1. Supplementary Material

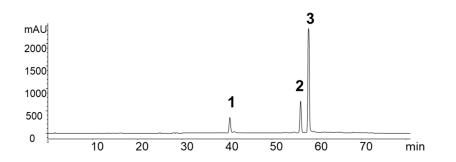


Figure S1. Reverse phase HPLC-UV analysis at 254 nm - optimized conditions for MPLC separation. Brachydin A (1), Brachydin B (2) and Brachydin C (3).

$$Kp = \frac{[S]_{upper phase}}{[S]_{lower phase}}$$

Figure S2. Coefficient of partition *K_p* equation (ARIZONA system).

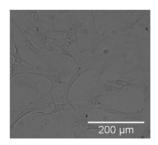


Figure S3. Human fibroblast like synoviocytes (optical microscope). Scale bar: 200 $\mu m.$

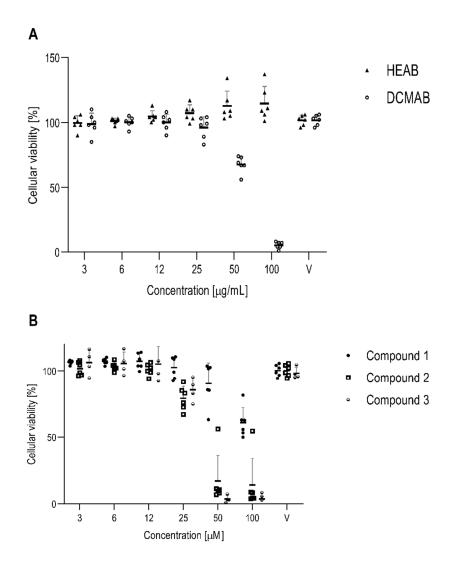


Figure S4. Scatter plot of Figure 3, showing cellular viability of human fibroblast like synoviocytes (HFLS) incubated with all tested compounds at increasing concentrations, after 24 h. Root extracts (**A**) and isolated compounds from dichloromethane extract (DCMAB) (**B**). Individual plotted values (n = 6) with mean values \pm S.D.. V = vehicle, 0.01 % dimethyl sulfoxide (DMSO).

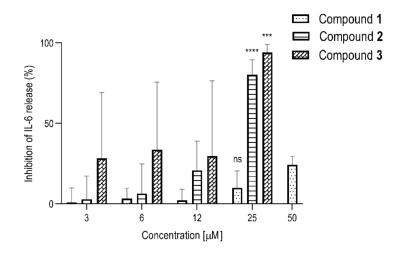


Figure S5. Inhibition percentage of IL-6 release normalized to 100 % activation, as a function of each compound concentration. Bars correspond to mean values \pm S.D.; n = 6.**** p <0.0001 and *** p = 0.0007 and ns = no significance.

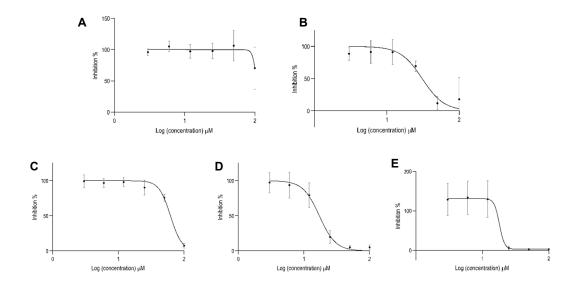


Figure S6. IC₅₀ calculations of hydroethanolic extract (HEAB) (**A**), dichloromethane extract (DCMAB) (**B**), **1** (**C**), **2** (**D**) and **3** (**E**). Points correspond to mean values ± S.D.; n = 6.

Table S1. Optimized MRM parameters for the quantification of active compounds by UHPLC-MS/MS.

Compound	Precursor Ion	SRM transition	DP (V)	EP (V)	CE (eV)	CXP (V)	Dwell time (ms)
1	525	525 → 271	30		10	7	
2	539	539 → 285	_	10			125
3	509	509 → 255	80	-	50	12	-

SRM, Selected reaction monitoring; DP, Declustering potential; EP, Entrance potential; CE, Collision energy; CXP, Collision cell exit.

Table S2. Calibration curves parameters for active compounds and determination of the LOD and LOQ in ng/mL.

Compound	Linear function	r ²	LOD (ng/mL)	LOQ (ng/mL)
1	$y = 816x + 6,06.10^3$		0.21	0.64
2	$y = 422x + 3,51.10^3$	0.998	0.47	1.41
3	y = 122x + 639		1.42	4.25

r², Correlation coefficient; LOD, Limit of Detection; LOQ, limit of quantification.