Mixture design for food packaging in a modified atmosphere

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Abstract: In the past, food packaging was primarily used to support product sales and protect food from contamination or environmental effects which would reduce the life of the food. According to the World Health Organization (WHO), in the United States, from 6.5 to 33 million cases of food borne diseases occur each year, and nearly 9000 of these result in death. For this reason, we use active and intelligent packaging to extend the shelf life of food and improve its quality. This article aims to create an optimisation scheme for an optimum blend preference in food packaging with an active atmosphere. This will be accomplished by setting criteria for the production of antimicrobial food packaging.

Key words: antimicrobial technology, shelf life of food, simplex lattice design, response optimiser

There has been significant development in active packaging systems over the past 20 years. Some packaging concepts belonging to this category were used previously, but in the last two decades, there has been systematic research. Two types of packaging systems with interactive functions are currently distinguished, as described below.

Intelligent packaging is used to monitor conditions in the surroundings of the packaged product, and thus provides information about the quality of the packed food during transportation and storage. Temperature indicators and indicators of internal composition of the atmosphere are currently commercially available, along with indicators for the measurement of freshness of the packaged product. A new category of intelligent packaging systems uses elements based on radio-frequency identification (RFID) technology.

Because the application portion of this paper deals with the optimisation of a three-component mixture in antimicrobial packaging, only a more detailed description of active packaging will be given below.

Active packaging materials are defined as materials that aim at extending the shelf life or maintaining or improving the conditions of packaged food (Hauser et al. 2014). They combine non-active barrier properties with the active function of extending products' shelf life and reducing the risk of pathogens (Hotchkiss 2002). The principal function of antimicrobial packaging is the release of antimicrobial substances onto the surface of the packaged foodstuff. Whilst agents from the packaging material slowly migrate to the product surface, the packaging can help to maintain high concentrations where necessary (Quintavalla and Vicini 2002). For this purpose, the active substances can either be incorporated directly in the packaging material or inserted in an additional active layer.

Active packaging is able to spontaneously change its properties in response to changes in conditions inside or outside the package. This is achieved via active package elimination of the negative effects of environmental conditions on the quality of the food product (e.g. extending the shelf life of food).

In recent years, a great deal of research has focused on traditional preservatives (Cuq 1997), including active substances like sorbic acid, which was added in wheat gluten and bees wax (Guillard et al. 2009) or polyvinyl acetate, and coated on low-density polyethylene (LDPE) film (Hauser and Wunderlich 2011). Potassium sorbate has also been incorporated directly in LDPE (Silveira et al. 2007). To avoid synthetic preservatives, antimicrobial packaging based on natural extracts from herbs like essential oils and oleoresins has been developed (Muriel-Galet et al. 2013). Although low concentrations of essential oil were found to be effective in in vitro studies, however, much higher concentrations were necessary in the food matrix to achieve comparable effects (Smid

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and Gorris 1999). In addition, the use of essential oils can have a great influence on the flavour of the preserved food (Burt 2004).

Maillard reaction products (MRPs) represent one possible form of antimicrobials. MRPs are created when carbonyl compounds (e.g. sugar) react with protein-bound amino acids. This can occur, for example, during the heat processing of bakery products or milk (Ledl and Schleicher 1990). In recent years, the antimicrobial effect of MRPs of mixture models (heated matter of amino acid plus reducing sugar) has been investigated. Thus, MRPs' antimicrobial activity against different type of bacteria was detected (e.g. *Salmonella typhimurium, Listeria monocytogenes,* Staphylococcus, etc.; Tauer et al. 2004; Ahmad et al. 2013).

Water content in food and the content of substances which are a source of energy (e.g. sugars, alcohols, amino acids) and nitrogen (proteins, amino acids) have a significant effect on the shelf life of food and the speed of reproduction of microorganisms. Generally, a food containing low molecular weight and a greater amount of water degrades faster, as low molecular substances metabolise microorganisms directly. Meanwhile, proteins and high molecular carbohydrates (starch, cellulose) must first be split into low molecular products by microbial exo-enzymes; following this, they are metabolised by endo-enzymes. The growth of microorganisms and their biochemical activity are strongly affected by pH in foods. Each microbial species can reproduce only in a certain pH range. For the optimal growth of most bacteria, the range is relatively narrow. Extreme pH can kill microorganisms.

pH has a value ranging from 0 to 14 and is defined as the negative logarithm of the concentration of oxonium cations: $pH = \log [H_3O+]$. Neutral pH is around pH 7; 'acidic' foods have lower values, while 'alkaline' foods have higher values. Solutions with a pH less than 7 7 pH has an influence on the reproduction of the bacteria in relation to the vitality and growth rate, as well as the intensity and character of the metabolism. The resistance of cells to increased temperatures is lower when there is increased deviation between the real and optimal pH values. This applies to both vegetative cells and spores. Most bacteria grow in a neutral or weakly alkaline pH (6.6 to 7.5).

