

REVIEW ARTICLE

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The multifactorial phenomenon of enzymatic hydrolysis resistance in unripe banana flour and its starch: A concise review

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Abstract

Unripe banana flour starch possesses a high degree of resistance to enzymatic hydrolysis, a unique and desirable property that could be exploited in the development of functional food products to regulate blood sugar levels and promote digestive health. However, due to a multifactorial phenomenon in the banana flour matrix—from the molecular to the micro level—there is no consensus regarding the complex mechanisms behind the slow enzymatic hydrolysis of unripe banana flour starch. This work therefore explores factors that influence the enzymatic hydrolysis resistance of raw and modified banana flour and its starch including the proportion and distribution of the amorphous and crystalline phases of the starch granules; granule morphology; amylose–amylopectin ratio; as well as the presence of nonstarch components such as proteins, lipids, and phenolic compounds. Our findings revealed that the relative contributions of these factors to banana starch hydrolytic resistance are apparently dependent on the native or processed state of the starch as well as the cultivar type. The inter-relativity of these factors in ensuring amylolytic resistance of unripe banana flour starch was further highlighted as another reason for the multifactorial phenomenon. Knowledge of these factors and their contributions to enzymatic hydrolysis resistance individually and interconnectedly will provide insights into enhanced ways of extraction, processing, and utilization of unripe banana flour and its starch.

KEYWORDS

amylolytic resistance, banana starch, enzymatic hydrolysis, granule morphology, unripe banana flour

1 | INTRODUCTION

Resistant starch (RS) can be described as the food starch component that cannot be digested in the mouth and small intestine of a healthy individual (Juarez-Garcia et al., 2006). Unripe banana flour is rich in RS (Thakorlal et al.,

2010) ranging from about 17% to 80% (Bi et al., 2017; Cahyana et al., 2019; Falodun et al., 2019) depending on the cultivar type and its native or processed state. RS consists of physically inaccessible starches classified as resistant starch 1 (RS1); ungelatinized starches classified as resistant starch 2 (RS2); retrograded starches classified as resistant

starch 3 (RS3); modified starches classified as resistant starch 4 (RS4); and starch nutrient complexes classified as resistant starch 5 (RS5) (Okumus et al., 2018; Tian & Sun, 2020). Recently, it was suggested that amylose–lipid complexes (Qin et al., 2019; Tang et al., 2022) and starch–polyphenol V-type complexes (Romero Hernández et al., 2022) should be classified as RS5 due to their ability to hinder the attachment of starch digestive enzymes to starch in the upper gastrointestinal tract. Unripe banana flour native RS is mainly categorized as RS2 (Li et al., 2020) and can be fermented by the microflora in the large intestine (Hernández-Nava et al., 2009).

Starch granules in different plant products, such as bananas, exist in a semicrystalline form and comprise both the amorphous regions, consisting of amylose and branched amylopectin, as well as the long-chain amylopectin crystalline regions that twist into H-bonded double helices (Butterworth et al., 2011). Classification of these starches according to their amylose/amylopectin results in normal starches with 25%–30% amylose and 70%–75% amylopectin; waxy starches with high amylopectin content of 98%–99%; and high-amylose starches with 50%–70% amylose content (Bello-Perez et al., 2020). This grouping ultimately influences the resistance of the starch to enzymatic hydrolysis. Wang et al. (2022) noted that factors including starch concentration, source, type, and concentration of enzymes, as well as amylose content, influence amylolysis. The RS fraction is therefore starch that resists alpha-amylase and pullulanase hydrolysis in vitro beyond 120 min compared to rapidly digestible and slowly digestible starch (Lockyer & Nugent, 2017). Thus, digestion of RS does not take place in the upper gastrointestinal tract, but RS is fermented by microbes in the colon, which produce short-chain fatty acids (SCFAs) such as acetate, butyrate, isobutyrate, isovalerate, and propionate (Koh et al., 2016; Meenu & Xu, 2019); gases such as methane, hydrogen, and CO₂; as well as alcohols including methanol and ethanol (Birt et al., 2013). Produced SCFAs are then utilized as a source of energy in the body and are of great importance to colonic health (Phothisoot et al., 2023). Butyrates formed through the fermentation of RS are the preferred matrix by colonocytes. The SCFA butyrates have also been implicated in increasing colonic blood circulation, reducing luminal pH, and preventing the formation of abnormal colonic cell populations (Lockyer & Nugent, 2017). Products of RS fermentation are absorbed from the gut into the blood flow where they are transported into the adipose and skeletal muscle tissue and undertake a unique role in improving insulin sensitivity (Lockyer & Nugent, 2017).

Currently, RS is classified as a prebiotic due to its fermentation by microbes in the colon and the corresponding production of SCFAs, which benefit the host (Moore et al., 2015). The major phyla that have been implicated in RS

fermentation include *Actinobacterium*, *Bacteroidetes*, and *Firmicutes* (Birt et al., 2013). Other RS hydrolyzing strains identified in fecal samples of humans include *Bifidobacterium* spp., implicated for lactate and acetate production; *Bacteroides* spp., implicated for acetate and propionate production; as well as *Fusobacterium* and *Butyrivibrio*, implicated for butyrate production (Meenu & Xu, 2019). Intake of RS in diets has also been linked to an increase in the number and activities of microbes such as *Lactobacillus*, *Bifidobacterium*, *Lachnospiraceae*, *Ruminococcaceae*, and *Clostridium* spp. (Zeng et al., 2017; Meenu & Xu, 2019). RS thus functions as both a prebiotic and a symbiotic by promoting the growth and actions of probiotic bacteria as well as interacting with other prebiotic dietary fibers such as beta-glucan (Raigond et al., 2015). Upon absorption, RS undertakes some of its probiotic roles by protecting ingested organisms on their way to the colon, thus increasing the amount of these beneficial organisms in the colon (Raigond et al., 2015).

The RS present in unripe banana flour can modulate metabolic activity and hence provide protective effects against illnesses such as cardiovascular diseases, cancer, diabetes, and obesity (2019Khoosani et al., 2019; Li et al., 2020; Sidhu & Zafar, 2018; Singh et al., 2016). RS has been shown to reduce the incidence of noncommunicable diseases, lowering GI and cholesterol, as well as exerting positive effects against cardiovascular diseases, colorectal cancer, diabetes, and obesity (Phothisoot et al., 2023). Furthermore, RS, which is classified as a dietary fiber, contributes to fecal bulking and hinders the growth of cancerous lesions of the bowel in vivo (Moore et al., 2015). Rabbani et al. (2009) showed that a diet containing RS from unripe banana had a positive effect on controlling shigellosis in children after 5 days of feeding, through a decrease in stool volume while increasing the concentrations of acetate, butyrate, and propionate in the feces of the children. In the United States, the daily recommended intake of RS is 6 g RS/meal. However, Americans are reported to consume an estimated daily intake of ~5 g RS/day (Birt et al., 2013). Despite the beneficial health effects derived from unripe banana flour-resistant starch (UBFRS), there exists a paucity of literature attempting to integrate and elucidate the mechanisms by which unripe banana flour starch resists enzymatic hydrolysis. Therefore, this review aims to elaborate on the intricacies of the resistance mechanisms of unripe banana flour starch to enzymatic hydrolysis.

