

Circulating TNF Receptors 1 and 2 Predict Mortality in Patients with End-stage
Renal Disease Undergoing Dialysis

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Supplementary Table 1.TNF-related biomarker added predictive benefit for prediction of all-cause mortality in hemodialysis patients -ROC curve analysis-

	AUC	Difference in AUC (95%CI)	P
hs-CRP	0.613		
TNF α & hs-CRP vs. hs-CRP	0.637	0.055 (-0.001, 0.055)	0.13
TNFR1 & hs-CRP vs. hs-CRP	0.632	0.064 (-0.027, 0.064)	0.43
TNFR2 & hs-CRP vs. hs-CRP	0.678	0.126 (0.003, 0.126)	0.04
Basic model ^a	0.733		
Basic model + TNF α vs. Basic model	0.733	0.005 (-0.006, 0.005)	0.91
Basic model + TNFR1 vs. Basic model	0.736	0.019 (-0.013, 0.019)	0.71
Basic model + TNFR2 vs. Basic model	0.744	0.034 (-0.014, 0.034)	0.40

ROC, receiver-operating characteristic; AUC, areas under the ROC

^aBasic model: age, previous history of CVD, SBP (categorical), maximum Δ SBP, and hs-CRP

Abbreviations used in this table are the same as in Table 1.

Supplementary Table 2. TNF-related biomarker added predictive benefit for prediction of all-cause mortality in hemodialysis patients -IDI and NRI analysis-

	IDI (95%CI)	Relative IDI	P	NRI (95%CI)	P
TNF α & hs-CRP vs. hs-CRP	0.004 (-0.003, 0.012)	0.160	0.19	0.173 (-0.071, 0.418)	0.17
TNFR1 & hs-CRP vs. hs-CRP	0.014 (0.001, 0.027)	0.509	0.04	0.389 (0.154, 0.623)	0.002
TNFR2 & hs-CRP vs. hs-CRP	0.041 (0.018, 0.064)	1.487	<0.001	0.425 (0.185, 0.665)	<0.001
TNF α & Basic model vs. Basic model ^a	0.001 (-0.002, 0.004)	0.005	0.66	0.077 (-0.168, 0.322)	0.54
TNFR1 & Basic model vs. Basic model	0.008 (-0.001, 0.018)	0.06	0.08	0.277 (0.038, 0.516)	0.03
TNFR2 & Basic model vs. Basic model	0.022 (0.006, 0.039)	0.165	<0.01	0.373 (0.133, 0.614)	<0.01

IDI, integrated discrimination improvement; NRI, net reclassification index

^aBasic model: age, previous history of CVD, SBP (categorical), maximum Δ SBP, hs-CRP

Abbreviations used in this table are the same as in Table 1.