

## Supplement table and figure

**Table S1. Complete blood count of rhesus macaques**

### Comparison CBC results at differential day after feeding for control group

Tests	Units	Day after feeding for control group						
		-21 day	-14 day	0 day	7 day	14 day	21 day	28 day
RBC	10 <sup>6</sup> /ul	4.79	4.76	4.83	5.01	4.93	5.12	5.02
Hct	%	38	38	39	40	39	41	40
Hb	gm%	12.0	12.2	12.2	12.7	12.3	12.7	12.5
WBC	cells/ul	3,700	4,500	4,200	7,500	4,100	4,700	3,300
Plt	cells/ul	154,000	235,000	233,000	173,000	265,000	206,000	251,000
MCV	fl	79.7	79.4	79.7	79.0	79.9	79.9	80.1
MCH	pg	25.1	25.6	25.3	25.3	24.9	24.8	24.9
MCHC	g/dl	31.4	32.3	31.7	32.1	31.2	31.1	31.1
Neu-s	%	35	63	58	51	57	49	41
N-b	%	0	2	2	2	2	3	3
Lymp	%	56	28	35	38	37	42	46
Bas	%	1	0	1	1	0	1	1
Eos	%	1	0	1	1	0	1	1
Mon	%	4	3	2	3	1	1	2

### Comparison CBC results at differential day after feeding for NHP-1

Tests	Units	Day after feeding for NHP-1						
		-21 day	-14 day	0 day	7 day	14 day	21 day	28 day
RBC	10 <sup>6</sup> /ul	6.38	5.99	6.26	6.16	6.12	6.06	6.18
Hct	%	44	41	43	42	42	41	42
Hb	gm%	14.5	14.0	14.7	14.2	14.1	13.8	14.2
WBC	cells/ul	5,400	6,800	7,900	6,700	6,600	8,800	7,900
Plt	cells/ul	303,000	266,000	359,000	341,000	339,000	340,000	376,000
MCV	fl	69.1	68.6	68.1	68.8	68.3	68.2	68.1
MCH	pg	22.7	23.4	23.5	23.1	23.0▼	22.8▼	23.0▼
MCHC	g/dl	32.9	34.1	34.5	33.5	33.7▼	33.4▼	33.7
Neu-s	%	42	57	65	46	53	49▼	50▼
N-b	%	1	2	2	1	2	0	1
Lymp	%	49	37	31	47	38	46▲	45▲
Bas	%	1	0	1	0	3	1	1
Eos	%	1	0	1	0	3	1	1
Mon	%	6	3	1	3	1▼	2	2

**Comparison CBC results at differential day after feeding for NHP-3**

Tests	Units	Day after feeding for NHP-3						
		-21 day	-14 day	0 day	7 day	14 day	21 day	28 day
RBC	10 <sup>6</sup> /ul	5.35	5.33	5.43	5.58	5.44	5.09	5.43
Hct	%	40	40	41	42	41	39	41
Hb	gm%	13.0	13.4	13.2	13.4	13.0	12.3	13.0
WBC	cells/ul	4,900	7,000	4,300	5,300	6,600	8,100	7,300
Plt	cells/ul	263,000	269,000	333,000	427,000	279,000	391,000	320,000
MCV	fl	75.0	75.2	75.3	76.0	76.1	76.2	74.3
MCH	pg	24.3	25.1	24.3	24.0	23.9▼	24.2▼	24.4▼
MCHC	g/dl	32.4	33.4	32.3	31.6	31.4▼	31.7▼	32.8
Neu-s	%	38	46	48	48	61	33▼	34▼
N-b	%	4	1	2	2	5	2	0
Lymp	%	53	46	41	46	28	58▲	60▲
Bas	%	0	0	0	0	0	0	0
Eos	%	0	0	0	0	0	0	0
Mon	%	2	3	1	4	0▼	3	3

‘▲’ means that the- values showed significantly increased after chigger challenge. ‘▼’ means that the values showed significantly decreased after chigger challenge.

