

Identification of candidate *Coxiella burnetii* T-cell epitopes for a novel human Q fever vaccine

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Background – Q fever

- Zoonosis primarily spread by small ruminants (goat, sheep) and cattle
- Neglected disease with world-wide prevalence; poor surveillance, studies indicate substantial contribution to “fever & infectious endocarditis of unknown origin”
- Caused by the obligate intracellular bacterium *Coxiella burnetii* (Cb) – infects macrophages
- Highly stable and contagious (potential biothreat agent)
- Long-term complications common (chronic fatigue, persistent infection = chronic Q fever)
- Outbreak in the Netherlands (2007-2010) affected more than 40,000 individuals
- T-cell produced IFNγ key in inducing phagosome/lysosome fusion to eliminate infection
- Current whole-cell Q fever vaccine (Q-VAX®) only licensed in Australia and reactogenic in individuals previously exposed to Cb
- Epitope-based subunit vaccines are expected to be non-reactogenic and show promise in murine studies

Objective & Approach

Identify target epitopes to develop a non-reactogenic T-cell-targeted vaccine to prevent Q fever disease in humans

- T-cell epitopes were predicted from
- Cb **type IV secretion system (T4SS) substrates**, expected to enter the class I antigen processing pathway and thus to trigger CD8+ T cell responses
 - Cb **sero-reactive proteins** known to elicit antibody responses in mice and humans

Epitopes were down-selected for cross-strain conservation, strong HLA binding, broad HLA coverage and low cross-reactivity with human/microbiome proteins. HLA binding was assessed in vitro, immunogenicity in HLA-DR3 transgenic mice and antigenicity in naturally exposed volunteers. Reactogenicity was excluded using a sensitized guinea pig model (data not shown).

In silico epitope selection

Results:
65 HLA class I epitopes (10 per HLA I supertype allele) and 50 promiscuous class II epitope clusters were down-selected (no more than 2 epitopes / source Ag)

HLA Class	Class I		Class II
Antigen source	T4SS Effector	Sero-reactive	Sero-reactive
Antigen count	53	40	40
Epitopes predicted	8,643	5,100	282
Conserved across Cb	3,971	4,578	188
Highly immunogenic	1,710	1,945	153
Different from human	1,511	1,558	98
w/o synthesis issues	1,108	1,163	81
Selected	30	35	50

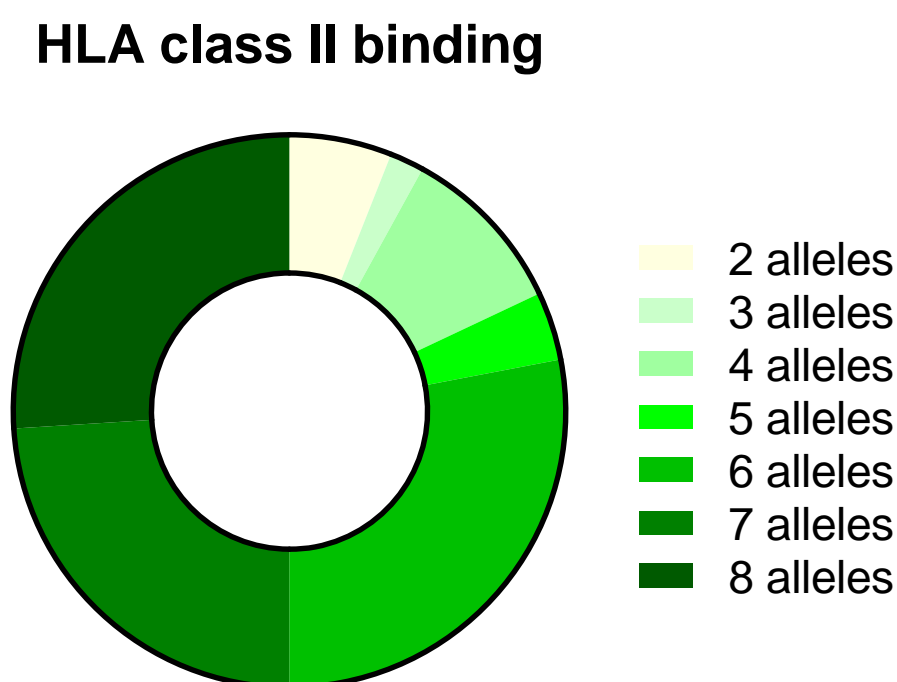
In vitro HLA binding

Results:
In vitro competition assays confirm that

- HLA class II peptides are promiscuous binders
- 89% of class I and 75% of class II predictions are correct

Peptides have potential for stimulating Cb-specific immune responses across a broad range of HLA types.

