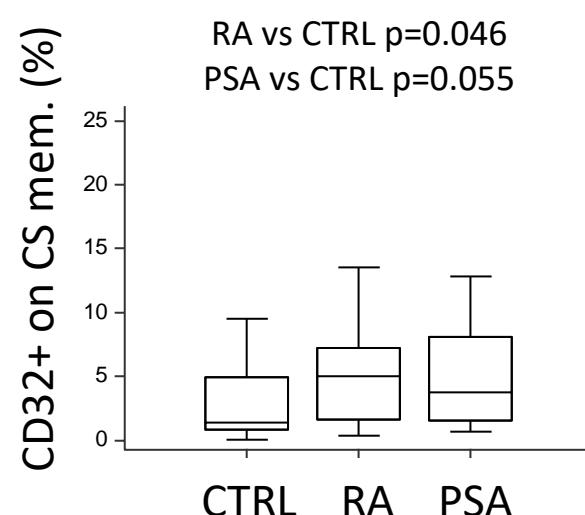
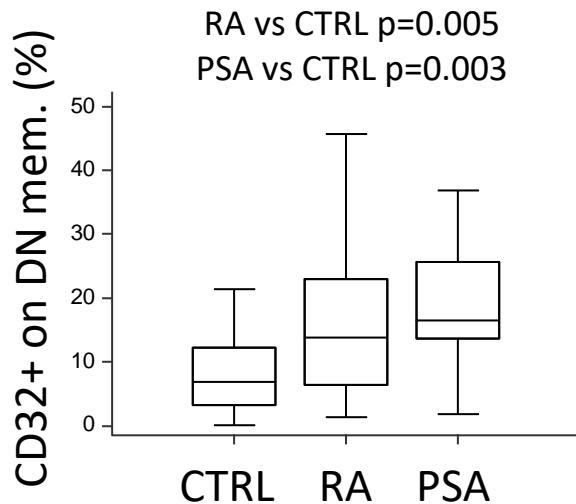
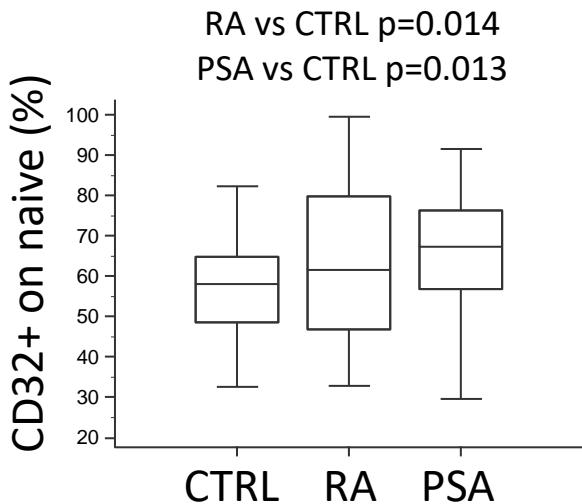
# Correlations of altered T lymphocyte populations with clinical parameters