2 | AMYLOLYSIS RESISTANCE IN STARCH

The enzymatic hydrolysis of starch has been shown to occur at the interface between the solid phase, which is the starch, and the liquid phase, which is the conveyed enzyme

acting on the starch (Wang et al., 2022). Naguleswaran et al. (2014) opined that the occurrence of amylolysis, which is the hydrolysis of starch granules by amylases, takes place via diffusion to a solid surface, adsorption, and catalysis. Furthermore, amylopectin molecules that possess greater levels of short chains and degree of branching but have more compact structure, high molecular density, high molar mass, and smaller molecular size are less susceptible to amylolytic attack (Naguleswaran et al., 2014). Enzymatic hydrolysis of amylose and amylopectin molecules is therefore determined by the structure of the granules or the arrangement of the crystallites within the granules (Dhital et al., 2015). In determining the mechanism of resistance to amylolytic attack, Ma et al. (2020) proposed that the presence of double helix order within or outside the crystal structure; the stabilization of the double helices into perfect crystalline structures; the presence of unevenly packed or isolated single helices; the entrapment of amorphous region within imperfect crystals; and the high molecular order formed by the attachment of short branch length amylopectin to the linear chain through covalent bonds are some mechanisms responsible for the resistance of starch to amylolytic attack.

Starch when heated in the presence of sufficient water undergoes gelatinization and increased digestibility. Upon cooling, high-amylose starches retrograde, forming crystalline regions that are inaccessible to enzymatic hydrolysis (Birt et al., 2013). The A-, B-, and C-type crystalline arrangement patterns of the different polymeric forms affect amylase hydrolysis (2018). It has been observed that the shorter A-type double helices and interior crystallites are more vulnerable to enzymatic hydrolysis than the longer chains and stable helices, which are seen to be more resistant (Singh et al., 2010). The surface properties of starch granules are other contributory factors to the enzymatic hydrolysis of starches (2018). Dhital et al. (2015) stated that cracks, pores, interior channels, and surface damage in starch granules further exacerbate enzymatic adsorption and binding to these molecules. Singh et al. (2010) showed that the presence of pores and pinholes in starches facilitates hydrolytic attack through penetration by amylases, thus resulting in endocorrosion, while starches (cooked or native) with smooth surfaces and without pores displayed very high resistance to enzymatic hydrolysis. However, starches from other food sources such as potatoes have been reported to be eroded through exocorrosion, which involves erosion of the granule surface during enzymatic hydrolysis.

Mass transfer limitations occurring due to the viscosity of the starch are known to influence the hydrolysis rate of starch (Singh et al., 2010). Available points for enzymatic attacks are increased due to a higher degree of branching. However, this higher degree of branching

may lead to an increase in steric hindrance (implicated in hindering enzymatic hydrolysis) and hence the mass transfer resistance. The molecular weight distribution of starches has also been reported as affecting enzymatic hydrolysis. Starches with higher molecular weight can experience sharp hindrances at the active centers of the enzyme (Singh et al., 2010). The large size of the starch granule compared to the alpha-amylase molecule provides avenues for attack by amylase by making available many possible sites for attachment of the enzyme (Butterworth et al., 2011). Similarly, the amylose-amylopectin ratio has been found to affect enzymatic hydrolysis as amylopectin is more easily digested than amylose in native starches. Das et al. (2022) showed that enzymatic treatment of banana flour led to increased amylose content in the modified flour due to amylopectin digestion by amylopullulanase. Amylopectin digestion has been attributed to its larger size with a greater surface area per molecule when compared to amylose (Singh et al., 2010). Furthermore, the glucose chains of amylose that are compactly held together by hydrogen bonds make amylose less susceptible to enzymatic hydrolysis than amylopectin with more branched glucose chains (Singh et al., 2010). Non-starch ingredients including proteins and lipids reduce the surface accessibility of the granules by blocking the adsorption sites of the binding enzymes (Singh et al., 2010). Amylase hydrolysis can only occur by the binding of glucan chains through some glucose units to their subsites located at the active center, with the number of these subsites in an active center varying from four to nine. Wang et al. (2022) stated that different enzymes including alpha-amylase, beta-amylase, isoamylase, glycosyl-transferase, and glucosidase are involved in the enzymatic hydrolysis of starch. However, the hydrolysis of alpha-(1,4) glycosidic bonds in starch polymers is catalyzed by alpha-amylase through endocorrosion.

3 | BANANA FLOUR RS CHARACTERISTICS AND AMYLOLYTIC RESISTANCE

3.1 | Starch molecular composition

Starch, which is made up of amylose and amylopectin, comprises a highly branched amylopectin molecule whose glucose bonds are joined by α -1,4 and α -1,6 linkages (Nasrin & Anal, 2014a). The α -1,6 branching points of amylopectin occur at every 24–30 glycosyl units and function as entry points for enzymes (Giacco et al., 2016; Nasrin & Anal, 2014a). The average number of UBFRS glycosyl units for amylopectin side chains was estimated to be 14.5 (Faisant et al., 1995; Langkilde et al., 2002), but

more recent estimates place them at 5.0–20.3 glucosyl units (Yee et al., 2021). It has been reported that the average degree of polymerization (DP_n) of banana RS was lower than that of jackfruit starch (Zhang et al., 2019), wheat amylopectin (Yoshio et al., 2011), and barley amylopectin (Tang et al., 2002). Furthermore, it was stated that UBFRS has less branching than other starches such as jackfruit and wheat starch (Bi et al., 2017; Das et al., 2022; Jiang et al., 2015; Yee et al., 2021). Longer chain branches of amylopectin produce more crystalline structures due to the presence of hydrogen bonds among such chains (Jane et al., 1997). However, Bi et al. (2017) and Chávez-Salazar et al. (2017) suggested that the amylopectin architecture can vary between cultivars. Hence, it could be imperative to consider the genome groupings during the selection of bananas for unripe banana flour production given their influence on the architecture of amylopectin chains.

The amylose content of native nonmodified banana starch ranges from 30% to 41.66% (Bi et al., 2017; Das et al., 2022). Pelissari et al. (2012) demonstrated that banana starch with 35.1% amylose content had higher RS content (50.3%) compared to flour with 23.1% amylose content (45.1% RS). Unripe banana flour starch amylose is a slightly branched molecule whose glucose bonds are joined mainly by α -1,4 linkages (Magallanes-Cruz et al., 2017). Amylose has very limited α -1,6 glucosyl bonds compared to amylopectin (Giacco et al., 2016); therefore, the presence of a limited number of α -1,6 glucosyl bonds could result in amylose undergoing slower hydrolysis compared to amylopectin. It has also been suggested that amylose is less digestible compared to amylopectin because it has a limited surface area exposed to digestion due to its ability to interact with other components such as lipids (Englyst & Cummings, 1985; Jenkins et al., 1988). Amylose–lipid complexes and amylopectin–lipid complexes (to a lesser extent) lead to higher resistance to hydrolysis due to the formation of more resistant V-type crystalline structures (RS5) (Wokadala et al., 2012). Hence, a higher amylose content would lead to a higher resistance to hydrolysis.

