

Precision Medicine και κατευθυντήριες οδηγίες: κινούνται προς την ίδια κατεύθ



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Σύγκρουση Συμφερόντων Καμία





Κλινική Ρευματολογίας και Κλινικής Ανοσολογίας ΠΓΝΛάρισα



Εργαστήριο Ρευματολογίας & Κλινικής Ανοσολογίας



Ιατρική Ακριβείας και κατευθυντήριες οδηγίες: κινούνται προς την ίδια κατεύθυνση;

Όχι



Terminology

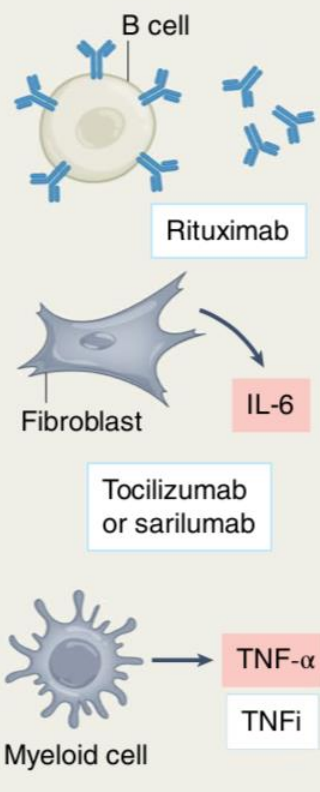
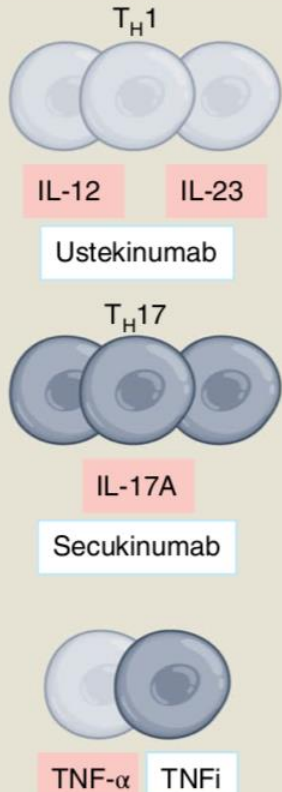
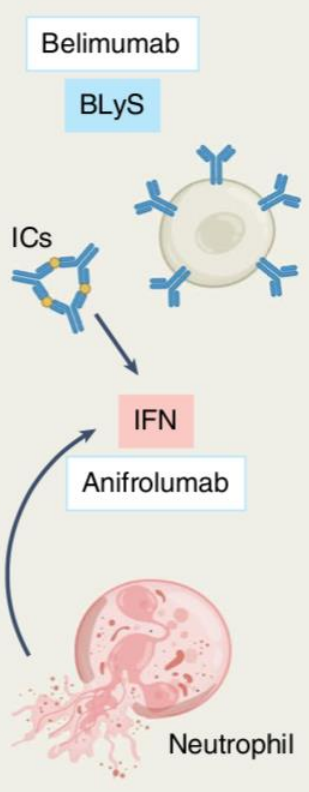
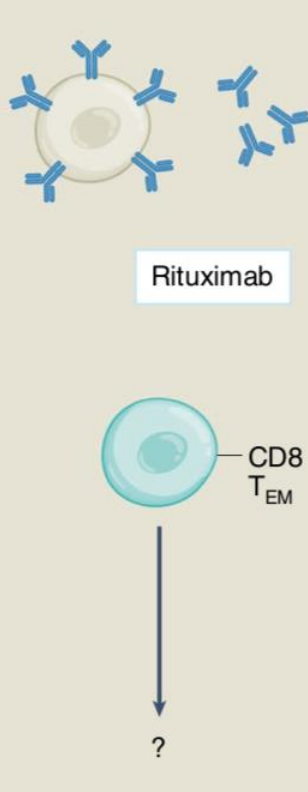

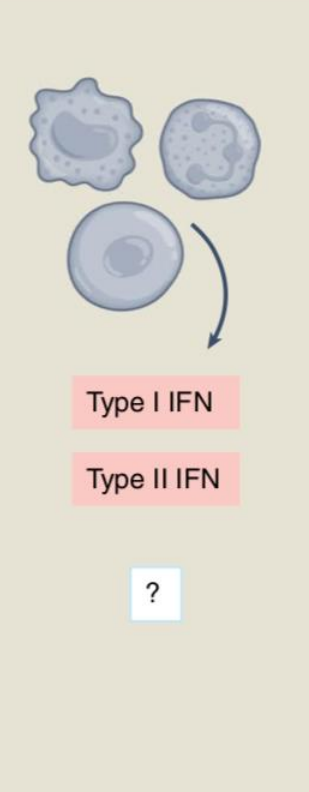
Clinical practice guidelines as "statements that include recommendations, intended to **optimize patient care**, that are informed by a systematic review of evidence and an assessment of the benefits and harms of alternative care options

Precision medicine, sometimes known as "personalized medicine" is an innovative approach to **tailoring disease prevention** and treatment that takes into account differences in people's genes, environments, and lifestyles.

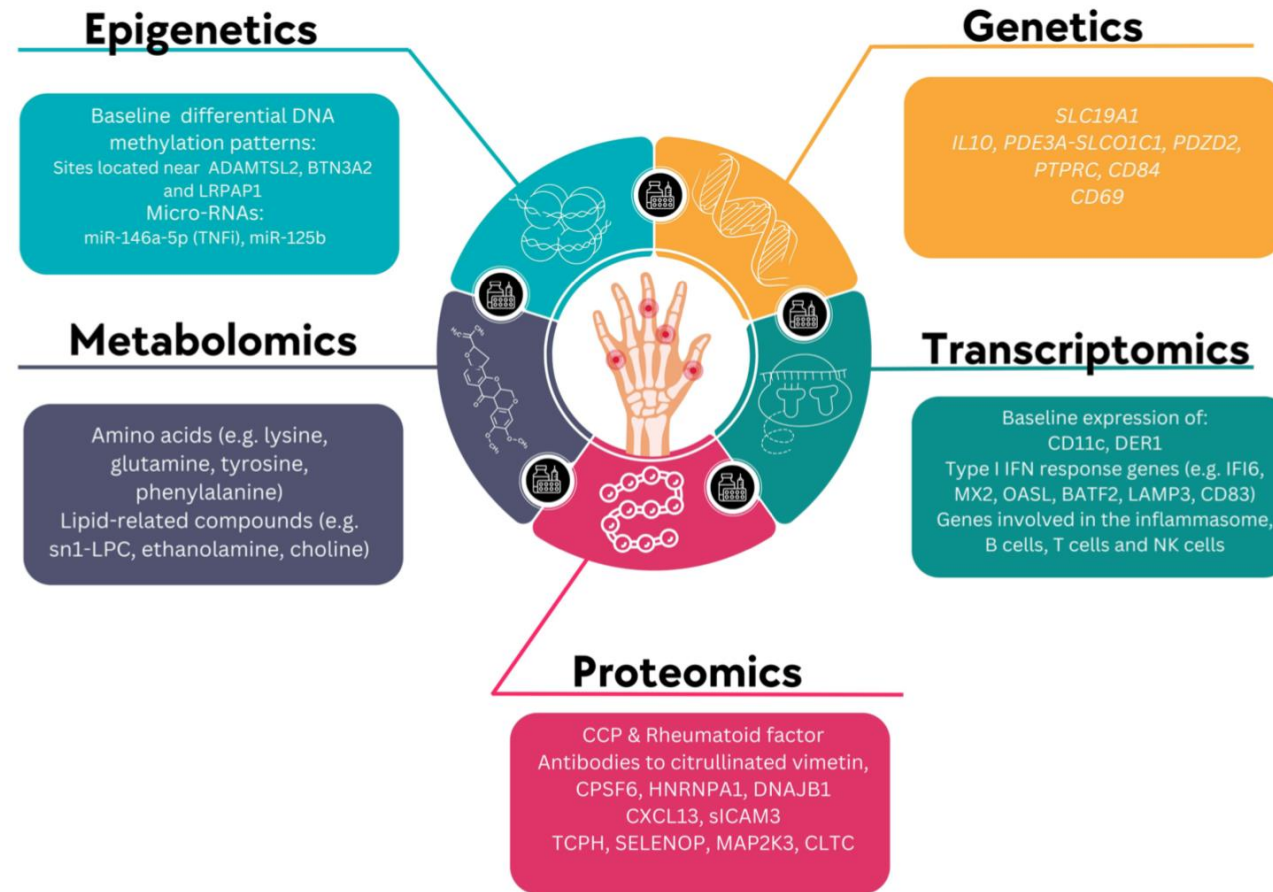
Ιατρική Ακριβείας και κατευθυντήριες οδηγίες: Βασίζονται στα ίδια (θεραπευτικά) εργαλεία;

