Bayesian Hierarchical Model of Width of Keratinized Gingiva

Veska Noncheva¹, Maria Dobreva¹, Ivan Chenchev²

¹(Faculty of Mathematics and Informatics, University of Plovdiv "Paisii Hilendarski", 24 Tzar Asen, 4000, Plovdiv, Bulgaria)

²(Department of Oral Surgery, Faculty of Dental Medicine, Medical University-Plovdiv, Bulgaria)

Abstract: The purpose of this paper is to offer a method for studying the treatment result of gingival recession. A parameter showing the success of the surgical treatment of gingival recessions is the keratinized gingival width. It was measured four times: at baseline, after 1 month, after 3 months and after 6 months. Every patient has data that can be described by an individual trend. Bayesian hierarchical model of the keratinized gingiva width's increase rate is built.

Keywords: Biostatistics, gingival recession, hierarchical modeling, longitudinal data, split-mouth randomized clinical trial

I. Introduction

The amount of keratinized gingiva is derived from the sum of the free gingiva and attached gingiva. Keratinized gingiva width (KGW) is measured from the mucogingival junction (MGJ) to the lowest point of the gingival margin in the middle of the crown of the tooth with gingival recession. In a healthy and intact periodontium, the length of keratinized attached gingiva includes coronally from the bottom of gingival sulcus to the mucogingival junction apically. The width of attached gingiva varies with the age. The keratinized attached gingiva provides the periodontium with increase resistance to external injury and stabilized the gingival margin against physical forces and helps patients plaque control measurements. An adequate amount of keratinized gingiva maintains gingival health by protecting the marginal gingiva from bacterial invasion [3,11], preventing an increase in gingival recession [4,22], facilitating plaque control [3,5,6,23], and improving denture stability [12,23].

Ainamo et al. [11] in different studies said that, MGJ remains stationary throughout life and changes in width of attached gingiva are caused by modification in position of coronal gingival margin. The width of attached gingiva increases with age and in supra-erupted teeth [7,15]. Lang and Löe [5] reported a study on the relationship between the gingival width and inflammation, in an effort to determine the adequate amount: In 100% of teeth with less than 2mm of keratinized tissue, inflammation and exudates was present; 76% of cases with greater than 2mm of keratinized tissue there was no exudates and was considered as clinically healthy. They concluded that 2mm of keratinized gingiva, with less than 1mm of attached gingiva is adequate to maintain gingival health. An adequate band of attached gingiva could be defined as that amount which is sufficient to prevent recession in opinion of individual practioners [16]. Thus no minimum width of attached gingiva has been established as standard necessary for gingival health. Another paper by Miyasato et al. [9] concluded that there is no relationship between inflammation and amount of attached gingiva whether or not plaque is present. According to Miller PD and Harris RJ one of the criteria to achieve complete root coverage is to achieve sufficient width of keratinized gingiva (KGW), [10,13]. In the dental researches increasing KGW is an important criterion for determining the success of the treatment of gingival recession.

II. Problem

The aim of this paper is to present an approach for studying the rate of increase of the keratinized gingiva width after surgery. This research involved 30 people at the age of 23 to 70 years with a total of 118 symmetrical Miller's Class I and Class II gingival recessions on different places of the jaws. All of the patients' gingival recessions were treated surgically and the results were monitored for six months postoperatively. The recessions on one side of the jaw were treated with coronally advanced flap (CAF) combined with platelet rich fibrin membrane, while the other side was treated with CAF combined with connective tissue graft [1]. An approach for the comparison of two methods of treatment of gingival recessions is described in [24]. The success of the operation was evaluated through the measurement of the keratinized gingival width in millimeters. KGW was measured 4 times: at baseline, after 1 month, after 3 months and after 6 months. Every patient has data that can be described by an individual trend. Because of it we will build hierarchical model (*see* [20,21]) of the rate of increase of the keratinized gingiva width. Hierarchical models of phenomena from oral surgery are described in [18,24].

