

# Insulin-like growth factor binding protein-7 concentrations in chronic heart failure: Results from the EMPEROR programme

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1	Insulin-like growth factor binding protein-7 concentrations in Chronic Heart Failure:
2	<b>Results from the EMPEROR Program</b>
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#### 41 Abstract

Aims: Insulin-like growth factor binding protein-7 (IGFBP7) is a biomarker of tissue
senescence with a role in cardio-renal pathophysiology. The role of IGFBP7 as a prognostic
biomarker across the full ejection fraction (EF) spectrum of heart failure (HF) remains less
well understood. We examined associations between IGFBP7 and risk of cardio-renal
outcomes regardless of EF and the effect of empagliflozin treatment on IGFBP7
concentrations among individuals with HF.

48 Methods and Results: IGFBP7 was measured in 1125 study participants from the

49 EMPEROR-Reduced and EMPEROR-Preserved Trials. Cox regression was used to study

associations with outcomes. Study participants with IGFBP7 levels in the highest tertile had a

51 higher-risk clinical profile. In Cox proportional hazards models adjusted for clinical

52 variables, N-terminal pro-B type natriuretic peptide and high-sensitivity cardiac troponin T,

53 baseline IGFBP7 values in the highest tertile predicted an increased risk of HF hospitalization

54 or CV death (HR: 2.00 95% CI 1.28-3.10, p=0.002, p for trend <0.001) and higher risk of

<sup>55</sup> renal composite endpoint (HR: 4.66, 95% CI: 1.61-13.53, p=0.005, p for trend = 0.001),

56 regardless of EF. Empagliflozin reduced risk for CV death/HF hospitalization irrespective of

57 baseline IGFBP7 (p-trend across IGFBP7 tertiles=0.26). Empagliflozin treatment was not

associated with meaningful change in IGFBP7 at 12 or 52 weeks.

59 Conclusion: Across the entire LVEF spectrum in the EMPEROR program, concentrations of
60 the senescence associated biomarker IGBP7 were associated with higher risk clinical status
61 and predicted adverse cardio-renal outcomes even in models adjusted for conventional
62 biomarkers. Empagliflozin did not significantly affect IGFBP7 levels over time.

Trial Registration: EMPEROR-Reduced (NCT03057977), EMPEROR-Preserved
 (NCT03057951)

65 Keywords: heart failure, IGFBP7, biomarker, empagliflozin, kidney function,

#### 66 Introduction

The progression of heart failure (HF) includes accelerated aging of the myocardium. Such 67 myocardial senescence results from permanent cell cycle arrest combined with loss of tissue 68 renewal. Senescent cells secrete a pool of molecules known as the senescence-associated 69 secretome that exert autocrine and paracrine to promote the aging process. IGFBP7 (insulin-70 like growth factor-binding protein-7) is a prominent member of the senescence-associated 71 72 secretome; among the effects of IGFBP7 is to augment cell cycle arrest, while enhancing oxidative stress, impairing DNA repair and promoting fibrosis. It does so in part by 73 74 interfering with the adaptive actions of foxhead box O3a (FOXO3a) to reduce cellular stress and induce autophagy(1-3). Although originally considered to be a member of the IGFBP 75 76 family, IGFBP7 differs from other members of the family in that it binds only weakly to 77 insulin-like growth factors(4-8).

Notably, studies have shown higher blood concentrations of IGFBP7 among patients 78 with acute or chronic HF syndromes are associated with abnormalities of diastolic filling and 79 increases in left atrial volume(9, 10). Furthermore, high concentrations of IGFBP7 may be 80 additive to those of N-terminal pro-B type natriuretic peptide (NT-proBNP) for diagnostic 81 82 accuracy in HF and in predicting an adverse prognosis in acute and chronic HF(11, 12). However, the true additive value of IGFBP7 for prognostication relative to conventional 83 84 biomarkers such as NT-proBNP or high sensitivity cardiac troponin (hs-cTn) has not been 85 explored across the entire range of left ventricular ejection fraction (LVEF). Furthermore, the effect of empagliflozin on concentrations of IGFBP7 is not described. We therefore sought to 86 more thoroughly evaluate this question among study participants in the Empagliflozin 87 88 Outcome Trial in Patients with Chronic Heart Failure (EMPEROR) program.

89 Methods

90 *Study sample* 

The design and results of EMPEROR-Reduced and EMPEROR-Preserved trials have been published(13, 14). This study consists of 9718 individuals who were initially enrolled in the EMPEROR-Reduced (n = 3730) and EMPEROR-Preserved (n = 5988) trials. In brief, the trials included individuals with class II-IV HF and elevated levels of natriuretic peptides across a broad spectrum of LVEF. Written informed consent was obtained from all patients and the study protocols were approved by the ethics committees of all participating institutions.

