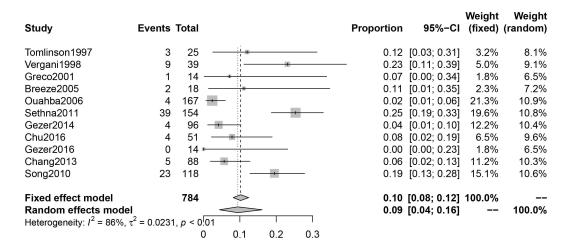
Supplementary Table1: Quality assessment of included studies evaluated by NOS.

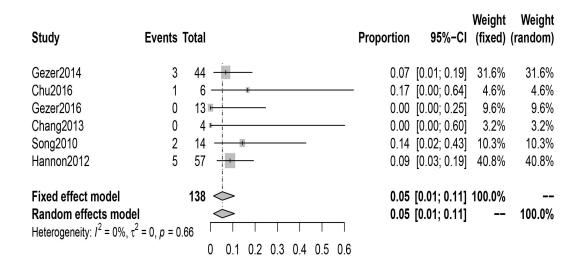
Study	Selection	Comparability	Outcomes	Total scores
Tomlinson ⁴	4	2	2	8
Vergani ⁵	4	1	2	7
Greco ⁶	4	2	2	8
Gaglioti ⁷	4	2	1	7
Breeze ⁸	4	1	2	7
Ouahba ⁹	4	2	2	8
Weichert ¹⁰	4	2	2	8
Madazli ¹¹	4	1	1	6
Sethna ¹²	4	2	2	8
Gezer ¹³	4	2	1	7
Chu ¹⁴	4	2	2	8
Gezer ¹⁵	4	1	1	6
Chang ¹⁶	4	1	1	6
Song ¹⁷	4	2	2	8
Zhang ²²	4	1	2	7
Li ¹⁹	4	2	2	8
Peng ²¹	4	2	2	8
Hannon ¹⁸	4	1	1	6
Shaffer ²³	4	1	2	7
Donnelly ²⁰	4	1	1	6
Hu^{24}	4	2	2	8
Bardin ²⁶	4	1	1	6
Duan ²⁵	4	1	1	6

NOS:The Newcastle-Ottawa scale. Selection: 1.Representative for the population; 2.Ascertainment of exposure; 3.Consecutive patients; 4.Outcome not present at the start of the study. Outcomes:1.Assessment of outcome well performed; 2.Chromosome abnormality well described; 3.Adequacy of patient



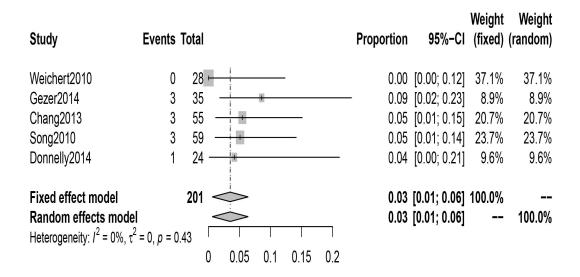
Supplementary Figure 1: Incidence of chromosomal abnormalities in mild VM.

CI: Confidence interval; VM: Ventriculomegaly.



Supplementary Figure 2: Incidence of chromosomal abnormalities in severe VM.

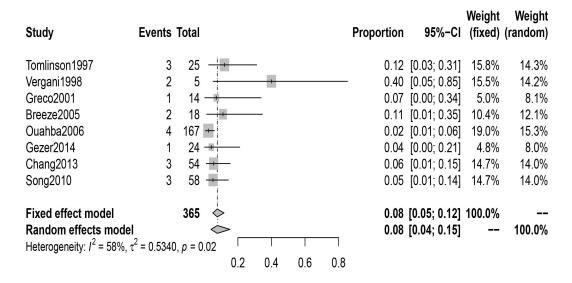
CI: Confidence interval; VM: Ventriculomegaly.



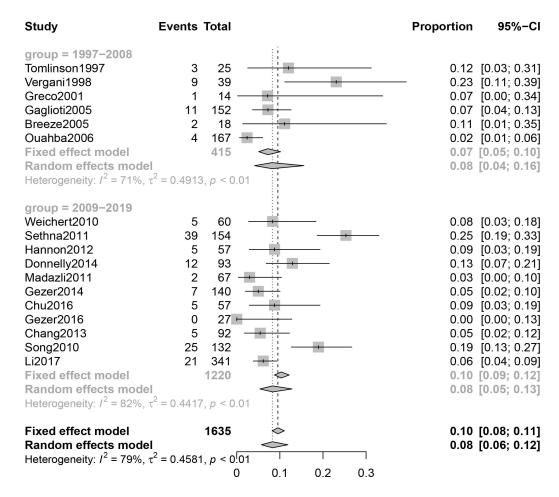
Supplementary Figure 3: Incidence of chromosomal abnormalities in isolated VM. CI: Confidence interval; VM: Ventriculomegaly.

