

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

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Short title: Hydrolytic resistance of unripe banana flour

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 non-starch 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-relatability 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 on enhanced ways of extraction, processing and utilization of unripe banana flour and its starch.

Keywords: amylolytic resistance, banana starch, enzymatic hydrolysis, granule morphology, resistant starch, starch granules, unripe banana flour

1. INTRODUCTION

Resistant starch 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 resistant starch (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. Resistant starch (RS) consist of physically inaccessible starch classified as resistant starch 1 (RS1); ungelatinized starch 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 gastro-intestinal tract. Unripe banana flour native resistant starch 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 produce such as banana, exist in semicrystalline form and comprises both the amorphous regions consisting of amylose and branched amylopectin, as well as the long chains crystalline 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 influences amylolysis. The RS fraction is therefore starch that resist alpha-amylase and pullulanase hydrolysis *in vitro* beyond 120 min when compared to the 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 (SCFA) 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; Meenu & Xu, 2018). Produced SCFA are then utilised 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 for increasing colonic blood circulation, reducing

luminal pH and for preventing the formation of abnormal colonic cell populations (Lockyer & Nugent, 2017). Products of RS fermentation is 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 are grouped as prebiotics **due to their fermentation** by microbes in the colon **and the corresponding production of SCFA which is of benefit** to 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, 2018). 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, 2018). 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 fibres 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 hence provide protective effects against illnesses such as cardiovascular diseases, cancer, diabetes, and obesity (Singh et al., 2016; Sidhu & Zafar, 2018; Khoozani et al., 2019; Li et al., 2020). Resistant starch has been shown to reduce the incidence of non-communicable 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 fibre, contributes to faecal 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 in controlling shigellosis in children after five days of feeding, through a decrease in stool volume while increasing the concentrations of acetate, butyrate and propionate in the faeces 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 at elaborating 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 of the solid phase being the starch and the liquid phase being the conveyed enzyme that acts on the starch (Wang et al., 2022). Naguleswaran et al. (2014) opined that occurrence of amylolysis which is the hydrolysis of starch granules by amylases, takes place via diffusion to solid surface, adsorption and catalysis. Furthermore, amylopectin molecules that possess greater levels of short chains and degree of branching but with more compact structure, high molecular density, molar mass and smaller molecular size are less susceptible to amyolytic attack (Naguleswaran et al., 2014). Enzymatic hydrolysis of amylose and amylopectin molecules are 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 amyolytic attack, Ma et al. (2020), proposed that the presence of double helix order within or outside the crystal structure; stabilization of the double helices into perfect crystalline structures; presence of unevenly

packed or isolated single helices; 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 affects amylase hydrolysis. 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 enzymatic hydrolysis of starches. Dhital et al. (2015) stated that cracks, pores, interior channels and surface damage in starch granules further exacerbates 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 hinderance (implicated for 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 hindrance at the active centres of the enzyme (Singh et al., 2010). The large size of the starch granule compared to the alpha-amylase molecule provide 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 are 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 makes 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 surface accessibility of the granules through blocking of 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 centre with the number of these subsites in an active centre varying from four to nine. Wang et al. (2022) stated that different enzymes including alpha-, 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 catalysed by alpha-amylase through endocorrosion.

3. BANANA FLOUR RESISTANT STARCH 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 glucosyl units and function as entry points for enzymes (Nasrin & Anal, 2014a; Giacco et al., 2016). The average number of UBFRS glucosyl units for amylopectin side chains were 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 resistant starch 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 (Jiang et al., 2015; Bi et al., 2017; Yee et al., 2021; Das et al., 2022). 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 non-modified banana starch ranges from 30 – 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 resistant starch (RS) content (50.3%) compared to flour with 23.1% amylose content (45.1% resistant starch). 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) leads to higher resistance to hydrolysis due to 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 (Soares et al., 2011; Vatanasuchart et al., 2012; Bi et al., 2017). 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 plantain 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 resistant starch 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 possess a characteristically lower degree of branching, higher crystallinity, and longer amylopectin chains compared to the A-type crystalline structure (Kim et al., 2015; Chávez-Salazar et al., 2017; 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 UBFS consists 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 (Wang et al., 2017; Cordoba et al., 2018). Cordoba et al. (2018) showed that Terra Plátan (AAB) displayed B-type crystalline structure whilst Catura cavendish (AAA) and Prata Ana (AAB) cultivar displayed the C-type crystalline structure. Likewise, *Musa* AAA Cavendish and *Musa* ABB Bluggoe resistant starch (RS2) displayed a C-type crystalline structure whilst *Musa* ABB Pisang Awak and *Musa* AA Pisang Mas resistant starch (RS2) displayed a B-type crystalline structure (Wang et al., 2017). The relative resistance in C-type non-modified UBFS compared to A-type crystallinity starches results from the fact that the C-type crystalline structures are structurally intermediate between the A- 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 (González-Soto et al., 2007; Das et al., 2022). 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 results 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 an intact, irregular, and ellipsoidal shaped starch granules with smooth surfaces (Figure 1), whose structures are dependent on the banana cultivar (Thakorlal et al., 2010; Wang et al., 2017; Cordoba et al., 2018). Pelissari et al. (2012) demonstrated that RS2 present in unripe banana flour and starch granules displayed irregularly shaped and compact granules with both elongated and spheroid form. 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 irregular shaped structure whilst plantain had small and ellipsoidal granules. Anyasi et al. (2017) reported that *Luvhele* and *Mabonde* banana flour had polygonal and oval shaped granules, whilst *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 structure whilst 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 its 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 (González-Soto et al., 2007; Das et al., 2022). 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 provide a pathway for endocorrosion. Hence, autoclaved, and enzymatically (amylopullulanase) modified banana resistant starch (RS3) samples have increased susceptibility to enzymatic hydrolysis compared to native banana resistant starch (RS2) due to differences in granule structure.