# Correlations of altered T lymphocyte populations with serum cytokine levels

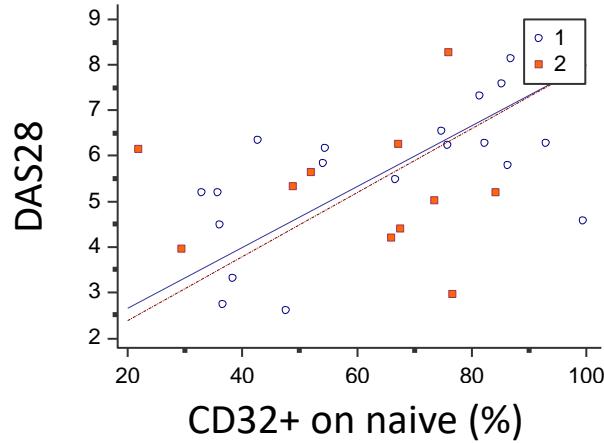


# CD32+ and CD86+ expression on B cell subpopulations

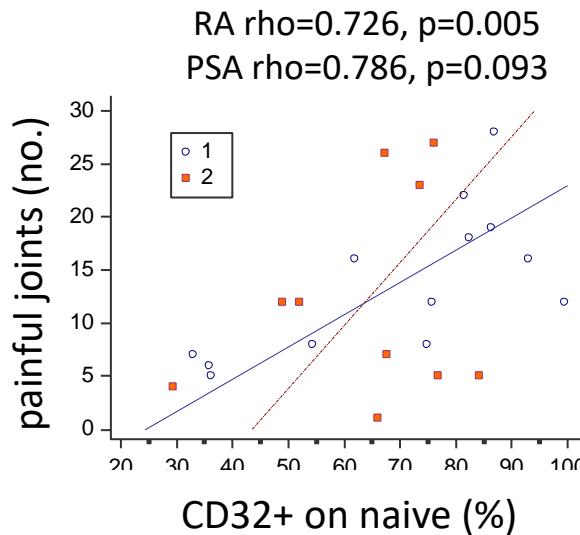


# Correlations of CD32+ and CD86+ expression on B lymphocytes populations with clinical parameters

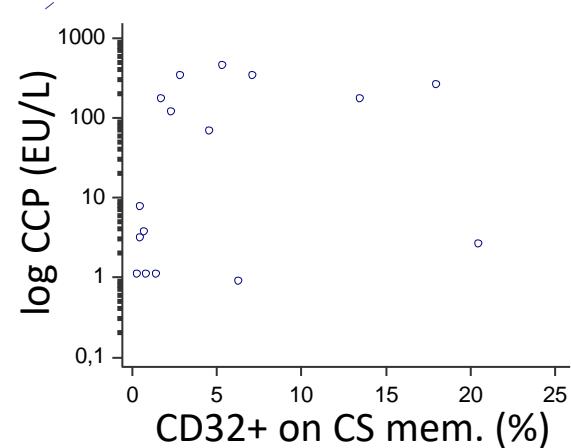
RA rho=0.558, p=0.013  
PSA rho=0.483, p=0.177



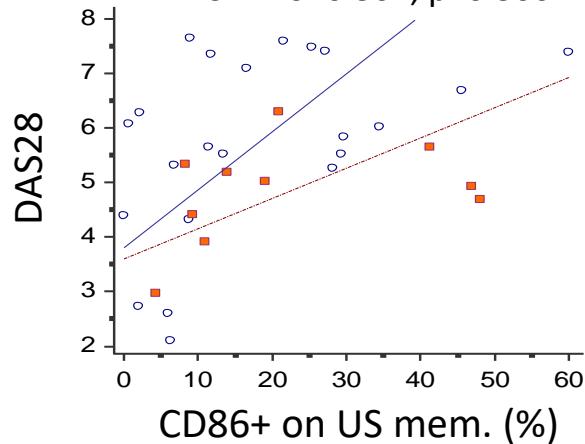
RA rho=0.726, p=0.005  
PSA rho=0.786, p=0.093



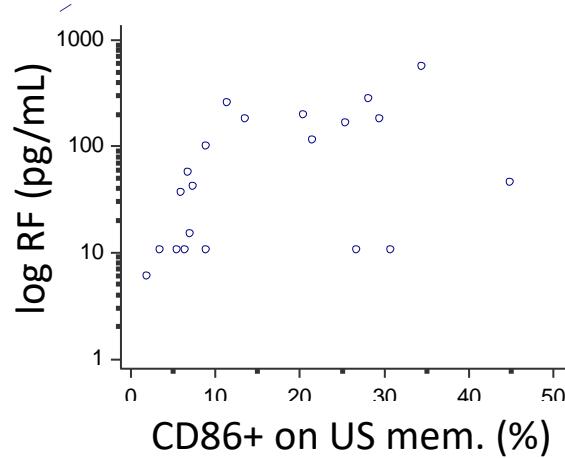
rho=0.502, p=0.040



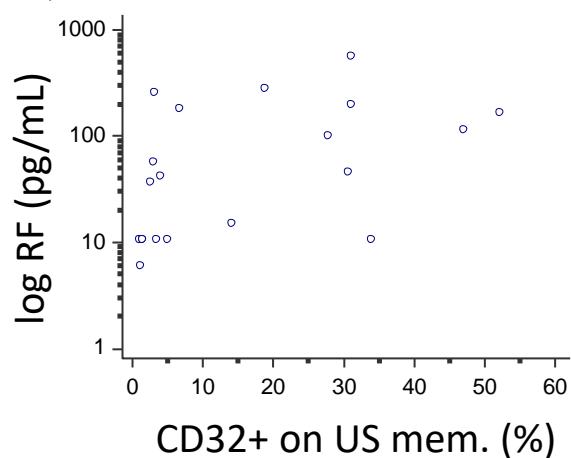
RA rho=0.471, p=0.027  
PSA rho=0.361, p=0.306



