

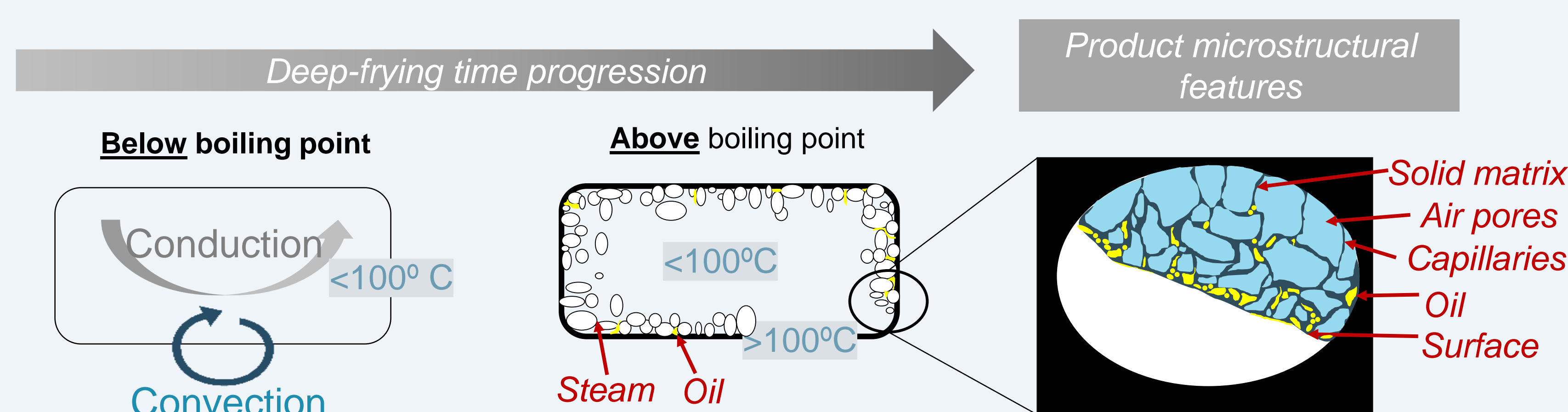
The contribution of wheat starch and wheat gluten to structure formation during deep-fat frying:

An objective tool to tailor fried food matrices of higher quality and nutrition?

Isabella M. Riley, Ujjwal Verma, Nand Ooms, Mieke A. Nivelde, Pieter Verboven, Bart Nicolai, Jan A. Delcour

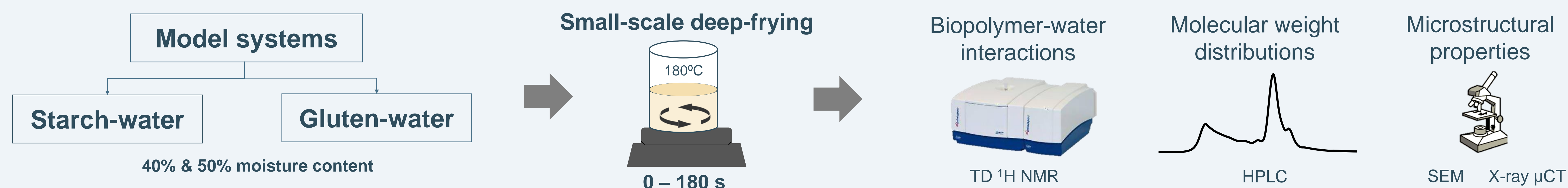
Research context

- Deep-fat frying is a popular cooking method across the world, commonly applied to various wheat-based food materials.
- This complex process involves heat exchange and the simultaneous transfer of oil and water. These contribute to unique textural and sensorial properties.
- Starch and gluten, the main constituents of wheat flour, are expected to contribute to deep-frying-induced structure formation.



Research question: What contribution do starch and gluten constituents have to the formation of a porous matrix and water loss/oil uptake dynamics of deep-fried wheat-based foods?

