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Physicochemical and Textural Optimization of Plant-Based Cheese Using Lentil–Gluten Protein Systems

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2 **Lentil–Gluten Protein Systems**

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8  
9  
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# 1 Physicochemical and Textural Optimization of Plant-Based Cheese Using

## 2 Lentil–Gluten Protein Systems

### 4 Abstract

5 The rising demand for sustainable foods has accelerated interest in plant-based cheese;  
6 however, replacing casein with plant proteins often compromises texture, meltability, and  
7 overall quality. This study investigated lentil–gluten protein systems in dairy-free cheese  
8 formulations and evaluated the effects of protein, modified tapioca starch, chitosan,  
9 maltodextrin, and oil using response surface methodology (RSM). A central composite design  
10 with 50 formulations was used to model effects on moisture, protein, fat, ash, color, meltability,  
11 and texture properties. Functional characterization showed that modified tapioca starch  
12 exhibited the highest swelling capacity (9.21 mL/g) and water retention capacity (WRC) (10.03  
13 g/g), while lentil protein showed the greatest water solubility index (WSI) (9.47%), and oil-  
14 holding capacity (OHC) (0.61 g/g). Linear models accurately described trends in moisture, ash,  
15 fat, and protein contents ( $R^2 = 0.95\text{--}0.99$ ). Protein ( $p < 0.01$ ), chitosan ( $p < 0.05$ ), and starch ( $p$   
16  $< 0.05$ ) significantly increased hardness, cohesiveness, and springiness, whereas oil enhanced  
17 meltability ( $p < 0.01$ ). Higher protein levels decreased lightness ( $L^*$ ) but increased  $a^*$  and  $b^*$   
18 values, reflecting natural pigments and Maillard reaction products. The optimized  
19 formulation—28.31% protein powder, 20% oil, 1.5% chitosan, 9.99% starch, and 2.03%  
20 maltodextrin—achieved high overall desirability and closely matched predicted values,  
21 confirming model validity. Overall, this study provides the first integrated quantitative  
22 assessment of lentil–gluten systems in plant-based cheese and identifies ingredient interactions  
23 that govern structure and melting behavior. The findings establish a data-driven formulation  
24 framework capable of producing high-protein, texturally stable dairy-free cheese with  
25 improved functional and sensory potential.

26

27 **Keywords:** Plant-based cheese; Lentil protein; Gluten protein; Response surface methodology;  
28 Texture; Ingredient interactions

29

## 30 **1. Introduction**

31 A transition from meat and dairy to plant-based diets has been prioritized as a strategy to meet  
32 the nutritional demands of a growing global population while reducing greenhouse gas  
33 emissions and mitigating the environmental impact of animal-based food production,  
34 particularly cheese (Grasso, Roos, Crowley, Arendt, & O'Mahony, 2021). Conventional  
35 cheese, produced from milk, starter cultures, salt, and enzymes through fermentation, remains  
36 one of the most widely consumed and nutrient-rich foods. Its popularity continues to grow at  
37 ~3% annually, largely due to its high protein, fat, and vitamin content (D. Zhang et al., 2024).

38 A number of plant-based food products—developed from ingredients such as nuts, grains, and  
39 legumes—have been designed to replicate the appearance, flavor, and texture of animal-based  
40 foods (Alehosseini, McSweeney, & Miao, 2025b; Boehm, Nicholson, & Baier, 2023). These  
41 alternatives can be rich in essential amino acids, phenolics, vitamins, minerals, and fiber, and  
42 may be consumed for health, environmental, ethical, and dietary reasons (McClements &  
43 Grossmann, 2023). The global plant-based food market is projected to reach \$113 billion by  
44 2031, positioning plant proteins from cereals, pulses, nuts, oilseeds, and pseudocereals as  
45 sustainable and cost-effective dairy replacements (Pereira et al., 2024; D. Zhang et al., 2024).  
46 However, many plant proteins exhibit limited solubility, restricting their suitability for cheese  
47 production (Wouters, Rombouts, Fierens, Brijs, & Delcour, 2016). Consumers may choose  
48 plant-based cheese due to lactose intolerance, milk allergy, or lifestyle choice, further driving  
49 innovation in this field (Aydar, Tutuncu, & Ozcelik, 2020).

50 Cheese analogues—products in which dairy proteins are partially or entirely substituted with  
51 plant-based ingredients—offer advantages such as shorter processing time, reduced cost, and  
52 nutritional flexibility (Kamath, Basak, & Gokhale, 2022). However, replacing casein with plant  
53 proteins often compromises texture and sensory quality, underscoring the need for a better  
54 understanding of ingredient functionality to improve dairy-free cheese formulations (Kamath  
55 et al., 2022; Masotti, Cattaneo, Stuknytė, & De Noni, 2018).

56 Among plant proteins, gluten and lentils hold particular promise for cheese applications  
57 (Naeem, Akhtar, Akram, Suleria, & Khalid, 2024; D. Zhang et al., 2024). Gluten, though  
58 largely insoluble in water, contains gliadins ( $\omega 5$ -,  $\omega 1,2$ -,  $\alpha$ -,  $\gamma$ -) and glutenins, which provide  
59 desirable cheese-like characteristics including water and fat absorption, viscoelasticity,  
60 emulsification, and foaming (D. Zhang et al., 2024). Lentil protein—an underutilized source  
61 rich in essential amino acids—contains 11S and 7S globulins similar to those in soy and  
62 exhibits valuable functional properties (Tang, Roos, & Miao, 2024). Legumes in general are  
63 nutrient-dense, providing proteins, vitamins, and minerals, although they are limited in sulfur-  
64 containing amino acids. This limitation is often addressed by combining complementary  
65 proteins (Alehosseini, McSweeney, & Miao, 2025a; Kovačević, Bechtold, & Pham, 2024).  
66 Lentils, in particular, are low in saturated fat and calories while offering high-quality protein,  
67 making them both nutritionally and environmentally sustainable (Kussmann, Abe Cunha, &  
68 Berciano, 2023; Naeem et al., 2024).

69 In dairy-free cheese, protein gelation is critical to texture formation. Heat treatment induces  
70 denaturation, aggregation, and network formation, enabling plant globulins to behave similarly  
71 to dairy proteins. These processes are influenced by pH and protein composition, which  
72 determine hydrophobic interactions, disulfide bonding, and hydrogen bonding within the gel  
73 network (Ge et al., 2023; Nicolai, 2019; Tang et al., 2024).

74 Despite growing consumer demand, many plant-based cheeses remain nutritionally  
75 unbalanced, with lower protein and higher saturated fat contents than conventional cheese,  
76 increasing the risk of deficiencies in nutrients such as calcium and vitamin B2 (Taeger &  
77 Thiele, 2024).

78 To date, no published study has systematically investigated the combined application of lentil  
79 and gluten proteins within dairy-free cheese matrices, despite their complementary structural  
80 and functional attributes (Naeem et al., 2024; D. Zhang et al., 2024). Given that lentil globulins  
81 form heat-induced particulate networks with high water-binding capacity, while gluten proteins  
82 develop extensible viscoelastic structures through gliadin-mediated flow and glutenin-driven  
83 disulfide crosslinking, their co-assembly offers a mechanistically grounded strategy for  
84 improving matrix cohesion, hydration stability, and fat entrapment in plant-based cheese  
85 systems (Chompoorat, Fasasi, Lavine, & Rayas-Duarte, 2022; Jarpa-Parra, 2017).

86 In this study, a novel lentil (legume-derived)–gluten (cereal-derived) protein system was  
87 formulated to leverage the complementary gelation pathways, hydration dynamics, as well as  
88 nutritional and functional properties of these proteins to enhance the physicochemical and  
89 textural properties of plant-based cheese. The research first evaluated the functional  
90 characteristics of the individual proteins and starch, followed by systematic optimization of  
91 cheese formulations using response surface methodology (RSM) to model the interactive  
92 effects of protein, chitosan, starch, maltodextrin, and oil concentrations on product quality. The  
93 specific objectives were to: (i) assess the swelling capacity, solubility index, and oil-holding  
94 capacity (OHC) of key ingredients; (ii) formulate and optimize lentil–gluten cheese alternatives  
95 using RSM; (iii) characterize the optimized formulation in terms of composition, color,  
96 meltability, and texture; and (iv) elucidate ingredient interactions that govern structural and  
97 functional performance. This integrated approach provides mechanistic insights into ingredient  
98 interactions and establishes a framework for the rational design of nutritionally balanced,

99 structurally robust, and sensory-appealing dairy-free cheese alternatives. The study's main  
100 strength lies in its innovative combination of complementary plant proteins and the application  
101 of RSM for data-driven formulation optimization. However, the work was conducted at  
102 laboratory scale and focused on instrumental analyses; storage stability was not evaluated.  
103 Future investigations should therefore include shelf-life assessments to substantiate the  
104 practical and commercial relevance of the developed formulations.

105

## 106 **2. Materials and methods**

### 107 **2.1. Materials**

108 The lentil protein powder (~81% protein) and modified tapioca starch were provided by HSN  
109 (Albolote, Spain) and 4mular (San Francisco, CA, US), respectively. Gluten from wheat (CAS  
110 Number: 8002-80-0,  $\geq 75\%$  protein), chitosan (low molecular weight, CAS Number: 9012-76-  
111 4,  $\geq 75\%$  deacetylated), and maltodextrin (CAS Number: 9050-36-6) were also supplied by  
112 Sigma-Aldrich. Moreover, coconut oil, Italian organic high oleic sunflower oil, acetic acid  
113 (CAS Number: 64-19-7, 99-100% purity), and salt were purchased from Coco loco (Naas,  
114 Ireland), Benvolio (Treviso, Italy), Lennox laboratory supplies ltd (Dublin, Ireland), and local  
115 supermarket (Fermoy, Ireland).

