



Expression, Purification, and Evaluation of Anti-IL-2R α Antibody Secreted by *Leishmania tarentolae*

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Abstract

Producing a functional anti-IL-2R α antibody in *Leishmania tarentolae*, a trypanosomatid protozoan non-pathogenic to human, is a cost-effective and safe alternative for the production of therapeutic recombinant proteins. Expression cassettes encoding heavy and light chains of the anti-IL-2R α antibody were cloned into two separate pLEXSY vectors. The plasmids were then used to transfect *L. tarentolae* laboratory strain p10 and stable expression of a recombinant humanized anti-IL-2R α antibody was obtained. The heavy and light chains successfully assembled to produce a tetrameric 150 kDa antibody. The antibody was glycosylated and functional as assessed by enzymatic deglycosylation and cell proliferation assays respectively. *L. tarentolae* can be used as an efficient, cost and labor-effective expression system for the production of therapeutic recombinant proteins.

Keywords Anti-IL-2R α antibody · *Leishmania tarentolae* · Monoclonal antibody

Introduction

Biopharmaceuticals constitute a considerable proportion of marketed drugs. Monoclonal antibodies (mAbs) and related products are the main biopharmaceuticals. MAbs are used as biomedical (Chon and Zarbis-Papastois 2011) research molecules for diagnostic, and therapeutic purposes (Stockwin and Holmes 2003; Majidi et al. 2009). The required therapeutic doses of mAbs however are much higher than that of other biological products (Jones et al. 2007; Iberg

and Hawiger 2019). Prokaryotic hosts constitute the most widely used protein expression systems. Albeit limitations to prokaryotic expression of antibodies, i.e., improper folding and assembly of the antibody fragments, improper disulfide bond formation, and lack of glycosylation have led to the employment of mammalian expression systems (De Marco 2009; Ke and Berkmen 2014; Dhara et al. 2018).

Glycosylation is a eukaryotic post-translational modification which is critical to antibody function (Jefferis et al. 1998; Jefferis 2005). *Leishmania tarentolae*, a trypanosomatid protozoan not pathogenic to humans (Breitling et al. 2002; Niimi 2012), has been developed as an alternative eukaryotic protein expression system for the production of recombinant proteins with mammalian-like posttranslational modifications. The advantages of this system include easy handling, faster growth rate, and cost-effectiveness (Niimi 2012; Fritsche et al. 2007). These features render it a promising system for the production of biopharmaceutical proteins.

Anti-IL-2R α antibodies prevent IL-2-induced clonal expansion of activated lymphocytes and decrease their survival through binding to the α chain of IL-2 receptor (Church 2003; Kandus et al 2010). This can be effective in treatment of leukemias and autoimmune diseases, such as rheumatoid arthritis (Ben-Ari 2004). The IgG1 antibody, has a single glycosylation site (Hossler et al. 2009; Del Val

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et al. 2016), Asp296 on the heavy chain. In the current study, *L. tarentolae* was used as a new host for the production of this antibody.

Materials and Methods

Several culture media, including LB, BHI, and RPMI₁₆₄₀, were used to prepare a compound LBR medium (Table 1). RPMI₁₆₄₀ was supplemented with 10% (v/v) FBS. To avoid bacterial contamination, penicillin, 50 IU and streptomycin 50 µg/ml were added.

Strains and Culture Conditions

Leishmania tarentolae laboratory strain p10 (Jena Bioscience, Germany) was cultured in RPMI₁₆₄₀ supplemented with 10% FBS at 26 °C while shaking at 70 rpm. New medium was added every 2 to 3 days. TOP10 *Escherichia coli* cells (CinnaGene, Iran) were used to propagate recombinant plasmids.

Plasmid Construction

The amino acid sequences of the two chains of humanized anti-IL-2R α antibody were obtained from NCBI, PDB, and the drug bank (DB00111) databases.

