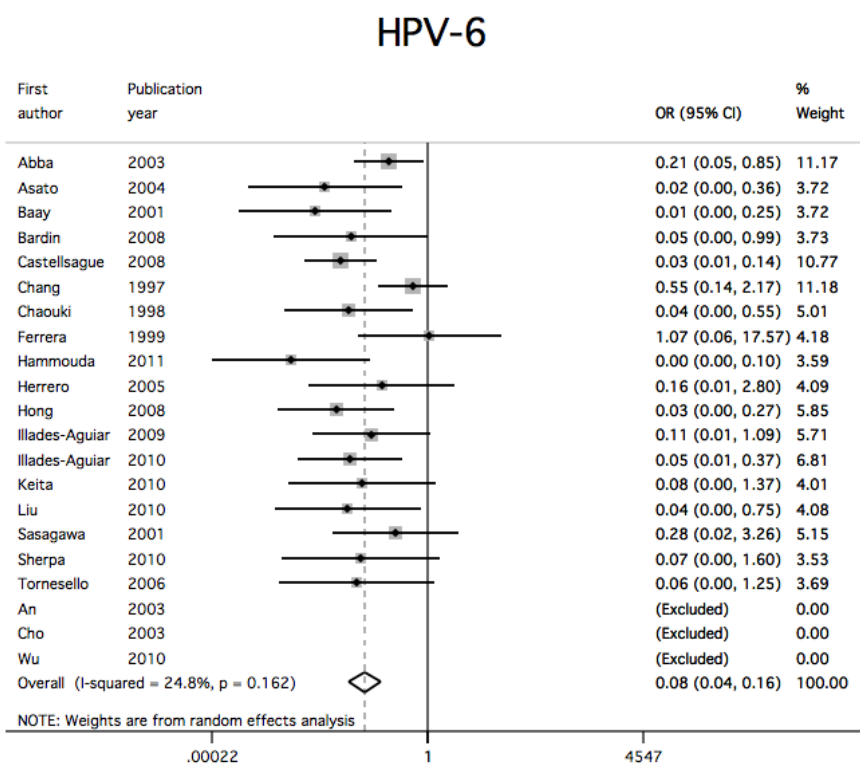
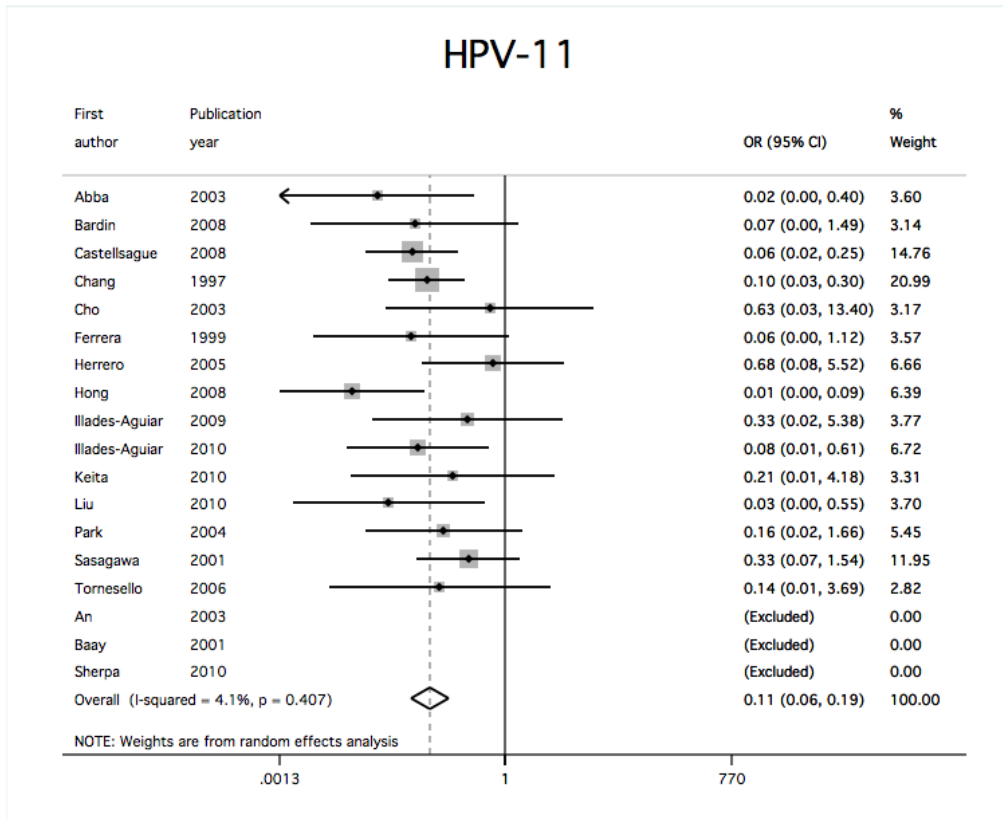


**Additional File 2. Meta-analyses assessing the relative oncogenic potential of each human papillomavirus (HPV) genotype (forest plots).** Studies are listed in alphabetical order. Each study is represented by a black cross, which corresponds to the odds ratio (OR) point estimate; a grey square, whose area reflects the weight each study contributes in the meta-analysis; and a horizontal line, which spans the 95% confidence interval (CI). The diamond at the bottom of the graph represents the combined OR and its 95% CI. The solid vertical line is an oncogenic potential equal to that of HPV-16 (OR 1.0) and the dotted vertical line indicates the value of the combined ORs from the random-effects model. The graphs were generated by Stata command metan.

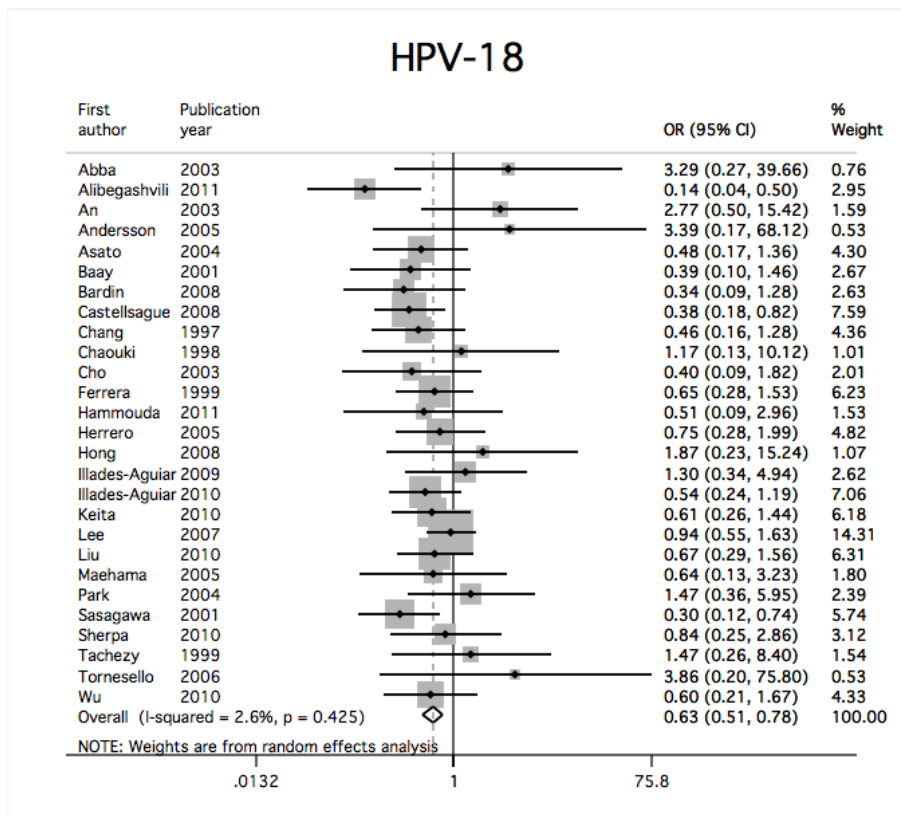
(a)