Study	Events Total	Proportion	95%-CI	Weight Weight (fixed) (random)
Weichert2010 Gezer2014 Chang2013	5 32 — — — — — — — — — — — — — — — — — —	0.04	[0.05; 0.33] [0.01; 0.09] [0.01; 0.18]	10.2% 18.1% 33.1% 21.7% 11.8% 18.7%
Song2010 Donnelly2014	22 73 — — — — — — — — — — — — — — — — — —	0.30	[0.20; 0.42] [0.08; 0.27]	
Fixed effect model Random effects model Heterogeneity: I^2 = 85%,		0.13	[0.09; 0.16] [0.04; 0.25]	100.0% 100.0%

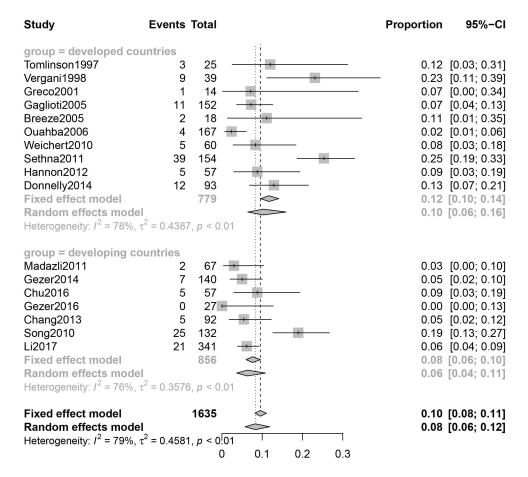
Supplementary Figure 4: Incidence of chromosomal abnormalities in non–isolated VM. CI: Confidence interval; VM: Ventriculomegaly.



Supplementary Figure 5: Incidence of chromosomal abnormalities in isolated mild VM. CI: Confidence interval; VM: Ventriculomegaly.

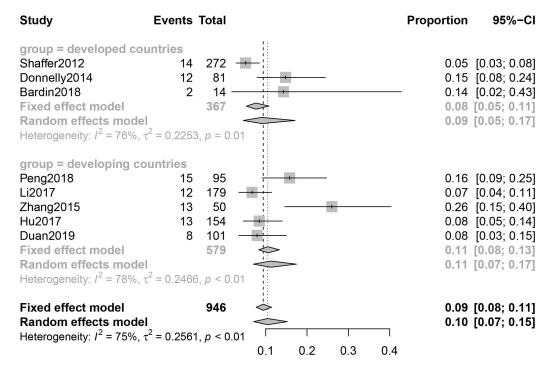


Supplementary Figure 6: Subgroup analysis of Karyotyping with different publication year. CI:Confidence interval.

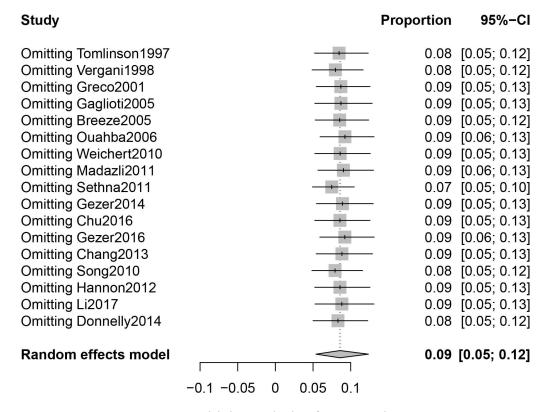


Supplementary Figure 7: Subgroup analysis of Karyotyping with different countries.

CI: Confidence interval.

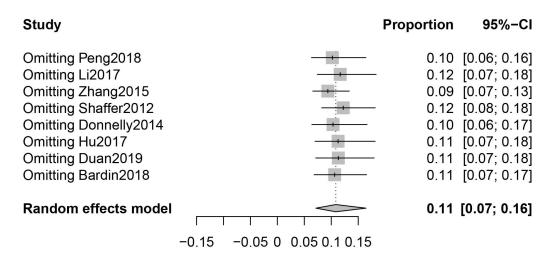


Supplementary Figure 8: Subgroup analysis of CMA. CI: Confidence interval; CMA: Chromosomal microarray analysis.



Supplementary Figure 9:Sensitivity analysis of Karyotypin

CI: Confidence interval.



Supplementary Figure 10: Sensitivity analysis of CMA. CI: Confidence interval;

CMA: Chromosomal microarray analysis.