

Scientific paper

# Estimation of Stability Constants of Cadmium(II) *bis*-Complexes with Amino Acids by Model Based on ${}^3\chi^v$ Connectivity Index

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## Abstract

Linear model for estimation of the second,  $K_2$ , and overall,  $\beta_2$ , stability constant of cadmium(II) binary and ternary *bis*-complexes with five aliphatic  $\alpha$ -amino acids based on valence connectivity index of the 3<sup>rd</sup> order ( ${}^3\chi^v$ ) was developed. Set of amino acids included glycine, alanine, 2-aminobutanoic, 2-aminopentanoic (norvaline) and 2-aminohexanoic acid (norleucine), which by bonding to the cadmium(II) gave 25  $K_2$  and 15  $\beta_2$  values. For estimation of  $\log \beta_2$ , the model gave  $r = 0.940$ , and the S.E.<sub>cv</sub> = 0.10, and for the two subsets of  $\log K_2$  constants the model yielded  $r = 0.936$  and 0.842, and S.E.<sub>cv</sub> = 0.09. The complex CdGG was excluded from all regressions.

**Keywords:** Stability of coordination compounds, Theoretical models, Topological indices

## 1. Introduction

Cadmium is due to its high production and environmental pollution ubiquitous in the biosphere.<sup>1,2</sup> It is toxic metal with deleterious effects on kidney, blood and blood vessels, liver, and bones, not to mention teratogenic effects.<sup>3</sup> Cadmium(II) binds preferently to nucleic acids and their components,<sup>4,5</sup> but its toxic effects are greatly modified by binding to metallothionein.<sup>6,7</sup> Despite its significant biological and environmental role, not many papers were published on cadmium(II) complexes with amino acids.

From the viewpoint of coordination chemistry  $\text{Cd}^{2+}$  is very plastic cation, meaning that distorted coordination geometry is frequent for complexes of cadmium(II) which have coordination number in the range from 2 to 8.<sup>8,9,10</sup> Although cadmium(II) is considerably «softer» cation than  $\text{Cu}^{2+}$  and  $\text{Zn}^{2+}$ , it binds oxalate and diamine with similar affinity as  $\text{Zn}^{2+}$ , but substantially weaker than  $\text{Cu}^{2+}$ .<sup>7,8,10</sup> Consequently, cadmium(II) stability constants for complexes with amino acids and peptides are smaller than copper(II) constants and very close to the constants of zinc(II).<sup>11–19</sup> Despite poor stability, cadmium(II) complexes with amino acids show antimicrobial activity,<sup>20</sup> and seem to participate in cadmium(II) toxicity.<sup>21,22</sup>

Because of biological and environmental significance of cadmium, we decided to check our models for

estimation of stability constants of coordination compounds, originally developed on copper(II) and nickel(II) chelates, on cadmium(II) complexes with amino acids. The main difficulties in applying topological indices for this purpose<sup>25</sup> stem from the fact that the constitutional formula (*i.e.* molecular graph) of a coordination compound is not as well defined as of an organic compound, and that graph theory can not deal with conformers and »classical« stereoisomers. Thus, the valence connectivity index of the 3<sup>rd</sup> order ( ${}^3\chi^v$ ) was calculated for different molecular species<sup>23,24</sup> and correlated to  $\log K_1$ ,  $\log K_2$ , and  $\log \beta_2$  of copper(II) and nickel(II) chelates with amino acids and their derivatives, diamines, triamines, and peptides (dipeptides to pentapeptides).<sup>25</sup> It was possible to obtain a good agreement between experiment and theory, and, moreover, to judge the quality of experimentally determined stability constants.<sup>26</sup> Fair estimates of the stability constants were obtained even from the regression functions developed on different class of compounds.<sup>27,28</sup> Also, by introduction of an indicator variable we succeeded to obtain a common model for estimation of the stability constants of copper(II) and nickel(II) chelates.<sup>29</sup>

In this paper we dealt with the estimation of the second,  $K_2$ , and overall stability constants,  $\beta_2$ , of cadmium(II) binary and ternary *bis*-complexes with five aliphatic  $\alpha$ -amino acids (glycine, alanine, 2-aminobutanoic, 2-aminopentanoic (norvaline) and 2-aminohexanoic acid