Apart from their usual treatment for HF, patients who met the eligibility criteria were 98 randomly assigned in a double-blind manner, with a 1:1 ratio, to receive either a placebo or 99 empagliflozin at a 10 mg dose per day. During study visits, outcomes related to HF were 100 101 assessed. In both trials, the primary outcome was a combination of cardiovascular death or hospitalization due to HF. The primary renal outcome was the combined risk of an estimated 102 glomerular filtration rate (eGFR) decrease of at least 40% over time, an eGFR decline to <10 103 or <15 ml/min/1.73m<sup>2</sup> (for patients with baseline eGFR < or  $\ge$  30 ml/min/1.73m<sup>2</sup>, 104 respectively), chronic dialysis, or renal replacement therapy. 105

Participation in the trials was not contingent on submitting samples of plasma, serum, 106 DNA, or urine for biobanking. Bio-samples were only obtained once a separate informed 107 consent was provided in compliance with local ethical and regulatory regulations. The 108 concepts and technique of sample collection for biobanking in the EMPEROR trials and the 109 list of nations that took part in the biobanking for biomarker sub-study have been published 110 111 elsewhere (15). Figure 1 details the study flow for the present analysis; for the purposes of this analysis, 594 study participants were included from EMPEROR-Reduced (LVEF <40%), 112 while 531 study participants were included from EMPEROR-Preserved (LVEF  $\geq$ 40%). 113

#### 114 Biomarker measurement

115 Biomarkers were measured in serum samples in the placebo and empagliflozin arms at

- 116 baseline (N= 561, 564), 12 weeks (N=558, 557) and 52 weeks (N=355, 358). The
- measurements of IGFBP7, as well as hs-cTnT, carbohydrate antigen 125 (CA-125), growth
- differentiation factor-15 (GDF-15), angiopoietin 2 (ANGPT-2), fatty acid-binding protein-3
- 119 (FABP-3) and bone morphogenetic protein 10 (BMP-10), were performed within the
- 120 Biomarker Research Agreement of Boehringer Ingelheim, the sponsor of the EMPEROR
- trials and Roche Diagnostics International Ltd. These measurements were made using an
- 122 electrochemiluminescence immunoassay (Elecsys®, Roche Diagnostics). NT-proBNP was
- 123 previously measured using similar methodology(16). Additionally, plasma samples from the
- same study participants were included in a plasma proteomics study using the Olink®
- 125 Explore 1536 platform previously reported(17).

#### 126 *Statistical analysis*

Mean (standard deviation), median (interquartile range), and count (frequency) were used to 127 describe the distribution of continuous and categorical variables as appropriate. Cox 128 129 proportional hazard regression models and negative binomial models were used to analyze the associations between baseline IGFBP7 and major HF outcomes, as appropriate. 130 Multivariable models included age, sex, geographical region, diabetes, LVEF, eGFR, and trial 131 (EMPEROR-Reduced or EMPEROR-Preserved) as covariates, and were additionally 132 adjusted for log-transformed baseline hs-cTnT and NT-proBNP, the most important 133 biomarker predictors of the primary outcome in the EMPEROR studies(18-21). Additional 134 135 adjustment was performed on GDF-15, another important prognostic marker in the EMPEROR-Program (22). Week 12 landmark analysis were performed to study the 136 associations between IGFBP7 changes and the subsequent occurrence of the primary 137

138 outcome. Heterogeneity of treatment effect on major HF outcomes across the groups defined by the tertiles of GDF-15 was investigated by fitting Cox regression and negative binomial 139 models, as appropriate, adjusted by the prespecified covariates except for hs-cTnT and NT-140 proBNP and including a IGFBP7-group by treatment interaction. With the mentioned 141 baseline covariates and baseline IGFBP7 levels considered, changes in IGFBP7 levels over 142 time were examined using mixed models for repeated measurements adjusted with same 143 covariates as multivariable models, excluding hs-cTnT and NT-proBNP. These models 144 compared the change in plasma concentrations for each protein from baseline to weeks 12 145 146 and 52 between treatment groups (empagliflozin vs. placebo). Correlation between IGFBP7 and proteins measured by Olink® Explore 1536 platform and the remaining proteins 147 measured by Roche Diagnostics were assessed by calculating Spearman correlation 148 149 coefficients. Statistical analyses were performed using the SAS software, version 9.4 (SAS 150 Institute, Inc.; Cary, NC). All tests were two-sided, and p-values < 0.05 were considered statistically significant. 151

152 **Results** 

Of 1125 people with HF included in this study, 361 (32%) were female. The mean (standard deviation) age of study participants was 70±10 years. Most study participants were White (n=948, 84%) and reside in Europe (n=632, 56%). Baseline characteristics of the study population stratified by thirds of IGFBP7 levels are detailed in **Table 1**. Study participants with the highest IGFBP7 concentrations were older, more likely to be male, had a higher prevalence of diabetes mellitus, atrial fibrillation, valvular heart disease, lower eGFR, and higher albumin/creatinine ratio, NT-proBNP and hs-cTnT compared to other tertiles.

The results of the Cox regression analysis are detailed in Figure 2. In multivariable
 model, considering study participants with IGFBP7 concentrations in tertile 1 as a reference,

162 participants with IGFBP7 concentrations in the highest tertile were at significantly higher risk for the primary endpoint of the EMPEROR Trial, HF hospitalization/CV death (hazard ratio 163 [HR]: 2.00 [95% confidence interval (CI): 1.28-3.10], p-value <0.001). Figure 3 depicts the 164 cumulative incidence function of the primary endpoint of the EMPEROR Trials for each 165 IGFBP7 tertile. Patients with IGFBP7 concentrations in the highest tertile had a significantly 166 higher rate of HF hospitalization/CV death compared to others (Log rank p-value <0.001). 167 Associations remained similar with additional adjustment on baseline GDF-15 levels: 168 EMPEROR-Pooled tertile 3 vs tertile 1 primary outcome HR 1.66, 95%CI 1.03-2.67, P 169 =0.03. 170

Associations for spline-transformed continuous IGFBP7 levels are presented in the
Supplementary Figure 1.

