

DETAILED DESCRIPTION OF THE *IN SITU* KINETICS AND SYNCHRONISM OF FEED FRACTIONS DEGRADATION

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Objectives

The valorisation of feeds by ruminants is largely dependent on their microbial degradation in the rumen and the partition between degradable and undegradable fractions. Nevertheless, the *in situ* degradation is rarely studied for all the components of dry matter (DM) and often dealt with only one or few of them.

The aims of this study were to **achieve a detailed description of the kinetics of degradation** of the main constituents of feeds and to **explain the observed variations between feeds and constituents** and to **elucidate the synchronism of their availability for ruminal micro-organisms**

Methods

➤ **10 feeds representative of the major types of kinetics of degradation** (Chapoutot, 1998):

barley (BAR)	maize (MAI)	pea (PEA)	dehydrated lucerne (DLU)	dried brewer's grains (BRG)	corn gluten meal (CGM)
dehydrated sugar beet pulp (SBP)	soyabean meal (SBM)	formaldehyde-treated soyabean meal (TSBM)			palmkernel meal (PKM)

➤ **Chemical composition of feeds:** crude protein (CP) and soluble N (Nsol), Cellwall components (NDF, ADF and ADL)

➤ ***In situ* degradation kinetics of DM, CP, structural carbohydrates (SC=NDF) and non-protein cytoplasmic constituents (npCC=DM-CP-NDF)** measured by Nylon bags methods (Michalet-Doreau *et al.*, 1987)

➤ **Cumulative synchronism index (CSI)** calculated over 9 consecutive time intervals from 0 to 72h:

[instantaneously (t=0), 0h-4h, 4h-8h, 8h-12h, 12h-16h, 16h-24h, 24h-36h, 36h-48h and 48h-72h]

$$CSI = \sum_j (Ndeg_j - 0.032 \times npDMdeg_j) \quad (j \text{ varying from 1 to 9})$$

• Ndeg_j and npDMdeg_j: quantity of nitrogen and total carbohydrates (npDM=100-CP) available for micro-organisms along the different time intervals (j) with a relative rate of particles outflow from the rumen of 0.06h⁻¹

• optimal requirement for micro-organisms: 32g N/kg of degraded carbohydrates (Sinclair *et al.*, 1991 and 1993)

Results and Discussion

➤ **Large variety of degradation (deg) patterns between different feeds and components** (figure 1)

CP and DM deg kinetics: fairly parallel, except for SBP and MAI (=very low CP deg during 10h or 20h), and BRG (=high proportion of undegradable CP
Cellwall deg: quite long lag phase (almost 16h) for TSBM and PKM

➤ **Variations of DM deg explained by different components** according to the incubation times: Nsol at 1h (R=0.94), CC at 12h (R=83) and SC at 72h (R=86)

➤ **Undegraded SC at 72h (g/kg DM) related to the ADL content of feeds (%DM):**

$$QSC72h = 40.8 ADL - 1.87 ADL^2$$

(n=10 ; R=0.98 ; RSD=20.9)

➤ **Large differences of synchronicity** between N and carbohydrate availability for microbes (figure 2):

- fairly **harmonious**: DLU and PKM
- **deficit of nitrogen**: BAR, SBP and MAI (-6, -10 and -12g N/kg feed)
- **excess of N**: BRG, TSBM and PEA (+7, +12 and +15g N/kg) and above all CGM (+35g N/kg) and SBM (+45g N/kg)