Nevertheless, the enzyme resistance capacity of banana starch has been attributed to the ratio of amylose and amylopectin present in unripe banana flour (Bi et al., 2017; Soares et al., 2011; Vatanasuchart et al., 2012). Conventionally, starch consists of 20%–30% amylose and 70%–80% amylopectin (Abdullah et al., 2018); however, the amylopectin content of unripe banana flour is generally higher than its amylose content (Kumar et al., 2019; Salazar et al., 2022). The amylose–amylopectin ratio influences the glycemic index (GI) of starchy foods (Björck et al., 1994); hence, a high amylose–amylopectin ratio is desirable as it improves the RS content of composite foods (Åkerberg et al., 1998). Ademosun et al. (2021) showed that the GI value of wheat noodles supplemented with unripe plan-

tain flour and orange peels (35.03–53.09) was significantly lower than that of the control samples without unripe banana flour (58.23–66.21). A higher amylose content in unripe banana flour used in the study could have resulted in an increased RS content, thus leading to the lowering of the GI value.

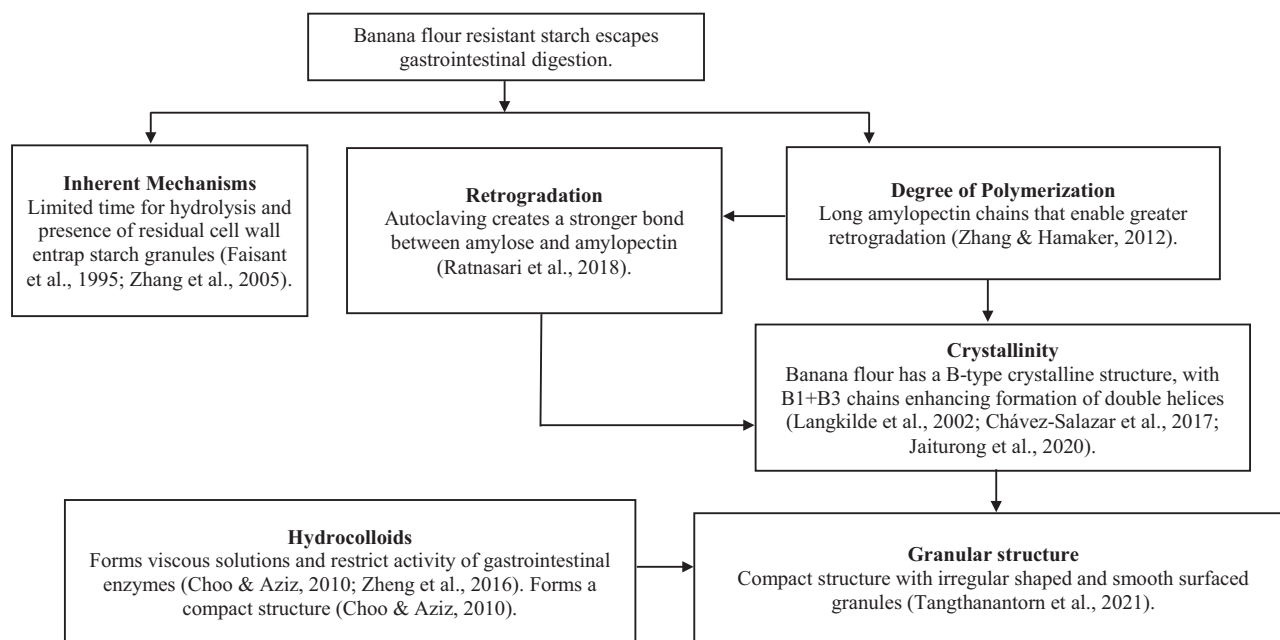
3.2 | Supramolecular structure

The structural properties of unripe banana flour and banana RS are summarized in Table 1. X-ray diffraction assessments showed that UBFRS (RS2) has a B-type crystalline structure (Faisant et al., 1995; Langkilde et al., 2002), which possesses a characteristically lower degree of branching, higher crystallinity, and longer amylopectin chains compared to the A-type crystalline structure (Chávez-Salazar et al., 2017; Kim et al., 2015; Martens et al., 2018). The longer amylopectin chains present in the B-crystalline structure produce a more crystalline structure and increased crystallinity compared to the A-type structure by enhancing the formation of double helical structures (Chávez-Salazar et al., 2017). Higher crystallinity results in the formation of a compact granular structure present in UBFRS, which hinders the starch granules (RS2) from being invaded by enzymes, resulting in reduced susceptibility to enzymatic hydrolysis (Ma & Boye, 2018; Tangthanantorn et al., 2021). Unlike the A-type crystalline structures, the B-type crystalline starch (Figure 1) tends to have a less porous structure because its branching sites are predominantly located in the amorphous zones (Kim et al., 2015). In addition, the longer B-type amylopectin chains are thought to further contribute to resistance through interactions with lipids and amylose, further preventing hydration and allowing for more extensive retrogradation (Zhang & Hamaker, 2012).

However, not all UBFSs consist of B-type crystalline structures (Table 1). Native RS2 prepared from Brazilian and Taiwanese banana cultivars have been reported to contain both B- and C-type crystalline structures (Cordoba et al., 2018; Wang et al., 2017). Cordoba et al. (2018) showed that Terra Plátan (AAB) displayed a B-type crystalline structure, while Catura cavendish (AAA) and Prata Ana (AAB) cultivars displayed a C-type crystalline structure. Likewise, *Musa* AAA Cavendish and *Musa* ABB Bluggoe RS (RS2) displayed a C-type crystalline structure, while *Musa* ABB Pisang Awak and *Musa* AA Pisang Mas RS (RS2) displayed a B-type crystalline structure (Wang et al., 2017). The relative resistance in C-type nonmodified UBFS compared to A-type crystallinity starches results from the fact that the C-type crystalline structures are structurally intermediate between the A-

TABLE 1 Degree of polymerization, crystallinity, and granular morphology of banana flour resistant starch.

