

A CYTOKINE SIGNATURE FROM MONOCYTE-DERIVED MACROPHAGES PREDICTS THE RESPONSE TO APREMILAST IN PATIENTS WITH PSORIATIC ARTHRITIS

BACKGROUND

- **Apremilast** is a **PDE4-inhibitor** approved for the treatment of psoriatic arthritis (PsA).
- The mechanisms of action of apremilast remain elusive, although data support the regulation of **innate immune cells**.
- The role of **macrophages** in the pathogenesis of PsA remains largely enigmatic

OBJECTIVE

- To investigate the **ex vivo** effects of **apremilast** on **monocyte-derived macrophages** in patients with PsA.

METHODS

Patients

- **23 PsA patients starting apremilast**
 - mean age 55±14.3; 10/23 women
 - disease duration 82 ± 95 months; 6 on MTX
- **21 knee osteoarthritis (OA) controls**
 - mean age 63 ± 8.7; 10/21 women

Peripheral blood (PB) monocytes

- isolated at **baseline** in subjects and after **4 months of apremilast** therapy in PsA
- differentiated into **M1- or M2-macrophages**

Gene expression (RT-qPCR) and supernatant levels (ELISA):

- **IL-1β, IL-23, and TNF-α** → **M1-macrophages**
- **IL-10, IL-1Ra, and TGF-β** → **M2-macrophages**.

Patients with PsA were coined as **responders** if at 4 months a decrease ≥4 points of DAPSA and any reduction in TJC or SJC occurred compared to baseline.

RESULTS

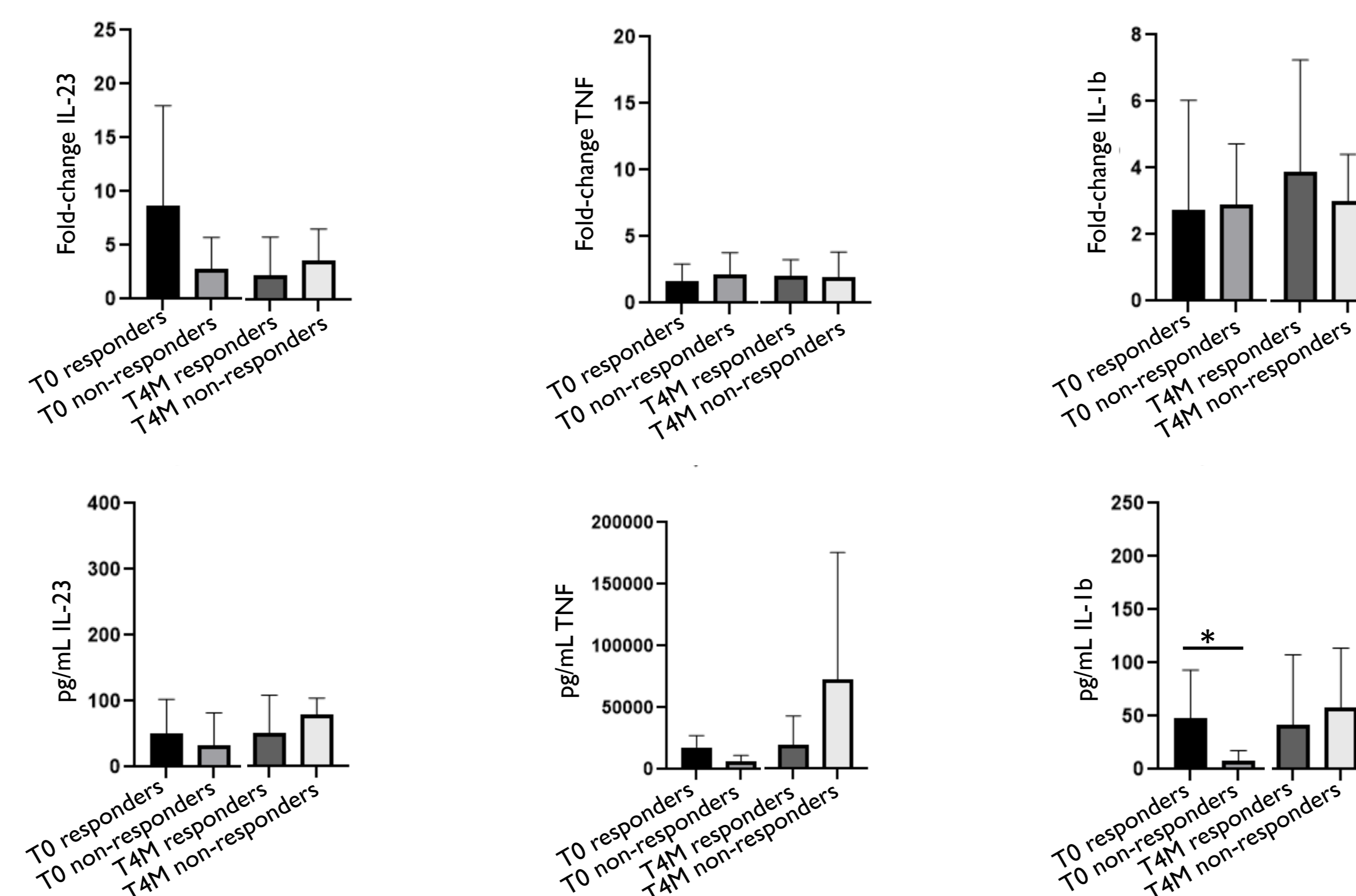
After receiving apremilast for 4 months:

