# CSF-1/CSF1R SYSTEM AS PREDICTOR OF LIVE BIRTH AFTER INDUCED PREGNANCY

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#### СИСТЕМА CSF-1/CSF1R КАК ПРЕДИКТОР живорождения ПОСЛЕ ИНДУЦИРОВАННОЙ БЕРЕМЕННОСТИ

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#### Abctract

The effectiveness of the IVF procedure is assessed not only by the indicators of the onset of clinical pregnancy, but also by the proportion of live births, since about a third of pregnancies end in termination, which is certainly determined by many factors. Colony-stimulating factors can be considered potential predictive markers of the induced pregnancy development and resulting live birth. To evaluate the predictive value of the CSF-1 system (serum CSF-1 concentration and carriage of CSF1R gene polymorphisms) in the setting of live birth after IVF-induced pregnancy in women with tuboperitoneal infertility. 88 patients undergoing IVF aged between 25 and 40 years were assigned to the following groups: Group I (n =32): patients whose post-IVF pregnancies resulted in live births at Week 24 or later; and Group II (n = 52): patients whose pregnancies did not occur or resulted in spontaneous miscarriages. ELISA determined the CSF-1 levels twice: on pre-IVF menstrual cycle Days 3-4, and on post-embryo-transfer Day 15. Genotyping according to the CSF1R gene rs3216780 and rs38669350 polymorphic markers used a polymerase chain reaction with Sanger sequencing. CSF-1 levels grew to postembryo-transfer Day 15 (3.9-fold in Group I and 1.8-fold in Group II); in the Group II pregnant patients, they were reliably higher than in the comparison group (p =0.0017). Most women whose pregnancy resulted in live births carried the del/G rs3216780 and TG/CA rs386693509 genotypes of the CSF1R gene. Univariate analysis identified the following predictors for the completion of induced pregnancy by delivery - peripheral blood CSF-1 levels from 121.3 to 314.8 pg/mL at the preconceptional stage and from 963.3 to 1682.8 pg/mL - on day 15 after embryo transfer, as well as carriage of del/G rs3216780 genotypes and TG/CA rs386693509 gene CSF1R.V combination, these predictors were observed in 66.7% of women from group I and in 7.7% from group II. The CSF-1 system plays an important role in the realization of a woman's reproductive function and the identified prognostic indicators - the pre-conception and post-ET Day 15 serum CSF-1 concentrations and the carriage of the CSF1R gene del/G rs3216780 and TG/CA rs386693509 genotypes can be used as predictors of the completion of induced pregnancy by childbirth.

**Keywords:** in vitro fertilization, live birth, induced pregnancy, cytokines, colony-stimulating factor-1, CSF1, CSF1R polymorphism, rs3216780, rs386693509

#### Резюме

Эффективность процедуры ЭКО оценивается не только показателями наступления клинической беременности, но и долей живорождений, поскольку около трети беременностей завершаются прерыванием, что безусловно, определяется многими факторами. Колониестимулирующие факторы можно рассматривать как потенциальные прогностические маркеры развития индуцированной беременности и последующего живорождения. Цель исследования - оценить прогностическую значимость системы CSF-1 (сывороточной концентрации CSF-1 и носительство полиморфизмов гена *CSF1R*) в рамках завершения ЭКО-индуцированной беременности родами у женщин с трубно-перитонеальным бесплодием. 88 пациенток, проходящих процедуру ЭКО, в возрасте от 25 до 40 лет были разделены на группы: І группа (n=32) - пациентки, беременность после процедуры ЭКО у которых завершилась родами в срок от 24 недели и более, II группа (n=52) - пациентки, у которых беременность не наступила или закончилась самопроизвольным иммуноферментного двукратно Методом выкидышем. анализа было определено содержание CSF-1: на 3-4 день менструального цикла, предшествующего процедуре ЭКО, и на 15 день после переноса эмбрионов. Генотипирование по полиморфным маркерам rs3216780 и rs386693509 гена было проведено методом полимеразной цепной реакции с CSF1R секвенированием по Сенгеру. Содержание CSF-1 увеличилось на 15 день после переноса эмбриона (в 3,9 раза для I группы и в 1,8 раза для II группы) и у пациенток, беременность из группы 2, содержание CSF-1 было достоверно выше, чем в группе сравнения (р=0,0017). Наибольший процент женщин, беременность которых закончилась родами, были носителями генотипа del/G rs3216780 и TG/CA rs386693509 гена CSF1R. При однофакторном анализе были выявлены следующие предикторы завершения индуцированной беременности родами - уровни CSF-1 в периферической крови от 121,3 до 314,8 пг/мл на преконцептивном этапе и от 963,3 до 1682,8 пг/мл - на 15 день после переноса эмбрионов, а также носительство генотипов del/G rs3216780 и TG/CA rs386693509 гена CSF1R. В сочетании данные предикторы отмечались у 66,7% женщин из I группы и у 7,7% из II группы. Система CSF-1 играет важную роль в реализации репродуктивной функции женщины и выявленные прогностические показатели - сывороточная концентрация CSF-1 преконцептивном этапе и на 15 день после переноса эмбрионов, а также носительство генотипов del/G rs3216780 и TG/CA rs386693509 гена CSF1R, использованы могут быть качестве предикторов завершения В индуцированной беременности родами.