rho=0.550, p=0.010

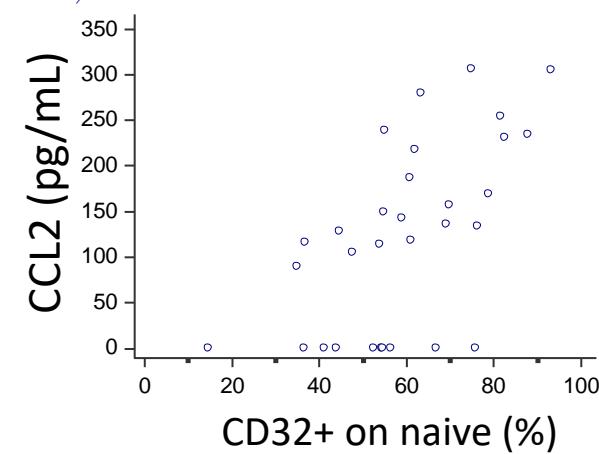


rho=0.578, p=0.006

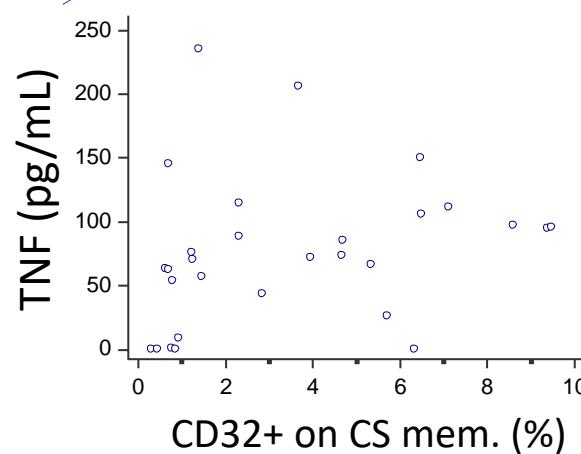


# Correlations of CD32+ and CD86+ expression on B lymphocytes populations with serum cytokine levels

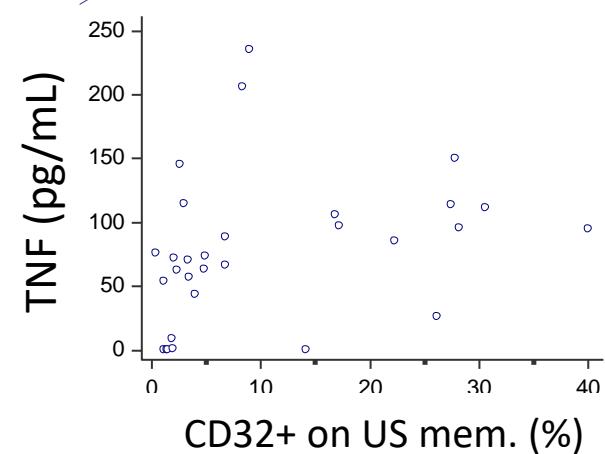
$\rho=0.504, p=0.005$



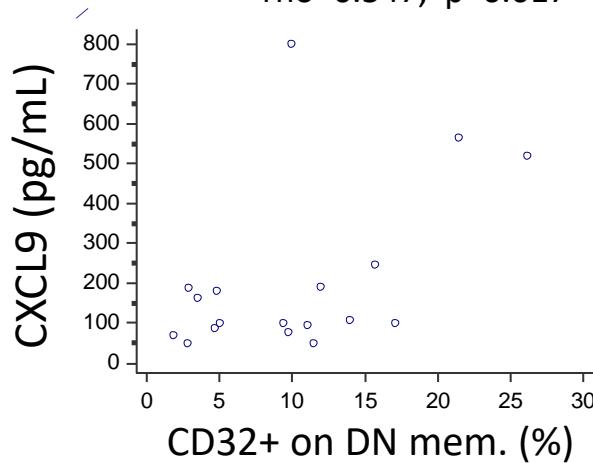
$\rho=0.549, p=0.002$



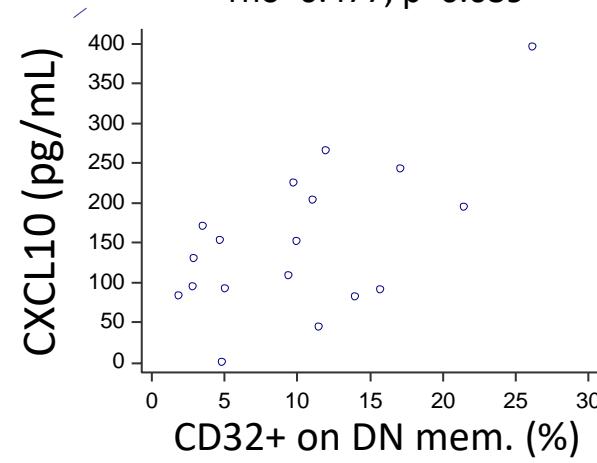
$\rho=0.472, p=0.008$



$\rho=0.547, p=0.017$

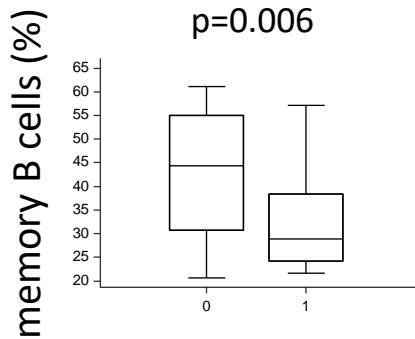


$\rho=0.477, p=0.039$

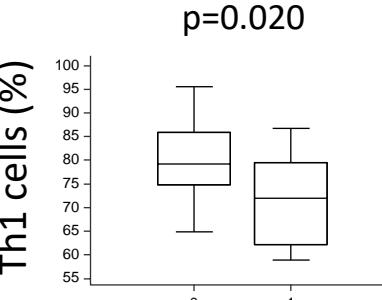


# Subpopulation differences in regards to therapy

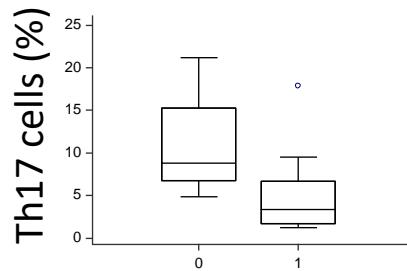
DMARD



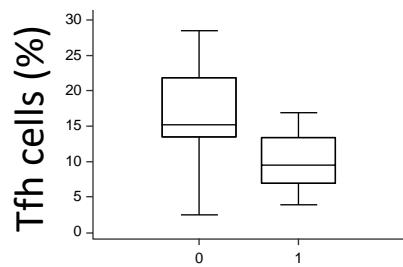
anti-TNF



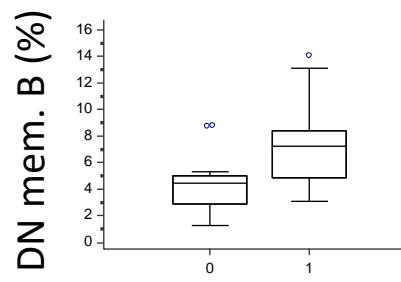
p=0.008



p=0.012



p=0.023



corticosteroids

# Conclusions

- memory T cells are expanded both in RA and PSA and correlate with RF and CCP levels
- Th17 and Tfh cells are found in lower frequency in RA and higher in PSA, while the Th17 frequency correlates with DAS28 positively in PSA
- CD32+ expression on naive and memory, and CD86+ on unswitched memory B cells is increased both in RA and PSA
- expression of CD32+ on naive and CD86+ on unswitched memory B cells correlates with DAS28 and RF in RA
- CD32+ expression on class switched and unswitched memory B cells correlates with RF, CCP and TNF levels
- these subpopulations may be of particular interest for further research, since their frequency is associated with disease activity and increased levels of proinflammatory and proresorptive cytokines



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Darja Flegar, MD

Antonio Markotić, MD

Sanja Ivčević

Katerina Zrinski Petrović

This work has been fully supported by Croatian Science Foundation under the project 5699.

## Acknowledgements