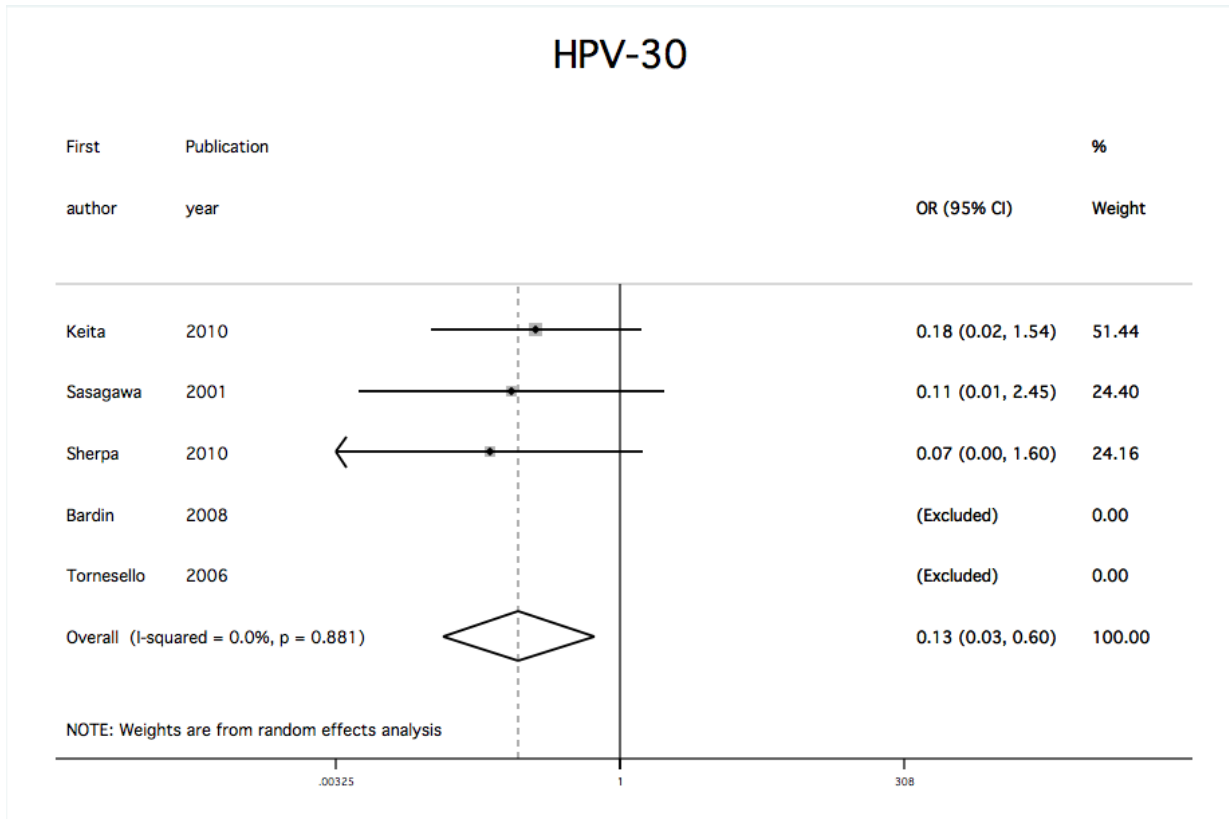
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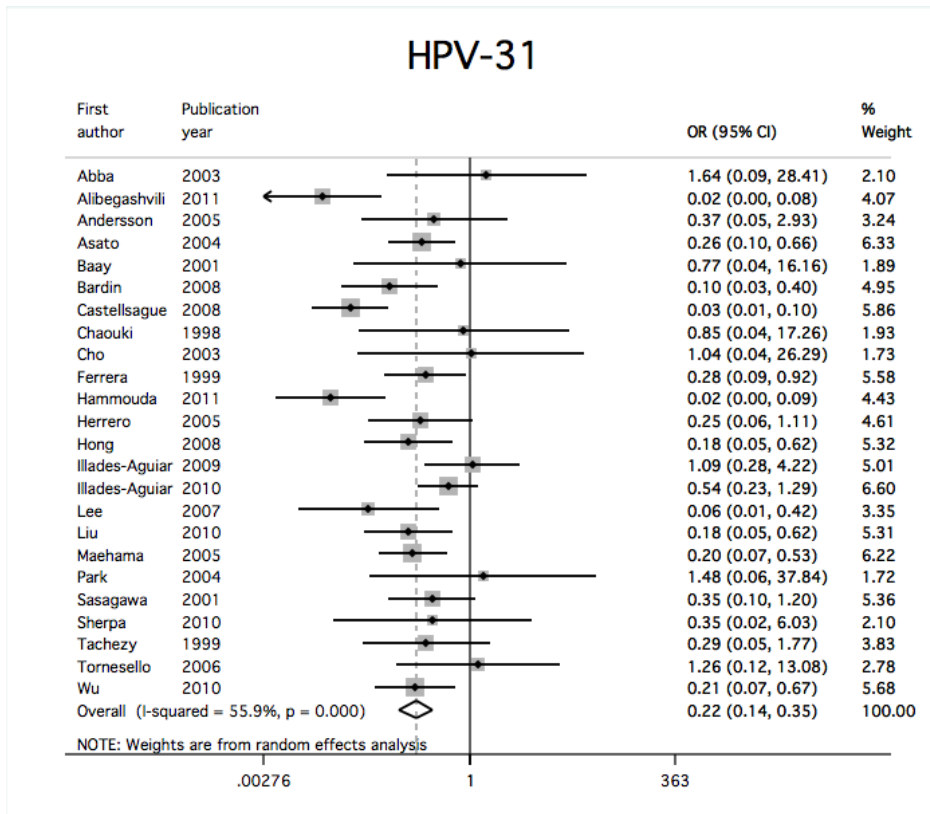
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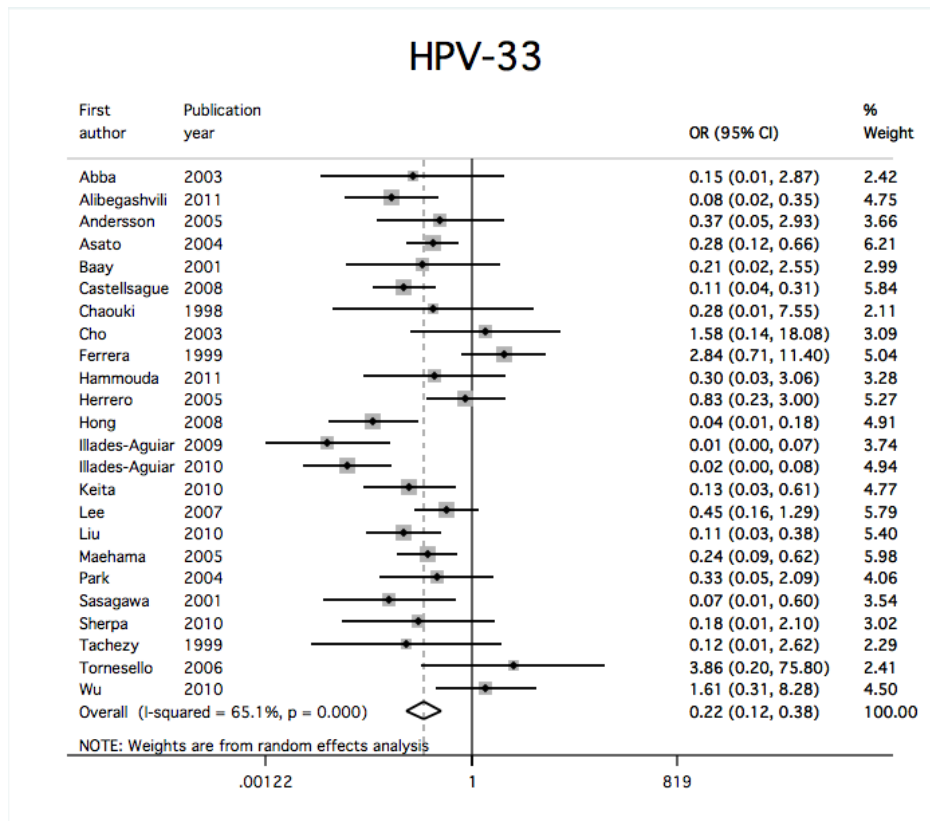
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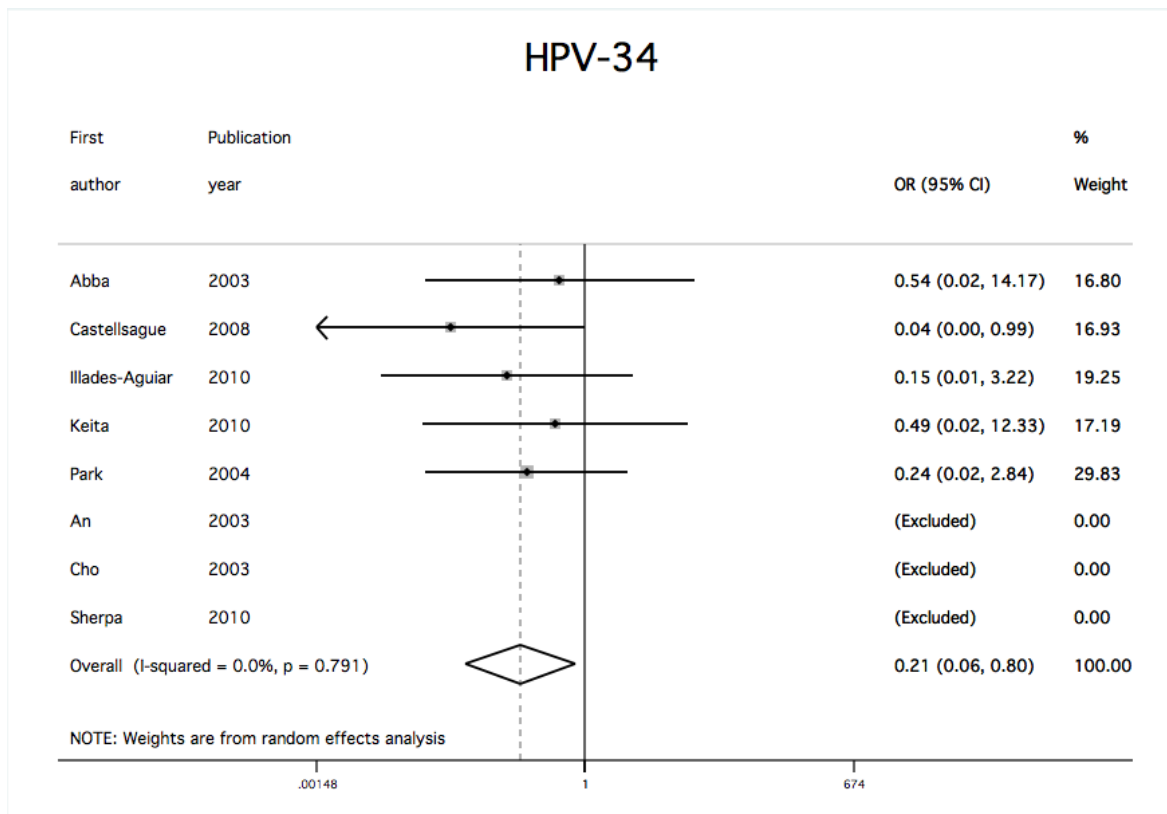
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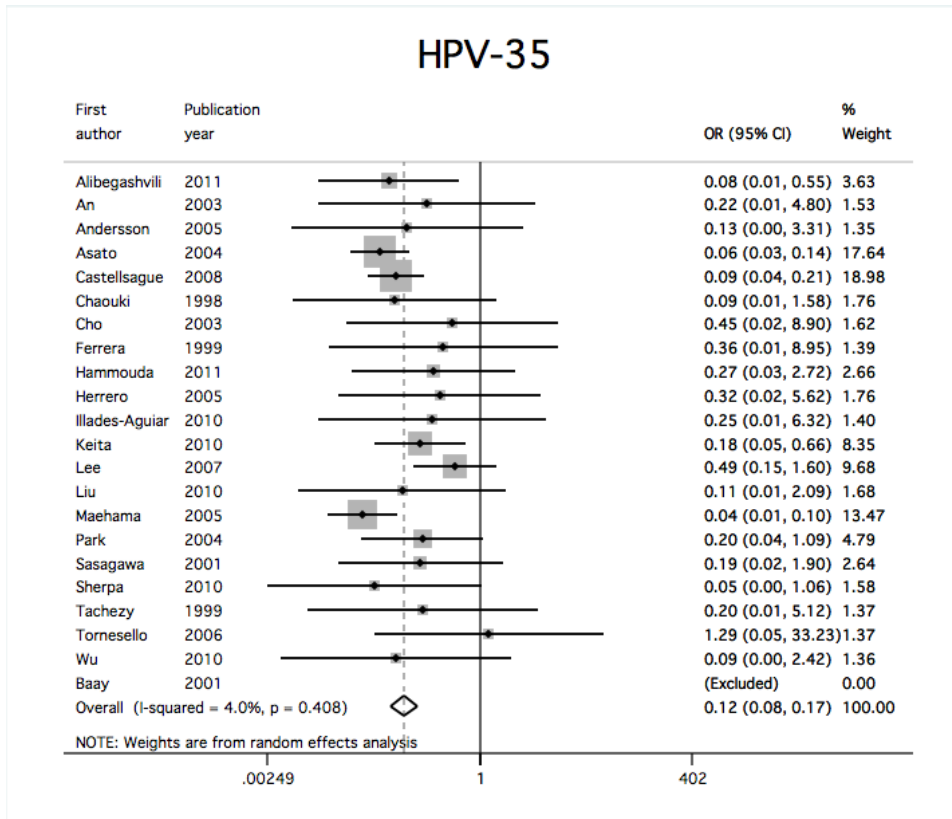
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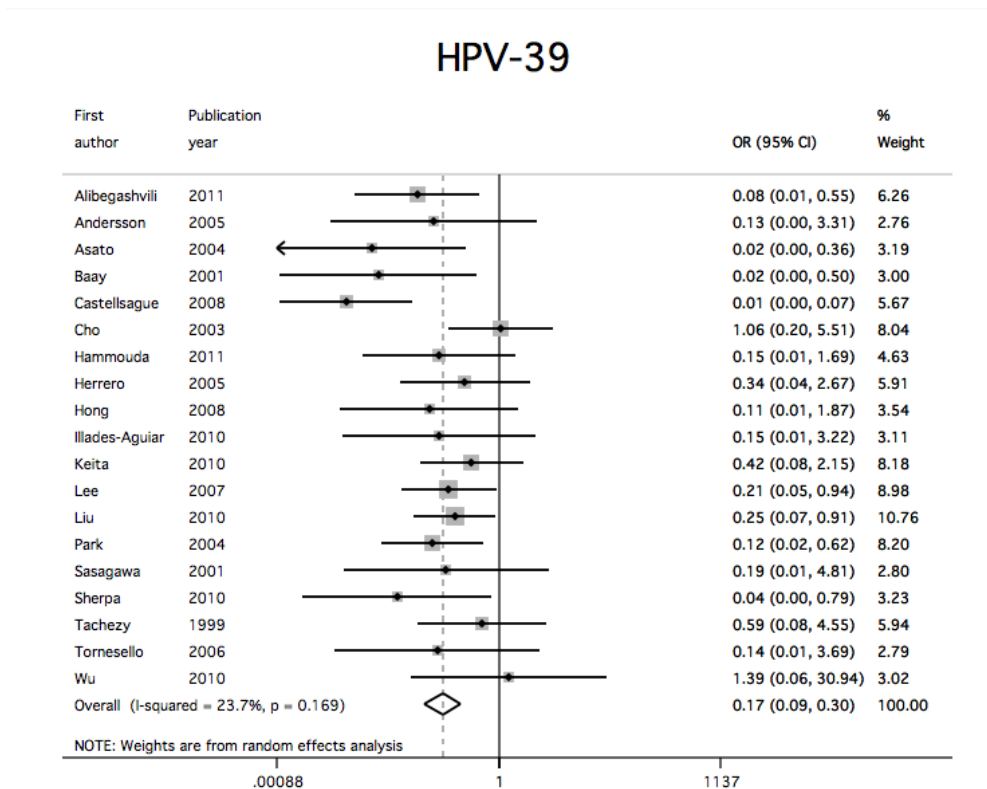
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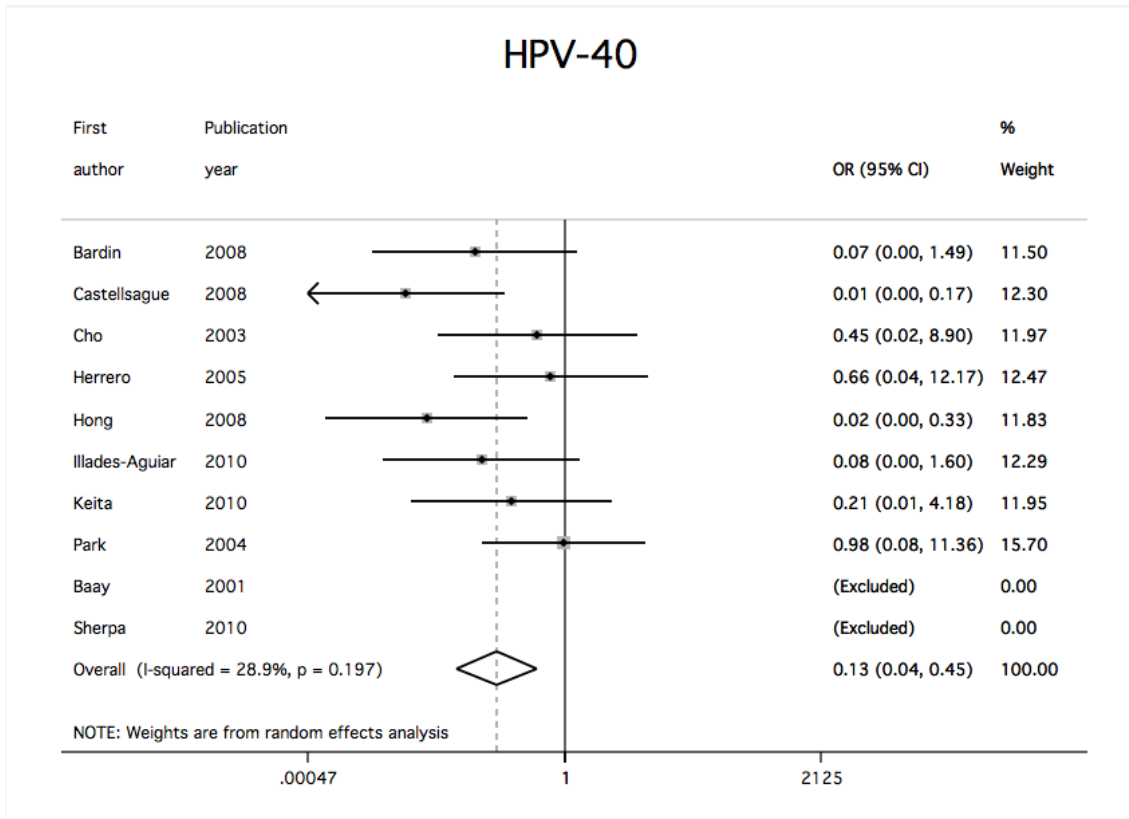
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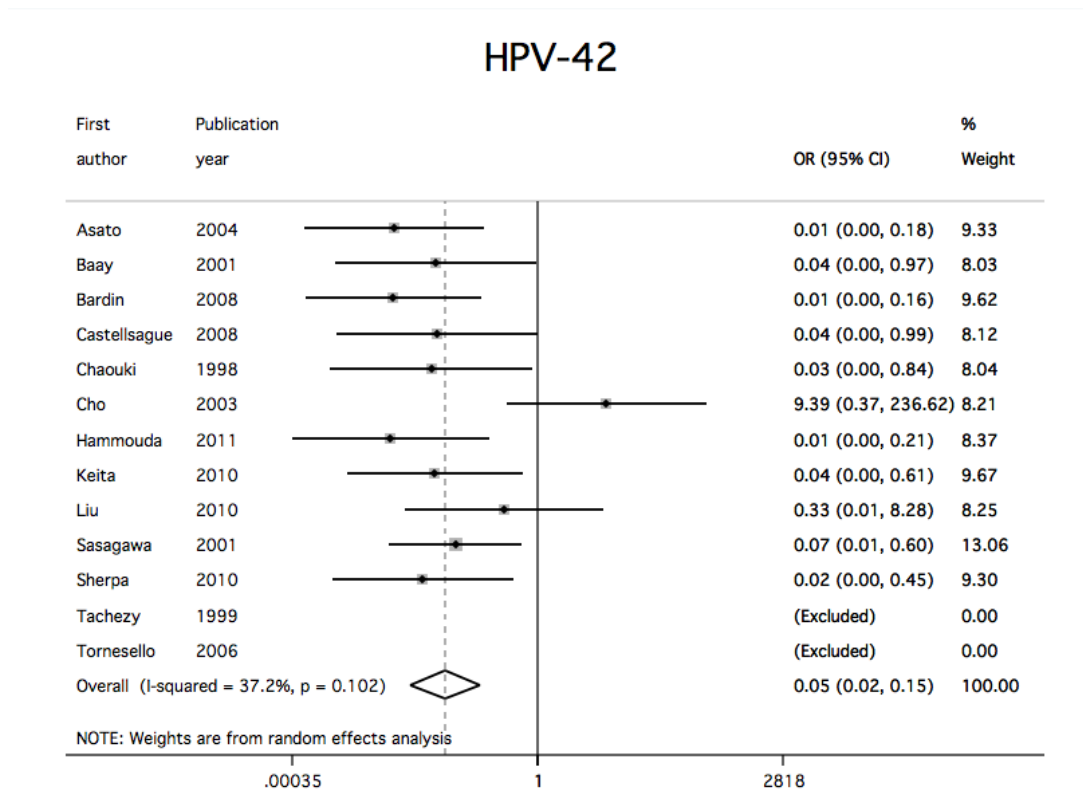
(i)



