



Development of a single droplet freezing apparatus for studying crystallisation in cocoa butter droplets



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ARTICLE INFO

Article history:

Received 14 November 2014

Received in revised form 5 February 2015

Accepted 8 February 2015

Available online 14 February 2015

Keywords:

Cocoa butter
Polymorphism
Crystallisation
Simulation
Heat transfer

ABSTRACT

The single droplet freezing apparatus described by Pore et al. (2009), which allows crystallisation to be monitored *in situ* by X-ray diffraction, was modified to allow rapid switching of coolant gas and monitoring by video microscopy. The apparatus was used to study drops of cocoa butter undergoing simulated spray freezing at high cooling rates, e.g. 130 K/min. The transformation of an Ivory Coast cocoa butter to the Form V polymorph was significantly faster in drops (~40 h) than in static bulk samples (10 days) crystallised under isothermal conditions. Phase transformation was observed from Forms I/II → III → IV → melt → V, with Form V crystallising directly from the melt at 28.6 °C. Numerical simulations of the temperature evolution within the droplet established that the drops are not isothermal, explaining why nucleation was initially observed in the lower (upstream) part of the droplet.

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1. Introduction

Cocoa butter is a naturally occurring, edible fat produced from the cacao bean, and is one of the major ingredients of chocolate. It consists of a mixture of triacylglycerides (TAGs) and exhibits polymorphic phase behaviour, existing in six distinct crystalline forms. These are labelled Forms I to VI following the Wille & Lutton nomenclature (Wille and Lutton, 1966) or γ , α , β' and β using the Larsson nomenclature (Larsson, 1966). Form I is the least thermodynamically stable polymorph and Form VI is the most stable. The different polymorphs can be identified via their sub-cell structure using X-ray diffraction (XRD) Clarkson et al., 1934. They can also be distinguished using differential scanning calorimetry (DSC). XRD offers unambiguous identification of the crystals present and is thus the preferred method of characterisation. Much of the chocolate literature employs the Roman classification, originally based on melting point data, and for this reason it is used here.

Real time XRD studies using laboratory X-ray diffractometers have been carried out to observe the evolving crystalline structure in cocoa butter in both static (Van Malsen et al., 1996, 1999, 1996a,b) and sheared crystallisation (Sonwai, 2002). Synchrotron XRD offers improved resolution, rapid data collection and simultaneous small (SAXS) and wide (WAXS) angle measurements, corresponding to the long and short range spacing in the crystal lattice

(Macmillan et al., 2002; Macmillan and Roberts, 2002; Van Langevelde et al., 2001). In most of these studies, the cocoa butter sample is held in temperature controlled sample holder (Van Malsen et al., 1996b), a thin capillary tube (Lopez and Ollivon, 2009), or a bespoke cell (Ueno et al., 1999). The cocoa butter samples are usually heated to 50–70 °C for 15–30 min to avoid memory effects (Van Langevelde et al., 2001), and then cooled, at a rate similar to that used in DSC (e.g. 0.5–20 K/min), or crash cooled at 50 K/min, to the required temperature. Droplet samples of cocoa butter have not previously been studied by XRD until the development of the single droplet freezing apparatus reported by Pore et al. (2009).

The use of a droplet suspended in a flowing medium is common in heat and mass transfer studies (Fuchs, 1959). The single droplet freezing apparatus (SDFA) was developed by (Hindmarsh et al., 2007) to study temperature transitions in freezing droplets of aqueous food solutions. The SDFA allows the thermal history of a droplet to be monitored by suspending it on a fine thermocouple and subjecting it to a cold, dry air stream. This apparatus was originally used with video imaging to study the freezing behaviour of dairy solutions. SDFA devices have also been constructed for use in MRI machines for mapping emulsion creaming and solidification within droplets (Hindmarsh et al., 2004).

Gwie et al. (2006) used an SDFA to study spray-freezing of tripalmitin and cocoa butter. Their SDFA was later modified to fit into a laboratory X-ray system by Pore et al. (2009), allowing *in-situ* real time monitoring of the freezing process. It was possible to observe the transformation from Form I to the higher melting polymorphs,

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