Supplementary Table S1. Baseline characteristics of flare (flare = loss of DAS28 remission) vs. sustained remission in the non-taper cohort (n=83).

Univariate analysis	Total n.	Flare (n=17)	Sustained Rem (n=66)	<i>p</i> -value	OR (95%CI)	AUC (95%CI) <i>p</i> -value	
Demographic variables Female, n (%)	83	14 (87.5%)	43 (65%)	0.076	3.912 (0.813 to 18.815)	0.386 (0.243 to 0.528) <i>p</i> =0.158	
Age (years)*	83	53.5 (19)	62 (17.8)	0.835	0.990 (0.945 to 1.038)	0.517 (0.355 to 0.679) <i>p</i> =0.836	
Disease duration (months)*	79	26.3 (23.2)	41 (44.6)	0.007	1.018 (0.998 to 1.038)	0.280 (0.125 to 0.434) <i>p</i> =0.007	
Remission duration (months)	83	8 (23.1)	12.5 (7.7)	0.352	1.010 (0.969 to 1.053)	0.425 (0.250 to 0.599) <i>p</i> =0.352	
RF+, n (%)	83	10 (62.5%)	40 (60.6%)	0.855	1.111 (0.360 to 3.430)	0.488 (0.329 to 0.646) <i>p</i> =0.877	
ACPA+, n (%)	83	10 (62.5%)	43 (65%)	0.874	0.913 (0.294 to 2.833)	0.511 (0.351 to 0.670) <i>p</i> =0.896	
Smoking, n (%) Never Ever	81	6 (35%) 11 (65%)	29 (44.6%) 35 (53%)	0.286	0.533 (0.166 to 1.713)	0.574 (0.419 to 0.728) <i>p</i> =0.363	
Clinical variables FJC28*	83	0 (0.75)	0 (1)	0.390	1.805 (0.619 to 5.263)	0.480 (0.312 to 0.649) p=0.826	
SJC28*	83	0 (0.75)	0 0)	0.055	0.210 (0.038 to 1.157)	0.551 (0.369 to 0.734) <i>p</i> =0.563	
CRP (mg/L)*	83	6.5 (<5 to 8.5)	0 (0)	0.003	0.941 (0.871 to 1.018)	0.634 (0.457 to 0.812) <i>p</i> =0.132	
ESR (mm/h)*	75	24.5 (28.8)	9.5 (8.7)	<0.0001	0.909 (0.857 to 0.963)	0.844 (0.751 to 0.937) <i>p</i> <0.0001	
EMS (mins)*	82	30 (60)	0 (6.3)	0.902	0.439 (0.159 to 1.212)	0.587 (0.333 to 0.841) <i>p</i> =0.477	
Patient-reported outcome variables VAS PGA*	82	22 (46.8)	8.5 (27)	0.809	1.004 (0.974 to 1.035)	0.499 (0.321 to 0.678) <i>p</i> =0.994	
VAS Pain*	69	17.5 (7)	9 (21.7)	0.910	1.015 (0.978 to 1.054)	0.500 (0.286 to 0.714) p=1	
VAS DA*	70	23.5 (42.5)	10.5 (32.2)	0.678	1.005 (0.891 to 1.030)	0.483 (0.263 to 0.703) <i>p</i> =0.887	
VAS Fatigue*	69	56.5 (56.8)	9.5 (43.3)	0.723	0.999 (0.978 to 1.021)	0.699 (0.532 to 0.866) <i>p</i> =0.102	
HAQ-DI*	65	0.56 (0.59)	0 (0.31)	0.322	0.526 (0.143 to 1.942)	0.864 (0.752 to 0.975) <i>p</i> =0.003	
RaQoL*	66	7 (8.7)	1 (2)	0.774	0.970 (0.878 to 1.072)	0.894 (0.795 to 0.993) <i>p</i> =0.001	
Ultrasound variables Fotal PD*	56	0 (3)	0 (0)	0.526	0.944 (0.598 to 1.490)	0.552 (0.307 to 0.797) <i>p</i> =0.669	
Total GS*	56	18.5 (10.2)	14 (11.5)	0.546	0.968 (0.875 to 1.072)	0.645 (0.437 to 0.853) <i>p</i> =0.233	
Immunological (T-cell variables) Normalised naïve*	56	19.11 (23.01)	12.22 (19.11)	0.405	1.021 (0.980 to 1.064)	0.498 (0.290 to 0.705) <i>p</i> =0.983	
Normalised Treg*	59	-3.16 (2.41)	-3.28 (6.4)	0.728	1.025 (0.753 to 1.394)	0.473 (0.306 to 0.640) <i>p</i> =0.784	
IRC*	59	2.00 (2)	2.40 (3)	0.802	1.060 (0.860 to 1.305)	0.468 (0.298 to 0.637) p=0.784	
Drug category Monotherapy, n (%) Combination therapy n (%)	48 35	7/17 (41%) 10/17 (59%)	39/66 (59%) 27/66 (41%)	0.185	0.539 (0.220 to 1.322)	$0.603  (0.449-0.758) \\ p=0.202$	