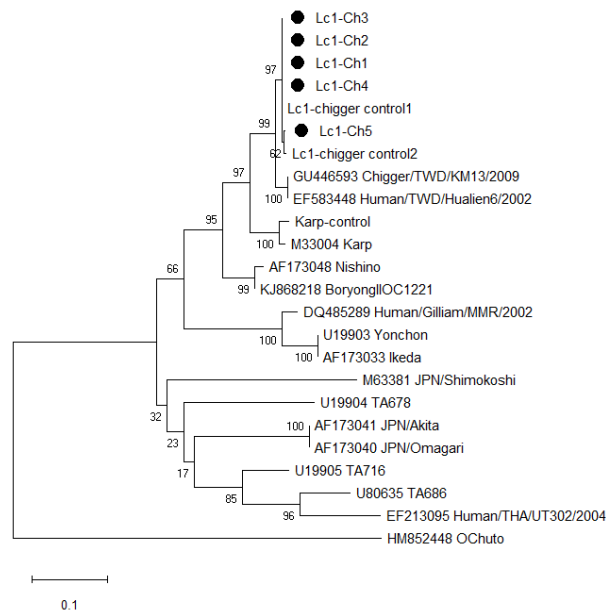
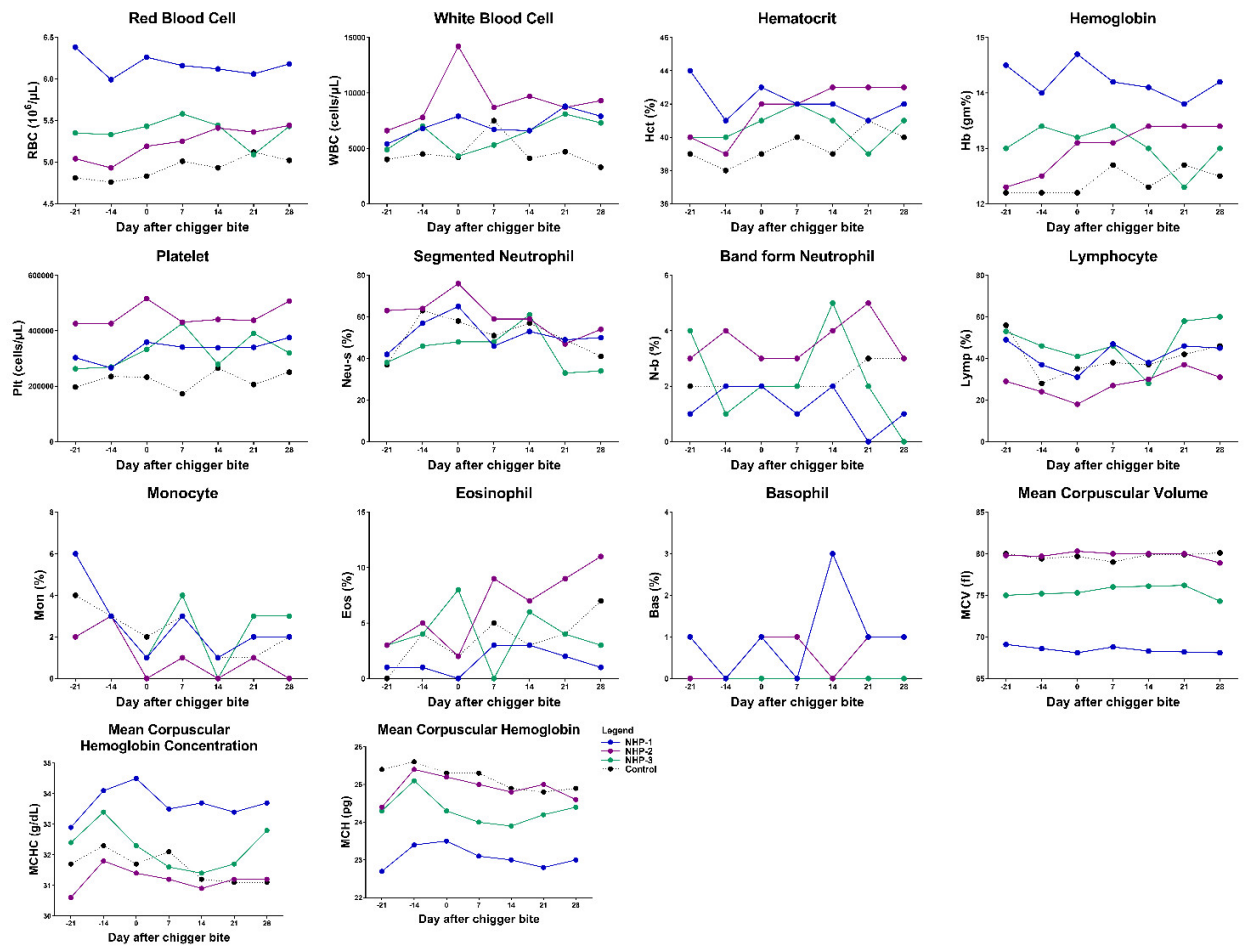


