



Background and research objective

Human gastric lipase (HGL) stereospecificity



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

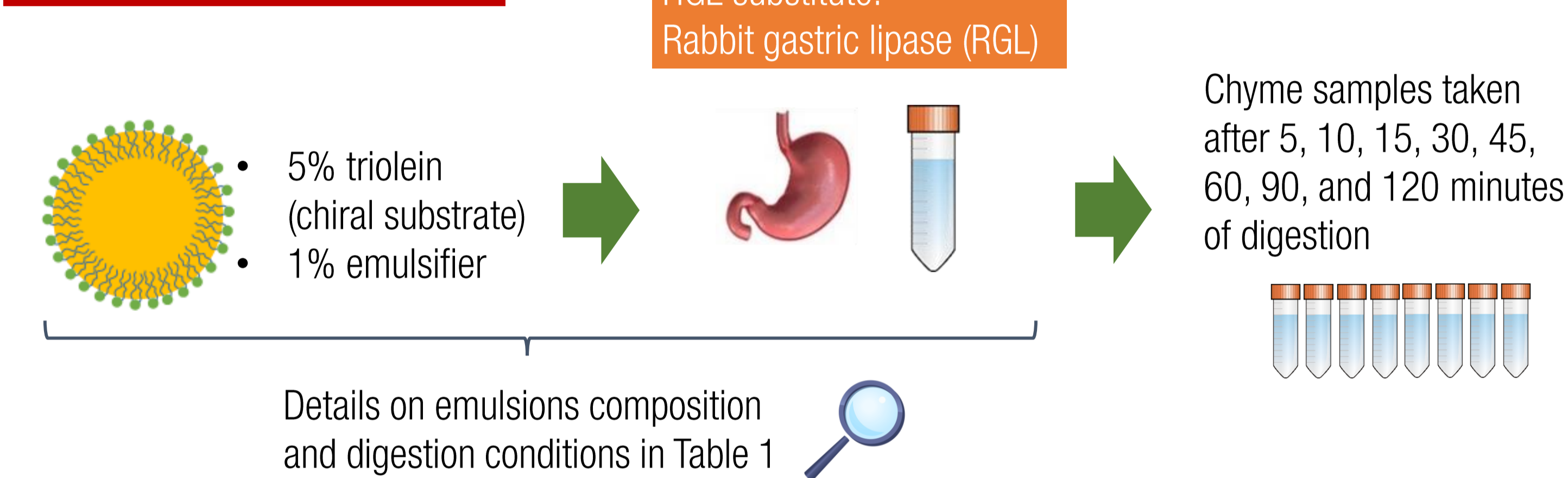
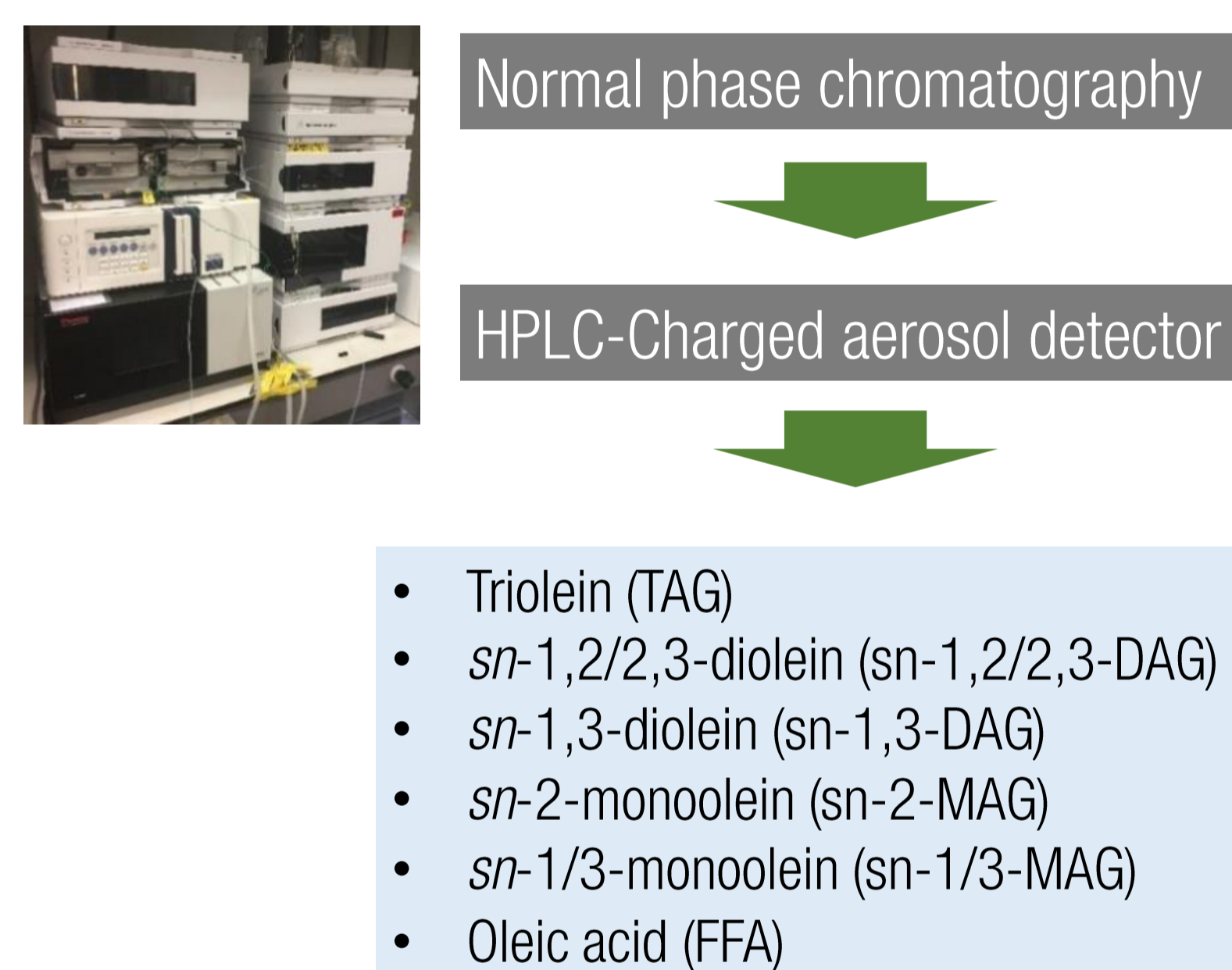


