

Power system for schoolchildren based on their genotypes

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ABSTRACT

The article discusses a decision support system based on the clustering of genotypes and used in managing personalized nutrition for schoolchildren. The results of solving the problem for specific objects that have passed laboratory studies of gene states are presented. Modern scientific research in the field of nutrigenomics has shown that the selection of the optimal nutrition plan, taking into account genetic characteristics, increases the effectiveness of such programs by 200-300% compared with traditional approaches. This report discusses the use of soft computing (the method of clustering multidimensional objects) to determine typical clusters for managing personalized customer nutrition based on their genotypes.

Keywords: Genotype; Personified nutrition; Clustering of multidimensional objects; Metrics of characteristics; Decision support system.

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INTRODUCTION

The most important period of a person's life is school age, the time is from 7 to 17 years, the time of physical, intellectual, moral formation and active development.

In the modern world, it is the students who take over and are forced to process the increasing pressure of the information flow that affects them not only at school, but also at home. In addition to the school curriculum, many children and adolescents are additionally engaged in circles and sports sections.

For the formation and maintenance of physical and mental health and the full assimilation of the school curriculum, it is important to correctly organize the nutrition of the student (Bekmansurov et al., 2019; Panfilova et al., 2020; Ivanova, et al., 2019; Voronkova et al., 2019; Chen et al., 2019; Frolova, et al., 2019).

Many parents believe that it is enough to rely on their own intuition and common sense in the matter of feeding a student. However, it is important to know and understand the principles of good nutrition, and the rules of food hygiene, the observance of which is key in maintaining the health of the child.

Eating disorders during this period can lead not only to functional disorders and chronic diseases of the gastrointestinal tract, but also to deviations in the functioning of almost all body systems.

Basic power features

The first function is to supply the body with energy. In this sense, a person can be compared with any machine that does work but requires fuel for this. Rational nutrition provides an approximate balance of energy entering the body and spent on supporting vital processes.

The second function of nutrition is to supply the body, primarily proteins, to a lesser extent - minerals, fats and to an even lesser extent – carbohydrates (Podoprigora et al., 2019; Movchan and Yakovleva, 2019).

Finally, the third function of nutrition is to supply the body with biologically active substances necessary for the regulation of vital processes. Enzymes and most hormones - the regulators of chemical processes in the body, are synthesized by the body itself. However, some hormones the human body can synthesize only from special precursors found in food. These precursors are vitamins found in foods (Poltarykhin, 2020; Almeida et al., 2019; Prodanova et al., 2019; Melnikov, 2019; Aleksandrova et al., 2019; Filatova et al., 2019; Akhmadeev et al., 2019; Dunets et al., 2019).

More recently, there has been evidence of the existence of another (fourth) nutrition function, which is to develop immunity. With insufficient nutrition, overall immunity decreases, and the body's resistance to various infections decreases. And vice versa, good nutrition with a sufficient content of proteins, fats, vitamins and calories enhances immunity, and increases resistance to infections.

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METHODS

Currently relevant are the directions in the development of specialized food products, as well as personalized food products and diets (based on the allelic polymorphism of individual “predisposition genes” associated with the absorption of certain food nutrients), aimed at improving the nutritional structure of the population. The indicated innovative directions for the development of science in the field of functional and personalized nutrition make it possible to solve the tasks set in the strategic program documents of the Government of the Russian Federation: “*Fundamentals of the state policy in the field of healthy nutrition for the period up to 2020*” and “*The concept of ensuring sanitary and epidemiological welfare of the population through development functional and specialized bakery in the Russian Federation until 2020 (Bread is health)*.” In the Order of the Government of the Russian Federation No. 1364-r dated 06/29/2016 “*Strategy for improving the quality of food products in the Russian Federation until 2030*” (www.government.ru) indicates the need to create conditions for the production of new-generation food products with specified quality characteristics, including specialized, functional and enriched, organic foods (Decree of the Government of the Russian Federation).

The social essence of personalized nutrition consists in the development of a modern industry and the logistics of providing the population with food, taking into account individual and group needs. Creating a flexible food quality management system, taking into account the individual characteristics of each person, will create the prerequisites for improving labour efficiency, quality of life, and increasing life expectancy.

