



Correction

Correction: Furnier et al. Translating Molecular Technologies into Routine Newborn Screening Practice. *Int. J. Neonatal Screen.* 2020, 6, 80

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In the original article [1], there was a mistake in Table 2 as published. For reference Vill et al. 2019, the entry “1 in 7096” has been corrected to “1 in 7524”. Also for reference Kay et al. 2020, the entry “No” under “SMN2 Inclusion” has been corrected to say “Real-time PCR assay to assess SMN2 copy number”. The corrected Table 2 appears below. The authors apologize for any inconvenience caused and state that the scientific conclusions are unaffected. The original article has been updated.



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Table 2. Selected spinal muscular atrophy newborn screening studies.

Reference	Region	Screening Method	SMN2 Inclusion	Number of Newborns Screened	Reported Incidence in Sample	Study Type
Chien et al. 2017 [33]	Taiwan	Real-time PCR <i>SMN1</i> assay to detect homozygous exon 7 deletion; verified by droplet digital PCR assay	Droplet digital PCR assay to assess <i>SMN2</i> copy number	120,267	1 in 17,181	Pilot
Boemer et al. 2019 [32]	Belgium	Real-time PCR <i>SMN1</i> assay to detect homozygous exon 7 deletion	No	Not applicable	Not applicable	Pilot
Vill et al. 2019 [36]	Germany	Real-time PCR <i>SMN1</i> assay to detect homozygous exon 7 deletion; verified by multiplex ligation-dependent probe amplification (MLPA)	MLPA to assess <i>SMN2</i> copy number	165,525	1 in 7524	Pilot
Kariyawasam et al. 2020 [34]	Australia	Real-time PCR <i>SMN1</i> assay to detect homozygous exon 7 deletion	Droplet digital PCR assay to assess <i>SMN2</i> copy number	103,903	1 in 10,390	Pilot
Kay et al. 2020 [35]	New York	Real-time PCR <i>SMN1</i> assay to detect homozygous exon 7 deletion	Real-time PCR assay to assess <i>SMN2</i> copy number	225,093	1 in 28,137	Routine

Reference

1. Furnier, S.M.; Durkin, M.S.; Baker, M.W. Translating Molecular Technologies into Routine Newborn Screening Practice. *Int. J. Neonatal Screen.* **2020**, *6*, 80. [[CrossRef](#)]