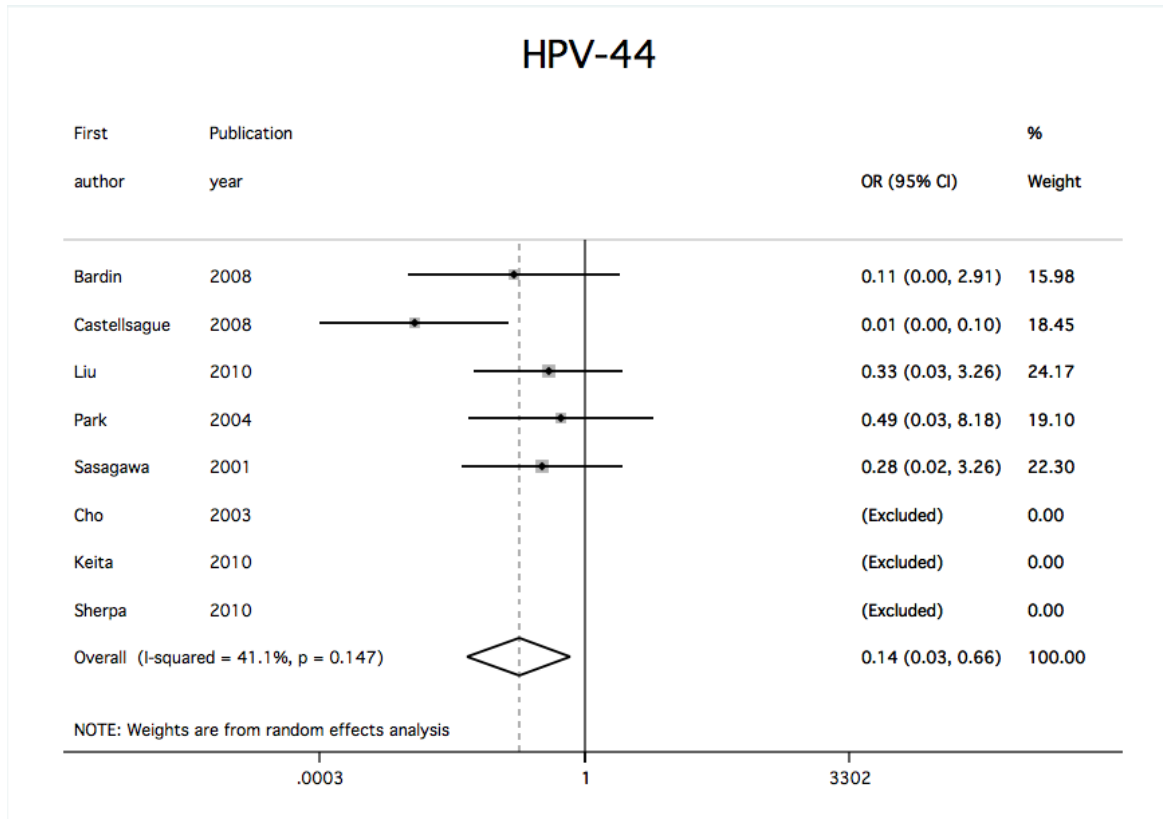
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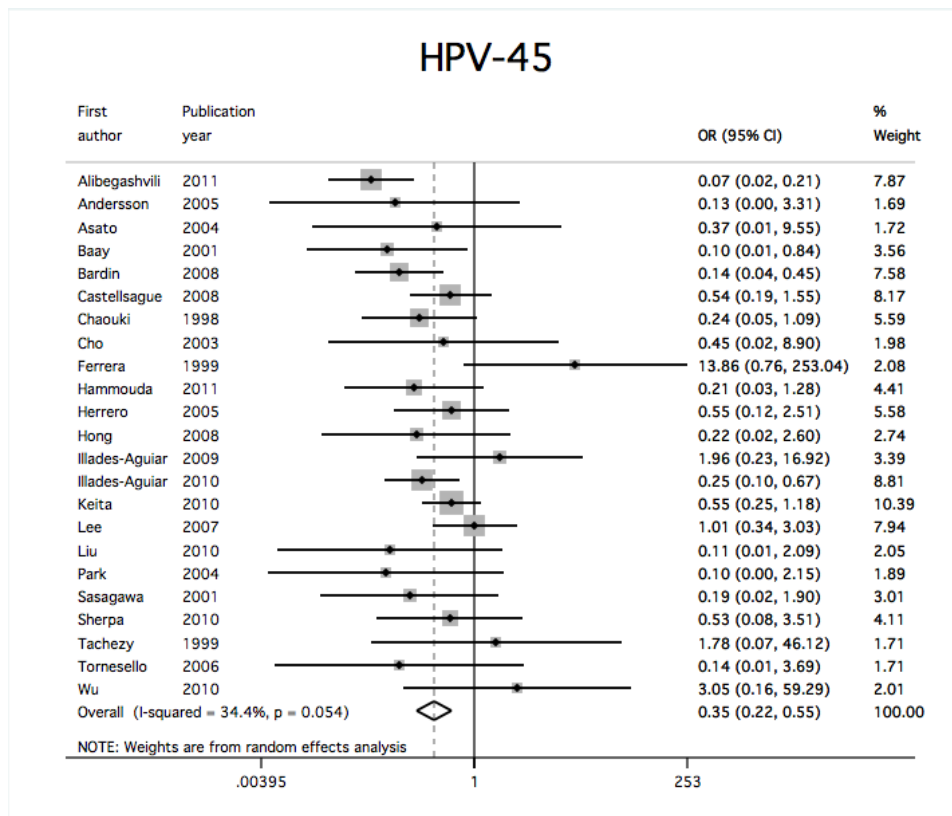
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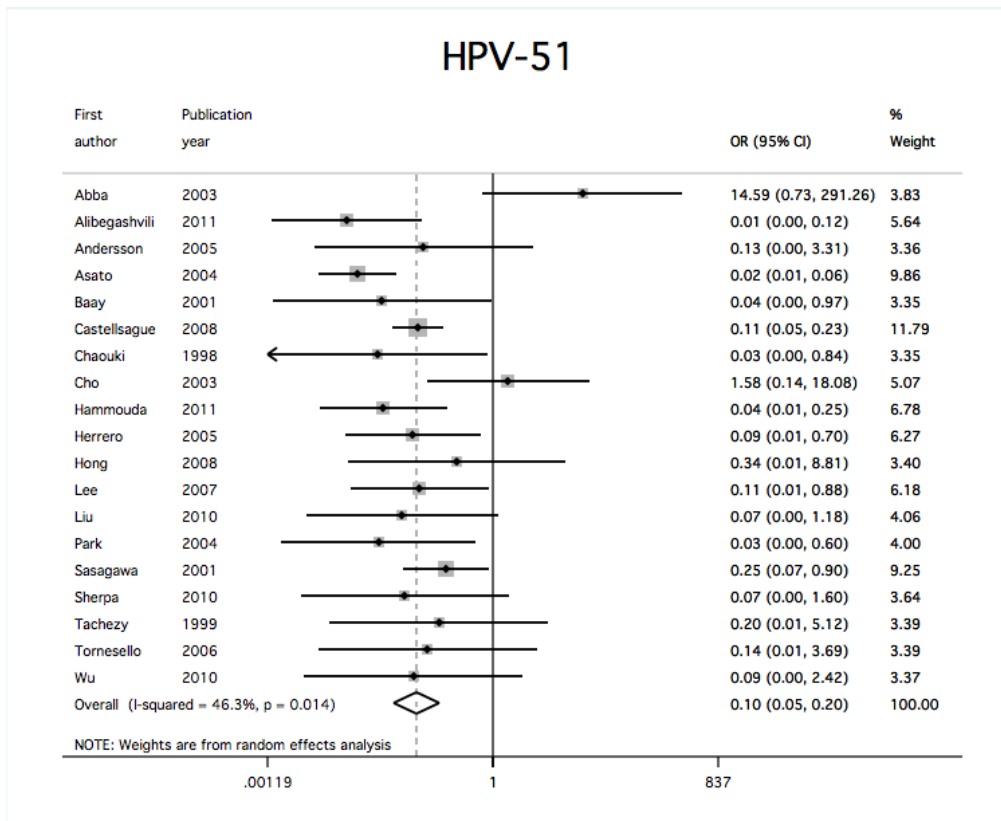
(1)



