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Meta-analysis: Insulin resistance and sustained virological response in hepatitis C

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The Editor: Alimentary pharmacology & therapeutic

Sevilla, Mayo 6th, 2011

Re: Meta-analysis: Insulin resistance and sustained virological response in hepatitis C.

Dear Editor,

Thanks for the acceptance of our manuscript. We have read, with interest, your minor modifications. In submitting this revised version of the manuscript, we have introduced the appropriate changes in the text (highlighted).

Yours sincerely,

Manuel Romero-Gómez, MD, PhD

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Meta-analysis: Insulin resistance and sustained virological response in hepatitis C.

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Abstract

Background&aim: Higher baseline homeostasis model assessment of insulin resistance (HOMA-IR) score predicted poorer sustained virological response (SVR) rate to peginterferon/ribavirin therapy in treatment-naïve chronic hepatitis C (CHC) patients. We conducted a meta-analysis to evaluate the impact of HOMA-IR on SVR in hepatitis C.

Methods: Relevant studies were identified by searching Medline and EMBASE. We identified 17 publications that addressed the influence of insulin resistance (IR) on SVR. The random effect model of Der Simonian and Laird method were used for heterogeneous studies using the Meta-Disc software 1.4, Madrid, Spain.

Results: Normal insulin sensitivity was associated with a higher rate of SVR (odds ratio (OR) 2.86 (95% CI: 1.97 to 4.16) in comparison with insulin resistance (IR). Moreover, in separate analysis by genotype selecting studies that used HOMA-IR>2 as cut-off defining IR, SVR was higher in patients with HOMAIR<2 in all genotypes: HCV-1, (O.R.:2.16(95%CI:1.51-3.08), HCV-2&3 (O.R.:3.06(95%CI:1.06-8.82) and HCV-4 (O.R.:6.65(95%CI:2.51-17.61). Studies reporting no association between HOMA and SVR included easy-to-cure cohorts, analyzed variables strongly related with IR like body mass index, steatosis, hyper γ GT, age and fibrosis, and reported differences in handling and interpretation of HOMA-IR.

Conclusion: Elevated HOMA-IR was associated with a lower cure rate of patients with hepatitis C treated with Peg-IFN- α /ribavirin irrespective of genotype, and the more difficult-to-treat cohort the better the HOMA-IR prediction. HOMA-IR is as a surrogate marker of IR susceptible to some biases derived from both handling and interpretation.

[Word count = 240]

Key words:

Hepatitis C virus, Insulin resistance, HOMA-IR, sustained virological response

Introduction

Worldwide approximately 200 million people are chronically infected with hepatitis C virus [¹]. Chronic HCV infection is characterized by a high rate of fibrosis progression, leading to cirrhosis and ultimately to hepatocellular carcinoma $[^{2,3}]$. Combination therapy with pegylated interferon alpha and ribavirin (Peg- FN/RBV) represents the current standard of care in chronic HCV $[^{4,5}]$. Despite the improvements in sustained virologic response (SVR) seen using combination of Peg- FN plus weightbased RBV, SVR rates in genotypes 1 and 4 populations have remained unsatisfactory with reported SVR rates of 28.9% to 54% [^{6,7}]. Even more unsatisfactory SVR rates on the order of 10% to 20% have been seen in difficult-to-treat populations such as African Americans, Hispanics, or insulin-resistant patients [⁶⁻⁸]. Studying the factors influencing SVR in patients with HCV treated with Peg- FN/RBV has been assiduously pursued over the past two decades. Now: there is a great interest by the importance of host factors such as fibrosis progression, genetic factors and metabolic disorders and these have been invoked to explain the reported differences in SVR $[^9]$. Insulin resistance (IR) is the main feature of metabolic syndrome and is characterised by hyperinsulinaemia in patients with normal fasting glucose. It has been associated with increased risk of developing diabetes mellitus (T2DM), cardiovascular disease and non-alcoholic fatty liver disease [¹⁰]. Accumulating large body of literature, including molecular, pathological, epidemiological, randomised controlled trials and observational studies, has highlighted the interaction between HCV and glucose metabolism (IR/T2DM) [^{11,12,13,14,15,16}]. Although, earlier works refereed to that a higher baseline homeostasis

