ONLINE SUPPLEMENTARY MATERIAL

Supplementary Table S1. Baseline and time-varying predictors of failure to apply **T2T-REM** (<u>according to CDAI and SDAI</u>) during 2-year follow-up in the RA BIODAM Cohort.

	CDAI		SDAI	
	Baseline model	Longitudinal model	Baseline model	Longitudinal model
	OR (95% CI)	OR (95% CI)	OR (95% CI)	OR (95% CI)
	(N=551)	(N=540)	(N=549)	(N=540)
Age (years)	¥	¥	¥	1.01 (1.00; 1.01)
Gender (female)	1.26 (1.05; 1.52)	1.30 (1.03; 1.64)	1.21 (1.01; 1.46)	1.28 (1.02; 1.60)
Disease duration (years)	¥	¥	¥	¥
Education (years)	0.97 (0.95; 0.99)	0.95 (0.93; 0.98)	0.96 (0.94; 0.98)	0.95 (0.93; 0.98)
Number of comorbidities	¥	¥	¥	¥
Current smoker	†	†	†	†
Type of centre (academic)	1.21 (1.02; 1.43)	¥	†	†
RF positivity	+	¥	†	†
ACPA positivity	¥	0.79 (0.64; 0.98)	¥	0.80 (0.65; 0.98)
RF and/or ACPA positivity	¥	†	¥	¥
PGA (0-10)	¥	¥	¥	¥
PhGA (0-10)	†	¥	¥	¥
Swollen joint count (0-44)	†	0.87 (0.85; 0.89)	†	0.88 (0.86; 0.90)
Tender joint count (0-53)	¥	¥	¥	¥
ESR (mm/h)	†	¥	†	¥
CRP (mg/L)	¥	¥	†	¥
HAQ	1.26 (1.12; 1.41)	¥	1.28 (1.14; 1.44)	¥
Number of previous csDMARDs	¥	1.13 (1.02; 1.24)	¥	¥
Previous treatment with any csDMARD	¥	¥	¥	¥

^{†:} not selected from the univariable analysis (p>0.20); ¥: not significant in the multivariable analysis (p>0.05). Country added as a covariate in all univariable models and in the final multivariable model. Final models also adjusted for type of treatment (csDMARDs and/or bDMARDs)

Supplementary Table S2. Baseline and time-varying predictors of failure to apply **T2T-LDA** (<u>according to CDAI and SDAI</u>) during 2-year follow-up in the RA BIODAM Cohort.

	CDAI		SDAI	
	Baseline model	Longitudinal model	Baseline model	Longitudinal model
	OR (95% CI)	OR (95% CI)	OR (95% CI)	OR (95% CI)
	(N=538)	(N=555)	(N=538)	(N=555)
Age (years)	¥	1.01 (1.00; 1.02)	¥	¥
Gender (female)	+	†	†	†
Disease duration (years)	¥	¥	¥	†
Education (years)	0.97 (0.95; 0.99)	¥	0.97 (0.95; 1.00)	¥
Number of comorbidities	¥	¥	¥	¥
Current smoker	†	†	†	†
Type of centre (academic)	¥	¥	¥	¥
RF positivity	¥	¥	¥	¥
ACPA positivity	0.62 (0.51; 0.75)	0.62 (0.52; 0.46)	0.63 (0.51; 0.77)	0.64 (0.53; 0.78)
RF and/or ACPA positivity	¥	¥	¥	¥
PGA (0-10)	¥	¥	¥	¥
PhGA (0-10)	¥	¥	¥	¥
Swollen joint count (0-44)	†	0.97 (0.95; 0.99)	+	0.97 (0.95; 0.99)
Tender joint count (0-53)	1.02 (1.01; 1.03)	1.02 (1.01; 1.04)	1.02 (1.01; 1.03)	1.02 (1.01; 1.03)
ESR (mm/h)	¥	†	+	†
CRP (mg/L)	¥	¥	+	+
HAQ	1.18 (1.02; 1.36)	¥	1.20 (1.04; 1.39)	1.40 (1.22; 1.61)
Number of previous csDMARDs	¥	¥	¥	¥
Previous treatment with any csDMARD	¥	¥	¥	¥

^{†:} not selected from the univariable analysis (p>0.20); ¥: not significant in the multivariable analysis (p>0.05). Country added as a covariate in all univariable models and in the final multivariable model. Final models also adjusted for type of treatment (csDMARDs and/or bDMARDs)