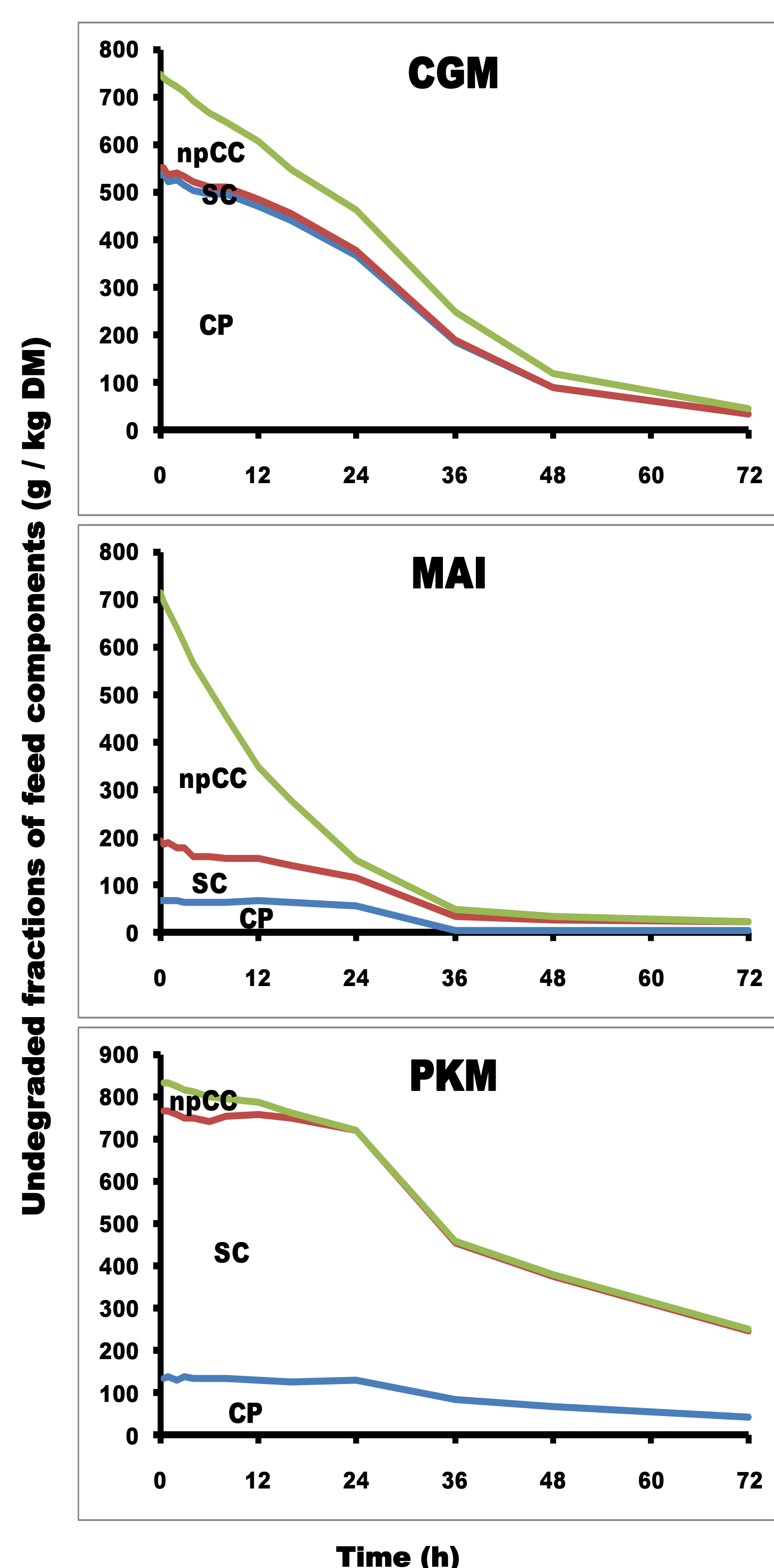


Figure 1: Examples of different degradation patterns

Conclusion

This study confirmed the **great variability of ruminal degradation patterns** between feeds and components which was **partially explained by chemical composition**.

The **feeds differed by their ability to provide, with synchronicity or not, nitrogen and carbohydrates for micro-organisms**. This could largely influence microbial synthesis efficiency.