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model assessment of insulin resistance (HOMA-IR) score predicted a poorer SVR rate to Peg-IFN- α /ribavirin in treatment-naïve chronic hepatitis C patients [^{8'17,18,19,20}]; this data was not universally confirmed by all studies [²¹]. In the view of the suboptimal response to the current standard in therapy; these postulated correlation between IR and SVR has gain a great importance, as one of the modifiable risk factor. Therefore, we performed a systematic review and meta-analysis to review the literature systematically to evaluate, quantify, and summarize the association between HOMA and SVR.

Methods

Study Selection Criteria

Inclusion and exclusion criteria were defined prior to commencement of the literature search. Studies were included if they met the following criteria: they included treatment-naive HCV patients aged older than 18 years regardless HCV genotype or the ethnic group; they provided information on SVR rates; and they were reported in English as full papers. Studies were excluded if they met the following criteria: they did not provide information on SVR; they did not allow correlation between the HOMA-IR and SVR; they included patients who were treated with nonpegylated interferons; they included patients with additional antivirals to Peg- FN/RBV (eg, amantadine); they included patients who were not treatment-naive; they included human immunodeficiency virus (HIV) coinfected patients; they included liver transplant recipients; letters/case reports, studies enrolling fewer than 10 subjects, or articles not reporting outcomes of interest or primary data (editorials, reviews).

Literature Search, Study Selection, and Data Extraction

We searched MEDLINE and EMBASE for studies published in English through December 2010 using a combination of the following terms: "Hepatitis C"[Mesh] OR "Hepacivirus"[Mesh] OR "Hepatitis C, Chronic"[Mesh]) AND "Insulin

resistance"[Mesh] OR "Homeostasis Model of Assessment"[Mesh] AND "Sustained virological Response"[Mesh]. Manual search of cited bibliographies was also performed. Only fully published articles were considered. Two researchers independently performed the literature search and data abstraction with regard to the inclusion and exclusion criteria by reading titles and abstracts. When reading titles and abstracts did not allow identification of eligible studies, articles were read in full. Relevant reviews and letters to the editor were excluded from the analysis, but read in full in order to identify potential relevant original studies. Disagreements between the 2 observers were resolved by discussion. Data from selected studies were extracted in a data sheet by MRG author and checked by a second author (ME). Disagreements could be resolved by discussion. The following data were extracted: year of publication, number of patients, cutoff of HOMA-IR to define IR, method of measurement of insulin, percentage of patients with normal insulin sensitivity and percentage of patients with advanced fibrosis.

End Point

The main endpoint was the SVR rate, defined as the percentage of patients with undetectable HCVRNA (<50 IU/ mL) 24 weeks after treatment completion of therapy.

Statistical analysis

Statistical analysis was performed by using the Meta-Disc software 1.4 (Zamora J, et al., BMC Medical Research Methodology 2006, 6:31.) has been used, considering: i) a summary of data from individual studies, ii) an investigation of the studies homogeneity graphical and statistically, iii) calculation of clustered indexes and iv) exploration of heterogeneity. Our assumption of heterogeneity was tested for each planned analysis using the Cochran-Q heterogeneity test. Random effects model by Der Simonian and Laird method was considered when heterogeneity was found. Only 2-sided tests with a

significance level of 0.05 were used. Confidence intervals (CIs) of individual studies were determined or approximated from the available data.