116

### 117 **2.2. Functional analysis of plant proteins and starch**

#### 118 **2.2.1. Swelling capacity**

119 The swelling capacity of protein and starch samples was determined following Talens et al.  
120 (2022) with some modifications. Briefly, 0.2 g of each sample was placed in a graduated  
121 cylinder containing 10 mL of distilled water at 25 °C. The sample was gently stirred to ensure  
122 uniform dispersion, then covered and left undisturbed at 25 °C for 18 h to allow full hydration  
123 and equilibration. After this period, the volume occupied by the swollen sample was recorded.

124 The swelling capacity was calculated using Equation 1 (Figure 1A). Measurements were  
 125 performed in triplicate and mean values are reported.

126

$$\text{Swelling Capacity (mL/g)} = \frac{\text{Volume occupied by sample}}{\text{Original sample dry mass}} \quad (1)$$

127

128

INSERT FIGURE 1 ABOUT HERE

129

### 130 **2.2.2. Water retention capacity (WRC)**

131 The WRC of the samples was determined according to Robertson et al. (2000). Briefly, 3 g of  
 132 sample was hydrated in 30 mL of distilled water at 25 °C and allowed to equilibrate for 18 h.

133 Following hydration, the samples were centrifuged at 3,000 × g for 20 min (SCRVALL LYNX

134 6000, Thermo Scientific, Waltham, MA, USA). The supernatant was carefully decanted by

135 inverting the tube, allowing the pellet to drain in place. The WRC was calculated using

136 Equation 2 (Figure 1B). All measurements were performed in triplicate and mean values are

137 reported.

138

$$\text{WRC (g/g)} = \frac{\text{Residue fresh mass} - \text{Residue dry mass}}{\text{Residue dry mass}} \quad (2)$$

139

### 140 **2.2.3. Water solubility index (WSI) and oil-holding capacity (OHC)**

141 To determine the WSI of the proteins (Equation 3), 2 g of each sample was dispersed in 100

142 mL of distilled water and heated in a water bath at 80 °C for 30 min. The mixture was then

143 centrifuged at 1100 × g for 10 min at 25 °C. The supernatant was carefully collected and dried

144 in an oven (FD 53, BINDER, Tuttlingen, Germany) at 103 ± 2 °C (Figure 1C) (Talens et al.,

145 2022).

146

$$\text{WSI (\%)} = \frac{\text{Mass of dissolved solid in supernatant}}{\text{Mass of dry solids}} \times 100 \quad (3)$$

147

148 The OHC was determined according to Garcia-Fontanals, Llorente, Valderrama, Bravo, and  
 149 Talens (2023) using Equation 4. Briefly, 1 g of the sample was mixed with 10 mL of sunflower  
 150 oil using a vortex mixer (FINEVORTEX, FINEPCR Co., Ltd. Gunpo-si, South Korea) for 30  
 151 min, then centrifuged at  $8000 \times g$  for 20 min. After centrifugation, the supernatant was decanted  
 152 and the remaining pellet was weighed (Figure 1D).

153

$$\text{OHC (g/g)} = \frac{\text{Mass of sample after oil absorption} - \text{Dry sample mass}}{\text{Dry sample mass}} \quad (4)$$

154

155 All measurements were performed in triplicate and mean values are reported.

156

### 157 **2.3. Formulation of dairy-free cheese alternatives**

158 The dairy-free cheese alternatives were formulated according to the procedure of by Dobson  
 159 and Marangoni (2023) with some modifications. All formulations corresponding to the 50  
 160 experimental runs of the central composite design were prepared under identical processing  
 161 conditions. A 5% (w/w) protein solution was prepared by dissolving the plant-based protein  
 162 blend (lentil protein 66.7 % and gluten 33.3 %) in the aqueous phase of deionized water (pH  
 163 5.1-5.4). The mixture was stirred at 400 rpm for 10 min using a magnetic stirrer (IKA<sup>®</sup> RET  
 164 basic, IKA-Werke GmbH & Co. KG, Staufen im Breisgau, Germany) to ensure proper protein  
 165 dispersion. The coconut oil was liquefied by heating to approximately 35-40 °C, mixed with  
 166 high oleic sunflower oil at a ratio of 75:25, and incorporated into the protein solution under  
 167 homogenization at 10,000 rpm for 5 min by an Ultra Turrax (IKA<sup>®</sup> T25, IKA-Werke GmbH &

168 Co. KG, Staufen im Breisgau, Germany). The emulsion was then combined with the remaining  
 169 dry ingredients—protein, modified tapioca starch, chitosan, maltodextrin, and salt—and mixed  
 170 until no visible clumps remained. The mixture was heated in a temperature-controlled water  
 171 bath (Medical Supply Co. Ltd, Dublin, Ireland) at 80 °C for 3 min, followed by blending at low  
 172 speed for 2 min in a laboratory blender (Waring Commercial, McCConnellsburg, PA, USA). The  
 173 heated mixture was maintained at 40 °C with gentle stirring for 10 min, poured into molds, and  
 174 stored at 4 °C for 24 h prior to analysis. Across all trials, the mean formulation yield was 96.92  
 175  $\pm$  2.86 %, calculated as the ratio of recovered solidified product mass to the initial formulation  
 176 mass. Ingredient ranges used in the dairy-free cheese alternatives are summarized in Table 1.

177  
 178 **Table 1.** Ranges of Ingredients Used in Dairy-Free Cheese Formulations

Ingredients	(%)
Lentil protein (66.7%) + gluten (33.3%)	(20, 25, and 30)
Coconut oil (75%) + sunflower oil (25%)	(10, 15, and 20)
Chitosan	(0.5, 1, and 1.5)
Modified tapioca starch	(5, 7.5, and 10)
Maltodextrin	(2, 3, and 4)
Salt	1

179

#### 180 **2.4. Moisture, ash, fat, and protein content**

181 The moisture and ash contents of dairy-free cheese alternatives were analyzed using a TGA-  
 182 701 thermogravimetric analyzer (LECO Instruments, St. Joseph, MI, USA). The total protein  
 183 of the samples was determined using an FP628 nitrogen analyzer (LECO, St. Joseph, MI, USA)  
 184 with a nitrogen to protein conversion factor of 6.25. Additionally, the fat content was  
 185 determined by SMART 6 + ORACLE systems (CEM Corporation, Matthews, NC, USA) (Fan  
 186 et al., 2023; Tang, Roos, & Miao, 2023).

187

#### 188 **2.5. Color profile determination of dairy-free cheese alternatives**

189 The colorimeter analysis of samples carried out by using a digital colorimeter (Chroma Meter  
 190 CR-400, Konica Minolta Sensing, Osaka, Korea) based on the CIE  $L^*a^*b^*$  system. The values  
 191 of  $L^*$  (lightness),  $a^*$  (redness/greenness), and  $b^*$  (yellowness/blueness) were determined.  
 192 Additionally, the parameters  $\Delta E$ ,  $WI$ , and  $C_{ab}^*$  were calculated using Equations 5–7,  
 193 respectively (A. Alehosseini et al., 2022; Garcia-Fontanals et al., 2023; Naeem et al., 2024).

194

$$\Delta E = \sqrt{(\Delta L^*)^2 + (\Delta a^*)^2 + (\Delta b^*)^2} \quad (5)$$

$$WI = 100 - \sqrt{(100 - L^*)^2 + a^{*2} + b^{*2}} \quad (6)$$

$$C_{ab}^* = \sqrt{a^{*2} + b^{*2}} \quad (7)$$

195

## 196 **2.6. Meltability**

197 A cylindrical dairy-free cheese alternatives (14 mm in height and 25 mm in diameter) were  
 198 stored in the refrigerator at 4 °C for 10 min, and then transferred to the oven (FD 53, BINDER,  
 199 Tuttlingen, Germany) preheated to 232 °C for 5 min. After removal, the samples were allowed  
 200 to cool for approximately 30 min before measuring the spread diameter at four different angles.  
 201 A paper template with a 100 mm diameter was printed, featuring concentric circles at 5 mm  
 202 intervals and lines at 45° angles, which were placed at the bottom of each glass plate, facing  
 203 upwards. The meltability was calculated by determining the percentage increase in the average  
 204 diameter from the initial diameter (Dobson & Marangoni, 2023).

205

## 206 **2.7. Texture profile analysis (TPA)**

207 The texture properties of dairy-free cheese alternatives were evaluated using a Texture  
 208 Analyser TA.XT.Plus (Stable Micro Systems, Godalming, UK). After 24 h of storage at 4 °C,  
 209 cylindrical samples (14 mm in height and 25 mm in diameter) were compressed at room  
 210 temperature to 30% of their original height in a double compression with a SMS P/75R probe

211 at a fixed speed of 1.0 mm/s. For each sample, textural parameters, including hardness,  
 212 cohesiveness, springiness, adhesiveness, and resilience, were calculated by the software  
 213 Texture Exponent (Version 6.2, Stable Micro Systems, Godalming, UK) (Grasso, Roos,  
 214 Crowley, & O'Mahony, 2024; Monga, Dev, & Singhal, 2022).

