Optimal Partner Wavelength Combination Method Applied to NIR Spectroscopic Analysis of Human Serum Globulin

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Abstract

Human serum globulin (GLB), which contains various antibodies in healthy human serum, is of great significance for clinical trials and disease diagnosis. In this study, the GLB in human serum was rapidly analyzed by near infrared (NIR) spectroscopy without chemical reagents. Optimal partner wavelength combination (OPWC) method was employed for selecting discrete information wavelength. For the OPWC, the redundant wavelengths were removed by repeated projection iteration based on binary linear regression, and the result converged to stable number of wavelengths. By the way, the convergence of algorithm was proved theoretically. Moving window partial least squares (MW-PLS) and Monte Carlo uninformative variable elimination PLS (MC-UVE-PLS) methods, which are two well-performed wavelength selection methods, were also performed for comparison.

The optimal models were obtained by the three methods, and the corresponding root-mean-square error of cross validation and correlation coefficient of prediction (SECV, RPCV) were 0.813 g·L⁻¹ and 0.978 with OPWC combined with PLS (OPWC-PLS), and 0.804 g L⁻¹ and 0.979 with MW-PLS, and 1.153 g L⁻¹ and 0.948 with MC-UVE-PLS, respectively. The OPWC-PLS and MW-PLS methods achieved almost the same good results. However, the OPWC only contained 28 wavelengths, so it had obvious lower model complexity. Thus it can be seen that the OPWC-PLS has great prediction performance for GLB and its algorithm is convergent and rapid. The results provide important technical support for the rapid detection of serum.

Introduction

Near infrared (NIR) spectroscopy is a green and developing analytical technique, which has been widely used in life sciences,¹⁻⁷ agricultural products and food,⁸⁻¹¹ soil,¹²⁻¹⁴ and other fields.^{15,16} For NIR spectroscopic analysis of complex system, wavelength selection is necessary and difficult. So far, many methods including continuous mode and discrete mode of wavelength selection have been successfully used in NIR spectroscopy analysis, but a general and effective method has not been found. Moving window partial least squares (MW-PLS) is a widely used and well performed wavelength selection method, which uses a moving window whose position and size can be changed to identify and select continuous wavebands in terms of the prediction effect, and such waveband can correspond to absorption of specific functional groups.^{13,15,16-19} This method can achieve high prediction effect on most spectral data sets, so it often presents as the comparison method of new method to evaluate the performance of the new method. However, it can be seen from the papers [17-19], as a traversal algorithm for continuous wavebands, all possible continuous bands are screened, this method is time-consuming when encountering a large dataset. Monte Carlo uninformative variable elimination by PLS (MC-UVE-PLS) is a popular method for discrete wavelength selection,²⁰ which creatively introduced noise to eliminate uninformative variables, but it cannot achieve satisfactory prediction results for some data sets.

Serum globulin (GLB), which is synthesized by human monocyte-phagocyte system, contains various antibodies in the serum of healthy people, so it can enhance the body's resistance to prevent infection. It

is mainly used for immunodeficiency diseases as well as prevention and treatment of viral infections and bacterial infections such as infectious hepatitis, measles, chickenpox, mumps and herpes zoster. In addition, it can also be used in asthma, allergic rhinitis, eczema and other endogenous allergic diseases. Therefore, the GLB in human serum is very important for clinical trials and disease diagnosis. In previous studies,^{21,22} FTIR/ATR spectroscopy was used for determination of GLB. The study found that for blood index, the NIR has higher quantitative analysis accuracy than the FTIR/ATR spectroscopy.^{6,23} The experimental results show that the molecular absorption information of GLB can be captured by NIR spectroscopy without reagent.

Optimal partner wavelength combination (OPWC) is a method of selecting discrete information wavelength by iteration. For the method, the best partner of each wavelength in a predetermined wavelength region was determined based on binary linear regression (BLR), and a partner wavelength subset (PWS) was obtained; then the best partner of each wavelength in the PWS was obtained with the same method. The iterative process may be continued until convergence was met, and the last obtained wavelength subset was called OPWC. On the basis of the OPWC, PLS model was established. In order to make full use of the samples, the leave-one-out cross validation (LOOCV) was adopted.

Because human serum is a complex multi-component system and the absorption interference of other components is very complex, it is difficult to extract the characteristic information of GLB. Therefore, OPWC-PLS method was employed to remove redundant wavelength and establish a high precision quantitative model. MW-PLS and MC-UVE-PLS methods were also performed for comparison. Experimental results showed that the OPWC-PLS has great prediction performance and the algorithm is convergent and rapid.

