CD14 inhibition mitigates myocardial damage and dysfunction following myocardial infarction



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Background

- MIR injury causes focal tissue scarring and initiates an excessive immune response leading to scar expansion, adverse remodelling and LV dysfunction, predisposing patients with poor LV function at admission (EF<40) to debilitating heart failure with reduced ejection fraction (HFrEF)¹
- CD14 is a master regulator of the macrophage response to damage associated molecular patterns (DAMPs) released with MIR
- Pro-inflammatory CD14⁺⁺ macrophages are instrumental to the propagation of adverse inflammatory signals and secondary damage¹
- Atibuclimab, a novel anti-CD14 monoclonal antibody, has been developed as a safe and selective inhibitor of CD14 without immune suppression

Findings

Clinically significant protection against left ventricular dysfunction at <u>7 days</u> post-MIR

Compared to SAL and ISO, at 7d post-MIR surgery (n=3grps x 15/grp), anti-CD14 treatment significantly improved LV EF by >30 % $(33\pm2 \text{ vs } 25\pm1 \text{ in SAL}, 24\pm2 \text{ \% in ISO}, p<0.001)$; stroke work by >38 % $(1662\pm69 \text{ vs } 1103\pm63 \text{ in SAL}, 1201\pm96 \text{ mmHg x ul in ISO}, p<0.001)$, -dV/dT by >24 % (-867±67 vs 563±30 in SAL, 700±59 ul / s in ISO, p<0.01) and peak power by >70 % (14.4±2.8 vs 6.7±0.7 in SAL, 8.4±1.1 mmHg x ml / s in ISO, p<0.05), see Fig 5A-D.



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Aims

To assess safety, immunological modulation and cardioprotection with CD14 inhibition given at reperfusion in rigorous murine studies of MIR injury

Hypothesis

Anti-CD14 treatment given at time of reperfusion protects against progressive left ventricular damage and dysfunction, is associated with modulation of macrophages, independently of acute damage

Project Overview

Animal ethics All experiments were approved by the ARA Animal Ethics Committee (E/7610/2019/B).

Study design, models and methods Adult male C57BI6 mice underwent MIR surgery to occlude the left anterior descending coronary artery for 60mins prior to reperfusion (Fig 1). Mice were randomised and coded, receiving IV saline (SAL, n=50), anti-CD14 antibody (n=50) or isotype control (ISO, n=50) at reperfusion and again at 24h. Assessments included infarct size (by Evan's blue/TTC), plasma troponin, bulk tissue RNA seq, flow cytometry, single cell RNA seq, histological fibrosis, immunohistochemistry, LV echocardiography (PLAX), and LV pressure-volume catheterisation at 1, 3 or 7 days post-MIR. Fig 5. LV function 7 days post-MIR A) Representative echocardiographic images at end-diastole and -systole, B) corresponding echocardiographic parameters, C) representative arterial blood pressures (upper) and pressure-volume loops (lower), and D) corresponding haemodynamic parameters. Scale bar 2mm.



Fig 6. Post-mortem tissue imaging A) From left – right: whole heart photgraphs, brightfield imaging of mid-ventricular section stained with picrosirius red, and immunofluorescent imaging of mid-ventricular section with channels for DAPI (blue), CD68 (green), troponin-I (pink) and merged, B) corresponding histological parameters lesion size (by picrosirius red staining) and CD68+ cell infiltrate abundance (immunohistochemistry). Scale bar 200µm.

Independent of acute infarct size or injury at <u>1 day</u> post-MIR

No differences were observed in area-at-risk or infarct size (measured by Evan's blue/TTC dual staining method, Fig 7A), or circulating cardiac troponin-I (cTnI, by



Anti-CD14 significantly treatment reduced scar size $(48.2\pm2.5 \text{ vs})$ 58.9±2.5 in SAL, 59.0±2.2 % total area in ISO, p<0.01) and CD68+ infiltrates $(22.2\pm0.8 \text{ vs } 26.6\pm1.0 \text{ in})$ SAL, 27.4 ± 1.5 % total cells in ISO, p<0.01). No differences were observed for heart, lung, kidney, or spleen weights at 7 days post-MIR. No rupture was observed in any mice in these studies, indicating consistently of reperfusion in model (equivalent to reestablishing vessel patency clinically Percutaneous Coronary with Intervention; PCI), as well as tx safety.





Fig 1. Three-phase preclinical trial design Hypothesis-driven studies were performed in series and parallel with independent cohorts for End-D1, D3 and D7 endpoints.