Humanized Anti-CD25 Heavy Chain 1

QVQLVQSGAEVKKPGSSVKVCSKASGYTFTSYRM-
HWVRQAPGQGLEWIGYINPSTGYTEY

NQKFKDKATITADESTNTAYMELSSLRSED-
TAVYYCARGGGVFDYWGQGTTLTVSSGSPV

FPLAPSSKSTSGGTAALGCLVKDYFPEPVTVS-
WNSGALTSGVHTFPAVLQSSGLYSLSSV

VTVPSSSLGTQTYICNVNHKPSNTKVDKKAEPK-
SCDKTHTCPPCPAPELGGPSVFLFPP

KPKDTLMISRTPEVTCVVDVSHEDPEVKFNWY-
VDGVEVHNAKTKPREEQYNSTYRVVSV

LTVLHQDWLNGKEYKCKVSNKALPAPIEK-
TISKAKGQPREPQVYTLPPSRDELTKNQVSL
TCLVKGFYPSDIAVEWESNGQPENNYKTTPPV-
LDSGDSFFLYSKLTVDKSRWQQGNVFC
SVMHEALHNHYTQKSLSLSPGK

> Humanized Anti-CD25 Light Chain 1

DIQMTQSPSTLSASVGDRVTITCSASSISYMHW-
YQQKPGKAPKLLIYTTSNLASGVPAR

FSGSGSGTEFTLTISSLQPDDFATYYCHQRSTYPLT-
FGSGTKVEVKRTVAAPSFIFFPS

DEQLKSGTASVCLLNNFYPREAKVQWKVD-
NALQSGNSQESVTEQDSKDYSLSTLTL

SKADYEKHKVYACEVTHQGLSSPVTKSFNR

Their molecular characteristics were adjusted based on Zenapax monograph (CAS Number: 152923-56-3: <https://www.clearsynth.com/en/CSO16306.html>). Protein sequences were reverse-translated to DNA using an online codon optimization tool (Integrated DNA Technologies. <https://www.euidtdnacom/CodonOpt>). Restriction enzymes sites for *Bgl*III and *Nhe*I were inserted at the 5' and 3' ends respectively. Vector pLEXSY (Jena Bioscience, Germany) was used as the backbone for designing the expression construct. The expression construct was designed as follows: a signal peptide sequence optimized for high-level protein expression in CHO cell line (SEQ ID No 9: <https://patents.google.com/patent/US20100240097A1/en>. Sequence: MMRPIVLVLLFATSALAQV (Young and Rance 2012) was placed upstream of the sequences encoding the humanized heavy or light chain of anti-IL-2R α antibody; ACC sequence (threonine amino acid) was placed between *Bgl*III restriction site and the signal peptide, to increase expression levels, according to the pLEXSY manufacturer guidelines.

Using signalp 4.1 server, in silico prediction of the signal peptide cleavage site in the constructs confirmed the expected cleavage site. The expression constructs were commercially synthesized and provided as inserts in recombinant pGH vectors (GenRay, China). These vectors were then digested by *Bgl*III and *Nhe*I restriction enzymes to either produce a 1407 bp fragment encoding the heavy chain or a 708 bp fragment encoding the light chain. The longer and shorter fragments were cloned between *Bgl*III and *Nhe*I restriction sites of pLEXSY-hyg2 and pLEXSY-neo2 (Jena Bioscience, Germany) vectors, respectively, generating pLEXSY-hyg2-H chain and pLEXSY-neo2-L chain plasmids. The vectors were then used to transform TOP10 *Escherichia coli* cells; the recombinant clones were selected and identified by colony PCR using P1442 (CCGACTGCAACAAGGTGTAG) and A264 (CATCTA TAGAGAAGTACACGTAAAAG) primers flanking the multiple cloning site of pLEXSY plasmid. The selected clones were confirmed by sequencing of the PCR products (Bioneer, Korea).

Table 1 Compound culture media which tested in this research

Name of culture medium	Composition
LBR	LB (40%), BHI (40%), RPMI ₁₆₄₀ (20%)
BHI-Hemin	BHI + hemin (5 mg/l)
LBR-FBS	LB (40%), BHI (40%), RPMI ₁₆₄₀ (20%) + 10% FBS
BHI-FBS	BHI + FBS (10%)