Table 1: Overview of data sets considered for the multi-response modeling

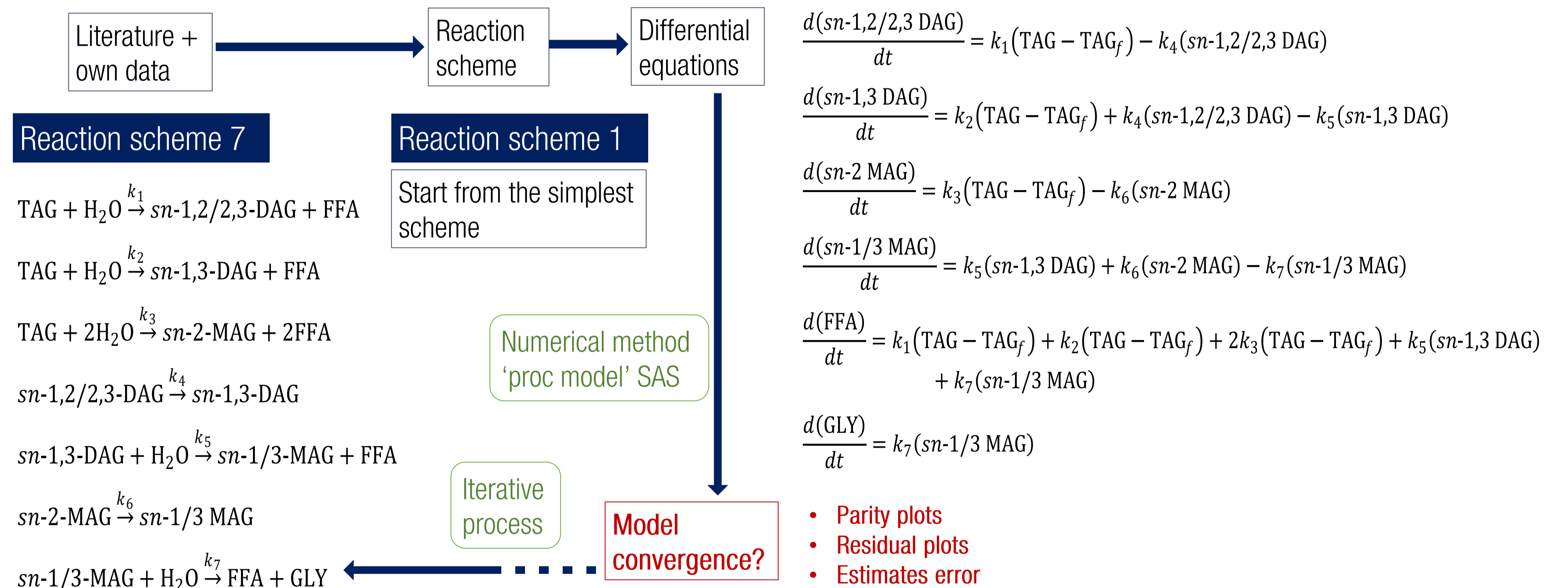
Emulsifier type	Emulsion initial droplet size $d(4,3)$ (μm)	Emulsion name	<i>In vitro</i> gastric conditions
Data set 1: Effect of emulsion droplet size ¹			
Sodium taurodeoxycholate	0.58 ± 0.03	FE	pH 5.5, RGLA 20 U/mL (Minekus <i>et al.</i> , 2014 and Sams <i>et al.</i> , 2016)
Sodium taurodeoxycholate	1.82 ± 0.02	ME	
Data set 2: Effect of emulsion interfacial composition ²			
Soy lecithin	1.38 ± 0.09	LEC	pH 3, RGLA 60 U/mL (Brodtkorb <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



Results and discussion

Model corresponding to reaction scheme 7 converged

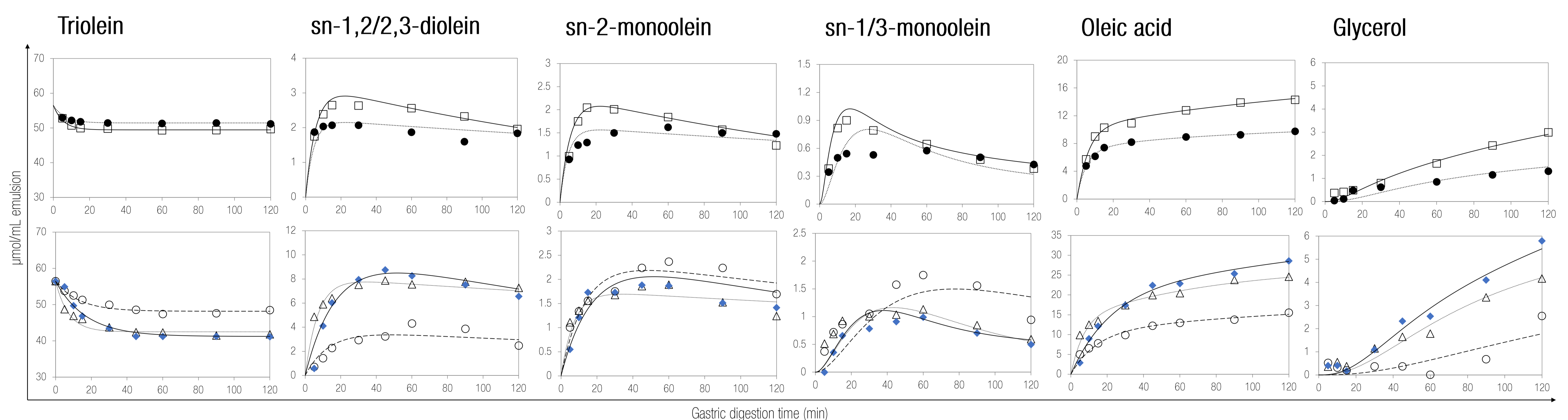


Table 2: Estimated parameters after subjecting data sets to multi-response modeling

	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.

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