



Unraveling the molecular mechanism of *in vitro* gastric lipolysis through the advanced multi-response modeling technique

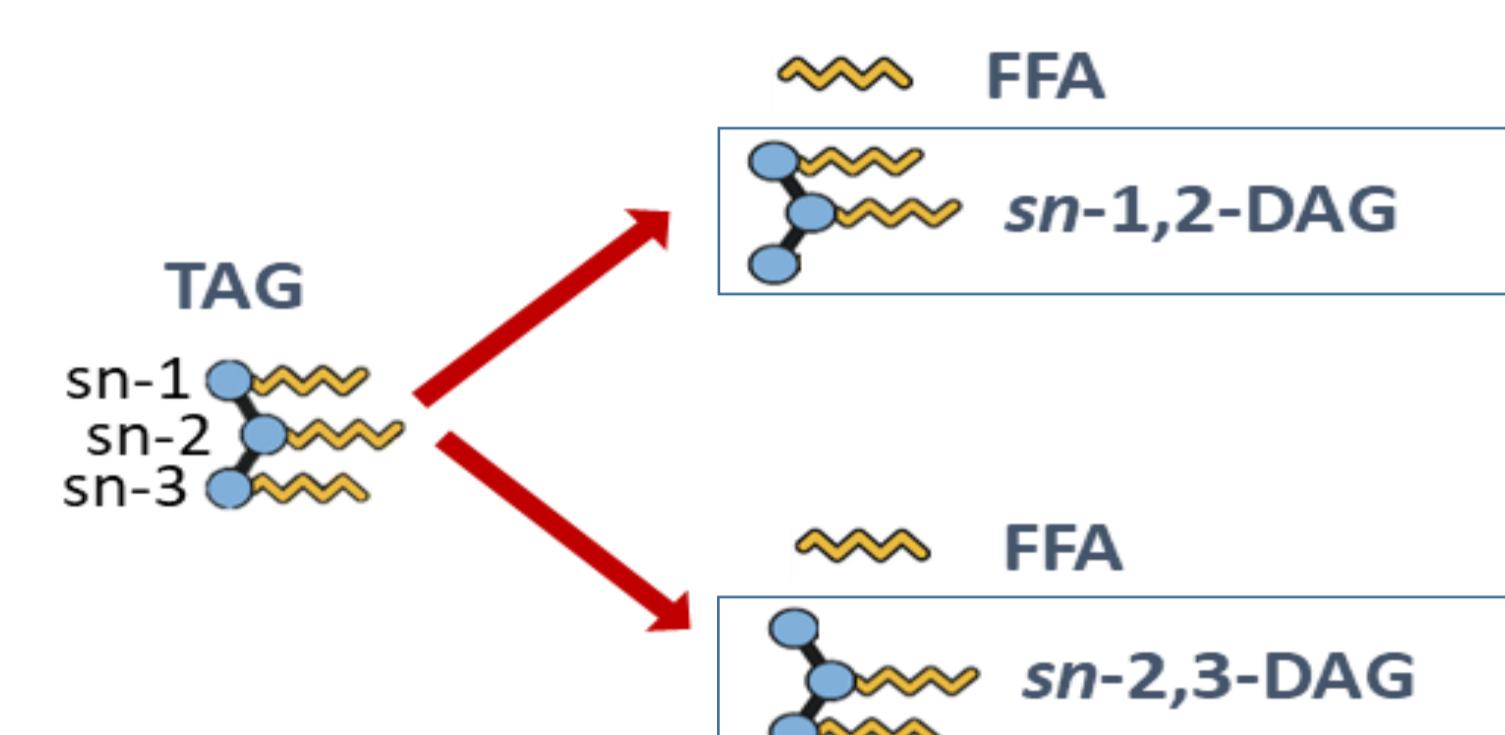
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Background and research objective

Human gastric lipase (HGL) stereospecificity



DAGs identified and quantified through expensive and laborious techniques

Research gap



Gastric lipid digestion mechanism?



- Lack of relevant and convenient substitute of HGL
- End-point quantification of digestion product(s)
- FFA release by titration
- Simple and reliable analytical platform is missing.

