

STATISTICAL DIAGNOSTICS OF METASTATIC INVOLVEMENT OF REGIONAL LYMPH NODES

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Abstract

The results of diagnostics of metastatic involvement of regional lymph nodes using discriminant analysis, support vector machine and robust methods are discussed. The method of statistical classification with indicating patients that require more detailed diagnostics is proposed and analysed.

1 Introduction

Melanoma of the skin is one of most aggressive human malignant tumors. In Europe melanoma takes 17th place in men and 8th place in women in rating of most frequently diagnosed cancer types [1]. Nevertheless local melanoma is not a disease with synonymous poor prognosis. In case of early diagnosis complete cure may be achieved in 95% of cases. During last decades the investigators are attempting to determine the factors, responsible for the course and prognosis of the disease. The significance of the factors is taken into consideration in determination of the stage of the disease, the importance of other factors is rejected during time.

Nowadays the problem of lymph node dissection combined with excision of primary focus of tumor remains disputable. The selection of group of patients with high risk of metastases in lymph nodes is very actual for the decrease of rate of lymph node dissection (often groundless). Although this group of patients is a very limited part of population, in a majority of cases right diagnostic decision is out of difficulties and requires working out special diagnostic programmes.

In clinical oncology parametric discriminant analysis [2] and support vector machine [3] are widely used for malignant neoplasms diagnostic in the case of presence of a priori information on a training sample. The mixture of observations should be separable in the feature space for successful classification.

2 Results of diagnostics using discriminant analysis and support vector machine

An attempt of working out indications to prophylactic lymphadenectomy in patients with clinically intact peripheral lymph nodes. Retrospective investigation of frequency of lymph node metastases depending on clinico-anatomical factors, characterizing primary tumor and organism of patient is carried out.

From 1970 to 2002, 982 melanoma patients treated at the State Institution “N.N. Alexandrov Research Institute of Oncology and Medical Radiology” (Minsk, Belarus). Class Ω_1 consisted of 205 patients with histological evidence of metastases in regional lymph nodes. Class Ω_2 included 777 patients with no regional lymph node metastases on histological examination.

A retrospective study was carried out to evaluate the rate of metastatic involvement of regional lymph nodes with regard to the following continuous and discrete clinico-anatomical features: age (AG), background (B), disease duration before the start of the treatment (DDST), tumor growth (TG), tumor square (TS), degree of tumor pigmentation (DTP), Clark invasion level (CIV), Breslow tumor thickness (BTH), ulceration (U), tumor-infiltrating lymphocytes (TIL), histological subtype (HS), growth phase (GP), satellites (ST), vascular invasion (VI), anatomic distribution of melanoma (ADM).

Quadratic discriminant functions were used to include information on difference in covariances between features of the classes; they increase the diagnostics accuracy in comparison with the linear decision rules [2]. Clinico-anatomical features DDST, BTH and TS have an abnormal observations (outliers). As it was mentioned in [2], the presence of such observations (which have to be used to construct diagnostics rules) causes a loss in the accuracy of the decision rules. Therefore robust decision rules based on the robust Huber M -estimators [2] were used.

We form two informative sets from the analyzed features 1) AG, B, DDST, TS, CIV, BTH; 2) AG, DDST, TS, DTP, CIV, BTH. We use robust quadratic decision rule for reclassification of the training sample. The rate of true classification using the first and the second informative sets is 65.22% and 62.07% accordingly for the patients from the class Ω_1 , and 63.88% and 62.70% accordingly for the patients from the class Ω_2 . So, the unconditional rate of true classification is 64.14% for the first informative set and 62.58% for the second informative set.

Standard discriminant analysis procedures are usually applied either to continuous observations, or to discrete data, and for the analysed data satisfactory values of error probabilities for the discriminant function can not be reached.

The support vector machine (SVM) technique gives the diagnostics performance of 58.02% for the first informative set and 57.73% for the second informative set.

Because of low performance of the robust decision rules and of the SVM method, and also because some extra diagnostics is available, we consider the decision rules with one more available decision.

3 Classification with detection of observations that require more detailed diagnostics

Let us construct a decision rule with 3 admissible decisions about an observation $x = x(\omega) \in \mathbf{R}^N$ on the patient $\omega \in \Omega = \Omega_1 \cup \Omega_2$. The decisions $d = 1$ and $d = 2$ mean that $\omega \in \Omega_1$ and $\omega \in \Omega_2$ respectively. The decision $d = 0$ corresponds to the situation, where ω needs more detailed diagnostics.

