

Association between HBsAg loss and long-term clinical outcome in chronic hepatitis B: a systematic review and meta-analysis

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Berkeley Public Health

ILC Late-Breaker Poster LBP-02

- Abstract was accepted as a Late-Breaker Poster
 - Reference number: LBP-02
- Please respect that the data presented here is confidential
- Will be on display during all three poster sessions at ILC

Background

- Studies have shown an association between hepatitis B surface antigen (HBsAg) loss and improved long-term clinical outcomes in chronic hepatitis B (CHB) patients
- Evaluation of HBsAg seroclearance as a ‘functional cure’ on a larger scale is crucial for drug development and regulatory decision making

Objectives

- To perform a systematic literature review and meta-analysis describing the association between functional cure (defined as HBsAg seroclearance with or without anti-HBs seroconversion) of CHB infection and long-term clinical outcome.

Methods

- Literature search in PubMed, EMBASE and Cochrane Library databases
- Full-text articles published in English between January 1990 – November 2018

Methods

- Inclusion criteria:
 - >50 CHB patients
 - ≥ 2 years of follow-up
 - HBsAg serostatus measured at baseline & during follow-up
 - Reported data on 1 or more long-term clinical outcomes (decompensation, HCC, liver transplantation (LT), and/or all-cause mortality)