Water activity, or a_w , is the partial vapour pressure of water (p) in a substance divided by the standard state partial vapour pressure of water (p_0) :

$$a_w = \frac{p}{p_0} \tag{1}$$

In the field of food science, the standard state is most often defined as the partial vapour pressure of pure water at the same temperature. Using this definition, pure distilled water has a water activity of exactly 1. As temperature increases, a_w typically increases, except in some products with crystalline salt or sugar. Pure water has an $a_w = 1.0$.

If the amount of water available for the microorganisms in the food is reduced, the value of aw < 1.0. The optimal value for most microorganisms aw > 0.98.

Substances with higher a_w tend to support more microorganisms. Bacteria usually require an a_w of at least 0.91, and fungi at least 0.7.

Reduction potential (Eh; also known as **redox potential**, **oxidation**) is a measure of the tendency of a chemical species to acquire electrons and thereby be reduced. It is measured in volts (V) or millivolts (mV). Each environment has a redox potential which given by the presence of oxidising agents (e.g. oxygen, nitrates, peroxides) or reducing agents (e.g. ferrous ions, hydrogen). The Eh value depends on the ratio of the oxidised and reduced substances, and is determined by the chemical composition of the food and the partial pressure of oxygen in the foodstuff.

MATERIALS AND METHODS

The term *experiment* is defined as the systematic procedure carried out under controlled conditions in order to discover an unknown effect, to test or establish a hypothesis or to illustrate a known effect. When analysing a process, experiments are often used to evaluate which process inputs have a significant impact on the process output, and what the target level of those inputs should be to achieve a desired result (output). Experiments can be designed in many different ways to collect this information (Birciakova et al. 2014). *Design of experiments* (DOE) is also referred to as *designed experiments* or *experimental design* – all of these terms have the same meaning.

Experimental design can be used at the point of greatest leverage to reduce design costs by speeding up the design process, reducing late engineering design changes and reducing product material and labour complexity (Stojanova and Tomsik 2014). Designed experiments are also powerful tools employed to

lower manufacturing costs by minimising process variation and reducing rework.

Mixture experiments comprise a special class of response surface experiments in which the product under investigation is made up of several components or ingredients (Mongomery 2012). Designs for these experiments are useful because many product design and development activities in industrial situations involve formulations or mixtures. In these situations, the response is a function of the proportions of the different ingredients in the mixture.

EXACT FORMULATION

Let $x = (x_1, x_2, ..., x_n)$ denote the vector of proportions of q mixing components and f(x) be the corresponding mean response. The factor space is simplex, given by:

$$Y = \left\{ x = (x_1, \dots, x_q) : x \ge 0; i = 1, 2, \dots, q; \sum_{i=1}^q x_i = 1 \right\} (2)$$

Scheffe introduced the following models in canonical forms of different degrees to represent the mean response function:

Linear:

$$f(x) = \sum_{i} \beta_{i} x_{i} \tag{3}$$

Quadratic:

$$f(x) = \sum_{i} \beta_{i} x_{i} \sum_{i < j} \beta_{ij} x_{i} x_{j}$$
(4)

In the above, we have used generic notations for the model parameters in different versions of mixture models. Using the identity $\sum x_i = 1$, model (4) can be converted to a canonical homogeneous quadratic model as follows:

$$f(x) = \sum_{i} \beta_{ii} x_i^2 \sum_{i < j} \beta_{ij} x_i x_j$$
(5)

In the simplex mixture experiment, the response (the quality or performance of the product based on some criterion) depends on the relative proportions of the components (ingredients) (Antony 2001). The amount of components, measured in weights, volumes or some other units, add up to a common total. In contrast, in a factorial design, the response varies depending on the amount of each factor.

Before carrying out our experiments, we needed to determine what design will be most appropriate for the experiment. The theoretical apparatus provides simplex centroid, simplex lattice and extreme vertices designs.

After we choose a design, we needed to carry out the following steps:

- Identify the components, process variables and mixture amounts are of interest;
- (2) Determine the model;
- (3) Ensure adequate coverage of the experimental region of interest; and
- (4) Determine the impact that other considerations have on the selected design.