Figure S1. Phylogenetic analysis of *O. tsutsugamushi* in *Leptotrombidium chaingraiensis* chiggers from the colonies. The evolutionary tree was inferred by using Maximum likelihood method and General Time Reversible model. A discrete Gamma distribution was used to model evolutionary rate differences among sites. The tree is drawn to scale, with branch lengths measured in the number of substitutions per site. Numbers at nodes indicate bootstrap value at 1000 replications.



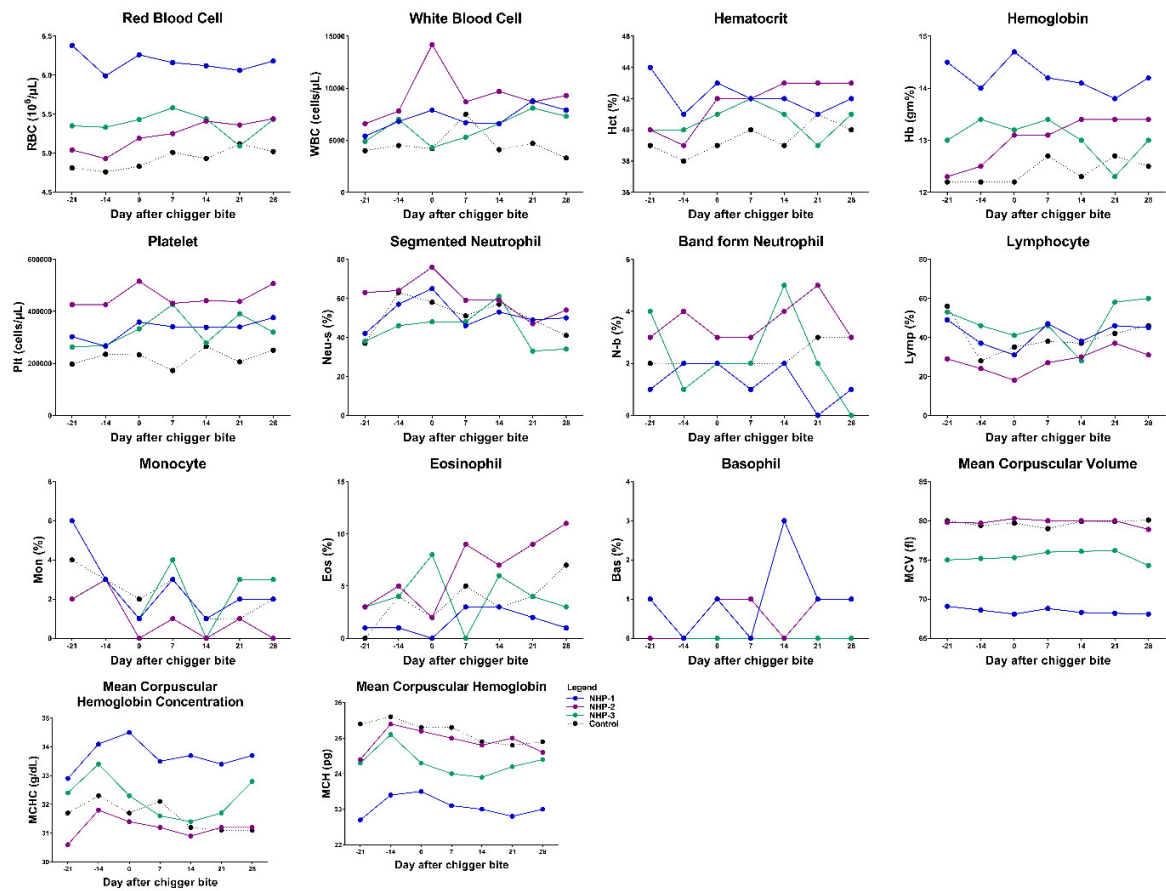


Figure S2. Hematological analysis of rhesus macaques after *O. tsutsugamushi* infection or non-infected chigger bite (control).