Results

Seventeen eligible studies fulfilled the inclusion and exclusion criteria are enrolled in the analysis (^{8,17-21,22,23,24,25,26,27,33,38,43}), three studies without SVR rate according to baseline HOMA-IR were further excluded (Figure 1). The characteristics of these studies are shown in Table 1. Five studies were from Italy, 3 from Japan, 2 from Taiwan, 1 from Spain, United Kingdom, USA, France, India and Egypt. The used cutoff values for diagnosis of IR ranged from HOMA>2 in 10 studies to HOMA > 2.5 in 1 study, HOMA>2.7 in 1 study, HOMA>3 in 1 study and HOMA>3.5 in 1 study and was not available data in three studies. Nine studies found a statistically a significant association between HOMA-IR and SVR, and the other 5 studies did not find a significant association between HOMA-IR and SVR (Figure 2 and Table 1). Three of enrolled studies did not show SVR rate according to HOMA-IR. Thus, pooled data included 2129 patients in 14 studies. The meta-analysis including all 14 studies demonstrated that insulin resistance was associated with a remarkably lower likelihood of SVR, using random effects model by Der Simonian and Laird method, odds ratio was 2.86 (95% CI: 1.97 to 4.16) regardless of genotype (Figure 2). The test of heterogeneity (Cochran-Q=36.92; df=13; p=0.0004) inconsistency $I^2 = 64.8\%$ and Tau-squared = 0.287. Sub-analysis including patients segregated by genotype 1, 2/3 or 4 demonstrated a negative and significant impact of increased HOMA-IR on SVR; three studies mixed genotypes were included and were separately analyzed (17, 21, 39) (Table 2, Figure 3, 4, 5). Even with limiting the analysis to include only studies using cut-off HOMA > 2for defining IR, the impact on SVR was maintained. Patients with HOMA-IR >2 had significantly lower rates of SVR compared with patients with HOMA-IR < 2, and this

difference was largely independent of HCV genotype: HCV-1 (O.R.:2.16;95%CI:1.51-3.08), HCV-2&3 (O.R.:3.06;95%CI:1.06-8.82) and HCV-4 (O.R.:6.65;95%CI:2.51-17.61). The difference in SVR rate in patients with IR compared to those with insulin sensitivity ranged from 3% to 41% in genotype 1, 5% to 57% in genotype 2/3 and 38% to 48% in genotype 4. The risk estimate for not achieving SVR from 10 studies using HOMA-IR > 2 as cutoff value for diagnosis of IR was 3.1 fold. Similarly, the pooled estimates from 11 studies enrolled patients infected with HCV genotype 1 indicate a statistically significant 2.16-fold decrease in the incidence of SVR among patients with IR. Some studies that found no association were characterized by including cohorts of patients with a mean HOMA-IR <3 and without advanced fibrosis. In these studies; multivariate analysis demonstrated independent variables associated with SVR strongly related to metabolic abnormalities showing a tight co-linearity with HOMA-IR like body mass index, steatosis, elevated γ -glutamyl-transpeptidase (GGT), age and fibrosis.

This meta-analysis of published studies indicates that IR assessed by HOMA-IR is associated with an approximately three fold increase in risk of failure to achieve SVR to Peginterferon+ribavirin in treatment naive patients with hepatitis C regardless of viral genotype. The discrepancy between the results of the cohorts in this review may be refereed to fact that the interaction between IR and SVR are very complex depends on numerous factors that need to be considered when interpreting the data: a) measurement and interpretation of HOMA-IR; b) colinearity between HOMA-IR and variables associated with metabolic derangements; and c) the characteristics of the cohort under study. HOMA-IR which was developed by Matthews et al [²⁸] as an indirect surrogate marker of IR hampered by many concerns, which may influence the extracted data. These factors includes; methods of measurement of insulin, handling of blood samples,