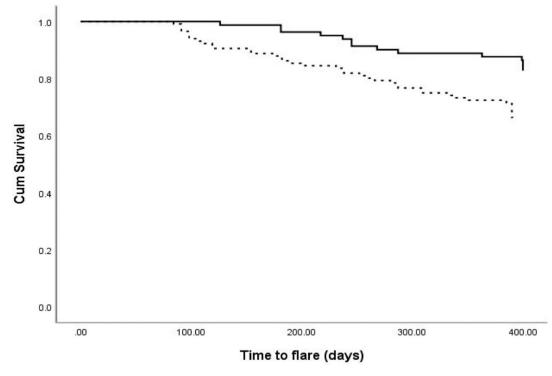
\*Median (IQR); CRP <5 mg/l = lowest detectable limit. RF: rheumatoid factor; CCP: anti-CCP antibody; TJC/SJC(28): number of tender and swollen joints out of 28; CRP: C-reactive protein, mg/l; ESR: erythrocyte sedi-mentation rate, mm/h; EMS: early morning stiffness, mins; VAS: visual analogue score; PGA: patient global assessment of disease; DA: disease activity; HAQ-DI: Health Assessment Questionnaire Disability Index; RAQoL: RA Quality of Life questionnaire score; PD: Power Doppler score; GS: Grey Scale synovial hypertrophy score; T-cell subsets (% of CD4<sup>+</sup>T-cells): naïve T-cells; IRC: inflammation related cells; Treg: T-regulatory cells.

## Tapering of cs-DMARD therapy in RA patients / H.L. Gul et al.

**Supplementary Table S2.** Modelling the prediction of sustained remission after 12 m in tapering cohort using Cox regression (n=117), missing data were imputed: HR (95% CI) and *p*-value.