4.2 Granule size

The diameter of unripe banana flour starch granules influence its susceptibility to enzymatic hydrolysis. Unripe banana flour starch granules have been reported to possess varied diameters ranging from 1.4 to 500 μm (Falade & Olugbuyi, 2010; Bezerra et al., 2013; Gao et al., 2016; Giraldo-Gómez et al., 2019; Khoozani et al., 2020; Thakaeng et al., 2021; Padhi & Dwivedi, 2022; Salazar et al., 2022). 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 (Jiang et al., 2015; Gao et al., 2016), whilst 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 molecular order of starch granules of 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-resistant starch (RS2) was greater than that of enzymatically modified UBFRS (RS3) (20 μm) (Das et al., 2022).

5. NON-STARCH COMPONENTS AND THE SUSCEPTIBILITY OF UNRIPE BANANA STARCH TO ENZYMATIC HYDROLYSIS

5.1 Native hydrocolloids (proteins and non-starch 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 the starch through prolonging the time required for effective hydrolysis (Zhang et al., 2005; Zhang & Hamaker, 2012).

The protein content of unripe banana flour are generally reported to be low <5% (Kumar et al., 2019; Rachman et al., 2021; Thakaeng et al., 2021) with higher concentrations of >5% reported to be present in the peels (de Angelis-Pereira et al., 2016; do Prado Ferreira & Teixeira Tarley, 2020) as well as 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 (Zhang et al., 2010; Shah et al., 2011). 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%) (Zhang et al., 2003; Campuzano et al., 2018), thus enabling to 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 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 (Milani & Maleki, 2012; Goff & Guo, 2019), and 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). Ng et al. (2020) suggested that the barrier formed by soluble fibre 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 (Haslinda et al., 2009; Dan, 2011; Menezes et al., 2011; Hoffmann Sardá et al., 2016; Kongolo et al., 2017). 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; Hoffmann 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 form a helical structure which reduces the binding activity of amylolytic enzymes and formation of enzyme-substrate complexes (Li, et al., 2021; Chi et al., 2022). However, the degree to which lipid complexation contributes to 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 et al., 2019a; Pico et al., 2019b). The polyphenols reported to be present in banana 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 (Bennett et al., 2010; Anyasi et al., 2018; Pico et al., 2019a; Pico et al., 2019b; Maseko et al., 2024). 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 (Liu et al., 2017; Gu et al., 2020). Furthermore, polyphenols can interact with starch molecules via covalent or non-covalent bonds to form starch-polyphenol complexes (Chi et al., 2022). The presence of starch-polyphenol complexes have been suggested to occur in unripe banana flour (Choo & Aziz, 2010; Sarawong et al., 2014; Pico et al., 2019b), 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

non-covalent 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; Kan et al., 2022; Ngo et al., 2022; Hernández 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 RESISTANT STARCH CONTENT OF UNRIPE BANANA FLOUR