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the conversion factor between SI units and mU/ml, the formula to calculate HOMA-IR and the threshold to define IR. For example: a) Insulin assays are not standardized and are not comparable between studies. Insulin inter-assay variations can be large, and values have varied considerably between different laboratories $[^{29,30}]$ b) careful handling of the blood samples is essential because hemolysis results in the degradation of insulin, while freezing the samples results in the degradation of C-peptide and glucose, with false-negative results being one of the consequences. c) Insulin should be reported in International Units (SI), but the conversion factor from microUnits/mL to SI units is not uniform among manufacturers; values ranging between 6 and 7.15 with no real scientific justification $[^{31}]$. d) HOMA-IR could be calculated using the formula from Matthews al. et or using the HOMA-2 calculator (available at http://www.dtu.ox.ac.uk/homacalculator/). However, results from both methods are quite different and could bias the final analysis in comparative studies. e) Since insulin secretion is pulsatile, the use of a single fasting blood sample to diagnose IR in CHC, although convenient, potentially underestimates the prevalence and degree of this disturbance. The mean value from three samples taken at 5-min intervals to compute HOMA is, theoretically, better than a single sample determination $[^{22}]$. However, in practice a single sample is often taken and, if population estimators are found, this is an acceptable compromise that yields a similar result for large datasets. No evidence is available in setting of CHC patients. f) Ultimately; till now; there is no international consensus regarding the threshold that defines IR by using HOMA-IR. Recently, Lam et al. showed that HOMA-IR >4 was the optimal value defining IR in hepatitis C setting when compared with steady-state plasma glucose $[^{32}]$. However, these data need to be further confirmed in different populations, using euglycaemic hyperinsulinaemic clamp as gold-standard, and taking ethnicity and obesity into account when evaluating changes

in HOMA-IR over time $[^{26}]$. Second; HOMA-IR shows a strong colinearity, with several variables implicated in metabolic disturbances such as obesity, hepatocyte steatosis, elevated GGT, fibrosis and older age. All these variables provide information, at least in part, on the metabolic abnormalities occurring in the individual, the final outcome of the multivariate analysis depending on the interactions between them. Interestingly in our review to the negative studies that failed to find an association between IR and SVR; the mean baseline HOMA-IR was <3 in all studies, and the prevalence of advanced fibrosis or cirrhosis was also lower or even these studies had excluded cirrhosis or had been under-powered. This observation supporting the idea that the more difficult-to-treat cohort the better the HOMA-IR prediction. In one of the negative studies, when HOMA-IR was excluded from the multivariate model, one of these variables emerged as independently associated with SVR. This highlights this type of statistical bias is very difficult to resolve when strongly related variables were included in the analysis. Indeed, Dai et al $\begin{bmatrix} 17 \end{bmatrix}$ demonstrated that the association of HOMAIR with SVR was stronger in cohorts including very difficult- to-treat patients. Ultimately; when more sophisticated methods to measure IR (such as whole-body insulin sensitivity) were used, a stronger association was reported between IR and SVR [³³]. Thirdly; several viral and host factors have an impact on response to treatment $[^{34,35}]$, these factors may influence the final outcome and explain the difference of results of these cohorts. Further study is needed to clarify the interaction of genotype, the degree of IR, gender, and genetics (such as the IL28ß mutation) and viral kinetics. This Meta-analysis has undoubtedly some limitations, since there is heterogeneity in the enrolled studies including characteristics cohort, methods for insulin measurement and HOMA-IR interpretation may have influenced on the concluded results.

In conclusion, this meta-analysis provides evidence that impaired HOMA-IR was associated with a lower cure rate of patients with hepatitis C treated with Peg- FN/RBV. The differences are greater in cohorts of patients who are difficult to cure (higher insulin resistance, higher prevalence of advanced fibrosis). The inherent limitations in the determination and calculation of HOMA-IR may explain, at least in part, the variability detected in the results. Better handling and interpretation of HOMA are mandatory until the development of new tools for assessment of insulin resistance.