Materials And Methods

Due to technical limitations, the Materials and Methods section is only available as a download in the supplementary files section.

Results And Discussion

3.1. Results with MW-PLS

The NIR spectra of 230 human serum samples in the scanning area (780-2498 nm) were shown in Figure 1. As can be seen from the figure, absorption at about 2000 nm and 2400 nm has obviously strong noise. In order to obtain satisfactory results, wavelength selection must be carried out to overcome noise interference. For comparison, PLS model of the full spectrum region was first established. The corresponding SECV and R_{pCV} were 1.423 g L⁻¹ and 0.935, respectively.

MW-PLS method was performed to optimize waveband and improve prediction accuracy. Depending on minimum SECV value, the optimal MW-PLS model was selected out. The corresponding waveband was

1504 to 1820 nm, located in the long-NIR region (1100 to 2498 nm). Prediction effects (SECV and R_{PCV}) and parameters of the above two methods were summarized in Table 2. The results showed that the predicted values were highly correlated with clinical measurements for the two methods, and comparing with optimal PLS model in full spectrum region, the optimal MW-PLS model achieved better prediction effect with fewer wavelengths.

3.2. Results with MC-UVE-PLS

The MC-UVE method was performed for eliminating the uninformative variables. Based on the parameter settings in section 2.4, 180 wavelengths were selected, and the SECV and R_{PCV} for the corresponding PLS models were 1.153 g L^{-1} and 0.948, respectively. Compared with the result of PLS in the full spectrum range, the prediction ability of this method was not significantly improved, which may be because it only eliminates non information variables without considering the influence of interference variables, while serum is a complex system with multiple interference variables.

3.3. Results with OPWC-PLS

The OPWC method was performed for screening information wavelength based on the steps mentioned in section 2.5. Firstly, 104 best partners for all 860 wavelengths were determined according to the results of LOOCV-BLR analysis, and $PWS^{(1)}$ with 104 wavelengths was obtained. Thus, the number of wavelengths was greatly reduced after the first projection. The correspondence between all 860 wavelengths and their best partners was shown in Figure 2. As shown in the figure, some wavelengths had the same best partner, such as the 2156 nm and 2190 nm as best partners of other wavelengths appeared 3 and 8 times, respectively, so projection was not a one-to-one mapping function in the whole spectral region. Obviously, was a subset of and the projection continues.

Based on the corresponding relationship determined above, the best partner of was easy to be selected, and the $PWS^{(2)}$ was obtained. Repeated the same process for $PWS^{(2)}$, and $PWS^{(3)}$ was obtained. As the projection progresses, the number of wavelengths decreased gradually until the number of wavelengths for $PWS^{(6)}$ no longer changed. The $PWS^{(6)}$ was the OPWC and it had only 28 wavelengths. Figure 3 showed the 28 wavelengths and their best partners. As the figure showed, the 28 wavelengths are divided into 14 groups, and the two wavelengths in each group are the best partners for each other.

Based on PLS, the LOOCVs were performed for every PWS, and the corresponding minimum SECV value and number of wavelengths ($N^{(s)}$) used are shown in Figure 4. As shown in the figure, the $N^{(s)}$ and minimum SECV values have almost the same trend. After the first projection, both of them decrease rapidly, and the remaining wavelengths are more important, so as the number of projections increases, they slowly decrease. This may be due to the removal of a large amount of noise and background information from the original spectrum after the first projection, so both the $N^{(s)}$ and minimum SECV values decrease rapidly. The partner wavelength subset of the original spectrum contains less redundant information, so the $N^{(s)}$ and minimum SECV values decrease slowly in the later projection iteration.

3.4. Comparison of OPWC-PLS and MW-PLS methods

Screening the information wavelengths of GLB in the human serum of a multi-component complex system is difficult and complicated. The wavelengths selected by the OPWC-PLS and MW-PLS methods, which correspond to the information of GLB, were shown in Figure 5. As indicated in Figure 5, the wavelengths selected by the OPWC method have a wider distribution range and partially coincides with the wavelengths selected by MW-PLS. This may be because the local characteristics of MW-PLS method make some wavelengths cannot be detected, which reflects the complexity of NIR model optimization and the commonness and difference of different methods.