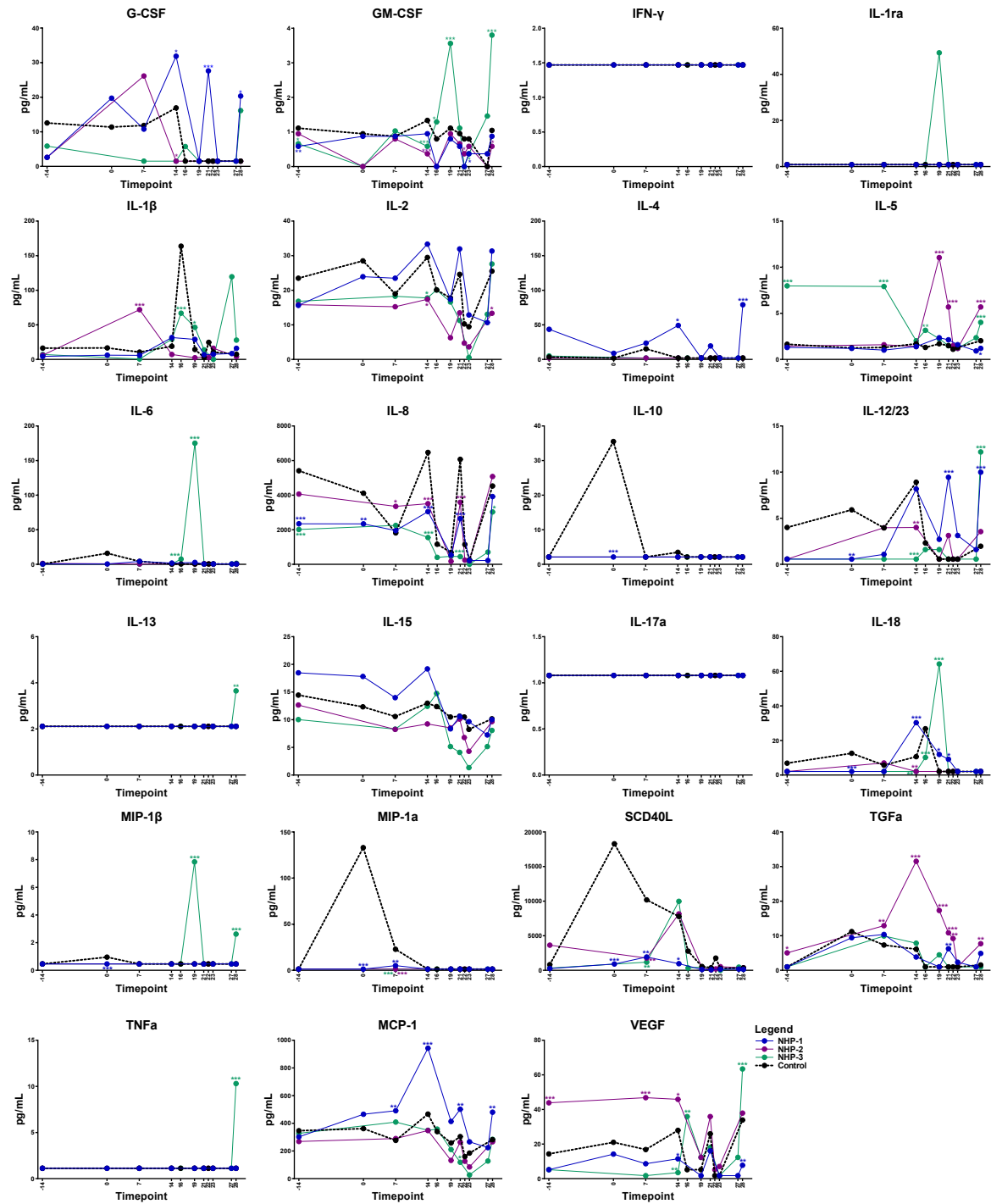