Type	Description	Degree of crystallinity	Type of crystallinity	Size	Shape	Degree of polymerization	Reference
RS2	Native (uncooked)		B-type			14.5 glucosyl units	Langkilde et al., 2002
RS2	Native (uncooked)				Intact, irregularly shaped granules with a smooth surface.		Thakorlal et al., 2010
RS2	Native (uncooked and centrifuged)	21.76–27.75	B- and C-type	29.07–39.17 μm	Ellipsoidal and flattened shape		Cordoba et al., 2018
RS3	Enzymatically modified		C _A -type		Tough and rigid.		Das et al., 2022
RS3	Autoclaved and debranched		C-type		Cavities and channels in starch granules		González-Soto et al., 2007
RS3	Modified (autoclaved, debranched, and enzymatically modified)				Irregular shaped		Khawas & Deka, 2017
RS2	Native (centrifuged/uncooked)		B- and C-type	5–90 μm	Elliptic, triangular shaped with smooth surfaces		Wang et al., 2017
RS2	Raw (uncooked)					18.9–20.3 glucosyl units	Yee et al., 2021

**FIGURE 1** Mechanisms of resistance of unripe banana flour starch to enzymatic hydrolysis.

and B-type and thus would have relatively higher resistance compared to type A (Ma & Boye, 2018). Modified (autoclaved, debranched, and enzymatically modified) UBFRS (RS3) displays C- and C_A-type crystalline structure (Das et al., 2022; González-Soto et al., 2007). The C_A-type

could therefore be more susceptible to enzymatic hydrolysis compared to the C-type crystalline structure due to the presence of short amylopectin chains, evidence of porous surfaces, and surface area effects, which result in high enzymatic hydrolysis.

4 | GRANULAR MORPHOLOGY AND ENZYMATIC HYDROLYSIS RESISTANCE

4.1 | Granule shape

There are diverse reports on the granular shape and surface structure of unripe banana flour starch (Table 1). The raw UBFRS-RS2 possesses intact, irregular, and ellipsoidal-shaped starch granules with smooth surfaces (Figure 1), whose structures are dependent on the banana cultivar (Cordoba et al., 2018; Thakorlal et al., 2010; Wang et al., 2017). Pelissari et al. (2012) demonstrated that RS2 present in unripe banana flour and starch granules displayed irregularly shaped and compact granules in both elongated and spheroid forms. Hoffmann Sardá et al. (2016) showed that raw unripe banana flour from Brazilian banana cultivars had spherical and elongated granules. Several reports provide evidence of the influence of cultivar type on the granular morphology of unripe banana flour starch. For example, Gao et al. (2016) reported that Cavendish had an irregular-shaped structure, while plantain had small and ellipsoidal granules. Anyasi et al. (2017) reported that *Luvhele* and *Mabonde* banana flour had polygonal and oval-shaped granules, while *Muomva-red* and *William* banana flour had elongated, polygonal, and spherical-shaped granules. Soares et al. (2011) found that desert banana cultivars (Pacovan and Mysore) had small and leaf-like structures, while plantains (Terra and Figo) had round and elongated structures.

Irregular- and ellipsoidal-shaped starch granules tend to be more susceptible to enzymatic hydrolysis compared to other starch granules because they have a larger surface area and volume (Gao et al., 2016). Likewise, leaf-like starch granules found in unripe bananas tend to have higher susceptibility to enzymatic hydrolysis relative to rounded granules due to the arrangement of amylopectin molecules (Soares et al., 2011). The presence of smooth surfaces with no signs of degradation has been reported on the surface of Terra, Figo, and Pacovan banana cultivars (Soares et al., 2011). The presence of smooth surfaces is thought to restrict the invasion of starch granules by α -amylase. However, this explanation may not be generally applicable given that pits have been reported on the surface of the granules of Mysore banana cultivars, yet they have similar resistance to digestion as cultivars with smooth surfaces (Soares et al., 2011).

The morphology of starch granules and their susceptibility to enzymatic hydrolysis can further be influenced by food processing techniques such as heating, milling, and baking (Anyasi et al., 2017). Autoclaved and enzymatically (amylopullulanase) modified UBFRS (RS3) samples displayed irregularly shaped granules with porous, tough, and rigid surfaces (Das et al., 2022; González-Soto et al.,

2007). Similar granular morphologies were reported for autoclaved, debranched, and enzymatically treated banana RS (Khawas & Deka, 2017). The presence of pores on the surface of processed starch granules provides a pathway for endocorrosion. Hence, autoclaved and enzymatically (amylopullulanase) modified banana RS (RS3) samples have increased susceptibility to enzymatic hydrolysis compared to native banana RS (RS2) due to differences in granule structure.

4.2 | Granule size

The diameter of unripe banana flour starch granules influences its susceptibility to enzymatic hydrolysis. Unripe banana flour starch granules have been reported to possess varied diameters ranging from 1.4 to 500 μm (Bezerra et al., 2013; Falade & Olugbuyi, 2010; Gao et al., 2016; Giraldo-Gómez et al., 2019; Khoozani et al., 2020; Padhi & Dwivedi, 2022; Salazar et al., 2022; Thakaeng et al., 2021). It has been suggested that small-sized starch granules are more susceptible to enzymatic hydrolysis compared to large-sized starch granules due to their large surface area to volume ratio (Gao et al., 2016; Jiang et al., 2015), while large rounded and elongated granules may show few signs of degradation (Soares et al., 2011).

Foam-mat, hot-air, oven, and sun drying tend to disrupt the molecular order of starch granules in banana flour due to high heat levels and longer drying times, resulting in relatively small-sized granules (Falade & Olugbuyi, 2010; Khoozani et al., 2020). Furthermore, it was also reported that the granule diameter (20–47.3 μm [Espinosa-Solis et al., 2009; Pelissari et al., 2012]; 5–90 μm [Wang et al., 2017]) of native (raw) unripe banana RS (RS2) was greater than that of enzymatically modified UBFRS (RS3) (20 μm) (Das et al., 2022).

5 | NONSTARCH COMPONENTS AND THE SUSCEPTIBILITY OF UNRIPE BANANA STARCH TO ENZYMATIC HYDROLYSIS

5.1 | Native hydrocolloids (proteins and nonstarch polysaccharides)

Hydrocolloids are long-chain protein or polysaccharide polymers with the ability to bind water and form gels or viscous dispersions (Woomer & Adedeji, 2021). Hydrocolloids interact with starch granules by forming physical barriers and viscous solutions around the surface of granules, resulting in the reduction of the activity of gastrointestinal enzymes (Faisant et al., 1995; Tester et al., 2006; Zheng

et al., 2016). Interactions of starch granules and non-starch polysaccharides have been shown to affect the rate and extent of digestion of starch by prolonging the time required for effective hydrolysis (Zhang & Hamaker, 2012; Zhang et al., 2005).

The protein content of unripe banana flour is generally reported to be low (<5%) (Kumar et al., 2019; Rachman et al., 2021; Thakaeng et al., 2021), with higher concentrations (>5%) reported in the peels (de Angelis-Pereira et al., 2016; do Prado Ferreira & Teixeira Tarley, 2020), as well as with increasing ripeness of the fruit (Campuzano et al., 2018). Miao et al. (2015) showed that the presence of proteins in unripe banana flour starch contributes to amylolytic resistance, as proteins interact with starch to reduce the rate of α -amylase hydrolysis. The interaction is thought to result in the formation of a three-way complex between fatty acids, amylose, and proteins (Shah et al., 2011; Zhang et al., 2010). This interaction occurs even with low protein concentrations (3.68%–8.02%), which could be up to 10 times less than starch (70.0%–84.4%) (Campuzano et al., 2018; Zhang et al., 2003), thus enabling, to a certain degree, resistance to enzymatic hydrolysis in unripe banana flour.