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Table 1: studies included in meta-analysis addressing the impact of HOMA on

sustained virological response

Table 1: studies included in meta-analysis addressing the impact of HOMA on sustained virological response

Author	year	HOMA Cut-off	SVR association	Insulin method	mean HOMA-IR (SD)	% patients with IS	Genotype	n	SVR in IS	non-SVR in IS	SVR in IR	non-SVR in IR	Delta %SVR IS - IR	O.R. (95%CI)
Akuta et al	2009	3.5	NO	NA	2.6 (2.6)	0,28	1	123	11	9	26	26	0,05	0.82(0.29-2.33)
Bortoletto et al	2010	3	YES	EIA	1.9 (1.1)	0,61	1	36	8	14	2	12	0,22	0.29(0.051-1.49)
Camma et al (*)	2006	NA	NO	NA	2.65 (2.01)		1	291	NA	NA	NA	NA		NA
Chu et al.	2008	2	YES	EIA	2.93 (0.14)	0,31	1	133	38	3	50	42	0,38	0.14(0.005-0.54)
Conjeevaram et al	2007	2	YES	NA	3.5 (5.0)	0,33	1	399	65	68	96	170	0,13	0.87(0.77-0.99)
Dai et al.	2009	2.5	YES	RIA	2.24 (2.46)	0,71	1	150	75	31	22	22	0,21	0.87(0.75-0.97)
Fattovich et al	2010	2	NO	ECLIA	2.7 (2.5)	0,44	1	181	37	43	44	57	0,03	0.9(0.5-1.64)
Grasso et al	2009	2	NO	EIA	2.6 (2.1)	0,77	1	90	34	35	8	13	0,11	0.63(0.23-1.72)
Miyaaki et al	2009	2	NO	NA	1.9 (0.87)	0,51	1	39	11	9	6	13	0,23	0.37(0.1-1.38)
Mizuta et al	2010	2	YES	NA	1.92 (1.92)	0,51	1	51	17	9	6	19	0,41	0.17(0.049-0.57)
Petta et al.	2009	2.7	NO	EIA	2.73 (1.69)	0,65	1	83	32	22	14	15	0,11	0.64(0.17-3.03)
Romero-Gomez et al.	2005	2	YES	ECLIA	3.01 (2.67)	0,38	1	113	26	17	23	47	0,28	0.55(0.33-0.93)
Walsh et al (*)	2006	NA	NO	NA	2.02 (0.19)		1	66	NA	NA	NA	NA		NA
Bortoletto et al	2010	3	YES	EIA	1.9 (1.1)	0,67	2&3	21	14	0	3	4	0,57	NC
Dai et al.	2009	2.5	NO	RIA	2.24 (2.46)	0,71	2&3	180	117	11	44	8	0,07	0.87(0.75-0.97)
Fattovich et al	2010	2	NO	ECLIA	2.7 (2.5)	0,57	2&3	209	93	26	66	24	0,05	0.8(0.42-1.54)
Poustchi et al.	2008	2	YES	RIA	3.34 (3.34)	0,41	2&3	82	32	2	31	17	0,30	0.15(0.03-0.77)
Walsh et al (*)	2006	NA	NO	NA	2.02 (0.19)		2&3	79	NA	NA	NA	NA		NA
Khattab et al.	2010	2	YES	ECLIA	2.82 (1.19)	0,40	4	131	47	6	32	46	0,48	0.07(0.01-0.43)
Moucari et al	2009	2	YES	ECLIA	3.7 (4.0)	0,49	4	108	39	20	14	35	0,38	0.19(0.07-0.51)
(*) Not included in meta-analysis due to lack of data HOMA: Homeostasis Model of Assessment						2565								
SVR: Sustained virologica	l response													
SD: Standard deviation														
O.R.: Odds ratio														
IS: Insulin sensitivity (HOI	MA under o	cut-off)												
E: Insulin resistance (HOMA above cut-off)														
NA: Non-available														
CCLIA: Electrochemiluminescence Immunoassay														
EIA:Enzyme Immunoassay														
RIA: Radio Immuno Assay														
NC: Calculation not allow	ed													

Table 2: Random effect Model for pooled data according to genotypes.