Ναι



	a RA	b PsA	c SLE	d AAV	e SSc	f pSS
Organs	Joints Lungs	Joints Skin Lungs	Joints Skin Kidneys	Lungs Kidneys	Lungs Kidneys Skin	Joints Skin Eyes Salivary glands
Autoabs	Cyclic citrullinated peptide (CCP) Rheumatoid factor (RF)	—————	dsDNA Smith (Sm) Cardiolipin	anti-neutrophil cytoplasmic (proteinase 3 and myeloperoxidase)	Diffuse SSc: Topoisomerase 1 RNA polymerase I and III Limited SSc: Centromere B	Ro/SSA La/SSB
HLA	HLA-DRB4 HLA-DRB1	HLA-B27	HLA-DRB2 HLA-DRB3	HLA-DRB1	HLA-DR52a	HLA-DR3
Potential tailored treatments	 <p>B cell Rituximab</p> <p>Fibroblast IL-6 Tocilizumab or sarilumab</p> <p>Myeloid cell TNF-α TNFi</p>	 <p>T_H1 IL-12 IL-23 Ustekinumab</p> <p>T_H17 IL-17A Secukinumab</p> <p>TNF-α TNFi</p>	 <p>Belimumab BlyS</p> <p>ICs</p> <p>IFN Anifrolumab</p> <p>Neutrophil</p>	 <p>Rituximab</p> <p>CD8 T_{EM}</p> <p>?</p>	 <p>Fibrosis</p> <p>VEGFR, FGFR, PDGFR Nintedanib</p> <p>Inflammatory cells IL-6 Tocilizumab</p>	 <p>Type I IFN</p> <p>Type II IFN</p> <p>?</p>

Precision Medicine



Μας ενδιαφέρουν τα «δακτυλικά αποτυπώματα» του ασθενούς



Box 1 | Current and emerging approaches for molecular characterization of patients with systemic autoimmune rheumatic disease

Current

- Autoantibodies
- Clinical imaging
- Clinical lab testing: complement levels and split products
- Soluble mediators: cytokines, chemokines, and soluble receptors
- Transcriptomics: molecular signatures
- Genetics: disease-associated variants
- Immunophenotyping: flow cytometry
- Tissue histology

Emerging

- Genetics: genetic load, polygenic risk scores, extended HLA haplotypes
- Transcriptomics: cell-specific expression/signatures (scRNA-seq)
- Immunophenotyping: single-cell proteomics (CyTOF), proteogenomics (CITE-seq), repertoire immunomics
- Perturbomics (multi-omic evaluation after stimulation or other perturbation conditions)
- Spatial tissue analytics: multiplex tissue imaging (CODEX, serial IHC)
- Imaging mass cytometry (Hyperion, IonPath)
- Epigenomics (sorted cell and single cell): DNA methylation, histone modification, chromatin conformation (ATAC-seq), protein-DNA interactions (CUT&RUN)
- Mass spectroscopy (biofluid) and imaging mass spectrometry (tissue): proteomics, metabolomics, lipidomics, and glycomics
- Environmental factors: microbiomics, exposomics