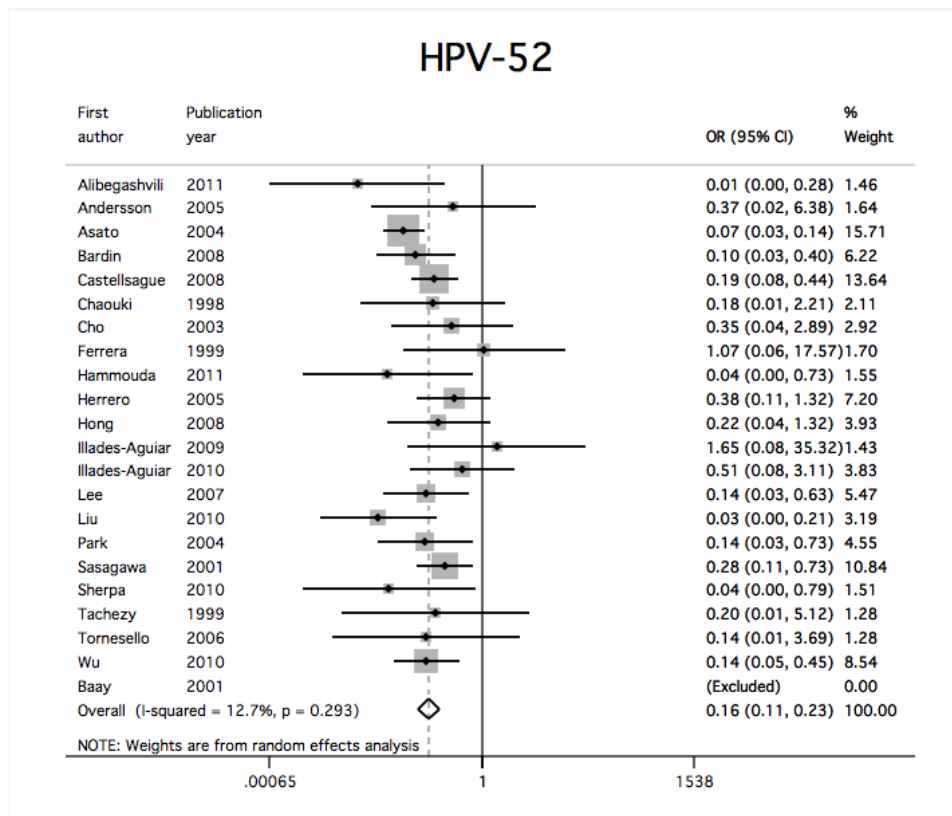
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(n)

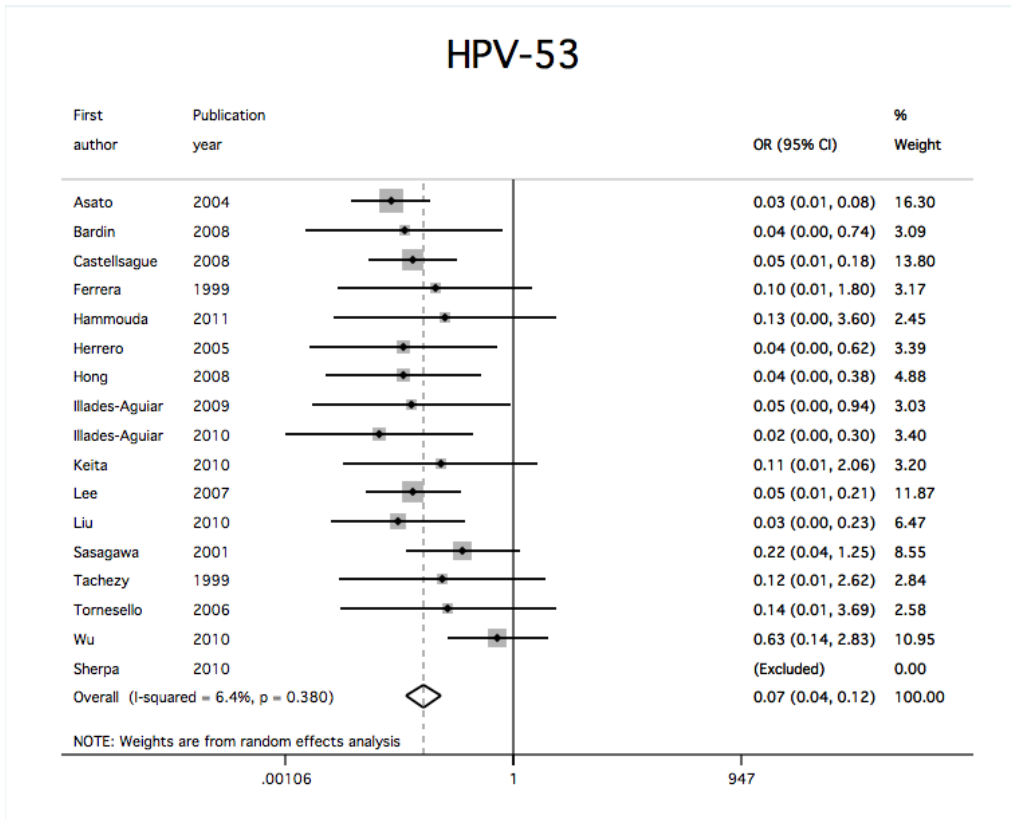


(o)