In week 12 landmark analysis, IGFBP7 changes (from baseline to week 12) were
strongly associated with the primary outcome: adjusted HR 2.67 for a fold-change in IGFBP7
from baseline to week 12 (HR 4.12, 95% CI 2.21-7.69 in the log-scale, P <0.0001), adjusting</li>
on the "extended model" plus baseline IGFBP7 levels.

As also shown in **Figure 2**, higher tertiles of IGFBP7 were associated with an array of other cardio-renal events, including significant adjusted risk for first HF event (HR 2.20 [95% CI 1.19-3.82]; p = 0.002), total HF events (adjusted event rate ratio: 2.13 [95% CI: 1.19-3.82], p-value = 0.004), CV death (HR: 1.84 [95% CI: 1.00-3.41]; p = 0.04) as well as sustained eGFR reduction  $\geq$ 40% or ESRD (HR: 4.66 [95% CI: 1.61-13.53], p-value = 0.001). **Supplemental Table 1 and 2** show the association between IGFBP-7 tertiles and

clinical outcomes among individuals with HFrEF and HFpEF respectively. No significant interaction was found between LVEF levels (<40% or  $\geq40\%$  according to populations of

EMPEROR-Reduced and EMPEROR-preserved trials) and IGFBP-7 concentrations for
prediction of clinical outcomes (p for interactions by study >0.1).

Table 2 details effect of empagliflozin on endpoints in the EMPEROR Trials as a 187 function of baseline IGFBP7. Two observations may be made: firstly, higher concentrations 188 of IGFBP7 were routinely associated with higher event rates regardless of treatment 189 assignment; second, in most cases, no interaction of treatment effect of empagliflozin by 190 baseline IGFBP7 was detected, however effect of empagliflozin to improve time to first HF 191 hospitalization may have been attenuated among those in the lowest IGFBP7 tertile. In Table 192 3, eGFR slope across IGFBP7 tertiles is detailed as is treatment effect of empagliflozin. 193 Similar to clinical events, decay of renal function (reflected in a more negative eGFR slope) 194 was generally worse in those with higher IGFBP7 concentrations. Empagliflozin had 195 196 generally similar effects on eGFR slope across tertiles of IGFBP7.

Supplemental Table 3 displays the top correlations between baseline IGFBP7 and
proteins measured through Olink proteomics or other automated immunoassays. At baseline,
most correlations were modest. The highest correlations of IGFBP7 (Spearman correlation>
0.60) were found with bone morphogenic protein-10, fatty acid binding protein-3, and growth
differentiation factor-15.

Table 4 shows the concentration of IGFBP7 among empagliflozin compared to
placebo groups at baseline and follow-up visits. Treatment with empagliflozin was associated
with a 3% greater reduction of IGFBP7 level versus placebo to week 12 (adjusted geometric
mean ratio: 0.97; 95% confidence interval [CI], 0.94-1.00, p=0.05) but not to week 52
(adjusted geometric mean ratio: 1.02, 95% CI 0.98-1.06; p=0.35) (Figure 4).
Discussion

208 Prior studies have suggested IGFBP7 measurement may inform cardio-renal risk in both acute and chronic HF states(12, 23, 24) but the value of the marker across the wide range of 209 LVEF remained unclear, the meaning of IGFBP7 to prognosticate cardiac and renal outcomes 210 even in presence of other biomarkers required further analyses, and the impact of SGLT2 211 inhibition on IGFBP7 was uncertain. In this study from the EMPEROR program, we showed 212 that elevated IGFBP7 levels were associated with more advanced age, a higher prevalence of 213 214 dysglycemia, and a greater burden of cardiac and kidney dysfunction. Higher concentrations of IGFBP7 or change in IGFBP7 were predictive of increased risk of HF hospitalization, 215 216 cardiovascular mortality, and progression of kidney disease even when adjusted for other relevant biomarkers, including NT-proBNP and hs-cTnT, the strongest overall prognostic 217 measures in the EMPEROR program (20, 21); similar findings were present when further 218 219 adjusting for GDF-15. IGFBP7 was a robust predictor of risk across the wide LVEF range in 220 the EMPEROR Trials and in most cases no interaction by baseline LVEF was present regarding both the ability of IGFBP7 to prognosticate outcomes or predict effect of 221 empagliflozin although effect of empagliflozin to improve time to first HF hospitalization 222 may have been attenuated among those in the lowest IGFBP7 tertile. Concentrations of 223 224 IGFBP7 were not meaningfully lowered by treatment with empagliflozin. In aggregate, these results demonstrate that IGFBP7 is an important cardiorenal risk marker across the spectrum 225 226 of LVEF in persons with HF and add considerably to available knowledge regarding the 227 biomarker in those treated with SGLT2 inhibitors(25).