Precision medicine vs guidelines

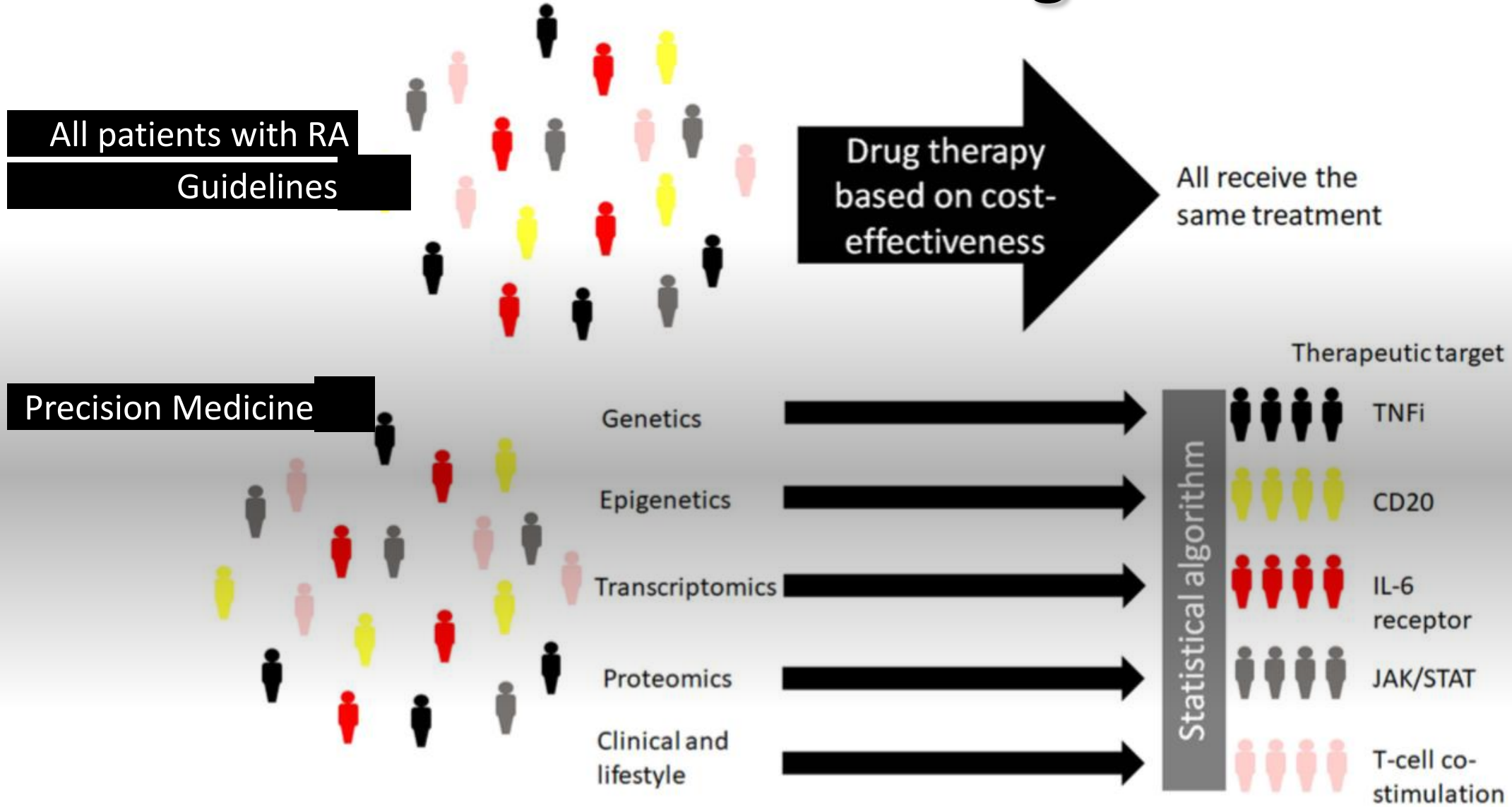


Figure 1 Illustration of how personalized medicine approaches using biomarkers and clinical predictors of treatment outcome can be applied to select a therapeutic target with an increased likelihood of response for the individual patient.

Box 1 | Current and emerging approaches for molecular characterization of patients with systemic autoimmune rheumatic disease

Current

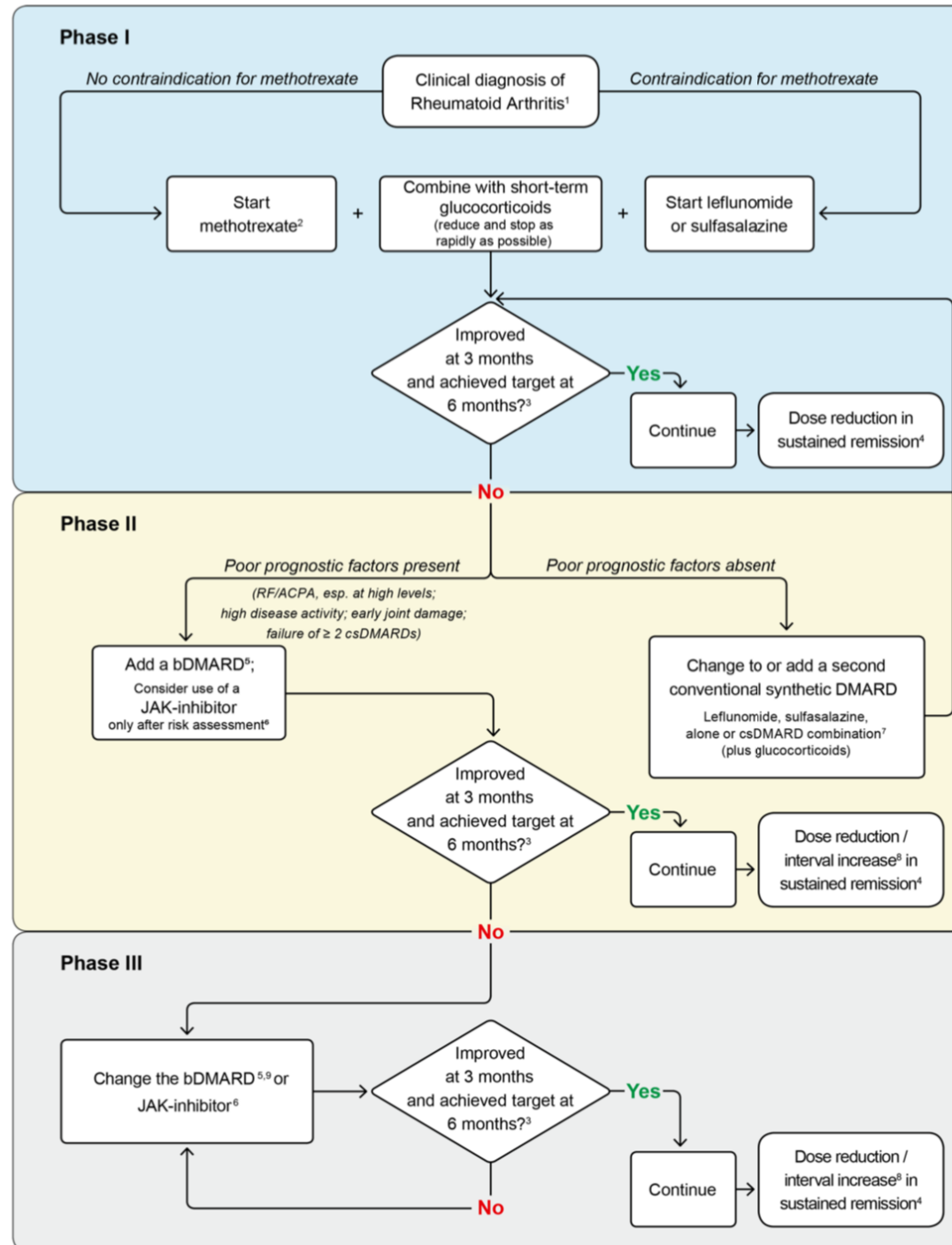
- Autoantibodies
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EULAR recommendations for the management of rheumatoid arthritis with synthetic and biological disease-modifying antirheumatic drugs: 2022 update



Ιατρική Ακριβείας με βάση τον ιστό (ανοσοφαινότυπο) και όχι τα *guidelines*

Rituximab versus tocilizumab in anti-TNF inadequate responder patients with rheumatoid arthritis (R4RA): 16-week outcomes of a stratified, biopsy-driven, multicentre, open-label, phase 4 randomised controlled trial



Frances Humby, Patrick Durez, Maya H Buch, Myles J Lewis, Hasan Rizvi, Felice Rivellese, Alessandra Nerviani, Giovanni Giorli, Arti Mahto, Carlomaurizio Montecucco, Bernard Lauwerys, Nora Ng, Pauline Ho, Michele Bombardieri, Vasco C Romão, Patrick Verschueren, Stephen Kelly, Pier Paolo Sainaghi, Nagui Gendi, Bhaskar Dasgupta, Alberto Cauli, Piero Reynolds, Juan D Cañete, Robert Moots, Peter C Taylor, Christopher J Edwards, John Isaacs, Peter Sasieni, Ernest Choy, Costantino Pitzalis, on behalf of the R4RA collaborative group