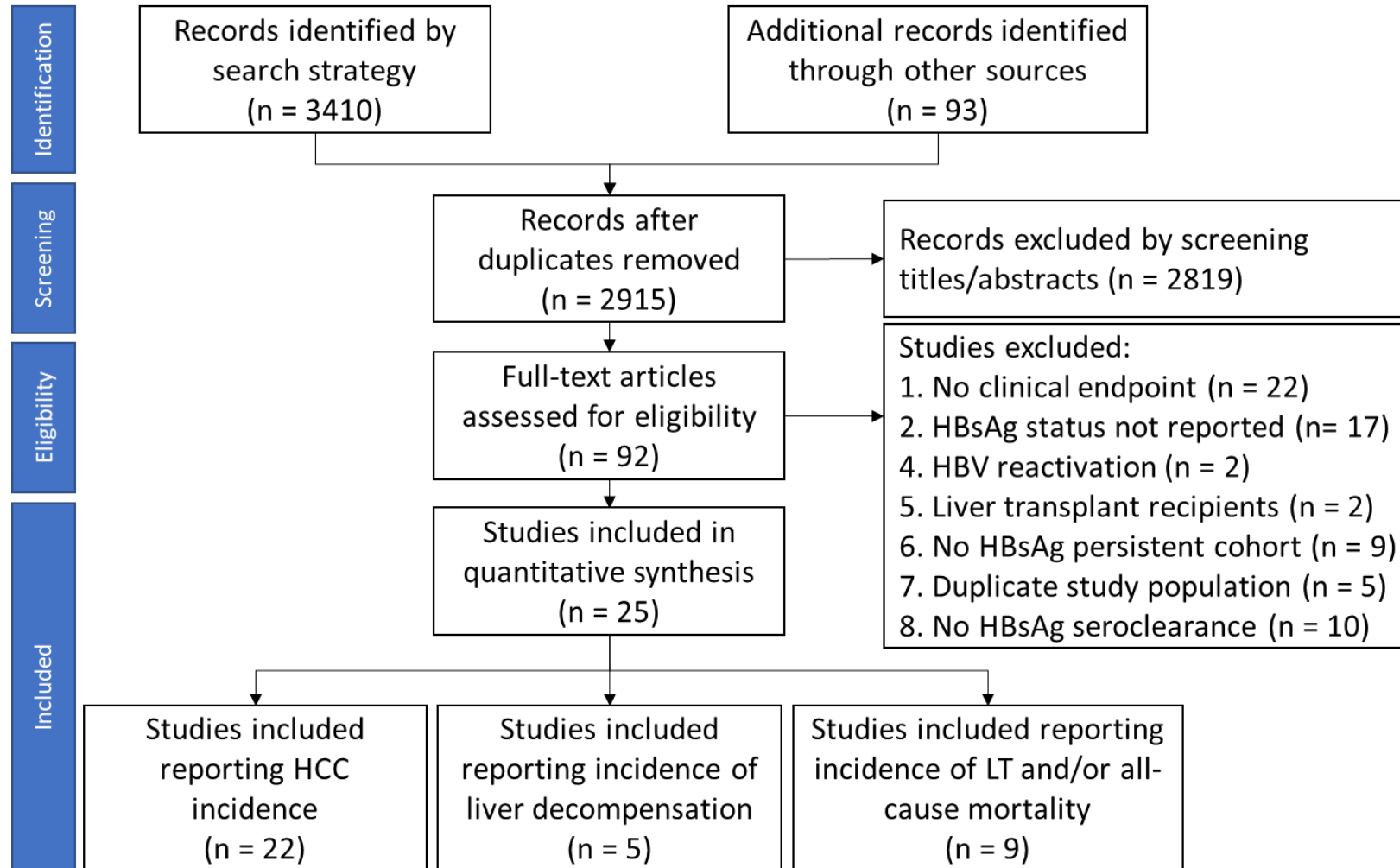
Methods

- **Exclusion criteria:**
 - Study duplication
 - HBV reactivation studies
 - Studies on LT patients
 - Co-infection with HCV, HIV, and/or delta*
 - Included 4 studies with minority subpopulations of co-infected individuals
 - Clinical events reported prior to HBsAg loss
 - No HBsAg-persistent comparison cohort
 - Zero clinical events reported for both cohorts

Methods

- Two investigators independently extracted and reviewed study data
- Comprehensive Meta-analysis Software was used to perform statistical analysis
 - Clinical events rates (number of clinical events in cohort/person-years follow-up in cohort) were used to calculate the rate ratio (RR) using a random effects model
 - Reciprocal continuity correction factors used for studies reporting zero-events for the HBsAg-loss cohort¹
 - Patient population demographic characteristics were categorized and analyzed in sub-group (Q-test for heterogeneity) and sensitivity analyses
 - General study information was used as covariates in a meta-regression analysis

