

# Reverse cardiac remodelling: VICTORIA for inflammation and cardiac metabolism

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*‘Continuous improvement is better than delayed perfection’.*  
 Mark Twain.

Cardiac structure and function may change over time reflecting the balance between ongoing cardiomyocyte injury and the beneficial effects of disease-modifying therapies.<sup>1</sup> A decline in left ventricular ejection fraction (LVEF) and an increase in left ventricular (LV) volumes, known as adverse remodelling (AR), is associated with worse prognosis. Conversely, recovery from LV dysfunction and dilatation, called reverse remodelling (RR), is usually characterized by improved symptoms, better quality of life, and lower risk of hospitalization or death.<sup>1</sup>

Following disease trajectories over time is crucial, as RR has become increasingly common with advances in treatment for heart failure (HF) with reduced ejection fraction (HFrEF), leading to HF with improved ejection fraction (HFimpEF). Cardiac remodelling, defined based on imaging findings, is accompanied by changes in laboratory values reflecting the severity of cardiac dysfunction and its impact on other organs and tissues through various mechanisms (hypoperfusion, activation of neurohormonal axes, and pro-inflammatory pathways). The relationship between cardiac remodelling and changes in biomarkers remains poorly explored.<sup>1</sup>

Prior studies have reported biomarker changes associated with RR in HFrEF. In the PROVE-HF study, reductions in N-terminal pro-B-type natriuretic peptide (NT-proBNP) correlated with improvements in LVEF, LV end-systolic volume index (LVESVI), left atrial volume index, and E/e' at 12 months.<sup>2</sup> Similarly, in the GUIDE-IT trial, LV volume reductions and LVEF recovery were proportional to the decrease in NT-proBNP.<sup>3</sup> PROTECT also showed that NT-proBNP could serve as a non-invasive indicator of cardiac structure and function in HFrEF.<sup>4</sup> A biomarker-assist score

(ST2-R2 score), including soluble suppression of tumorigenesis-2, was developed to predict RR and mortality.<sup>5</sup>

In VICTORIA, a phase 3 randomized placebo-controlled trial, the oral soluble guanylate cyclase stimulator vericiguat reduced the primary composite outcome of HF hospitalization (HFH) or cardiovascular death in 5050 patients with LVEF <45% and a high risk of new HFH.<sup>6</sup> An echocardiographic substudy in 419 patients showed significant improvements in LV structure and function over 8 months in both vericiguat and placebo groups.<sup>7</sup>

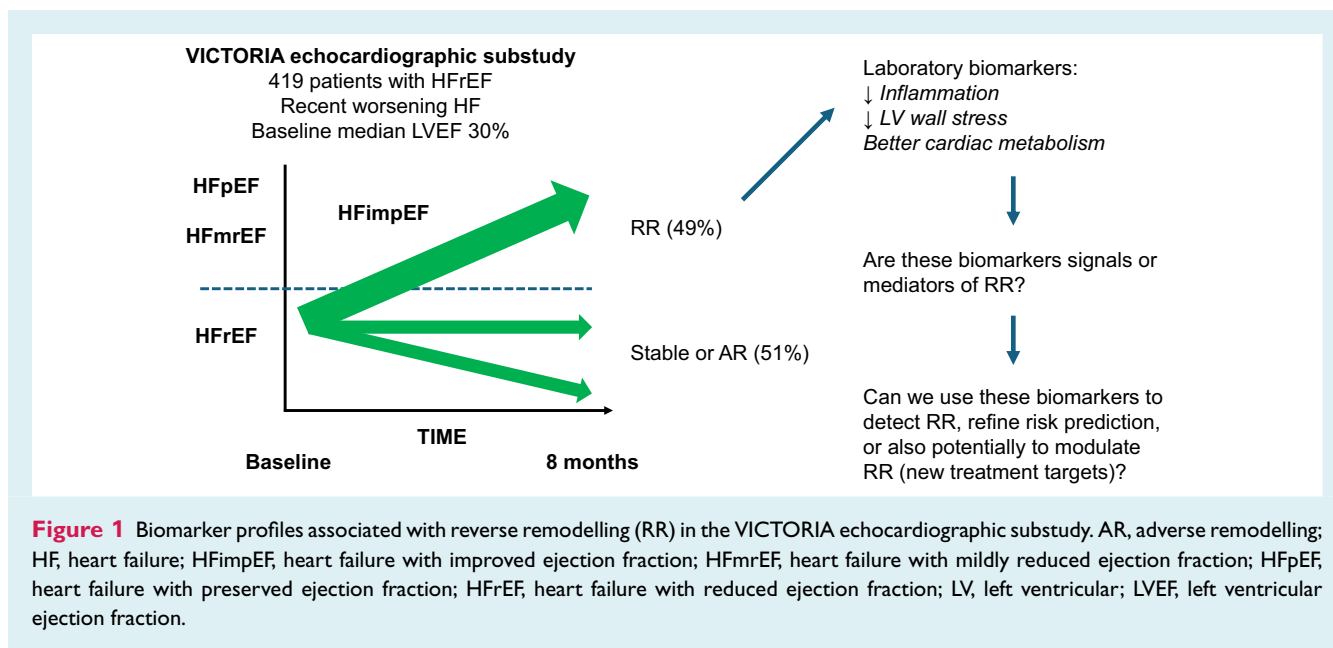
This current analysis of VICTORIA reports on biomarker profiles associated with RR in 419 patients with HFrEF over 8 months (Figure 1).<sup>8</sup> RR, defined as >5% absolute increase in LVEF or a >15% relative decrease in LVESVI, was seen in nearly half of patients (49%). The remainder (51%) showed no change or worsening of LVEF or LVESVI between baseline and 8 months. AR was defined as >5% absolute decrease in LVEF or a >15% relative increase in LVESVI. HFimpEF was defined as LVEF ≤40% at baseline with ≥10% point increase in LVEF and second LVEF measurement at 8 months >40%.<sup>8</sup> A total of 92 circulating protein biomarkers were collected at baseline and 8 months. Using weighted co-expression network analysis, clusters of highly correlated biomarkers were identified based on their temporal changes. The relationship between these clusters and remodelling was then assessed.