Lancet 2021



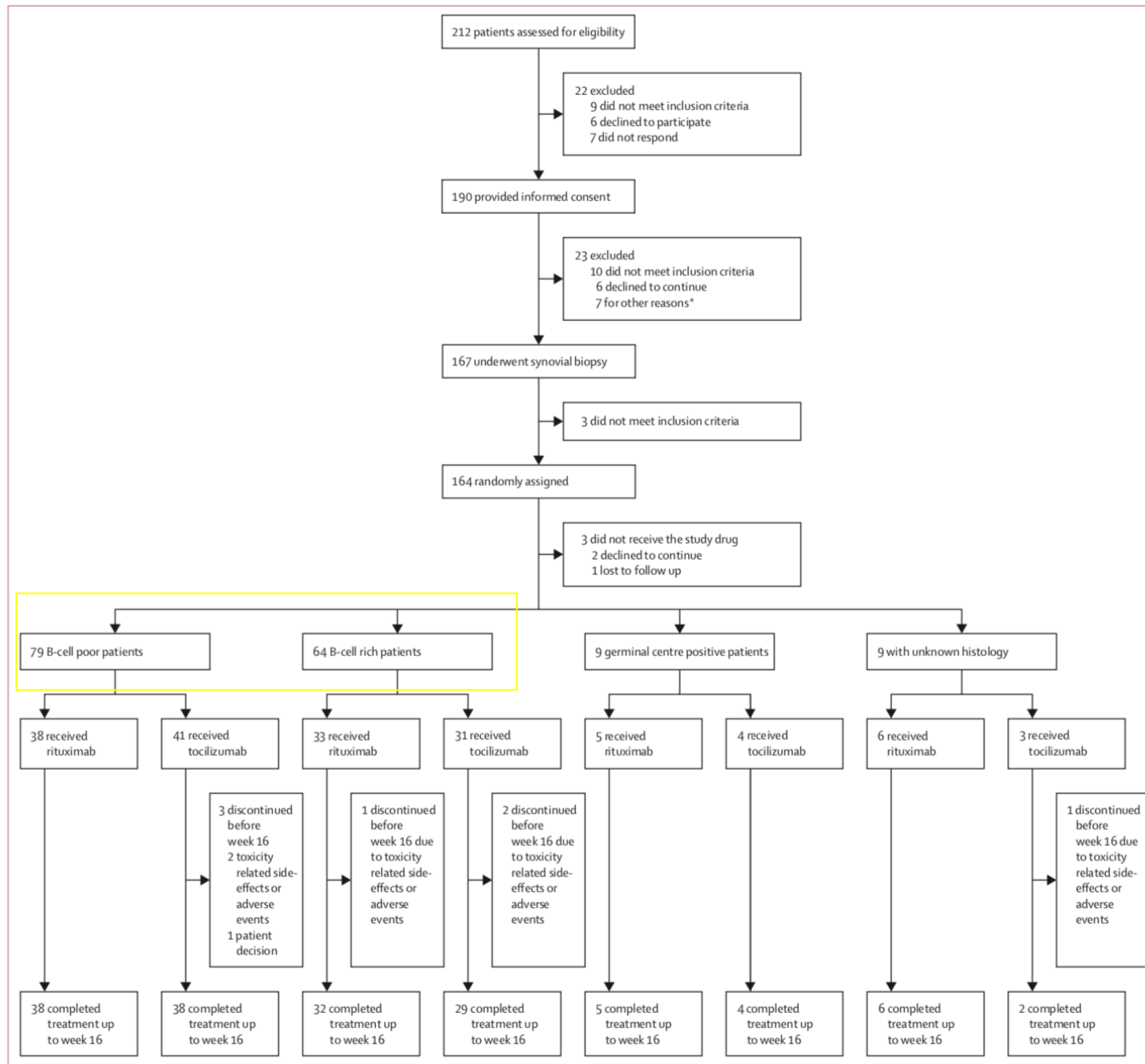


Figure: Trial profile

*Six patients did not have suitable joints at biopsy and one for clinical reasons unrelated to rheumatoid arthritis.

Η R4RA κλινική δοκιμή

- Συνολικά 164 ασθενείς με ρευματοειδή αρθρίτιδα υποβλήθηκαν σε βιοψία αρθρικού αρθρικού υγρού πριν από τη θεραπεία και ταξινομήθηκαν ιστολογικά είτε ως φτωχοί σε B-κύτταρα είτε ως πλούσιοι σε B-κύτταρα
- Στη συνέχεια, κατανεμήθηκαν τυχαία στην ομάδα tocilizumab ή στην ομάδα rituximab), με ιστολογική ταξινόμηση ως παράγοντα στρωματοποίησης
- Μετά την ταξινόμηση με αλληλουχία RNA, το ποσοστό απόκρισης CDAI50% ήταν σημαντικά υψηλότερο στην ομάδα του tocilizumab σε σύγκριση με την ομάδα του rituximab



ΣΥΝΕΠΕΙΕΣ

- Η κλινική δοκιμή R4RA αντιπροσωπεύει ένα ορόσημο στη μηχανιστική διερεύνηση σε επίπεδο ιστού ασθένειας της σχέσης μεταξύ του τρόπου δράσης του φαρμάκου και της κλινικής απόκρισης
- Σε σύγκριση με την τρέχουσα κλινική προσέγγιση, το R4RA δείχνει ότι σε ασθενείς με χαμηλή ή απουσία υπογραφής έκφρασης Β-λεμφοκυττάρων στον αρθρικό ιστό με αλληλούχιση RNA, η θεραπεία με αναστολή του υποδοχέα IL-6 δηλαδή με τοσιλιζουμάμπη— είναι ανώτερη από τη στόχευση Β-λεμφοκυττάρων με rituximab