The theoretical support of the scientific direction “*Personalized Nutrition*” consists in developing a theory of the informational influence of food quality on the quality of human life processes. One of the possible tools for this effect is nutrigenomics - the science of the influence of diets on the expression of human genes. In this area, it is necessary to develop a theory for controlling the expression of favourable and unfavourable genes through nutrients (Aleksandrovich et al., 2018). A theory is also needed that describes a possible correlation of the consumer’s emotional preferences and the objective needs of his body (Poltarykhin, Dibrova, 2020; Dzhavatov et al., 2018; Jafarpour et al., 2019; Ivanova et al., 2019; Bondarenko et al., 2019; Masood et al., 2019).

The methodology for the development of technical specifications for the design of an innovative consumer facility of personalized food consists in taking into account 3 groups of characteristics of the designed facility. Identification of psycho-emotional preferences of consumers is carried out by sequential refinement and detailing during the preferences surveys based on the PATTERN heuristic examination method. The method involves detailing “*from general to particular*”. In this case, descriptors of a higher level are split into descriptors of a lower level. Detailing is carried out to obtain the final (indivisible) descriptors. The result of such a heuristic examination is a “*target tree*”, including descriptors and their significance levels. Depending on the needs, using such a tree, various trajectories can be calculated: elements (nodes), branches (lines) and families.

Identification of the objective needs of the body according to the results of the analysis of the genome is based on a special technique. According to this technique, the genome is analysed for the presence of unfavourable gene alleles in it. If such panels are found, nutrigenomic factors are included in the project to prevent or compensate for possible nutrient deficiencies. The result of gene research should be a list of descriptors that have an inhibitory or stimulating effect on the expression of borderline gene alleles.

Despite the wide range of possible competencies of target food products, their number is still assumed to be final. A number of technologies should be developed that allow implementing a project for the production of food products that meet the solution of standard tasks. Such products should provide the quantitative and qualitative need of the individual organism for nutrients.

Moreover, the same competencies can be assigned to food products of different food commodity groups. For example, antioxidant activity can be provided with fish, dairy, confectionery products, as well as extreme nutrition products. For a qualitative and quantitative assessment of the moisture contained in food products, various methods for its instrumental assessment have been developed and applied in practice.

One of the types of target nutrition is food with a prolonged shelf life. Caramel with a significantly reduced ability to get wet can act as such a product. To increase the shelf life of caramel, it is proposed to introduce surfactants into the caramel mass formulation. Due to the property of reducing surface tension at the interface, the surfactant in the process of glass transition of caramel mass is placed on the periphery of the product, thereby reducing its hygroscopicity.

Assessment of merchandising characteristics of bioorganic objects by nuclear magnetic resonance.

Moisture is a critical feature in the design of food products with specified consumer properties. It is often impossible to differentiate between the three states of water identified during the study (equilibrium humidity, humidity above and below equilibrium). Incorrect identification of the moisture state leads to the loss of food competencies planned during the design process.

Formulation of the problem:

Given: C_0 – the original set of multidimensional objects,

$C_0 = \{S_n\}$, $n = 1, \dots, N$

$M_p(M)$ – performance metric, $M_p(i)$ – weight coefficient i specifications, $i = 1, \dots, M$

$X(n, i)$ – i state characteristic i facility n , $n=1, \dots, N$, $i=1, \dots, M$

Table 1 presents the results of the analysis of the state of genes for one person. Column 4 indicates the status of the gene named in column 1. For digital processing, points 0,1,2 is put in column 4. When scoring 0, 1, 2, it is assumed that 1 and 2 are the average and high risk of predisposition to the disease or condition, 0 means no risk to the disease or condition. In addition, fourteen genes were selected by experts to cluster the states. The metric $M_p(M)$ is normalized (see formula 1).

$$\sum_{i=1}^N M_p(i) = 1 \quad (1)$$

Source set C_0 need to be divided into multiple clusters C_k (see formulas 2 and 3):

$$C_0 = \{C_k\} \quad k=1, \dots, K \quad (2)$$

$$C_k = \{S_z\}, \quad z=1, \dots, N_k \quad (3)$$

Any pair of clusters does not have common elements, i.e., any object can only be in one cluster (see formula 4);

$$\forall C \in C \forall C_1 \in C_0: C \cap C_1 = \emptyset \quad (4)$$

where C_k is the set of objects of the k th class; k is the class number;

It is required to determine those C_k that maximize the U criterion (see formula 5):

$$U(K) = \max\{U_1(K) - U_2(K)\} \quad \text{for } K = N, N-1, \dots, 2 \quad (5)$$

where $U(K_0)$ – optimal value of the clustering quality criterion;

$U_1(K)$ – class compactness with K clusters;

$U_2(K)$ – measure of class proximity with K clusters.