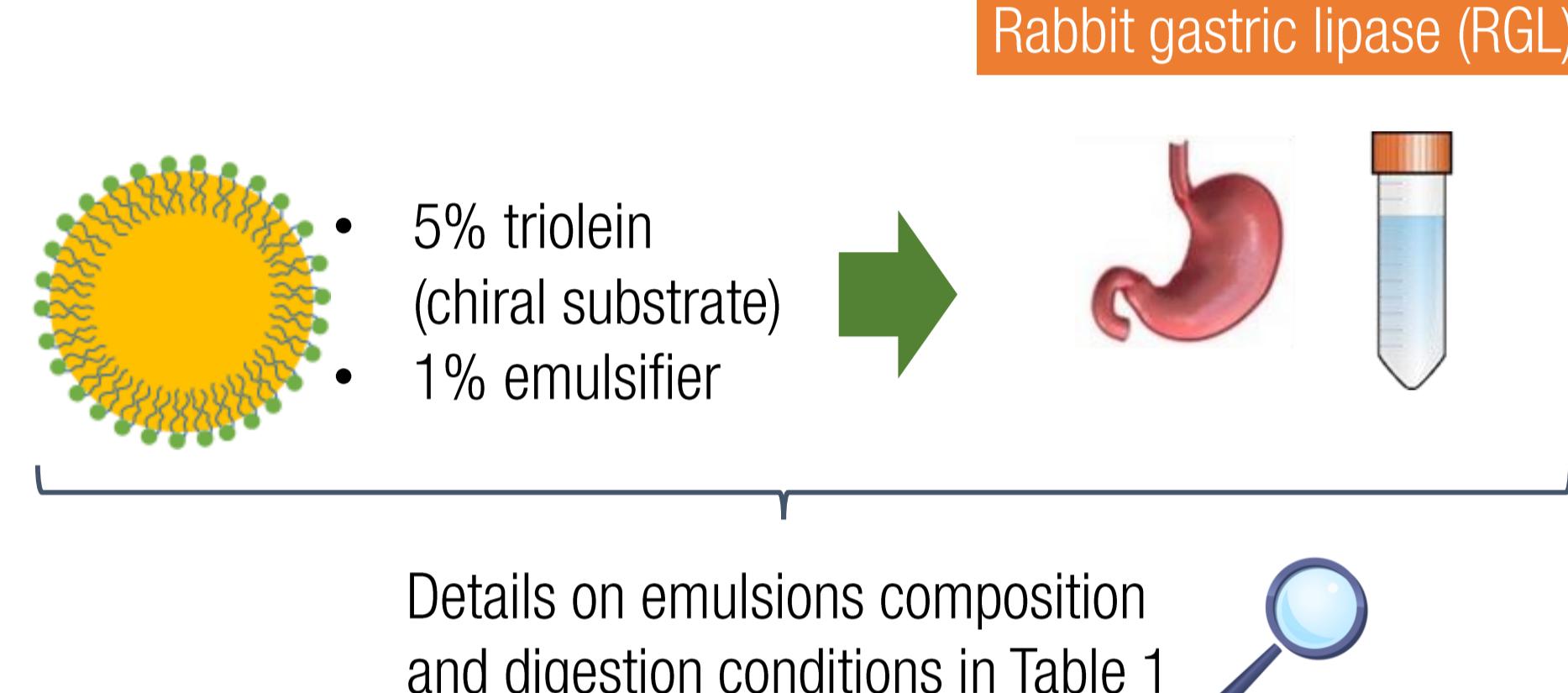
Research objective



To elucidate the molecular mechanism of *in vitro* gastric lipid digestion by integrating multiple lipolysis responses quantified as a function of digestion time.