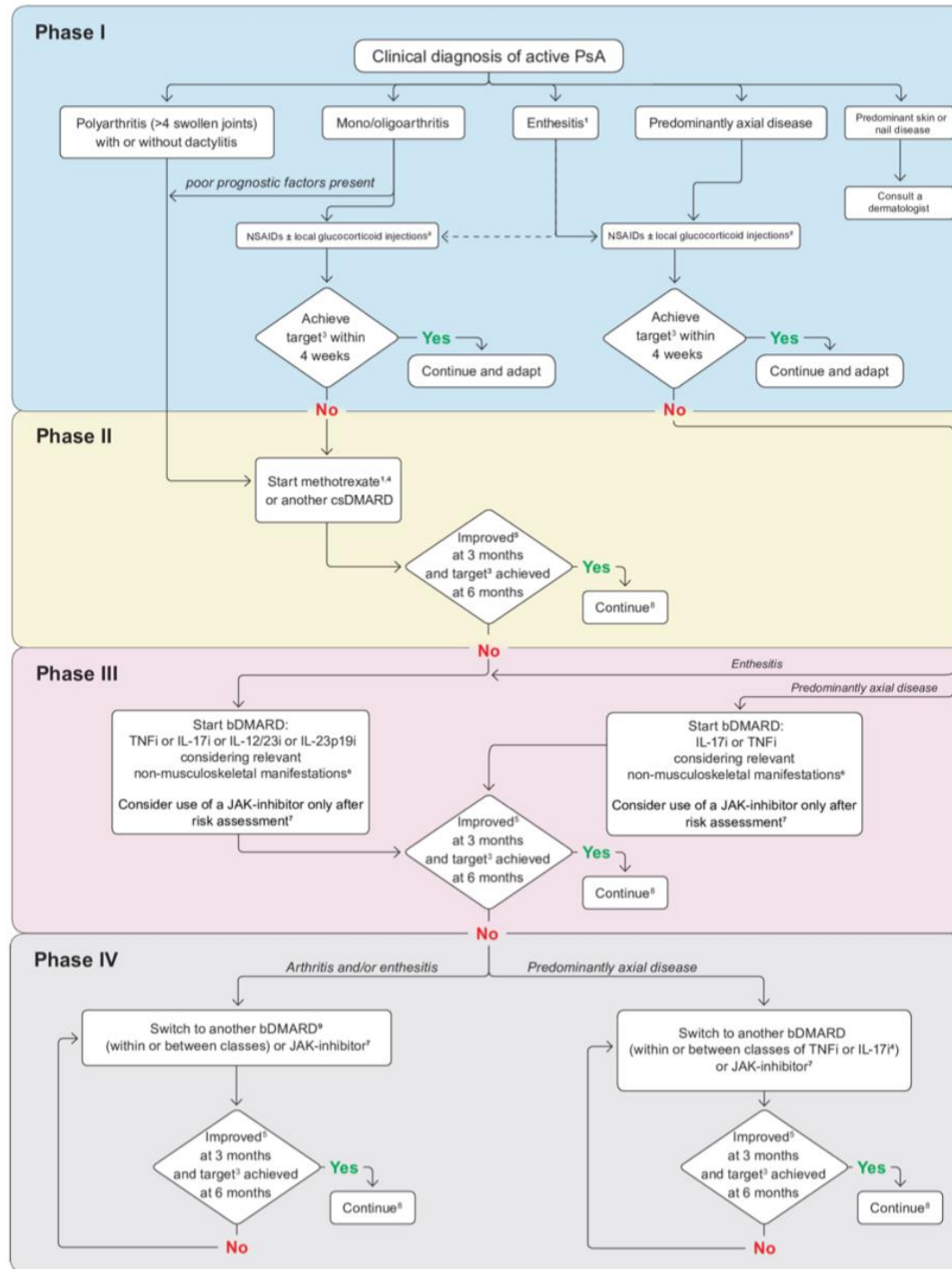


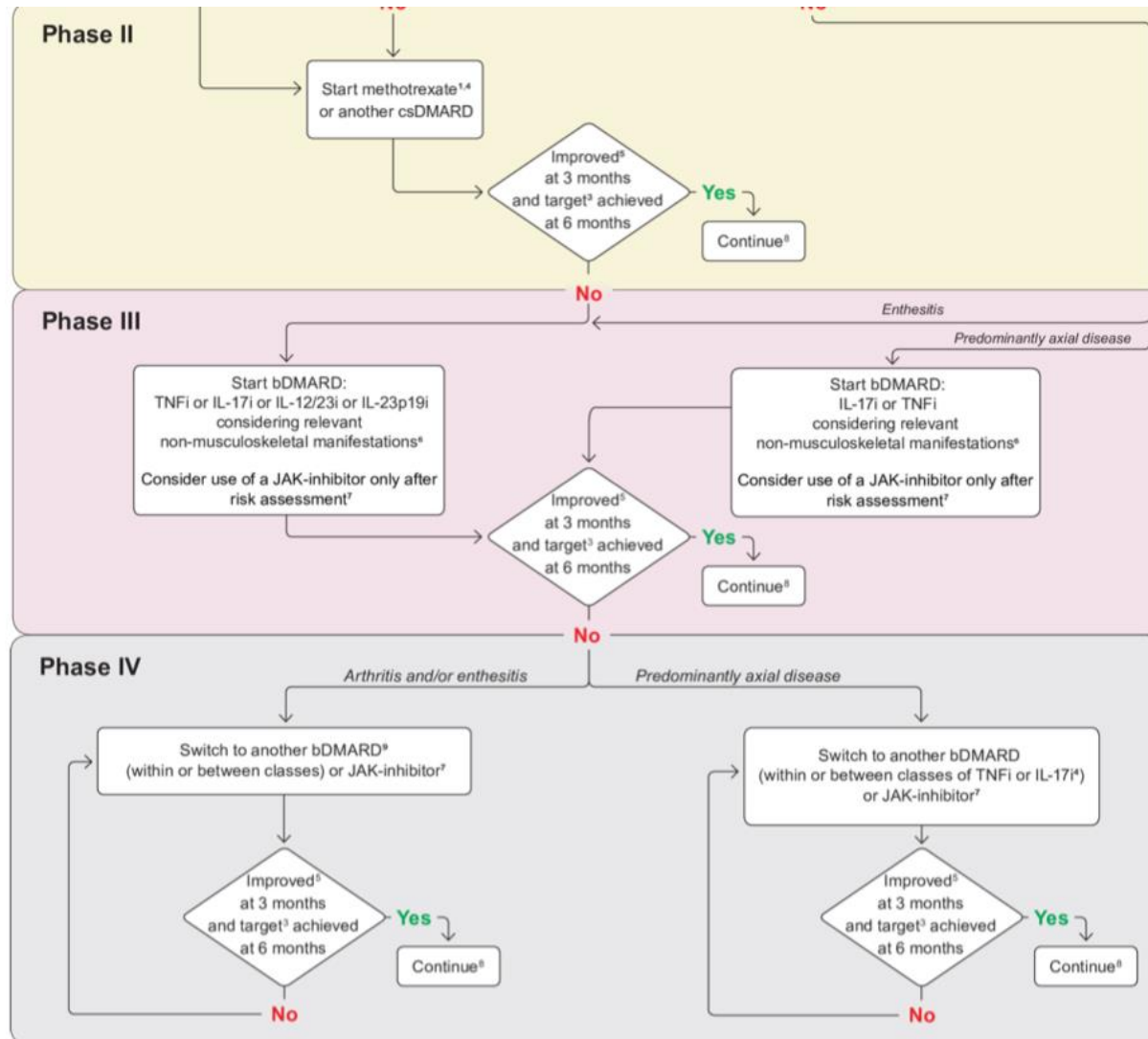


OPEN ACCESS

EULAR recommendations for the management of psoriatic arthritis with pharmacological therapies: 2023 update

Laure Gossec ,^{1,2} Andreas Kerschbaumer ,³ Ricardo J O Ferreira ,^{4,5} Daniel Aletaha ,³ Xenofon Baraliakos ,⁶ Heidi Bertheussen,⁷ Wolf-Henning Boehncke,⁸ Bente Appel Esbensen ,^{9,10} Iain B McInnes,¹¹ Dennis McGonagle,^{12,13} Kevin L Winthrop ,¹⁴ Andra Balanescu,¹⁵ Peter V Balint,¹⁶ Gerd R Burmester ,¹⁷ Juan D Cañete ,^{18,19} Pascal Claudepierre,^{20,21} Lihi Eder ,²² Merete Lund Hetland ,^{23,24} Annamaria Iagnocco ,²⁵ Lars Erik Kristensen,^{26,27} Rik Lories,^{28,29} Rubén Queiro ,^{30,31} Daniele Mauro ,³² Helena Marzo-Ortega ,^{12,13} Philip J Mease ,^{33,34} Peter Nash ,³⁵ Wendy Wagenaar,^{36,37} Laura Savage,³⁸ Georg Schett ,³⁹ Stephanie J W Shoop-Worrall ,⁴⁰ Yoshiya Tanaka ,⁴¹ Filip E Van den Bosch ,⁴² Annette van der Helm-van Mil,⁴³ Alen Zabotti ,⁴⁴ Désirée van der Heijde ,⁴³ Josef S Smolen³





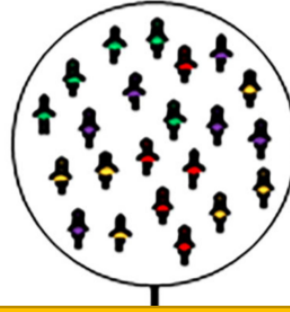
Precision medicine using different biological DMARDs based on characteristic phenotypes of peripheral T helper cells in psoriatic arthritis