	Univariate	Unadjusted	Model 1 Clin only	Model 2 Clin + US	Model 3 Clin + PRO	Model 4 Clin + T-cells	Model 5 Clin + PRO + US	Model 6 Clin + PRO + T-cells	Model 7 Clin + US + T-cells	Model 8 Clin + PRO + US + T-cells
Clinical variabl	es									
TJC28	0 (0-0) [0-5] p=0.001	1.465 (1.126-1.905) <i>p</i> =0.004	1.534 (1.178-1.997) <i>p</i> =0.001	1.561 (1.183-2.056) <i>p</i> =0.002	1.475 (1.116-1.949) <i>p</i> =0.006	1.439 (1.154-1.924) <i>p</i> =0.005	1.505 (1.117-2.028) <i>p</i> =0.007	Not selected by model	1.452 (1.106-1.907) <i>p</i> =0.007	1.511 (1.140-2.002) <i>p</i> =0.004
CRP	0 (0-0) [0-34] <i>p</i> =0.011	0.014 (1.011-1.103) <i>p</i> =0.04	1.065 (1.019-1.113) <i>p</i> =0.005	1.064 (1.016-1.114) P=0.009	Not selected by model	Not selected by model	Not selected by model 1	Not selected by model	Not selected by model	Not selected by model
EMS	0 (0-5) [0-180] <i>p</i> =0.011	1.009 (1.002-1.017) <i>p</i> =0.019	Not selected by model	Not selected by model	Not selected by model	Not selected by model	Not selected by model	Not selected by model	Not selected by model	Not selected by model
PROs										
VAS Pain	5 (2-15) [0-80] <i>p</i> =0.004	1.044 (1.021-1.068) <i>p</i> <0.0001	NA	NA	Not selected by model	NA	Not selected by model	Not selected by model	NA	Not selected by model
VAS PGA	10 (5-20) [0-77] <i>p</i> =0.058	1.030 (1.010-1.050) <i>p</i> <0.003	NA	NA	Not selected by model	NA	Not selected by model	Not selected by model	NA	Not selected by model
VAS DA	5 (2-14) [0-98] <i>p</i> =0.002	1.018 (1.005-1.031) <i>p</i> =0.007	NA	NA	Not selected by model	NA	Not selected by model	Not selected by model	NA	Not selected by model
VAS Fatigue	10 (4-24.5) [0-96] <i>p</i> =0.006	1.019 (1.007-1.030) <i>p</i> =0.001	NA	NA	Not selected by model	NA	Not selected by model	Not selected by model	NA	Not selected by model
HAQ-DI	0 (0-0.25) [0-2.88] <i>p</i> <0.0001	2.729 (1.755-4.241) <i>p</i> <0.0001	NA	NA	2.455 (1.171-3.260) <i>p</i> <0.001	NA	2.705 (1.718-4.259) <i>p</i> <0.001	Not selected by model	NA	2.063 (1.161-3.665) <i>p</i> =0.014
RaQoL	0 (0-3) [0-23] <i>p</i> =0.0001	1.120 (1.063-1.181) <i>p</i> <0.0001	NA	NA	Not selected by model	NA	Not selected by model	1.097 (1.032-1.166) <i>p</i> =0.003	NA	Not selected by model
Ultrasound vari	ables									
Total PD	0 (0-1.00) [0-23] <i>p</i> =0.010	1.138 (1.043-1.242) <i>p</i> =0.004	NA	1.127 (1.032-1.231) <i>p</i> =0.008	NA	NA	1.127 (1.031-1.232) <i>p</i> =0.008	NA	1.163 (1.062-1.274) <i>p</i> =0.001	1.158 (1.057-1.267) <i>p</i> =0.002
Total GS	15 (11.25-250 [0-45] <i>p</i> =0.046	1.029 (0.998-1.061) <i>p</i> =0.066	NA	Not selected by model	NA	NA	Not selected by model	NA	Not selected by model	Not selected by model
TT-cell Variable	s									
IRC	1.25 (0.50-1.75) [0.20-17.00] <i>p</i> <0.0001	1.28 (1.190-1.380) <i>p</i> <0.0001	NA	NA	NA	1.291 (1.193-1.399) <i>p</i> <0.0001	NA	1.233 (1.142-1.331) <i>p</i> <0.0001	1.311 (1.211-1.419) <i>p&lt;</i> 0.001	1.244 (1.142-1.355) <i>p</i> <0.0001
AUC (95% CI)			0.474 (0.466-0.482)	0.584 0.580-0.595	0.559 (0.556-0.571)	0.675 (0.669-0.685)	0.651 (0.646-0.658)	0.692 (0.679-0.697)	0.729 (0.724-0.738)	0.761 (0.755-0.769)

RF: rheumatoid factor; CCP: anti-CCP antibody; TJC/SJC(28): number of tender and swollen joints out of 28; CRP: C-reactive protein, mg/l; ESR: erythrocyte sedimentation rate, mm/h; EMS: early morning stiffness, mins; VAS: visual analogue score; PGA: patient global assessment of disease; DA: disease activity; HAQ-DI: Health Assessment Questionnaire Disability Index; RAQoL: RA Quality of Life questionnaire score; PD: Power Doppler score; GS: Grey Scale synovial hypertrophy score; T-cell subsets (% of CD4<sup>+</sup>T-cells): naïve T-cells; IRC: inflammation related cells; Treg: T-regulatory cells.



Supplementary Fig. S1. Kaplan Meier Plot showing time to flare in the tapering (dashed line) and stable therapy (bold line) cohorts.