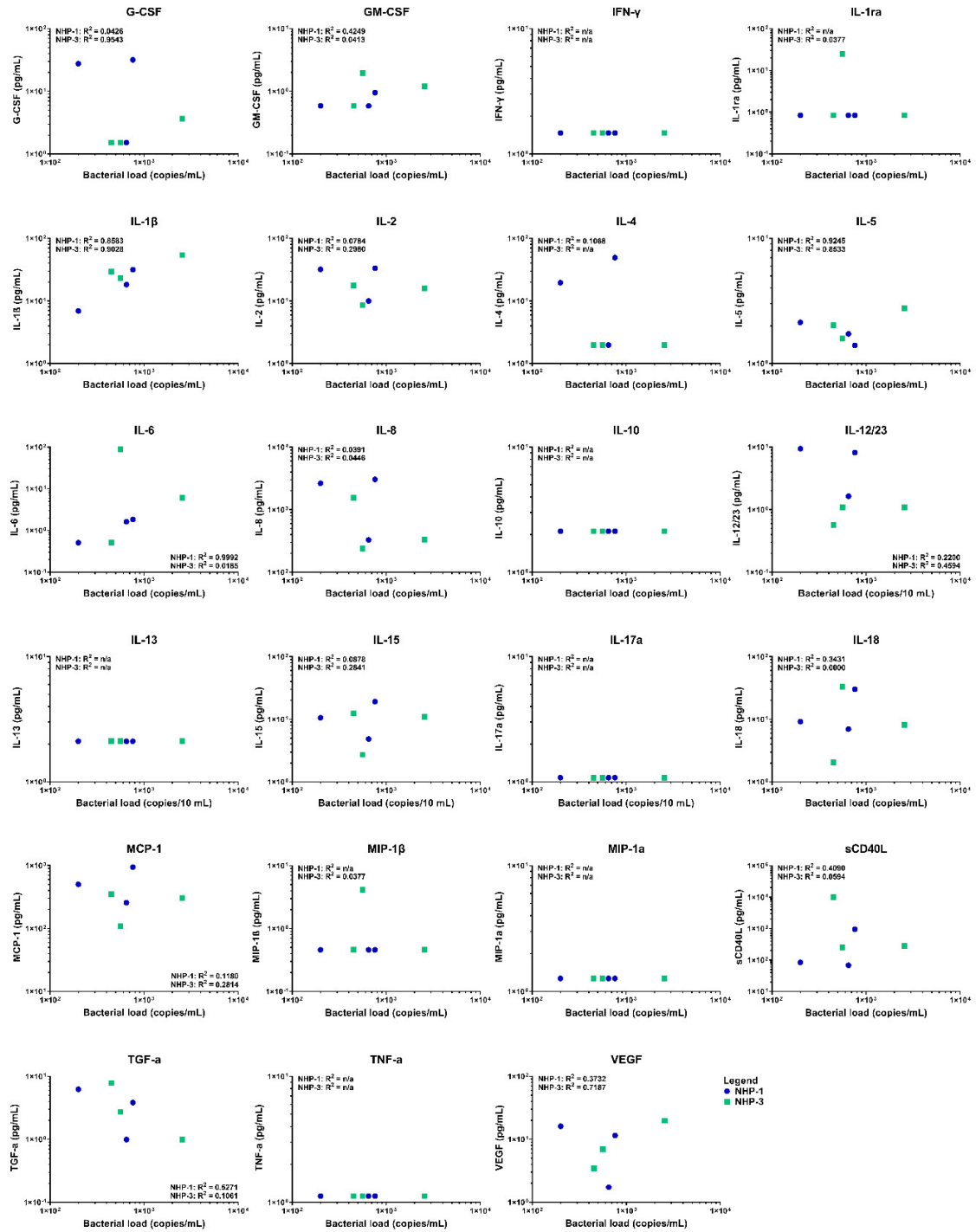