Ippei Miyagawa¹, Shingo Nakayamada¹, Kazuhisa Nakano¹, Satoshi Kubo¹, Shigeru Iwata¹, Yusuke Miyazaki¹, Maiko Yoshikawa¹, Hiroko Yoshinari¹ and Yoshiya Tanaka¹



Precision Medicine
Stratification (divide into subgroups)
and use of targeted therapy

PsA



IL-17
inhibitors

IL-12/23(p40)
inhibitor

TNF or IL-17
inhibitors

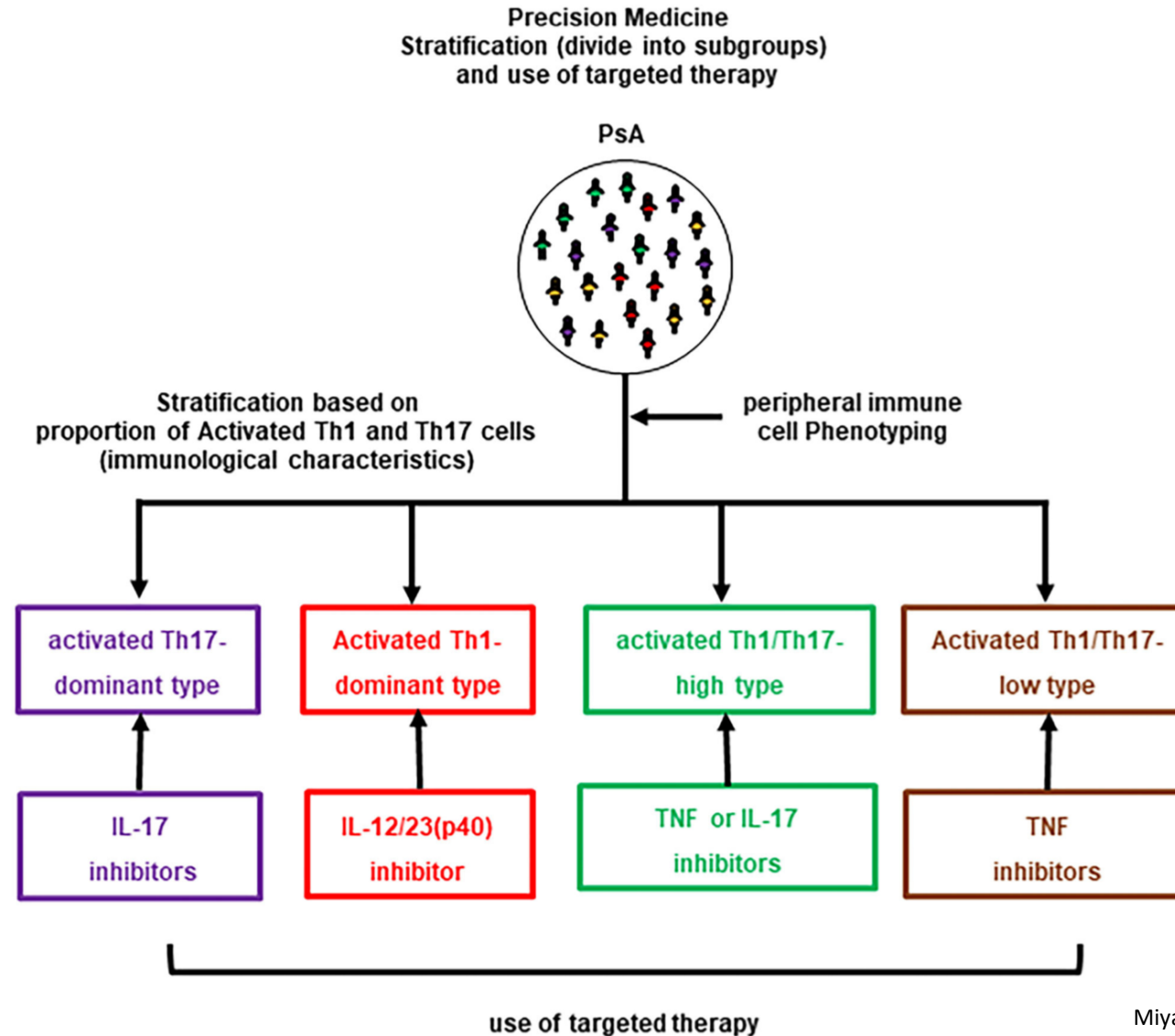
TNF
inhibitors



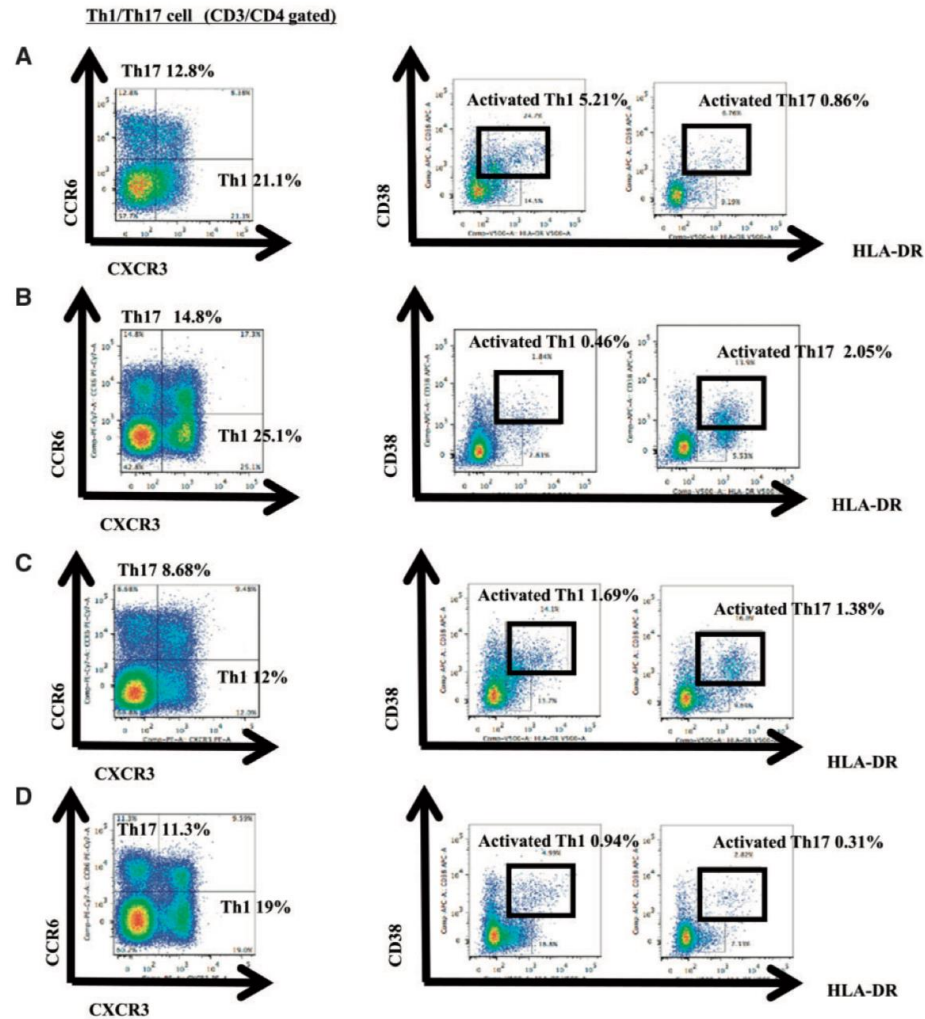
use of targeted therapy



Standard biologics therapy (Guidelines) vs **Strategic** Biologics Therapy