215

## 216 **2.8. Statistical analysis**

217 For the functional properties of proteins and starch, all analyses were performed in triplicate.  
 218 Mean comparisons were conducted using analysis of variance (ANOVA), with significance  
 219 determined at  $P < 0.05$ . Post hoc multiple comparisons were assessed using Tukey's test in  
 220 JMP® Pro software (Version 18.0.1, SAS Institute Inc, Cary, NC, USA). For dairy-free cheese  
 221 alternatives, the experimental design and analysis were performed using central composite  
 222 design (CCD) of RSM with 50 runs and 8 replications at the central point through ANOVA for  
 223 a statistical significance  $P < 0.05$ , by using Design Expert software (Version 23, State-Ease,  
 224 Minneapolis, MN, USA). In this study, five independent variables—the amounts of plant-based  
 225 proteins, oil, chitosan, modified tapioca starch, and maltodextrin—were considered with three  
 226 levels. The selected variable ranges were determined based on pre-tests and previous research.  
 227 For each point, moisture, ash, fat, protein, color parameters, meltability, and texture properties  
 228 of the dairy-free cheese alternatives were determined as responses. A second order polynomial  
 229 equation was also applied to determine responses (Equation 8).

230

$$\begin{aligned}
 Y = & a_1X_1 + a_2X_2 + a_3X_3 + a_4X_4 + a_5X_5 + a_{11}X_1^2 + a_{22}X_2^2 + a_{33}X_3^2 + a_{44}X_4^2 \\
 & + a_{55}X_5^2 + a_{12}X_1X_2 + a_{13}X_1X_3 + a_{14}X_1X_4 + a_{15}X_1X_5 + a_{23}X_2X_3 \\
 & + a_{24}X_2X_4 + a_{25}X_2X_5 + a_{34}X_3X_4 + a_{35}X_3X_5 + a_{45}X_4X_5 + a_0
 \end{aligned} \tag{8}$$

231

232 Where  $a_0$  is a constant and  $a_i$ ,  $a_{ii}$ , and  $a_{ij}$  are also the linear, quadratic, and interactive  
233 coefficients, and  $X_1$ ,  $X_2$ ,  $X_3$ ,  $X_4$ , and  $X_5$  are the protein powder, oil, chitosan, starch, and  
234 maltodextrin concentrations. For formulation optimization using RSM, fat content and  
235 hardness were set to be maximized, with pure protein content fixed at approximately 25%, and  
236 all other responses constrained within acceptable ranges.

237

### 238 **3. Results and discussion**

#### 239 **3.1. Functional analysis of plant proteins and starch**

240 Swelling capacity is strongly influenced by the structural characteristics and composition of  
241 biopolymers (Hoover, 2001). As shown in Table 2, the swelling capacities of modified tapioca  
242 starch, gluten, and lentil protein were 9.21, 5.32, and 4.83 mL/g, respectively. The superior  
243 swelling of modified starch likely results from structural alterations (e.g., cross-linking,  
244 substitution, or depolymerization) that disrupt the native granular organization and increase the  
245 exposure of hydrophilic hydroxyl groups. This enhances water penetration and retention within  
246 amorphous regions of starch granules, leading to pronounced swelling (BeMiller, 2018;  
247 Hoover, 2001). By contrast, proteins such as gluten and lentil protein possess compact tertiary  
248 and quaternary structures stabilized by hydrophobic interactions, disulfide bonds, and  
249 hydrogen bonding. These conformations restrict water accessibility, thereby limiting swelling.  
250 Gluten, composed mainly of glutenin and gliadin, forms an elastic, cohesive network upon  
251 hydration that resists expansion, while lentil protein, though slightly more soluble, remains a  
252 globular protein with constrained swelling compared to polysaccharides like starch  
253 (Damodaran, Parkin, & Fennema, 2007; Jarpa-Parra, 2017; Wieser, 2007).

254

255 **Table 2.** Swelling capacity, water retention capacity (WRC), water solubility index (WSI), and oil-  
256 holding capacity (OHC)

Sample	Swelling capacity (mL/g)	WRC (g/g)	WSI (%)	OHC (g/g)
Gluten	5.32 <sup>b</sup> ± 0.31	1.50 <sup>c</sup> ± 0.02	2.32 <sup>b</sup> ± 0.41	0.54 <sup>b</sup> ± 0.01
Lentil protein	4.83 <sup>b</sup> ± 0.06	2.98 <sup>b</sup> ± 0.11	9.47 <sup>a</sup> ± 0.02	0.61 <sup>a</sup> ± 0.06
Modified tapioca starch	9.21 <sup>a</sup> ± 0.87	10.03 <sup>a</sup> ± 0.03	–	–

257

258 These differences in swelling behavior have direct implications for cheese-matrix  
 259 development. Modified starch, with its high swelling capacity, contributes to the formation of  
 260 a continuous, hydrated phase capable of controlling serum separation and enhancing gel  
 261 firmness—effects widely documented for physically or chemically modified starches used in  
 262 processed cheese systems. In contrast, the modest swelling observed for lentil and gluten  
 263 proteins suggests their role is less water immobilization and more network formation, where  
 264 protein–protein contacts (non-covalent interactions for lentil globulins and disulfide-mediated  
 265 elasticity for gluten) provide structural reinforcement rather than bulk hydration. This  
 266 distinction highlights why combining starch with complementary proteins is mechanistically  
 267 advantageous in plant-based cheese design: each biopolymer contributes unique functional  
 268 attributes that together regulate water distribution, porosity, and viscoelasticity (Dobson,  
 269 Laredo, & Marangoni, 2022; Jarpa-Parra, 2017; Trivedi et al., 2008; Wieser, 2007).

270 Lentil protein exhibited a significantly higher WRC than gluten, likely due to the presence of  
 271 residual non-protein components in lentil protein, including dietary fibers and non-gluten  
 272 polysaccharides, which provide additional hydrophilic binding sites. Moreover, the globular  
 273 structure of lentil protein may expose polar amino acid residues during isolation, enhancing its  
 274 ability to bind water (Boye, Zare, & Pletch, 2010; Karaca, Low, & Nickerson, 2011). The  
 275 higher WRC of lentil protein can also be attributed to its greater content of charged amino acids  
 276 and water-accessible side chains, which increase water binding compared to gluten. Lentil  
 277 globulins undergo partial unfolding during extraction, exposing hydrophilic residues that  
 278 increase water uptake and capillary-driven retention within the protein matrix (Jarpa-Parra et

279 al., 2014). In contrast, gluten's polymeric glutenin network is stabilized by extensive  
280 intermolecular disulfide bonds that reduce segmental mobility and limit the formation of water-  
281 binding cavities (Wieser, 2007). These mechanistic differences explain the superior water-  
282 holding performance of lentil protein and indicate that lentil protein may help mitigate  
283 syneresis tendencies in dairy-free cheese formulations—an observation supported by previous  
284 work with legume-based gels (Boye, Aksay, et al., 2010; Lam, Can Karaca, Tyler, &  
285 Nickerson, 2018; Stone, Karalash, Tyler, Warkentin, & Nickerson, 2015).

286 The WSI, defined as the fraction of protein dissolved in water or buffer, was 2.32% for gluten  
287 and 9.47% for lentil protein (Table 2), indicating the latter's significantly greater solubility.  
288 Protein solubility depends on the balance between hydrophobic protein–protein and  
289 hydrophilic protein–solvent interactions and is affected by ionic strength, pH, temperature, and  
290 amino acid composition. Solubility is typically lowest near the isoelectric point (pI), where  
291 reduced electrostatic repulsion favors aggregation (Lam et al., 2018; Tang et al., 2023). Jarpa-  
292 Parra et al. (2014) reported solubility values of ~2–11% for lentil protein near pH 4–5, with  
293 higher solubility under acidic or alkaline conditions. Similarly, Hanley, Dobson, Stobbs, and  
294 Marangoni (2025) observed lentil protein isolate solubility of 3.7% at pH 5 and 6.2% at native  
295 pH. The higher WSI observed in this study may be attributed to the relatively high abundance  
296 of polar amino acids in lentil protein (e.g., glutamine, asparagine, arginine, lysine, glutamic  
297 acid, and aspartic acid), which enhance hydrogen bonding and electrostatic interactions with  
298 water (Jarpa-Parra et al., 2014). By contrast, gluten contains a higher proportion of hydrophobic  
299 residues, with many polar residues engaged in intra- and intermolecular bonding, which  
300 restricts solubility (Wieser, 2007). The solubility contrast is highly relevant to cheese  
301 structuring. Lentil protein's higher WSI suggests that it can disperse more uniformly during  
302 hydration and thermal treatment, forming finer particulate aggregates and contributing to a  
303 more homogeneous gel network. In contrast, gluten's low solubility and tendency to form

304 cohesive, elastic strands support the development of larger-scale network continuity rather than  
305 fine particulate dispersion. This complementary behavior offers a mechanistic rationale for  
306 pairing these two proteins: lentil protein contributes water-distributed particulate domains,  
307 while gluten provides elasticity and network coherence—an interaction that has not been  
308 previously reported in plant-based cheese matrices (Boye, Aksay, et al., 2010; Dobson et al.,  
309 2022; Wieser, 2007).