A potential role for IGFBP7 in HF was originally identified in a proteomics screen in an animal model of cardiomyopathy(26), and numerous studies have since linked concentrations of IGFBP7 to worse systolic and diastolic function in patients with HF(12, 23, 24). As part of the senescence-associated secretome, IGFBP7 has a broad range of effects including induction of cell cycle arrest, interference with DNA repair and detoxification of

233 reactive oxygen species, and promotion of fibrosis, potentially mediated by an action to suppress FOXO3a signaling(27, 28). Gene knock-out of IGFBP7 in HF models is associated 234 with a reduction in proinflammatory and profibrotic pathways, resulting in preservation of 235 236 cardiac function(9). Furthermore, RNA-mediated silencing of IGFBP7 or neutralization with monoclonal antibodies blocked the actions of IGFBP7 on FOXO3a signaling and mitigated 237 pressure-overload induced HF(29). A similar role has been identified for IGFBP7 in the 238 239 kidney, where it acts as a marker of renal stress and a potential mediator of the progression of chronic kidney disease(30). These results implicate IGFBP7 as a potential target for future 240 241 therapies to slow the progression of HF and nephropathy.

Notably, baseline IGFBP7 concentrations or changes in IGFBP7 were strongly 242 prognostic for major cardio-renal outcomes in EMPEROR, adding strong prognostic value 243 244 across the entire LVEF spectrum. The fact that IGFBP7 remained a robust predictor of outcomes in the trial even when adjusted for NT-proBNP and hs-cTnT results is noteworthy: 245 these two biomarkers have been found to be dominant for predicting outcomes in the trial 246 program, almost routinely superseding other biomarkers, so the finding of prognostic 247 meaning with IGFBP7 in this study is quite meaningful. Given the primacy of NT-248 249 proBNP(16) and hs-cTnT(19) for prognosis in the EMPEROR program(21), the results of 250 this study further solidify the robustness of IGFBP7 for prognostic assessment. We extend the 251 results to also show IGFBP7 remains strongly prognostic when further adjusted for GDF-15 252 concentrations in addition to NT-proBNP and hs-cTnT. As few biomarkers substantially add to information from NT-proBNP and hs-cTnT, the results are important and suggest that 253 254 IGFBP7 adds orthogonal biologic information to measures of myocardial wall stress and 255 cardiomyocyte necrosis. It is reasonable to consider the paradigm of wall stress (NT-proBNP) 256 triggering a cascade of senescence signaling (IGFBP7), culminating in necrosis (hs-cTnT).

Further translational research is needed to understand links between IGFBP7, relatedbiomarkers, and clinical outcomes in HF.

In EMPEROR, across nearly all outcomes evaluated, treatment with empagliflozin 259 was associated with generally similar event reductions regardless of baseline IGFBP7 260 concentration; effect of empagliflozin on time to first HF hospitalization might have been 261 262 attenuated in the lowest IGFBP7 tertile. Taken in aggregate, these results suggest that IGFBP7 measurement should not be used to decide on whether to utilize an SGLT2 inhibitor 263 in HF, although those with highest IGFBP7 values had large absolute risk reduction. In 264 contrast to treatment of HF with sildenafil(31), sacubitril/valsartan(12) or heart 265 transplantation(32), where such treatments may lower concentrations of IGFBP7, in this 266 study, treatment with empagliflozin did not meaningfully reduce concentrations of the 267 268 biomarker during the duration of study, similar to results from analyses of canagliflozin in persons with type 2 diabetes (16, 33). This implies factors leading to rise of IGFBP7 may be 269 independent of effects of SGLT2 inhibition. 270

It is noteworthy that despite the multiple cardioprotective effects of SGLT2 inhibitors 271 (34), concentrations of the prognostic IGFBP7 were reduced by 3% at week 12, but not at 272 week 52 of the study. These results are consistent with prior evidence suggesting 273 canagliflozin did not reduce IGFBP7 concentrations in those with diabetes and CV risk 274 treated for a year (30). In contrast, another study of individuals with diabetic kidney disease 275 suggested canagliflozin reduced IGFBP7 by approximately 2% over a similar time course of 276 treatment (35). More in-depth understanding regarding how biomarkers like IGFBP7 might 277 reflect benefit of therapeutics including SGLT2 inhibitors is needed. 278

A particular strength of the EMPEROR Trial is the depth of biomarker data available
for cross-correlation. Our study reveals correlations between the level of IGFBP7 and certain

281 plasma proteins that are linked to key processes in cardiovascular disease. Whereas most of the correlations were modest in nature, several proteins with links with myocardial disease 282 processes, including cardiac remodeling showed at least moderate correlation with 283 284 IGFBP7.(36) (37). The top correlations with IGFBP7 (Rho >0.6) are Bone Morphogenic Protein (BMP)-10, Fatty Acid Binding Protein (FABP)-3, and GDF-15. In experimental 285 models BMP10 stimulated cardiomyocyte proliferation and repaired cardiac function after 286 287 heart injury(37), BMP10 has also been described as a marker of adverse atrial remodelling (38). FABP3, is a small protein found intracellularly within cardiac or skeletal muscles that 288 289 become released into circulation and excreted into the urinary system following injury to skeletal muscle, which may occur secondary to PAD-related tissue ischemia. FABP3 has been 290 implicated in metabolic syndrome and mitochondrial dysfunction (39, 40). The implication of 291 292 GDF-15 in the EMPEROR program has been thoroughly discussed (22), however the 293 correlation with IGFBP7 and additive value for prognosis of these two biomarkers is noteworthy. The investigation into the relationship between IGFBP7 and myocardial disease 294 processes requires additional research. 295