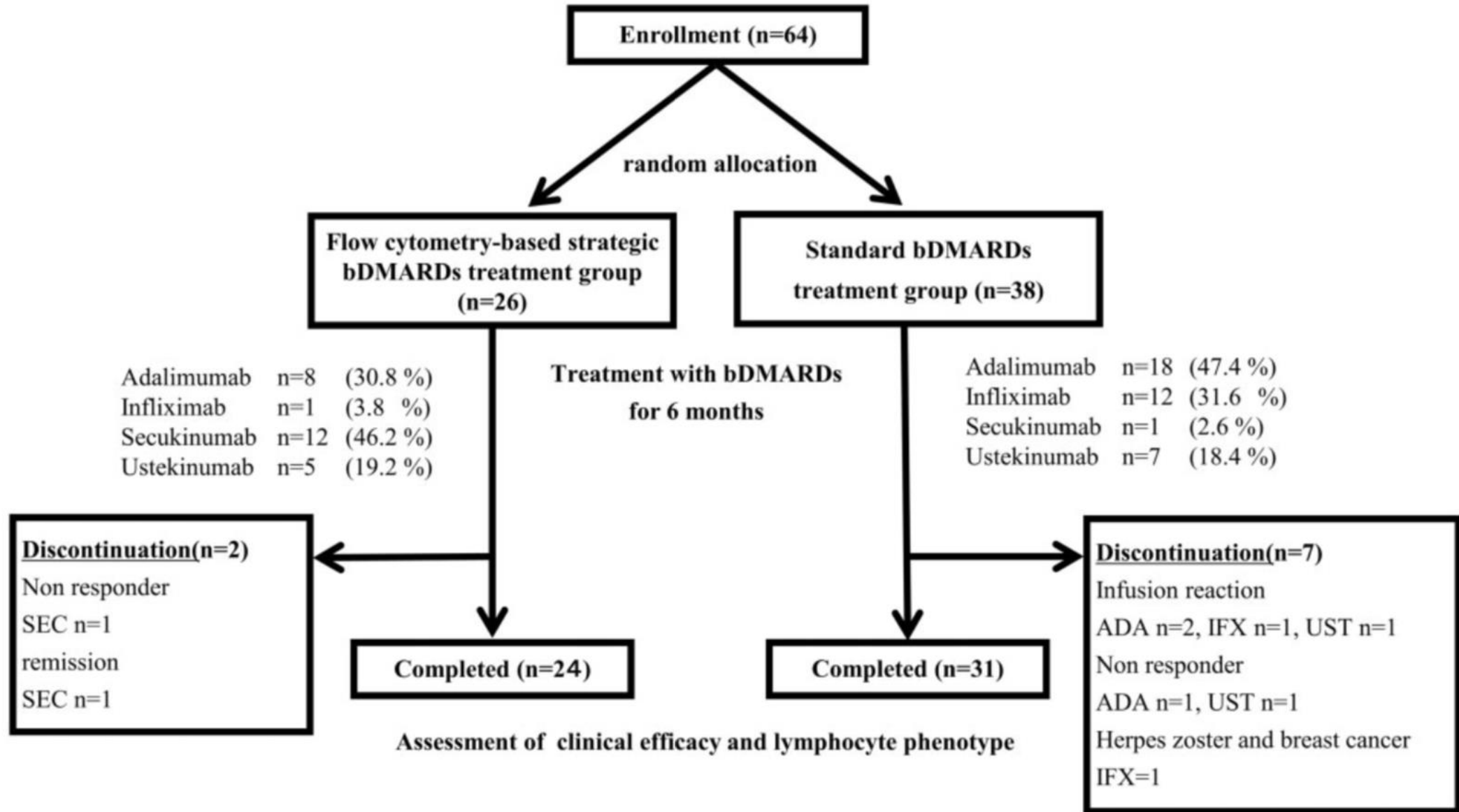
PsA was classified into four types by peripheral blood lymphocyte analysis



Twenty-six patients with PsA were classified into the following four types according to the phenotype of CD4⁺T cells in peripheral blood: (A) activated Th1-dominant (6 cases); (B) activated Th17-dominant (10 cases); (C) activated Th1/Th17-high (4 cases); and (D) activated Th1/Th17-low (6 cases). Results are representative of each group.



