## **Description of Additional Supplementary Files**

File Name: Supplementary Data 1

Description: Metabolite reporting standards including quality control metrics.

File Name: Supplementary Data 2

Description: Whole genome sequencing analysis of plasma metabolome (519 locus-metabolite

associations).

File Name: Supplementary Data 3

Description: Unknown metabolite annotations using bioinformatic tools (SIRIUS, GNPS, HMDB, In-house

library).

File Name: Supplementary Data 4

Description: Associations between Unknown Peaks and Clinical Phenotypes (Jackson Heart Study).

File Name: Supplementary Data 5

Description: Phenotypic associations with sentinel SNP's using Phenoscanner V2.

File Name: Supplementary Data 6

Description: MS/MS fragmentation data for high condifence metabolite predictions. DDA: data

dependent acquisition; RT: retention time.

File Name: Supplementary Data 7

Description: Metabolite ID-GNPS Node Map.