

Serum autoantibody profiling of primary Sjögren's syndrome patients reveals novel biomarkers associated with the disease, disease activity, and clinical response to VAY736

Petra Budde¹, Julie Doucet², Hans-Dieter Zucht¹, Rémi Kazma², Paul Maguire², Alexandre Avrameas², Marie-Anne Valentin², Stephen Oliver², Peter Schulz-Knappe¹, Alessandra Vitaliti²

¹Protagen AG, Dortmund, Germany; ²Novartis Institutes for Biomedical Research, Basel, Switzerland

Introduction

- Overexpression of B cell activating factor (BAFF) in salivary glands contributes to the pathogenesis of primary Sjögren's syndrome (pSS) by promoting autoantibody (AAB) production.
- Treatment of pSS patients with VAY736, an anti-BAFF receptor mAb, appears promising and was associated with a depletion of circulating B cells and a positive therapeutic effect [1].
- In addition to the classical anti-SS-A/Ro and anti-SS-B/La, a broader set of AABs may reflect B cell disturbances in pSS and could serve as markers during clinical development of novel pSS therapeutics.

Objectives

- The study was undertaken to explore novel AABs in pSS and their associations with the disease, disease activity, and clinical response to VAY736.

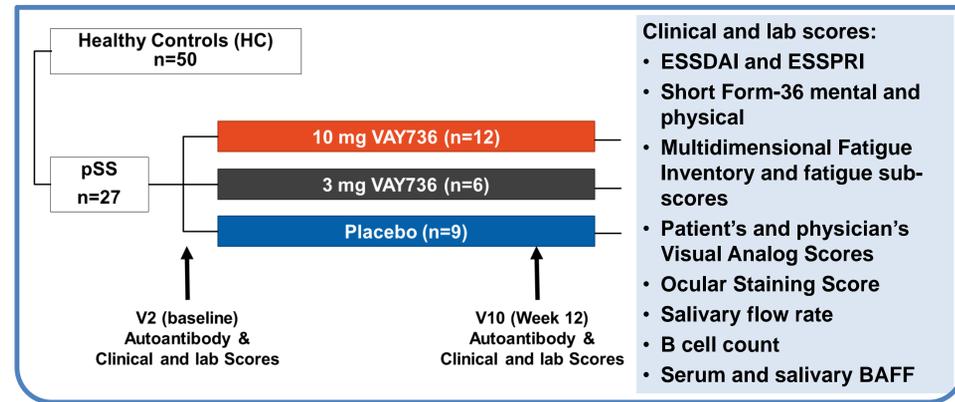


Fig. 1: Study design

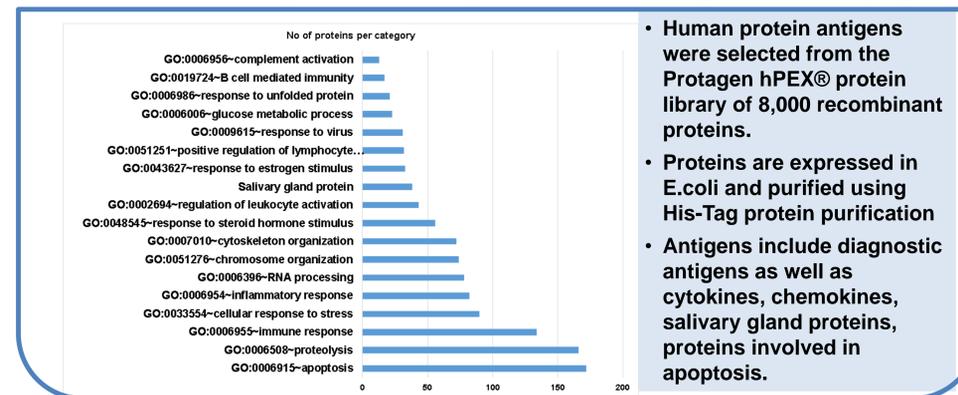
- Serum samples were collected from pSS patients pre- and post-VAY736 administration, and from age and gender-matched HCs (Fig. 3). Statistical analysis was conducted as detailed in Table 1.

Goal	Statistical approach
Identification of AABs associated with pSS	Parameters: AAB levels in HCs and in pSS patients at baseline Tests: Wilcoxon rank sum test, significance analysis of microarrays, and comparison of the 90 th quantiles between groups.
Identification of AABs associated with pSS activity	Parameters: AAB levels in pSS patients at baseline and week 12, as well as relative changes Tests: Pearson correlation with clinical and lab scores
Assessment of VAY736 treatment-specific changes in AAB levels	Parameters: AAB levels in pSS patients at baseline and Week 12 Test: linear mixed-effects models adjusting for dosage, age, and gender effects

Table 1: Statistical approach

Methods

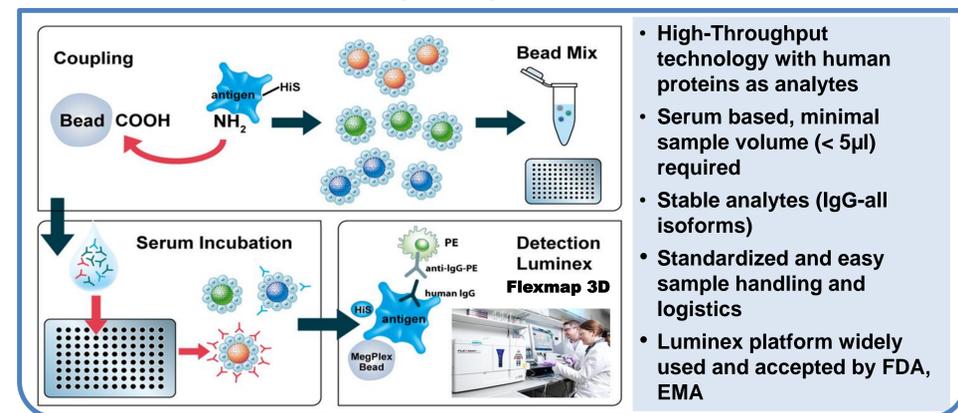
- A discovery screen was designed comprising 1,596 antigens, which are directly relevant to pSS associated processes (Fig.2).