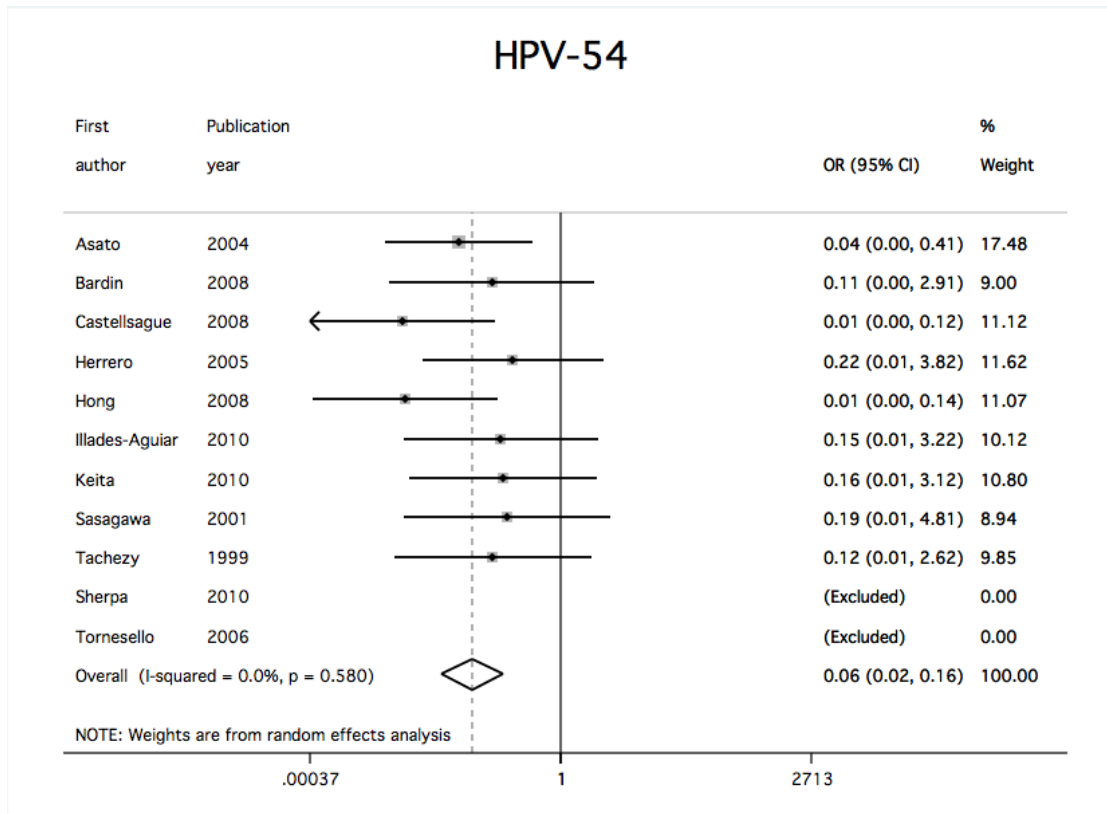




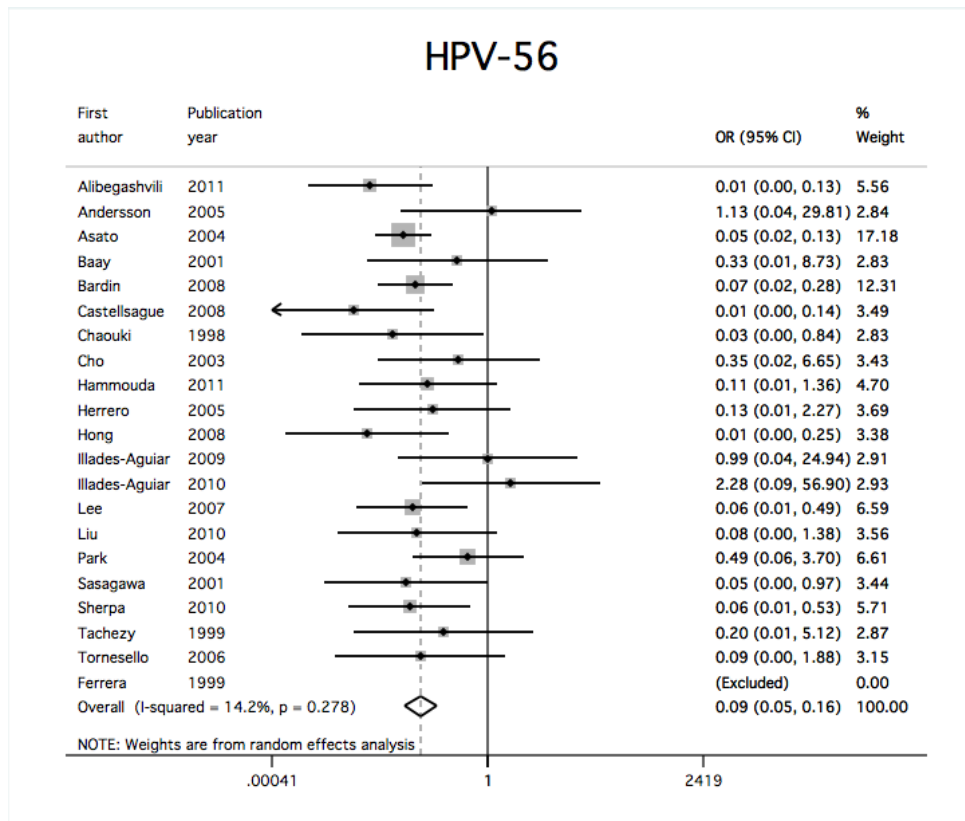
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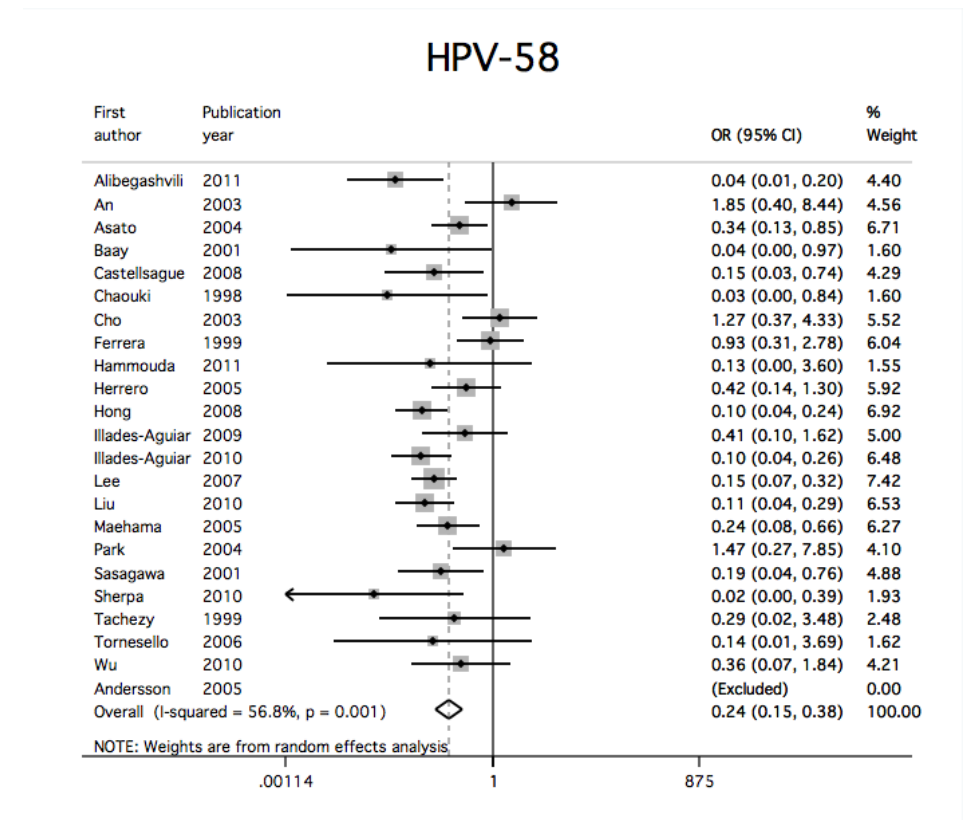
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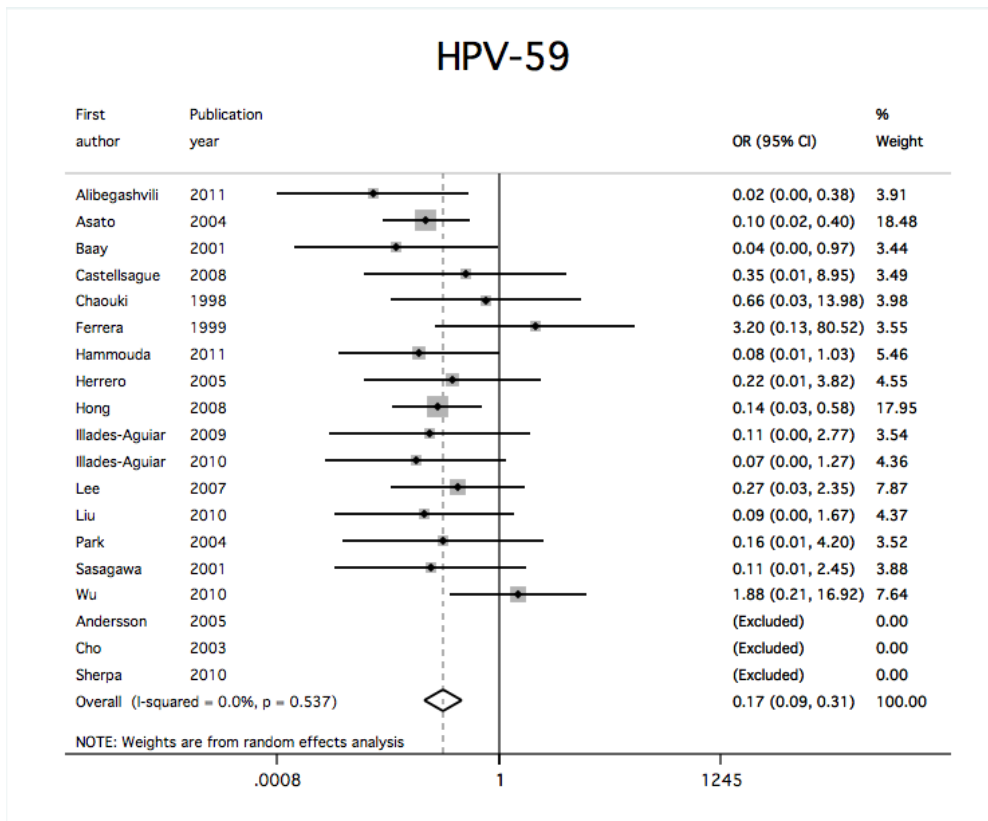
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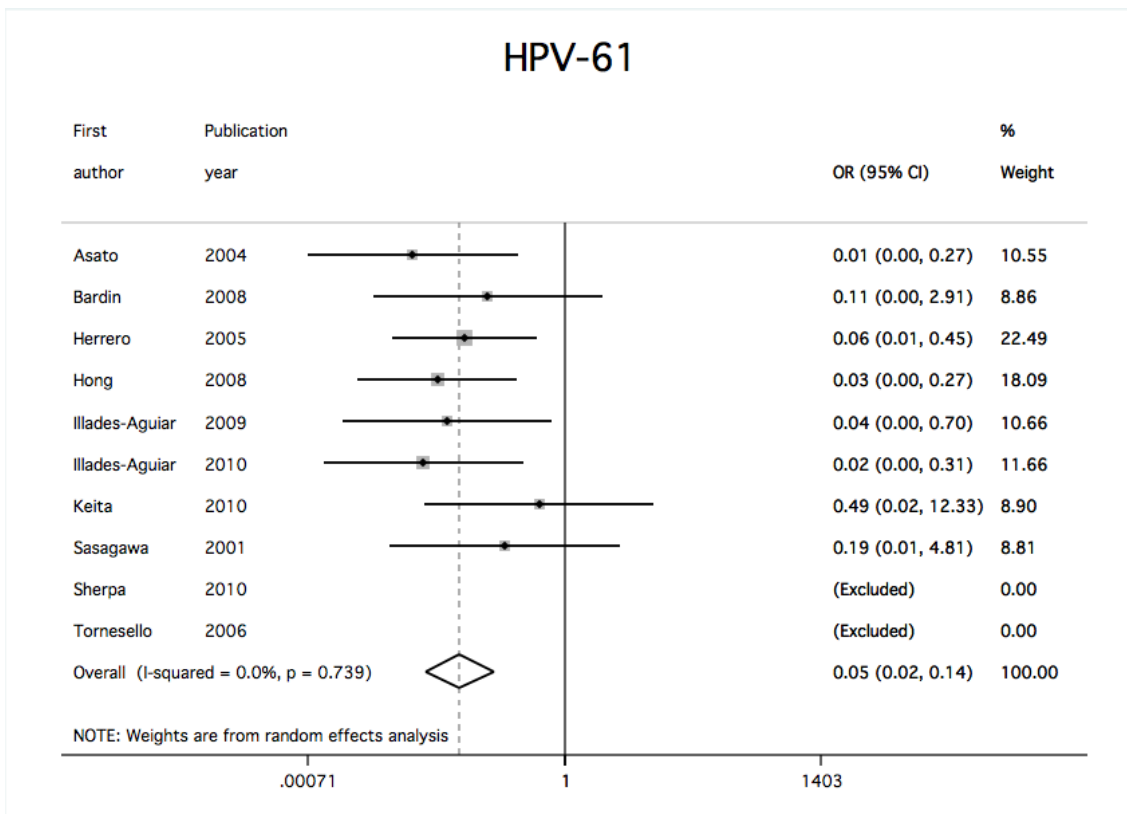