The measure of similarity between two objects is determined based on a potential function $f(S_i, S_j)$:

$$f(S_i, S_j) = 1 / (1 + D^2(S_i, S_j));$$

$$D(S_i, S_j) = \sqrt{\sum_{m=1}^N (X_{i,m} - X_{j,m})^2}$$

$$U_1(K) = \frac{1}{K} \sum_{k=1}^K 2 / (k * (N_k - 1)) * \sum_{S_i \in C_k} \sum_{S_j \in C_k, i \neq j} f(S_i, S_j)$$

where K is the number of classes at the current classification step; C_k – k feature class; N_k – the number of objects in the class C_k ; $f(S_i, S_j)$ – potential function of two objects S_i and S_j ; $D(S_i, S_j)$ - distance between objects S_i and S_j in the space of characteristics X , taking into account the metric.

$$U_2(K) = 2 / (K(K-1)) * \sum_{k=1}^K N_k$$

where C_p – many clusters obtained as a result of solving the clustering problem,

$$F(C_k, C_l) = 1 / (N_k * N_l)$$

Table 1. Input information for each object for processing.

Gene number	Genes	Polymorphism / mutation	wt/wt	Genotype
1	CYP1A1	wt, 2A, 2B, *4: 4887C>A, 4889A>G, 6235T>C)	0	wt/wt
2	CYP1A2	intron 1 -164 C>A (*1A>*1F)	2	*1A/*1A
3	GSTT1	wt, del	0	+
4	GSTM1	wt, del	0	+
5	NAT2	*4(wt) (T341C), 481C>T *5, 590G>A *6, 857G>A *7 S1, S2, S3 (*5, *6, *7) – slow acetylators	2	S1/S2
6	MTHFR	677 C>T	0	C/C

7	ADRB2	48A>G, 81C>G	2	G/G, C/G
8	SR(HTR2A)	102 T>C	1	T/C
9	PPARD	294 T>C	0	T/T
10	NBPF3(ALPL)	rs4654748	2	C/C
11	FUT2	rs602662	0	A/A
12	BCMO1	rs7501331	0	C/C
13	GC(D)	rs2282679	0	
14	APOA5	INTERGENIC Rs12272004	2	C/C

Thus, optimal clustering involves maximizing the criterion (Takenaka, 2012). In essence, such a statement means that related objects are collected in each cluster, and there are significant differences between the objects of different clusters. In its class, this problem relates to soft computing problems solved by integer mathematical programming methods. To solve it, we used a complex of programs for assessing the quality of multidimensional objects (Integrated Quality Assessment).

RESULT

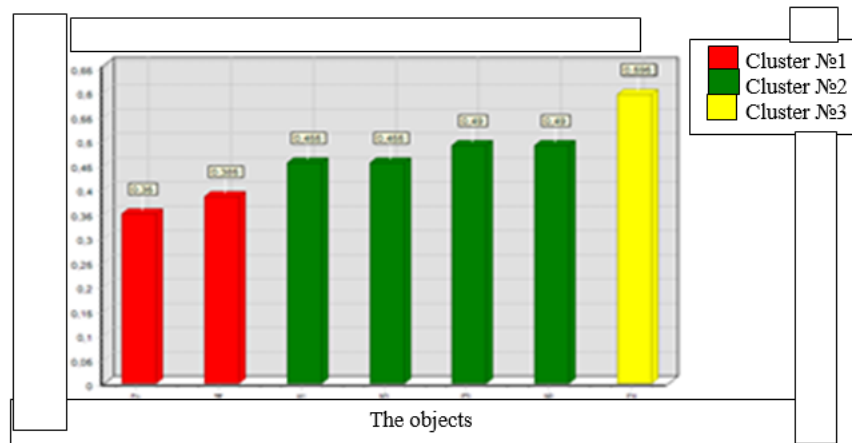


Figure 1. The results of the clustering of profiles of girls.

