

Supplementary data

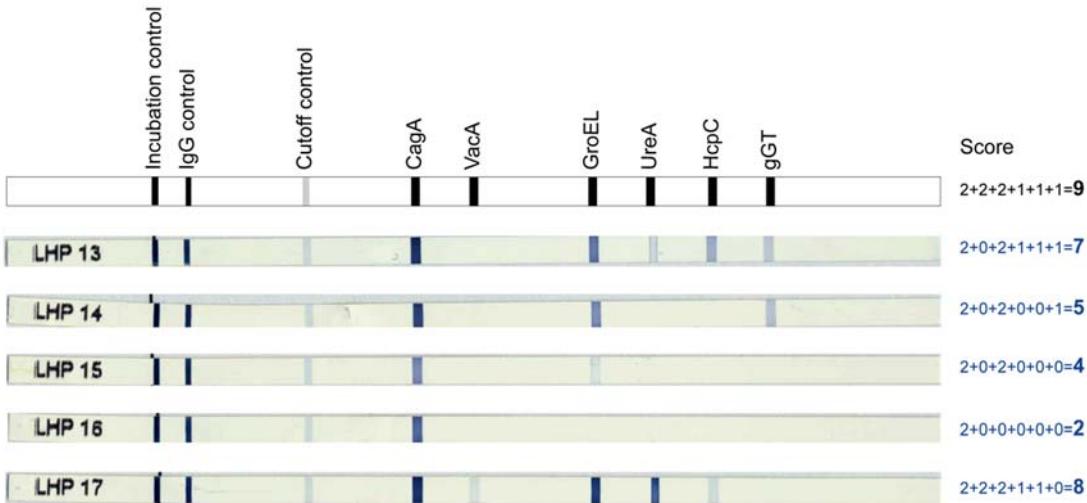
**Validation of a novel immuno-line assay for patient stratification according to virulence of the infecting *Helicobacter pylori* strain and eradication status**

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Histology	GERMANY (D)				NETHERLANDS (NL)			
	positive	negative	eradicated	total	positive	negative	eradicate d	total
N	409	447	435	1291	35	91	102	228
%	31.7	34.6	33.7	100	15.4	39.9	44.7	100
mean age	52.7 (± 16.1)	48.2 (± 16.6)	59.3 (± 13.7)	53.6 (± 16.2)	62.8 (± 12.6)	61.1 (± 10.6)	61.5 (± 11.7)	62.2 (± 11.3)
Gender (f/m)	0.53/0.47	0.55/0.45	0.59/0.41	0.55/0.45	0.63/0.37	0.49/0.51	0.40/0.60	0.48/0.52
Serology	409	447	281	1137	35	91	102	228
mean recomLine score	4.8 (± 1.9)	0.24 (± 0.8)	3.2 (± 1.8)	N/A <sup>1</sup>	5.1 (± 2.1)	2.7 (± 2.2)	3.4 (± 2.1)	2.5 (± 2.3)
recomLine +	400 <sup>4</sup>	40	210 <sup>4</sup>	650	32	14	65 <sup>4</sup>	111
% recomLine +	98.3	9.0	74.7	57.4	91.4	15.4	84.4	54.7
recomLine -	7	404	71	482	3	77	12	92
% recomLine -	1.7	91.0	25.3	42.6	8.6	84.6	15.6	45.3
Excluded for analysis	2 <sup>2</sup>	3 <sup>2</sup>	154 <sup>3</sup>	159	35 <sup>1</sup>	91 <sup>1</sup>	25 <sup>3</sup>	151

Supplementary Table 1. Study Populations

<sup>1</sup>not applicable; <sup>2</sup>borderline; <sup>3</sup>no documentation; <sup>4</sup>included in analysis



Supplementary Figure 1. The result of five representative *recomLine* tests incubated with individual patient sera are shown (LHP13 – LHP17). The assignment of the bands is given in the cartoon above. The test score calculation is given beside the *recomLine* tests. LHP 16 shows only a CagA band and thus represents a typical test result for an eradicated patient.