Fiber increases the viscosity of foods and limits enzymatic digestion by reducing access of gastrointestinal enzymes to the substrates and thereby reducing the conversion rate of starch into glucose (Abutair et al., 2016). The pectin content of unripe banana flour has been reported to range between 3.29% and 5.61% (Bi et al., 2017). Pectin is regarded as a hydrocolloid alongside gelatin, alginate, carrageenan, gellan, and agar (Goff & Guo, 2019; Milani & Maleki, 2012; 2010), as well as a soluble fiber (Martínez et al., 2015). Insoluble dietary fiber components including lignin, cellulose, and hemicellulose have been reported to be present in unripe banana flour (Silva et al., 2020; 2014). Ng et al. (2020) suggested that the barrier formed by soluble fiber protects starch granules against water penetration, which limits the swelling and gelatinization capacity of starch and reduces the ability of enzymes to access the starch due to limited entry or attachment points.

5.2 | Lipids

The rate and extent of digestion of starch are also influenced by its lipids composition and extent of phosphorylation (Qi & Tester, 2016; Qi et al., 2018). The presence of lipids in unripe banana flour has been reported by several authors and is said to be $\leq 2.0\%$ (w/w) (Dan, 2011; Haslinda et al., 2009; Hoffmann Sardá et al., 2016; Kongolo et al., 2017; Menezes et al., 2011). Banana peel possesses higher lipid content when compared to the pulp; hence, the lipid content of unripe banana flour with peels is higher than without peels (Haslinda et al., 2009; Hoff-

mann Sardá et al., 2016). Interaction of amylose and lipids through amylose–lipid complex formation reduces susceptibility to hydrolysis by α -amylase (Wokadala et al., 2012). Lipids may also reduce α -amylase hydrolysis complexing with longer amylopectin chains (Tester et al., 2006). It has been suggested that the interaction of lipid and amylose induces a change in the torsion angles of glycosidic bonds and forms a helical structure, which reduces the binding activity of amylolytic enzymes and the formation of enzyme–substrate complexes (Chi et al., 2022; Li et al., 2021). However, the degree to which lipid complexation contributes to the hydrolysis resistance of banana flour starch is not clear and requires further investigation.

5.3 | Polyphenols

Numerous studies have reported the presence of polyphenols in unripe banana flour (Anyasi et al., 2018; Pico, Corbin, et al., 2019; Pico, Xu, et al., 2019). The polyphenols reported to be present in bananas include anthocyanins (cyanidin, delphinidin, and their derivatives in both fruit pulp and peel), chlorogenic acid, ferulic acid, flavan-3-ols (catechins, epicatechins, and gallic acid), flavonoids (myricetin and quercetin), and tannins (Anyasi et al., 2018; Bennett et al., 2010; 2006 Maseko et al., 2024; 2015 Pico, Corbin, et al., 2019; Pico, Xu, et al., 2019). Phenolic compounds have been reported to be present in both banana peel and pulps (Castelo-Branco et al., 2017), although banana peels contain more polyphenols compared to the pulp (Agama-Acevedo et al., 2016; Castelo-Branco et al., 2017; Maseko et al., 2024).

Polyphenols facilitate the escape of starch from digestion in the small intestine by inhibiting the activity of α -amylase and glucosidase, resulting in intact starch being transported into the colon (Gu et al., 2020; Liu et al., 2017). Furthermore, polyphenols can interact with starch molecules via covalent or noncovalent bonds to form starch–polyphenol complexes (Chi et al., 2022). The presence of starch–polyphenol complexes has been suggested to occur in unripe banana flour (Choo & Aziz, 2010; Pico, Xu, et al., 2019; Sarawong et al., 2014); hence, unripe banana flour is considered a natural source of starch–polyphenol complexes (Hernández et al., 2022). Polyphenols contain hydroxyl groups that enable them to interact with starch via noncovalent bonds such as hydrogen bonds and electrostatic and ionic interactions (Chi et al., 2017; Giuberti et al., 2020; Ngo et al., 2022).

Polyphenols can reduce the activity of enzymes and starch digestibility by binding to starch-digesting enzymes and forming V-type inclusion complexes through hydrophobic interaction and hydrogen bonding with amylose and amylopectin side chains (Han et al.,

2020; Hernández et al., 2022; Kan et al., 2022; Ngo et al., 2022). The extent to which the polyphenol inhibition mechanism contributes to the overall resistance to enzymatic hydrolysis is, however, difficult to ascertain (Hernández et al., 2022). Providing more information on the mechanisms responsible for starch–polyphenol complexes will enhance the exploitation of polyphenols in increasing the resistance of unripe banana flour starch. Furthermore, optimizing processing conditions to maintain the structure of the starch–polyphenol complexes present in unripe banana flour and increasing the production and application of banana peel flour as a standalone functional ingredient or composited with banana pulp flour will enhance the application of polyphenols for increasing the resistance of unripe banana flour starch.

6 | GENOME GROUPINGS AND THE RS CONTENT OF UNRIPE BANANA FLOUR

The effects of the genomic and subgenomic group and cultivar type on the RS content of unripe banana flour are summarized in Table 2. Kumar et al. (2019) found that the RS content of unripe banana flour prepared from cultivars belonging to the ABB genomic group (44.4%–46.5%) was greater than those in AAA (32.7%) and AAB genomic group (31.3%–35.6%). Vatanasuchart et al. (2012) showed that the RS content of unripe banana flour from the BBB (68.1%) genomic group was greater than the AA (52.2%–57.2%), AAA (57.0%), and ABB (56.6%–61.8%) genomic groups. It has been suggested that the B genomic group tends to have more RS content compared to the A genomic group due to its high starch and protein content, larger granule size, and higher amylose content (Annor et al., 2016; Gibert et al., 2014; Narayana et al., 2017)

However, recent studies have suggested that the genomic group has no significant effect on the RS content of unripe banana flour. Chang et al. (2022) found that there was no significant difference in the RS content of the total starch composition of unripe banana flour from cultivars belonging to the AA (90.38% dry basis of total starch), AAA (89.20%–90.86% dry basis of total starch), and ABB (90.58% dry basis of total starch) genomic groups. This conforms with the findings of Reis et al. (2019), who suggested that the genomic group had no effect on the RS content of unripe banana flour. Kongolo et al. (2017) found that for cultivars grown under the same agronomic conditions, the RS content varied. The RS content of the AA genomic group (39.6%–47.3%) was similar to that of the AAA (38.6%–47.1%) and AAB (39.2%) genomic groups but slightly higher than the AAAB genomic group (19.9%–47.4%). Cordoba et al. (2018) also found that there

was an overlap in the RS content of AAB (24.79%–34.85%) and AAA (31.92%) genomic groups. Other studies on unripe banana flour suggested that cultivars from the AAA genomic group with RS content of 4.83%–57.49% (Campuzano et al., 2018; Khoozani et al., 2019; Menezes et al., 2011; Rayo et al., 2015; Rosado et al., 2020; Tian et al., 2020) had RS content that overlapped with that of the ABB cultivar types, which ranged from 8.60% to 68.8% (Haslinda et al., 2009; Nasrin et al., 2015; Rodríguez-Ambriz et al., 2008; Vatanasuchart et al., 2012). The effect of the genomic group could therefore be attributed to differences in agronomic conditions, maturity at harvest, as well as postharvest processing.