Transfection of *L. tarentolae* with the Expression Constructs

The recombinant vectors were extracted from *E. coli* using a plasmid extraction kit (Bioneer, Korea) and digested with *Swa*I restriction enzyme (Fermentas, Lithuania), yielding the following fragments: 2864 bp (the same fragment length for both vectors), and 5679 bp (pLEXSY-neo2-L chain) or 6600 bp (pLEXSY-hyg2-H chain). After electrophoresis, the respective expression cassettes (5679 bp and 6600 bp) were eluted from the gel (AccuPrep® Gel Purification kit- Bioneer, Korea) for subsequent transfections of *L. tarentolae*. *L. tarentolae* P10 was cultured in RPMI₁₆₄₀ medium supplemented with 10% (v/v) FBS for 48 h and the cell suspension was adjusted to 1×10^8 parasites/ml in electroporation buffer (Eppendorf, Germany). Electroporation (Eppendorf, Germany) was performed using 5–10 µg of DNA (1300 V/5 ms). About 24 h after culturing in RPMI₁₆₄₀ supplemented with 10% (v/v) FBS, hygromycin B (25 µg/ml) and neomycin G480 (25 µg/ml) were added. Stringent selection of colonies were performed by increasing the concentration of hygromycin up to 200 µg/ml and neomycin up to 100 µg/ml, over 10 days. To evaluate the insertion of the expression cassettes into the *ssu* locus of *L. tarentolae*, following DNA extraction, the cassettes were amplified using F3001 and A264 primers (Table 2).

Expression of Recombinant Anti-IL-2R α Antibody

Ice-cold supernatant of the selected positive transfected cells grown for 48 h in agitated cultures in LBR medium was precipitated with ice-cold trichloroacetic acid (TCA, final concentration 10%, v/v) for 30 min and centrifuged at 10,000×g and 4 °C for 5 min. The pellet was washed with ethanol, dried, and resuspended in SDS-PAGE loading buffer. Reduced and non-reduced samples were prepared and run on a 12% SDS PAGE gel. Western blotting analysis was performed using mouse polyclonal anti-human IgG conjugated with horseradish peroxidase (HRP) (Dako, Denmark). Densitometric measurement of the protein yield was performed using ImageJ.1.50a software, and the yield calculated based on a standard curve.

Table 2 Oligonucleotide sequences

Name of primer	Sequence (5' to 3')
P 1442	CCGACTGCAACAAGGTGTAG
A 264	CATCTATAGAGAAGTACACGTAAG
F 3001	GATCTGGTTGATTCTGCCAGTAG

Purification of the Humanized Anti-IL-2R α Antibody

The cultured transfected *L. tarentolae* promastigotes were centrifuged at 1000×g for 5 min, and the pellet resuspended in fresh medium. The supernatant was collected and the centrifugation repeated, at 8000×g for 5 min, to obtain a second supernatant. The second supernatant was used for IgG1 mAb purification using Protein G affinity chromatography (Amersham, UK) (Sugino and Niimi 2012). The concentration of the purified antibody was measured using a biophotometer (Eppendorf, Germany).

In-Cell ELISA Assay (ICE Assay)

Human peripheral blood lymphocytes (PBLs) were isolated from blood samples obtained from healthy volunteers who provided informed consent, according to the guidelines of the local ethics committee. PBLs were isolated on Ficoll/Hypaque gradients (Lymphodex, Germany) and were cultured at a concentration of 1×10^6 cells/ml in the presence of T cell mitogen phytohemagglutinin (PHA, 5 µg/ml), in a 96-well microplate. After 72 h, ICE assay (Mitoscience company, https://www.ispybio.com/search/protocols/elisa_protocol_26.pdf) was performed using PHA-activated T cells expressing the IL-2 α receptor and non-activated T cells as a control group in duplicate. The anti-IL-2R α antibody was used as the primary antibody (5 µg/ml) and HRP-conjugated mouse anti-human antibody (1:1000 dilution) as the secondary antibody. The blank wells (duplicate) did not include the primary antibody. After incubation with the substrate TMB (3,3',5,5'-tetramethylbenzidine), the colorimetric signal was measured within 20 min using a microplate reader, at 650 nm.

Immunocytochemistry (ICC Assay)

PHA-activated and non-activated T cells were stained using an ICC procedure similar to the ICE method, except that the water-insoluble substrate DAB (3,3'-diaminobenzidine) was used for visualizing antibody binding to the target.