- Human protein antigens were selected from the Protagen hPEX® protein library of 8,000 recombinant proteins.
- Proteins are expressed in E.coli and purified using His-Tag protein purification
- Antigens include diagnostic antigens as well as cytokines, chemokines, salivary gland proteins, proteins involved in apoptosis.

Fig. 2: Overview of antigens included in this study

- IgG antibodies were measured using the bead-based Luminex® xMAP® platform SeroTag® (Fig.3).



- High-Throughput technology with human proteins as analytes
- Serum based, minimal sample volume (< 5µl) required
- Stable analytes (IgG-all isoforms)
- Standardized and easy sample handling and logistics
- Luminex platform widely used and accepted by FDA, EMA

Fig. 3: Schematic SeroTag workflow of bead-based assays

Results

- Compared to HC, pSS patients had increased IgG AAB levels to 36 antigens, including the known SSA and SSB (p<0.05). Examples are shown in Fig.4.

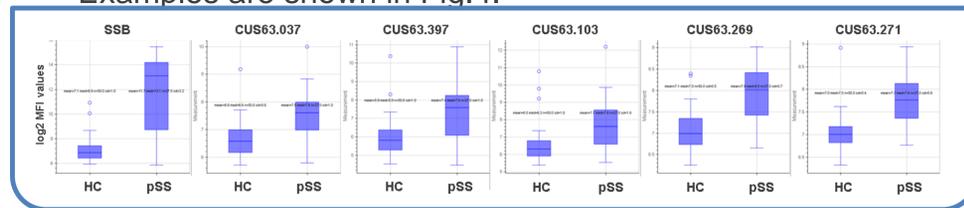


Fig. 4: Box and Whisker plots showing novel autoantibodies in HC and pSS

Results (continued)

- SS-A/Ro and SS-B/La AABs were not associated with disease activity or response to treatment.
- When combining all treatment arms, 48 AABs were significantly correlated with different clinical outcome measures (p<0.05, |r|>0.46). Fig. 5A shows AABs associated with ESSDAI.
- 12 baseline AABs correlated with change in clinical outcome measures over 12 weeks in VAY736-treated patients (Fig 5B).

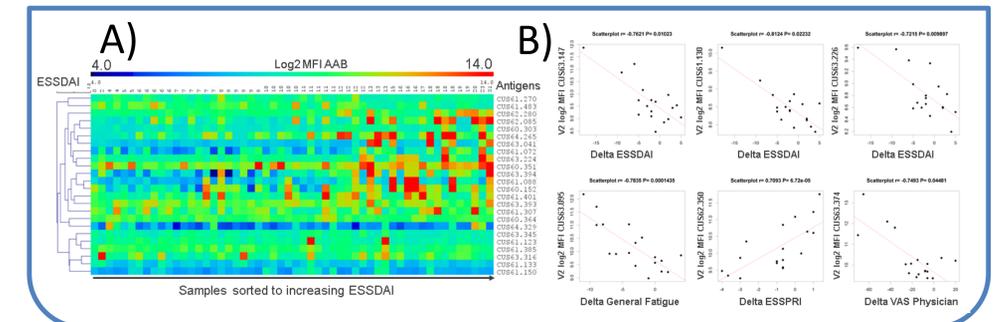


Fig. 5: Autoantibodies correlating with clinical outcome scores

- 51 serum AAB levels correlated with salivary BAFF levels, but not BAFF serum levels (p<0.05, |r|>0.55).

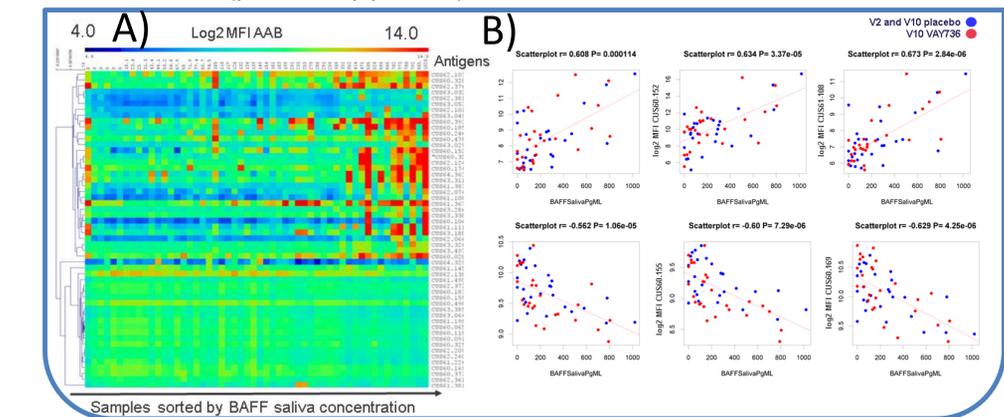


Fig. 6: Autoantibodies correlating with salivary BAFF levels

- No reduction in AABs levels was observed in response to VAY736 at week 12.

Conclusions

The genes encoding novel antigens are involved in apoptotic, anti-viral, metabolic, inflammatory, blood coagulation and B-cell processes, suggesting a possible link to the disease pathology. Further large-scale studies are needed to confirm the value of these markers.

References

1. Dörner T et al. Arthritis Rheum 2016; 68(suppl S10):4051