



# Effects of heat moisture treatment on the physicochemical properties of starch nanoparticles



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## ABSTRACT

In this study, the effect of heat moisture treatment (HMT) on the properties of waxy maize starch nanoparticles (SNPs) was investigated. The SNPs were adjusted to 20% and 30% moisture levels and heated at 90 °C and 110 °C for 4 h. Transmission electron microscopy, X-ray diffractometry, differential scanning calorimetry, and Fourier transform infrared spectroscopy were used to characterize the morphology and crystal structure of the SNPs after HMT. The research found that the morphology of SNPs did not change significantly, keeping nanoscale size. When the SNPs were subjected to HMT at 110 °C and 30% moisture content, the crystalline structures changed from B-type to A-type, and the crystallinity of the SNPs increased significantly. HMT significantly increased the onset temperature, peak temperature, final temperature, and enthalpy value of the SNPs. As the HMT temperature and SNP moisture content increased, the hydrogen bonds between the starch molecular chains in the SNPs became stronger.

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## 1. Introduction

Starch nanoparticles (SNPs), a novel product derived from starch, are nano-sized in at least one dimension. In recent years, many publications have reported the preparation, structure analysis, and functional performance of SNPs (Chin, Pang, & Tay, 2011; Ma, Jian, Chang, & Yu, 2008; Santander-Ortega et al., 2010; Singh, Dartois, & Kaur, 2010; Singh, Singh, Pandey, & Sanghi, 2010). SNPs are usually obtained by one of three ways: physical methods, acid hydrolysis, and enzymatic hydrolysis. Physical methods such as precipitation (Ma et al., 2008) and microfluidization (Liu, Wu, Chen, & Chang, 2009) can produce SNPs 10–20 nm in size under a pressure of 207 MPa. With hydrolysis using H<sub>2</sub>SO<sub>4</sub>, the amorphous domains of semicrystalline starch granules are disrupted, leaving platelet-like starch nanocrystals (LeCorre, Bras, & Dufresne, 2011). The obtained starch nanocrystals have a lamellar morphology with a thickness of 4–8 nm and a diameter of 50–120 nm. Using selective enzymatic hydrolysis, nanoscale starch particles with an average diameter of 500 nm have been prepared from waxy rice starch (Kim, Park, & Lim, 2008). A new method using enzymolysis and retrogradation has also been used to prepare SNPs. Waxy maize starch with high amylopectin content (98% amylopectin) was usually debranched with pullulanase and the ensuing branched

short linear glucans could form nano-scale starch particles during the retrogradation process. This method has the advantage of being quite rapid and presenting a higher yield. The particle sizes of SNPs prepared from proso millet starch and waxy maize starch are 20–100 nm and 60–120 nm, respectively (Sun, Gong, Li, & Xiong, 2014; Sun, Li, Dai, Ji, & Xiong, 2014). In recent years, SNPs have been widely used in food packaging (Chang, Jian, Yu, & Ma, 2010; Rhim, Park, & Ha, 2013), drug delivery (Simi & Emilia, 2007; Wim & Paul, 2008), and paper fabrication, due to their high mechanical properties and renewable nature (Simi & Emilia, 2007; Yoon & Deng, 2006). However, the SNPs do not always have the physical or chemical properties appropriate for certain types of processing; thus, some modifications are commonly used to produce SNPs with special properties. There are three strategies for modifying SNPs – chemical, enzymatic, and physical methods – which could promote specific functional properties. Angellier, Molina-Boisseau, Belgacem, and Dufresne (2005) chemically modified the surfaces of starch nanocrystals using alkenyl succinic anhydride and phenylisocyanate. Chakraborty, Sahoo, Teraoka, Miller, and Gross, (2005) investigated the chemical modification of starch-derived nanocrystals with ethylene glycol methyl ether and stearic acid chloride. Namazi and Dadkhah (2008) also observed the surface modification of starch nanocrystals through the ring-opening polymerization of  $\epsilon$ -caprolactone. Heat moisture treatment (HMT) is one of the important physical methods used to modify starch using simple and environmentally safe processes at low cost and without chemical reagent byproducts

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(Adebowale, Afolabi, & Olu-Owolabi, 2005; Sun, Wang, Xiong, & Zhao, 2013). Consumers tend to be more receptive to such physical modifications, as they avoid traditional chemical means to modify starch. To the best of our knowledge, there are no reports regarding the modification of SNPs by HMT. Therefore, in this study, SNPs were treated with moisture content (20% and 30%) at 90 °C and 110 °C for four hours, and the effect of this HMT on the physico-chemical and morphological properties of the SNPs were studied by using transmission electron microscopy (TEM), X-ray diffractometry (XRD), differential scanning calorimetry (DSC), and Fourier transform infrared spectroscopy (FTIR). The main objective was to study the influence of HMT on the property and structure of the SNPs.

## 2. Materials and methods

### 2.1. Materials

Waxy maize starch (98% amylopectin) was obtained from National Starch & Chemical Co., Ltd. (Guangdong, China). Pullulanase (E.C.3.2.1.41, 6000 ASPU/g, 1.15 g/mL), where ASPU is defined as the amount of enzyme that liberates 1.0 mg glucose from starch in 1 min at pH 4.4 and 60 °C) was supplied by Novozymes Investment Co. Ltd. (Beijing, China). All reagents used were analytical grade.