Biological Activity Assay (BrdU Assay)

Inhibition of PHA-activated T cell proliferation by the anti-IL-2R α antibody was assessed using Roche BrdU assay kit (cat. no. 11647229001). Isolated PBLs were diluted to 1×10^6 cells/ml in RPMI₁₆₄₀ supplemented with 10% (v/v) FBS, and cultured in 96-well microtiter plates (100 µl of the cell suspension/well; the blank wells were empty; all measurements were performed in triplicate). Wells containing untreated cells were designated as control group 1 (cell proliferation control) and those treated with PHA only as control group 2. The experimental group was treated with

both PHA(5 µg/ml) and anti-IL2R α . All groups were incubated at 37 °C and 5% CO₂. 500 to 700 µg of the purified anti-IL-2R α antibody was added to the experimental group after 2 h of incubation. BrdU was added to the cell cultures after 48 h of incubation and the cells were incubated for an additional 24 h. Harvested cells were stained with antibodies included in the kit, followed by absorbance measurements (370 nm/492 nm) according to the manufacturer's instructions (<https://www.sigmaaldrich.com/content/dam/sigma-aldrich/docs/Roche/Bulletin/1/11647229001bul.pdf>).

Glycosylation Analysis

Deglycosylation of asparagine-linked glycans was done with N-glycosidase F (PNGase F, Sigma) and the SDS-PAGE electrophoretic mobility of the protein before and after the treatment was compared.

Statistical Analysis

Data were analyzed using SPSS.17 software by two-sided unpaired Student's *t*-test. The level of statistical significance was set at $p < 0.05$.

Results

The heavy and light chain genes of anti-IL2R α antibody were synthesized and cloned into pLEXSY vectors. Figure 1 shows the schematic design of gene construction for expression in *L. tarentolae*.

Recombinant pLEXYS vectors were digested by *SawI* restriction enzyme. The linear fragment transfected into *L. tarentolae* contained the genes for antibody, antibiotics and the homologous arms. The integration of expression cassettes into the *L. tarentolae ssu* gene was verified by PCR (Fig. 2).

Confirmation of Protein Expression in *Leishmania* Using SDS-PAGE and Western Blot

The expression of the recombinant 150 kDa anti-IL-2R α antibody was confirmed by SDS-PAGE and western blot analysis under non-reducing conditions (Fig. 3).

Forty eight hours after culture of cells in a medium without FBS the cultures were centrifuged and the proteins in the supernatant were precipitated with TCA and run on 12% SDS-PAGE gel and stained with Coomassie Brilliant Blue G250. The total amount of protein produced was estimated

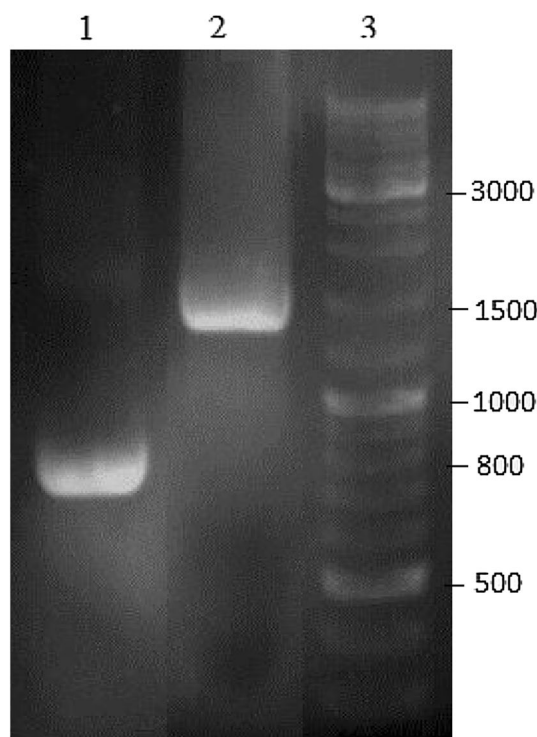


Fig. 2 Confirmation of genomic integration of the anti-IL-2R α antibody heavy and light chain genes by PCR. Diagnostic PCR was performed using F3001 forward primer hybridizing to the *ssu* gene of *L. tarentolae* and A264 reverse primer hybridizing within the construct. *Lane 1*, PCR products from transfected cells, 1900 bp and 2600 bp fragments, confirming integration of light and heavy chains, respectively; *lane 2*, 10,000 bp DNA ladder

using the analysis of bands by ImageJ.1.50a software and compared with the standard sample.