There are several study limitations that should be considered when interpreting our 296 297 findings. Prior studies have implicated IGFBP7 as a cardio-renal prognostic biomarker; this 298 study considerably extends the understanding of this biomarker in numerous ways: first, we 299 evaluate its prognostic meaning across the wide EF range in the EMPEROR program, 300 thoroughly evaluating it relative to both NT-proBNP and hs-cTnT; second, we evaluated impact of empagliflozin on IGFBP7 concentrations; lastly, we provide cross-correlations with 301 302 other markers identified in a proteomics screen. A second limitation is that IGFBP7 is not 303 cardiac specific and may be expressed in multiple tissues. For example, recent studies have 304 implicated expression of the peptide in renal tubular epithelium and concentrations of the biomarker are strongly prognostic for renal outcomes in chronic diabetes mellitus(16, 33). 305

Our study involved the measurement of IGFBP7 levels in blood. The concentration of
IGFBP7 in the heart may not be directly related to its level in the bloodstream. Furthermore,
most of the study population were white males. Future research might explore the comparison
of IGFBP7 levels among various races and ethnicities. This post hoc analysis included
analyses without adjusting for multiple comparisons. Finally, the study participants were only
monitored for 3 years. As a result, additional research is needed to determine the long-term
effects of IGFBP7 levels on HF prognosis.

In conclusion, this study provides evidence regarding the importance of IGFBP7 313 314 among patients with HF. More elevated baseline IGFBP7 concentrations were associated with a higher risk phenotype and were correlated with key biomarkers associated with myocardial 315 316 processes pivotal in HF. Elevated values of IGFBP7 were linked to heightened adjusted risk 317 of HF hospitalization, diminished survival, and higher risk of kidney failure across the entire LVEF range in HF. IGFBP7 remained a robust predictor of risk regardless of adjustment with 318 NT-proBNP or hs-cTnT. Lastly, concentrations of IGFBP7 were not meaningfully affected by 319 therapy with empagliflozin. Our results, demonstrated across the wide LVEF range in the 320 EMPEROR program, further emphasize IGFBP7 as an important biomarker in HF. 321

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### 474 Figure legends

475 **Figure 1:** Study flowchart

476 **Figure 2:** Cox regression and negative binomial regression results by tertile of IGFBP7 at

477 baseline. Multivariable models adjusted for age, sex, baseline eGFR, baseline LVEF, log

478 (NT-proBNP), log (HS Troponin T), region, diabetes, treatment arm (empagliflozin vs.

479 placebo), study (EMPEROR-Reduced vs. EMPEROR-Preserved). Results for total HHF are

480 reported as adjusted rate ratios from a negative binomial model with the addition of log(days

481 at risk) as an offset. Abbreviations: CV: cardiovascular, HF: heart failure, ESRD: end-stage

482 renal disease, HHF: hospitalization for heart failure, IGFBP7: insulin growth factor binding

- 483 protein 7.
- 484 Figure 3: Cumulative incidence function of HF hospitalization/CV death according to
  485 IGFBP7 Tertile group. T denotes: tertile.
- 486 **Figure 4:** IGFBP7 concentration across study visits stratified by treatment arm

487	Table 1. Base	line characteristic	es of study	population	stratified by	tertile of IGFBP7
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488 concentration.