7 | EFFECT OF MODIFICATIONS ON THE RS CONTENT OF UNRIPE BANANA FLOUR

The effect of physical and chemical modifications on the RS content of unripe banana flour is summarized in Table 3 and Figure 2.

7.1 | Physical modifications

Heat–moisture treatment (HMT) generally reduces the RS content of unripe banana flour. Heat disrupts the molecular order of starch granules resulting in irreversible changes such as granule swelling, melting of native crystalline structure, starch solubilization, and loss of birefringence (Anyasi et al., 2017). Zhang and Hamaker (2012) demonstrated that cooking at 100°C for 10–20 min reduced the RS content of banana flour. De la Rosa-Millan et al. (2014) showed that the RS content of flour prepared from bananas that had been cooked for 5 min was 61.5% compared to 66.5% in raw banana flour. Similarly, Rodríguez-Damian et al. (2013) found that the RS content of banana flour cooked for 5 min (58.5%) was greater than that of banana flour samples cooked for 15–25 min (23.3%–23.7%). Given that the RS content of unripe banana flour tends to decrease with cooking time (De la Rosa-Millan et al., 2014; Rodríguez-Damian et al., 2013), shorter cooking times could be applied to limit the destruction of the structural integrity of starch granules and thus minimize the reduction in RS content of banana flour.

Annealing and HMT followed by storage have been shown to improve the RS content of unripe banana flour (2010). The RS content of unripe banana flour subjected to annealing and storage was greater than that of cooked banana flour samples (De la Rosa-Millan et al., 2014). Cahyana et al. (2019) demonstrated that annealing produced higher RS content in unripe banana flour

TABLE 2 Effect of genomic and subgenomic group classification on unripe banana flour (UBF)-resistant starch content.

Genomic group	Subgenomic group	Cultivar	Resistant starch content (%)	References
AA	Inarnibal	Pisang Lemark	47.30	Kongolo et al., 2017
AA	Sucrier	Lady Finger	39.60	
AAA	Cavendish	Gross Michel	37.90	
AAA	Cavendish	IBP. 5.61	47.10	
AAA	Sucrier	Khai Thong	42.10	
AAA	Red	Green red	38.90	
AAB	Pome	Pome	39.20	
AAAB	Pome synthetic hybrid	FHIA 18	19.90	
AAAB	Goldfinger selection	Goldfinger	47.20	
AAAB	Goldfinger selection	PKZ	47.40	
AAA	Cavendish	Grand Naine	32.70	Kumar et al., 2019
ABB		Monthan	44.35	
ABB		Saba	46.50	
AAB		Nendran	35.55	
AAB		Populu	31.28	
ABB	Plantain	Mzuzu	90.58	Chang et al., 2022
AAA	Cavendish	Malindi	90.86	
AA	Pisane litin	Mshale	90.38	
AAA	Cavendish	Bukoba	90.69	
	Dwarf red banana	<i>Guihongjiao No. 1</i>	86.15	Bi et al., 2017
ABB	Pisang Awak	<i>Jinfen No. 1</i>	85.88	
AAA	Cavendish	<i>Williams B₆</i>	90.46	
AAA	Cavendish		57.48 (standard UBF)	Hoffmann Sardá et al., 2016
AAA	Cavendish		43.25 (standard peel UBF)	
AAA		Kluai Hom	57.20	Vatanasuchart et al., 2012
AA		<i>Kluai Khai</i>	52.20	
AA		<i>Kluai Lembuenang</i>	57	
ABB		<i>Kluai Namwa</i>	56.60	
ABB		<i>Kluai Hakmak</i>	61.80	
BBB		<i>Kluai Hin</i>	68.80	
AAB		<i>Terra Pla'tano</i>	34.85	Cordoba et al., 2018
AAA		<i>Caturra Cavendish</i>	31.92	
AAB		<i>Prata Ana</i>	24.79	
AAA	Cavendish	<i>Nanicão</i>	40.90–58.50	Tribess et al., 2009
AAA	Cavendish	<i>Nanica</i>	32.20–46.72	Khoozani et al., 2019

(Continues)

TABLE 2 (Continued)

Genomic group	Subgenomic group	Cultivar	Resistant starch content (%)	References
AAA	Cavendish	<i>Nanicão</i>	48.99	Menezes et al., 2011
AAA	Cavendish	<i>Nanicão</i>	53.95–57.49 (unmodified)	Rayo et al., 2015
AAA	Cavendish		33.86	Bezerra et al., 2013
AAA	Cavendish	<i>Pei-Chiao</i>	30.30	Liao & Hung, 2015
	Cavendish	<i>Nanicão</i>	38.92–44.27	La Fuente & Tadini, 2017
AAA	Cavendish		22.80	Rosado et al., 2020
AAA			11.12–38.28	Campuzano et al., 2018
AAA	<i>Musa acuminata</i> Colla		4.33	Tian et al., 2020
AAA		<i>Nanicão</i>	8.20	Menezes et al., 2010
AAA	Gros Michel	<i>Kluai Hom Thong</i>	41.74–46.35	Virulchatapan & Luangsakul, 2020
AAA		<i>Nanicão</i>	8.20	Menezes et al., 2010
AAB			30.40	Rodríguez-Ambriz et al., 2008
AAB	Harton		31	Gutiérrez, 2018
ABB	<i>Sapientum</i>	<i>Kluai Nam Wa</i>	32.26–48.88	Moongngarm et al., 2014
	<i>Musa paradisiaca</i> L.		17.50	Juarez-Garcia et al., 2006
	<i>Musa paradisiaca</i>	<i>Terra</i>	49.50	Pelissari et al., 2012
ABB	Awak		39.50	Haslinda et al., 2009
			39.50	
ABB	<i>Musa paradisiaca</i>		67	
		<i>Pei Chiao</i>	42.23	Li et al., 2020
		<i>Fomosana</i>	34.00	
		<i>Tai-Chiao</i>	30.03	
		<i>Nam</i>	40.25	Ramos et al., 2009
		<i>Fhia 01</i>	10.01	
		Chifre de Vaca	61.05	Reis et al., 2019
		Comprida	60.79	
		Curare Enano	68.50	
		Mongolo	65.92	
		Pinha	53.12	

