

Rapport: Fosterdiagnostik med mikroarray för utökad analys av kromosomer, 246

| Sid | Stod | Ska stå | Korrigeringsdatum |
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| 29 | Denna utvärdering omfattar en genomgång av sammanfattningar till 1 969 vetenskapliga artiklar, varav 316 | Denna utvärdering omfattar en genomgång av sammanfattningar till 1 969 vetenskapliga artiklar, varav 319 | 160411 |
| 42 | Två kvalitativa [47,48] och två kvantitativa [49,50]. | s. Två kvantitativa [47,48] och två kvalitativa [49,50]. | 160411 |
| 94 | Hillman 2013 [30] | Hillman 2013 [48] | 160411 |
| 43 | Totalt inkluderades åtta studier där olika typer av ultraljudsavvikelse följts upp med mikroarray [24,27,33,38–40,42,43]. | Totalt inkluderades 13 studier där olika typer av ultraljudsavvikelse följts upp med mikroarray [21,24,27,28,32-34, 37-39,41,42]. | 160411 |
| 96 | 21. Hillman SC, Pretlove S, Coomarasamy A, McMullan DJ, Davison EV, Maher ER, et al. Additional information from array comparative genomic hybridization technology over conventional karyotyping in prenatal diagnosis: a systematic review and meta-analysis. <i>Ultrasound Obstet Gynecol</i> 2011;37:6-14. | 21. Hillman SC, McMullan DJ, Hall G, Togneri FS, James N, Maher EJ, et al. Use of prenatal chromosomal micro-array: prospective cohort study and systematic review and meta-analysis. <i>Ultrasound Obstet Gynecol</i> 2013;41: 610-20. | 160411 |
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| 97 | 31. Kan AS, Lau ET, Tang WF, Chan SS, Ding SC, Chan KY, et al. Whole-genome array CGH evaluation for replacing prenatal karyotyping in | 31. Liao C, Fu F, Li R, Xie GE, Zhang YL, Li J, et al. Implementation of high-resolution SNP arrays in the investigation of fetuses with ultrasound malformations: 5 years of | 160411 |

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| | Hong Kong. PLoS One 2014;9:e87988. | clinical experience. Clin Genet 2014;86:264-9. | |
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