Experimental design

1. Static *in vitro* digestion



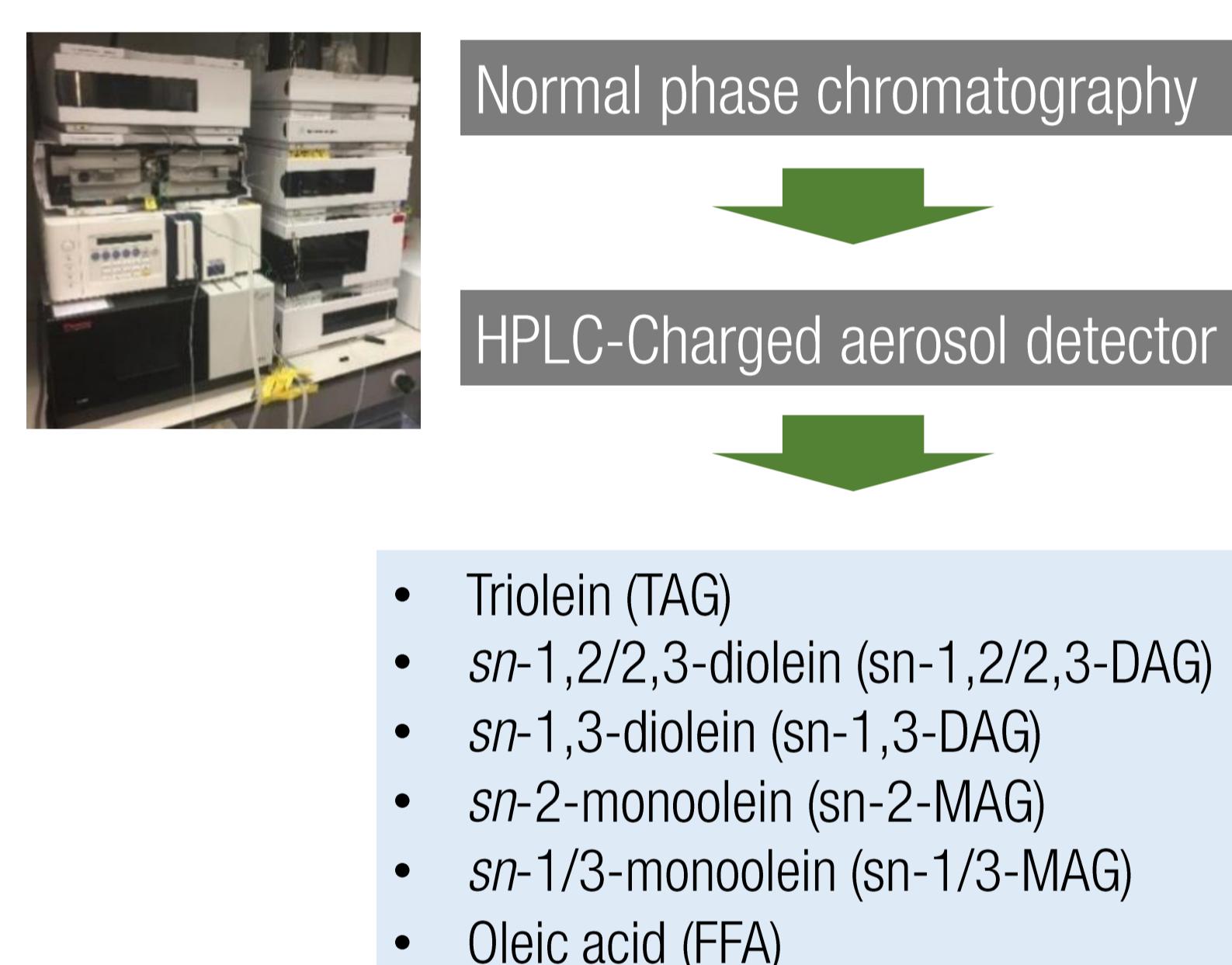
Details on emulsions composition and digestion conditions in Table 1

Table 1: Overview of data sets considered for the multi-response modeling

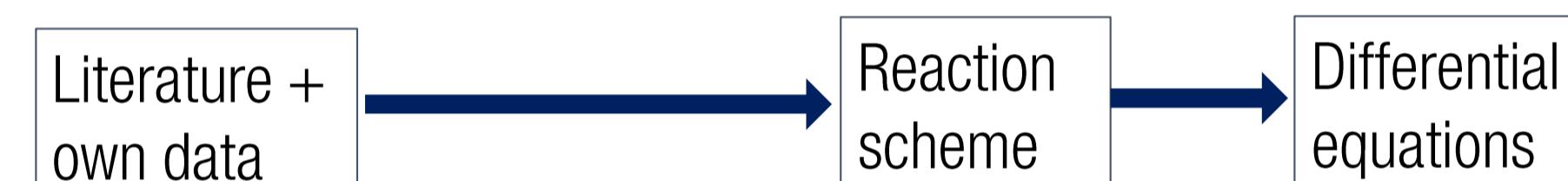
| Emulsifier type | Emulsion initial droplet size $d(4,3)$ (μm) | Emulsion name | <i>In vitro</i> gastric conditions |
|---|--|---------------|---|
| Data set 1: Effect of emulsion droplet size¹ | | | |
| Sodium taurodeoxycholate | 0.58 ± 0.03 | FE | pH 5.5, RGLA 20 U/mL (Minekus <i>et al.</i> , 2014 and Sams <i>et al.</i> , 2016) |
| Sodium taurodeoxycholate | 1.82 ± 0.02 | ME | |
| Data set 2: Effect of emulsion interfacial composition² | | | |
| Soy lecithin | 1.38 ± 0.09 | LEC | pH 3, RGLA 60 U/mL (Brodkorb <i>et al.</i> , 2019) |
| Soy protein isolate | 1.36 ± 0.02 | SPI | |
| Citrus pectin | 1.17 ± 0.01 | CP | |