310 OHC reflects the ability of proteins or biopolymers to physically entrap or bind oil and is  
311 primarily governed by hydrophobic interactions between nonpolar lipid chains and exposed  
312 hydrophobic amino acid side chains. Structural features such as porosity, surface area, and  
313 flexibility of the protein matrix, as well as oil properties (e.g., viscosity, polarity, and droplet  
314 size), also influence OHC (Kaur & Singh, 2007; Lam et al., 2018; Tang et al., 2023). In this  
315 study, lentil protein showed a slightly higher OHC (0.61 g/g) than gluten (0.54 g/g), likely due  
316 to greater surface hydrophobicity and a more open structural conformation, which facilitate  
317 lipid interactions (Boye, Zare, et al., 2010). These results are consistent with previous findings;  
318 for example, De Angelis et al. (2021) reported oil absorption values for red lentil flour between  
319 0.32 and 0.45 g/g, depending on particle size and pre-treatment. The slightly higher OHC of  
320 lentil protein has important implications for fat immobilization in cheese analogues. Proteins  
321 with higher OHC can stabilize dispersed oil droplets more effectively during emulsification  
322 and heating, reducing free-oil formation and enhancing meltability—critical attributes in  
323 cheese analogues (McClements, 2015). The OHC values observed in this study exceed those  
324 reported for native lentil flour, likely due to differences in protein purity, particle size, and  
325 processing conditions during isolate production (Boye, Zare, et al., 2010; Klupšaitė &  
326 Juodeikienė, 2015).

327 Together, the WRC, WSI, and OHC data demonstrate that lentil protein presents a  
328 multifunctional profile (high hydration + moderate solubility + moderate OHC), while gluten

329 provides network elasticity—highlighting the scientific rationale for combining these two  
330 proteins in dairy-free cheese systems.

331

### 332 **3.2. Moisture, ash, fat, and protein content**

333 The moisture content of the plant-based cheese formulations ranged from 29.68% to 60.20%.

334 A significant negative linear relationship was observed between moisture content and the levels

335 of protein, oil, modified tapioca starch, and maltodextrin (Equation 9), indicating that

336 increasing the proportion of these ingredients reduced the amount of free water within the

337 matrix. This trend reflects both hydrophilic water-binding and hydrophobic water-excluding

338 effects. Lentil protein and gluten exhibit moderate to high water-binding capacity due to their

339 polar amino acid side chains and partially unfolded structures; however, when incorporated at

340 higher levels, they increase total solids and consequently decrease the proportion of free water

341 (Boye, Zare, et al., 2010; McClements, 2015). Similarly, lipids—particularly coconut and high-

342 oleic sunflower oils—are hydrophobic and displace water within the emulsion, limiting overall

343 moisture retention (McClements, 2015). Modified tapioca starch and maltodextrin also reduce

344 moisture by binding water through hydrogen bonding and forming viscous or gel-like

345 structures that immobilize water molecules (BeMiller, 2018). The pronounced negative effect

346 of both ingredients suggests competitive water immobilization among matrix components.

347 Modified starch granules swell extensively during heating, reducing the availability of free

348 water, while maltodextrin lowers water activity by forming hydrogen-bonded water clusters

349 that contribute to a denser matrix microstructure. Such competition for water is characteristic

350 of starch–protein systems and explains the sharper moisture decline observed at higher

351 inclusion levels (Hoobin et al., 2013; Liu, Xie, Yu, Chen, & Li, 2009).

352 As expected, increasing oil and plant protein levels directly elevated fat and protein contents,

353 respectively, reflecting their compositional contributions, as predicated by Equations 11 and

354 12 (Alehosseini et al., 2025a). The protein content of the formulations ranged from 17.67% to  
355 27.63%, while the ash content ranged from 1.02% to 2.98%, reflecting the intrinsic mineral  
356 composition of the raw materials. Lentil protein, rich in potassium, phosphorus, and iron, was  
357 the primary contributor to ash, while the gluten powder likely contributed only trace minerals,  
358 originating from residual salts or processing residues (Iqbal, Khalil, Ateeq, & Sayyar Khan,  
359 2006). Chitosan, a deacetylated derivative of chitin, may also contribute trace minerals such as  
360 calcium, magnesium, and phosphorus, thereby elevating ash levels even at low concentrations  
361 (Younes & Rinaudo, 2015). In contrast, refined lipids (i.e., coconut and sunflower oils) are  
362 essentially mineral-free, and carbohydrate-based ingredients such as maltodextrin and tapioca  
363 starch contain negligible ash due to their high purification levels (BeMiller, 2018; Iqbal et al.,  
364 2006; Stevens, 2020). The relatively narrow variation in ash compared with moisture and  
365 protein suggests that mineral-bearing ingredients played a secondary but consistent structural  
366 role. This is consistent with observations in other legume-based systems, where minerals  
367 contribute to ionic bridging and mild protein aggregation, subtly affecting firmness and water  
368 mobility rather than significantly altering total ash content (Shand, Ya, Pietrasik, &  
369 Wanasundara, 2007; Stone et al., 2015). These observations align with previous studies. Grasso  
370 et al. (2024) reported moisture levels of 55.4–56.6% in plant-based cheese systems containing  
371 zein, chickpea protein concentrate, and tapioca starch. Similarly, Ndatsu et al. (2023) observed  
372 protein levels of 31.2–39.2%, moisture contents of 42.3–54.3%, fat levels of 13.4–23.5%, and  
373 ash contents of 4.4–6.4% in tofu produced from soybeans and Bambara groundnut. By contrast,  
374 lentil milk-based soft cheeses typically exhibit much higher moisture (72.44–79.58%) and  
375 lower protein contents (3.16–6.74%), reflecting differences in protein concentration, matrix  
376 structure, and processing methods (Naeem et al., 2024). Compared to previous plant-based  
377 cheese studies, the present formulations exhibit noticeably higher protein content. Gluten forms  
378 a viscoelastic network that can expel free water during heating, while lentil globulins undergo

379 partial aggregation and exhibit limited swelling capacity (Cornet, van der Goot, & van der  
380 Sman, 2020; Hong, Wanasundara, Nickerson, & Shand, 2012). Together, these behaviors  
381 produce a more compact, protein-dense matrix, consistent with the observed reductions in  
382 moisture and the higher protein concentration. This mechanistic interaction distinguishes the  
383 lentil–gluten system from plant cheeses based on single-protein sources, where either water  
384 retention or network formation dominates.

385 ANOVA results confirmed that the linear models accurately captured the experimental trends  
386 in composition, with coefficients of determination ( $R^2$ ) of 0.95 (moisture), 0.95 (ash), 0.99 (fat),  
387 and 0.97 (protein) (Table 3). Most formulation variables—including protein, oil, modified  
388 starch, and maltodextrin—had significant effects on moisture ( $p < 0.01$ ). The high  $R^2$  values  
389 indicate strong model predictability, with moisture, fat, protein, and ash responding as expected  
390 to their primary ingredients. Moisture exhibited multicomponent competition, reflecting the  
391 complex interactions among proteins, starches, and polysaccharides, whereas fat and ash  
392 behaved largely additively.

393

**Table 3.** Analysis of variance (ANOVA) for moisture, ash, fat, and protein content

Source	Moisture content		Ash		Fat content		Protein content	
	Sum of Squares	p-value	Sum of Squares	p-value	Sum of Squares	p-value	Sum of Squares	p-value
<b>Model</b>	2038.36	< 0.0001	18.17	< 0.0001	843.41	< 0.0001	581.13	< 0.0001
Protein ( $X_1$ )	767.12	< 0.0001	17.87	< 0.0001	0.0467	0.4800	580.18	< 0.0001
Oil ( $X_2$ )	986.26	< 0.0001	0.0482	0.1415	842.91	< 0.0001	0.0241	0.7924
Chitosan ( $X_3$ )	0.3380	0.7131	0.2423	0.0016	0.2406	0.1131	0.2119	0.4370
Modified tapioca starch ( $X_4$ )	217.12	< 0.0001	0.0036	0.6842	0.1943	0.1534	0.6533	0.1754
Maltodextrin ( $X_5$ )	67.51	< 0.0001	0.0068	0.5773	0.0170	0.6696	0.0568	0.6866
<b>Residual</b>	108.59		0.9459		4.05		15.15	
Lack of Fit	97.92	0.2294	0.9217	0.0605	3.65	0.2274	14.28	0.0612
Pure Error	10.67		0.0242		0.3964		0.8722	
<b>Cor Total</b>	2146.96		19.12		847.46		596.28	

394

395 To provide a more quantitative description of these effects, predictive regression models were  
 396 developed (Equations 9-12). These models quantify how protein powder ( $X_1$ ), oil ( $X_2$ ), chitosan  
 397 ( $X_3$ ), modified tapioca starch ( $X_4$ ), and maltodextrin ( $X_5$ ) influence the compositional attributes  
 398 of the formulations. The sign and magnitude of each coefficient indicate the direction and  
 399 strength of each ingredient's effect. For example, negative coefficients in the moisture model  
 400 highlight water displacement or immobilization by solids, whereas positive coefficients in the  
 401 ash and protein models reflect compositional enrichment from mineral- and protein-rich  
 402 ingredients. The fat model, meanwhile, shows a nearly one-to-one increase in fat content with  
 403 oil addition, consistent with its direct contribution to the formulation. Collectively, these  
 404 equations confirm and extend the experimental observations, while offering predictive  
 405 capability for future formulation optimization.

406

$$\text{Moisture (\%)} = -0.95 X_1 - 1.08 X_2 - 1.01 X_4 - 1.41 X_5 + 97.32 \quad (9)$$

$$\text{Ash (\%)} = 0.14 X_1 + 0.17 X_3 - 1.74 \quad (10)$$

$$\text{Fat (\%)} = 0.99 X_2 + 0.55 \quad (11)$$

$$\text{Protein (\%)} = 0.83 X_1 + 1.51 \quad (12)$$

407

### 408 3.3. Color profile determination of dairy-free cheese alternatives

409 The color of dairy-free cheese alternatives is influenced by intrinsic factors (e.g., ingredient  
 410 type and concentration, fat content, and functional additives) and extrinsic factors (e.g.,  
 411 processing, packaging, and storage), all of which influence visual appeal and consumer  
 412 acceptance (Bekele, Hansen, Eshetu, Ipsen, & Hailu, 2019).