The default data for the experiment are listed in Table 1. Here, we chose a mixture design.

We used the following contour plot to help you visualise the response surface. Contour plots are useful for establishing desirable response values, mixture blends and operating conditions.

StdOrder	RunOrde	PtTyp	Blocks	x1 (N2)	x2 (CO2)	x3 (O2)	Y (colony count)
10	1	-1	1	0.16667	0.16667	0.6667	4900
2	2	2	1	0.50000	0.50000	0.00000	6300
9	3	-1	1	0.16667	0.66667	0.16667	5100
1	4	1	1	1.00000	0.00000	0.00000	8400
5	5	2	1	0.00000	0.50000	0.50000	4800
4	6	1	1	0.00000	1.00000	0.00000	8200
6	7	1	1	0.00000	0.00000	1.00000	7800
7	8	0	1	0.33333	0.33333	0.33333	4900
8	9	-1	1	0.66667	0.16667	0.16667	6500
3	10	2	1	0.50000	0.00000	0.50000	6600

Table 1. Default data of randomised trials of mixture design

Source: own calculation



Figure 1. Mixture contour plot for modified atmosphere packaging

Source: own calculation

The contour plot in Figure 1 shows how a response variable (colony count) relates to three components (the percentage of components of the modified at-



Figure 2. Result of the simplex lattice design for modified atmosphere packaging

Source: own calculation

mosphere, which are N2, CO2 and O2) based on a model equation. Points which exhibit the same response are connected to produce the contour lines

Table 2. Regression model for mixtures: Y (colony count versus x1 [N₂], x2 [CO₂], x3 [O₂])

Estimated regression co	efficients for Y (cold	ony count: cfu/obje	ct) (component pi	coportions)	
Term	Coef	SE	Coef	Т	Р
VIF					
x1 (N ₂)	8 548	378.2	*	*	1.964
x2 (CO ₂)	8 148	378.2	*	*	1.964
x3 (O ₂)	7 702	378.2	*	*	1.964
x1 (N ₂)*x2 (CO ₂)	-8 595	1 743.3	-4.93	0.008	1.982
x1 (N ₂)*x3 (O ₂)	-6 686	1 743.3	-3.84	0.019	1.982
x2 (CO ₂)*x3 (O ₂)	-13 886	1 743.3	-7.97	0.001	1.982
S = 392,187 PRESS =	9 926 598				

R-Sq = 96.58% R-Sq(pred) = 44.81% R-Sq(adj) = 92.30%

Analysis of variance for Y (colony count: cfu/object) (component proportions)

DF	Seq SS	Adj SS	Adj MS	F	Р
5	17 369 756	17 369 756	3 473 951	22.59	0.005
2	$1\ 774\ 444$	393 515	196 758	1.28	0.372
3	15 595 311	15 595 311	5 198 37	33.80	0.003
1	3 630 911	3 738 979	3 738 979	24.31	0.008
1	2 205 254	2 262 459	2 262 459	14.71	0.019
1	9 759 146	9 759 146	9 759 146	63.45	0.001
4	615 244	615 244	153 811		
9	17 985 000				
	DF 5 2 3 1 1 1 4 9	DFSeq SS517 369 75621 774 444315 595 31113 630 91112 205 25419 759 1464615 244917 985 000	DFSeq SSAdj SS517 369 75617 369 75621 774 444393 515315 595 31115 595 31113 630 9113 738 97912 205 2542 262 45919 759 1469 759 1464615 244615 244917 985 000	DFSeq SSAdj SSAdj MS517 369 75617 369 7563 473 95121 774 444393 515196 758315 595 31115 595 3115 198 3713 630 9113 738 9793 738 97912 205 2542 262 4592 262 45919 759 1469 759 1469 759 146917 985 000500500	DFSeq SSAdj SSAdj MSF517 369 75617 369 7563 473 95122.5921 774 444393 515196 7581.28315 595 31115 595 3115 198 3733.8013 630 9113 738 9793 738 97924.3112 205 2542 262 4592 262 45914.7119 759 1469 759 1469 759 14663.454615 244615 244153 811513 811

Source: own calculation

Table 3. Results	s from the \circ	optimiser (of microorganisms [*]	<i>colony count/object</i>
		1	0	

Parameters							
Y (colony co Minimum	Goal	Lower	Target	Upper	Weight	Import	
	4000	4000		6000	1	1	
Global solution							
Components							
$x1(N_2) = 0.0101010$							
$x2 (CO_2) = 0.479951$							
$x3 (O_2) = 0.509948$							
Predicted responses							
Y (colony co = 4449.77, Co	mposite desird	ability = 0.77511	15				

Source: own calculation

of constant responses. Because a contour plot only shows three components at a time, whilst holding any other components and process variables at a constant level, contour plots are only valid for fixed levels of the extra variables. If the holding levels are changed, the response surface of the colony count changes as well, sometimes drastically.

