



Supplementary Materials

Tyrosine-Modification of Polypropylenimine (PPI) and Polyethylenimine (PEI) Strongly Improves Efficacy of siRNA-Mediated Gene Knockdown

Sandra Noske, Michael Karimov, Achim Aigner * and Alexander Ewe *

Rudolf-Boehm-Institute for Pharmacology and Toxicology, Clinical Pharmacology, Leipzig University, Faculty of Medicine, Leipzig, 04107, Germany; Sandra.noske@medizin.uni-leipzig.de (S.N.); michael.karimov@medizin.uni-leipzig.de (M.K.)

* Correspondence: achim.aigner@medizin.uni-leipzig.de (A.A.); alexander.ewe@medizin.uni-leipzig.de (A.E.); Tel.: +49-(0)341-9724660 (A.A.)

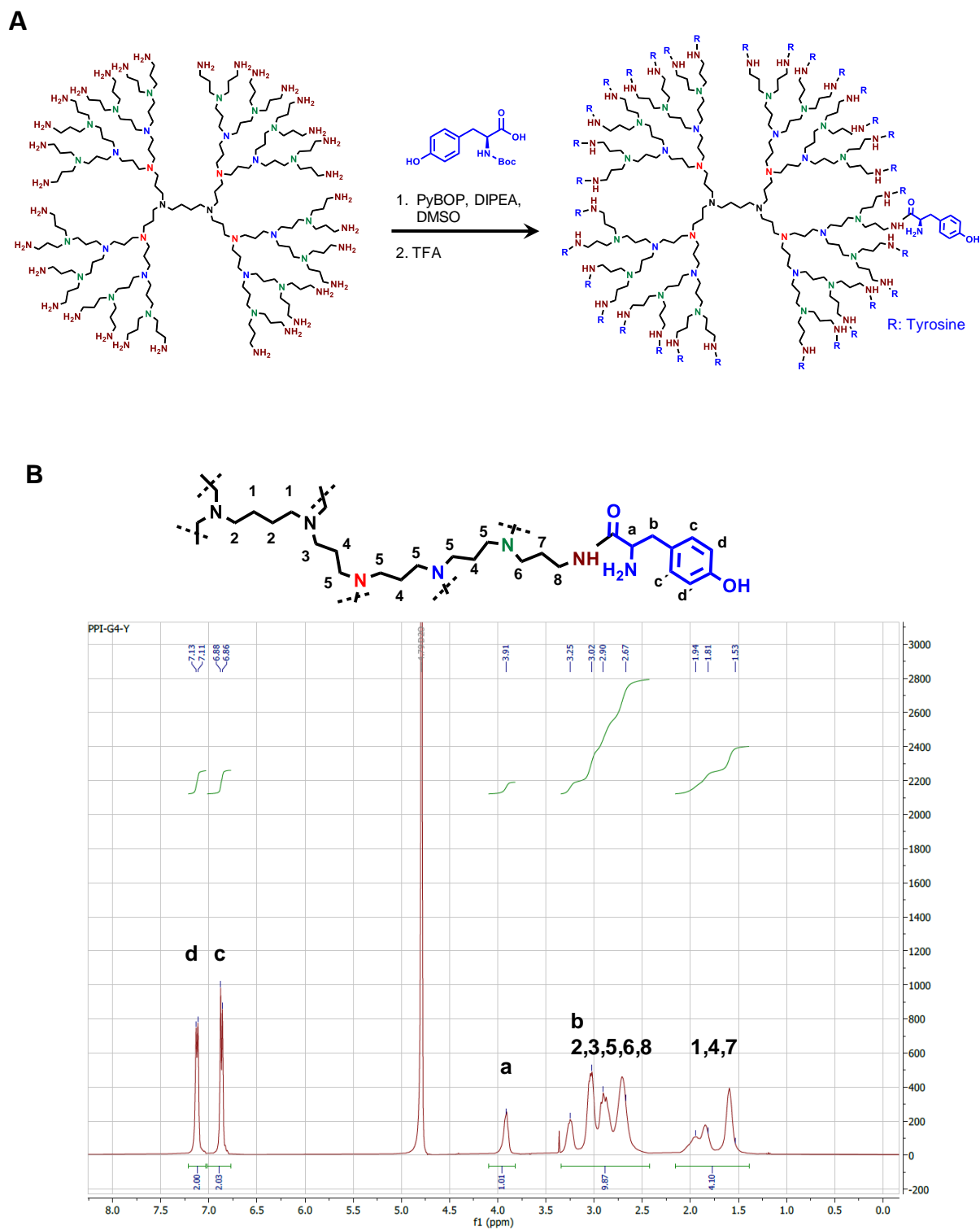


Figure S1. (A) Synthesis scheme for the generation of PPI-G4-Y. (B) $^1\text{H-NMR}$ analysis of PPI-G4-Y. Relevant signals are highlighted.