413 In this study,  $L^*$ ,  $a^*$ , and  $b^*$  values ranged from 58.28 to 83.32,  $-0.06$  to  $1.11$ , and  $16.04$  to  
 414  $23.97$ , respectively, indicating noticeable variability in brightness and chromaticity due to  
 415 formulation differences. ANOVA results (Table 4) showed that linear models provided

416 excellent fits for  $L^*$ ,  $a^*$ ,  $\Delta E$ , and  $WI$ , with  $R^2$  values of 0.91, 0.83, 0.91, and 0.93, respectively.  
417 In contrast, two-factor interaction (2FI) models better explained variation in  $b^*$  and  $C_{ab}^*$ ,  $R^2$   
418 values of 0.96 for both. These results emphasize the importance of ingredient interactions on  
419 color attributes. Protein ( $p < 0.01$ ), oil ( $p < 0.01$ ), and modified starch ( $p < 0.05$ ) significantly  
420 affected  $L^*$ ,  $b^*$ ,  $\Delta E$ ,  $WI$ , and  $C_{ab}^*$ . Chitosan had a significant effect on  $L^*$ ,  $\Delta E$ , and  $WI$  ( $p <$   
421  $0.01$ ), while protein and modified starch strongly influenced  $a^*$  ( $p < 0.01$ ). Maltodextrin also  
422 significantly affected  $L^*$ ,  $b^*$ ,  $WI$ , and  $C_{ab}^*$  ( $p < 0.05$ ). Among interactions, protein–oil ( $X_1X_2$ )  
423 had a pronounced effect on  $b^*$  and  $C_{ab}^*$  ( $p < 0.01$ ).

424

**Table 4.** Analysis of variance (ANOVA) for  $L^*$ ,  $a^*$ ,  $b^*$ ,  $\Delta E$ ,  $WI$ , and  $C_{ab}^*$ 

Source	$L^*$		$a^*$		$b^*$		$\Delta E$		$WI$		$C_{ab}^*$	
	Sum of Squares	p-value	Sum of Squares	p-value	Sum of Squares	p-value	Sum of Squares	p-value	Sum of Squares	p-value	Sum of Squares	p-value
<b>Model</b>	1187.71	< 0.0001	3.87	< 0.0001	174.14	< 0.0001	944.52	< 0.0001	970.51	< 0.0001	175.24	< 0.0001
Protein ( $X_1$ )	421.41	< 0.0001	3.62	< 0.0001	139.83	< 0.0001	218.44	< 0.0001	542.80	< 0.0001	140.96	< 0.0001
Oil ( $X_2$ )	568.26	< 0.0001	0.0157	0.3507	25.91	< 0.0001	571.95	< 0.0001	280.80	< 0.0001	25.84	< 0.0001
Chitosan ( $X_3$ )	127.92	< 0.0001	0.0013	0.7874	0.3050	0.2288	104.30	< 0.0001	89.33	< 0.0001	0.3068	0.2282
Modified tapioca starch ( $X_4$ )	56.86	< 0.0001	0.2240	0.0009	0.9456	0.0381	41.89	< 0.0001	44.92	< 0.0001	0.9256	0.0404
Maltodextrin ( $X_5$ )	13.24	0.0271	0.0003	0.8978	1.25	0.0182	7.94	0.0626	12.66	0.0080	1.25	0.0184
Protein $\times$ Oil ( $X_1 \times X_2$ )					3.68	0.0002					3.71	0.0002
Protein $\times$ Chitosan ( $X_1 \times X_3$ )					0.3983	0.1704					0.4028	0.1689
Protein $\times$ Modified tapioca starch ( $X_1 \times X_4$ )					0.2869	0.2428					0.2945	0.2376
Protein $\times$ Maltodextrin ( $X_1 \times X_5$ )					0.3383	0.2055					0.3383	0.2063
Oil $\times$ Chitosan ( $X_2 \times X_3$ )					0.0520	0.6161					0.0553	0.6059
Oil $\times$ Modified tapioca starch ( $X_2 \times X_4$ )					0.0259	0.7233					0.0226	0.7413
Oil $\times$ Maltodextrin ( $X_2 \times X_5$ )					0.1263	0.4358					0.1288	0.4322

Chitosan × Modified tapioca starch ( $X_3 \times X_4$ )					0.0332	0.6887					0.0358	0.6779
Chitosan × Maltodextrin ( $X_3 \times X_5$ )					0.8033	0.0548					0.8033	0.0552
Modified tapioca starch × Maltodextrin ( $X_4 \times X_5$ )					0.1639	0.3753					0.1668	0.3721
<b>Residual</b>	111.40		0.7750		6.90		95.70		72.06		6.93	
Lack of Fit	111.01	0.0600	0.7102	0.1588	6.82	0.0752	95.40	0.0701	71.70	0.0619	6.85	0.0899
Pure Error	0.3868		0.0648		0.0800		0.3042		0.3568		0.0800	
<b>Cor Total</b>	1299.11		4.64		181.05		1040.22		1042.57		182.17	

426

427 Predictive models describing the effects of formulation variables on color attributes are  
 428 presented in Equations (13–18). For example, negative coefficients in  $L^*$  indicate factors that  
 429 decrease brightness, while positive coefficients in  $a^*$  or  $b^*$  highlight contributions to red or  
 430 yellow hues, respectively—consistent with how color parameters respond to ingredient  
 431 proportions in mixture design models of cheese analogue systems (Garcia-Fontanals et al.,  
 432 2023).

433

$$L^* = -0.70 X_1 - 0.82 X_2 - 3.88 X_3 + 0.52 X_4 + 0.62 X_5 + 97.15 \quad (13)$$

$$a^* = +0.06 X_1 + 0.03 X_4 - 1.45 \quad (14)$$

$$b^* = +0.04 X_1 - 0.44 X_2 - 0.28 X_4 - 1.05 X_5 + 0.01 X_1 X_2 + 23.43 \quad (15)$$

$$\Delta E = +0.51 X_1 + 0.82 X_2 + 3.50 X_3 - 0.44 X_4 + 4.39 \quad (16)$$

$$WI = -0.80 X_1 - 0.57 X_2 - 3.24 X_3 + 0.46 X_4 + 0.61 X_5 + 89.36 \quad (17)$$

$$C_{ab}^* = +0.04 X_1 - 0.44 X_2 - 0.28 X_4 - 1.05 X_5 + 0.01 X_1 X_2 + 23.44 \quad (18)$$

434

435 These results are consistent with previous findings. Naeem et al. (2024) reported  $L^*$ ,  $a^*$ , and  $b^*$   
 436 values of 39.61–41.64, 0.13–7.97, and 11.43–15.01, respectively, in lentil milk-based soft  
 437 cheese analogs, while Garcia-Fontanals et al. (2023) reported  $L^* = 62.57$ ,  $a^* = -0.25$ , and  $b^* =$   
 438 17.23 in hybrid faba bean–insect protein cheeses. Differences in protein type, pigment  
 439 composition, and fat content likely explain the observed variability across studies (Chudy,  
 440 Bilska, Kowalski, & Teichert, 2020; Sharan et al., 2021).

441 As confirmed by the predictive models (Equations 13–15), in the present study, increasing  
 442 protein powder (lentil and gluten) decreased  $L^*$  while increasing  $a^*$  and  $b^*$ , reflecting  
 443 contributions from natural pigments and heat-induced browning reactions, including Maillard  
 444 reaction products formed during thermal processing (Martins, Jongen, & van Boekel, 2000;  
 445 Sharan et al., 2021). Conversely, modified starch and maltodextrin, being white and highly  
 446 dispersible, increased  $L^*$  while reducing  $b^*$  by diluting pigment dispersion and enhancing light

447 scattering. Modified starch also increased  $a^*$ , likely by improving pigment dispersion within  
448 the matrix. Oils reduced  $L^*$  and  $b^*$ , possibly by diminishing light scattering and diluting water-  
449 soluble pigments. Variations in  $b^*$  may also reflect differences in carotenoid levels (e.g.,  $\beta$ -  
450 carotene, lutein, zeaxanthin) inherent to protein and oil components (Chong & Wong, 2017;  
451 Chudy et al., 2020; Naeem et al., 2024; Wannasin & McClements, 2023). Overall, these results  
452 not only quantify color changes but also provide causal insights into ingredient interactions and  
453 pigment distribution, offering guidance for designing visually appealing dairy-free cheeses.

454

### 455 **3.4. Meltability**

456 Meltability, a key functional property of cheese, describes the extent and uniformity of  
457 softening and flow upon heating. In plant-based cheese, meltability is influenced by ingredient  
458 composition (e.g., type and ratio of proteins, fats, polysaccharides, and additives), as well as  
459 processing conditions, which together shape the microstructure, water mobility, and fat  
460 dispersion (Alehosseini et al., 2025b; Grasso et al., 2021; Rodríguez, 2017).