- 17/23 patients were responders; 10/17 had DAPSA down-stage
- Response was independent of concomitant MTX, systemic glucocorticoids or baseline DAPSA

M1-macrophages

	OA	PsA baseline	PsA 4 months
IL-23 expression (fold change)	1.09 (1.63)	6.43 (7.88)*	2.50 (3.46)
		Responders	8.68 (9.31)
		Non-responders	2.26 (3.60)**
IL-23 ELISA (pg/mL)	89.0 (101.1)	43.9 (49.6)	54.4 (54.5)
		Responders	50.4 (52.3)
		Non-responders	50.8 (57.3)
TNF-α expression (fold change)	1.36 (1.30)	1.86 (1.38)	2.03 (1.28)
		Responders	1.68 (1.27)
		Non-responders	2.04 (1.21)
TNF-α ELISA (pg/mL)	17 262 (23 479)	13 252 (9 483)	18 285 (22 854)
		Responders	17 844 (9 629)
		Non-responders	19 007 (23 667)
IL-1β expression (fold change)	2.33 (2.46)	2.80 (2.72)	3.74 (3.10)
		Responders	2.75 (3.28)
		Non-responders	3.89 (3.38)
IL-1β ELISA (pg/mL)	20.6 (33.8)	31.1 (39.8)	44.7 (62.9)
		Responders	47.7 (45.5)
		Non-responders	41.9 (65.6)
TGF-β expression (fold change)	1.34 (1.30)	1.01 (0.66)	0.72 (0.28)
		Responders	1.03 (0.75)
		Non-responders	0.98 (0.56)
TGF-β ELISA (pg/mL)	578.1 (70.9)	592.2 (199.3)	656.7 (74.5)**
		Responders	679.8 (41.1)
		Non-responders	649.3 (76.9)
IL-1Ra expression (fold change)	2.62 (7.00)	1.52 (1.11)	1.46 (0.87)
		Responders	1.27 (0.49)
		Non-responders	1.43 (0.91)
IL-1Ra ELISA (ng/mL)	32.2 (31.6)	25.0 (16.7)	25.5 (19.6)
		Responders	24.8 (18.7)
		Non-responders	25.2 (20.3)