(Continues)

TABLE 2 (Continued)

Genomic group	Subgenomic group	Cultivar	Resistant starch content (%)	References
		Red Yade	43.44	
		Samura B	52.33	
		Terra Ana Branca	46.26	
		Terra Ponta Aparada	56.13	
		Terra Sem Nome	21.15	
		Tipo Velhaca	55.91	
		Trouis Vert	60.59	
		Terra Maranhão	62.10	
		D'Angola	46.35	
		Terrinha	40.14	
		BRS SCS Belluna	59.14	
		BRS Platina	67.43	
	Grand Naine		51.37	
		Pacovan	70.06	
		Prata Anã	58.30	
	Plantain	<i>Red Essong</i> (French bunch)	33.30–50.23	Udomkun et al., 2021
	Plantain	<i>Mbouroukou 3</i> - <i>False Horn</i> bunch	35.36–53.11	
	Plantain hybrids	<i>PITA 14</i>	32.72–39.67	
	Plantain hybrids	<i>PITA 27</i>	30.91–49.89	
	<i>Musa sapientum</i> , triploid hybrid banana		7.90–8.90	Chaipai et al., 2018
	<i>Musa paradisiaca</i>		24.60	Hernández- Nava et al., 2011
		<i>Kluai Namwa</i> <i>plantain</i>	8.60	Nasrin et al., 2015

(80%–85%) compared to HMT and dual retrogradation (45%–60%). Annealing has been shown to increase the RS content of unripe banana flour by preserving the compact granule surface and increasing the crystallinity of the starch (Cahyana et al., 2019). Rodríguez-Damian et al. (2013) showed that the RS content of banana flour (24.0%–29.0%) subjected to HMT and storage for 25 min was greater than that of banana flour (23.3%) cooked for 25 min. Bello-Perez et al. (2015) further demonstrated that the RS content of banana flour spaghetti stored for 5 days increased from 3.2% to 7.0% (w/w). HMT minimizes starch granule damage, while storage facilitates the retrogradation of the starch granules by

enabling the reassociation of amylose and the formation of a more crystalline structure, which results in improved RS yields. The increase in crystallinity can also be attributed to the strong bond formed by amylose and amylopectin due to the autoclaving process (González-Soto et al., 2007).

7.2 | Chemical and enzymatic modifications

Gutiérrez (2018) reported that phosphated plantain flour recorded the highest RS content (80%) compared to the acetylated, methylated, and oxidized plantain flour

TABLE 3 Effect of modification type on unripe banana flour-resistant starch content.

Modification type	Modifying agent	Modifying conditions	Resistant starch content (%)	References
Physical modifications				
Cooking	H ₂ O	100°C	10.50–61.50	Zhang & Hamaker, 2012
Cooking	H ₂ O	95°C	12.10–61.50	De la Rosa-Millan et al., 2014
Cooking and annealing	H ₂ O	64°C for 24 h	18.70–64.10	
Cooking, annealing, and storage	H ₂ O	64°C for 24 h; 4°C for 7 days (storage: refrigeration)	23.20–64.30	
Cooking	H ₂ O	95°C for 15–25 min	23.34–58.50	Rodríguez-Damian et al., 2013
Cooking and heat–moisture treatment	H ₂ O	30% distilled water; 120°C for 24 h (heating/drying)	22.90–55.83	
Cooking, heat–moisture treatment, and storage (refrigeration)	H ₂ O	30% distilled water; 120°C for 24 h (heating/drying); –20°C for 7 days (storage: refrigeration)	21.70–59.40	
Heat–moisture treatment	H ₂ O	30% distilled water, 100°C for 8 h (heating/drying), and 4–5°C for 24 h (refrigeration)	45–50	Cahyana et al., 2019
Annealing		70% distilled water (w/w) and water bath at 55°C for 12 h	80–85	
Dual retrogradation		Water suspension (1:5.5 [w/v]); 100°C for 30 min (heating); and 4°C for 48 h (refrigeration)	50–60	
Autoclaving		121°C for 30 min	25.25–30.30	Liao & Hung, 2015
Blanching		100°C for 1 min	38.40–52.21	
Blanching	H ₂ O	100°C for 1 min	37.33–55.11	
Blanching	H ₂ O	100°C for 1 min	34.32–52.35	
Blanching	H ₂ O		32.53–52.67	
Cooked			12.40	
Heat treated (uncooked)			81.19 g	
Cooked			10.90	
Cooked			93.68	
Chemical modifications				
Native			31	Gutiérrez, 2018
Acetylation	Acetic anhydride (CH ₃ CO) ₂ O	7.5 mL = 0.079 mol	39.10	
Carboxymethylation	Monochloroacetic acid (ClCH ₂ CO ₂ H)	4.7 mL = 0.079 mol	39	
Methylation	Dimethyl sulfate (CH ₃) ₂ SO ₄	7.5 mL = 0.079 mol	78	
Oxidation	Hydrogen peroxide (H ₂ O ₂)	6.2 mol—30% (0.079 mol)	23	
Phosphation	Sodium trimetaphosphate (STMP—Na ₃ P ₃ O ₉)		82.10	
Native			40.69	Almanza-Benitez et al., 2015

(Continues)

TABLE 3 (Continued)

Modification type	Modifying agent	Modifying conditions	Resistant starch content (%)	References
Chemical	Hydrochloric acid		44.87	
Native	Thermoplastic plantain flour films		1.99	Gutiérrez & Álvarez, 2016
Crosslinking	Aloe vera gel		2.12–3.05	
Native (Red Essong French bunch)			33.30–50.23	Udomkun et al., 2021
Chemical	Citric acid	10 g/L citric acid for 10 min	37.65–54.32	
Native (Mbouroukou 3 - False Horn bunch)			35.36–53.11	
Chemical	Citric acid	10 g/L citric acid for 10 min	39.44–56.37	
Native (PITA 14)			32.72–39.67	
Chemical	Citric acid	0 g/L citric acid for 10 min	36.28–53.63	
Native (PITA 27)			30.91–49.89	
Chemical	Citric acid	0 g/L citric acid for 10 min	34.40–53.97	
Native (uncooked)			85.67 g	Sánchez-Rivera et al., 2017
Chemical (uncooked)	Citric acid		93.87 g	