We also added new experiment data to refine the solution area. For this purpose, we used a simplex lattice design, as illustrated by Figure 2.

Both plots show how the component proportions are related to the colony count of microorganisms/ objects. To minimise the colony count of bacteria, we would choose proportions for the components in the lower centre of the design space where the colony count ratings are the lowest. In both plots, the lightest green contour represents the lowest in the



Figure 3. Result of the binary form of the design for modified atmosphere packaging

Source: own calculation

design space. The blend which produces the lowest colony count rating is at the vertex, which comprises of the following:

These values are from the interval of admissible values (i.e. the percentages relate to coded values of the mixed factors). The regression model is shown in Table 2. Result of the binary form of the design is shown in Figure 3.

The results of the analysis are shown in Table 3. For the significance test, it was decided to select significance levels of a = 5% (0.05). If the p-value was less than the significance level (0.05), the factor or interaction effect was then regarded to be statistically significant. For the present experiment, the main effects of N₂, CO₂ and O₂ were statistically significant. The calculated effect factor in the coded values (response factor changing from -1 to +1) is shown in the first column of Table 3. The second column represents the regression coefficient (i.e. a half effect of each factor). Table 3 also shows that the quadratic model is more suitable than a linear model. The regression model therefore estimates the predicted response: Y (colony) = 4449.77 cfu/object. We used overlaid contour plots in binary form to jointly evaluate multiple responses. Overlaid contour plots can help to identify component settings which optimise a single response or set of responses. This plot showed us the function membership of a feasible set of responses in binary form (the range of responses was from 4000 to 6000 cfu/object).

The optimum outcome provided by the response optimiser of microorganisms' *colony count/object*



Figure 4. Response optimiser of microorganisms' colony count/object

Source: own calculation

was very close to what was determined by visual inspection of the contour and surface plot. Table 3 shows that responses from the optimiser of microorganisms' colony count/object have the limitations when it comes to seeking the best possible result.

The proportions of components must be selected in such a manner that they sum to 1. We compared the plots showing concentrations of N_2 , CO_2 and O_2 in relation to temperature to see which level of the process variable resulted in a low count of colony microorganisms. The microorganism's colony count/ object ratings in the lower centre of the design space and the optimum response belonged to the following



Figure 5. Response trace plot for a modified atmosphere of food packaging

Source: own calculation

settings of modified atmosphere: low concentration of N_2 and middle concentrations of CO_2 and O_2 (Figure 4).

Figure 5 shows a response trace plot (also called a component effects plot) which illustrates how each component affects the response relative to a reference blend. If the design contains process or amount variables, they must be held at a fixed level.

CONCLUSION

Food packaging data obtained under a modified atmosphere had three components and one process variable. Previous investigation showed that the microorganism colony count/object was the best when x1 (N₂) was set at 10.1%. Therefore, CO₂ was fixed at 48% (low level) and O₂ at 50% (medium level). For the modified atmosphere data, the reference blend was the centre point. The trace plot provided the information below about the component effects.

Starting at the location corresponding to the reference blend:

As the proportion of N_2 (black curve) in the mixture:

- Increased (and the other mixture components decreased), the microorganisms' colony count/object rating decreased;
- Decreased (and the other mixture components increased), the microorganisms' colony rating increases.

As the proportion of CO_2 (red curve) in the mixture:

Increased (and the other mixture components decreased), the microorganisms' colony count/

object rating decreased rapidly. This was valid in proportion from a 0.1 value of deviation from the reference blend;

 Decreased (and the other mixture components increased), the microorganisms' colony count/ object rating increased rapidly. This was valid in proportion from a 0.1 value of deviation from the reference blend.

As the proportion of O_2 (green curve) in the mixture:

- Increased (and the other mixture components decreased), the microorganisms' colony count/ object rating decreased rapidly. This was valid in proportion from a 0.2 value of deviation from the reference blend;
- Decreased (and the other mixture components increased), the microorganisms' colony count/object rating increased rapidly. This was valid in proportion from a 0.2 value of deviation from reference blend.

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