Ключевые слова: экстракорпоральное оплодотворение, роды, индуцированная беременность, цитокины, колониестимулирующий фактор-1, макрофагальный колониестимулирующий фактор, CSF1, полиморфизм CSF1R, rs3216780, rs386693509.

## 1 1 Introduction

In vitro fertilization (IVF) is one of the most popular infertility treatment 2 method. However, when assessing its effectiveness, on average, about a third of the 3 protocols lead to pregnancy, and of these, about 70% result in live births [2]. This 4 problem stimulates researchers to identify new informative factors influencing the 5 success of IVF. As the factors of the immune system are directly involved in the 6 realization of reproductive functions, its components can be considered potential 7 predictive markers of the induced pregnancy development and resulting live birth 8 [1, 11]. 9

Colony-stimulating factors form a family of cytokines that regulate the 10 differentiation and proliferation of hematopoietic progenitor cells. Thus, colony-11 stimulating factor-1 (CSF-1) regulates the proliferation, differentiation, and survival 12 of macrophages; supports the growth and proliferation of extravillous trophoblastic 13 cells; plays an important role in embryo implantation in the uterus and subsequent 14 placental growth [3, 12, 14]. A reduced number of cells or poor development of 15 trophoblast can induce alterations in the placental structure and nutrient transport 16 function-these lead to pregnancy loss or disturbed development [3]. In addition, 17 CSF-1 is an embryotrophic factor: it favors the development of the blastocyst by 18 increasing the number of blast cells and gene expression [4]. The CSF-1's biologic 19 effects are implemented via its binding with the specific CSF1R receptor; this leads 20 to its dimerization, activation of the tyrosine kinase and signaling cascade [5, 8, 12, 21 13]. Trophoblast expresses CSF and CSF1R more than any other non-macrophage-22 derived cell types do. The placental development depends upon the CSF signaling, 23 as CSF stimulates the proliferation of the placental cells, the differentiation of the 24 cytotrophoblastic cells, and regulates the pre-implantation division of the 25 blastocystic cells. The total role of CSF in the development of the induced pregnancy 26 is under study, yet [3]. 27

The objective of this study is to evaluate the predictive value of the CSF-1 system (serum CSF-1 concentration and carriage of *CSF1R* gene polymorphisms) components in the setting of live birth after IVF-induced pregnancy in women with tuboperitoneal infertility.

32 2 Material and Methods

#### 33 Ethical considerations

The Institutional Review Board at a large public university approved the research. All procedures performed in studies involving human participants were in accordance with the ethical standards of the institutional and/or national research committee and with the 1964 Helsinki Declaration and its later amendments or comparable ethical standards. Written informed consent was obtained from all individual participants involved in the study. The study was approved by the local ethics committee of Federal State Budgetary Educational Institution of Higher CSF-1/CSF1R AND IVF CSF-1/CSF1R И ЭКО

10.46235/1028-7221-17199-CCS

41 Education "National Research Ogarev Mordovia State University" protocol No. 68

42 of 31.03.2015.