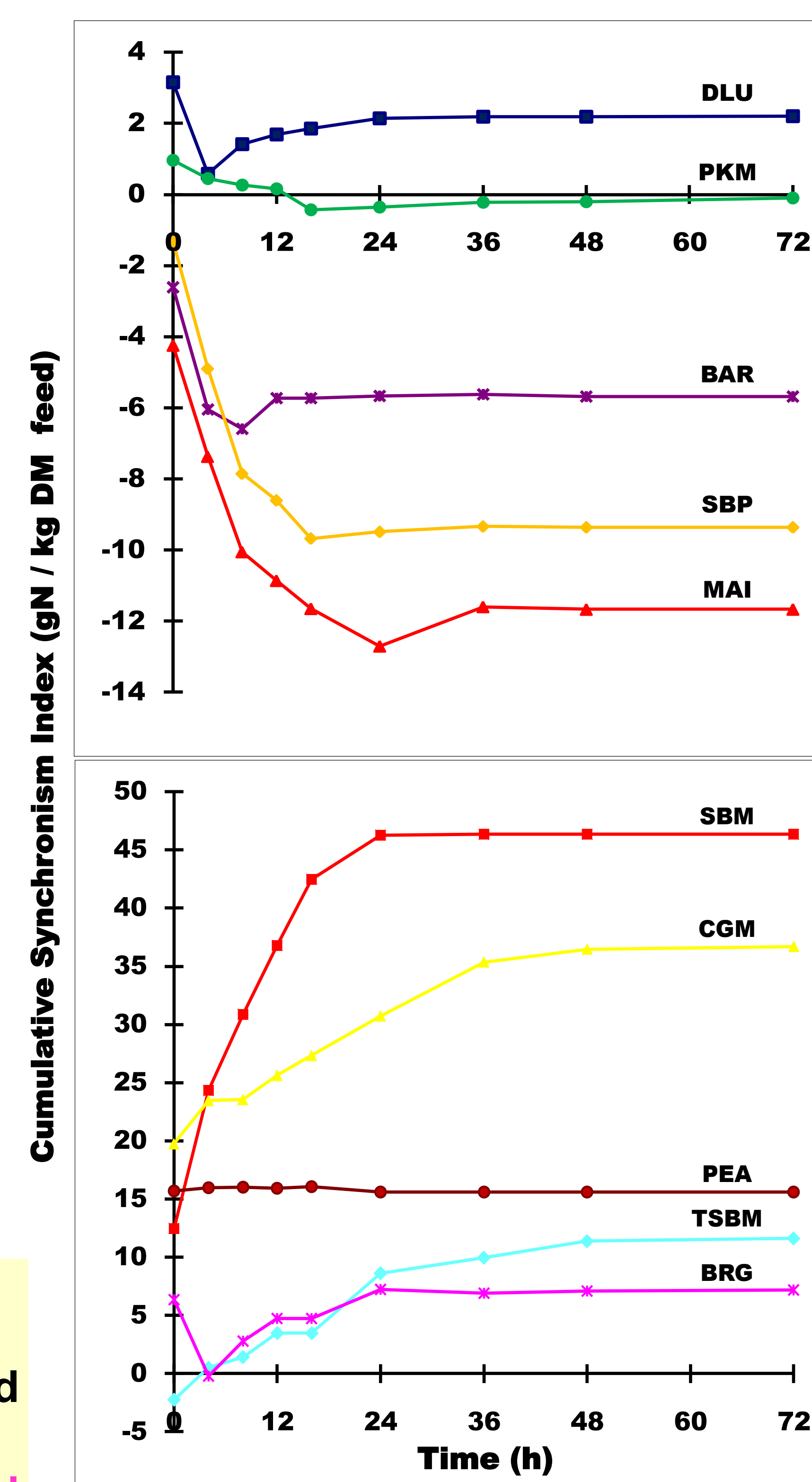


Figure 2: Evolution of the cumulative synchronism Index of feeds