(norleucine). Both set of constants,  $K_2$  ( $N = 25$ ) and  $\beta_2$  ( $N = 15$ ), were determined at the same experimental conditions ( $T = 298$  K,  $I$  (LiClO<sub>4</sub>) = 3 mol L<sup>-1</sup>).<sup>10,16,17</sup>

## 2. Methods

### 2.1. Calculation of Topological Indices

We calculated topological indices using a program system E-DRAGON, developed by R. Todeschini and coworkers,<sup>30</sup> which is capable of yielding 119 topological indices in a single run, along with many other molecular descriptors.<sup>31,32</sup> Connectivity matrices were constructed with the aid of *Online SMILES Translator and Structure File Generator*.<sup>33</sup>

All models were developed by using  ${}^3\chi^v$  index (the valence molecular connectivity index of the 3<sup>rd</sup> order), which was defined as:<sup>34–36</sup>

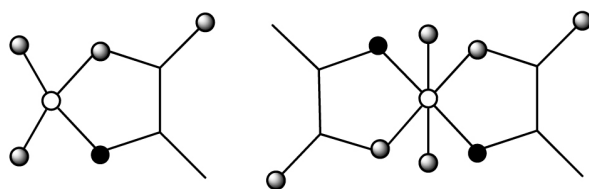
$${}^3\chi^v = \sum_{\text{path}} [\delta(i) \delta(j) \delta(k) \delta(l)]^{-0.5} \quad (1)$$

where  $\delta(i)$ ,  $\delta(j)$ ,  $\delta(k)$ , and  $\delta(l)$  are weights (valence values) of vertices (atoms)  $i$ ,  $j$ ,  $k$ , and  $l$  making up the path of length 3 (three consecutive chemical bonds) in a vertex-weighted molecular graph. Valence value,  $\delta(i)$ , of a vertex  $i$  is defined by:

$$\delta(i) = [Z^v(i) - H(i)]/[Z(i) - Z^v(i) - 1] \quad (2)$$

where  $Z^v(i)$  is the number of valence electrons belonging to the atom corresponding to vertex  $i$ ,  $Z(i)$  is its atomic number, and  $H(i)$  is the number of hydrogen atoms attached to it. For instance,  $\delta$  values for primary, secondary, tertiary and quaternary carbon atoms are 1, 2, 3, and 4, respectively; for oxygen in the OH group it equals 5, and for NH<sub>2</sub> group  $\delta(N) = 3$ . It has to be pointed out that  ${}^3\chi^v$  is only a member of the family of valence connectivity indices  ${}^n\chi^v$ , which differ between each other by the path length, *i.e.* the number of  $\delta$ 's in the summation term, Eq. 1.

The  ${}^3\chi^v$  indices for cadmium(II) *mono*- and *bis*-complexes were calculated from the graph representations of the *aqua* complexes with two water molecules (Fig. 1), assuming that Cd(II) in *mono*-complexes is tetraordinated, and in *bis*-complexes hexacoordinated, as for copper(II) chelates.<sup>23,37</sup> This is supported by X-ray structures



**Figure 1.** The graph representations for cadmium(II) *mono*- (CdL) and *bis*-complex (CdLA) with alanine. Heteroatoms are marked with O(Cd), ●(N), and ●(O).

of cadmium(II) *bis*-complexes with glycine<sup>38</sup> and alanine,<sup>39</sup> and some mixed Cd(II) *bis*-complexes.<sup>40,41,42</sup> Moreover, the alternative assumption, that both, CdL and CdLA complexes are tetraordinated yielded bad results.

### 2.2. Regression Calculations

Regression calculations, including the leave-one-out procedure of cross validation, *cv*, were done using the CROMRsel program.<sup>43</sup> The standard error of cross validation estimate is defined as:

$$S.E._{cv} = \sqrt{\sum_i \frac{\Delta X_i^2}{N}} \quad (3)$$

where  $\Delta X$  and  $N$  denotes *cv* residuals and the number of reference points, respectively.

## 3. Results and Discussion

### 3.1. Estimation of the Overall Stability Constant $\beta_2$

Firstly we estimated the overall stability constant,  $\beta_2$ :



where M denotes Cd<sup>2+</sup>, and L and A denote  $\alpha$ -amino acids. Model was developed on 15 cadmium(II) binary and ternary *bis*-complexes with five aliphatic  $\alpha$ -amino acids (Table 1). Beside naturally occurring glycine (G) and alanine (A), there were 2-aminobutanoic (B), 2-aminopentanoic (P) and 2-aminohexanoic acid (H) in the set.