## 43 Study population

Based on the inclusion, non-inclusion, and exclusion criteria, this study 44 involved 88 women with tuboperitoneal infertility aged 25 to 35 years, with an 45 average age of  $28.7 \pm 2.9$  years, who were treated from June 2022 to December 2023 46 at the Department of Assisted Reproductive Technologies of the Perinatal Center of 47 the State Budgetary Healthcare Institution of the Republic of Mordovia "Mordovian 48 Republican Central Clinical Hospital" (Saransk). Criteria for inclusion were: signed 49 informed consent for enrollment; current tuboperitoneal infertility factor; normal 50 ovarian reserve; regular menstrual cycle; normal karyotype of both spouses; fertile 51 semen in partner/husband; and transfer of only-good-quality embryos. Criteria for 52 non-inclusion were: patients with other forms of infertility, acute genital or 53 extragenital diseases, as well as exacerbated chronic diseases. Criteria for exclusion 54 were: voluntary withdrawal of patient from study at any stage. 55

All the patients received the "short-protocol" superovulation stimulation, 56 which involved using the recombinant follicle-stimulating hormone and 57 gonadotropin-releasing hormone antagonists. All the patients had only-good-quality 58 fresh embryos transferred into the uterine cavity under ultrasonic control on Days 59 5–6 of cultivation. Diagnosis of pregnancy was based on determining the serum 60 concentration of the human chorionic gonadotropin β-subunit on post-embryo-61 transfer (ET) into uterine cavity Day 15 and ultrasounding on Day 21 to assess for 62 the presence of one or two fetal eggs in the uterine cavity. Live birth was defined as 63 the child born alive after 24 weeks of pregnancy and having survived more than 64 1 month. The facts of live birth were to be confirmed by personal observation and 65 data from the clinical records of the postpartum women. Miscarriage was defined as 66 the loss of pregnancy before 28 weeks. 67

<sup>68</sup> Depending on the outcomes the patients were assigned to two groups. Group I <sup>69</sup> consisted of the 36 women whose pregnancies resulted in live births, mean age 28.6 <sup>70</sup>  $\pm$  3.1 years. Group II consisted of the 52 patients whose pregnancies did not occur <sup>71</sup> or resulted in spontaneous miscarriages, mean age 29.3  $\pm$  2.7 years. The study groups <sup>72</sup> were comparable in terms of age, reproductive function and past gynecological <sup>73</sup> diseases (p > 0.05).

## 74 Serological testing

To study the serum concentration of cytokines, venous blood was collected aseptically by venipuncture from the cubital vein in the morning on an empty stomach in a volume of 10 ml, obtained using a VACUETTE® vacuum system (GreinerBio-One, Austria) into tubes with a coagulation activator. The blood was centrifuged for 15 minutes at 2000 g, the supernatant liquid was collected into Eppendorf tubes and frozen at a temperature of - 30° C. Storage at a temperature of CSF-1/CSF1R AND IVF CSF-1/CSF1R И ЭКО

10.46235/1028-7221-17199-CCS

-40-60° C in the refrigerator for up to 6 months. The samples were not thawed and
re-frozen.

Blood from the peripheral vein was sampled twice during the IVF procedure. The first sampling was performed on pre-IVF menstrual cycle Days 3–4 (preconception stage), while the second sample was taken on post-embryo-transfer (ET) into uterine cavity Day 15. Serum CSF-1 concentration was determined by ELISA using kits from *R&D Systems* (USA) with a Personal Lab (*Adaltis*, Italy) immunoassay analyzer.

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## Sequence-based genotyping

The material for the genetic study was venous blood in a volume of 5 ml, obtained using a VACUETTE® vacuum system (GreinerBio-One, Austria) into tubes with EDTA (ethylenediaminetetraacetate). Storage of whole blood - 80°C, transportation at -20°C.

Genomic DNA was isolated from whole blood with the help of a QIAamp 94 DNA Mini Kit at a QIAcube automated nucleic acid isolation station (QIAGEN, 95 Germany). The rs3216780 and rs386693509 dinucleotide polymorphisms in the 96 CSF1R gene UTR-3 region were examined with Sanger sequencing. The primers for 97 the CSF1R gene amplification were selected using the Primer-BLAST Internet 98 resource (http://www.ncbi.nlm.nih.gov/tools/primer-blast) and the NG\_012303.1 99 nucleotide from reference sequence the **NCBI** Database 100 (http://www.ncbi.nlm.nih.gov/nuccore/NG\_012303.1). The sequencing of the 101 purified PCR products used an ABI PRISM 3500 analyzer (Thermo Fisher 102 Scientific, USA) with a Big Dye<sup>TM</sup> Terminator v3.1 Cycle Sequencing Kit (Applied 103 Biosystems, USA). The nucleotide sequences were analyzed using the Sequence 104 Scanner v.1.0. Peak Trace, Vector NTI Advance 10, and Chromas Lite 2.1.1 105 software packages. 106

