Supplementary Material for Predictive Modeling of Short-Term Recidivism: A Mixture Cure Rate Approach with Diverse Link Functions

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1 Additional Simulation Results

This supplementary material provides additional simulation results to support the findings presented in the main text. Specifically, it includes results for scenarios where the true latency distribution follows a lognormal or a loglogistic model, complementing the Weibull based simulations discussed in the main manuscript. Each case evaluates the performance of the four link functions, with true value recovery, coverage, and model assessment evaluation.

In additional, we present Cox-Snell residual plots of the Weibull based simulation that further assess model fit. These plots compare the fit of each link function when used to estimate data generated under each of the other link functions, offering insight into goodness-of-fit and the flexibility of the proposed link structures. These figures and tables are organized to match the structure of the main manuscript and provide supporting evidence for the flexibility of the proposed modeling framework.

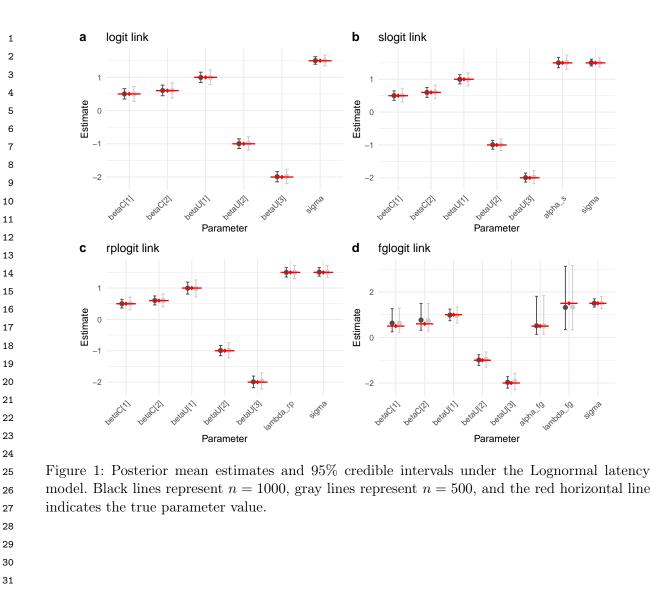
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Table 1: Coverage probabilities for each parameter for lognormal and loglogistic latency across difference link functions, and sample sizes.

N	Distribution	Parameter	Logit	SLogit	RPLogit	FGLogit
500	Lognormal	$\beta_C[1]$	0.939	0.926	0.941	1.000
		$\beta_C[2]$	0.959	0.936	0.938	1.000
		$\beta_U[1]$	0.948	0.947	0.944	0.946
		$\beta_U[2]$	0.946	0.951	0.952	0.946
		$\beta_U[3]$	0.937	0.953	0.952	0.932
		α	_	0.934	_	1.000
		λ	_	_	0.947	1.000
		σ	0.953	0.945	0.954	0.950
	Loglogistic	$\beta_C[1]$	0.932	0.941	0.938	1.000
		$\beta_C[2]$	0.948	0.953	0.952	1.000
		$\beta_U[1]$	0.949	0.950	0.952	0.934
		$\beta_U[2]$	0.945	0.945	0.943	0.942
		$\beta_U[3]$	0.946	0.932	0.945	0.938
		α	_	0.935	_	1.000
		λ	_	_	0.953	1.000
		σ	0.949	0.934	0.945	0.953
	Lognormal	$\beta_C[1]$	0.954	0.936	0.947	1.000
1000		$\beta_C[2]$	0.949	0.941	0.951	1.000
		$\beta_U[1]$	0.937	0.935	0.957	0.953
		$\beta_U[2]$	0.944	0.940	0.940	0.945
		$\beta_U[3]$	0.937	0.933	0.943	0.945
		α	_	0.946	_	1.000
		λ	_	_	0.951	1.000
		σ	0.943	0.952	0.949	0.948
	Loglogistic	$\beta_C[1]$	0.945	0.943	0.936	1.000
		$\beta_C[2]$	0.954	0.951	0.944	1.000
		$\beta_U[1]$	0.939	0.940	0.943	0.959
		$\beta_U[2]$	0.940	0.953	0.934	0.949
		$\beta_U[3]$	0.936	0.946	0.934	0.947
		α	_	0.941	_	1.000
		λ	_	_	0.942	1.000
		σ	0.951	0.941	0.954	0.949

Table 2: Number of times each model was selected under different true models with lognormal and loglogistic latency

	Criterion	True slogit	True rplogit	True fglogit
	LOOIC	91	85	78
Lognormal	WAIC	91	86	78
	DIC	92	89	87
	LOOIC	91	87	79
Loglogistic	WAIC	91	87	78
	DIC	88	81	71



model. Black lines represent n = 1000, gray lines represent n = 500, and the red horizontal line