	Tertile 1 N=375	Tertile 2 N=375	Tertile 3 N=375
Age, year	67.1 (10.2)	71.2 (9.1)	71.7 (10.1)
Sex			× /
Male	239 (63.7)	262 (69.9)	263 (70.1)
Female	136 (36.3)	113 (30.1)	112 (29.9)
Race			
White	310 (82.7)	318 (84.8)	320 (85.3)
Black	9 (2.4)	9 (2.4)	7 (1.9)
Asian	28 (7.5)	35 (9.3)	33 (8.8)
Other	24 (6.4)	10 (2.7)	11 (2.9)
Ethnicity	× /		
Not Hispanic / Latino	285 (76.0)	296 (78.9)	273 (72.8)
Hispanic / Latino	75 (20.0)	68 (18.1)	95 (25.3)
Region			,
North America	46 (12.3)	39 (10.4)	71 (18.9)
Latin America	72 (19.2)	62 (16.5)	83 (22.1)
Europe	221 (58.9)	233 (62.1)	178 (47.5)
Asia	25 (67)	30 (8 0)	31 (8 3)
Other	11(2.9)	11 (2.9)	12(32)
Smoking status	11 (2.9)	11 (2.7)	12 (3.2)
Never smoked	167 (44 5)	165(440)	166 (44 3)
Ex–smoker	154 (41.1)	168 (44 8)	179(47.7)
Currently smokes	43(11.5)	33 (8 8)	26 (6 9)
Drinking alcohol	15 (11.5)	33 (0.0)	20 (0.7)
Does not drink alcohol	218 (58 1)	228 (60.8)	223 (59 5)
Drinks alcohol	1/3 (38.1)	137 (36 5)	144 (38.4)
Changes in NVHA during 3m	145 (50.1)	157 (50.5)	144 (30.4)
Improved	<i>A</i> <b>0</b> (13 1)	13 (11 5)	<i>A</i> 3 (11 5)
No change	(13.1) 310 (82 7)	$\frac{43}{11.3}$	-43(11.3) 205 (78 7)
Unstable	2(0.5)	0	293(10.1)
Worsened	2(0.3) 14(3.7)	18(48)	0(1.0) 21 (8 2)
Time since HF diagnosis y	14(3.7)	10(4.0)	51(6.3)
UE hospitalization within 1	4.0 (3.2)	4.8 (3.2)	5.5 (5.9)
vear	108 (28.8)	92 (24 5)	108 (28.8)
Diabetes status	100 (20.0)	)2(21.5)	100 (20.0)
Diabetic	164 (43 7)	182 (48 5)	195 (52 0)
Non-diabetic	211 (56 3)	102 (10.5)	199(32.0) 180(48.0)
LVEF. %	395(14.8)	407(148)	40.2(15.4)
LVEF	57.5 (14.0)	40.7 (14.0)	40.2 (13.4)
<50 %	263 (70.1)	252 (67.2)	257 (68 5)
50  to  < 60%	53(141)	65 (17 3)	67 (16 5)
> 60%	59(15.7)	58 (15 5)	56(14.9)
SRP mmHg	1282(13.7)	128.2(15.3)	10(14.7)
DRP mmHg	120.2(10.2) 75 0 (10 7)	120.2(13.0) 7/6(10.6)	127.3(13.4) 71.1(10.4)
Heart rate hom	(10.7)	74.0(10.0) 71 2(12 2)	74.4 (10.4) 70 7 (11 2)
<b>BMI</b> $kg/m^2$	20.3(11.0)	(1.2(12.3))	10.1 (11.3) 20.60 (5.77)
<b>aCFP</b> mI $/min/1$ 72 m <sup>2</sup>	27.33 (3.47) 70 0 (19 7)	27.00 (J. / J)	27.00(3.77)
$\geq 60\%$ SBP, mmHg DBP, mmHg Heart rate, bpm BMI, kg/m <sup>2</sup> eGFR, mL/min/1.73 m <sup>2</sup>	59 (11.1) 59 (15.7) 128.2 (16.2) 75.0 (10.7) 69.5 (11.8) 29.35 (5.47) 70.9 (18.7)	58 (15.5) 128.2 (15.8) 74.6 (10.6) 71.2 (12.3) 29.06 (5.73) 60.1 (18.8)	56 (14.9) 127.3 (15. 74.4 (10.4 70.7 (11.3 29.60 (5.7 50.3 (18.4

Albumin/Cr ratio*, mg/g	10.61 (5.30-30.00)	15.03 (7.07-48.62)	35.18 (9.72-131.00)
Atrial fibrillation/flutter	129 (34.4)	189 (50.4)	231 (61.6)
Atrial fibrillation	122 (32.5)	180 (48.0)	226 (60.3)
Hypertension	302 (80.5)	307 (81.9)	323 (86.1)
Hypercholesterolemia	257 (68.5)	252 (67.2)	274 (73.1)
ACS within 6 months	19 (5.1)	22 (5.9)	18 (4.8)
MI	146 (38.9)	130 (34.7)	140 (37.3)
CAD	152 (40.5)	150 (40.0)	159 (42.4)
CABG or PCI	146 (38.9)	134 (35.7)	151 (40.3)
Valvular heart disease	57 (15.2)	53 (14.1)	73 (19.5)
Invasive EP procedure	37 (9.9)	26 (6.9)	33 (8.8)
Left Bundle Branch Block	49 (13.1)	45 (12.0)	38 (10.1)
Cardiac Biomarker			
NT-proBNP, pg/mL	1048 (582-1746)	1268 (725-2239)	1968 (1050-3410)
hs-cTnT, ng/L	15.50 (10.80-22.40)	17.80 (12.50-27.35)	25.95 (17.50-37.50)

489 Abbreviations: ACS: acute coronary syndrome, BMI: body mass index, CABG: coronary

490 artery bypass graft, CAD: coronary artery disease, DBP: diastolic blood pressure, eGFR:

491 estimated glomerular filtration rate, EP electrophysiology, HF: heart failure, hs-cTnT: high

492 sensitive cardiac troponin T, LVEF: left ventricular ejection fraction, MI: myocardial

493 infarction, NYHA: New York heart association, NT-proBNP: N terminal pro b type

494 natriuretic peptides, PCI: percutaneous coronary intervention, SBP: systolic blood pressure.

**Table 2:** Effect of empagliflozin on endpoints in the EMPEROR Trials as a function of baseline IGFBP7. In most cases, no interaction by

 baseline IGFBP7 was detected, however effect of empagliflozin to improve time to first HF hospitalization may have been attenuated among

 those in the lowest IGFBP7 tertile.

