

**S9 Table. Associations between inflammatory biomarkers and intima media thickness at the right common carotid artery (N=728)**

Biomarker	Model A*		Model B†		Model C‡	
	β	(95% CI)	β	(95% CI)	β	(95% CI)
<b>sCD163</b>						
Quintile 1	Ref	Ref	Ref	Ref	Ref	Ref
Quintile 2	0.00	[-0.03, 0.03]	-0.01	[-0.04, 0.02]	-0.01	[-0.04, 0.02]
Quintile 3	0.02	[-0.01, 0.06]	0.02	[-0.01, 0.05]	0.02	[-0.01, 0.05]
Quintile 4	0.03	[-0.00, 0.06]	0.02	[-0.01, 0.05]	0.02	[-0.02, 0.05]
Quintile 5	0.03	[-0.00, 0.06]	0.01	[-0.02, 0.04]	0.01	[-0.03, 0.04]
<b>sCD14</b>						
Quintile 1	Ref	Ref	Ref	Ref	Ref	Ref
Quintile 2	0.00	[-0.03, 0.03]	0.00	[-0.03, 0.03]	0.01	[-0.02, 0.04]
Quintile 3	0.02	[-0.01, 0.06]	0.02	[-0.01, 0.05]	0.01	[-0.02, 0.04]
Quintile 4	0.02	[-0.01, 0.05]	0.02	[-0.01, 0.05]	0.03	[-0.00, 0.06]
Quintile 5	0.03	[-0.01, 0.06]	0.01	[-0.02, 0.05]	0.01	[-0.02, 0.05]
<b>ICAM-1</b>						
Quintile 1	Ref	Ref	Ref	Ref	Ref	Ref
Quintile 2	-0.02	[-0.05, 0.01]	-0.02	[-0.05, 0.01]	-0.01	[-0.04, 0.02]
Quintile 3	0.00	[-0.03, 0.03]	0.00	[-0.03, 0.03]	0.01	[-0.02, 0.04]
Quintile 4	0.01	[-0.02, 0.04]	0.00	[-0.03, 0.03]	-0.01	[-0.04, 0.02]
Quintile 5	0.02	[-0.01, 0.06]	0.02	[-0.02, 0.05]	0.00	[-0.03, 0.04]
<b>CCL2</b>						
Quintile 1	Ref	Ref	Ref	Ref	Ref	Ref
Quintile 2	0.00	[-0.04, 0.03]	0.00	[-0.04, 0.03]	0.00	[-0.03, 0.03]
Quintile 3	0.02	[-0.01, 0.05]	0.01	[-0.02, 0.04]	0.01	[-0.02, 0.04]
Quintile 4	0.02	[-0.01, 0.06]	0.01	[-0.02, 0.04]	0.00	[-0.03, 0.03]
Quintile 5	0.03*	[0.00, 0.07]	0.02	[-0.01, 0.05]	0.01	[-0.02, 0.04]
<b>CRP</b>						
Quintile 1	Ref	Ref	Ref	Ref	Ref	Ref
Quintile 2	-0.01	[-0.04, 0.03]	-0.01	[-0.04, 0.02]	-0.02	[-0.05, 0.01]
Quintile 3	0.02	[-0.01, 0.05]	0.02	[-0.01, 0.05]	0.00	[-0.03, 0.03]
Quintile 4	0.02	[-0.01, 0.05]	0.01	[-0.02, 0.04]	-0.01	[-0.04, 0.02]
Quintile 5	0.05**	[0.02, 0.08]	0.03*	[0.00, 0.06]	0.02	[-0.02, 0.05]
<b>IL-6</b>						
Quintile 1	Ref	Ref	Ref	Ref	Ref	Ref
Quintile 2	0.03	[-0.00, 0.06]	0.02	[-0.01, 0.05]	0.00	[-0.03, 0.03]
Quintile 3	0.02	[-0.01, 0.05]	0.01	[-0.02, 0.04]	-0.01	[-0.04, 0.02]
Quintile 4	0.07***	[0.03, 0.10]	0.05**	[0.02, 0.08]	0.03	[-0.00, 0.06]