## 107 Statistical analysis

The statistical analysis of the findings used the Stat Soft Statistica 12.0 (USA) 108 application software package. The distribution of quantitative data corresponded to 109 the normal distribution, and therefore were expressed as the arithmetic mean with 110 the standard deviation, and the Student's t-test was used to analyze the differences 111 between groups. The qualitative values are presented as the absolute values (n) and 112 percentages. To assess the associations of the factors with the procedure outcome, 113 the Pearson's  $\gamma^2$  test and the odds ratio (OR) with a 95% confidence interval (CI) 114 were used. The significance level of the revealed differences and correlations in all 115 the types of analysis was assumed to be p < 0.05. The distribution of the allele 116 frequencies and genotypes of polymorphic variants complied with the Hardy-117 Weinberg equilibrium. 118

#### 119 3 **Results**

The pre-conception CSF-1 levels did not significantly differ between the groups (p = 0.730); nevertheless, the over-time CSF-1 assessment found that on post-ET Day 15 women in both groups showed a reliably higher CSF-1 (3.9-fold in Group I (p < 0.0001) and 1.8-fold in Group II (p = 0.012), while the Group I patients had a reliably higher CSF-1 than those of the comparison group (p < 0.0001).

To assess the possible effects of the serum CSF-1 level on the post-IVF live births, an interquartile analysis was performed. Table 1 shows the isolated CSF-1 quartiles (Me (Q 0.25–Q 0.75). The CSF-1 concentrations corresponding to the 1st and 2nd pre-conception quartiles and the 3rd and 4th post-ET Day 15 quartiles associated with the maximal numbers of the post-IVF live births.

To evaluate the predictive value of the CSF-1 concentrations, the odds ratio 130 with a 95% confidence interval was calculated, and the sensitivities and specificities 131 of the identified predictors of live births were determined. Patients with 121.3 to 132 314.8 pg/mL pre-conception peripheral blood CSF-1 had 6.18-fold higher odds of 133 live birth at induced pregnancy (95% CI = 2.386-15.990, p < 0.001, sensitivity: 134 75%, specificity: 67.3%). Post-ET Day 15 CSF-1 963.3 to 1682.8 pg/mL supposes 135 71.4-fold higher odds of post-IVF live birth (95% CI = 14.646-348.073, p < 0.05, 136 Complies with Fisher's exact test, sensitivity: 94.4%, specificity: 80.8%). 137

In the setting of the analysis of the gene CSF1R rs3216780 (del/G) and rs386693509 (TG/CA) polymorphisms, the distribution of genotypes was assessed for compliance with the Hardy-Weinberg equilibrium—no abnormalities were detected. Taking into account the findings, the multiplicative and general inheritance models were applied to these polymorphisms to identify the post-IVF live birth predictor.

The analysis of the *CSF1R* gene polymorphism alleles and genotypes distribution showed that 53.4% of the examined women carried the del allele, while 46.6% carried the *G* allele. The *G* allele and the *del/G* genotype were predominant in the favorable outcome group of women. Therefore, the carriage of the *CSF1R* gene *del/G* rs3216780 genotype acts as a predictor of live birth after induced pregnancy (OR = 11.18, 95% CI = 4.09–30.58, sensitivity: 75%, specificity: 78.8%) (Table 2).

The analysis of the *CSF1R* gene rs386693509 polymorphism distribution showed that 73.3% carried the *TG* allele, while 26.7% carried the *CA* allele. The *TG* allele was predominant in both groups. In Group I, the *TG/CA* was predominant (75%), versus 15.4% in Group II women. Therefore, the carriage of the *CSF1R* gene *TG/CA* rs386693509 genotype acts as a predictor of post-IVF live births (OR = 16.5, 95% CI = 5.68–47.92, sensitivity: 75%, specificity: 84.6%) (Table 2).