		Serology (recomLine)		total
		positive	negative	
Histology	positive	400	7	407
	negative	40*	404	444
total		440	411	851

Supplementary Table 2. Cross Table Histology vs. Serology (recomLine)

Sensitivity (recomLine): 98.3%

Specificity (recomLine): 91.0%

\*Corrected Specificity (recomLine): 95.5%

(\*from 40 false positives, 21 presenting isolated CagA were considered as eradicated)

**Chronicity of inflammation**

N	mild	moderate	severe	Total	p
	64	297	28	389	
<b>CagA positivity (%)</b>	59.4	70.4	89.3	69.9	0.015
<b>VacA positivity (%)</b>	17.2	20.2	46.4	21.6	0.004
<b>GroEL positivity (%)</b>	90.6	85.5	89.3	86.6	0.506
<b>UreA positivity (%)</b>	23.4	30.3	42.9	30.1	0.173
<b>HcpC positivity (%)</b>	45.3	44.4	50.0	45.0	0.851
<b>gGT positivity (%)</b>	46.9	54.9	57.1	53.7	0.473

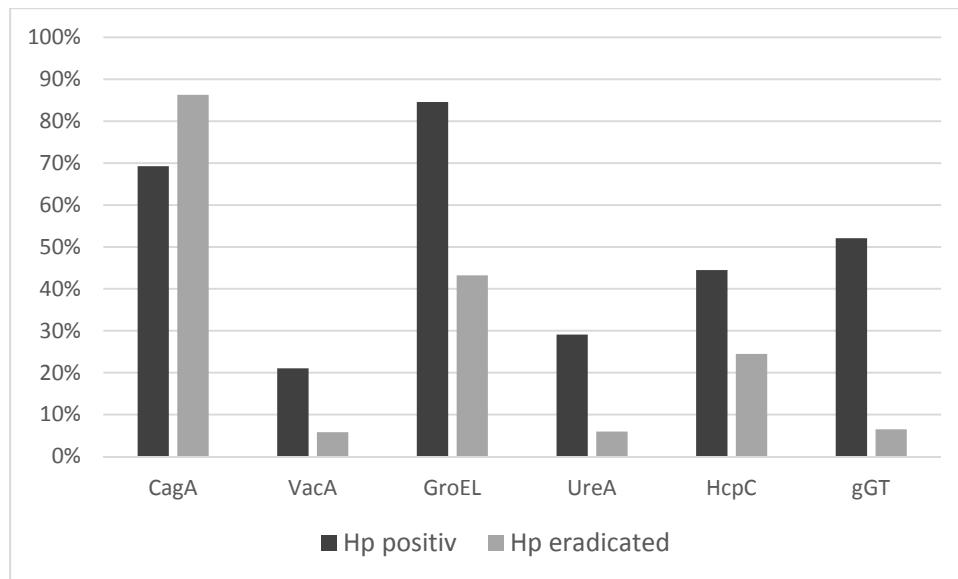
**Activity of inflammation**

	mild	moderate	severe	Total	
<b>N</b>	229	138	20	387	p
<b>CagA positivity (%)</b>	63.8	78.3	85.0	70.0	0.0044
<b>VacA positivity (%)</b>	20.5	21.7	35.0	21.7	0.3226
<b>GroEL positivity (%)</b>	87.3	84.1	95.0	86.6	0.3533
<b>UreA positivity (%)</b>	29.7	29.0	45.0	30.2	0.3336
<b>HcpC positivity (%)</b>	46.7	44.9	30.0	45.2	0.3536
<b>gGT positivity (%)</b>	52.0	58.7	50.0	54.3	0.4228

Supplementary Table 3. Association between gastritis parameters and single *H. pylori* antigen positivity

	Signs of pathology						
N	absent	atrophy	metaplasia	ulcer	GC	Total	p
<b>CagA positivity (%)</b>	299	22	53	16	3	393	0.001
<b>VacA positivity (%)</b>	63.9	81.8	88.7	87.5	100.0	69.5	0.157
<b>GroEL positivity (%)</b>	19.7	31.8	24.5	31.3	66.7	21.9	0.746
<b>UreA positivity (%)</b>	87.3	86.4	81.1	87.5	100.0	86.5	0.333
<b>HcpC positivity (%)</b>	29.8	18.2	34.0	43.8	0.0	30.0	0.373
<b>gGT positivity (%)</b>	47.5	31.8	35.8	43.8	33.3	44.8	0.885
	53.2	63.6	52.8	56.3	66.7	53.9	

Supplementary Table 4. Association between most severe lesions and single antigen *H. pylori* positivity



**Supplementary Figure 2.** Antigen frequencies of *H. pylori* positive and eradicated patients  
CagA was significantly ( $p<0.01$ ) more often positive in eradicated patients (86.3%) than in *H. pylori* positive patients (69.2%), while all other immune responses were lower in the eradicated cohort