Figure 1: HLA class II peptide binding was assessed to the main supertypes (DRB1*0101, *0301, *0401, *0701, *0801, *1101, *1301, *1501 – each peptide against each allele)

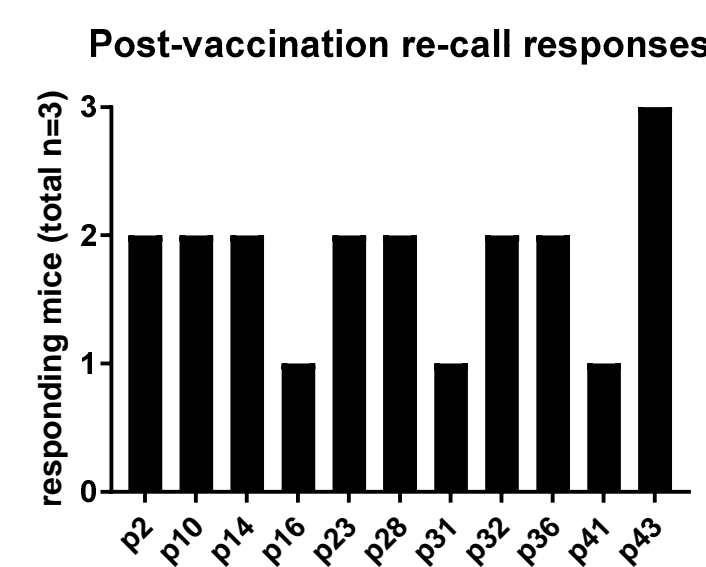


In vivo immunogenicity

Results:
11/50 HLA class II peptides were immunogenic in tgHLA-DR3 mice. All 11 were predicted to bind HLA-DR3, and all except one (p36) also showed binding to DRB1*0301 in vitro.

Peptides show capacity to elicit de novo responses in vivo.

Figure 2: 5 groups of mice (n=3 each) were immunized with 10 epitopes each by heterologous DNA/DNA/peptide/ peptide prime-boost immunizations. Re-call responses were assessed by IFNγ ELISpot. Response were defined positive when spot counts reached (i) a stimulation index of at least 2 above the matched negative control wells (ii) >50 spot/million splenocytes (above background) and were (iii) statistically higher (p<0.05) than in mock immunized mice (Student's t-test)



Human cohort

Antigenicity of the peptides was assessed in naturally Cb-exposed individuals from the Dutch village of Herpen. Volunteers were grouped by clinical Q fever history (2007-2010 outbreak) and cellular responses to whole cell Cb.

Group A: IFA and IGRA-; no clinical Q fever history
Group B: IGRA+, no clinical Q fever history (asymptomatic)
Group C: IGRA+, past symptomatic Q fever episode

Two partially overlapping subgroups of n=77 individuals each were selected for class I and class II peptide screening by cultured ELISpot.