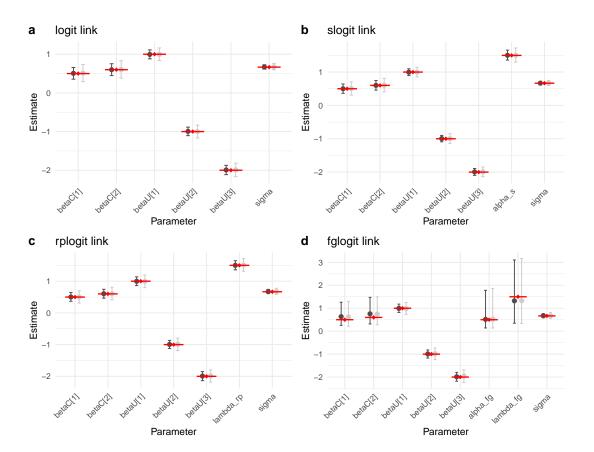


Figure 2: Posterior mean estimates and 95% credible intervals under the Loglogistic latency model. Black lines represent n=1000, gray lines represent n=500, and the red horizontal line indicates the true parameter value.

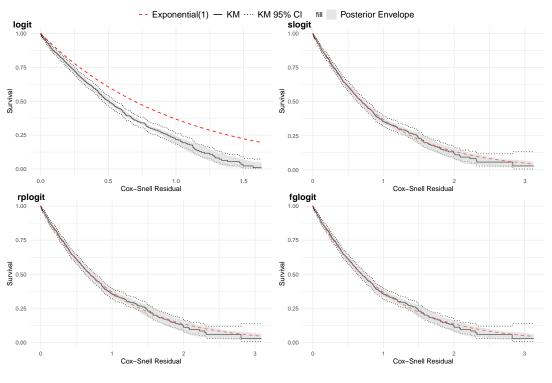


Figure 3: Cox-Snell residual plot with Weibull slogit link being true model

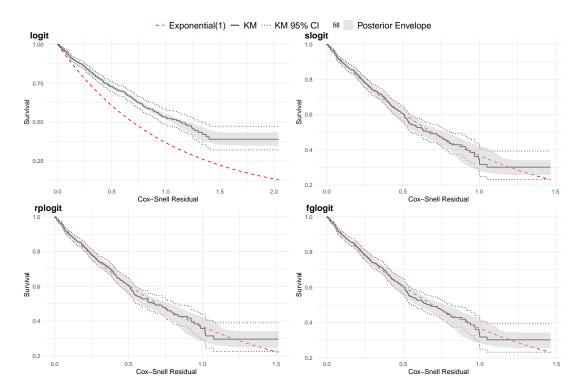


Figure 4: Cox-Snell residual plot with Weibull rplogit link being true model

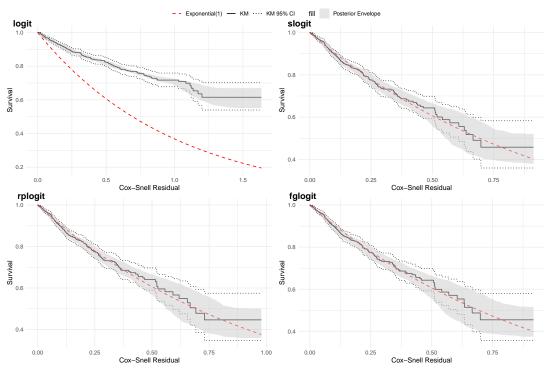


Figure 5: Cox-Snell residual plot with Weibull fglogit link being true model

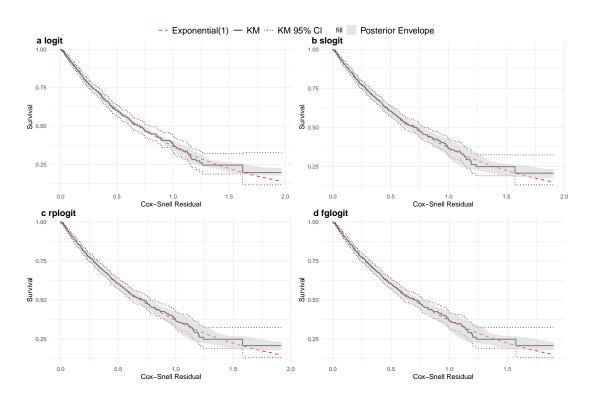


Figure 6: Cox-Snell residual plot with Lognormal logit link being true model

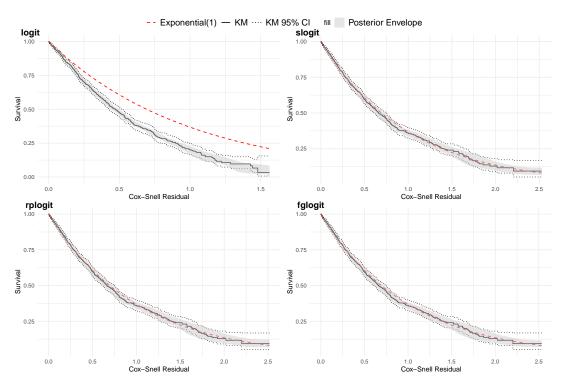