Materials and methods



Results

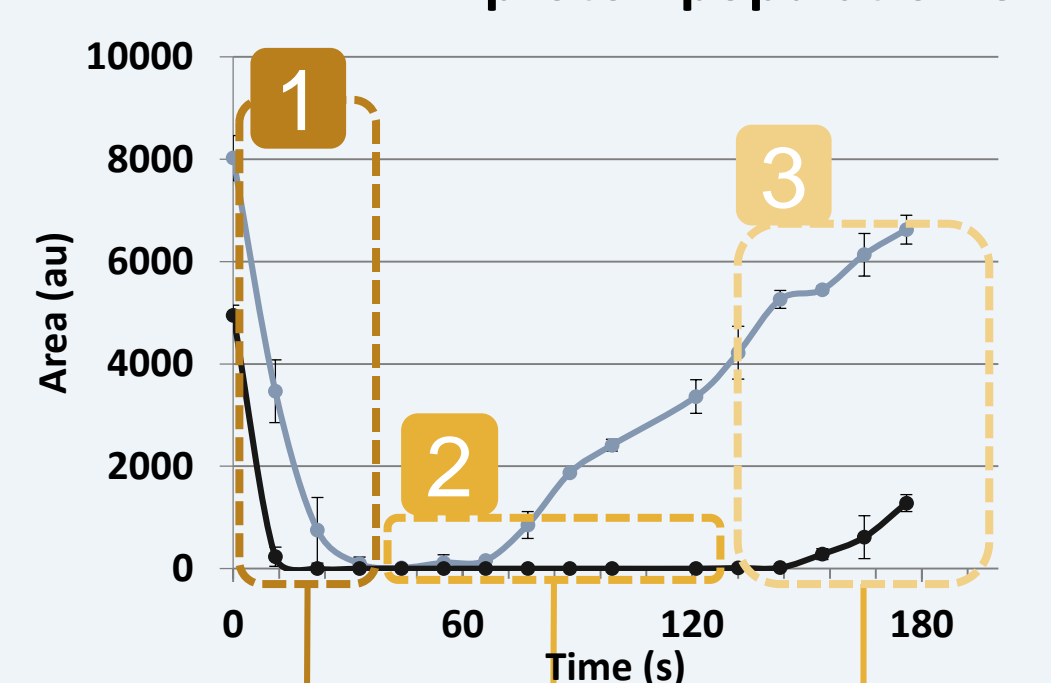
WHEAT STARCH

Changes in the physical state of wheat starch (WS)

Temperature-controlled TD ¹H NMR may be used to evaluate proton dynamics [free induction decay (FID) and Carr-Purcell-Meiboom-Gill (CPMG)] during simulated deep-frying conditions at ca. 125° C.

WS 40% Hydration (blue line) WS 50% Hydration (black line)

FID proton population evolution



FID: Representing rigid, fast-relaxing protons (e.g. crystalline amylopectin, amorphous starch in the glassy state).

1 Loss of crystalline amylopectin due to starch gelatinization.

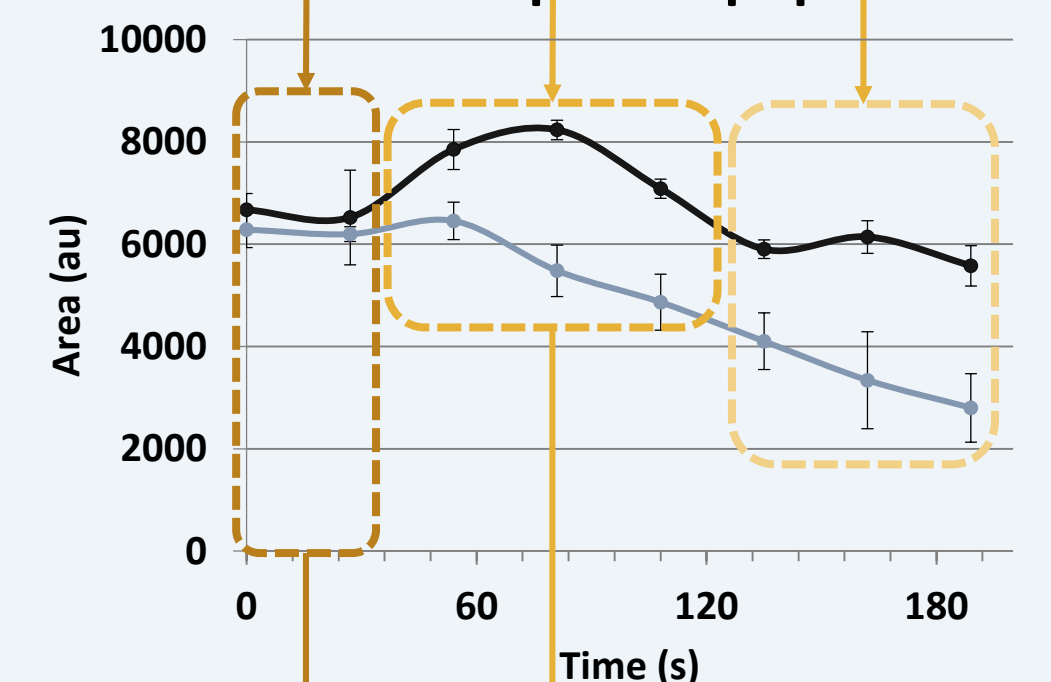
- Starch granules cluster together thereby forming pores for oil to enter, however, starch granular integrity is maintained.