Although at first it seemed there is no correlation between  $\log \beta_2$  and  ${}^3\chi^v$ (CdLA), after closer inspection of Fig. 2 we noticed some kind of the order. As  $\beta_2$  constant is independent on sequence of ligand bonding to the metal, complexes were such named that the first ligand (L) was always smaller or equal to the second ligand (A). Thus we were able to see from Fig. 2 that stability constant  $\beta_2$  of *bis*-complexes with identical second ligand (A) depends on the first ligand (L), *i.e.*  $\log \beta_2$  linearly decreases along the homologous series of the first amino acid (with an exception of PH in the hexanoate series). Also, for the mixed complexes with identical first ligand,  $\log \beta_2$  linearly increases along the homologous series of the second amino acid, with an exception of GG and GA in the glycine series. From Fig. 2 it is evident that slopes of the descending lines drop and the slopes of the ascending lines rise along the homologous series.

Subsequently, we developed a linear function based on the ascending lines:

$$\log \beta_2 = a_1[{}^3\chi^v(\text{CdLA})] + a_2[{}^3\chi^v(\text{CdL})] + a_3[{}^3\chi^v(\text{CdA})/{}^3\chi^v(\text{CdL})] + b \quad (5)$$

**Table 1.** Experimental overall,  $\beta_2$ , and the second,  $K_2$ , stability constants and  ${}^3\chi^v$  indices calculated for Cd(II) aminoacids

| (Complexes<br>(CdLA or CdAL))* | $\log \beta_2$ | $\log K_2$ | ${}^3\chi^v(\text{CdLA})$ | ${}^3\chi^v(\text{CdL})$ | ${}^3\chi^v(\text{CdA})$ |
|--------------------------------|----------------|------------|---------------------------|--------------------------|--------------------------|
| CdGG                           | 7.49           | 3.48       | 6.933                     | 2.436                    | 2.436                    |
| CdGA <sup>a</sup>              | 7.47           | 3.46       | 7.292                     | 2.436                    | 2.921                    |
| CdGB <sup>a</sup>              | 7.34           | 3.33       | 7.496                     | 2.436                    | 3.125                    |
| CdGP <sup>a</sup>              | 7.41           | 3.40       | 7.656                     | 2.436                    | 3.285                    |
| CdGH <sup>a</sup>              | 7.43           | 3.42       | 7.925                     | 2.436                    | 3.554                    |
| CdAA <sup>ab</sup>             | 6.93           | 3.24       | 7.651                     | 2.921                    | 2.921                    |
| CdAB <sup>a</sup>              | 7.13           | 3.44       | 7.855                     | 2.921                    | 3.125                    |
| CdAP <sup>a</sup>              | 7.15           | 3.46       | 8.015                     | 2.921                    | 3.285                    |
| CdAH <sup>a</sup>              | 7.24           | 3.55       | 8.284                     | 2.921                    | 3.554                    |
| CdBB <sup>ab</sup>             | 6.88           | 3.24       | 8.059                     | 3.125                    | 3.125                    |
| CdBP <sup>a</sup>              | 7.05           | 3.41       | 8.219                     | 3.125                    | 3.285                    |
| CdBH <sup>a</sup>              | 7.2            | 3.56       | 8.488                     | 3.125                    | 3.554                    |
| CdPP <sup>ab</sup>             | 7.01           | 3.29       | 8.379                     | 3.285                    | 3.285                    |
| CdPH <sup>a</sup>              | 7.31           | 3.59       | 8.648                     | 3.285                    | 3.554                    |
| CdHH <sup>ab</sup>             | 7.03           | 3.29       | 8.917                     | 3.554                    | 3.554                    |
| CdAG <sup>b</sup>              |                | 3.78       | 7.292                     | 2.436                    | 2.921                    |
| CDBG <sup>b</sup>              |                | 3.70       | 7.496                     | 2.436                    | 3.125                    |
| CdBA <sup>b</sup>              |                | 3.49       | 7.855                     | 2.921                    | 3.125                    |
| CdPG <sup>b</sup>              |                | 3.69       | 7.656                     | 2.436                    | 3.285                    |
| CdPA <sup>b</sup>              |                | 3.43       | 8.015                     | 2.921                    | 3.285                    |
| CdPB <sup>b</sup>              |                | 3.33       | 8.219                     | 3.125                    | 3.285                    |
| CdHG <sup>b</sup>              |                | 3.69       | 7.925                     | 2.436                    | 3.554                    |
| CdHA <sup>b</sup>              |                | 3.50       | 8.284                     | 2.921                    | 3.554                    |
| CdHB <sup>b</sup>              |                | 3.46       | 8.488                     | 3.125                    | 3.554                    |
| CdHP <sup>b</sup>              |                | 3.57       | 8.648                     | 3.285                    | 3.554                    |