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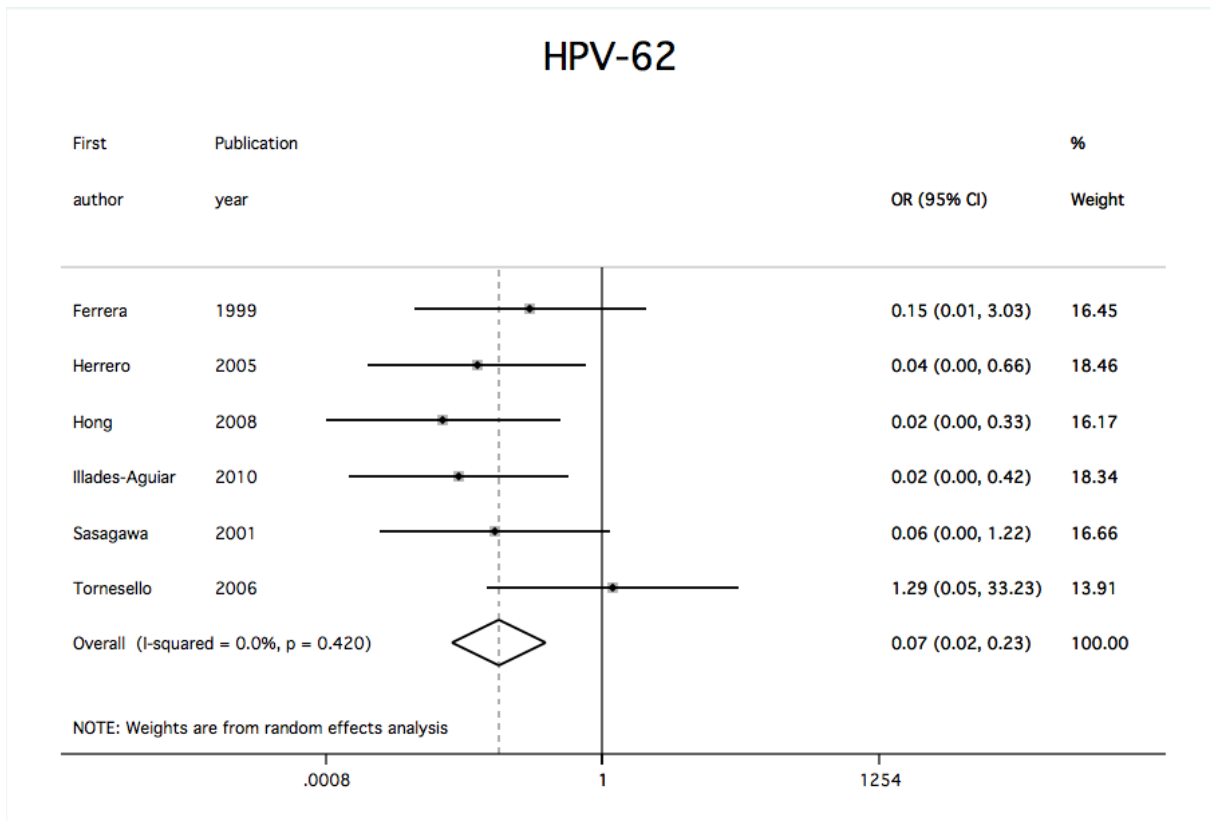
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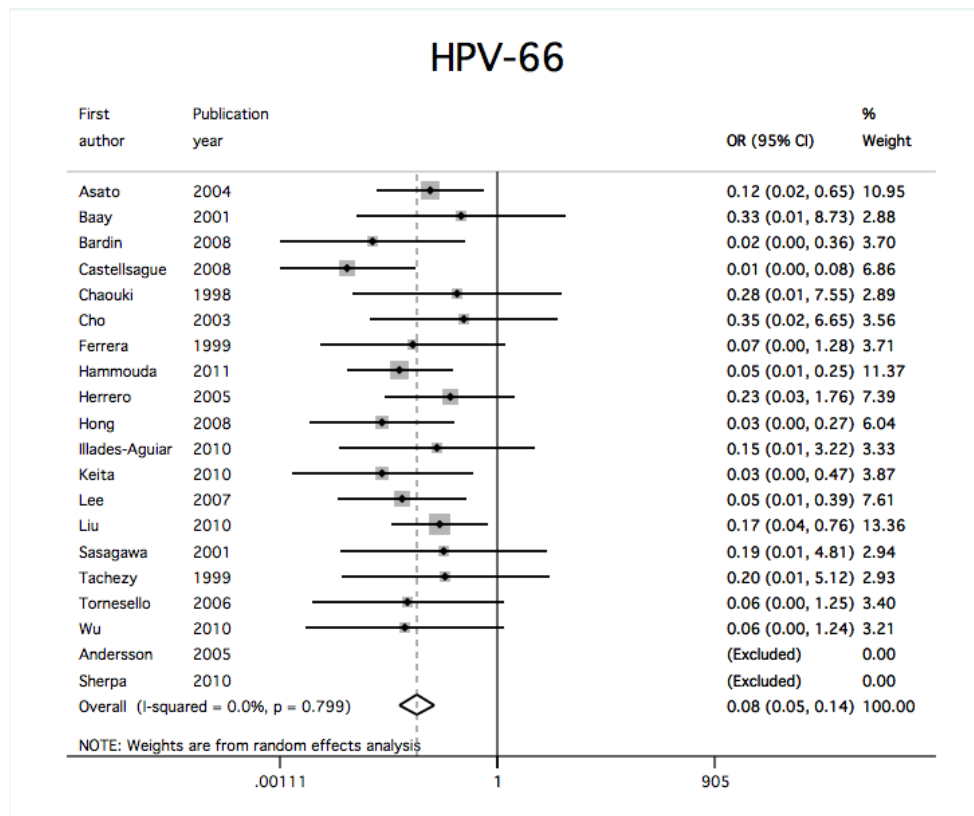
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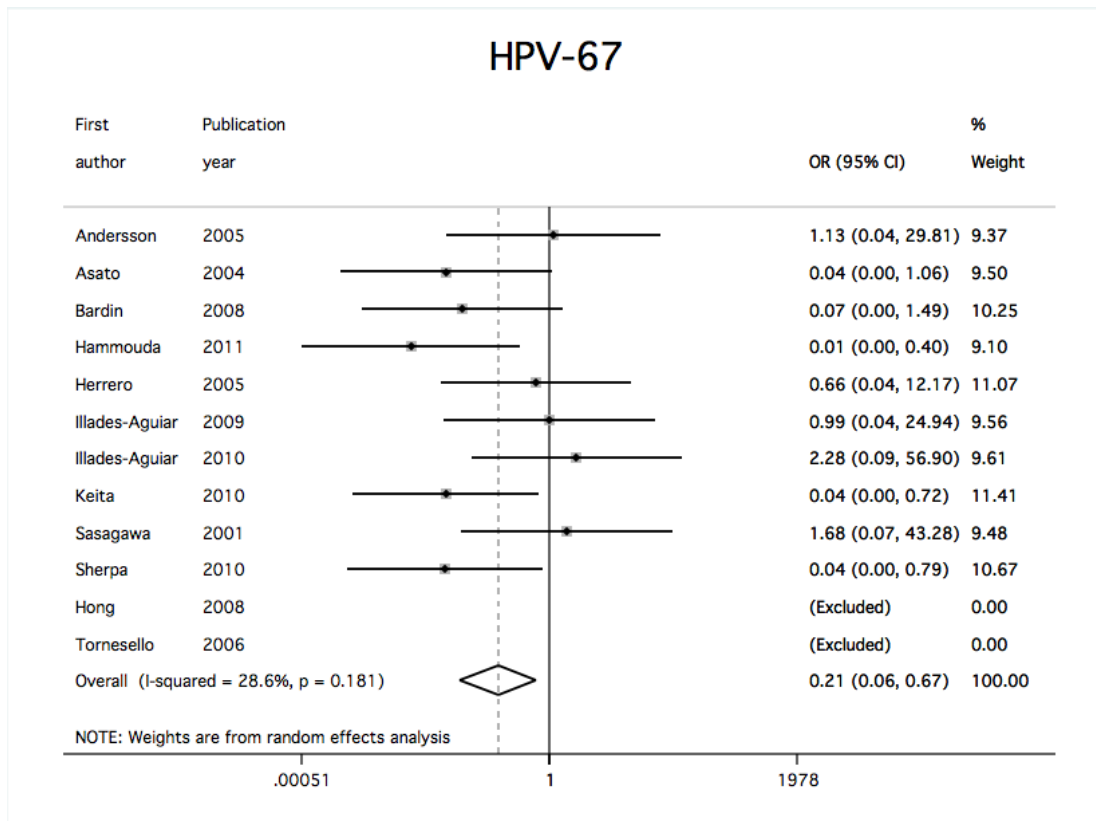
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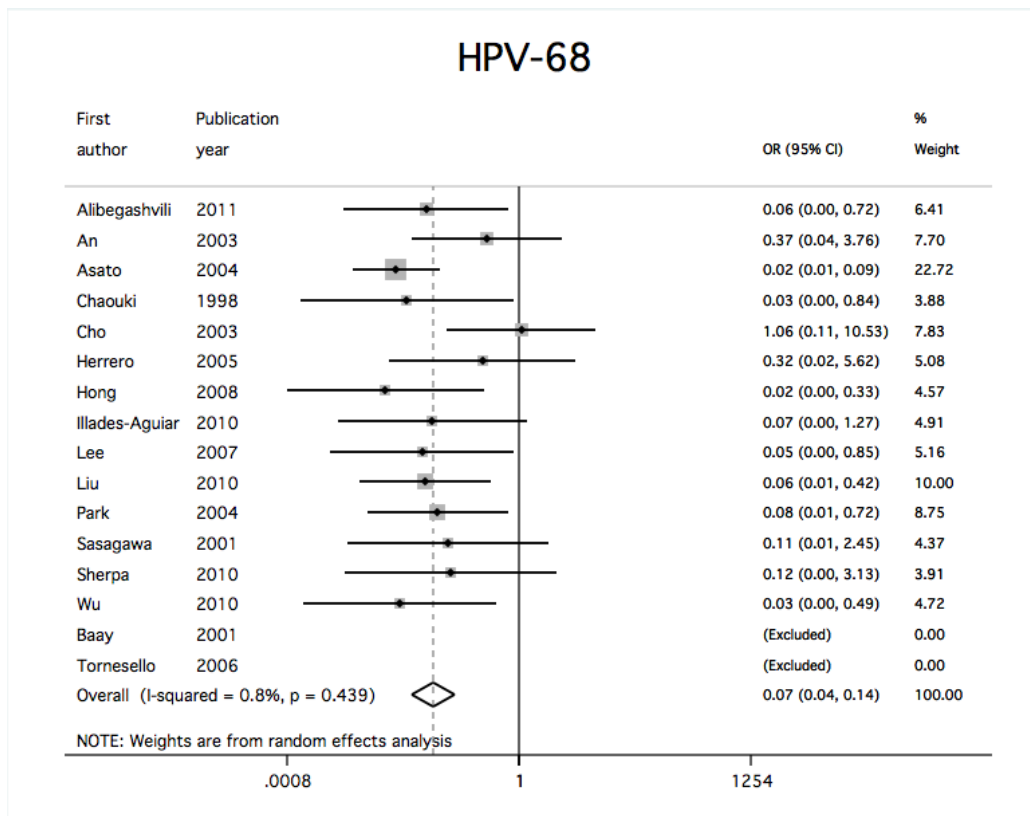
(w)