2 The increased interaction of previously crystalline starch chains with water.

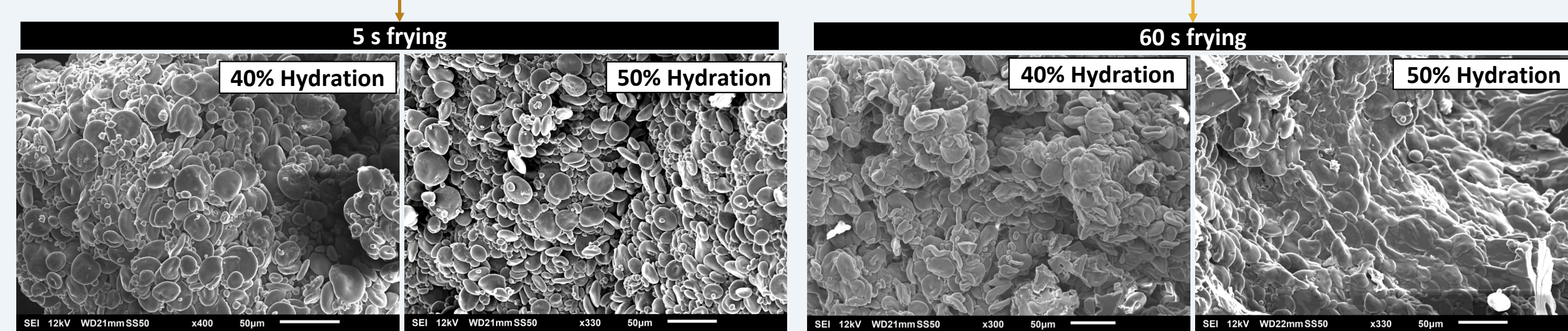
- Starch granules lose integrity, causing the microstructure to be predominantly characterized by a merged mass of starch granules.

3 Progressive increase of starch chains in the glassy state.

CPMG proton population evolution



CPMG: Representing slowly-relaxing protons such as those belonging to water domains (e.g. water interacting strongly or weakly with starch)



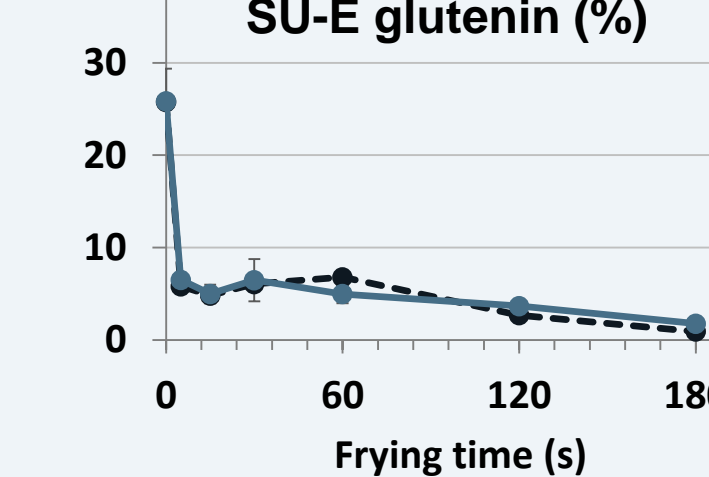
WHEAT GLUTEN

1 Non-covalent interactions

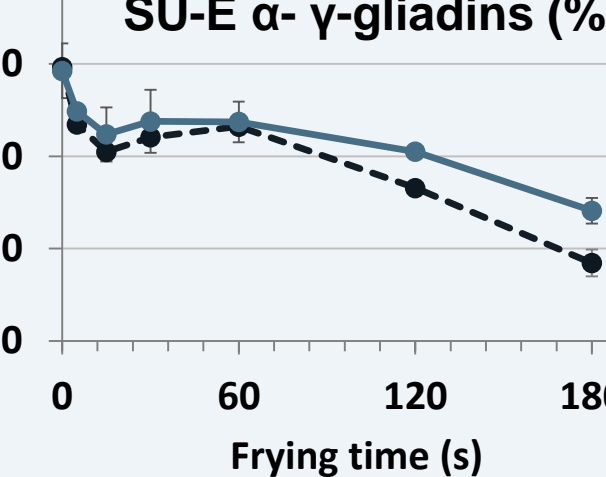
Extractability of fried gluten (FG) protein in media containing urea (2.0M) and 2% (w/v) SDS – (SU-E)