Genotype	Studies	Pooled data	Heterogeneity	Inconsistency			
	number	O.R.(95%CI)	(Cochran-Q)	inconsistency			
All	14	0.35 (0.24- 0.51)	36.92	I ² =64.8%			
		0.00 (0.21 0.01)	p=0.0004				
Genotype 1	11	0.46 (0.32 – 0.66)	18.05	I ² =44.6%			
			p=0.05	1 - 1 10 /0			
Genotype	4	0.33 (0.11-0.94)	8.78	I ² =65.8%			
2&3	4	0.55 (0.11-0.94)	p=0.03	1 -05.070			
Genotype 4	2	0.15 (0.05 - 0.40)	2.35	$I^2 = 57.4\%$			
ounotype i			p=0.12	1 0/11/0			
HOMA cut-	10	0.33 (0.20 - 0.53)	35.9	$I^2 = 72.1\%$			
off 2	10	0.55 (0.20 - 0.55)	P=0.0001	1 - 72.170			
			0,				

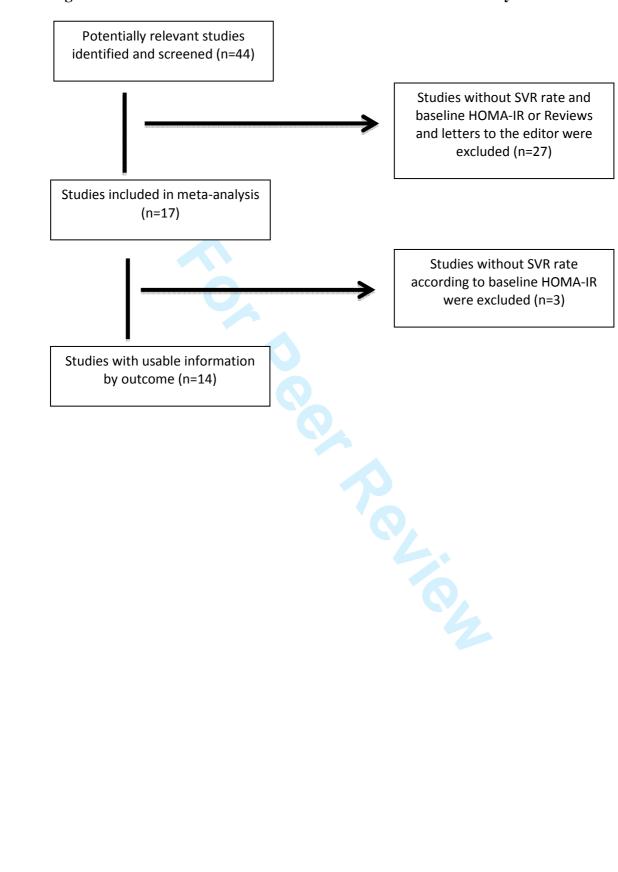
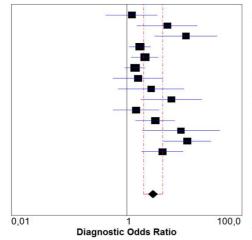


Figure 1. Chart flow of studies screened and included in meta-analysis

Figure 2. Odd Ratios and 95% CI for the association between insulin resistances assessed by HOMA-IR and sustained virological response to antiviral therapy in chronic hepatitis C patients treated with interferon + ribavirin in 14 studies. Black boxes indicate point estimates for OR.



Akuta et al Bortoletto et al Chu et al
Conjeevaram et al
Dai et al
Fattovich et al
Grasso et al
Mayaaki et al
Mizuta et al
Petta et al
Romero-Gómez et al Poustchi et al
Khattab et al
Moucari et al

Random Effects Model Pooled Diagnostic Odds Ratio = 2,86 (1,97 to 4,16) Cochran-Q = 36,92; df = 13 (p = 0,0004) Inconsistency (I-square) = 64,8 % Tau-squared = 0,2877

Diagnostic OR (95% CI)

1,22 (0,43 - 3,44) 5,03 (1,50 - 16,82) 10,64 (3,06 - 36,95)

(1,11 - 2,58)

(1,20 - 3,59) (0,92 - 2,09) (0,58 - 4,29)

(0,72 - 9,80)

(1,76 - 20,32)