	Placebo incidence rate (Events per 100 patient years; 95% CI)	Empagliflozin 10 mg incidence rate (Events per 100 patient years; 95% CI)	Hazard ratio (95% CI)	Interaction trend p-value
CV death/HF hospitalization				
Tertile 1	5.67 (3.36-8.57)	6.65 (4.06-9.87)	1.15 (0.61-2.19)	0.26
Tertile 2	8.81 (5.76-12.51)	5.16 (3.00-7.88)	0.58 (0.31-1.08)	0.20
Tertile 3	19.51 (14.62-25.10)	14.67 (10.58-19.43)	0.69 (0.45-1.04)	
First HF hospitalization				
Tertile 1	2.52 (1.09-4.54)	4.99 (2.79-7.81)	1.95 (0.83-4.62)	0.07
Tertile 2	6.10 (3.62-9.23)	3.64 (1.88-5.97)	0.57 (0.27-1.19)	0.07
Tertile 3	15.09 (10.83-20.05)	11.53 (7.94-15.78)	0.68 (0.42-1.08)	
CV death				
Tertile 1	3.73 (1.93-6.12)	2.86 (1.31-5.00)	0.77 (0.32-1.83)	0.76
Tertile 2	4.16 (2.21-6.70)	2.04 (0.82-3.81)	0.55 (0.22-1.38)	0.70
Tertile 3	7.06 (4.43-10.31)	6.09 (3.67-9.12)	0.84 (0.45-1.56)	
Total HF hospitalizations*				
Tertile 1	3.73	7.03	1.78 (0.75, 4.22)	0.17
Tertile 2	8.95	5.57	0.68 (0.31, 1.50)	0.17
Tertile 3	24.16	18.62	0.80 (0.43, 1.47)	
First intensification of diuretic				
Tertile 1	8.76 (5.72-12.44)	7.50 (4.70-10.94)	0.79 (0.44-1.40)	0.51
Tertile 2	12.38 (8.58-16.88)	7.96 (5.15-11.37)	0.63 (0.38-1.06)	0.51
Tertile 3	18.32 (13.55-23.79)	12.46 (8.68-16.92)	0.61 (0.39-0.95)	

CV denotes: cardiovascular; HF denotes: heart failure; CI denotes: confidence intervals; hazard ratios based on Cox regression models and \*adjusted event rate ratio based on negative binomial regression with terms for age, baseline eGFR (CKD-EPI), baseline LVEF, region, diabetes at baseline, sex, study, treatment assignment, tertiles of IGFBP7 at baseline and treatment by tertiles of IGFBP7 at baseline interaction.

Additionally, the negative binomial regression model included log(days at risk) as an offset. Too few events were present to analyze sustained eGFR reduction  $\geq$ 40% or ESRD in this analysis.

**Table 3:** Adjusted eGFR change slope from baseline by tertiles of IGFBP7 at baseline in A) all study participants and B) by treatment assignment.

#### A)

	β	95 % CI	p-value
Lowest tertile slope* [/year]	-1.771	(–2.5450.997)	<0.001
Middle tertile slope* [/year]	-1.044	(-1.7840.303)	0.006
Highest tertile slope* [/year]	-2.179	(-2.9771.382)	<0.001
Middle vs Low	0.727	(-0.346 – 1.800)	0.18
High vs Low	-0.409	(-1.567 – 0.750)	0.49

B)

Category	Placebo slope (/year)	Empagliflozin 10 mg slope (/year)	Treatment by time interaction p-value	Subgroup interaction trend p-value
Tertile 1	-2.551 (-3.5901.512)	-1.609 (-2.6720.545)	0.21	
Tertile 2	-2.462 (03.5141.411)	-0.036 (-1.037 – 0.966)	0.001	0.93
Tertile 3	-2.152 (-3.2331.070)	-1.312 (-2.3740.250)	0.28	

Model includes baseline eGFR (CKD-EPI), age, baseline LVEF, log (NT-proBNP) [pg/mL] at baseline, log (hs-cTnT) [ng/L] at baseline as linear covariate(s) and region, diabetes at baseline, sex, treatment, study, baseline eGFR (CKD-EPI) by Time interaction, Time by tertiles of IGFBP7 at baseline interaction, tertiles of IGFBP7 at baseline as fixed effect(s). Intercept and slope allowed to vary randomly between patients. \* Slope measured at baseline eGFR (CKD-EPI) = 60.45.

Statistic	Placebo	Empagliflozin 10mg				
Baseline						
Values at visit						
Number of patients	558	561				
gMean (gCV [%])	133.7 (37.3)	135.2 (38.2)				
Week 12						
Values at visit						
Number of patients	555	554				
gMean (gCV [%])	137.2 (37.5)	134.5 (37.5)				
Adjusted* gMean	137.5	133.1				
95% confidence interval	(134.4, 140.6)	(130.2, 136.2)				
Relative change from baseline		. , ,				
gMean ratio (gCV [%])	1.02 (32.81)	0.99 (29.54)				
Adjusted* gMean ratio	1.03	0.99				
95% confidence interval	(1.00, 1.05)	(0.97, 1.02)				
Comparison vs Placebo (ratio of						
relative change)						
Adjusted* gMean ratio	referent	0.97				
95% confidence interval		(0.94, 1.00)				
p-value		0.05				
Week 52						
Values at visit						
Number of patients	352	356				
gMean (gCV [%])	135.2 (37.4)	138.0 (36.7)				
Adjusted* gMean	136.6	139.3				
95% confidence interval	(132.6, 140.7)	(135.3, 143.5)				
Relative change from baseline						
gMean ratio (gCV [%])	1.01 (33.6)	1.04 (34.6)				
Adjusted* gMean ratio	1.02	1.04				
95% confidence interval	(0.99, 1.05)	(1.01, 1.07)				
Comparison vs Placebo (ratio of						
relative change)						
Adjusted* gMean ratio	referent	1.02				
95% confidence interval		(0.98, 1.06)				
p-value		0.35				