III. Model

We suppose that every individual *j* contributes multiple observations of $(x_{i|j}, y_{ij})$ pairs, where the subscript notation *i|j* means the *i*-th observation within the *j*-th individual (patient). The observations $y_{i|j}$, *i=1*, ..., *118*, *j=1,2,3*, are the values of KGW measured at 1st, 3rd and 6th month after, and $x_{i|j}$ are the values of KGW measured before, at baseline, 1st and 3rd month respectively. With these longitudinal data, we will estimate a regression curve for every individual. Our goal is to describe each individual with a linear regression, and simultaneously to estimate the typical slope and intercept of the group overall. A key assumption for our analysis is that each individual is representative of the group. Therefore, every individual slopes and intercepts. Thereby we get sharing of information across individuals, and shrinkage of individual estimates toward the overarching mode.



Figure 1.KGW data 3 months and 6 months after surgery.

The model assumes homogeneity of variance. Homogeneity of variance is less easy to identify visually when the predictor's values are not uniformly distributed. The apparent vertical spread of the data seems to be larger at x = 1.5 than at x = 3.5 in Figure 1. Despite this deceiving appearance, the data do respect homogeneity of variance. The reason for the apparent violation is that for regions in which x is sparse, there is less opportunity for the sampled y values to come from the tails of the noise distribution. In regions where the predictor is dense, there are many opportunities for the predicted variable to come from the tails. The data have outliers, and the use of heavy-tailed noise distributions is straight forward in contemporary Bayesian software. Student's t-distribution has the following three parameters: μ that controls its mean, the "scale" parameter σ that controls its width, and a third parameter v that controls the heaviness of its tails, called "normality" parameter. The parameter v is also known as "degrees of freedom". The normality parameter can range continuously from 1 to ∞ . When v = 1 the t distribution has very heavy tails and it is a convenient descriptive model of data with outliers. When v approaches ∞ the t distribution becomes much closed to normal.

1.1 Hierarchical dependency structure

The hierarchical dependencies in the model are presented in Figure 2.



Figure 2. A model of dependencies. The standard deviation of the noise within each subject is the same for all subjects.

The datum y_{ijj} is the *i*-th datum within the *j*-th subject. It has a Student's *t*-distributed random value around the central tendency $\mu_{ij} = \beta_{0,j} + \beta_{1,j} x_{ijj}$, j=1,2,3, i=1,...,118. The intercept and slope are subscripted with *j*, meaning that every subject has its own slope and intercept. The slopes across the individuals all come from a higher-level normal distribution. The model assumes that $\beta_{1,j} \sim \text{normal} (\mu_1, \sigma_1)$, where μ_1 describes the typical slope of the individuals and σ_1 describes the variability of those individual slopes. Analogous considerations apply to the intercepts $\beta_{0,j} \sim \text{normal} (\mu_0, \sigma_0)$. The group-level parameters are given generic vague priors. This model assumes that the standard deviation of the noise within each subject is the same for all subjects. The tdistribution's scale parameter σ is given a noncommittal uniform prior, and the normality parameter *v* is given a broad exponential prior. The intercept and slope are given broad normal priors that are vague on the scale of the data. The two group-level parameters β_0 and β_1 have separate normal priors. Our goal is to determine what combinations of the parameters are credible, given the data. The solution comes from Bayes' rule and its implementation in JAGS software [25]. Our task is to specify sensible priors and to make sure that the MCMC process generates a trustworthy posterior sample that is converged and well mixed.