The effects of the genomic, sub-genomic group, and cultivar type on the RS content of unripe banana flour are summarized in Table 2. Kumar et al. (2019) found that 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 group. 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 (Gibert et al., 2014; Annor et al., 2016; 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 group. This conforms with the findings of Reis et al. (2019), who suggested that 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% (Menezes et al., 2011; Rayo et al., 2015; Campuzano et al., 2018; Khoozani et al., 2019; Rosado et al., 2020; Tian et al., 2020) had RS content that overlapped with that of the ABB cultivar types with RS content of 8.60 – 68.8% (Rodríguez-Ambriz et al., 2008; Haslinda et al., 2009; Vatanasuchart et al., 2012; Nasrin et al., 2015). 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 RESISTANT STARCH CONTENT OF UNRIPE BANANA FLOUR

The effect of physical and chemical modifications on the resistant starch 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 the 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 to 25 min (23.3 - 23.7%). Given that the RS content of unripe banana flour tends to decrease with cooking time (Rodríguez-Damian et al., 2013; De la Rosa-Millan et al., 2014), 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. 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 (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 the RS content 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. Heat moisture treatment minimizes starch granule damage while storage facilitates retrogradation of the starch granules by enabling 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 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 (23 – 78%). The highest RS value of phosphated flour was attributed to the cross-linking 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 were 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 non-modified plantain flour (40.69%) (Almanza-Benitez et al., 2015). Increase in the RS level of unripe plantain flour treated with hydrochloric acid was attributed to hydrolysis of the amorphous zones of starch and 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. Several studies have shown that the gelatinization temperature of debranched and enzymatically modified banana flour resistant starch (103 – 106 °C) was greater than the unmodified one (69.1 - 72.5 °C) (Cordoba et al., 2018; Yee et al., 2021; Das et al., 2022). Similarly, the enthalpy of gelatinization (4.4 - 188.2 J/g) and peak temperature (68.5 – 147 °C) of the debranched, enzymatically modified and litherized banana RS were greater than the unmodified UBFRS (6.1 - 13.9 J/g and 37.6 - 72.5 °C) respectively (Lehmann et al., 2002; Aparicio-Saguilán et al., 2005; Aparicio-Saguilán et al., 2008; Nasrin & Anal, 2014b; Cordoba et al., 2018; Yee et al., 2021; Das et al., 2022). 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 resistant starch content.

8. CONCLUSION

The resistance to enzymatic hydrolysis of UBFRS can be attributed to a multi-phenomenon mechanism which 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 as well as the presence of other native non-starch 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 DP_n, degree of amylopectin branching, and amylose-amylopectin ratio was 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 a 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 Dibakoane: Literature review; writing – review & editing. **Laura Suzanne Da Silva:** Writing – review & editing. **Belinda Meiring:** Writing – review & editing. **Tonna A. Anyasi:** Supervision; writing – review & editing. **Victor Mlambo:** Supervision; writing – review & editing. **Obiro Cuthbert Wokadala:** Conceptualization; supervision; writing – review & editing.

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CONFLICT OF INTEREST STATEMENT

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.