### 2.2. Preparation of SNPs

The SNPs were prepared using the method described by Sun, Li, et al. (2014). Waxy maize starch (15 g, db) was dispersed in 100 mL of disodium hydrogen phosphate and citric acid buffer solution (pH 5.0), and the starch slurry was cooked in boiling water, with vigorous stirring, for 30 min to fully gelatinize the starch. The temperature of the cooked waxy maize starch was adjusted to 58 °C and pullulanase (30 ASPU/g of dry starch) was added. After an 8 h incubation period at 58 °C, the reaction was stopped by heating the starch at 100 °C for 30 min to inactivate the pullulanase, followed by cooling to room temperature. Then, the solutions were stored at 4 °C for 8 h. The suspensions were washed several times with distilled water until neutrality and then freeze dried to obtain SNPs.

### 2.3. Heat moisture treatment (HMT)

HMT of the SNPs was performed according to the method described by Hormdok and Noomhorm (2007), with some modifications. The moisture content of the SNPs was adjusted to 20% and 30%, after which the samples were placed in sealed glass tubes and heated at 90 °C and 110 °C for 4.0 h.

### 2.4. Transmission electron microscopy (TEM)

Transmission electron micrographs of the SNP samples were taken with a Hitachi 7650 (Tokyo, Japan) transmission electron microscope with an acceleration voltage of 80 kV. The samples were deposited on a carbon-coated grid and freeze dried.

### 2.5. X-ray diffraction (XRD)

The crystalline structures of the SNP samples were studied using an X-ray diffractometer (AXS D8 ADVANCE; Bruker, Karlsruhe, Germany). The samples were stored in a sealed container in a saturated solution (1000 mL) of NaCl to standardize moisture content, with Cu K $\alpha$  radiation ( $\lambda = 1.543$ ). Signals of the reflection angle of  $2\theta$  from 4° to 40° were recorded. Samples crystallinity was determined by plotting the peaks baseline on the diffractogram and calculating

the area using the software spectrum viewer (Version 2.6) according to the method described by Jivan, Madadlou, and Yarmand (2013). The area above and under the curve corresponded to crystalline domains and amorphous regions, respectively. The ratio of upper area to total area was taken as the crystallinity degree:

Crystallinity percentage

$$= \text{Area under the peaks} / \text{Total curve area} \times 100 \quad (1)$$

### 2.6. Differential scanning calorimetry (DSC)

Gelatinization parameters of the SNP samples were measured using a differential scanning calorimeter (DSC1; Mettler Toledo, Schwerzenbach, Switzerland) equipped with a thermal analysis data station and data recording software (STAR@ SW 9.20), as described by Chanvrier et al. (2007), with some modifications. Using a microsyringe, deionized water (20  $\mu$ l) was added to the SNPs (10.0 mg db) in an aluminum pan. The pan was hermetically sealed before DSC analysis. The scanning temperature range and the heating rates were 25–120 °C and 10 °C/min, respectively. In all measurements, the thermogram was recorded with an empty aluminum pan as a reference. During the scans, the space surrounding the sample chamber was flushed with dry nitrogen to avoid condensation. The transition temperatures reported are the onset ( $T_o$ ), peak ( $T_p$ ), and conclusion ( $T_c$ ). The enthalpy of gelatinization ( $\Delta H$ ), estimated by integrating the area between the thermogram and a baseline under the peak, was expressed in Joules per gram of dry starch; three replicates per sample were analyzed.

### 2.7. Fourier transform infrared spectroscopy analysis (FTIR)

The infrared spectra of the SNP samples were recorded on an FTIR spectrophotometer (NEXUS-870; Thermo Nicolet Corporation, Madison, WI, USA) as described by Kunal, Banthia, and Majumdar (2008). All of the samples were mixed with KBr and pressed into pellets, which were then subjected to attenuated total reflectance spectroscopy in a 4000–400  $\text{cm}^{-1}$  range. Intensity measurements were performed on the spectra by recording the height of the absorbance bands from the baseline.

### 2.8. Statistical analysis

Each measurement was carried out using at least three fresh, independently prepared samples. The results were reported as the mean value and standard deviation. The data were subjected to analysis of variance (ANOVA) using the SPSS V.12 statistical software package (SPSS Inc., Chicago, IL). Duncan's multiple range test was also used to determine the difference of the means from the ANOVA, using a significance test level of 5% ( $p < 0.05$ ).

## 3. Results and discussion

### 3.1. Morphology of SNPs

The morphologies of unmodified and HMT SNPs under different heat treatment conditions were examined by TEM (Fig. 1). As shown in Fig. 1, the shapes of the SNPs before and after modification were nearly spherical or elliptical and the particle sizes of the SNPs were 50–100 nm wide and 80–120 nm long. It appeared that the HMT did not change the sizes or shapes of the SNPs to any obvious extent. Watcharatwinkul, Puttanlek, Rungsardthong, and Uttapap (2009) reported that HMT did not alter the shape or size of canna starch, and similar observations were reported for heat-moisture-treated finger millet (Adebowale et al., 2005), rice (Khunae, Tran,

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