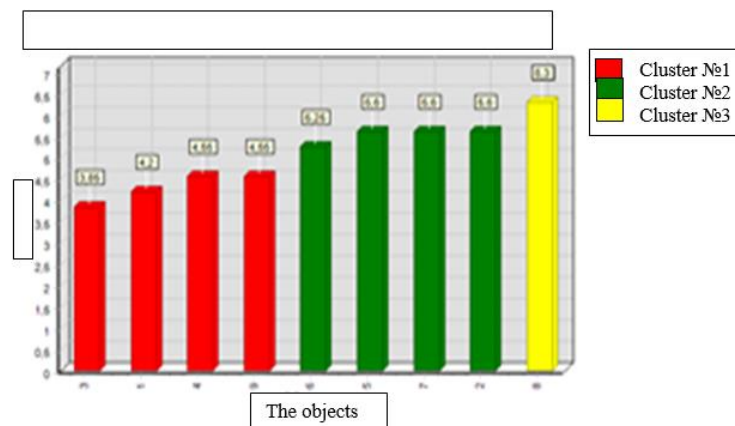


Figure 2. Clusterization results of boys' profiles.

Figure 1 shows the results of solving the clustering problem for questionnaires for girls. Figure 2 shows the results of solving the clustering problem by questionnaires for boys. In both cases, three clusters were obtained. The interpretation and characteristics of these clusters are presented in Table 1.

Figure 3 shows the structural-functional model of the decision-making subsystem in the process of managing personalized food. Block 1 receives a stream of customers for personalized catering. At the output of this block is information about the state of the client genes, which is used in block 2 to solve the clustering problem. Next, for each cluster in block 3, an optimal diet is developed.

Table 2. Interpretation of clusters resulting from clustering.

Diseases / Conditions	Cluster №1	Cluster №2	Cluster №3
Oxidative stress	Oxidative stress (low or moderate predisposition to oxidative stress)	Oxidative stress (medium or high susceptibility to oxidative stress)	Oxidative stress high
Antioxidant status	Antioxidant status high	Antioxidant status medium	Low antioxidant status
Obesity	No (not predisposed to obesity)	No or there (no predisposition or moderate predisposition to obesity)	there is (revealed an average predisposition to obesity)
Nutrition / Diet Enriched	Balanced diet	Low Carbohydrate, Vitamin Enriched Nutrition	Low Carbohydrate, High Vitamin Eating

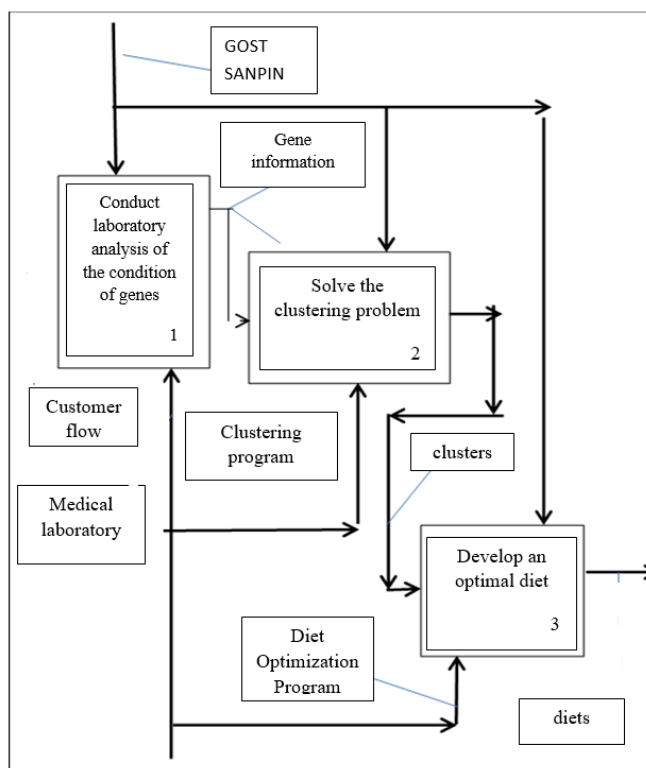


Figure 3. Structural-functional model of the decision-making subsystem in managing personalized nutrition for schoolchildren.

CONCLUSION

This article discusses the statement of the problem of clustering information about the state of customer genes for their personalized nutrition. A program is proposed to solve this problem. The result of solving the clustering problem for real customers who have undergone special medical analyses of the conditions of 14 genes is presented. A methodology has been developed for consumer clustering within a given social group based on threats to their disease state, determined according to the genome. The technique allows to identify clusters with different levels of resistance to hereditary diseases and disease states. A structural and functional model of the decision-making system is developed for managing personalized nutrition for schoolchildren.

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