RGLA: rabbit gastric lipase activity

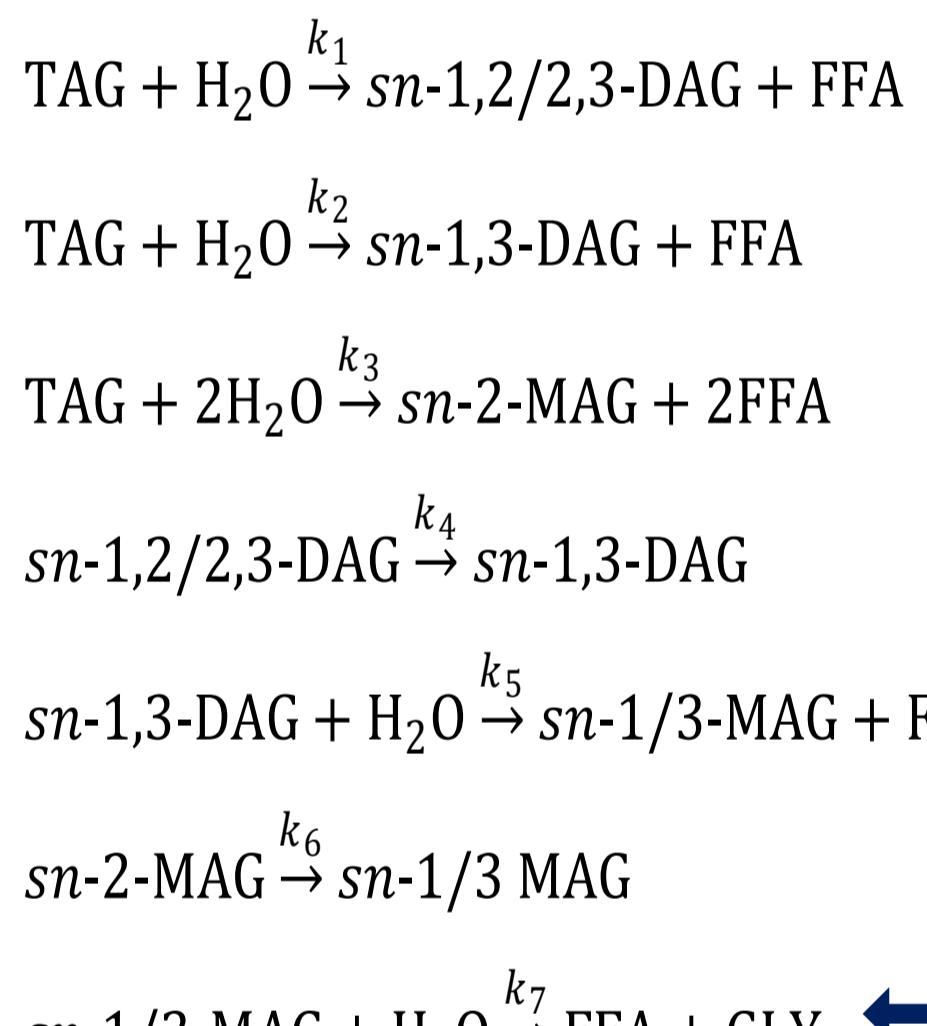
2. Analytical platform



3. Multi-response kinetic modeling



Reaction scheme 7



Reaction scheme 1

Start from the simplest scheme

Numerical method 'proc model' SAS

Iterative process

Model convergence?

$$\begin{aligned} \frac{d(\text{TAG})}{dt} &= -(k_1 + k_2 + k_3)(\text{TAG} - \text{TAG}_f) \\ \frac{d(\text{sn-1,2/2,3 DAG})}{dt} &= k_1(\text{TAG} - \text{TAG}_f) - k_4(\text{sn-1,2/2,3 DAG}) \\ \frac{d(\text{sn-1,3 DAG})}{dt} &= k_2(\text{TAG} - \text{TAG}_f) + k_4(\text{sn-1,2/2,3 DAG}) - k_5(\text{sn-1,3 DAG}) \\ \frac{d(\text{sn-2 MAG})}{dt} &= k_3(\text{TAG} - \text{TAG}_f) - k_6(\text{sn-2 MAG}) \\ \frac{d(\text{sn-1/3 MAG})}{dt} &= k_5(\text{sn-1,3 DAG}) + k_6(\text{sn-2 MAG}) - k_7(\text{sn-1/3 MAG}) \\ \frac{d(\text{FFA})}{dt} &= k_1(\text{TAG} - \text{TAG}_f) + k_2(\text{TAG} - \text{TAG}_f) + 2k_3(\text{TAG} - \text{TAG}_f) + k_5(\text{sn-1,3 DAG}) + k_7(\text{sn-1/3 MAG}) \\ \frac{d(\text{GLY})}{dt} &= k_7(\text{sn-1/3 MAG}) \end{aligned}$$