461 The meltability of the dairy-free cheese alternatives ranged from 1.20% to 24%, consistent with  
462 values reported for commercial plant-based cheeses (5.59–21.00%) by Grasso et al. (2021), but  
463 substantially lower than dairy cheeses, which exhibit greater meltability due to their casein  
464 matrix and emulsified fat structure (Kovačević et al., 2024). This difference highlights the  
465 limited ability of plant proteins to form heat-sensitive networks, in contrast to casein, which  
466 readily rearranges upon heating to enable flow, as demonstrated by rheological and  
467 microstructural studies of dairy cheese systems (Fox, Guinee, Cogan, & McSweeney, 2017).  
468 ANOVA results (Table 5) showed that meltability was best described by a quadratic model ( $R^2$   
469 = 0.93), indicating strong nonlinear effects of formulation variables on this property.

470

**Table 5.** Analysis of variance (ANOVA) for meltability, hardness, cohesiveness, springiness, and adhesiveness

Source	Meltability		Hardness		Cohesiveness		Springiness		Adhesiveness		Resilience	
	Sum of Squares	p-value	Sum of Squares	p-value	Sum of Squares	p-value	Sum of Squares	p-value	Sum of Squares	p-value	Sum of Squares	p-value
<b>Model</b>	1450.83	< 0.0001	5857.31	< 0.0001	0.3434	< 0.0001	12686.16	< 0.0001	61.55	< 0.0001	1047.60	< 0.0001
Protein ( $X_1$ )	266.56	< 0.0001	3560.10	< 0.0001	0.1725	< 0.0001	12229.54	< 0.0001	28.25	< 0.0001	409.16	< 0.0001
Oil ( $X_2$ )	569.90	< 0.0001	1202.34	< 0.0001	0.1274	< 0.0001	178.52	0.0015	3.34	0.0015	525.49	< 0.0001
Chitosan ( $X_3$ )	215.51	< 0.0001	580.32	< 0.0001	0.0122	0.0150	197.26	0.0009	5.97	< 0.0001	23.51	0.0089
Modified tapioca starch ( $X_4$ )	15.83	0.0448	472.34	< 0.0001	0.0246	0.0008	64.72	0.0482	0.1109	0.5269	64.81	< 0.0001
Maltodextrin ( $X_5$ )	5.12	0.2424	42.20	0.0865	0.0068	0.0648	16.11	0.3161	2.81	0.0031	24.62	0.0075
Protein $\times$ Oil ( $X_1 \times X_2$ )	41.40	0.0020							3.22	0.0017		
Protein $\times$ Chitosan ( $X_1 \times X_3$ )	0.0450	0.9117							6.73	< 0.0001		
Protein $\times$ Modified tapioca starch ( $X_1 \times X_4$ )	0.0050	0.9705							0.0490	0.6736		
Protein $\times$ Maltodextrin ( $X_1 \times X_5$ )	0.4050	0.7397							2.16	0.0085		
Oil $\times$ Chitosan ( $X_2 \times X_3$ )	99.40	< 0.0001							0.1867	0.4129		
Oil $\times$ Modified tapioca starch ( $X_2 \times X_4$ )	1.81	0.4845							0.5248	0.1742		
Oil $\times$ Maltodextrin ( $X_2 \times X_5$ )	2.20	0.4401							0.0655	0.6263		

Chitosan × Modified tapioca starch ( $X_3 \times X_4$ )	8.40	0.1373							0.0300	0.7414		
Chitosan × Maltodextrin ( $X_3 \times X_5$ )	9.25	0.1198							1.81	0.0150		
Modified tapioca starch × Maltodextrin ( $X_4 \times X_5$ )	0.6050	0.6848							0.0113	0.8398		
Protein <sup>2</sup> ( $X_1$ ) <sup>2</sup>	14.15	0.0569							0.0596	0.6422		
Oil <sup>2</sup> ( $X_2$ ) <sup>2</sup>	6.39	0.1929							0.2898	0.3092		
Chitosan <sup>2</sup> ( $X_3$ ) <sup>2</sup>	6.27	0.1972							0.0115	0.8379		
Modified tapioca starch <sup>2</sup> ( $X_4$ ) <sup>2</sup>	9.82	0.1094							0.0034	0.9111		
Maltodextrin <sup>2</sup> ( $X_5$ ) <sup>2</sup>	0.0912	0.8746							0.1978	0.3995		
<b>Residual</b>	104.37		604.17		0.0834		689.48		7.84		137.90	
Lack of Fit	104.07	0.1022	603.79	0.0650	0.0831	0.0609	665.12	0.1500	7.83	0.0600	137.38	0.0670
Pure Error	0.3000		0.3771		0.0003		24.36		0.0169		0.5194	
<b>Cor Total</b>	1555.20		6461.47		0.4269		13375.64		69.40		1185.49	

473 Protein, oil, and chitosan ( $p < 0.01$ ), along with modified starch ( $p < 0.05$ ), significantly  
 474 influenced meltability. Notably, interactions between protein and oil ( $X_1X_2$ ) and oil and  
 475 chitosan ( $X_2X_3$ ) were also significant ( $p < 0.01$ ). The significant interaction terms reflect the  
 476 complex network formation and plasticization mechanisms in plant-based cheese matrices  
 477 (Lyu, Sala, & Scholten, 2023). To quantitatively assess the effects of formulation variables on  
 478 meltability, a predictive regression model was developed, with coefficients indicating the  
 479 direction and relative strength of each ingredient's influence (Equation 19):

$$\begin{aligned} \text{Meltability (\%)} = & -2.02 X_1 + 4.29 X_2 - 13.87 X_3 - 5.89 X_4 - 0.04 X_1X_2 - \\ & 0.70 X_2X_3 + 100.44 \end{aligned} \quad (19)$$

481  
 482 Protein had a strong negative effect on meltability. Higher levels of lentil protein and gluten  
 483 likely formed dense, thermally stable networks reinforced by hydrogen bonds and disulfide  
 484 bridges, limiting flow upon heating. Heat-induced Maillard cross-linking further reduced  
 485 thermal flow (Grasso et al., 2022; Mattice & Marangoni, 2020; Sharan et al., 2021). Chitosan  
 486 also decreased meltability, probably by forming electrostatic interactions with negatively  
 487 charged proteins and starch phosphates, thereby reinforcing the gel network and restricting  
 488 protein and fat mobility (Alehosseini, Shahiri Tabarestani, Kharazmi, & Jafari, 2022).  
 489 Modified tapioca starch had a similar negative impact: upon heating, starch granules  
 490 gelatinized and retrograded, forming semi-crystalline networks resistant to deformation  
 491 (Hanley et al., 2025; Sutter, Assad-Bustillos, & Windhab, 2023; Ye, Hewitt, & Taylor, 2009).  
 492 This effect explains why formulations with higher starch content exhibited lower meltability  
 493 despite adequate oil content, as the crystalline starch domains act as physical barriers to flow.  
 494 In contrast, oil content significantly enhanced meltability. Fats such as coconut oil and high-  
 495 oleic sunflower oil acted as plasticizers, reducing matrix rigidity and lowering the melting point

496 by interrupting protein–starch interactions (McClements & Grossmann, 2023; Sivakami,  
497 2021). These data confirm prior findings that fat phase continuity facilitates melt flow in plant-  
498 based matrices (Dobson & Marangoni, 2023; Sanders, Dobson, & Marangoni, 2024).  
499 Processing conditions further influenced meltability: protein hydration prior to starch addition  
500 allowed partial unfolding and early network formation, controlled heating at 80 °C minimized  
501 excessive denaturation, and high shear mixing promoted starch swelling and fat dispersion.  
502 Nonetheless, high starch and protein concentrations still restricted meltability in certain  
503 formulations, consistent with prior observations (Dobson & Marangoni, 2023). These results  
504 suggest that meltability is not solely determined by individual ingredients but also by their  
505 interactions and processing-induced microstructural arrangements, emphasizing the  
506 importance of integrated formulation–processing strategies.

507

### 508 **3.5. Texture profile analysis (TPA)**

509 TPA is a widely used method for evaluating the mechanical and sensory properties of food,  
510 simulating the first two bites of mastication by compressing samples to a defined deformation  
511 level (Dobson & Marangoni, 2023). In this study, hardness, cohesiveness, springiness,  
512 adhesiveness, and resilience were measured to characterize the textural properties of the dairy-  
513 free cheese alternatives. ANOVA results (Table 5) and the predictive model (Equation 20)  
514 indicated that hardness increased significantly with higher levels of plant protein, oil, chitosan,  
515 and modified tapioca starch, while maltodextrin had no significant effect. This suggests that  
516 network formation and structural reinforcement primarily govern firmness, rather than simple  
517 water-binding effects (Dobson & Marangoni, 2025). Lentil protein (rich in globulins) and  
518 gluten (gliadin and glutenin) synergistically formed a viscoelastic network via protein–protein  
519 interactions, reinforced by thermal denaturation during processing (Gasparre, van den Berg,  
520 Oosterlinck, & Sein, 2022; Jo, Huang, & Chen, 2020). Coconut oil, owing to its high saturated

521 fat content and sharp melting point, enhanced firmness through crystalline fat structuring upon  
522 cooling, while high-oleic sunflower oil contributed minimal softening. This aligns with prior  
523 findings that saturated fats can act as active structuring agents, whereas unsaturated fats may  
524 act primarily as lubricants, highlighting formulation-dependent fat–protein interactions  
525 (Sanders et al., 2024). Chitosan, a cationic biopolymer, likely increased hardness by forming  
526 electrostatic protein–polysaccharide interactions (Xie et al., 2024). Modified tapioca starch  
527 gelatinized during heating and retrograded upon cooling, providing rigidity and acting as a  
528 filler in the protein–polysaccharide matrix (N.-N. Zhang et al., 2022). The combined effect of  
529 starch and protein emphasizes the importance of non-covalent interactions and filler  
530 reinforcement in plant-based cheese matrices (Lyu, Sala, & Scholten, 2022). Comparable  
531 trends have been reported by Dobson and Marangoni (2023), although other systems (e.g., soy,  
532 pea) have shown the opposite effect, underscoring the formulation-dependent and ingredient-  
533 specific nature of textural outcomes (Mefleh et al., 2022; Rinaldoni, Palatnik, Zaritzky, &  
534 Campderrós, 2014).