FG 40% Hydration (black line) FG 50% Hydration (blue line)

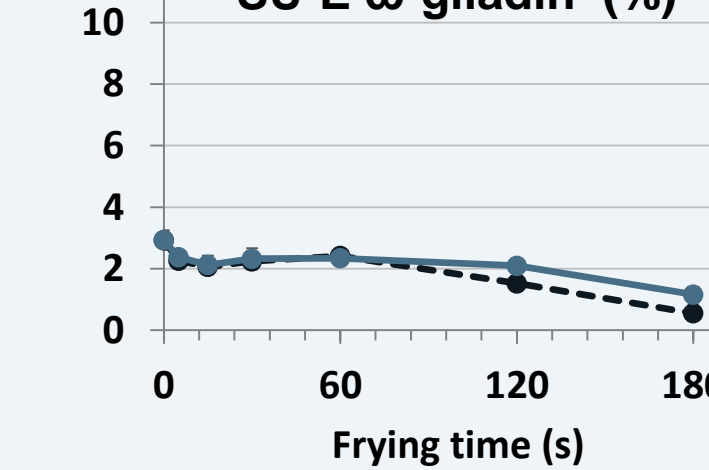
SU-E gluten (%)



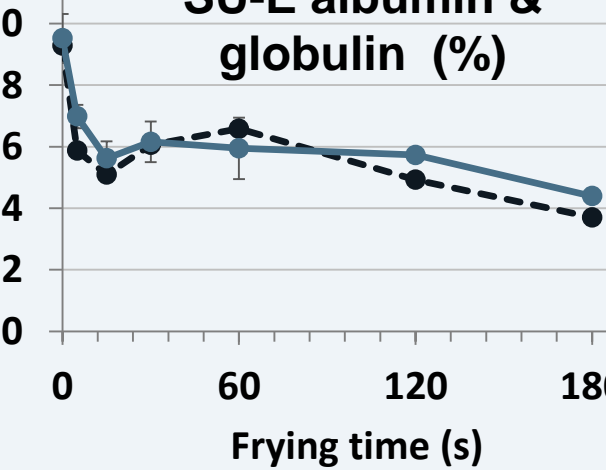
SU-E α-γ-gliadins (%)



SU-E ω-gliadin (%)