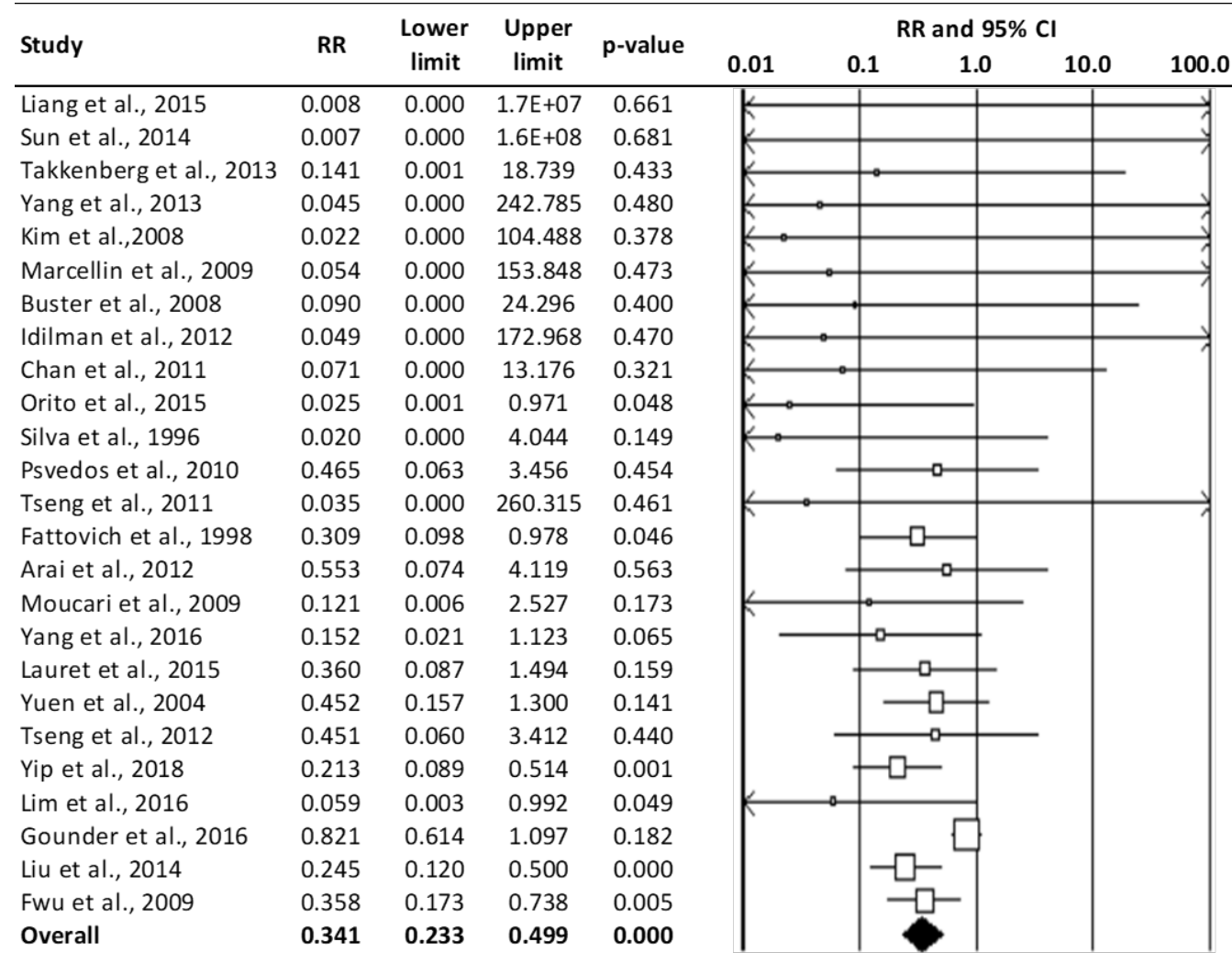
Study Selection Flow Chart



“First Clinical Event” Endpoint

- Composite endpoint of the first reported adverse clinical event
 - Decompensation, HCC, liver transplant, or death

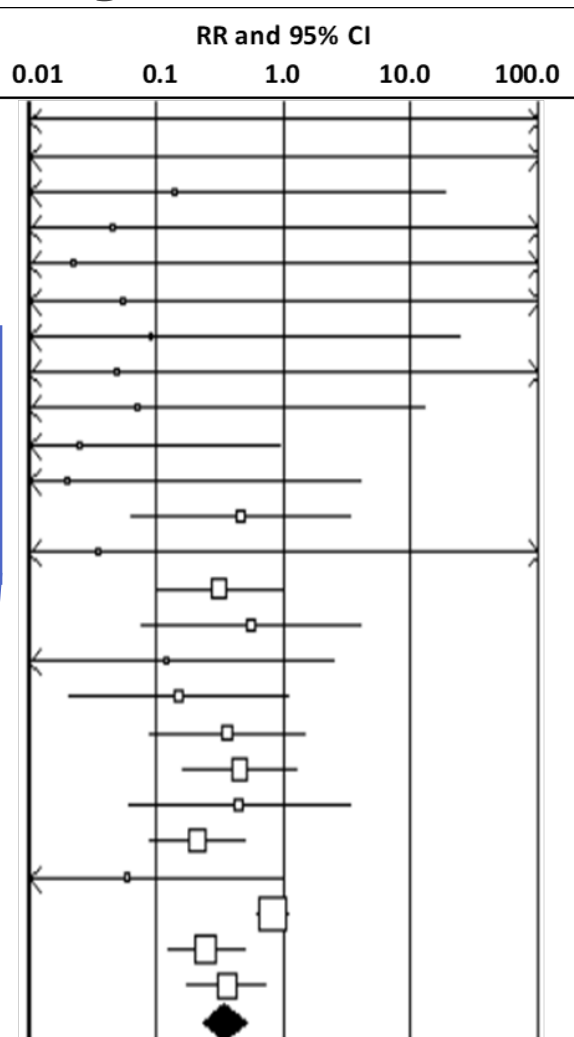
First Clinical Event: Meta-Analysis







First Clinical Event: Meta-Analysis

Study	RR	Lower limit	Upper limit	p-value	RR and 95% CI				
					0.01	0.1	1.0	10.0	100.0
Liang et al., 2015	0.008	0.000	1.7E+07	0.661					
Sun et al., 2014	0.007	0.000	1.6E+08	0.681					
Takkenberg et al., 2013	0.141	0.001	18.739	0.433					
Yang et al., 2013	0.045	0.000	242.785	0.480					
Kim et al., 2008	0.022	0.000	104.488	0.378					
Marcellin et al., 2009	0.054	0.000	153.848	0.473					