Biomarker	Model A*		Model B†		Model C‡	
	$\beta$	(95% CI)	$\beta$	(95% CI)	$\beta$	(95% CI)
Quintile 5	0.05**	[0.02, 0.08]	0.02	[-0.01, 0.06]	0.00	[-0.03, 0.04]
<b>sTNF-<math>\alpha</math>R1</b>						
Quintile 1	Ref	Ref	Ref	Ref	Ref	Ref
Quintile 2	0.00	[-0.03, 0.03]	0.00	[-0.03, 0.03]	0.01	[-0.02, 0.04]
Quintile 3	0.02	[-0.01, 0.05]	0.02	[-0.01, 0.05]	0.02	[-0.01, 0.05]
Quintile 4	0.02	[-0.01, 0.05]	0.02	[-0.01, 0.05]	0.02	[-0.01, 0.05]
Quintile 5	0.05**	[0.01, 0.08]	0.03	[-0.00, 0.06]	0.02	[-0.01, 0.05]
<b>sTNF-<math>\alpha</math>R2</b>						
Quintile 1	Ref	Ref	Ref	Ref	Ref	Ref
Quintile 2	0.02	[-0.01, 0.05]	0.01	[-0.02, 0.04]	0.01	[-0.02, 0.04]
Quintile 3	0.01	[-0.02, 0.04]	0.00	[-0.03, 0.03]	0.01	[-0.02, 0.04]
Quintile 4	0.01	[-0.02, 0.05]	0.00	[-0.03, 0.04]	0.01	[-0.02, 0.04]
Quintile 5	0.05***	[0.02, 0.09]	0.03	[-0.00, 0.06]	0.03	[-0.00, 0.06]
<b>Fibrinogen</b>						
Quintile 1	Ref	Ref	Ref	Ref	Ref	Ref
Quintile 2	0.01	[-0.02, 0.04]	0.01	[-0.02, 0.04]	0.01	[-0.02, 0.04]
Quintile 3	0.05**	[0.01, 0.08]	0.04*	[0.01, 0.07]	0.03	[-0.00, 0.06]
Quintile 4	0.04*	[0.00, 0.07]	0.02	[-0.01, 0.05]	0.01	[-0.02, 0.04]
Quintile 5	0.07***	[0.04, 0.11]	0.06***	[0.02, 0.09]	0.04*	[0.01, 0.07]
<b>D-dimer</b>						
Quintile 1	Ref	Ref	Ref	Ref	Ref	Ref
Quintile 2	0.00	[-0.03, 0.04]	-0.01	[-0.04, 0.02]	0.00	[-0.03, 0.03]
Quintile 3	0.00	[-0.03, 0.03]	-0.02	[-0.05, 0.01]	-0.02	[-0.05, 0.01]
Quintile 4	0.02	[-0.01, 0.06]	0.00	[-0.03, 0.03]	0.00	[-0.03, 0.03]
Quintile 5	0.00	[-0.03, 0.03]	-0.03*	[-0.06, -0.00]	-0.04*	[-0.06, -0.01]

Abbreviations: sCD163, cluster of differentiation 163; sCD14, cluster of differentiation 14; CCL2, chemokine (C-C motif) ligand 2; ICAM-1, intercellular cell adhesion molecule-1; CRP, C reactive protein; IL-6, interleukin-6; sTNF- $\alpha$ R1, tumor necrosis factor-alpha receptor 1; sTNF- $\alpha$ R2, tumor necrosis factor-alpha receptor 2. Results presented as  $\beta$  coefficients and 95% confidence intervals, \*  $p < 0.05$ , \*\*  $p < 0.01$ , \*\*\*  $p < 0.001$

\*Model A: Unadjusted.

†Model B: Adjusted for HIV serostatus, age, race, baseline education, center, cohort.

‡Model C: Adjusted for variables in model B along with cumulative pack years, alcohol consumption since last visit, HCV, BMI, SBP (per 10mm Hg), total cholesterol (per 5mg/dl), HDL (5mg/dl), glucose levels (per 10 mg/dl), and use of medication for hypertension, diabetes and high cholesterol.