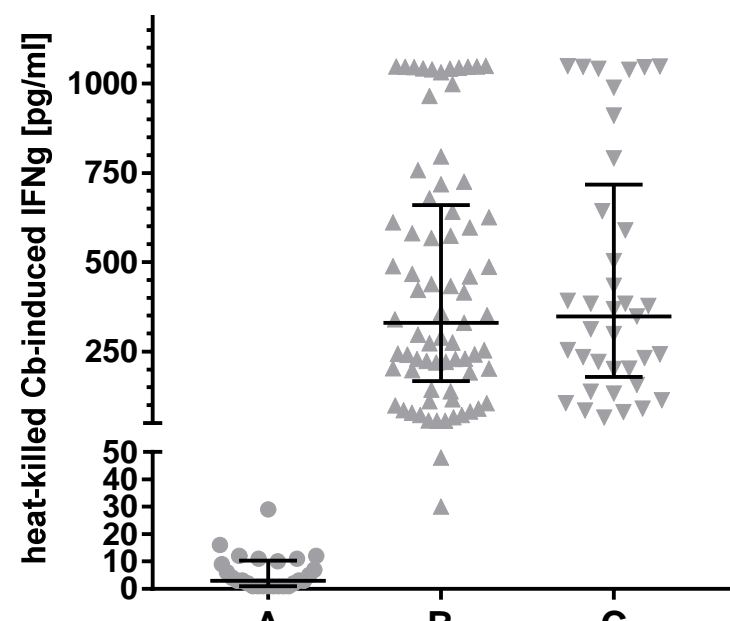


Figure 3: Cellular responses to whole cell heat-killed Cb (strain Cb2009-02629) were assessed for 136 volunteers from Herpen in a whole blood IFNγ release assay (Q-detect™ IGRA)

Discussion / Conclusions

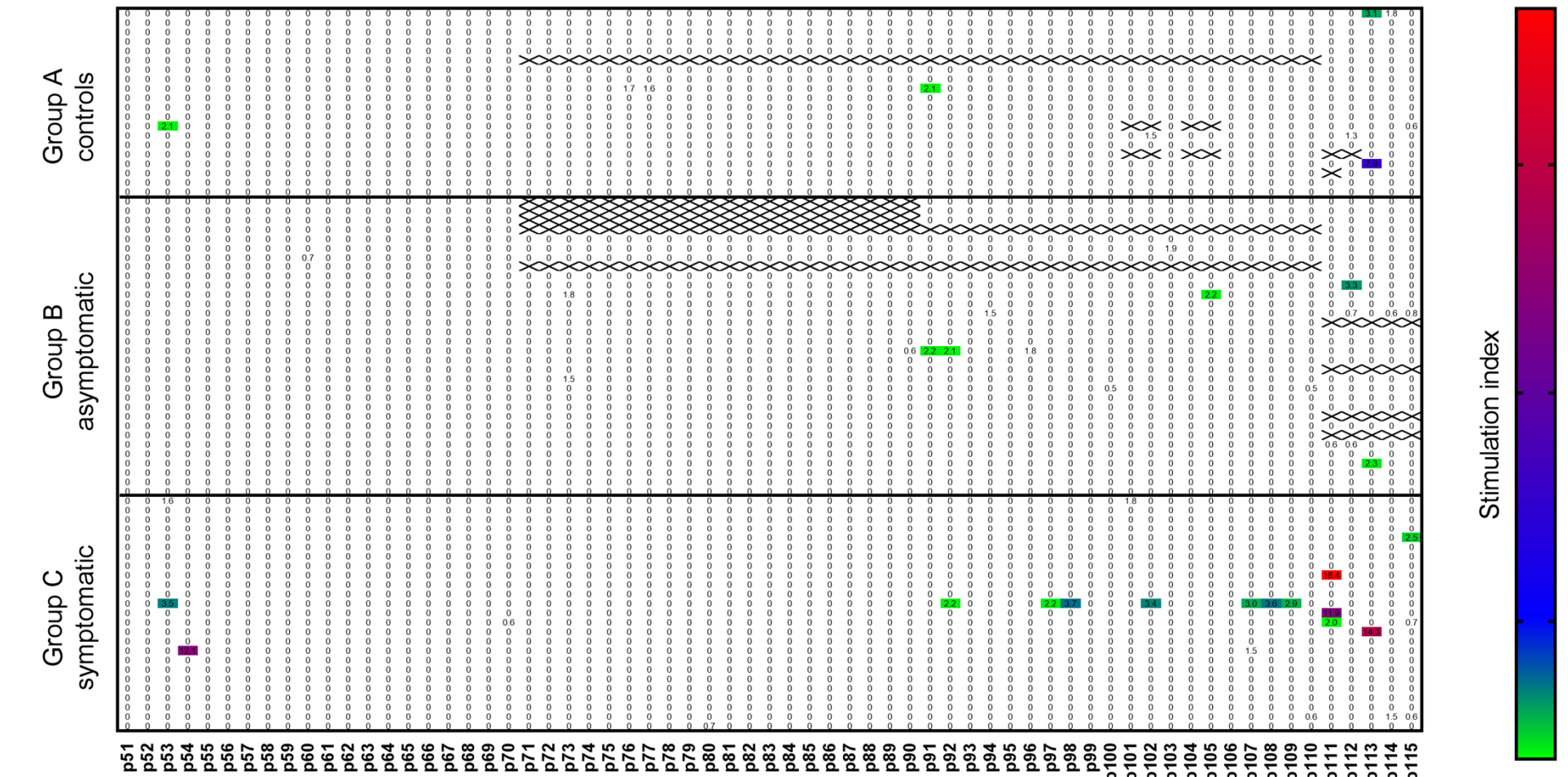
- Sero-reactivity guided screening approach is very efficient in identifying relevant source antigens for HLA class II (21/50 = 42% highly antigenic)
- Natural exposure to Cb induces long-lived responses to immunodominant, promiscuous and conserved HLA class II T-cell epitope clusters
- Weak responses to T4SS HLA class I epitopes
 - Greater number of binding opportunities for class II clusters than for individual class I epitopes
 - CD4 repertoire larger than CD8 repertoire?
 - CD8 responses more short-lived?
- **Class II epitope clusters are key candidates for an epitope-based Q fever vaccine to elicit sustained T-cell memory that can be boosted and recalled by natural exposure**