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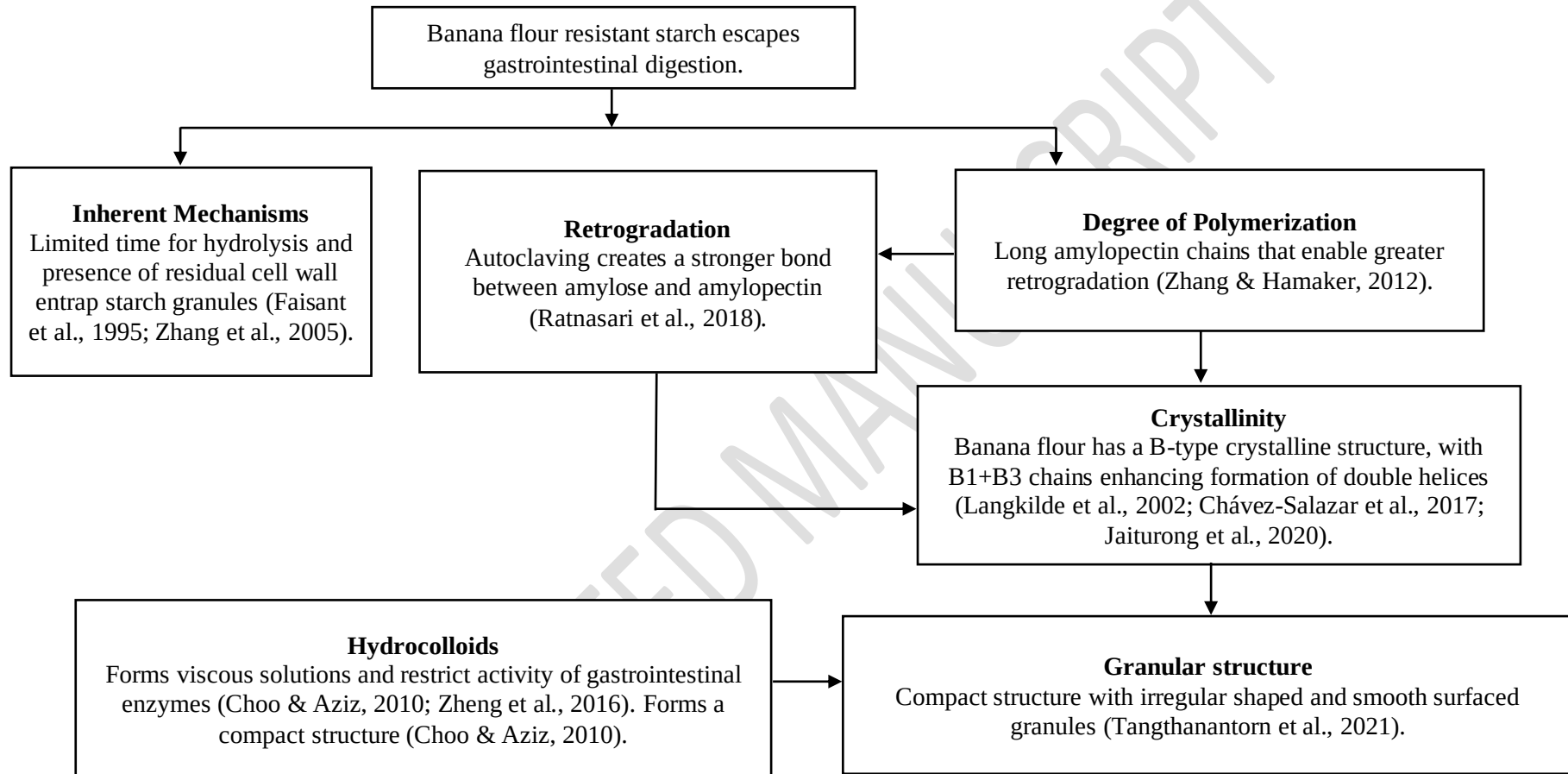


FIGURE 1 Mechanisms of resistance of unripe banana flour starch to enzymatic hydrolysis

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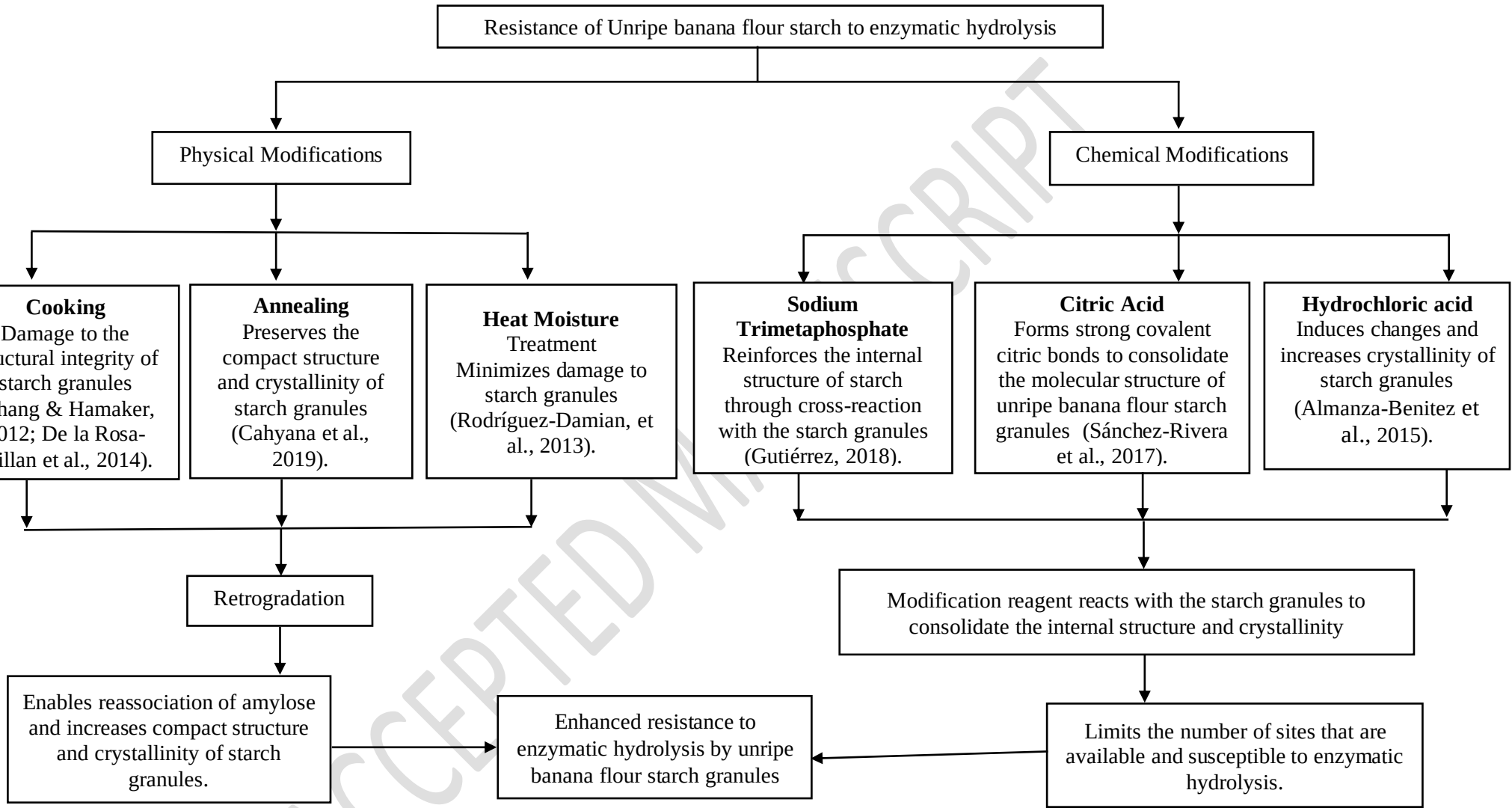