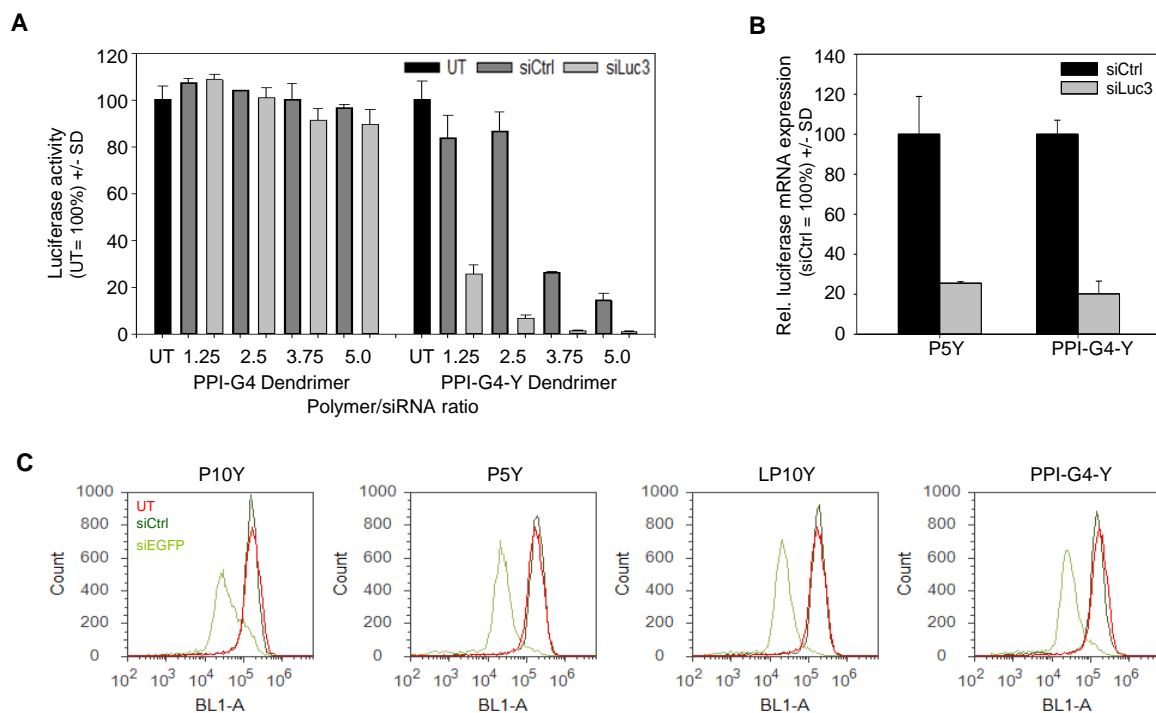


Figure S2. (A) Knockdown efficacies of PPI-G4 dendrimer-based siRNA complexes with (right) or without (left) tyrosine modification, as determined by luciferase knockdown in PC3-EGFP/Luc reporter cells. Bars show results upon transfection with complexes containing negative control siRNA (black) or luciferase-specific siRNA (grey), respectively, with bars normalized for untreated (UT) negative control. (B) Luciferase knockdown on the mRNA level in PC3-EGFP/Luc cells, as determined by RT-qPCR. (C) Knockdown efficacies of various tyrosine-modified PPI- or PEI-based siRNA complexes targeting EGFP in PC3-EGFP/Luc cells (original flow cytometry data from Figure 3A).