Peptide re-call responses in exposed human volunteers

- Results:**
- HLA class I peptides recall only a few infrequent responses
 - HLA class II peptides recall a wide range of responses, particularly amongst IGRA+ volunteers
 - 21 of the 50 HLA class II peptides are highly antigenic = recognized by 10-28% of IGRA+ volunteers
 - Amongst these 21 peptides are 5 epitope pairs where both peptides originate from the same source antigen
 - Q-detect™ IGRA responses to whole Cb correlate with number of Cb peptides recognized per volunteer (Spearman p=0.026)
 - Past Q fever asymptomatic and symptomatic volunteers do not differ in their cumulative peptide response

Promiscuous HLA class II peptides are antigenic and recall long-lasting memory responses from natural exposed IGRA+ volunteers following, irrespective of their clinical Q-fever history.

A – HLA class I peptide screen



B – HLA class II peptide screen

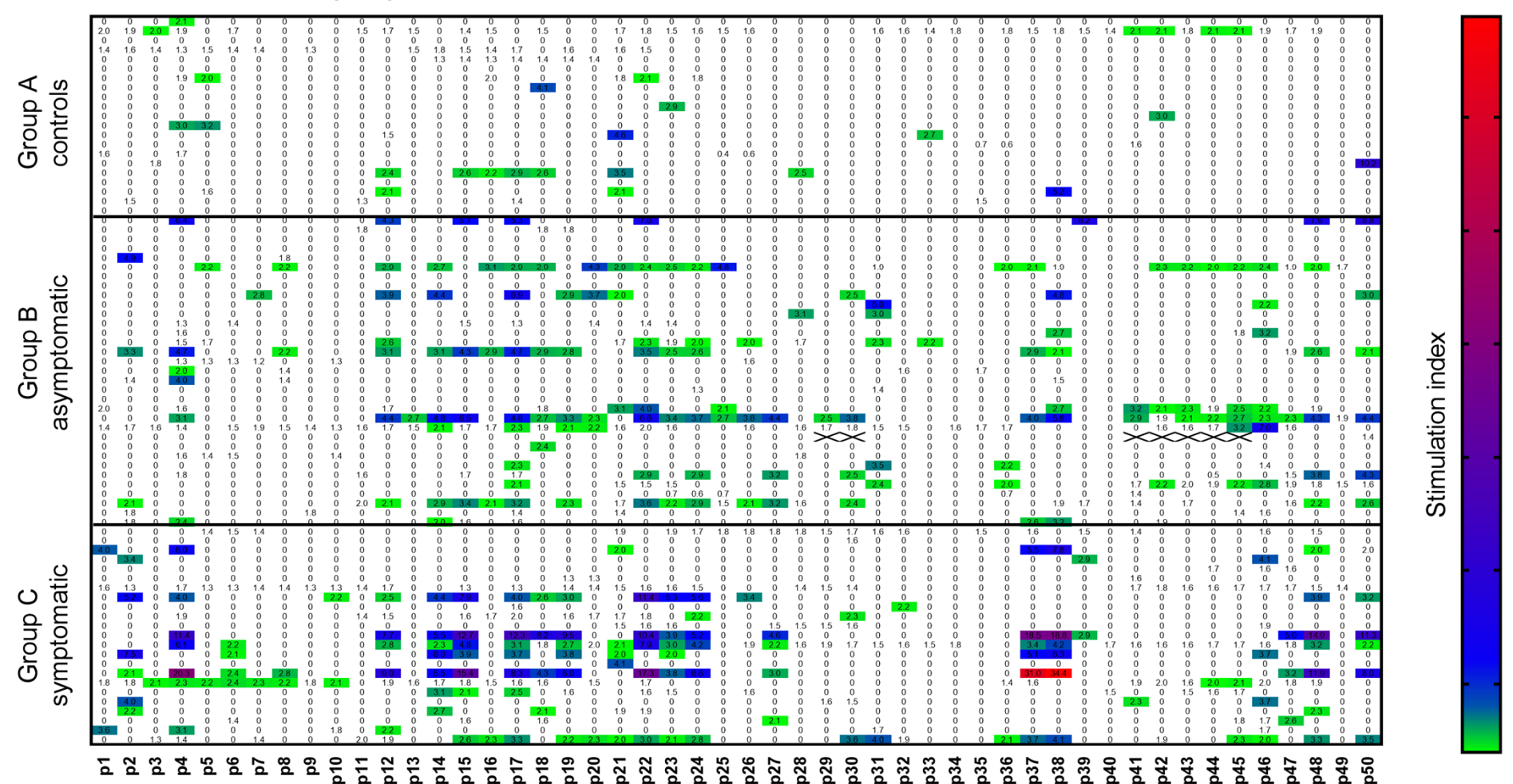


Figure 4: Re-call responses to HLA class I (A) and II peptides (B) assessed by cultured ELISpot for group A, B and C volunteers. Responses were defined positive when (i) significantly higher than spot counts in matched negative control wells from the same expansion culture by one-way ANOVA with Holm-Šidák multiple comparison correction post-hoc test, reached (ii) a stimulation index of at least 2 above the matched negative control wells and (iii) an absolute cut-off of >10 SFU/well.