(0,58 - 3,64)

(1,42 - 6,88) (1,87 - 41,18)

4,18 (1,82 - 9,61)

(4,30 - 29,47)

1,69

2,08

1,39 1,58

2,65 5,98

1,45

3,13 8,77

11,26



Figure 3. Odd Ratios and 95% CI for the association between insulin resistances assessed by HOMA-IR and sustained virological response to antiviral therapy in genotype 1 chronic hepatitis C patients treated with interferon + ribavirin in 11 studies. Black boxes indicate point estimates for OR

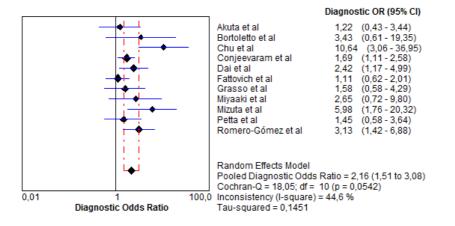


Figure 4. Odd Ratios and 95% CI for the association between insulin resistances assessed by HOMA-IR and sustained virological response to antiviral therapy in genotype 2 and 3 chronic hepatitis C patients treated with interferon + ribavirin in 4 studies. Black boxes indicate point estimates for OR

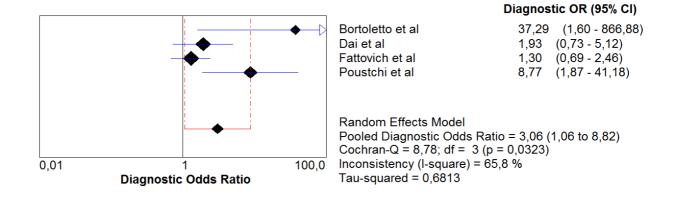


Figure 5. Odd Ratios and 95% CI for the association between insulin resistances assessed by HOMA-IR and sustained virological response to antiviral therapy in genotype 4 chronic hepatitis C patients treated with interferon + ribavirin in 2 studies. Black boxes indicate point estimates for OR

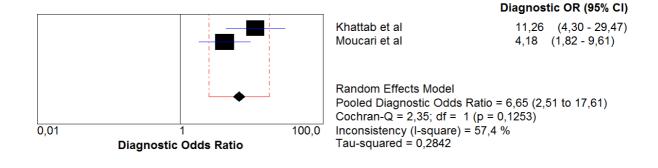
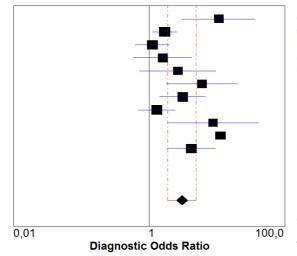


Figure 6. Odd Ratios and 95% CI for the association between insulin resistances assessed by HOMA-IR and sustained virological response to antiviral therapy in chronic hepatitis C patients treated with interferon + ribavirin in 10 studies using HOMA-IR > 2 as cut-off defining insulin resistance. Black boxes indicate point estimates for OR



	Diagnostic OR (95% C				
Chu et al	10,64 (3,06 - 36,9	5)			
Conjeevaram et al	1,69 (1,11 - 2,58)				
Fattovich et al	1,11 (0,62 - 2,01)				
Grasso et al	1,58 (0,58 - 4,29)				
Mayaaki et al	2,65 (0,72 - 9,80)				
Mizuta et al	5,98 (1,76 - 20,32)			
Romero-Gómez et al	3,13 (1,42 - 6,88)				
Fattovich et al	1,30 (0,69 - 2,46)				
Poustchi et al	8,77 (1,87 - 41,18))			
Khattab et al	11,26 (4,30 - 29,4	7)			
Moucari et al	4,18 (1,82 - 9,61)				

Random Effects Model Pooled Diagnostic Odds Ratio = 3,05 (1,88 to 4,93) Cochran-Q = 35,90; df = 10 (p = 0,0001) Inconsistency (I-square) = 72,1 % Tau-squared = 0,4363

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