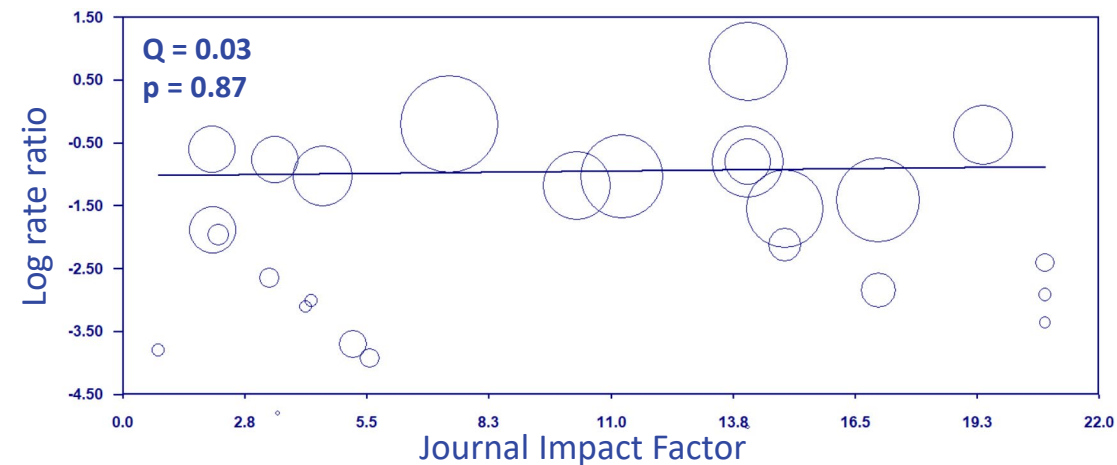
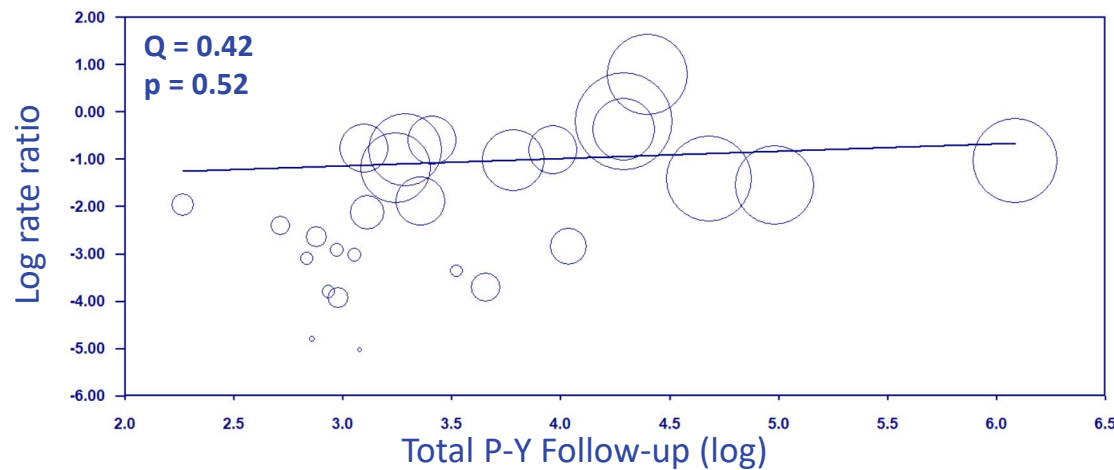
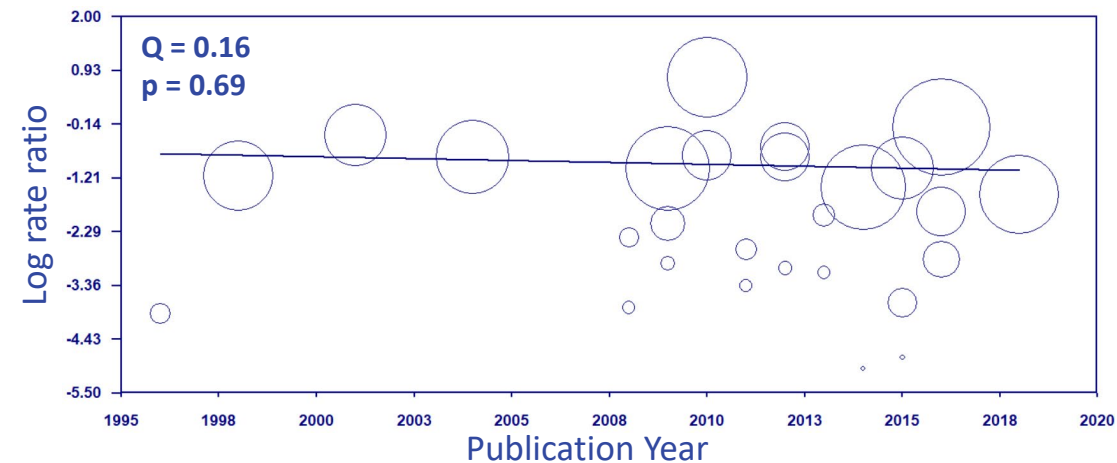
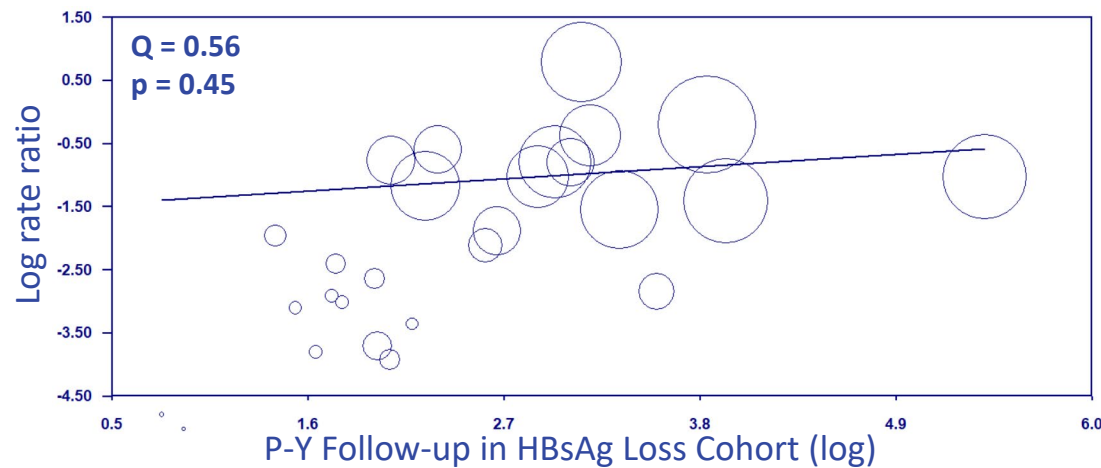
	RR	Lower limit	Upper limit	p-value
Overall	0.341	0.233	0.499	0.000
Fattovich et al., 1998	0.309	0.098	0.978	0.046
Arai et al., 2012	0.553	0.074	4.119	0.563
Moucari et al., 2009	0.121	0.006	2.527	0.173
Yang et al., 2016	0.152	0.021	1.123	0.065
Lauret et al., 2015	0.360	0.087	1.494	0.159
Yuen et al., 2004	0.452	0.157	1.300	0.141
Tseng et al., 2012	0.451	0.060	3.412	0.440
Yip et al., 2018	0.213	0.089	0.514	0.001
Lim et al., 2016	0.059	0.003	0.992	0.049
Gounder et al., 2016	0.821	0.614	1.097	0.182
Liu et al., 2014	0.245	0.120	0.500	0.000
Fwu et al., 2009	0.358	0.173	0.738	0.005
Overall	0.341	0.233	0.499	0.000