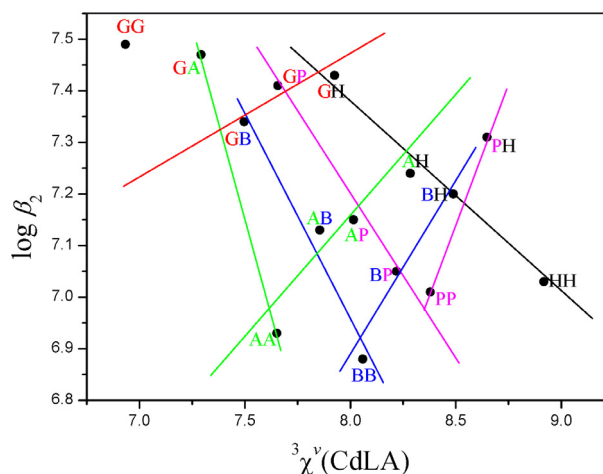
\* L always denotes smaller ligand and A the bigger one.

<sup>a</sup> Complexes in the set for estimation of  $\beta_2$ , and in the second subset for estimation of  $K_2$ .

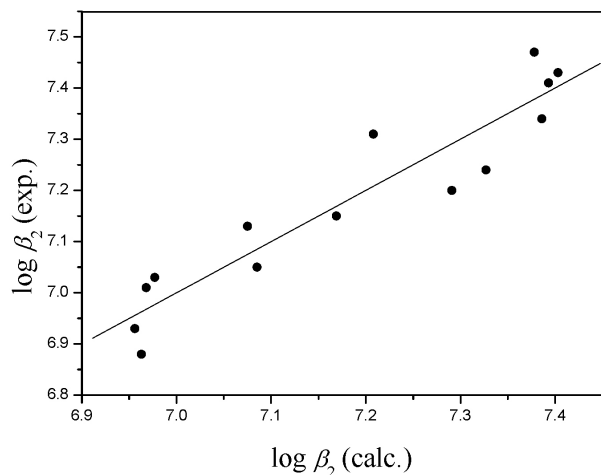
<sup>b</sup> Complexes in the first subset for estimation of  $K_2$ .

The regression gave  $r = 0.940$ , and the  $\text{S.E.}_{\text{cv}} = 0.10$  (Table 2, Fig. 3). The complex CdGG was not included into regression because it does not belong to any family of lines.

The  ${}^3\chi^v(\text{CdA})$  index was included in the function because ascending lines show linear dependence of  $\beta_2$  on the variation of the second ligand. Also,  ${}^3\chi^v(\text{CdA})$  was divided with  ${}^3\chi^v(\text{CdL})$  to compensate different slope of the lines, which depends on the first ligand (Fig. 2). On the other hand, the second term of Eq. 5,  ${}^3\chi^v(\text{CdL})$ , was introduced as a compensation for difference in intercepts of the ascending lines (Fig. 2). They decrease in the homologous series of the first ligand.