535 Cohesiveness, which reflects the strength of internal bonding (Grasso et al., 2022), increased  
536 with protein, chitosan, and starch, but decreased with oil addition (Equation 21). Gluten  
537 contributed elasticity, lentil protein increased structural density, and chitosan strengthened the  
538 network via electrostatic and hydrogen bonding, while starch gelation and water-binding  
539 further supported cohesiveness (Boeck, Zannini, Sahin, Bez, & Arendt, 2021; N.-N. Zhang et  
540 al., 2022). In contrast, higher oil levels disrupted protein–polysaccharide continuity, leading to  
541 dispersed fat domains, consistent with previous observations in low-fat cheese analogues (Lim,  
542 Easa, Karim, Bhat, & Liang, 2011). This provides mechanistic insight into why cohesiveness  
543 decreases despite the presence of network-forming components.

544 Springiness followed the same trend: it was enhanced by protein, chitosan, and starch but  
545 reduced by oil (Equation 22). Elastic recovery was attributed to the viscoelastic protein

546 network, further reinforced by chitosan gelation and starch retrogradation (Grasso et al., 2022;  
547 Mattice & Marangoni, 2020). Oils, particularly coconut oil, acted as plasticizers, weakening  
548 intermolecular linkages and reducing elastic recovery (Bhasney, Patwa, Kumar, & Katiyar,  
549 2017; Dobson et al., 2022; Mattice & Marangoni, 2020).

550 Based on significant main effects in the fitted model (Equation 23), adhesiveness decreased  
551 with higher protein and chitosan but increased with oil and maltodextrin. A dense protein  
552 network, particularly with gluten cross-linking, limited water mobility and reduced stickiness  
553 (Ushkalova, Zhao, Gu, Wang, & Zhang, 2025). Chitosan further decreased adhesiveness by  
554 stabilizing gel structures (Xie et al., 2024). In contrast, oils reduced friction at the probe–sample  
555 interface, while maltodextrin’s water-binding enhanced lubrication, producing a stickier  
556 texture (Akshit, Poswal, Kaushik, Deshwal, & Huppertz, 2025).

557 Resilience, the ability of the matrix to recover after deformation, was enhanced by protein,  
558 chitosan, and starch, but reduced by oil and maltodextrin (Equation 24). Protein–chitosan  
559 interactions promoted elasticity, while starch retrogradation strengthened recovery capacity.  
560 Conversely, oil disrupted network integrity, and maltodextrin diluted the protein phase,  
561 lowering resilience (Scott & Awika, 2023; Xie et al., 2024). This observation is novel in dairy-  
562 free cheese alternatives and suggests that resilience can be tailored by optimizing  
563 polysaccharide–protein ratios, an aspect not fully explored in previous plant-based cheese  
564 studies.

565 Collectively, these results indicate that desirable cheese-like textures in plant-based products  
566 rely on carefully balancing moisture retention, phase continuity, and non-covalent interactions  
567 among proteins, polysaccharides, and fats (Mattice & Marangoni, 2020; Naeem et al., 2024).  
568 This study highlights the unique role of chitosan as both a hardness and cohesiveness enhancer  
569 while modulating adhesiveness and resilience, an aspect that could inform future formulation  
570 strategies for dairy-free cheeses.

571 ANOVA results (Table 5) confirmed that hardness, cohesiveness, springiness, and resilience  
 572 were best described by linear models ( $R^2 = 0.91, 0.80, 0.95,$  and  $0.88,$  respectively), while  
 573 adhesiveness fit a quadratic model ( $R^2 = 0.89$ ). Plant protein ( $p < 0.01$ ), oil ( $p < 0.01$ ), and  
 574 chitosan ( $p < 0.05$ ) significantly influenced all TPA parameters. Modified tapioca starch  
 575 significantly affected hardness ( $p < 0.01$ ), cohesiveness ( $p < 0.01$ ), springiness ( $p < 0.05$ ), and  
 576 resilience ( $p < 0.01$ ), while maltodextrin significantly influenced adhesiveness and resilience  
 577 ( $p < 0.01$ ). Several interactions, including protein  $\times$  oil, protein  $\times$  chitosan, protein  $\times$   
 578 maltodextrin, and chitosan  $\times$  maltodextrin, also significantly shaped adhesiveness. These  
 579 interactions indicate synergistic and antagonistic effects between ingredients, providing  
 580 mechanistic understanding beyond simple main effects, which supports predictive formulation  
 581 modeling. Predictive models were developed, with coefficients indicating the relative effect of  
 582 protein, oil, chitosan, starch, and maltodextrin on hardness, cohesiveness, springiness,  
 583 adhesiveness, and resilience (Equations 20–24).

584

$$\text{Hardness (N)} = 2.05 X_1 + 1.19 X_2 + 8.26 X_3 + 1.49 X_4 - 65.99 \quad (20)$$

$$\text{Cohesiveness} = 0.01 X_1 - 0.01 X_2 + 0.04 X_3 + 0.01 X_4 + 0.21 \quad (21)$$

$$\text{Springiness (\%)} = +3.79 X_1 - 0.46 X_2 + 4.82 X_3 + 0.55 X_4 - 30.09 \quad (22)$$

$$\begin{aligned} \text{Adhesiveness (N.s)} = & -0.31 X_1 + 0.05 X_2 - 3.90 X_3 + 0.17 X_5 - 0.01 X_1 X_2 + \\ & 0.18 X_1 X_3 - 0.05 X_1 X_5 - 0.47 X_3 X_5 + 16.29 \end{aligned} \quad (23)$$

$$\text{Resilience (\%)} = +0.69 X_1 - 0.79 X_2 + 1.66 X_3 + 0.55 X_4 - 0.85 X_5 + 12.05 \quad (24)$$

585

### 586 3.6. Formulation optimization of dairy-free cheese alternatives

587 Optimization analysis showed that maximizing fat content and hardness, with protein content  
 588 fixed at 25% and all other parameters constrained within acceptable ranges, yielded an optimal  
 589 formulation consisting of 28.31% protein powder, 20% oil, 1.5% chitosan, 9.99% modified

590 tapioca starch, and 2.03% maltodextrin. This combination produced a dairy-free cheese  
591 alternative with a high overall desirability (Figure 2).

592

593 INSERT FIGURE 2 ABOUT HERE

594

595 When compared with existing literature, the optimized formulation exhibits higher protein  
596 content than many commercial dairy-free cheeses (e.g., plant-based cheeses with 0.1-3%  
597 protein) while maintaining desirable textural properties, highlighting the innovation of using a  
598 combination of protein powder, chitosan, and modified starch to balance texture and nutritional  
599 value (Ali, O'Mahony, O'Sullivan, & Kerry, 2025; Grasso et al., 2021). These findings not  
600 only provide a scientifically grounded formulation strategy but also offer mechanistic insights  
601 into how the interplay of protein, fat, and polysaccharides governs the functional properties of  
602 dairy-free cheese alternatives, as evidenced by the significant main and interaction effects  
603 identified in the RSM models.

604 Validation of the RSM was conducted by preparing three replicates of the predicted optimal  
605 formulation and measuring their properties. The agreement between the experimental data and  
606 the predicted values also confirmed the predictability of the RSM model resulting from the  
607 optimal formulation (Table 6).

608

**Table 6.** Experimental validation data of the optimal formulation

	Experimental	Predicted		Experimental	Predicted		Experimental	Predicted
Moisture (%)	36.22	35.87	$b^*$	17.00	16.50	Cohesiveness	0.50	0.45
Ash (%)	2.59	2.48	$\Delta E$	36.28	36.08	Springiness (%)	79.50	80.73
Fat (%)	20.58	20.35	$WI$	57.98	56.29	Adhesiveness (N.s)	4.23	4.69
Protein (%)	24.77	25.00	$C_{ab}^*$	17.01	16.51	Resilience (%)	22.24	22.04
$L^*$	61.58	61.57	Meltability (%)	6.00	5.76			
$a^*$	0.63	0.55	Hardness (N)	43.72	43.12			

