

## Supplementary Materials

### Optimizing PCR Detection of West Nile Virus from Body Fluid Specimens to Delineate Natural History in an Infected Human Cohort

**Table S1.** Individual and average (expected) Ct values and respective concentrations of WNV RNA generated by repeated testing of the spiked reference control replicates.

Replicates	Set 1		Set 2		Set 3		Average		SD	
	Ct	cp/ $\mu$ l	Ct	cp/ $\mu$ l	Ct	cp/ $\mu$ l	Ct	cp/ $\mu$ l	Ct	cp/ $\mu$ l
LO 1	32.2	9.1	32.4	8.1	32.4	8.0	32.4	8.2	0.1	0.5
LO 2	32.3	8.6	32.4	7.6	32.3	8.8				
LO 3	32.4	8.2	32.4	8.0	32.5	7.6				
MED 1	26.2	646.0	26.5	497.3	26.4	538.3	26.4	571.6	0.1	56.0
MED 2	26.1	659.1	26.3	578.5	26.4	563.0				
MED 3	26.3	604.9	26.4	541.6	26.5	515.9				
HI 1	20.8	29,560.3	21.1	24,108.8	21.1	22,248.6	21.0	25,622.1	0.2	3181.2
HI 2	20.7	30,088.5	21.0	25,329.0	21.1	22,251.0				
HI 3	20.8	28,062.9	20.9	26,903.1	21.2	22,046.9				

SD, standard deviation; cp/ $\mu$ l, copies per microliter; LO, low spiking load; MED, medium spiking load; HI, high spiking load.

**Table S2.** Individual Ct values and percent recovery of WNV RNA in whole blood and urine specimens using original and optimized extraction protocols.

	Rep 1 Ct	Rep 2 Ct	Rep 3 Ct	Expected Ct	#Rep within 1 Ct	% recovery	Average % recovery
WB LO	34.2	34.4	34.2	32.4	0/3	26	19
WB MED	29.1	29.2	29.0	26.4	0/3	14	
WB HI	23.4	23.8	23.6	21.0	0/3	16	
WB optimized LO	31.6	31.6	31.7	32.4	3/3	163	133
WB optimized MED	26.2	26.0	26.0	26.4	3/3	125	
WB optimized HI	20.8	20.8	20.8	21.0	3/3	112	
UR A LO	37.9	37.4	38.4	32.4	0/3**	2	2
UR A MED	33.7	31.4	32.6	26.4	0/3**	1	
UR A HI	26.2	26.5	26.4	21.0	0/3**	2	
UR A optimized LO	32.1	31.9	32.0	30.1*	3/3**	25	27
UR A optimized MED	25.9	25.9	25.8	24.1*	3/3**	28	
UR A optimized HI	20.7	20.6	20.4	18.7*	3/3**	26	
UR B LO	33.3	33.2	33.8	32.4	2/3	49	52
UR B MED	27.4	27.3	27.4	26.4	3/3	49	
UR B HI	21.7	21.7	21.7	21.0	3/3	59	
UR B optimized LO	30.7	30.8	30.6	30.1*	3/3	65	58
UR B optimized MED	24.8	24.8	24.8	24.1*	3/3	60	
UR B optimized HI	19.5	19.5	19.9	18.7*	2/3	50	

\* Expected Ct decreased by 2.3 due to 5-fold higher volume of spiked specimen (1 ml vs. 0.2 ml).

\*\* Number of biological replicates within 2 Ct units from expected Ct.

Rep, biological replicate; WB, whole blood; UR, urine; LO, low spiking load; MED, medium spiking load; HI, high spiking load.