**Figure 2.** Plot of  $\log \beta_2$  vs.  ${}^3\chi^v(\text{CdLA})$  reveals sequences of linear dependences. The ligand and the line that indicate linear dependence of that ligand series are coloured with the same colour.**Table 2.** Linear regressions (Eq. 5) for the estimation of the overall,  $\beta_2$ , and second,  $K_2$ , stability constants

| Regr. No. | N  | Dependent variable | Slope (S.E.)             |                         |   | Intercept (S.E.) | r     | S.E. | S.E. <sub>cv</sub> |
|-----------|----|--------------------|--------------------------|-------------------------|---|------------------|-------|------|--------------------|
|           |    |                    | Independent variable     |                         |   |                  |       |      |                    |
|           |    |                    | ${}^3\chi^v(\text{MLA})$ | ${}^3\chi^v(\text{ML})$ | $\frac{{}^3\chi^v(\text{MA})}{{}^3\chi^v(\text{ML})}$ |                  |       |      |                    |
| 1         | 14 | $\log \beta_2$     | 3.33(89)                 | -6.6(18)                | -8.0(24)  | 8.85(92)         | 0.940 | 0.06 | 0.10               |
| 2         | 14 | $\log K_2$         | 3.08(89)                 | -7.7(24)                | -6.3(18)  | 5.71(93)         | 0.936 | 0.06 | 0.09               |
| 3         | 14 | $\log K_2$         | 3.50(86)                 | -6.8(17)                | -8.9(23)  | 5.06(89)         | 0.842 | 0.06 | 0.09               |



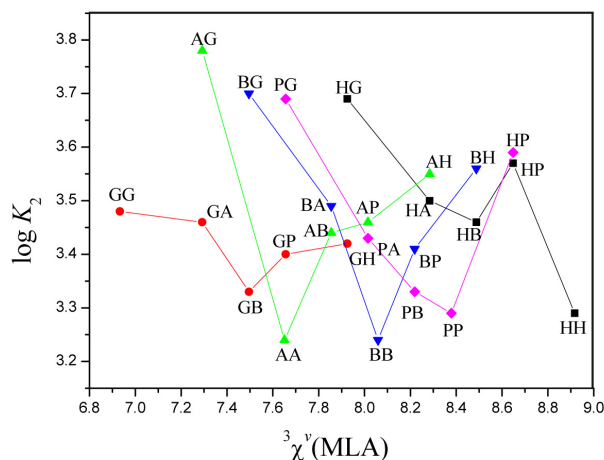
**Figure 3.** Experimental vs. calculated values of  $\log \beta_2$  for cadmium(II) complexes with five aliphatic  $\alpha$ -amino acids ( $N = 14$ ), Table 2, No. 1;  $r = 0.940$ ,  $S.E._{cv} = 0.10$ .

### 3. 2. Estimation of the Second Stability Constant $K_2$

The values of the second stability constant,  $K_2$ :



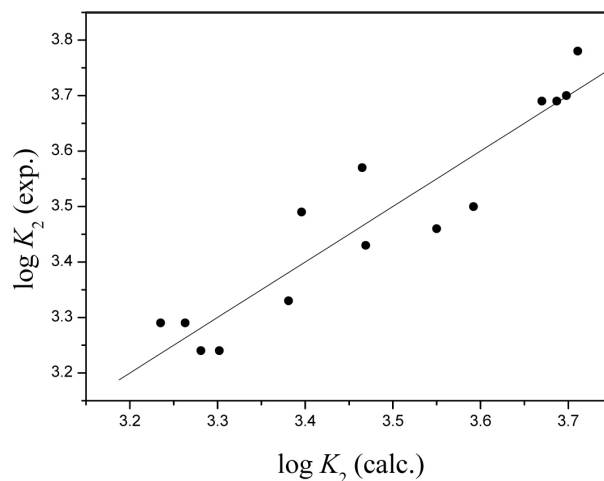
of the cadmium(II) complexes (Table 1), with the same ligands as in the case of the  $\beta_2$  constant, were also estimated by the Eq. 5. However, because of the dependence of  $K_2$  on the sequence of ligand bonding to the Cd(II), there were 25  $K_2$  values which we divided into two subsets (Table 1). The dependence of  $\log K_2$  on  ${}^3\chi^v(\text{CdLA})$  (Fig. 4) shows that for every first ligand series of bis-complexes the binary complex has the minimum  $K_2$  value. The exception was again the glycine series. Thus we