What are the key findings? First, all tested biomarkers at baseline did not predict future RR. Second, reductions in biomarkers associated with inflammation, LV wall stress, and cardiac metabolism – especially NT-proBNP, growth differentiation factor-15 (GDF-15), insulin-like growth factor-binding protein 1 (IGFBP1) and 7 (IGFBP7), chitinase-3-like protein 1 (CHI3L1), tumour necrosis factor ligand superfamily member 13B (TNFSF13B), osteoprotegerin (OPG), cathepsin D – were observed in patients who experienced RR, but not in those who did not, after adjustment for possible confounders. Third, no biomarker clusters or individual biomarkers were associated with AR or HFimpEF after correction for confounders.<sup>8</sup>

As for possible study limitations, less than 10% patients from VICTORIA were included in this analysis, and the cohort included

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just 27% of women, which may limit result generalizability. The assessment of remodelling was performed concurrently with changes in biomarkers, rather than using changes in biomarkers as predictors of future remodelling and final outcomes. Possible confounding factors when exploring cardiac remodelling include patient demographics, comorbidities, concurrent medications, genetic predispositions, baseline LVEF and LVESVI, measurement variability, and study design and methodology. Furthermore, only one follow-up time interval at the arbitrary time-point of 8 months was studied. Even the definition of RR was just one of many proposed criteria for RR.<sup>9</sup>

In this study, RR was observed on the background of HF therapy including 76% of patients on angiotensin-converting enzyme inhibitors or angiotensin receptor blockers, 16% on angiotensin receptor–neprilysin inhibitors, 95% on beta-blockers, 79% on mineralocorticoid receptor antagonists, and 12% having a biventricular pacemaker.<sup>8</sup> Sodium–glucose cotransporter 2 (SGLT2) inhibitors have also been shown to promote RR in HFrEF.<sup>10,11</sup> However, only 15 patients were on SGLT2 inhibitors, as the VICTORIA trial was conducted before SGLT2 inhibitors became a pillar of HFrEF treatment.

Future studies should examine biomarker profiles over time according to aetiology, HF duration, and rhythm (sinus rhythm vs. atrial fibrillation). For example, shorter HF duration and non-ischaeamic aetiology are predictors of RR, while genetic or familial cardiomyopathies are less likely to recover.<sup>12</sup> Beyond LVESVI and LVEF, future studies could assess other LV parameters (wall thickness, mass, strain, diastolic function) and other cardiac chambers (left atrium, right heart) and utilize cardiac magnetic resonance (e.g. the absence of late gadolinium enhancement predicts RR).<sup>12</sup>

The interplay between biomarkers and RR, from association to cause–effect relationships, is investigated through dedicated mechanistic studies, potentially identifying new risk predictors. If biomarkers not only signal but also contribute to cardiac

remodelling, they could serve as new treatment targets. For instance, the GARDEN-TIMI 74 (NCT05492500) trial is assessing ponesegromab, a monoclonal antibody directed against GDF-15, in patients with HFrEF and elevated baseline GDF-15.

If reduced inflammatory activation and improved cardiac metabolism partially mediate RR, anti-inflammatory therapies and treatments that enhance cardiac metabolism become highly interesting for inducing RR in HFrEF. In the DREAM-HF trial, mesenchymal precursor cell therapy, which has anti-inflammatory properties, increased LVEF at 12 months compared to sham controls, especially in patients with inflammation.<sup>13</sup> Furthermore, the SGLT2 inhibitor empagliflozin has been shown to promote RR in HFrEF,<sup>10,11</sup> as well as to reduce inflammatory biomarkers.<sup>14</sup>

Changes in biomarkers closely linked to the remodelling process and predictive of outcomes could serve as surrogate endpoints in HFrEF clinical trials.<sup>15</sup> Serial biomarker monitoring could guide the timing and frequency of clinical and imaging assessments. Several key questions remain: How many biomarkers are needed for optimal prediction of RR? Does a biomarker-guided strategy improve patient outcomes? Will such a strategy be cost-effective? How feasible is assessing biomarker profiles (single biomarkers, multiple biomarkers, biomarker panels, proteomic, and other multi-omics data) in clinical practice and research?

Overall, this analysis of VICTORIA provides novel insights by demonstrating that RR is associated with reductions in biomarkers associated with inflammation, wall stress, and dysregulated cardiac metabolism. These findings suggest potential targets for new HF therapies. What remains uncertain is whether these are signals or mediators of RR, as these associations may not imply causation. Future studies should define whether we can use these biomarkers to detect RR, refine risk prediction, or also potentially to modulate RR.

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