Figure S3. Serum levels of 23 cytokine/ chemokine in rhesus macaques during *O. tsutsugamushi* infection via chigger bite.



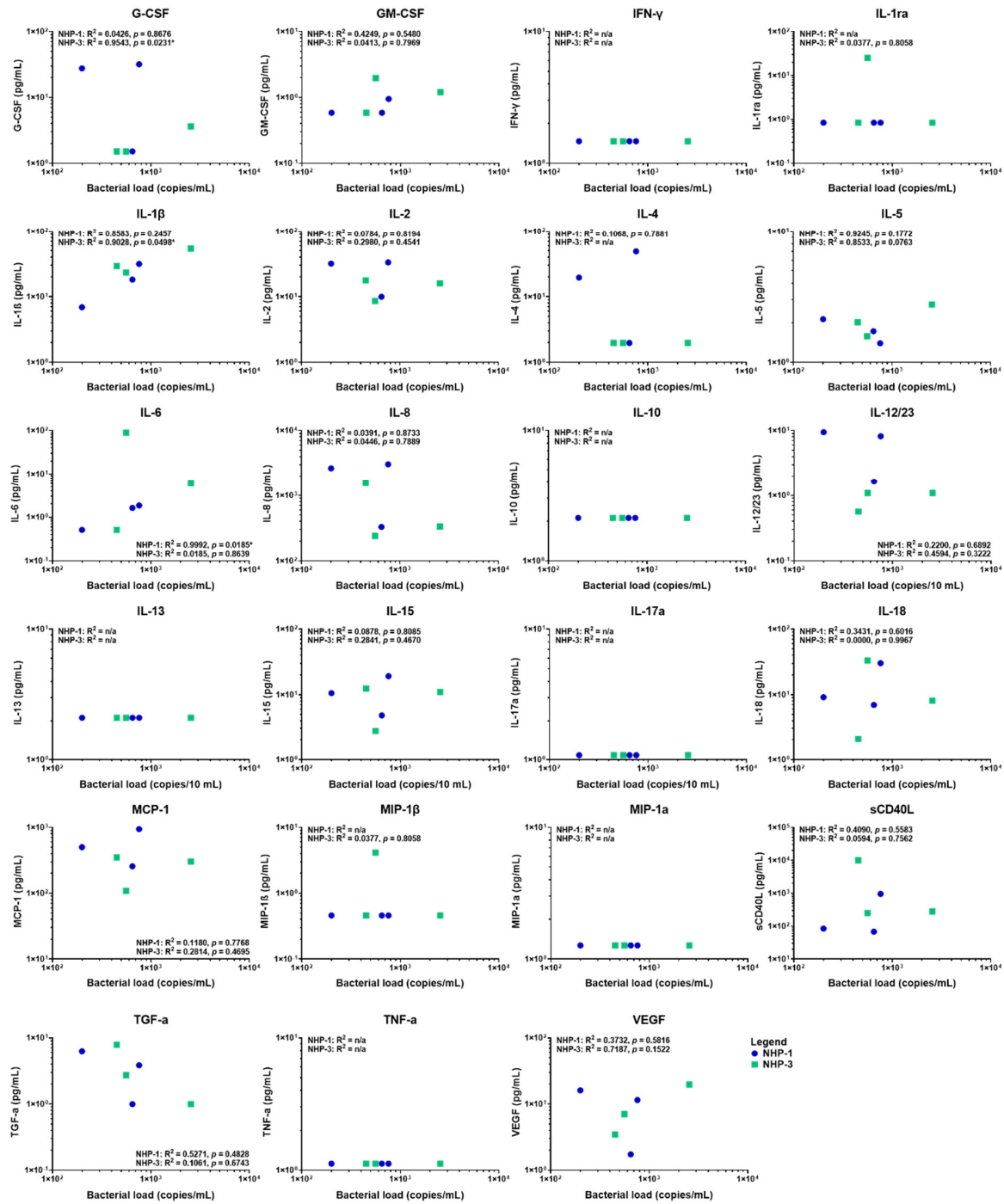


Figure S4. Immunological correlation analysis of serum cytokine/ chemokine of all macaques and bacteremia during acute phase of infection. Correlation efficiency and P value were calculated based on the duplication of each individual samples.