IV. Solution

The upper-right panel of Figure 3 shows the posterior distribution of the jointly credible values of the slope $\beta_{I,3}$ and the intercept $\beta_{0,3}$. The correlation of the credible values of the slope and intercept is extremely strong. This narrow diagonal posterior distribution is difficult for Gibbs sampling algorithms to explore, resulting in extreme inefficiency in the chains. The two parameter values change very slowly, and the MCMC chain is highly auto correlated. In practice, this requires us to wait too long before getting a suitably representative sample from the posterior distribution. In order to make the sampling more efficient, the data are standardized before being sent to the model. In this way credible regression lines do not suffer such strong correlation between their slopes and intercepts.



Figure 3. Marginal posterior distributions on parameters.



1.2 MCMC diagnostics

Our first task is to check whether the chains appear to be well mixed and suitably representative of the posterior distribution. Figure 4 shows diagnostic information for the standardized $\beta_{1,3}$ parameter. The density plot in the lower-right shows that the three sub chains are well super-imposed which is echoed by the Gelman-Rubin statistic in the lower-left panel being very close to 1.0. The autocorrelation plot in the upper-right panel shows essentially zero autocorrelation. The diagnostic information for other parameters is similar.

| | Mean | Median | Mode | 95% | 95% |
|----------|---------|---------|---------|----------|---------|
| | | | | HDIlow | HDIhigh |
| beta0[1] | 1.40000 | 1.40000 | 1.60004 | 1.19982 | 1.60015 |
| beta0[2] | 0.00000 | 0.00000 | 0.00000 | -0.00023 | 0.00022 |
| beta0[3] | 0.00000 | 0.00000 | 0.00001 | -0.00024 | 0.00025 |
| beta1[1] | 0.41111 | 0.40000 | 0.40007 | 0.33320 | 0.50013 |
| beta1[2] | 1.00000 | 1.00000 | 1.00000 | 0.99990 | 1.00010 |
| beta1[3] | 1.00000 | 1.00000 | 1.00000 | 0.99989 | 1.00011 |
| Sigma | 0.00064 | 0.00063 | 0.00063 | 0.00063 | 0.00065 |
| Nu | 0.20606 | 0.20563 | 0.20416 | 0.18079 | 0.23264 |

Table 1. Estimates of the model parameters v, σ , $\beta_{0,k}$ and $\beta_{1,k}$, k=1,2,3.

1.3 Interpreting the posterior distributions

Now we will discuss the results of the analysis. The parameter estimates in Table 1 indicates that the most credible value of the slope $\beta_{1,1}$ is about 0.4, which means that KGW in 1st month after surgery increases about 0.4 mm for every 1-mm increase in KGW at baseline. The posterior distribution of $\beta_{1,2}$ shows that the most credible value of the slope $\beta_{1,2}$ is about 1, which means that KGW in 3rd month after surgery increases about 1 mm for every 1-mm increase in KGW in 1st month. The slope $\beta_{1,3}$ has modal estimate of about 1. It means that KGW in 6th month after surgery increases about 1 mm for every 1-mm increase in KGW in 1st month. The slope $\beta_{1,3}$ has modal estimate of about 1. It means that KGW in 6th month after surgery increases about 1 mm for every 1-mm increase in KGW in 3rd month. One way to summarize the uncertainty is by marking the span of values that are most credible and cover 95% of the distribution. This is the *highest density interval* (HDI). Values within the 95% HDI are more credible (i.e., have higher probability "density") than values outside the HDI, and the values from a slope of about 0.33 mm per mm to a slope of about 0.5 mm per mm. The 95% HDI for both slopes $\beta_{1,2}$ and $\beta_{1,3}$ are very precise. The individuals seem to have nearly the same slope and σ_1 is estimated to be small (see Table 1). In fact, the 95% HDIs on the slopes are far from zero. Thus, there is no doubt that there is a linear trend in these KGW data.

V. Conclusion

We have studied the behavior of the clinical periodontal parameter keratinized gingiva width (KGW) during the gingival recession therapy. This study presents an approach to study the rate of increase of the keratinized gingiva width (KGW) during treatment. The slope parameters in the model are meaningful because they describe tendencies in the KGW data.

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