

**TABLE S2.** Associations of serum albumin concentration with annual change\* in eGFRcr in HIV-infected WIHS participants.

	<b>Unadjusted Estimate (95% CI)</b>	<b>P Value</b>	<b>Adjusted without ACR ** Estimate (95% CI)</b>	<b>P Value</b>	<b>Adjusted with ACR ** Estimate (95% CI)</b>	<b>P Value</b>
<b>Continuous Predictors</b>						
Serum Albumin (per -0.5 g/dL)	-0.59 (-0.79, -0.40)	p<0.001	-0.52 (-0.71, -0.32)	p<.0001	-0.48 (-0.68, -0.29)	p<0.001
ACR (per doubling)	-0.22 (-0.33, -0.12)	p<0.001			-0.24 (-0.34, -0.15)	p<0.001
Baseline eGFRcr (per ↑10 ml/min)	-0.72 (-0.82, -0.62)	p<0.001	-0.82 (-0.92, -0.72)	p<.0001	-0.84 (-0.94, -0.75)	p<0.001
<b>Dichotomized Predictors</b>						
Serum Albumin < 3.8 g/dL	-0.64 (-1.02, -0.26)	p=0.0010	-0.40 (-0.78, -0.016)	p=0.041	-0.32 (-0.69, 0.063)	p=0.10
ACR ≥ 30 mg/g	-0.86 (-1.43, -0.29)	p=0.0033			-1.30 (-1.84, -0.76)	p<.0001
Baseline eGFRcr<60 ml/min	4.66 (3.71, 5.61)	p<.0001	4.65 (3.74, 5.57)	p<.0001	5.14 (4.21, 6.07)	p<.0001

\* Results reported as estimated annual change (95% confidence interval) calculated from multivariable linear mixed models.

\*\* Adjusted models control for age, race, HIV RNA, CD4, HCV, DM, ARV use, baseline eGFR, SBP, DBP, BMI, and study site. For ACR as a predictor instead of adjusting for ACR the full model adjusts for serum albumin concentration.

\*\*\*ARV included current combination antiretroviral therapy (cART) use, current nucleoside reverse transcriptase inhibitor (NRTI) use, current non-nucleoside reverse transcriptase inhibitor (NNRTI) use, and current protease inhibitor (PI) use.