609

#### 610 4. Conclusion

611 This study provides a mechanistic and quantitative assessment of how lentil–gluten protein  
612 systems, in combination with modified tapioca starch, chitosan, maltodextrin, and blended  
613 vegetable oils, determine the composition, meltability, color, and texture of dairy-free cheese  
614 alternatives. Functional characterization revealed distinct and complementary ingredient  
615 behaviors: modified tapioca starch exhibited the highest swelling capacity (9.21 mL/g) and  
616 WRC (10.03 g/g), forming the primary hydration matrix, while lentil protein demonstrated  
617 markedly higher WSI (9.47%) and OHC (0.61 g/g) compared with gluten, supporting  
618 emulsification and uniform protein dispersion. The systematic application of RSM enabled  
619 precise modeling of both independent and interactive effects among formulation variables.  
620 Increasing protein from 20% to 30% increased hardness and cohesiveness, but reduced  
621 meltability due to the formation of thermally stable lentil–gluten networks reinforced by  
622 hydrogen bonding and disulfide cross-linking. Oil content, in contrast, improved meltability by  
623 plasticizing the matrix and disrupting protein–starch interactions, while starch and chitosan  
624 significantly enhanced hardness, cohesiveness, springiness, and resilience through  
625 gelatinization, retrogradation, and electrostatic interactions. Color parameters were strongly  
626 formulation-dependent, with increasing protein levels decreasing  $L^*$  but increasing  $a^*$  and  $b^*$ ,  
627 an effect attributed to the concentration of lentil-derived carotenoids and phenolic pigments, as  
628 well as the formation of Maillard reaction chromophores during thermal processing. The  
629 optimized formulation—28.31% protein powder, 20% oil, 1.5% chitosan, 9.99% starch, and  
630 2.03% maltodextrin—achieved a high desirability score and showed excellent agreement  
631 between predicted and experimental values, confirming the robustness and predictive utility of  
632 the RSM model. The study also demonstrates, for the first time, that lentil proteins form  
633 hydrated particulate structures while gluten contributes viscoelastic extensibility, enabling a  
634 protein-dense, cohesive matrix uncommon in legume-only cheese analogues. Overall, this

635 work advances the scientific basis for formulating high-protein plant-based cheeses by  
636 establishing quantitative, ingredient-specific structure–function relationships. The findings  
637 offer actionable strategies for tailoring meltability, texture, and compositional quality and  
638 underline the potential of lentil–gluten systems for developing nutritionally balanced and  
639 structurally robust dairy-free cheese products. Future research should integrate sensory  
640 validation, explore targeted hydrocolloid substitutions to enhance meltability, and assess  
641 nutrient bioaccessibility and processing scalability to facilitate commercial application.

642

#### 643 **CRedit authorship contribution statement**

644 -

645

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650

#### 651 **Declaration of competing interest**

652 The authors declare that they have no known competing financial interests or personal  
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654

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**Figure 1.** Schematic representation of the procedures for assessing swelling capacity, water retention capacity (WRC), water solubility index (WSI), and oil-holding capacity (OHC)

**Figure 2.** Three dimensional (3D) plots of changes in desirability at different levels of (A) protein powder and oil, (B) oil and chitosan, (C) oil and modified tapioca starch, and (D) oil and maltodextrin

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Figure 1.

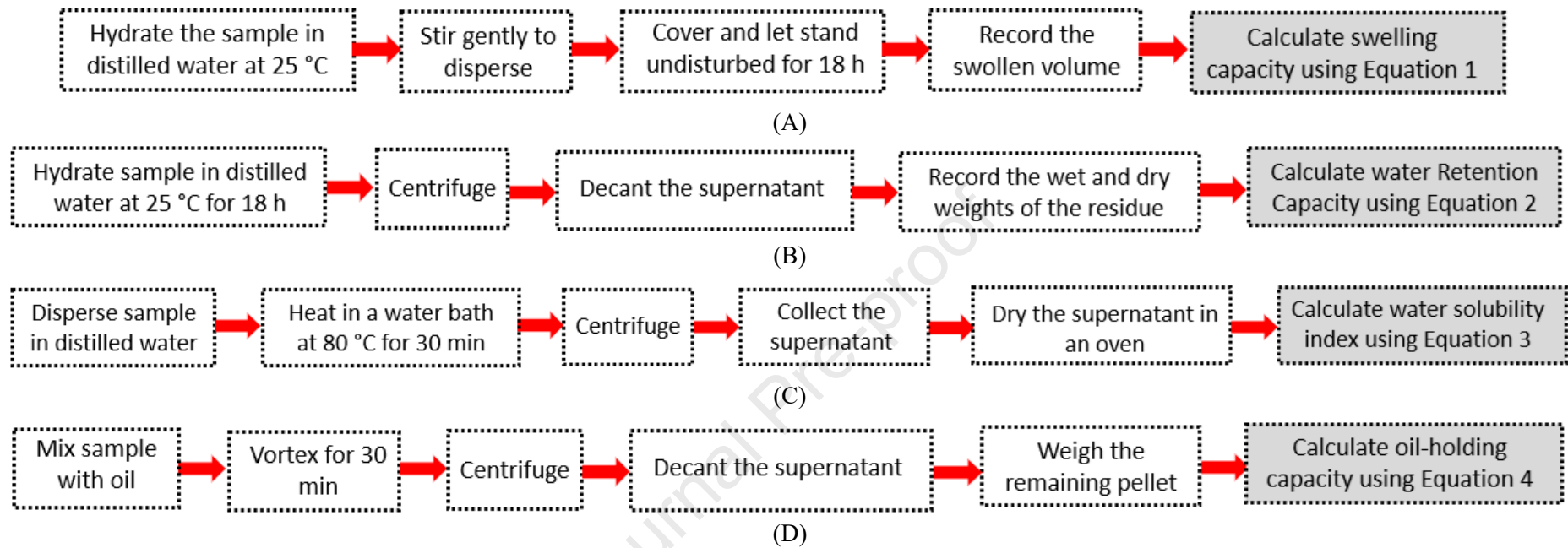
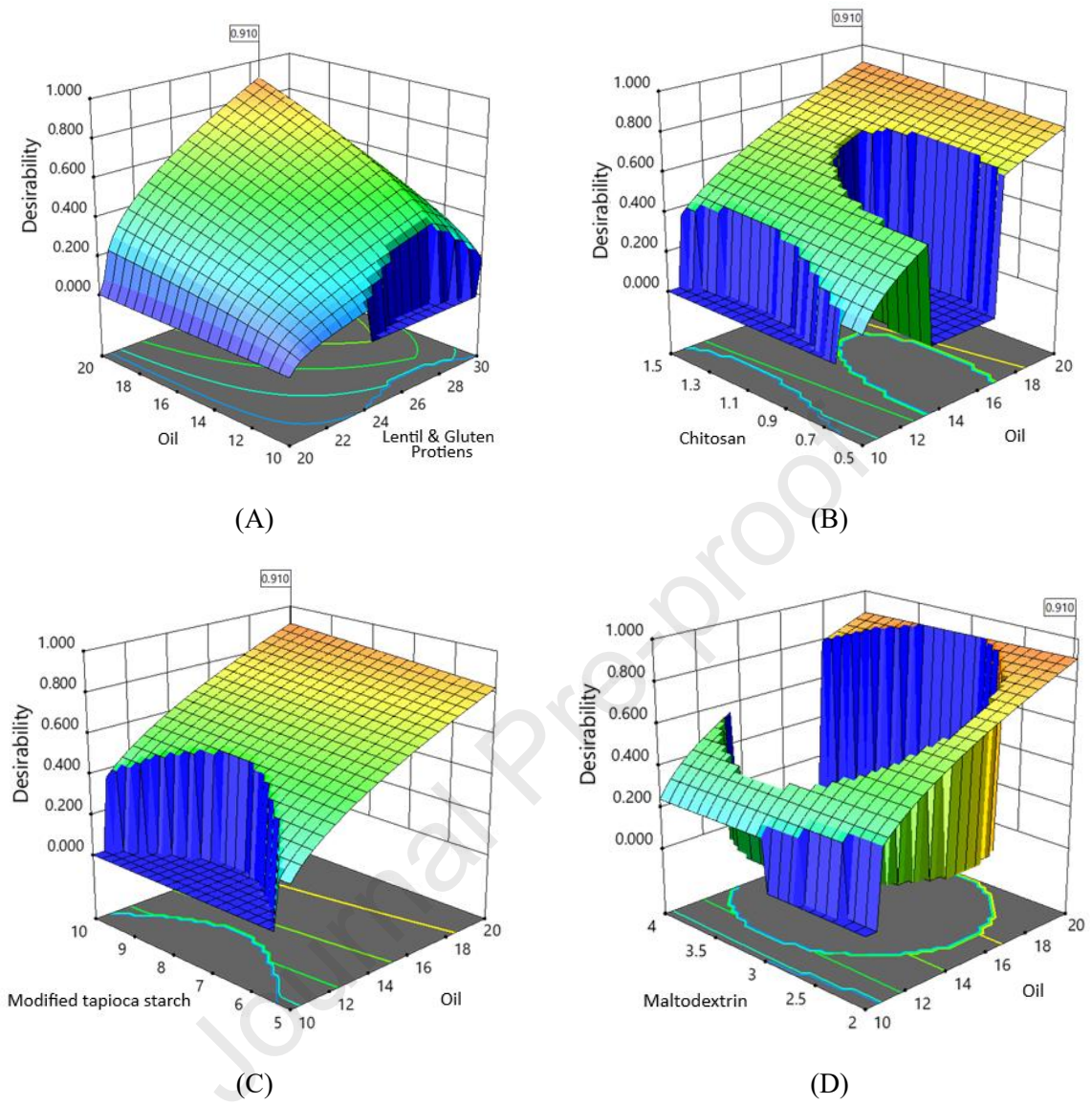


Figure 2.



**Highlights**

- Lentil and gluten proteins form viscoelastic matrices in dairy-free cheese.
- Chitosan and modified tapioca starch enhance hardness and textural stability.
- Oil type modulates meltability by altering protein–starch network dynamics.
- Protein and starch interactions significantly influence color,  $L^*$ ,  $a^*$ , and  $b^*$  values.
- RSM modeling predicts physicochemical and textural properties of formulations.

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**Declaration of interests**

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

The authors declare the following financial interests/personal relationships which may be considered as potential competing interests:

Song Miao reports financial support was provided by European Commission. If there are other authors, they declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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