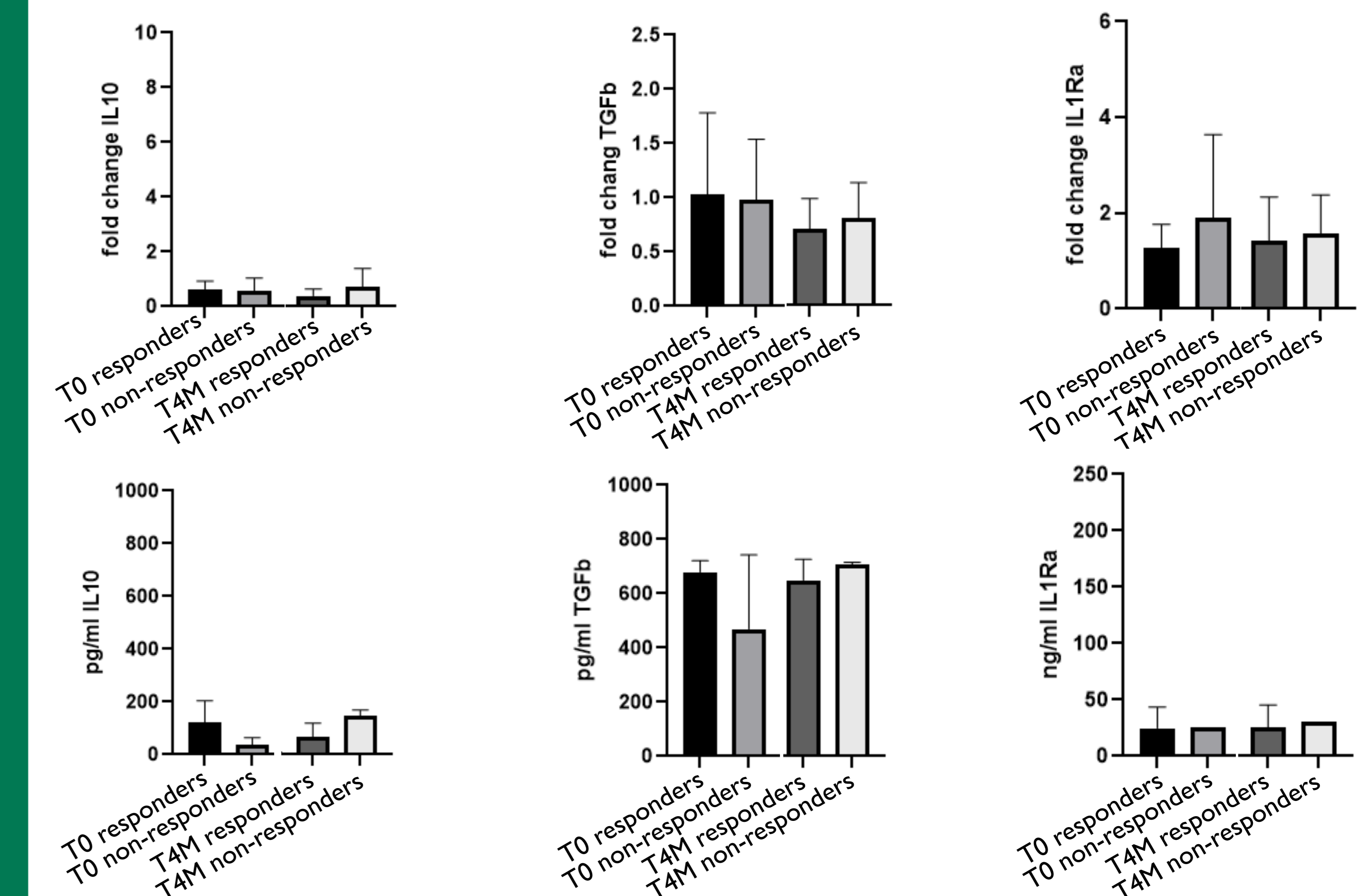
* $p < 0.05$ PsA baseline vs. OA; ** $p < 0.05$ PsA at 4 months vs. PsA baseline; # $p < 0.05$ non-responders vs. responders at baseline



M2-macrophages

	OA	PsA baseline	PsA 4 months
IL-10 expression (fold change)	4.03 (9.24)	0.62 (0.35)	0.46 (0.34)
		Responders	0.64 (0.29)
		Non-responders	0.40 (0.24)
IL-10 ELISA (pg/mL)	91.1 (82.6)	84.4 (79.2)	77.4 (56.8)
		Responders	120.7 (83.0)
		Non-responders	66.5 (52.6)
TGF-β expression (fold change)	1.34 (1.30)	1.01 (0.66)	0.72 (0.28)
		Responders	1.03 (0.75)
		Non-responders	0.98 (0.56)
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		Responders	24.8 (18.7)
		Non-responders	25.2 (20.3)

** $p < 0.05$ PsA at 4 months vs. PsA baseline



CONCLUSIONS

- An **IL-23 signature** characterizes PsA PB-derived M1 macrophages;
- Baseline **IL-1β, TNF-α, IL-23** predict **response** to apremilast
- IL-23 decreases in responders; **lack of response** is linked to an **increased myeloid inflammatory signature**