Denote by $\lambda(x)$ the discriminant function constructed for the problem of distinguishing between Ω_1 and Ω_2 . In the case where discrete and continuous features are supposed independent, and the Fisher model describes the sub-vector of continuous features, the discriminant function of the adaptive decision rule based on the maximum likelihood criterion, has the form:

$$\lambda(x) = \ln \left(\hat{P}_2\{x_{discr}\} / \hat{P}_1\{x_{discr}\} \right) + b^T x_{cont} - H;$$

$b = \hat{\Sigma}_{cont}^{-1}(\hat{\mu}_{2cont} - \hat{\mu}_{1cont})$; $H = \frac{1}{2} \left(\hat{\mu}_{2cont}^T \hat{\Sigma}_{cont}^{-1} \hat{\mu}_{2cont} - \hat{\mu}_{1cont}^T \hat{\Sigma}_{cont}^{-1} \hat{\mu}_{1cont} \right)$, where the vector x_{discr} is the subvector of x , consists of the values of discrete features; x_{cont} is the subvector of values of continuous features; $\hat{P}_i\{\cdot\}$ is the estimate of the probability to observe the value x_{discr} under the condition that the observation x belongs to the class Ω_i ; $\hat{\mu}_{icont}$ are estimates of the vectors of mathematical expectations of x_{cont} for the observations from Ω_i , $i = 1, 2$; $\hat{\Sigma}_{cont}$ is the estimate of the covariance matrix for x_{cont} .

Introduce the decision rule:

$$d = d(x) = 2 \cdot \mathbf{1}_{(B, +\infty)}(\lambda(x)) + \mathbf{1}_{(-\infty, A)}(\lambda(x)), \quad (1)$$

where x is the observed value of the vector of features, $A < 0$, $B > 0$ are parameters. For the choice of A , B , the following two criteria are proposed.

3.1 Risk-oriented approach

Introduce the notation: $\pi_1 \in (0, 1)$ is the prior probability of the random event $\{\omega \in \Omega_1\}$; $P_{ij}(A, B)$ is the probability of the decision $d = j$ provided the observation comes from Ω_i ; $w_{ij} \geq 0$ is the cost of the correspondent decision. The values A , B are the solutions of the risk (expected losses) minimization problem:

$$R(A, B) = \pi_1 (w_{12}P_{12}(A, B) + w_{10}P_{10}(A, B)) + (1 - \pi_1) (w_{21}P_{21}(A, B) + w_{20}P_{20}(A, B)) \rightarrow \min_{A < 0, B > 0}. \quad (2)$$

Consider the case where observations from the class Ω_i have the Gaussian probability distribution:

$$\mathcal{L}\{x(\omega)\} = \mathcal{N}_N(\mu_i, \Sigma), \quad \omega \in \Omega_i, i = 1, 2. \quad (3)$$

Denote: $\Phi(\cdot)$ is the distribution function of $\mathcal{N}_1(0, 1)$; $\Delta = \sqrt{(\mu_2 - \mu_1)^T \Sigma^{-1} (\mu_2 - \mu_1)}$, $b = \Sigma^{-1}(\mu_2 - \mu_1)$, $H = \frac{1}{2} (\mu_2^T \Sigma^{-1} \mu_2 - \mu_1^T \Sigma^{-1} \mu_1)$, $m_1 = b^T \mu_1 - H$.

Theorem 1. *If the model (3) is valid, then the minimization problem (2) is equivalent to the pair of separate minimization problems:*

$$R(A) = (1 - \pi_1)(w_{21} - w_{20})\Phi\left(\frac{A - m_1}{\Delta} - \Delta\right) - \pi_1 w_{10}\Phi\left(\frac{A - m_1}{\Delta}\right) \rightarrow \min_{A < 0},$$

$$R(B) = \pi_1(w_{10} - w_{12})\Phi\left(\frac{B - m_1}{\Delta}\right) + (1 - \pi_1)w_{20}\Phi\left(\frac{B - m_1}{\Delta} - \Delta\right) \rightarrow \min_{B > 0}.$$

3.2 Frequency-oriented approach

Let the maximal possible values α, β for the error probabilities be given:

$$P_{12}(A, B) = \alpha, P_{21}(A, B) = \beta. \quad (4)$$

Solving (4) w.r.t. A, B , we get the values to be used in the decision rule (1).

Denote by $\Phi^{-1}(\gamma)$ the quantile of the level $\gamma \in (0, 1)$ for the standard normal probability distribution $\mathcal{N}_1(0, 1)$.

Theorem 2. *If the model (3) holds, then the solution of (4) is*

$$A = m_1 + \Delta \cdot \Phi^{-1}(\beta) + \Delta^2, B = m_1 + \Delta \cdot \Phi^{-1}(1 - \alpha).$$

3.3 Numerical results

Let us present briefly the results of statistical data analysis with the proposed algorithm. The vector x_{discr} was consisted of four components: TG, DTP, CIV, U; the vector x_{cont} was formed by three features: DDST, TS, BTH. The thresholds were $A = -0.25, B = 0.1$.

Using the decision rule (1) the following estimates of error have been obtained for the observations from the training sample: $P_{12} = 0.125, P_{21} = 0.257$; 25 patients of 137 (18.25%) from the class Ω_1 and 92 of 574 (16.03%) were indicated as patients requiring extra diagnostics.

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