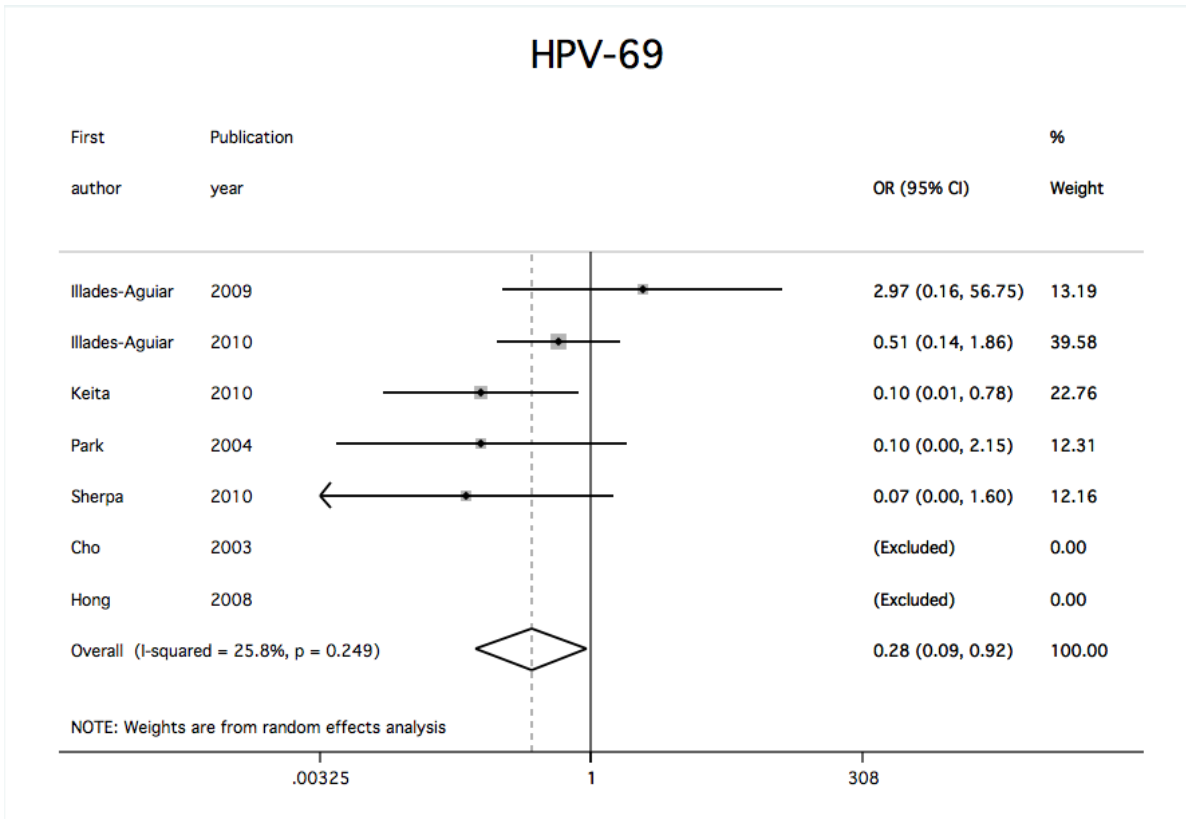
(x)



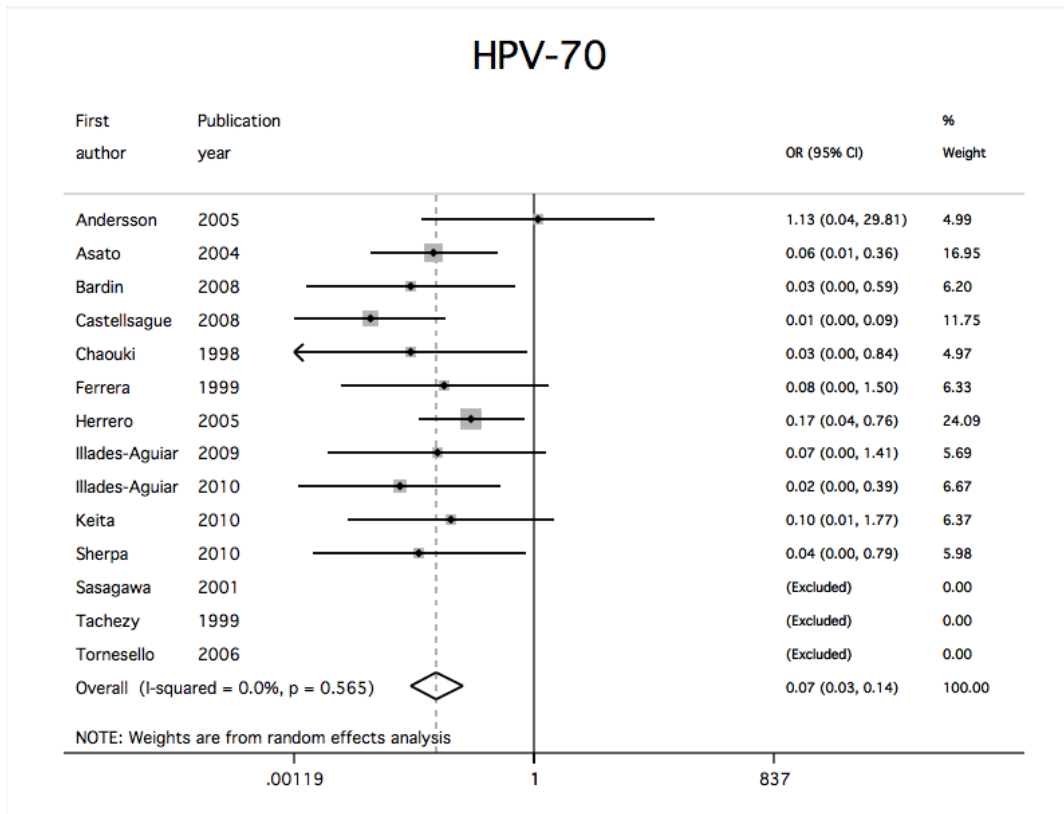
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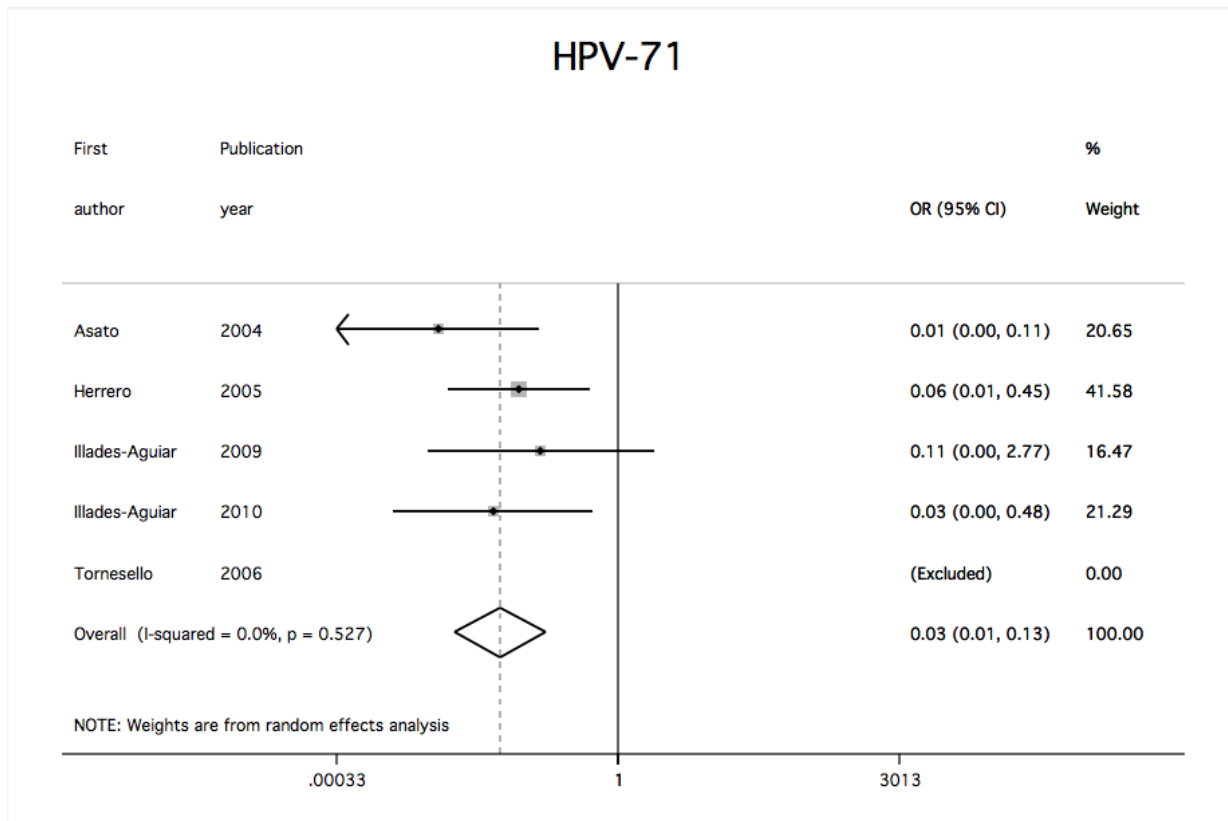
(z)