The Specificity of Anti-IL-2R α Antibody

The specificity of anti-IL-2R α antibody was examined by ICE. Student's *t*-test analysis of the mean absorbance values indicated significant differences between the experimental (activated T cells) and control group (untreated T cells) ($p = 0.005$). The results revealed that the recombinant anti-IL-2R α antibody specifically binds to the IL-2 α receptor expressed on the activated T cells but not to untreated T cells (Fig. 4).

ICC Evaluation and Activity of Anti-IL-2R α Antibody

ICC analysis verified binding of the recombinant anti-IL-2R α antibody to the surface of proliferating T cells

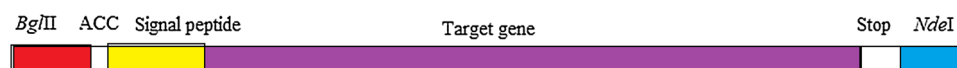


Fig. 1 Schematic design of gene construction

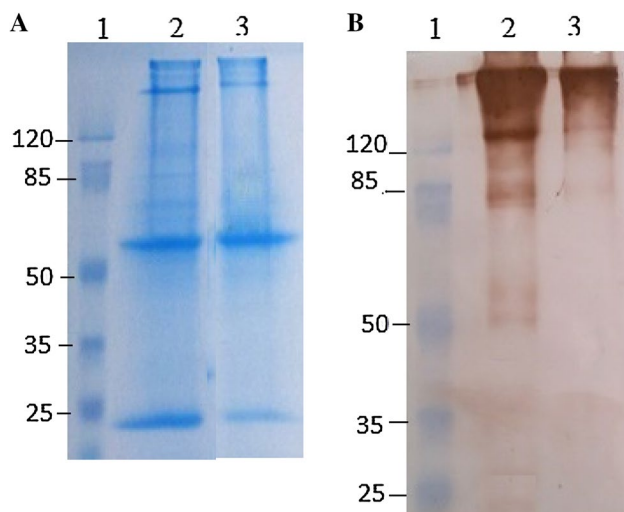


Fig. 3 Analysis of the expressed recombinant 150 kDa anti-IL-2R α antibody by SDS-PAGE (a) and western blotting (b). **a** Lane 1, purified recombinant anti-IL-2R α antibody; lane 2, protein size marker (kDa). **b** Lane 1, human IgG as control; lane 2, purified recombinant anti-IL-2R α antibody; lane 3, protein size marker (kDa)

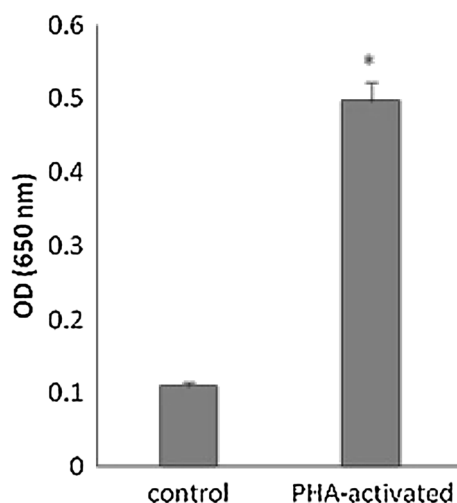


Fig. 4 ICE assay with the expressed recombinant anti-IL-2R α antibody. The experiment was performed using T cells activated by PHA, with T cells not activated by PHA serving as a negative control. Mean absorbance values: negative control, 0.11 ± 0.008 ; experimental group: 0.497 ± 0.013 (mean \pm SD, $n=2$). Student's t -test analysis of the mean absorbance values indicated significant differences between the experimental and control groups ($p=0.005$)

expressing IL-2 α receptor but not to non-expressing T cells (Fig. 5).

To assess the overall function of the antibody we used inhibition of PHA-activated T cell proliferation by the anti-IL-2R α antibody test using Roche BrdU assay kit (Fig. 6).

The proliferation of activated T cells was clearly inhibited by the anti-IL-2R α antibody. Cell proliferation of the

antibody-treated group (experimental; + PHA, + antibody) was almost the same as in control group 1 (no PHA or antibody activation; $p=0.29$). The proliferation of PHA-activated control group (control group 2; +PHA, no antibody) was significantly higher than those of the experimental and control groups ($p=0.0002$). As anticipated, N-glycan removal by PNGase F resulted in an SDS-PAGE mobility shift of the heavy chain of the antibody (Fig. 7).