Primary Clinical Endpoints: Independent Meta-analyses

Endpoint	RR	Lower limit	Upper limit	p-value	RR and 95% CI		
					0.0	0.5	1.0
Decompensation (n=5)	0.313	0.150	0.654	0.002			
HCC (n = 22)	0.307	0.214	0.440	0.000			
LT/All-cause Mortality (n = 9)	0.751	0.569	0.991	0.043			
First Clinical Event (N = 25)	0.341	0.233	0.499	0.000			

First Clinical Event: Regression Analysis



First Clinical Event: Subgroup/Sensitivity Analysis

Treatment

HBV Genotype

Co-infection

Follow-up in HBsAg-loss cohort

Average follow-up

Baseline HBeAg status

Natural history

Treatment

Both

A/D

B/C

Yes

No

<50 person-years

50-999 person-years

>1000 person-years

<5 years

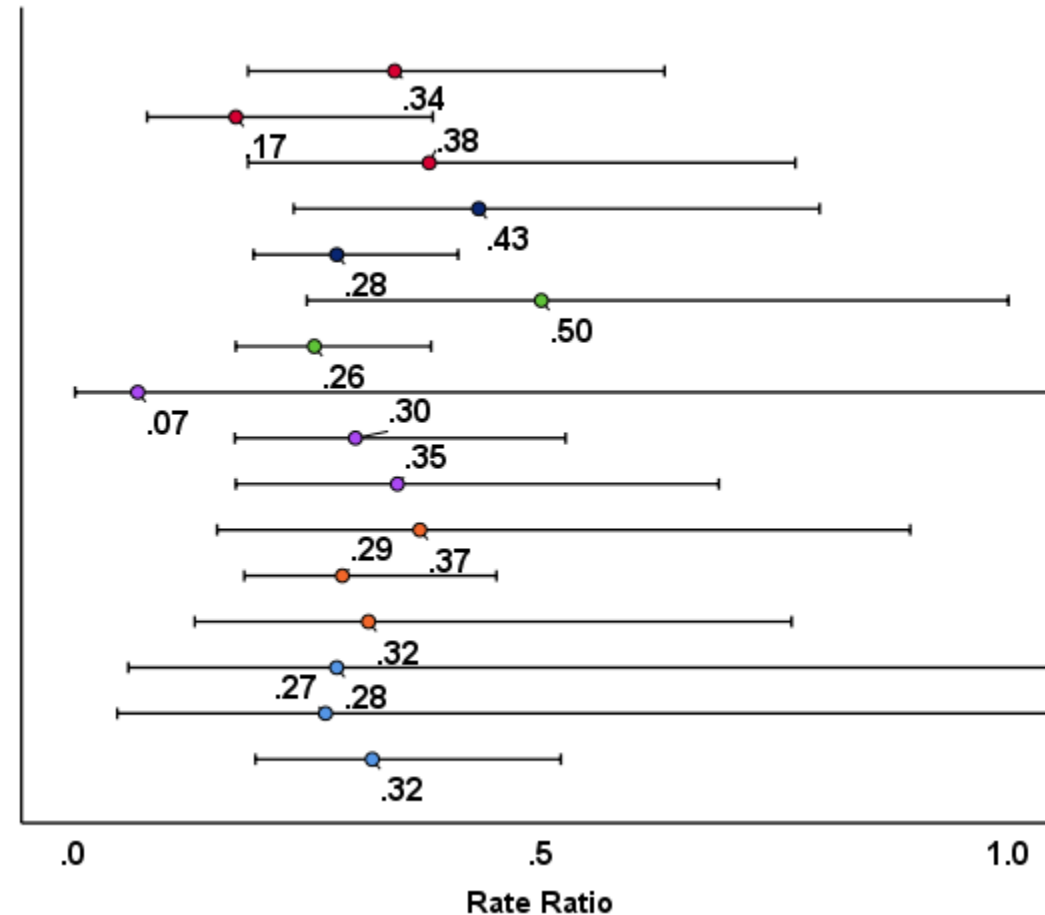
5-10 years

>10 years

Positive

Negative

Mix



First Clinical Event: Subgroup Analysis

Treatment

Natural history

HBV Genotype

Treatment

Co-infection

Both

Follow-up in