- Parity plots
- Residual plots
- Estimates error

Results and discussion

✓ Model corresponding to reaction scheme 7 converged

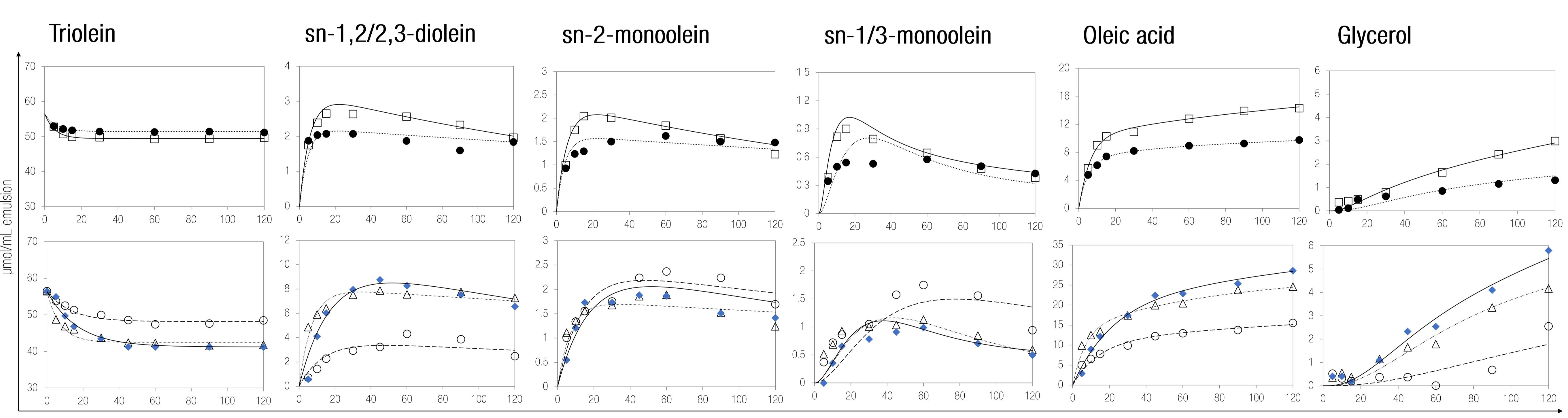


Table 2: Estimated parameters after subjecting data sets to multi-response modeling

| Data set 1 | | Data set 2 | | | |
|------------|-------------------|-------------------|-------------------|-------------------|-------------------|
| FE | ME | LEC | SPI | CP | |
| k_1 | 0.081 ± 0.004 | 0.098 ± 0.005 | 0.035 ± 0.003 | 0.039 ± 0.003 | 0.080 ± 0.005 |
| k_2 | 0.034 ± 0.003 | 0.051 ± 0.007 | 0.019 ± 0.004 | 0.010 ± 0.002 | 0.040 ± 0.005 |
| k_3 | 0.058 ± 0.003 | 0.071 ± 0.004 | 0.023 ± 0.002 | 0.009 ± 0.001 | 0.017 ± 0.001 |
| k_4 | 0.004 ± 0.000 | 0.002 ± 0.001 | 0.003 ± 0.001 | 0.003 ± 0.001 | 0.001 ± 0.001 |
| k_5 | 0.541 ± 0.285 | 0.097 ± 0.027 | 0.033 ± 0.013 | 0.086 ± 0.036 | 0.024 ± 0.005 |
| k_6 | 0.004 ± 0.000 | 0.002 ± 0.001 | 0.003 ± 0.001 | 0.003 ± 0.001 | 0.001 ± 0.001 |
| k_7 | 0.037 ± 0.003 | 0.024 ± 0.003 | 0.018 ± 0.003 | 0.059 ± 0.004 | 0.043 ± 0.005 |
| TAG_f | 49.68 ± 0.06 | 51.44 ± 0.05 | 48.36 ± 0.19 | 41.18 ± 0.29 | 42.47 ± 0.27 |

$k_{1,3,5,7}$: sn-1/3 positions cleavage

k_2 : sn-2 position cleavage

$k_{4,6}$: isomerization reactions

$k_1 > k_2 > k_{4,6}$

Conclusions

- The reaction mechanism was the same for all conditions. This demonstrates the validity of the mechanistic multi-response model under different static conditions (e.g. different pH and lipase activity values).
- These findings may be the starting point to develop *in silico* models to predict the gastric lipolysis based on emulsion design properties.