Discussion

MAbs and related products comprise a major proportion of the biopharmaceutical market (Chon and Zarbis-Papastoitsis 2011) but their high therapeutic doses necessitate a large-scale and cost-effective manufacturing process (Jones et al. 2007). Although recombinant protein production in prokaryotic expression systems is cost-effective, the production of full-length monoclonal antibodies in these systems presents a number of challenges due to the size and complexity of the expressed molecules. Consequently, the mammalian cells are the primary choice for the production of mAbs, with the CHO cell lines the most commonly used expression system. Proper folding and human-like post-translational modifications of the recombinant proteins comprise the main advantages of these systems (Khan 2013). However, recombinant protein production in the mammalian cell lines requires more expensive equipment and materials than other expression systems, e.g., *E. coli*. Moreover, these expression systems are at a risk of contamination with animal viruses (Berting et al. 2010). Considering the above, a novel protein expression system based on *L. tarentolae*, was developed (Fritsche et al. 2007; Niimi 2012).

Here, we produced a recombinant humanized anti-IL-2R α antibody in *L. tarentolae* cultured in a low-cost medium (LBR). Identification of a full-length 150 kDa antibody in the culture medium by SDS-PAGE and western blotting under non-reducing conditions indicated correct assembly of the four chains, suggesting proper formation of disulfide bridges. This was in accordance with the findings of Sugino and Niimi (Sugino and Niimi 2012) who produced functional human laminin (LM)-332, a large heterotrimeric glycoprotein, in *L. tarentolae*. It was also consistent with the observations of Jørgensen et al. (2014) who expressed scFv-Fc in this system. No accumulation of antibody fragments was detected in cell lysates (data not shown), which implies that the CHO-specific signal peptide incorporated in the expression constructs was processed by *L. tarentolae*. A similar successful secretion of truncated human TPA protein with its native signal sequence in this expression system was reported by Nazari and Davoudi (2011). ICE and ICC assays revealed that the produced recombinant antibody specifically binds to its endogenous target (i.e., the

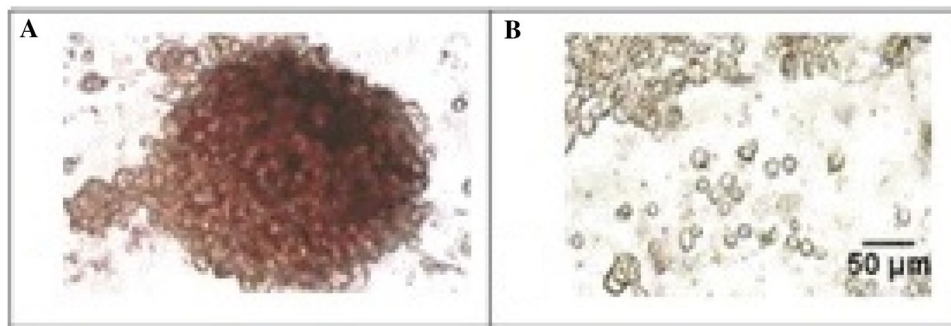


Fig. 5 ICC assay with PHA-activated lymphocytes (a) and non-activated lymphocytes as a control (b), using the expressed recombinant anti-IL-2R α antibody. The recombinant anti-IL-2R α antibody was bound to the IL-2R α expressed on T lymphocytes activated by PHA,

and identified by a secondary antibody in a colorimetric reaction. The control group, which did not express IL-2R α , was not recognized by the anti-IL-2R α antibody and no colorimetric reaction was observed