Therefore, the univariate analysis has identified the following predictors of the live births at induced pregnancy: 121.3 to 314.8 pg/mL pre-conception CSF-1 in

**Russian Journal of Immunology (Russia)** 

peripheral blood and 963.3 to 1682.8 pg/mL post-ET Day 15 CSF-1, as well as the
carriage of the *CSF1R* gene *del/G* rs3216780 and *TG/CA* genotypes. Combinations
of these predictors were reported in 66.7% of Group I and 7.7% of Group II women.

## 162 4 **Discussion**

Evaluating the IVF procedure's effectiveness implies not only the 163 development of pregnancy but also its live-birth outcome, as about a third of 164 pregnancies end in losses; these are, certainly, determined by multiple factors. 165 Embryo implantation and subsequent placentation depend on the consistent 166 embryo/endometrium interaction regulated by multiple neurohumoral factors. CSF1 167 is a major regulator of the macrophage-phagocytic system development and 168 homeostasis and, accordingly, a key factor in the control of trophoblastic cells 169 development and homeostasis [3, 12]. Earlier studies in women with tuboperitoneal 170 infertility revealed a relationship between the CSF-1 concentration and the post-IVF 171 pregnancy onset [6]. Literature sources show, however, only not numerous studies 172 that have assessed the association between the CSF-1 system and post-IVF 173 pregnancies resulting in live births. 174

This is consistent with the findings of other study showing an association between the grown CSF-1 levels and the higher probability of pregnancy [5]. Patients with recurrent miscarriage are also known to have inhibited endometrial CSF-1 production [9]. Hence, the CSF-1 expression can be a potential factor improving the IVF outcomes.

The mechanism by which CSF-1 influences the rates of implantation and 180 placentation is not understood so far. It is known, however, that adding CSF-1 to a 181 culture of human trophoblastic cells leads to their differentiation into 182 syncytiotrophoblastic cells and stimulates the production of the placental lactogen. 183 Moreover, adding CSF-1 to a culture of murine blastocystic cells causes the growth 184 of the trophoblast [7]. It is worth noting that in experimental work using 185 osteopetrotic mutant mice that lacked CSF-1, it was shown that the fetus can 186 synthesize its own CSF-1, compensating for its lack in the mother [10]. 187

We believe that the identified predictive value of the CSF-1 system can be due to the *CSF1R* gene polymorphism, which ensures its sensitivity and the implementation of the cytokine's biologic functions. An additional explanation is the longer CSF1R expression on the cell surface at the *del/G* and *TG/CA* variants this favors the stimulation of trophoblastic cell growth, angioprotective effect, and faster placental vascularization.

## 194 5 Conclusions

The results of our study allow for the conclusion that CSF-1 plays a major role in the implementation of the female reproductive function. The pre-conception and post-ET Day 15 serum CSF-1 concentrations and the carriage of the *CSF1R* gene del/G rs3216780 and *TG/CA* rs386693509 genotypes are of predictive value in the setting of the post-IVF live births. The findings are of interest from the point of view of completing the data on the contributions of both cytokines and gene polymorphisms in the induced pregnancy development. A more accurate assessment requires further research and analysis of the other factors that may have an effect on the expression of the genes involved in the embryo/endometrium interaction and on the live births.

# таблицы

| Factor                    | Pre-conception CSF-1, pg/mL |            |              |              |
|---------------------------|-----------------------------|------------|--------------|--------------|
|                           | 1st quartile                | 2nd        | 3rd quartile | 4th quartile |
|                           | 121.3–                      | quartile   | 314.9–       | 422.2–725.7  |
|                           | 257.5                       | 257.6-     | 422.1        | (n = 22)     |
|                           | (n = 22)                    | 314.8      | (n = 22)     | ()           |
|                           |                             | (n = 22)   |              |              |
| Live births, n (%)        | 11 (50%)                    | 16 (72.7%) | 5 (22.7%)    | 4 (18.2%)    |
| Spontaneous miscarriages, | 11 (50%)                    | 6 (27.3%)  | 17 (77.3%)   | 18 (81.8%)   |
| n (%)                     |                             |            |              |              |
|                           | Post-ET Day 15 CSF-1, pg/mL |            |              |              |
|                           | 1st quartile                | 2nd        | 3rd quartile | 4th quartile |
|                           | 205.9-                      | quartile   | 963.3-       | 1251.4-      |
|                           | 691.2                       | 691.3-     | 1251.4       | 1682.8       |
|                           | (n = 22)                    | 963.2      | (n = 22)     | (n = 22)     |
|                           |                             | (n = 22)   |              |              |
| Live births, n (%)        | _                           | 2 (9%)     | 14 (63.6%)   | 20 (91%)     |
| Spontaneous miscarriages, | 22 (100%)                   | 20 (91%)   | 8 (36.4%)    | 2 (9%)       |
| n (%)                     |                             |            |              |              |

