Time (min)	HCl Inflow 1*		SGF + Pepsin Inflow 2*		Total liquid	Gastric
	Amount released (mL)	Rate (mL/min)	Amount released (mL)	Rate (mL/min)	 entered into HGS (mL) 	emptying (mL)
10	21	0.7	69	2.3	30	23
20					30	51
30					30	46
60	25	0.8	81	2.7	106	114
90	14	0.5	39	1.3	53	90
120	13	0.4	18	0.6	31	49
150	12	0.4	10	0.3	22	24
180	9	0.3	9	0.3	18	23
Total	94		226		320	420

Supplementary Table 1. Liquid Inflows and gastric emptying in the human gastric simulator

*The two inflows were maintained to mimicked the gastric secretions and dilution of food in the human stomach fed a solid meal, adapted from Malagelada et al., (1976).

Malagelada J-R, Longstreth GF, Summerskill WHJ, Go VLW. Measurement of gastric functions during digestion of ordinary solid meals in Man. *Gastroenterology*, 1976, **70**(2), 203-10.

Peptide ID	Mass	CS Polarity		Start	End	NCE	
	(<i>m/z</i>)	[<i>z</i>]	Polarity	(min)	(min)	NCL	
P7	808.8944	2	+	0.3	0.8	22	
P6	813.9051	2	+	0.3	0.8	22	
P8	1265.6046	2	+	2.2	2.9	16	
Р9	1164.9482	3	+	2.8	3.9	16	
P1	784.9270	2	+	4.2	4.6	15	
P1H	787.9303	2	+	4.2	4.6	15	
Р3	1032.5437	3	+	4.45	4.8	15	
P2	1132.0989	2	+	4.5	4.85	16	
P4	1029.5433	3	+	4.65	5.05	14	
P5	1304.0170	3	+	4.85	5.2	15	

Supplementary Table 2. Inclusion list for parallel reaction monitoring.

Order of list based on their relative retention time.

P1, LQLQPFPQPQLPY; P2, LQLQPFPQPQLPYPQPQPF; P3, LQLQPFPQPQLPYPQPHLPYPQPQPF; P4, LQLQPFPQPQLPYPQPQLPYPQPQPF; P5, LQLQPFPQPQLPYPQPQLPYPQPQLPYPQPQPF (33mer); P6, RPQQPYPQPQPQY; P7, RPQQPYPQSQPQY; P8, FQPSQQNPQAQGSFQPQQLPQF; P9, VRVPVPQLQPQNPSQQQPQEQVPLVQQQQF.

			<u> </u>			
	Peptide ID	Mass	Target Peak 1	Confirming	Target peak 2	
	replice ib	(<i>m/z</i>)	Talget Feak I	peak (T1C1)		
	P7	808.8944	1210.5965	1182.5973	407.1925	
	P6	813.9051	1220.6172	1192.6212	407.1925	
	P8	1265.6046	2141.0195	857.4502	391.1969	
	P9	1164.9482	1254.1615	1017.6191	987.5244	
	P1	784.9270	1290.7208	617.3293	279.1341	
	P1H	787.9303	1293.7296	618.3329	279.1341	
	P3	1032.5437	1417.7468	1305.1910	263.1391	
	P2	1132.0989	2002.0605	713.3612	488.2505	
	P4	1029.5433	713.3608	1550.8344	263.1389	
	P5	1304.0170	713.3613	2374.2735	263.1390	

Supplementary Table 3. Trace finder peptide quantitation ions settings.

Order of list based on their relative retention time.

Time	GE, %
10	7.1
20	22.9
30	35.7
60	59.3
90	77.6
120	85.5
150	88.5

Supplementary Table 4: Gastric emptying (GE) rates at different digestion times.

Adapted from (Malagelada, 1977; Malagelada, Longstreth, Summerskill, & Go, 1976).

Malagelada, J.-R. (1977). Quantification of Gastric Solid-Liquid Discrimination During Digestion of Ordinary Meals. *Gastroenterology*, 72(6), 1264-1267. doi:10.1016/S0016-5085(77)80024-3
Malagelada, J.-R., Longstreth, G. F., Summerskill, W. H. J., & Go, V. L. W. (1976). Measurement of Gastric Functions During Digestion of Ordinary Solid Meals in Man. *Gastroenterology*, 70(2), 203-210. doi:10.1016/S0016-5085(76)80010-8

Calculation 1. Example of the calculation used to determine the free amino groups present in the small intestine

The calculation of the amount of free amino groups present in the small intestine at 30 min is used as an example to show how the observed data in this study and the gastric emptying values in Supplementary Table 3 were used in the calculation.

Based on Figure 1, the amount of free amino groups present in the small intestine at 30 min are the free amino groups in the 10 min gastric and 20 min small intestinal digested sample and free amino groups in the 20 min gastric and 10 min small intestinal digested sample. The free amino groups of each sample were calculated using the equation described in Calculations of Materials and Methods, and the following information was obtained:

- Amount of free amino groups in the 10 min gastric and 20 min small intestinal digested sample was 5,603 μ g/g of bread
- Amount of free amino groups in the 20 min gastric and 10 min small intestinal digested sample was 5,794 μg/g of bread
- The gastric emptying at 10 min 7.1% (GE₁₀)
- The gastric emptying at 20 min 22.9% (GE₂₀)

The calculation was done based on one gram of bread (DM).

Amount of bread (mg) released from the stomach at 0-10 min

$$1,000 x \frac{GE_{10}}{100}$$
$$1,000 x \frac{7.1}{100} = 71.0$$

Amount of bread (mg) released from the stomach at 10-20 min

$$1,000 x \frac{GE_{20} - GE_{10}}{100}$$
$$1,000 x \frac{(22.9 - 7.1)}{100} = 158.8$$

Amount of free amino groups present in the bread released from the stomach at 10 min

= 0.071 g x 5,603 μ g/g of bread = 397.81 μ g/g of bread

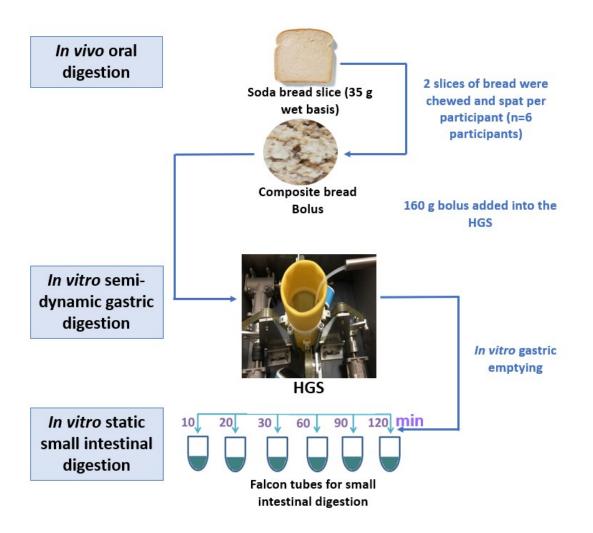
Amount of free amino groups present in the bread released from the stomach at 20 min

= 0.158 g x 5,794 μ g/g of bread = 915.45 μ g/g of bread

Total free amino groups present in the small intestine at 30 min

= 397.81 + 915.45 = 1,313.26 μg/g of bread

The same equations were used to determine the amount of epitopes in the small intestine at 30 min.



Supplementary Figure 1. Graphical illustration of dynamic *in vivo* and *in vitro* gastrointestinal digestion conducted during the experiment.

HGS, human gastric simulator