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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 *SMN1* assay to detect homozygous exon 7 deletion; verified by droplet digital PCR assay	Droplet digital PCR assay to assess *SMN2* copy number	120,267	1 in 17,181	Pilot
Boemer et al. 2019 [32]	Belgium	Real-time PCR *SMN1* assay to detect homozygous exon 7 deletion	No	Not applicable	Not applicable	Pilot
Vill et al. 2019 [36]	Germany	Real-time PCR *SMN1* assay to detect homozygous exon 7 deletion; verified by multiplex ligation-dependent probe amplification (MLPA)	MLPA to assess *SMN2* copy number	165,525	1 in 7524	Pilot
Kariyawasam et al. 2020 [34]	Australia	Real-time PCR *SMN1* assay to detect homozygous exon 7 deletion	Droplet digital PCR assay to assess *SMN2* copy number	103,903	1 in 10,390	Pilot
Kay et al. 2020 [35]	New York	Real-time PCR *SMN1* assay to detect homozygous exon 7 deletion	Real-time PCR assay to assess *SMN2* 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]