HBsAg-loss cohort

<50 person-years

50-999 person-years

>1000 person-years

Average follow-up

<5 years

5-10 years

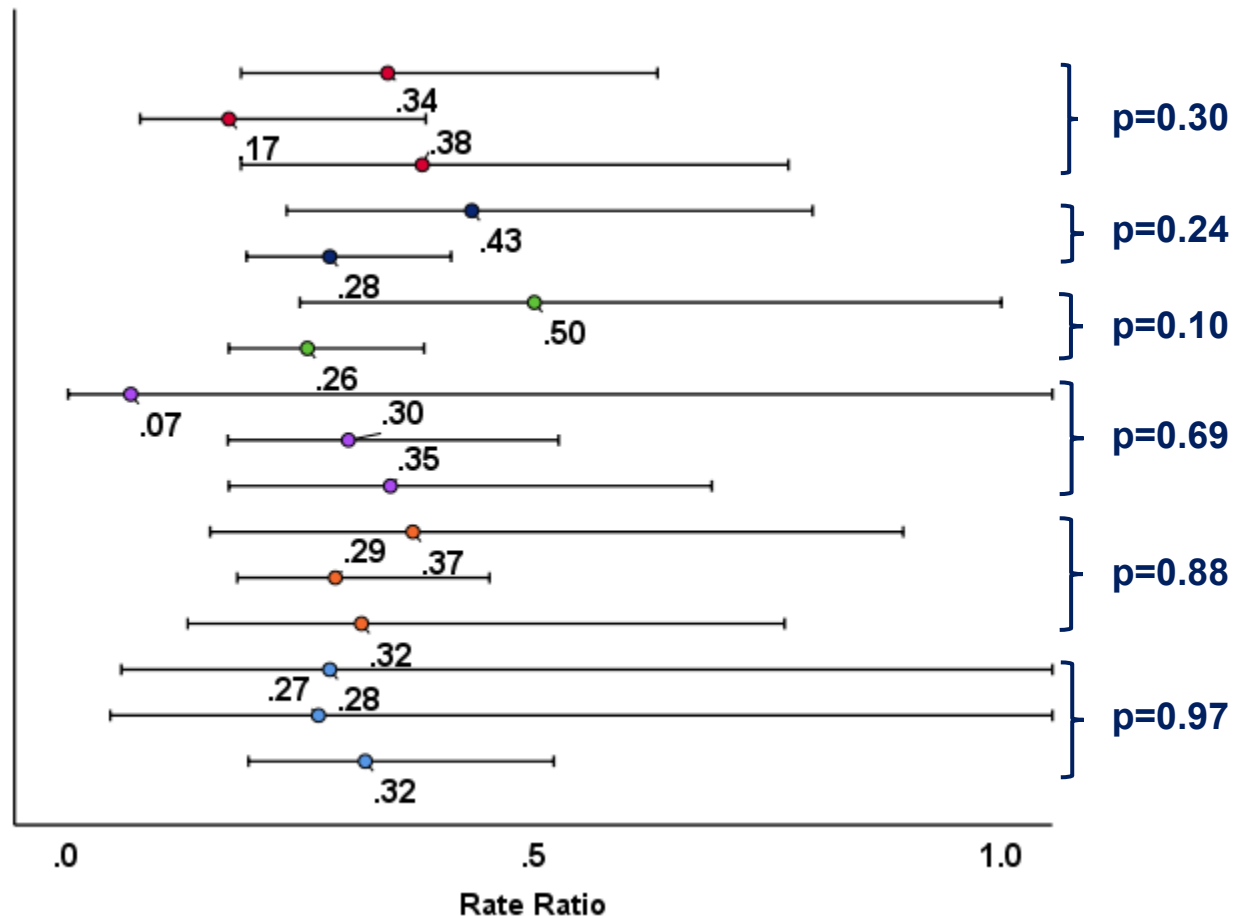
>10 years

Baseline HBeAg
status

Positive

Negative

Mix



First Clinical Event: Sensitivity Analysis

Treatment

Natural history

HBV Genotype

Treatment

Coinfection

Both

Follow-up in
HBsAg-loss cohort

A/D

B/C

Average follow-up

Yes

No

<50 person-years

50-999 person-years

>1000 person-years

Baseline HBeAg
status

<5 years

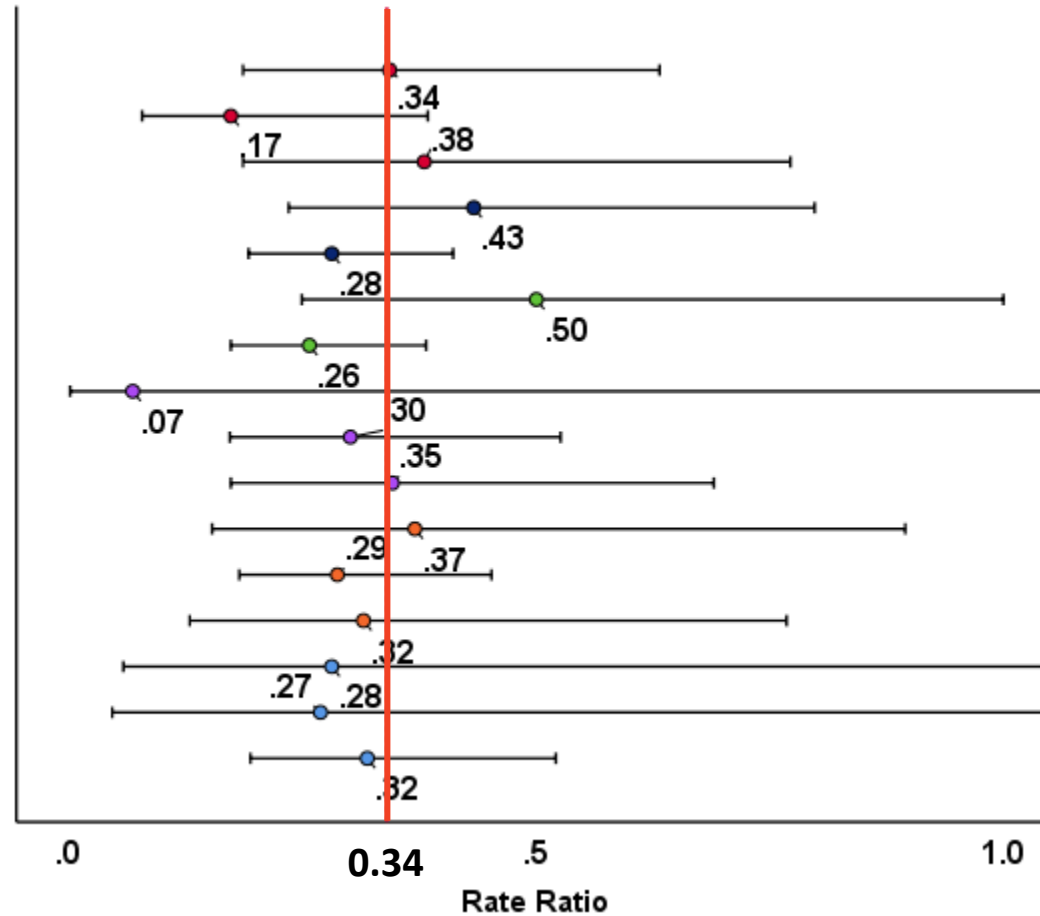
5-10 years

>10 years

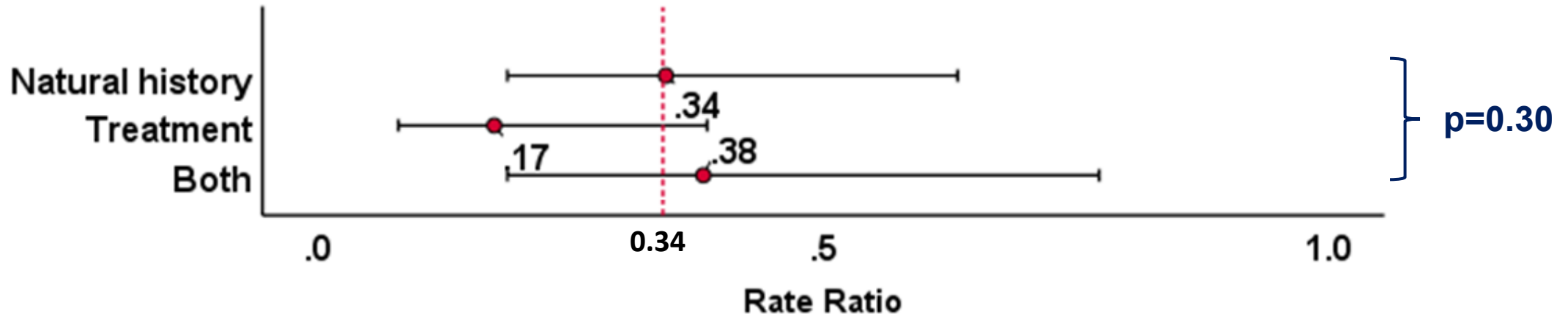
Positive

Negative

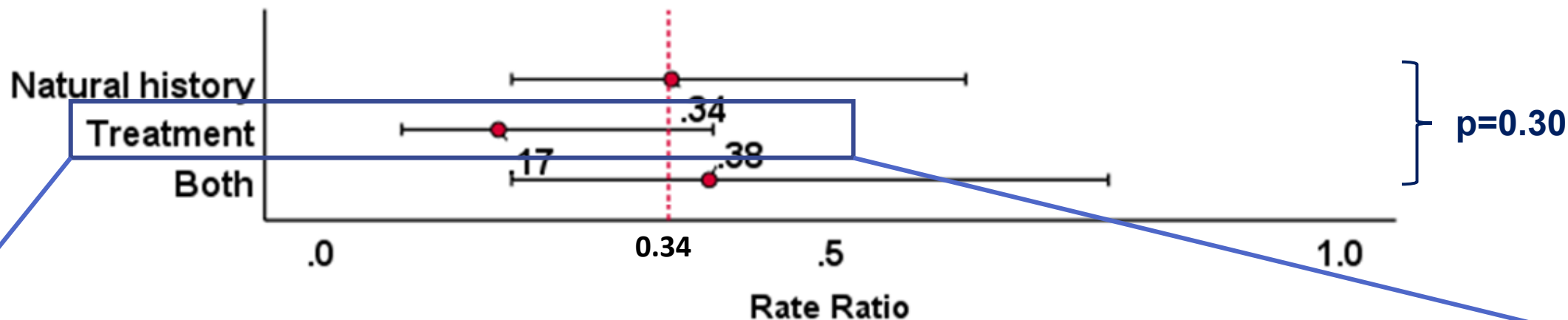
Mix




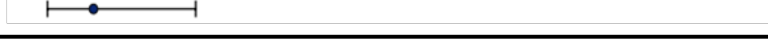


First Clinical Event: Treatment Subgroups



First Clinical Event: Treatment Studies Sub-analysis



Endpoint	RR	Lower limit	Upper limit	p-value	RR and 95% CI			
					0.0	0.5	1.0	1.5
IFN (n = 1)	0.121	0.006	2.527	0.173				
Nuc (n = 5)	0.185	0.079	0.433	0.000				
Nuc+IFN (n = 3)	0.102	0.004	2.873	0.180				
Overall (n = 9)	0.174	0.079	0.385	0.000				

Conclusion

- The results of our analysis support the use of HBsAg seroclearance as a surrogate marker for improved clinical outcome in CHB patients
 - Significantly reduced risk of adverse clinical events in patients who achieved HBsAg loss
 - No difference in incidence of long-term clinical outcome between patients who cleared HBsAg on-treatment and those who cleared HBsAg spontaneously

Next Steps

- We have reached out to corresponding authors for additional 15 articles that fulfilled our inclusion criteria, but did not differentiate endpoints based on HBsAg status
- Manuscript for peer review publication in preparation

Thank you!

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