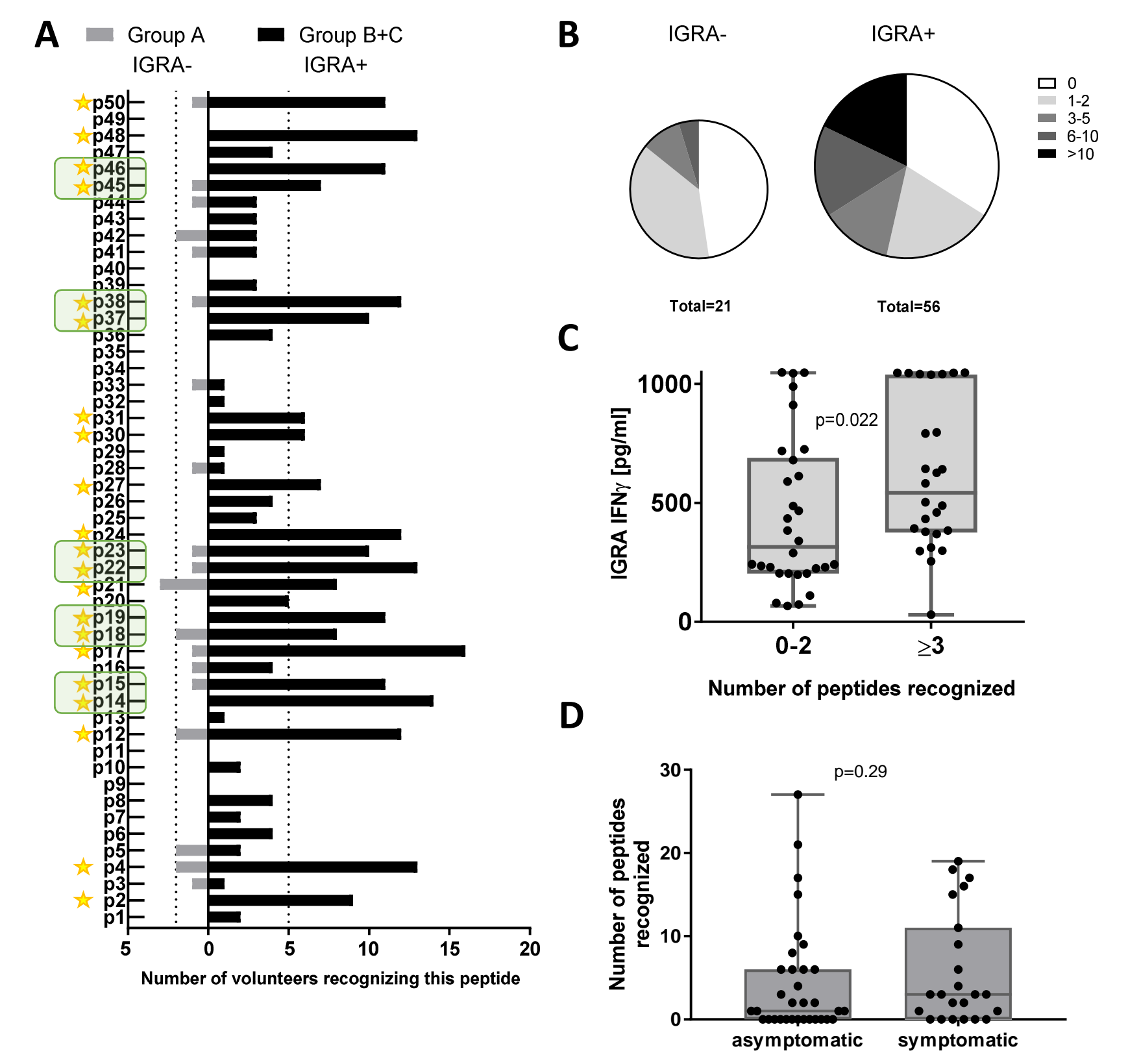


Figure 5: Re-call responses to HLA class II peptides: (A) Proportion of responding volunteers amongst group A controls (n=21) and group B+C IGRA+ volunteers (n=56) **Yellow stars** in (A) indicate highly antigenic peptides recognized by >10% of IGRA+ volunteers, and **green boxes** indicate epitope pairs from the same source antigen (for highly antigenic peptides only).

(B) Number of peptides recognized by individual volunteers in both groups.

(C) IGRA responses to whole cell heat-killed Cb for volunteers recognizing none or few (0-2) or a larger number of peptides (≥3).

(D) Number of peptides recognized by individual volunteers compared for those IGRA+ volunteers that experienced either asymptomatic Cb infection or a clinical Q fever episode during the 2007-2010 epidemic.

Disclosures

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Ethics statement

Animal research protocols for studies with HLA-DR3 transgenic mice performed by EpiVax were reviewed and approved by TGA Sciences Incorporated Institutional Animal Care and Use Committee (P07-10R20-EV69, P07-10R20-EV71). Animal research protocols for guinea pig experiments were reviewed and approved by the Colorado State University Institutional Animal Care and Use Committee (14-5305A, 16-6844A). The human study was reviewed and approved by the Medical Ethical Committee Brabant (Tilburg, Netherlands, NL51305.028.15) and all donors provided written informed consent.