Supplementary Information for:

High prevalence of hepatitis delta virus in Cameroon

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Figure S1

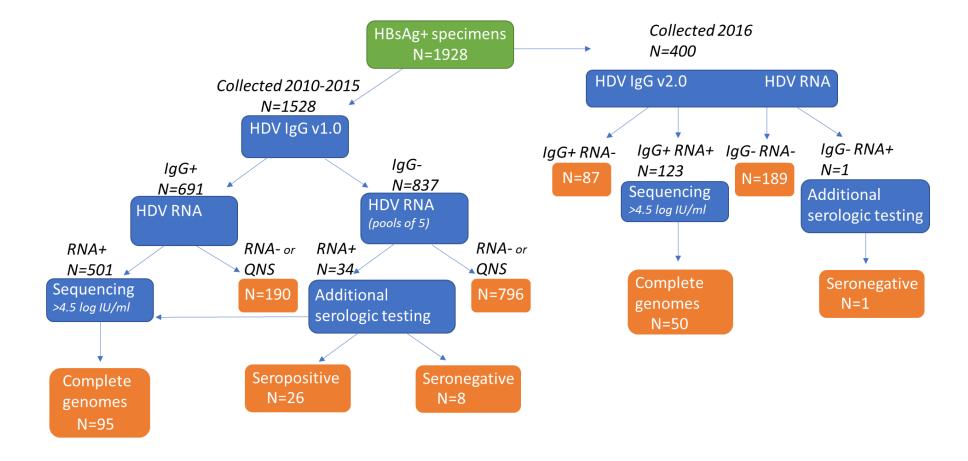


Figure S1. Testing workflow: Testing algorithms for specimens collected in 2010-2015 (left) and 2016 (right) are shown. For the 2010-2015 cohort, a total of 34 HDV RNA positive seronegative samples were identified. With additional serologic testing, 26 of these 34 were determined to be HDV seropositive: 3 samples were positive for HDV IgM only and 23 samples had IgG antibodies detectable using recombinant HDAg. Eight of the 34 seronegative samples were HDV RNA positive with no evidence of HDV antibodies (IgM or IgG) in the 2010-2015 cohort. Given the improvements in serodetection by testing with the recombinant HDAg, a revised assay incorporating both HDAg peptides and recombinant protein (version 2.0) was used to screen and additional set of HBsAg samples collected in 2016, with parallel testing for HDV RNA. One RNA+ seronegative specimen was identified in the 2016 cohort with viral load of 1.14 log IU/ml, additional serologic testing determined there was no evidence of HDV antibodies (IgM or IgG) in this specimen. *Quantity not sufficient (QNS)*

Table S1

					HDV RNA			HDV supplemental	
Sample	Year	Donor risk group	Site	HIV	HDV VL		HDV IgG	HDV IgM	HDV IgG
Sample	Teal	Dollor risk group	Site	status	(log	HDV GT	s/co	_	(rAg)
					IU/ml)			S/N ^a	S/N ^a
Evidence of HDV IgM antibodies only									
1	2013	Unknown illness	Kribi	NEG	0.44	VLNS	0.02	34.45	1.33
2	2011	VTC	Kribi	NEG	1.63	VLNS	0.08	12.08	QNS
3	2013	Unknown illness	Ngovagang	POS	8.05	1	0.17	39.01	QNS
Evidence of HDV IgG									
4	2013	Blood donor	Yaounde	QNS	1.82	VLNS	0.28	100.47	77.26
5	2013	Unknown illness	Djoum	NEG	4.69	1	0.98	31.89	81.64
6	2013	Unknown illness	Sangmelima	POS	5.27	1	0.48	8.01	87.18
7	2014	Unknown illness	Bipindi	POS	4.04	1	0.86	3.32	86.57
8	2013	Unknown illness	Ebolowa	NEG	6.78	6	0.92	1.30	77.61
9	2010	Chest clinic	Yaounde	POS	3.82	VLNS	0.78	1.02	84.23
10	2015	Antenatal clinic	Ebolowa	POS	1.94	VLNS	0.72	1.79	10.47
11	2014	Unknown illness	Kribi	NEG	5.26	1	0.52	3.16	37.97
12	2015	Antenatal clinic	Ebolowa	POS	4.12	QNS	0.41	0.89	49.20
13	2014	Unknown illness	Lolodorf	NEG	6.07	1	0.19	1.86	8.60
14	2011	VTC	Sangmelima	NEG	7.31	7	0.52	4.79	30.60
15	2013	Unknown illness	Ebolowa	POS	8.41	1	0.66	0.51	39.70
16	2015	VTC	Sangmelima	NEG	5.82	1	0.57	1.02	44.00
17	2015	Blood donor	Yaounde	NEG	0.58	VLNS	0.06	0.83	9.27
18	2014	Unknown illness	Kribi	POS	0.66	VLNS	0.29	1.14	92.10
19	2015	Chest clinic	Yaounde	POS	2.64	VLNS	0.42	3.37	83.77
20	2012	Unknown illness	Ngovagang	NEG	3.18	7	0.12	3.25	91.98
21	2011	Antenatal clinic	Ebolowa	NEG	2.94	VLNS	0.74	2.39	100.59
22	2013	Chest clinic	Yaounde	QNS	5.33	1	0.10	0.54	50.81
23	2010	Chest clinic	Yaounde	POS	5.78	1	0.08	2.10	98.89
24	2011	VTC	Kribi	NEG	6.99	7	0.17	0.85	98.89
25	2012	Hospital patient	Yaounde	POS	6.75	1	0.06	1.06	6.60
26	2011	Blood donor	Yaounde	POS	3.46	7	0.08	6.92	26.80
No evidence of HDV seroconversion									
27	2013	Unknown illness	Ebolowa	NEG	0.48	VLNS	0.02	3.29	1.58
28	2013	Unknown illness	Kribi	NEG	1.55	VLNS	0.01	4.98	1.06
29	2015	Blood donor	Yaounde	NEG	0.75 ^b	VLNS	0.35	0.85	5.48
30	2015	Blood donor	Yaounde	NEG	0.79 ^b	VLNS	0.18	0.55	0.61
31	2015	Blood donor	Yaounde	NEG	0.81 ^b	VLNS	0.02	1.42	0.58
32	2015	Blood donor	Yaounde	NEG	1.23 ^b	VLNS	0.05	0.84	0.16
33	2013	Chest clinic	Yaounde	QNS	2.25	VLNS	0.10	1.42	0.83
34	2014	Unknown	unknown	POS	3.75	1	0.04	0.44	0.90
^a Shown are signal to noise values, S/N>10 considered as evidence of HDV antibodies. Original cut-off in peptide IgG									
ascau was 25*NC Values >6 are considered elevated indicated in italics									

assay was 35*NC. Values >6 are considered elevated, indicated in italics.

Four seronegative blood donor specimens with viral load below <1.5 mentioned in the text are indicated in bold

^b Not determined due to insufficient volume, tested with recombinant HDAg in IgM assay. QNS: quantity not sufficient VLNS: viral load not sufficient

Figure S2

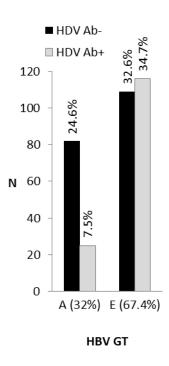


Figure S1. HBV genotype data. HBV genotype data was available for 334 samples in the study cohort; 2 HBV GT AE samples were HDV negative. Results for HBV GT A and E are given above by HDV seroprevalence.