**Table 2.** Multiplicative general models of inheritance of CSF1R gene rs3216780 (*del/G*) and rs386693509 polymorphisms.

| CSF1R ge ne | Allele/genoty | Group<br>I | Group<br>II | <b>x</b> <sup>2</sup> | р      | (         | DR        |
|-------------|---------------|------------|-------------|-----------------------|--------|-----------|-----------|
| пе          | ре            | (n = 36)   | (n = 52)    |                       |        | Valu<br>e | 95%<br>CI |
|             |               | )          |             |                       |        |           |           |
| rs3216780   | G             | 0.597      | 0.375       | 8.44                  | 0.004  | 2.47      | 1.33-     |
| (del/G)     |               |            |             |                       |        |           | 4.58      |
|             | del           | 0.403      | 0.625       |                       |        | 0.40      | 0.22-     |
|             |               |            |             |                       |        |           | 0.75      |
|             | del/del       | 0.028      | 0.519       | _                     | < 0.05 | 0.03      | 0.00-     |
|             |               |            |             |                       | *      |           | 0.21      |
|             | del/G         | 0.750      | 0.212       | 25.13                 | < 0.00 | 11.1      | 4.09–     |
|             |               |            |             | 9                     | 1      | 8         | 30.58     |
|             | G/G           | 0.222      | 0.269       | 0.251                 | 0.617  | 0.78      | 0.29–     |
|             |               |            |             |                       |        |           | 2.10      |
| rs3866935   | TG            | 0.597      | 0.827       | 11.47                 | 0.000  | 0.31      | 0.16–     |
| 09          |               |            |             |                       | 7      |           | 0.62      |
| (TG/CA)     | CA            | 0.403      | 0.173       |                       |        | 3.22      | 1.61–     |
|             |               |            |             |                       |        |           | 6.44      |
|             | TG/TG         | 0.222      | 0.750       | 23.81                 | < 0.00 | 0.10      | 0.03-     |
|             |               |            |             |                       | 1      |           | 0.26      |
|             | TG/CA         | 0.750      | 0.154       |                       | < 0.00 | 16.5      | 5.68-     |
|             |               |            |             | 2                     | 1      | 0         | 47.92     |
|             | CA/CA         | 0.028      | 0.096       | _                     | > 0.05 | 0.27      | 0.03–     |
|             |               |            |             |                       | *      |           | 2.40      |

\*Complies with Fisher's exact test

## ТИТУЛЬНЫЙ ЛИСТ\_МЕТАДАННЫЕ

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#### Блок 3. Метаданные статьи

CSF-1/CSF1R SYSTEM AS PREDICTOR OF LIVE BIRTH AFTER INDUCED PREGNANCY СИСТЕМА CSF-1/CSF1R КАК ПРЕДИКТОР ЖИВОРОЖДЕНИЯ ПОСЛЕ ИНДУЦИРОВАННОЙ БЕРЕМЕННОСТИ

#### Сокращенное название статьи для верхнего колонтитула:

CSF-1/CSF1R AND IVF CSF-1/CSF1R И ЭКО

Ключевые слова: экстракорпоральное оплодотворение, роды, индуцированная беременность, цитокины, колониестимулирующий фактор-1, макрофагальный колониестимулирующий фактор, CSF1, полиморфизм CSF1R, rs3216780, rs386693509.

**Keywords:** in vitro fertilization, live birth, induced pregnancy, cytokines, colonystimulating factor-1, CSF1, CSF1R polymorphism, rs3216780, rs386693509.

Иммунологические чтения в Челябинске 2025. Количество страниц текста – 7, Количество таблиц – 2, Количество рисунков – 0. 30.03.2025

## СПИСОК ЛИТЕРАТУРЫ

| Порядковый<br>номер<br>ссылки | Авторы, название<br>публикации и источника, где<br>она опубликована, выходные<br>данные  | публикации и источника   | Полный интернет-адрес (URL)<br>цитируемой статьи или ее doi |
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