

Gefahr in Verzug: Plasmid-DNA-Kontamination bis zu 145mal höher als die zulässigen TGA-Sicherheitswerte

Researcher	Affiliation, Country	Pharma Company	# of Vials	First reported	Methods	DNA/dose (limit 10ng)	DNA/RNA ratio (limit 1/3030)	Concerns	Publication
McKernan, K.	Medicinal Genomics, US	Pfizer, Moderna	a dozen	2023-02	Electrophoresis (Agilent) Fluorometer (Qubit) qPCR/RT-qPCR	2250ng – 3390ng [1] 312ng – 843ng [1] 12ng	1/8 – 1/2 1/47 – 1/8 1/161 – 1/43	Adverse events, Gene integration	Reported to FDA Presented at WCH Reported by Epochtimes Found gene integration in OvCar3 cancer cells transfected by Kämmerer Found SV40 in tumors
Nitta, T.	Tokyo Univ, Japan	Moderna	1 or 2	2023-03	qPCR/RT-qPCR	0.1ng [2]	1/1,000,000	No Problem	No publication
Buckhaults, P. J.	University of South Carolina, US	Pfizer, Moderna Pfizer 2020, Pfizer 2023, Moderna 2020, Moderna 2023	some	2023-07	qPCR	0.6ng – 18.7ng 7.7ng (SV40, Pfizer) 4.5 – 5.5ng (Neo/Kan, Pfizer) 1.5 – 9.0ng (ORI, Pfizer) 2.5 – 18.7ng (Spike, Pfizer) 0.002 – 0.004ng (ORI, Moderna 2023)		Adverse events, Gene integration	Presented in South Carolina Senate Presented gene integration to normal human epithelial stem cells
König, B. Kirschner, J. O.	MMD, Germany Indep., Germany	Pfizer, Moderna	4	2024-04	qPCR	3600ng – 5340ng	1/12 – 1/7	Adverse events, Gene integration	Reported to the German Ministry of Health Reported by local public broadcasting MDPf peer-reviewed paper
Speicher, D. J.	University of Guelph, Canada	Pfizer, Moderna	27	2023-10	Fluorometer (Qubit) qPCR	1896ng – 5100ng 0.22ng – 2.43ng (Spike) 0.01ng – 4.27ng (ORI)		Adverse events, Gene integration	Presented at WCH
Speicher, D. J.	University of Guelph, Canada	Pfizer, Moderna (Australia)	3	2024-06	Fluorometer (Qubit) qPCR	451ng – 1420ng (after RNaseA/DNaseA) 6.46ng – 163.68ng (Spike) 0.54ng – 12.97ng (ORI) 3.70ng – 14.69ng (SV40, Pfizer)		Adverse events, Gene integration	Reported to Therapeutic Goods Administration (TGA) Under litigation
Raoult, D.	Aix-Marseille Univ (Former Prof), France	Pfizer	some	2024-11	Fluorometer (Qubit)	216ng (Avg) 5160ng (Avg, after Triton-X-100)		Gene integration	Preprint
Kämmerer, U.	Univ. Hospital of Würzburg, Germany	Pfizer	4	2024-12	Fluorometer (Qubit)	2712 – 3683ng (after Triton-X-100) 32.71 – 42.09ng (after Triton-X-100/RNaseA)		Adverse events, Gene integration	Peer-reviewed paper of Science, public health policy, and the law Transfection to HEK293 cells
Wang, T. J. Kim, A., Kim, K. (FDA scientists helped them)	Centerville High School, US (executed at FDA White Oak Campus)	Pfizer	2 of 4	2024-12	Fluorometer (Qubit)	41.4 – 109.5ng (extracted by Monarch Plasmid DNA Miniprep, which includes RNaseA)		Gene integration	Peer-reviewed paper of Journal of High School Science

[1] Multiplied the value by 300 for ul

[2] From the description of DNA 44x10fg to mRNA 400ng, the calculation for Moderna 1-dose as mRNA 100ug

<https://porthedlandmotion.info/>

Antrag auf Aussetzung der Covid-19-Impfstoffe (Pfizer und Moderna) DNA Kontamination – **Port Hedland AUSTRALIEN**

Quellen:

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https://x.com/P_J_Buckhaults/status/1684569351998038017

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<https://julesonthebeach.substack.com/p/australian-dna-contamination-confirmed>

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<https://publichealthpolicyjournal.com/biontech-rna-based-covid-19-injections-contain-large-amounts-of-residual-dna-include-an-sv40-promoter-enhancer-sequence/>

<https://jhss.scholasticahq.com/article/127890-a-rapid-detection-method-of-replication-competent-plasmid-dna-from-covid-19-mrna-vaccines-for-quality-control>