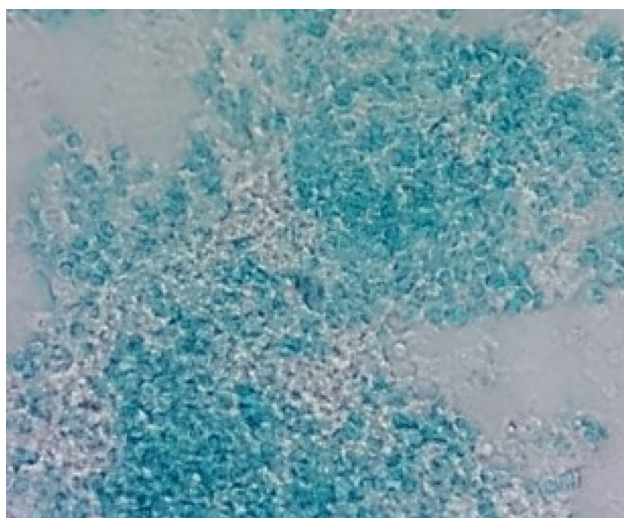


Fig. 6 The results of the BrdU assay. The mean reading for non-activated lymphocytes (control 1) was 0.505 ± 0.1 . The mean readings for PHA-activated lymphocytes that were either treated with the expressed recombinant anti-IL-2R α antibody or untreated with the antibody (control 2) were 0.594 ± 0.07 and 2.153 ± 0.19 , respectively. The difference in the rates of cellular proliferation between the experimental group and control group 1 was not significant ($p=0.29$; two-sided unpaired Student's *t*-test). The proliferation rate of control group 2 was significantly greater than that of the experimental group ($p=0.0002$). The data are presented as the mean \pm SD ($n=3$)

alpha chain of IL-2 receptor). Glycosylation of the heavy chain, an important eukaryotic post-translational modification, was also verified. The average dose of the expressed anti-IL-2R α antibody required for the inhibition of IL-2-dependent lymphocyte proliferation was 5–7 mg/ml, which is comparable with the required amount of commercial daclizumab (5–10 mg/ml). The recombinant protein yield in the *L. tarentolae* (Lai et al. 2019) expression system was 2 mg/l after 48 h culture in LBR medium, which was comparable with similar studies that reported 0.1 to 6 mg of secreted

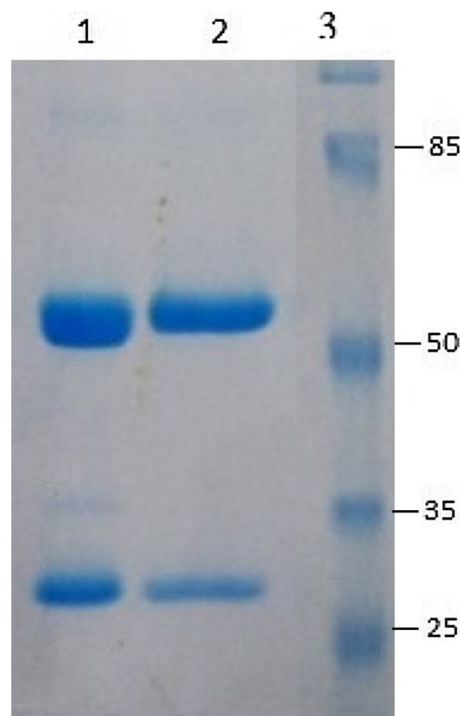


Fig. 7 Enzymatic deglycosylation of the heavy chain of the recombinant anti-IL-2R α antibody under reducing conditions. N-glycosylation of the expressed recombinant antibody was evaluated by PNGase F treatment. Electrophoretic mobility of the purified heavy chain of the recombinant antibody treated with the enzyme (lane 1) and without enzymatic treatment (lane 2) is shown. Lane 3, protein size marker (kDa)

protein per ml (Basile and Peticca 2009). The expression levels of mAbs in mammalian cell systems were initially on the order of 100–500 mg/l (Jones et al. 2007). However, recent advances in mammalian cell line development, and media and bioreactor optimizations reportedly resulted in up to 10 g protein/l yields and cell densities of over 20 million cells/ml in fed-batch processes (Hunter et al 2018). To date,

no genetic manipulations of *L. tarentolae* LEXSY host P10 to adapt this expression system to large-scale production of recombinant proteins have been reported. In summary, we expressed, purified, and evaluated human anti-IL-2R α antibody in *L. tarentolae*. *L. This protozoan* can thus be used as a host for the production of therapeutic monoclonal antibodies. The low-cost growth medium and equipment, easy handling, low risk of contamination by viruses or prions, and mammalian-like posttranslational modifications constitute main advantages of this expression system.

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Compliance with Ethical Standards

Conflict of interest There are no conflict of interests for authors.

Ethics Approval The ethics process was approved by Shahid Beheshti University of Medical Sciences Ethical Committee with IR.SBMU.REC.1394.99 number.

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