Figure 6 showed the relationship between the predicted and measured GLB values based on the MW-PLS and OPWC-PLS methods, respectively. The prediction effect and corresponding parameters N and F were summarized in Table 2. The SECV and R_{pCV} were 0.813 g L^{-1} and 0.978 with OPWC-PLS, and 0.804 g L^{-1} and 0.979 with MW-PLS, respectively. The results show that, like MW-PLS, the prediction effect of OPWC-PLS was also obviously better than that of the whole spectrum PLS, and the OPWC is an effective method for screening wavelengths. The phenomenon conveys that better prediction results can be achieved with fewer wavelengths. Thus one can conclude that it is very necessary to first perform wavelength selection before building a calibration model. The two methods had achieved almost the same good prediction results (SECV and R_{pCV}). However, the optimal OPWC-PLS model adopted only 28 wavelengths, while the other adopted 159 wavelengths. Therefore, the OPWC method has great prediction performance for wavelength selection.

The differences in prediction of the OPWC-PLS and MW-PLS methods for GLB illustrate that MW-PLS can achieve higher prediction accuracy, but it is time-consuming and employs more wavelengths, while OPWC-PLS can achieve similar prediction results with MW-PLS in less time. In addition, MW-PLS, as a continuous wavelength screening method, is more suitable for determining the object with relatively concentrated molecular absorption bands; while OPWC-PLS, as a discrete wavelength screening method, may be more suitable for determining the object with relatively fragmented molecular absorption bands.

Conclusion

The change of GLB content in human serum has important reference value for clinical trial and disease diagnosis. In this study, the OPWC-PLS method was employed for rapid analysis of GLB based on NIR spectroscopy. MW-PLS and MC-UVE-PLS methods were also employed for comparison. The results indicate that, OPWC-PLS and MW-PLS methods achieved satisfactory prediction results, while the MC-UVE-PLS method was not suitable for the data set of this study, and the prediction effect of the model is not significantly improved. The optimal OPWC-PLS model adopted 28 wavelengths, and corresponding SECV and R_{pCV} were 0.813 g L^{-1} and 0.978, respectively. The optimal MW-PLS model adopted 159 wavelengths, and corresponding SECV and R_{pCV} were 0.804 g L^{-1} and 0.979, respectively. The OPWC-PLS achieved almost the same prediction effect as MW-PLS with faster speed and fewer wavelengths. Therefore, OPWC is an efficient approach for information wavelength selection.

The predicted GLB values obtained by MW-PLS and OPWC-PLS were highly correlated with the reference values. Compared with traditional method, the method based on NIR spectroscopy has the merits of rapidity, simplicity and no chemical reagent. Therefore, the results have important reference value for the rapid determination of GLB. In addition, the wavelengths selected by the two methods are partially the same, reflecting the commonness and difference of different methods.

List Of Abbreviations

GLB: Globulin; NIR: Near infrared; OPWC: Optimal partner wavelength combination; MW-PLS: Moving window partial least squares; MC-UVE: Monte Carlo uninformative variable elimination; SECV: Root-mean-square error of cross validation of prediction; R_{PCV} : Correlation coefficient of prediction; BLR: Binary linear regression; PWS: Partner wavelength subset; LOOCV: Leave-one-out cross validation; SD: Standard deviation.

Declarations

Availability of data and materials

The datasets used and/or analysed during the current study are available from the corresponding author on reasonable request.

Competing interests

The authors declare that they have no competing interests.

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Authors' contributions

HY analyzed the spectral data of human serum samples and optimized the wavelength model, and was a major contributor in writing the manuscript. ZY and ZHH carried out the spectrum experiment. KXS performed model validation. All authors read and approved the final manuscript.

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Tables

Table 1 Statistical analysis of measured GLB values of 230 samples

Indicator	Number	Min	Max	Mean	SD
GLB(g L ⁻¹)	230	18.70	41.60	27.47	73.95

Table 2 Prediction effects of three methods

Methods	Adopted wavelengths (nm)	N	F	SECVR _{PCV}
PLS	780-2498	860	151.42	30.935
MW-PLS	1504-1820	159	100.80	40.979
OPWC-PLS	1410, 1534, 1536, 1538, 1542, 1676, 1678, 1698, 1732, 1734, 1738, 1742, 1744, 1746, 1750, 1870, 2128, 2132, 2218, 2220, 2222, 2228, 2254, 2258, 2306, 2310, 2318, 2340	28	7	0.8130.978

Figures

Figure 1

NIR spectra of 230 human serum samples in the scanning area (780-2498 nm).

Figure 2

Best partners of 860 wavelengths in the full spectrum region.

Figure 3

Best partners of the selected 28 wavelengths.

Figure 4

Number of wavelengths and minimum SECV value for each projection.

Figure 5

Position of the selected wavelengths with MW-PLS and OPWC-PLS located the average spectrum.

Figure 6

Relationship between the predicted values and measured values of GLB based on (a) MW-PLS and (b) OPWC-PLS methods.

Supplementary Files

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