 Table 4: IGFBP7 concentration (ng/mL) among placebo and empagliflozin arms across study visits

\* Model includes age, baseline eGFR (CKD–EPI), baseline LVEF as linear covariates and study, region, baseline diabetes status, sex, treatment by visit interaction, log(baseline IGFBP7) by visit interaction as fixed effects.

# Sampling for biobanking and IGFBP -7 measurements



\*Random sampling stratified by treatment and availability of data, originally selected for OLINK measurement, same patients us

ed for IGFBP -7 measurement.

## Figure 2:

Endpoint	Events/100 py (95% Cl)	HR (95% CI)		p-value for trend
CV death or first I	HHF			0.0008
<t1< th=""><th>6.05 (4.26, 8.15)</th><th>Referent</th><th></th><th></th></t1<>	6.05 (4.26, 8.15)	Referent		
≥T1 and <t2< td=""><td>6.78 (4.89, 8.98)</td><td>1.03 (0.66, 1.63)</td><td>⊢≢⊸</td><td></td></t2<>	6.78 (4.89, 8.98)	1.03 (0.66, 1.63)	⊢≢⊸	
≥T2	17.03 (13.76, 20.64)	2.00 (1.28, 3.10)	<b>⊢-≣</b>	
First HHF				0.0016
<t1< td=""><td>3.76 (2.38, 5.45)</td><td>Referent</td><td></td><td></td></t1<>	3.76 (2.38, 5.45)	Referent		
≥T1 and <t2< td=""><td>4.68 (3.14, 6.54)</td><td>1.12 (0.64, 1.96)</td><td>⊢⊢∎−−−</td><td></td></t2<>	4.68 (3.14, 6.54)	1.12 (0.64, 1.96)	⊢⊢∎−−−	
≥T2	13.23 (10.37, 16.43)	2.20 (1.28, 3.78)	·∎	-
Total HHF (first an	nd recurrent)			0.0041
<t1< td=""><td>7.79 (4.66, 13.02)</td><td>Referent</td><td></td><td></td></t1<>	7.79 (4.66, 13.02)	Referent		
≥T1 and <t2< td=""><td>7.41 (4.54, 12.09)</td><td>0.95 (0.53, 1.72)</td><td>⊢<b></b></td><td></td></t2<>	7.41 (4.54, 12.09)	0.95 (0.53, 1.72)	⊢ <b></b>	
≥T2	16.59 (10.97, 25.09)	2.13 (1.19, 3.82)	<b>⊢</b>	-
First event of inte	nsification of diuretic ther	apy		0.1856
<t1< td=""><td>8.05 (5.91, 10.50)</td><td>Referent</td><td></td><td></td></t1<>	8.05 (5.91, 10.50)	Referent		
≥T1 and <t2< td=""><td>9.94 (7.55, 12.65)</td><td>1.09 (0.73, 1.63)</td><td>┝──╋──┥</td><td></td></t2<>	9.94 (7.55, 12.65)	1.09 (0.73, 1.63)	┝──╋──┥	
≥T2	15.30 (12.19, 18.76)	1.31 (0.87, 1.99)	ı∔∎—ı	
CV death				0.0389
<t1< td=""><td>3.18 (1.94, 4.71)</td><td>Referent</td><td></td><td></td></t1<>	3.18 (1.94, 4.71)	Referent		
≥T1 and <t2< td=""><td>3.08 (1.88, 4.58)</td><td>0.90 (0.48, 1.71)</td><td>⊢<b></b></td><td></td></t2<>	3.08 (1.88, 4.58)	0.90 (0.48, 1.71)	⊢ <b></b>	
≥T2	6.49 (4.64, 8.65)	1.84 (1.00, 3.41)		
Sustained eGFR r	eduction ≥40% or ESRD			0.0011
<t1< td=""><td>0.97 (0.32, 1.99)</td><td>Referent</td><td></td><td></td></t1<>	0.97 (0.32, 1.99)	Referent		
≥T1 and <t2< td=""><td>1.11 (0.41, 2.17)</td><td>1.08 (0.32, 3.60)</td><td></td><td>4</td></t2<>	1.11 (0.41, 2.17)	1.08 (0.32, 3.60)		4
≥T2	5.22 (3.41, 7.41)	4.66 (1.61, 13.53)	, <u> </u>	
		· · · · · · · · · · · · · · · · · · ·		
		0.25	0.5 1 2	4 8 16
			of overt	+ d rick of overt
		Decreased fisk	or event increase	u lisk of event





Figure 4:

