- Supplemental Figure 1: The correlation coefficients determined from spearman
 correlation analysis between hs-CRP and forearm muscle protein turnover components
 using regular scale for hs-CRP among whole study population (n=129).
- 730 Supplemental Figure 2: The rates of forearm muscle protein components
- 731 according to hsCRP concentrations grouped by quintiles among non-diabetic
- 732 **MHD patients.** We examined protein turnover markers are presented as median
- 733 (interquartile range) after categorizing patients into quartiles based on their hs-CRP

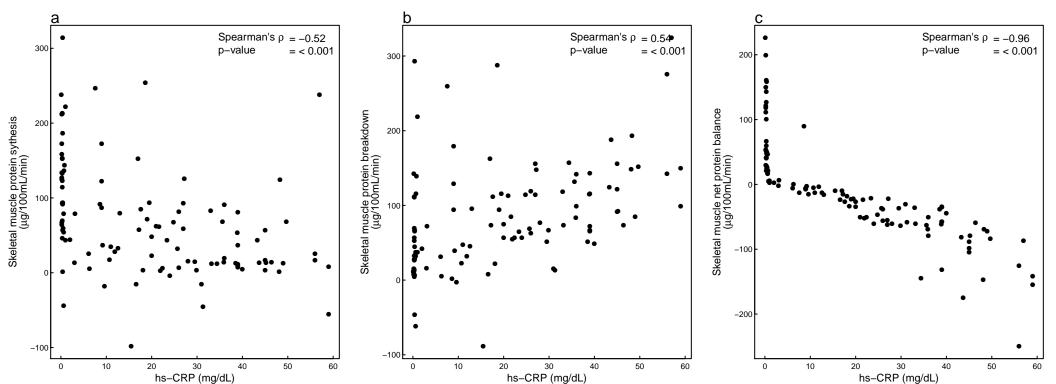
- values. There were statistically significant associations between hs-CRP quintiles and
- forearm skeletal muscle protein synthesis (S2A), protein degradation (S2B) and net
- forearm skeletal muscle protein balance (**S3C)** (p<0.001 for all). The unadjusted
- comparisons of four groups defined by hs-CRP quintiles and various skeletal muscle
- and whole body protein turnover components were performed using the Kruskall-Wallis
- test and the data are depicted as box plots.

740Supplemental Figure 3: The correlation coefficients between hs-CRP and forearm

741 muscle protein turnover components in non-diabetic MHD patients (n=100).

- 742 Spearman correlation analysis showed an inverse (negative) correlation between hs-
- CRP (log scale) and forearm skeletal muscle protein synthesis (S3A) whereas a direct
- 744 (positive) correlation was observed with protein degradation (S3B). The inverse
- (negative) correlation between hs-CRP and net protein balance was remarkably robust(S3C).
- 747 Supplemental Figure 4: The correlation coefficients between serum interleukin-6
- 748 (IL-6) concentrations and forearm muscle protein turnover components in MHD
- 749 **patients (n=56).** Spearman correlation analysis showed an inverse (negative)
- correlation between IL-6 (log scale) and forearm net skeletal muscle protein balance
- 751 although this did not reach statistical significance.

752



Supplemental Figure 2