Statistics One-way ANOVA and Tukey (parametric) or Kruskal-Wallis (non-parametric) post-hoc tests were applied while blinded. All graphed data presented as mean \pm SD, in-text as mean \pm SE.

Preclinical Trial Rigour & Reproducibility

All mice were **randomised** and each study **independently blinded** (treatments coded to A, B, C) throughout all *in vivo* and post-mortem procedures, data acquisition, data analysis and reporting. **Exclusion criteria** were predetermined to be model insufficiency (lack of ST-segment elevation during surgery [Fig 2]

or area-at-risk at 24h <35% [Fig 3]) or technical insufficiency of endpoint (e.g. ruptured vessel during catheterisation).



ruptured vessel during catheterisation). Fig 2. Example of ST-segment Elevation (STE) after left anterior descending coronary artery ligation in mice



Fig 3. Assessment of area-at-risk (AAR) by echocardiographic strain at 24h post-MIR surgery A) Example tracing of the endocardium, B) relative displacement map with dotted pink line denoting ischemic zone,² C) 3-dimensional representation of displacement between cardiac cycles (parallel waveforms) commercial ELISA kit, Fig 7B) at 1 day post-MIR surgery.

Fig 7. Acute cardiac injury 1 day post-MIR A) Representative images of Evan's blue and triphenyl tetrazolium chloride (TTC) co-stained heart sections. Non-ischemic myocardium is indicated by the blue-stained, ischemic myocardium (Area-at-Risk; AAR) is indicated by the sum of pink and white/beige, and infarct size (IS) is indicated by white/beige areas B) Corresponding AAR, IS and cTnI comparisons.

Associated with modulation of macrophage population phenotypes at <u>3 days</u> post-MIR

No differences were observed in area-at-risk, fibrosis, relative abundances of viable leukocyte sub-populations counted by either flow cytometry or immunohistochemistry of CD68+ cells in whole left mid-ventricular sections; or any measure of echocardiographic LV function at 3 days post-MIR (all ANOVA p>0.05), data omitted for space. **However**, principal component analysis of bulk tissue RNA sequencing (RNAseq) data indicated deviations from saline and isotype controls with anti-CD14 treatment, further explored with single cell RNAseq (corrected for false discovery rate, p<0.01), shown below.



Fig 8. Single Cell RNA Sequencing at 3 days post-MIR A) Marker gene analysis for cell type (left) and macrophage (right) clustering, B) cell type proportions, C) tSNE cellular atlas plots of sorted macrophage subpopulations.

Significantly differentially expressed (DE) marker genes for each cell subset are shown (Fig 8A). Cell type proportions were similar between groups, with Mac 1 cells representing approximately 75% of total sorted cells in each group (Fig 8B and 8C).



Anti-CD14 treatment resulted in over 1,000 significantly DE genes in macrophage populations. The differences in expression of the top 200 DE genes (ranked by sig.) was predominantly in Mac1 cells (Fig 9). Functional enrichment analysis revealed significant upregulation of genes



The **primary endpoint** for each separately delivered study was pre-specified as infarct size (end-D1 study), immune cell phenotype by single cell RNA seq (D3), and LV ejection fraction (EF%) by ultra-high



for quality control of project primary endpoint ejection fraction (EF%) by echocardiography³

frequency echocardiography (D7), for relevant power calculations.

References

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Fig 9. Heatmaps indicating relative expression Top 200 significantly differentially expressed genes in Mac 1 cells (adjusted p<0.01).

associated with mRNA processing and chromosome/chromatin modification (omitted for space) and downregulation of large clusters of genes associated with inflammatory responses to damage and other stimuli; immune cell activation, differentiation and migration/chemotaxis; and responses to/production of proinflammatory cytokines (Fig 10).

and responses to/production of pro- Fig 10. Functional enrichment analysis indicating only inflammatory cytokines (Fig 10).

Conclusions & Future Translation

Anti-CD14 treatment given at reperfusion:

- significantly protects against progressive left ventricular injury and dysfunction 7 days post-MIR, independently of acute infarct size
- is associated with modulation of macrophage phenotype at 3 days and subsequent macrophage abundance at 7 days post-MIR

These data support that CD14 antagonism with novel antibody therapies like Atibuclimab may find clinical utility in the protection against immunological drivers of HFrEF in acute MI patients, particularly for those with poor LV function following PCI.

This work formed the basis of successful application for IND and Study May Proceed Letter from the FDA for Phase I Clinical Trials