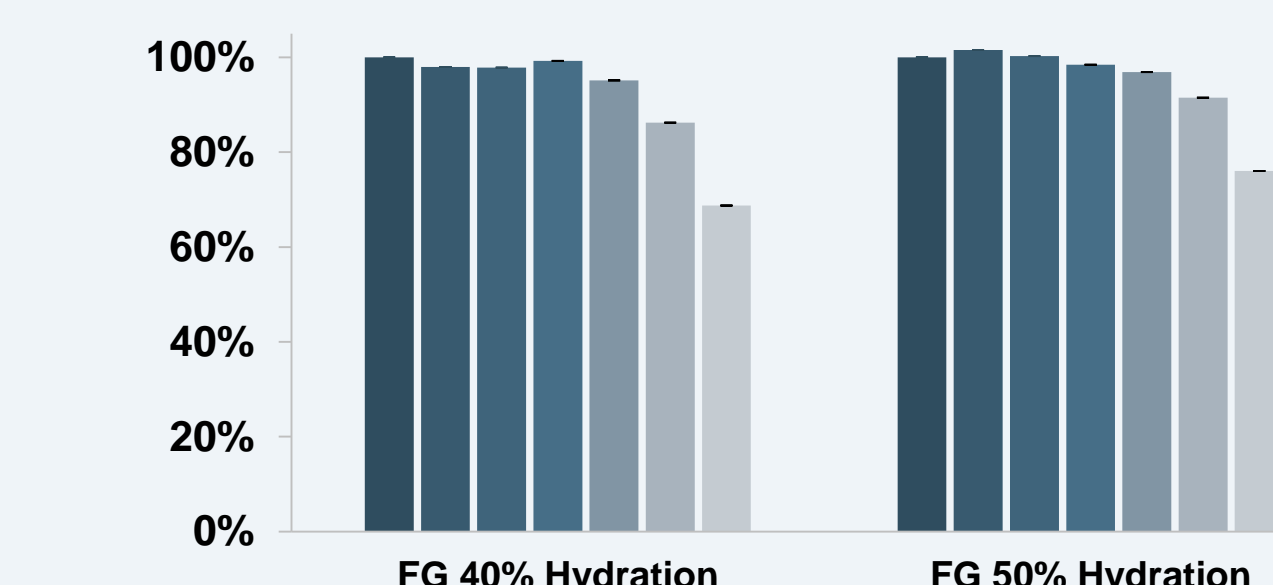


SU-E albumin & globulin (%)



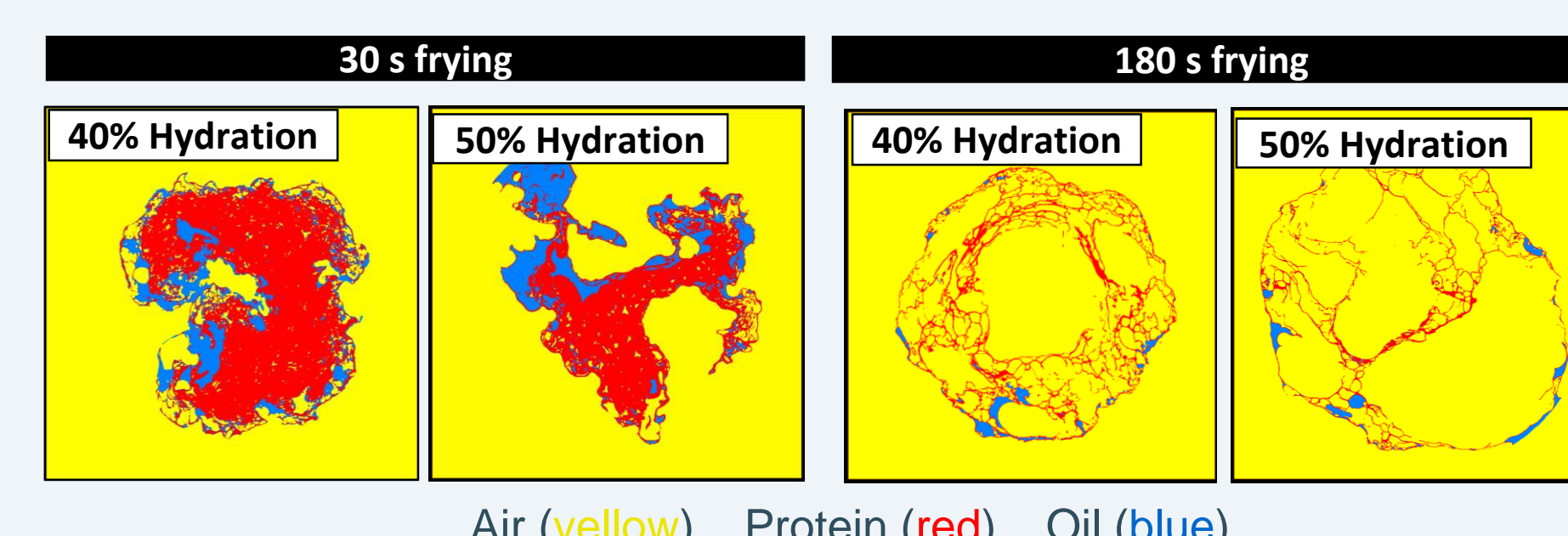
0 s 5 s 15 s 30 s 60 s 120 s 180 s

SUD-E (%)



- Non-covalent interactions contribute to the gluten network based on a decrease in SU-E.
- SU-E α-γ-gliadins are incorporated into the gluten network with longer deep-frying (60-180 s).
- Covalent reactions are dominated by disulfide bonds. However, non-disulfide bonds contribute to the gluten network (decrease in SUD-E) after 60 s of frying.
- Substantial expansion occurs later in the frying process.

3 Microstructural properties



Air (yellow) Protein (red) Oil (blue)

Conclusions

1
Wheat starch immediately gelatinizes upon deep-frying and the gluten network formed is dominated by non-covalent interactions.

2
Wheat starch (and presumably wheat gluten) progressively enter(s) into the glassy state during deep-frying depending on the initial moisture present in the system.

3
Non-disulfide cross-links contribute to the gluten network with longer deep-frying times.