Figure 7: Cox-Snell residual plot with Lognormal slogit link being true model

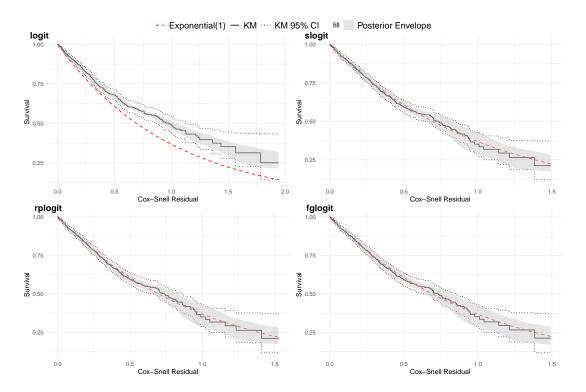
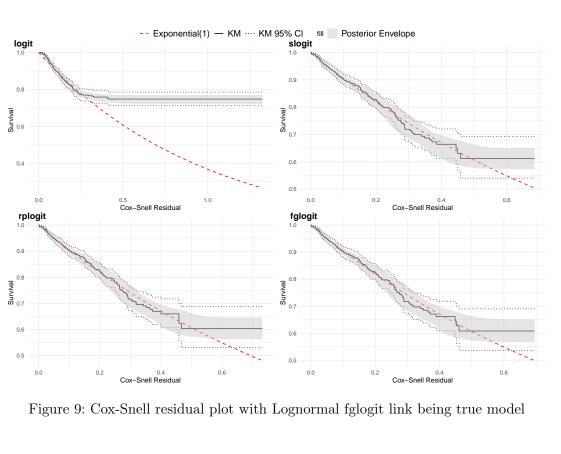


Figure 8: Cox-Snell residual plot with Lognormal rplogit link being true model



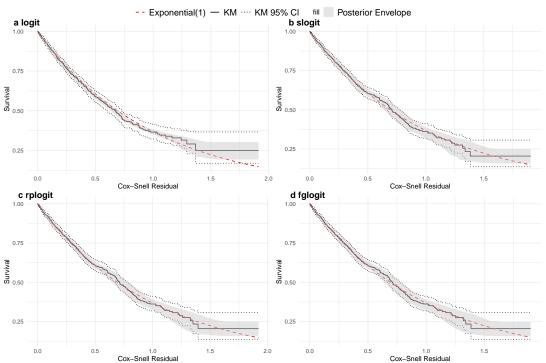


Figure 10: Cox-Snell residual plot with Loglogistic logit link being true model

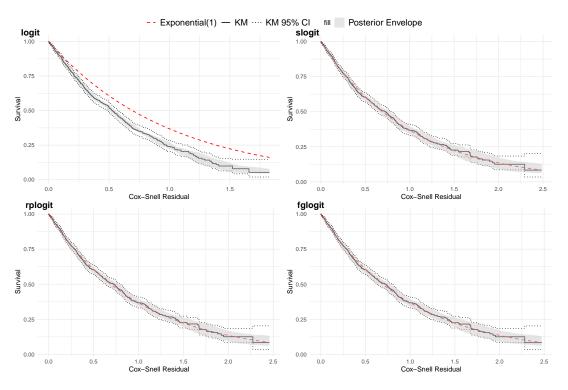


Figure 11: Cox-Snell residual plot with Loglogistic slogit link being true model

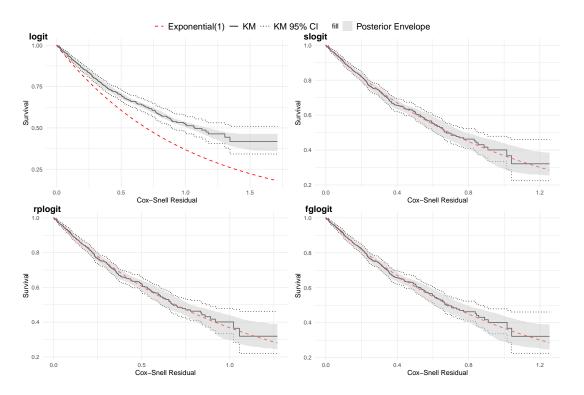
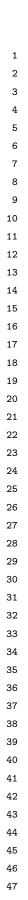


Figure 12: Cox-Snell residual plot with Loglogistic rplogit link being true model



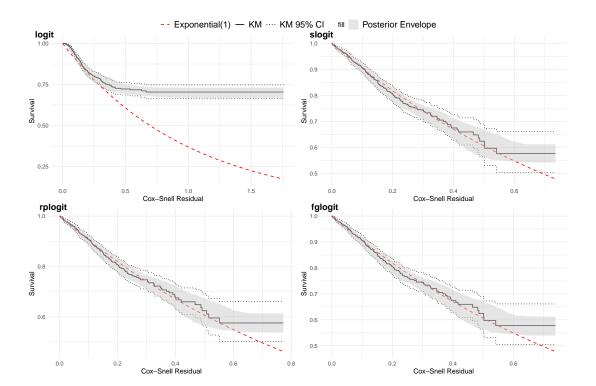


Figure 13: Cox-Snell residual plot with Loglogistic fglogit link being true model