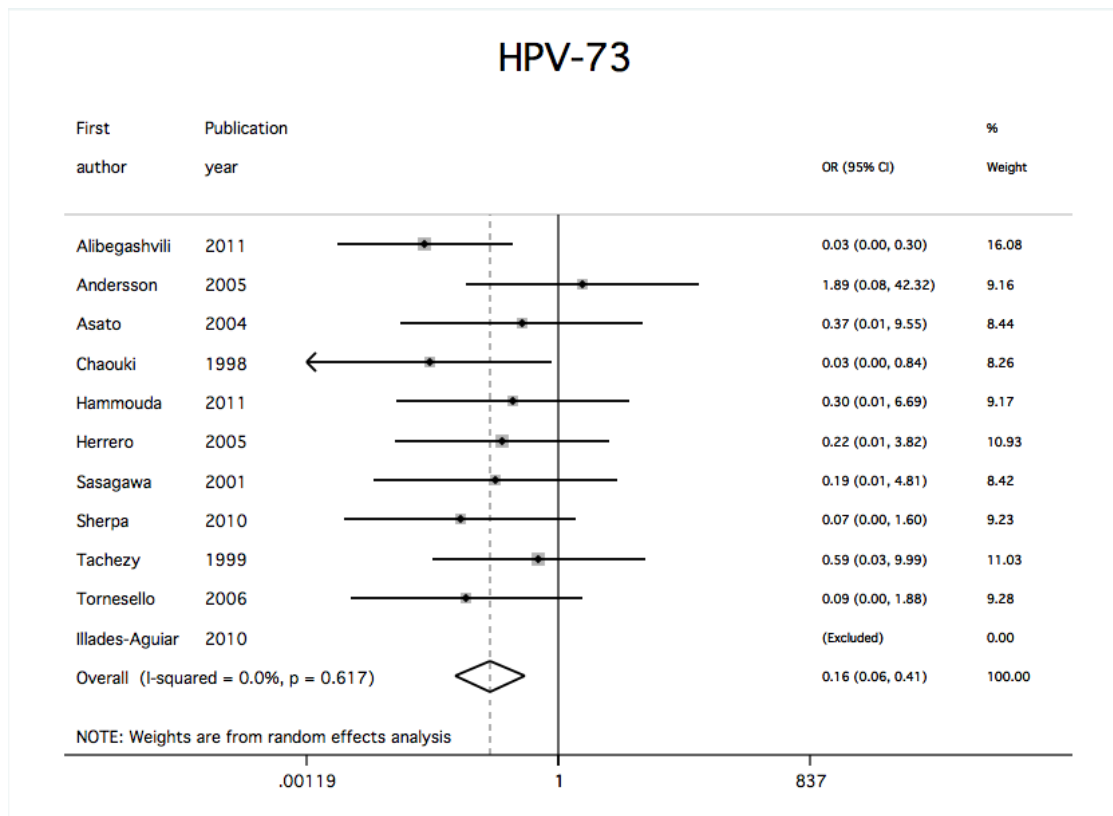
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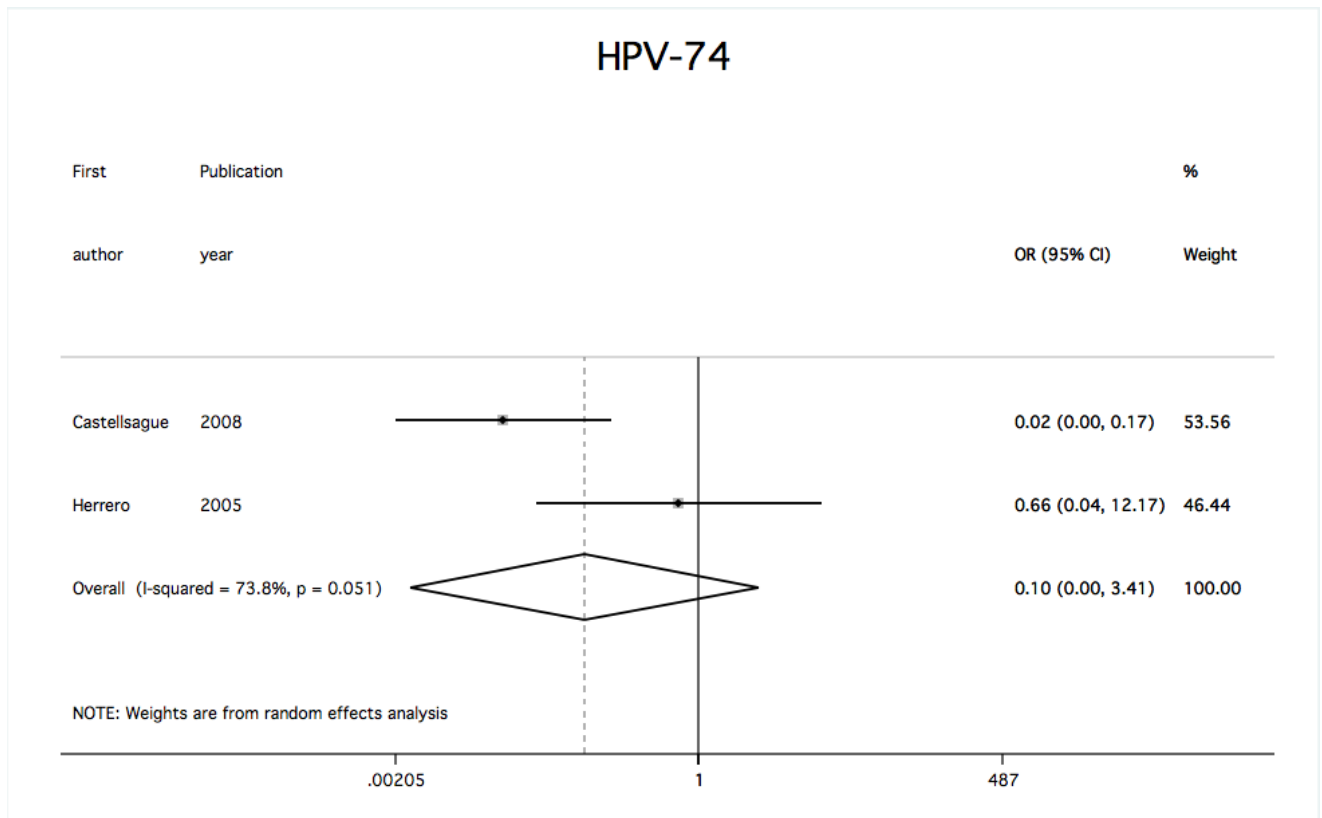
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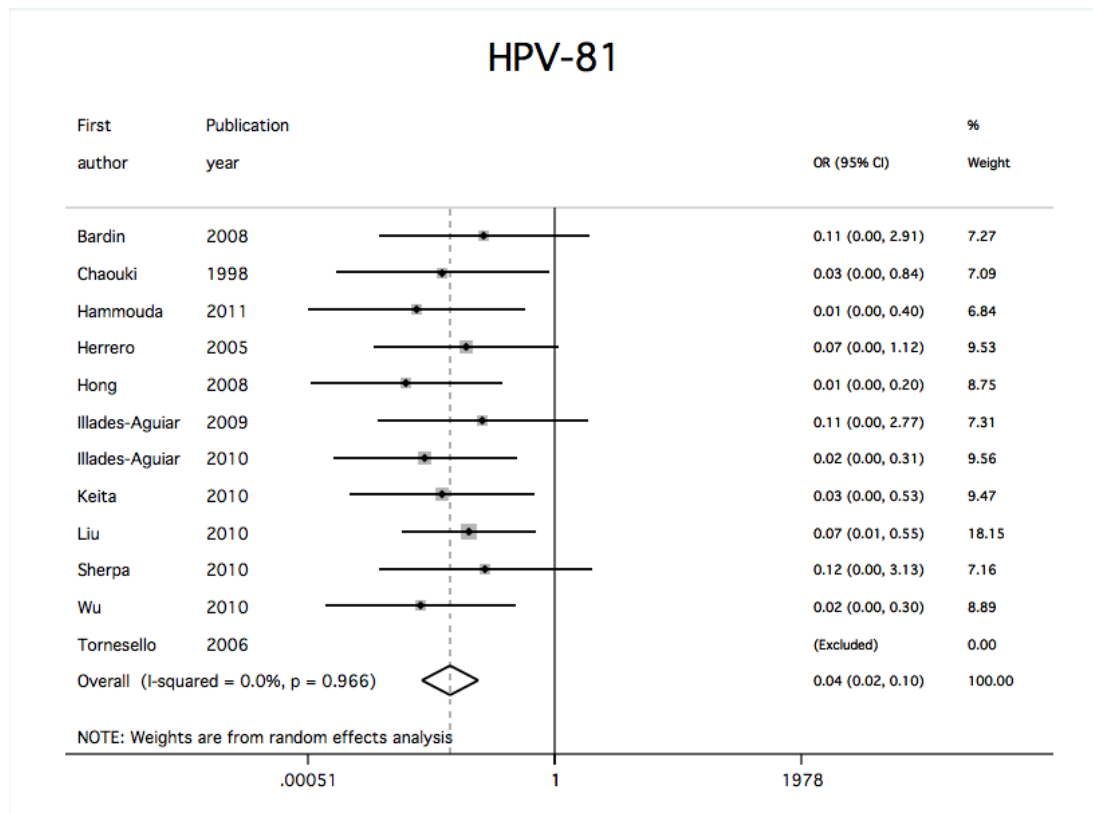
(ac)



(ad)



(ae)





(af)