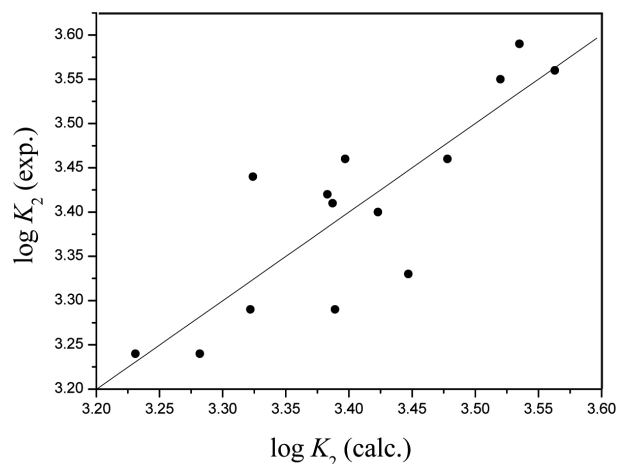
**Figure 4.** Plot of  $\log K_2$  vs.  ${}^3\chi^v(\text{CdLA})$ . The ligand and the curve that connect the complexes of that ligand series are coloured with the same colour.

divided the data to make the first subset consisted of complexes preceding the minimum and the second subset consisted of complexes to go after the minimum (Fig. 4). In the second subset were complexes with the first ligand smaller or equal to second, same as in the case of  $\beta_2$  constant, and the opposite was true for the first subset (Table 1). For both subsets the pattern similar to  $\log \beta_2$  pattern (Fig. 2) was observed, although with somewhat larger deviations for the second set. Consequently, Eq. 5 was also used to estimate  $\log K_2$ . The only difference was that for  $K_2$  constant, L denoted the smaller and A the bigger ligand, regardless the order of their bonding to the cadmium(II) (Eq. 6).

For the first subset ( $N = 14$ ) the regression gave  $r = 0.936$ , and  $S.E._{cv} = 0.09$  (Table 2, Fig. 5), and for the second subset ( $N = 14$ )  $r = 0.842$ , and  $S.E._{cv} = 0.09$



**Figure 5.** Experimental vs. calculated values of  $\log K_2$  for the first subset of cadmium(II) complexes (Table 1) with five aliphatic  $\alpha$ -amino acids ( $N = 14$ ), Table 2, No. 2;  $r = 0.936$ ,  $S.E._{cv} = 0.09$ .



**Figure 6.** Experimental vs. calculated values of  $\log K_2$  for the second subset of cadmium(II) complexes (Table 1) with five aliphatic  $\alpha$ -amino acids ( $N = 14$ ), Table 2, No. 3;  $r = 0.842$ ,  $S.E._{cv} = 0.09$

(Table 2, Fig. 6) were yielded. From both subsets, as for the estimation of  $\beta_2$ , the complex CdGG was excluded because it does not belong to any series of lines.

## 4. Conclusion

In this paper, for the first time we developed the regression model for the estimation of stability constants of cadmium(II) complexes. The model is valid for the estimation of second,  $K_2$ , and overall,  $\beta_2$ , stability constant of cadmium(II) bis-complexes with aliphatic amino acids. It is based on the observation that stability constants of bis-complexes with the same smaller ligand linearly increase along the homologous series of the bigger ligand, and that stability constants of bis-complexes with the same bigger ligand linearly decrease along the homologous series of the smaller ligand.

In further research, we will try to apply this model to other metal complexes to see if it is generally valid.

## 5. Acknowledgement

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## Povzetek

Razvili smo linearni model za oceno druge in splošne konstante stabilnosti,  $K_2$  in  $\beta_2$ , za kadmijeve(II) binarne in ternarne bis-komplekse s petimi alifatskimi  $\alpha$ -amino kislinami; model je osnovan na valenčnem indeksu povezanosti tretjega reda ( ${}^3\chi^v$ ). Niz aminokislin je vseboval glicin, alanin, 2-aminobutanoično, 2-aminopentanoično (norvalin) in 2-aminoheksanoično kislino (norlevcin), ki se vežejo na kadmij(II) in tako določimo 25  $K_2$  and 15  $\beta_2$  vrednosti. Napovedi modela za  $\log \beta_2$  so ocenjene z  $r = 0.940$  in  $S.E._{cv} = 0.10$ , medtem ko so statistični parametri ocene modela za  $\log K_2$  dveh delnih nizov  $r = 0.936$  in  $0.842$ , in  $S.E._{cv} = 0.09$ . Kompleks CdGG je bil izločen iz vseh regresijskih modelov.