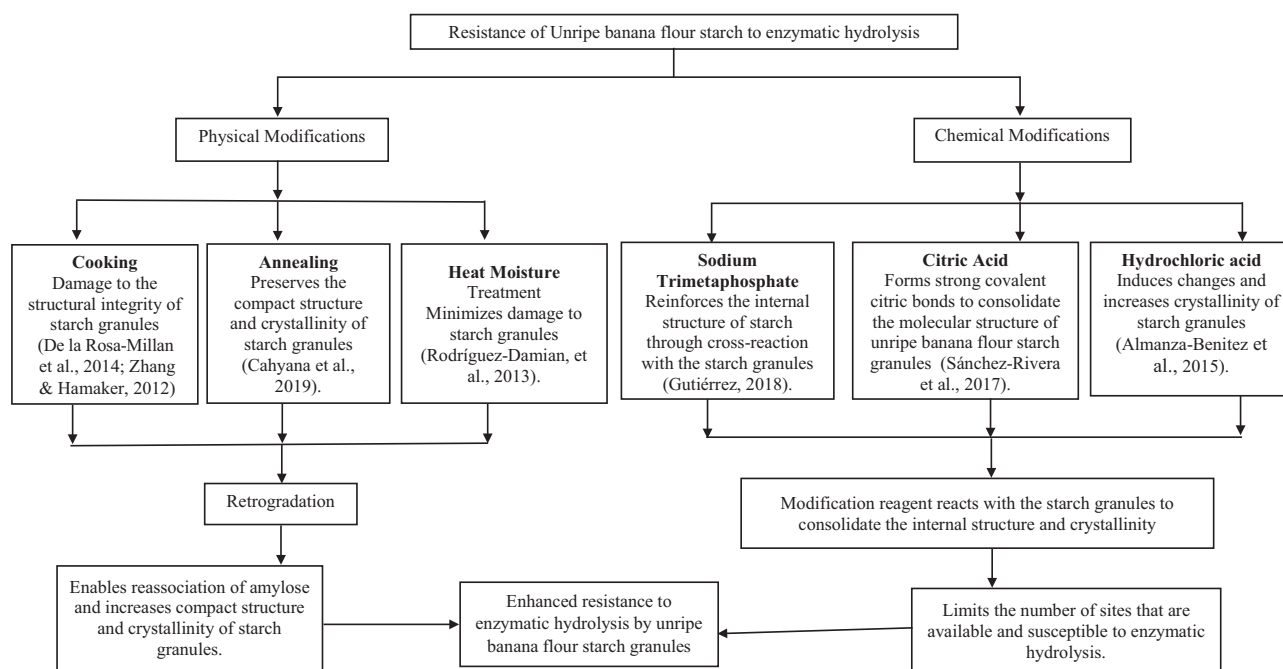


FIGURE 2 Effects of physical and chemical modifications and their associated mechanisms on the resistance of banana flour starch granules to enzymatic hydrolysis.

(23%–78%). The highest RS value of phosphated flour was attributed to the crosslinking reaction of granules with sodium trimetaphosphate (STMP), which was thought to reinforce the internal structure of the starch (Gutiérrez, 2018). Furthermore, Sánchez-Rivera et al. (2017) demonstrated that the RS content of raw and cooked plantain

flour treated with citric acid was greater than native and heat-treated raw and cooked plantain flour. Similarly, Udomkun et al. (2021) found that the RS content of citric acid-treated plantain flours was higher than blanched and native plantain flours. The highest RS content of citric acid-treated plantain flours was attributed to its citric

acid crosslinking effect, which facilitated the formation of strong covalent citric bonds to consolidate the molecular structures of unripe banana flour and limit the number of sites susceptible to enzymatic attacks (Sánchez-Rivera et al., 2017). Moreover, unripe plantain films substituted with *Aloe vera* gel possessed higher RS content compared to native plantain films, thus suggesting that plantain flour films are susceptible to crosslinking with the citric acid present within *Aloe vera* gel (Gutiérrez & Álvarez, 2016). It has been suggested that unripe plantain flour modified with citric acid and phosphate can be used as an ingredient to develop functional gluten-free products due to the high RS content (Gutiérrez, 2018; Udomkun et al., 2021). The RS of plantain flour (44.9%) modified with hydrochloric acid was found to be greater than nonmodified plantain flour (40.69%) (Almanza-Benitez et al., 2015). The increase in the RS level of unripe plantain flour treated with hydrochloric acid was attributed to the hydrolysis of the amorphous zones of starch and an increase in the crystallinity and change in the structure of the starch granules (Almanza-Benitez et al., 2015).

The susceptibility of unmodified unripe banana flour starch to enzymatic hydrolysis is related to changes in thermal properties induced by processing (2010). Several studies have shown that the gelatinization temperature of debranched and enzymatically modified banana flour RS (103–106°C) was greater than the unmodified one (69.1–72.5°C) (Cordoba et al., 2018; Das et al., 2022; Yee et al., 2021). Similarly, the enthalpy of gelatinization (4.4–188.2 J/g) and gelatinization temperature (68.5–147°C) of the debranched, enzymatically modified, and lintnerized banana RS were greater compared to the unmodified UBFRS (6.1–13.9 J/g and 37.6–72.5°C) (Aparicio-Saguilán et al., 2005, 2008; Cordoba et al., 2018; Das et al., 2022; Lehmann et al., 2002; Nasrin & Anal, 2014b; Yee et al., 2021). The RS content of enzymatically modified banana flour using amylopullulanase treatment was observed to be higher at 68.99% when compared to the unmodified banana flour with an RS content of 38.5% (Das et al., 2022). The enzymatic treatment was seen to lead to an increase in amylopectin degradation, thus increasing the amylose content and resulting in a more ordered crystalline state. These findings suggest that chemical and enzymatic modification methods can be employed in developing unripe banana flour and starch-functional foods with high RS content.

8 | CONCLUSION

The resistance to enzymatic hydrolysis of UBFRS can be attributed to a multi-phenomenon mechanism that is primarily dependent on the status of the banana flour: native

or processed. Factors such as the proportion of crystalline and amorphous phases, type, and their distribution; starch granule shape, size, and structure; amylose–amylopectin ratios; and the presence of other native nonstarch components such as proteins, lipids, and polyphenols have all been implicated as other mechanisms known to influence the resistance of both native and modified UBFRS. The DPn, degree of amylopectin branching, and amylose–amylopectin ratio were observed to have more influence on the enzymatic resistance of processed UBFRS compared to the native one. Furthermore, unmodified UBFRS was seen to be less susceptible to enzymatic hydrolysis due to its B-type crystalline structure, irregularly shaped granules, and compact granular structure. However, recent findings demonstrated that modification of unripe banana starch through autoclaving, debranching, and enzymatic treatment results in greater susceptibility to enzymatic hydrolysis due to the formation of a C_A- and C-type crystalline structure as well as porous structures on the granule surface. With differences in banana genome groupings contributing to the RS content in unripe banana flour, further research is needed on the contributions of these factors to enzymatic hydrolysis resistance in the different genome groups, both individually and interconnectedly.

AUTHOR CONTRIBUTIONS

Siphosethu R. Dibakoane: Writing—original draft; writing—review and editing. **Laura Suzanne Da Silva:** Writing—review and editing. **Belinda Meiring:** Writing—review and editing. **Tonna A. Anyasi:** Supervision; writing—review and editing. **Victor Mlambo:** Supervision; writing—review and editing. **Obiro Cuthbert Wokadala:** Conceptualization; supervision; funding acquisition; writing—review and editing.

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CONFLICT OF INTEREST STATEMENT

The authors declare no conflicts of interest.

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