FIGURE 2 Effects of physical and chemical modifications and their associated mechanisms on the resistance of banana flour starch granules to enzymatic hydrolysis

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1160 **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)	21.76 - 27.75	B-type	29.07 - 39.17 μ m	Ellipsoidal and spherical	14.5 glucosyl units	(Cet al., 1995)
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)		B- and C-type		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 μ m	Elliptic, triangular shaped with smooth surfaces		(Wang et al., 2017)
RS2	Raw (uncooked)					18.9 - 20.3 glucosyl units	(Yee et al., 2021)

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1162 **TABLE 2** Effect of genomic and sub-genomic group classification on unripe banana flour resistant starch content

Genomic Group	Sub-genomic group	Cultivar	Resistant starch content (%)	References
AA	Inarnibal	Pisang Lemark	47.30	(Kongolo et al., 2017)
AA	Sucier	Lady Finger	39.60	
AAA	Cavendish	Gross Michel	37.90	
AAA	Cavendish	IBP. 5.61	47.10	
AAA	Sucier	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	(Kumar et al., 2019)
AAAB	Goldfinger selection	PKZ	47.40	
AAA	Cavendish	Grand Naine	32.70	
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	(Bi et al., 2017)
AAA	Cavendish	Bukoba	90.69	
	Dwarf red banana	<i>Guihongjiao No. 1</i>	86.15	
ABB	Pisang Awak	<i>Jinfen No. 1</i>	85.88	
AAA	Cavendish	<i>Williams B₆</i>	90.46	
AAA	Cavendish		57.48 (standard UBF)	
AAA	Cavendish		43.25 (standard peel UBF)	
				(Hoffmann Sardá et al., 2016)

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)
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, 2015b)
	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> <i>Colla</i>		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 L.</i>		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	
		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		1163
Plantain	<i>Red Essong</i> (French bunch)	33.30 – 50.23	(Udomkun et al., 2021)	1164
Plantain	<i>Mbouroukou 3 - False Horn bunch</i>	35.36 – 53.11		1165
Plantain hybrids	<i>PITA 14</i>	32.72 – 39.67		1166
Plantain hybrids	<i>PITA 27</i>	30.91 – 49.89		1167
<i>Musa sapientum</i> , triploid hybrid banana		7.90 – 8.90	(Chaipai et al., 2018)	1168
<i>Musa paradisiaca</i>		24.60	(Hernández-Nava et al., 2011)	1169
	<i>Kluai Namwa plantain</i>	8.60	(Nasrin et al., 2015)	1170
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1183 **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, 2015a)
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)
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 ⁻¹ 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 ⁻¹ citric acid for 10 min	39.44 – 56.37	
Native (PITA 14)			32.72 – 39.67	
Chemical	Citric acid	0 g L ⁻¹ citric acid for 10 min	36.28 – 53.63	
Native (PITA 27)			30.91 – 49.89	

Chemical	Citric acid	0 g L ⁻¹ 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	

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