FIG. 1 Study design

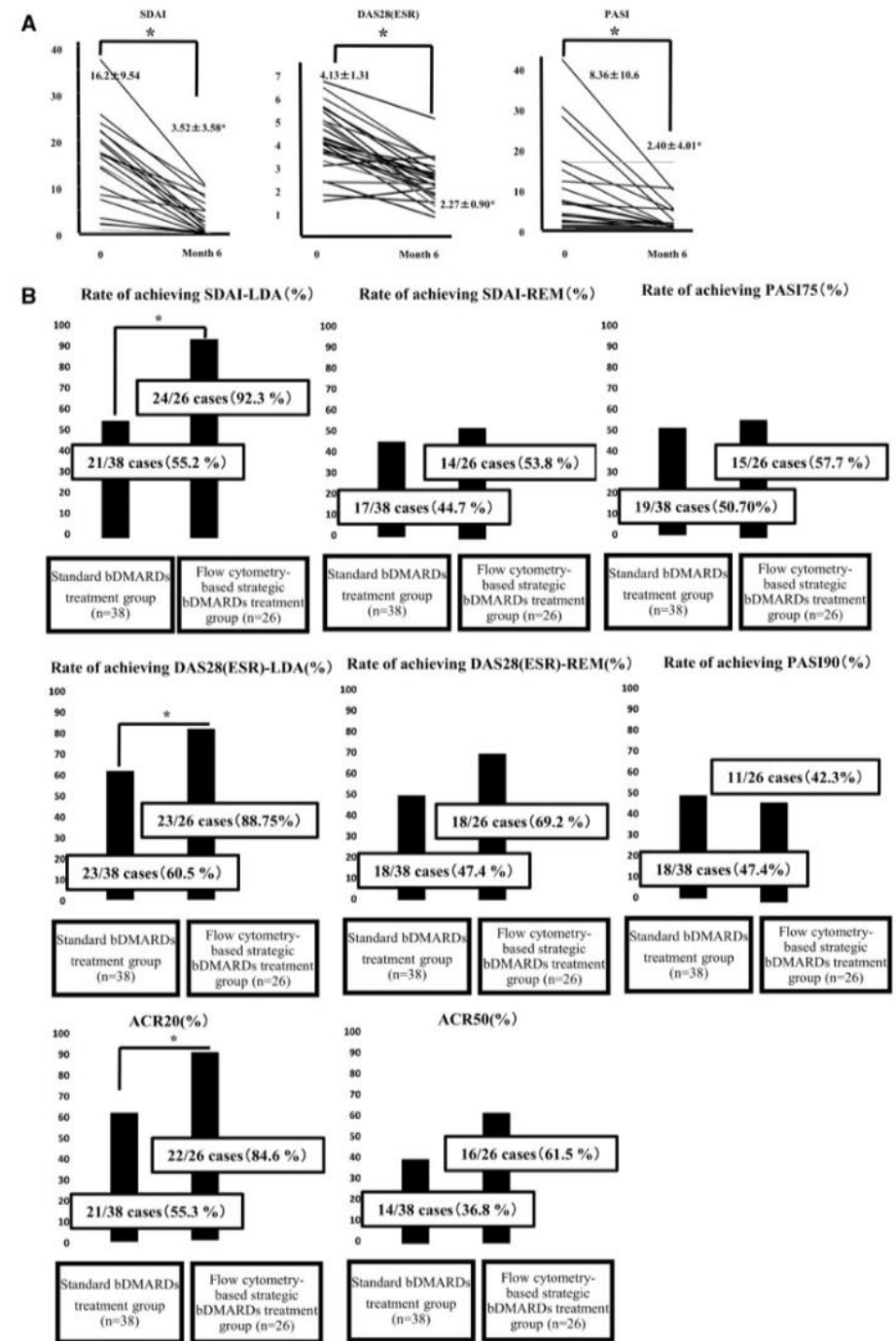


Results

- The therapeutic efficacy of strategic biologics therapy was compared with that of standard biologics therapy.
- The therapeutic efficacy was higher in the strategic bDMARDs treatment group than in the standard bDMARDs treatment group, indicating the value of precision medicine in the selection of specific bDMARDs in the treatment of PsA.



Strategic choice of biologic products based on lymphocyte phenotype and their efficacy



What is Autoprediction?



AutoPrediction©

A disease-specific, parameter-dependent algorithm, which predicts customized response to individualized treatment

- Flexible (not all parameters needed to predict)
- Semi-quantitative
- Includes demographic, clinical, laboratory and/or histological parameters
- Immunological (autoantibody, serum cytokines, chemokines, cell subsets (Th1, Th17, Tregs, Bregs etc, surface cell subsets i.e CD24, CD27, CD39)



Systemic Sclerosis

87 patients with Systemic Sclerosis,
who responded to treatment

92 patients with Systemic Sclerosis,
who did not respond to treatment

Serum/plasma

Blood

DNA/RNA

PBMC

Saliva

Urine

Stool Sample

534 analytes

(17 autoab specificities,
45 cytokines,
19 angiogenic factors,
17 fibrogenic factors
27 cell-surface markers,
19 chemokines, ...)



Systemic Sclerosis

87 patients with Systemic Sclerosis,
who responded to treatment

92 patients with Systemic Sclerosis,
who did not respond to treatment

Type of Treatment

IVIG

MMF

CYC

Cyclosporin

MTX

AZT

Rituximab

Anti-TNF

Infliximab

....

Specific treatment for Pulmonary arterial hypertension, digital ulcers, etc



Immunogenetic, Genetic, Pharmacogenetics,

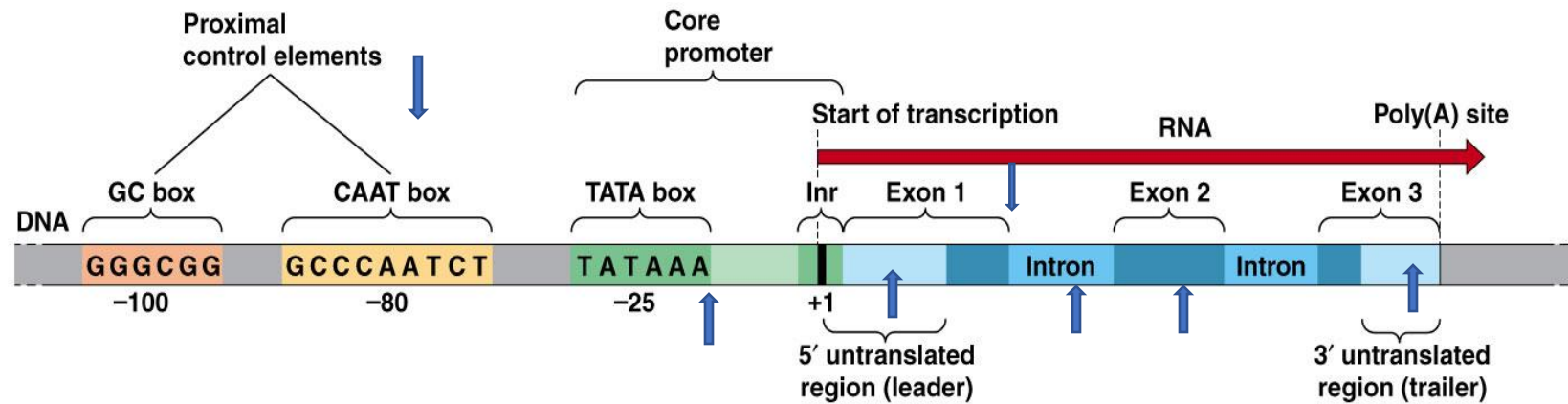
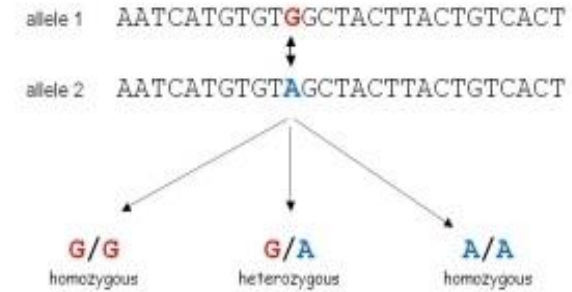
- **594,322 SNPs (single nucleotide polymorphisms)**

Infinium Global Screening Array-24 v2.0 BeadChip,
Illumina



Genetic variation- SNPs

- **Single Nucleotide Polymorphism** : the most common form of genetic variation
- Change in **one** nucleotide
- Usually one common (major) and one rare (minor) allele
- **Millions** of SNPs across the genome
- Different effects based on their location



AutoPrediction -examples

- Prediction algorithm for response of SSc patients to individual treatment-
- **infliximab** (27parameters 14 with positive predictive value and 13 with negative predictive value)



General Comments

- **Less than 30%** of the predictive score can be attributed to **autoantibodies** (mono or poly-specificity or combination)
- **More than 40%** of the total predictive score is to be attributed to **cell-subset phenotyping**
- **Less than 20%** of the predictive score can be based on serum cytokines
- **Less than 10%** of the scoring is due to **genotyping**



Example A

Algorithm % prediction (0-92%)= ZxPre-Predict algorithm
(3.2autoab1)+(9.1autoab2)-(6.7autoab3)-
(1.2 autoab levels/cut off autoab4)+
(3.1cytokine1)+ (1.7cytokine2)+..... +
(3.9 xCD4+CD27+) (+2.2CD4-CD39+)+
+(9.2 B10)+ (1.3CD4C+D25+).....

Prediction algorithm for response of SSc to infliximab
(27 parameters 14 with positive predictive value &
13 with negative predictive value



The roadmap to Autoprediction and Precision Medicine is not an easy task





DO NOT PUT THE CAR
BEFORE THE HORSE

GUIDELINES και πάλι guidelines μέχρι.....