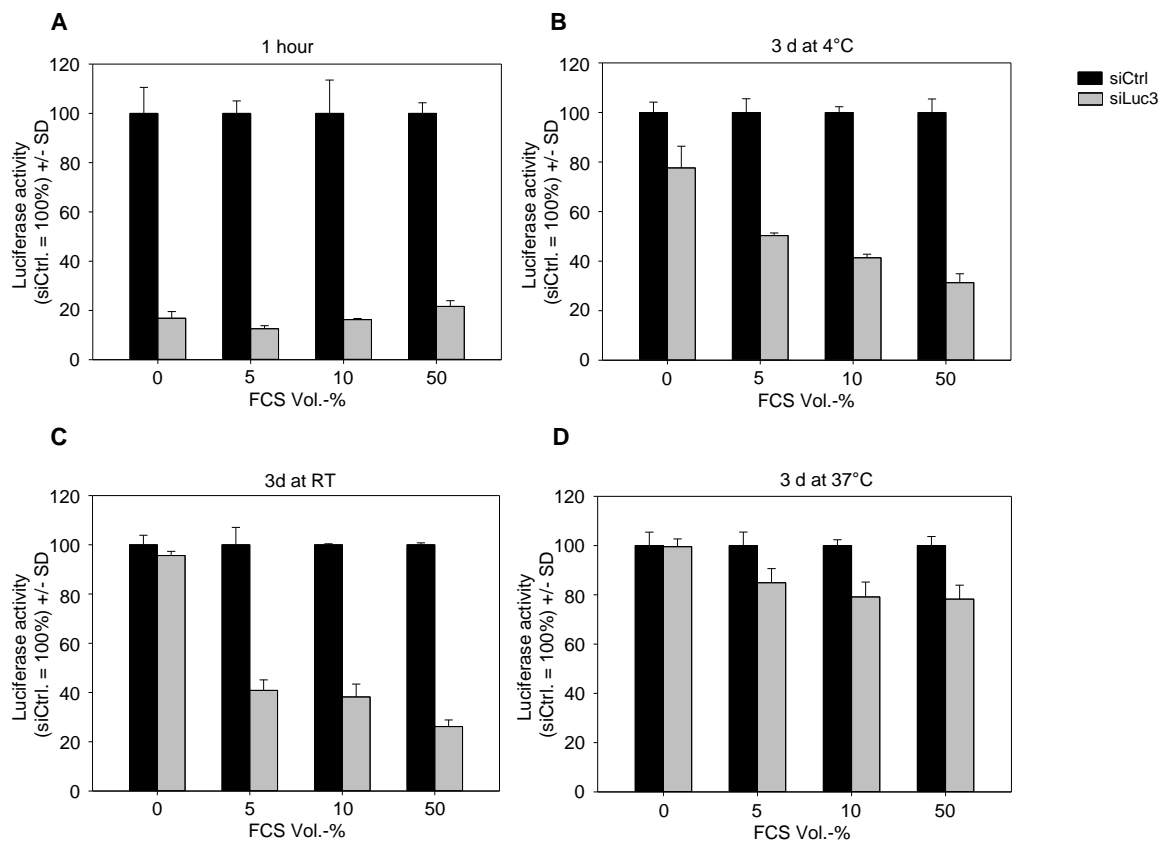


Figure S3. Knockdown efficacies of PPI-G4 dendrimer-based siRNA complexes upon storage for three days at different temperatures, in the absence or presence of FCS at the indicated concentrations, as determined by luciferase activity in PC3-EGFP/Luc cells.

Table S1. siRNAs used in this study.

siRNA		Sequence (5' – 3')	Vendor
siLuc 2	<i>sense</i>	CGUACGCGGAAUACUUCGA dTdT	Dharmacon Lafayette, CO, USA
	<i>antisense</i>	UCGAAGUAUCCGCGUACG dTdT	
siLuc 3	<i>sense</i>	CUUACGCUGAGUACUUCGA dTdT	Eurogentec, Seraing, Belgium
	<i>antisense</i>	UCGAAGUACUCAGCGUAAG dTdT	
siGAPDH	<i>sense</i>	CCUCAACUACAUGGUUUAC dTdT	Eurogentec Seraing, Belgium
	<i>antisense</i>	GUAAACCAUGUAGUUGAGG dTdT	
siEGFP	<i>sense</i>	GCAGCACGACUUCUUCAAG dTdT	Eurogentec Seraing, Belgium
	<i>antisense</i>	CUUGAAGAAGUCGUGCUGC dTdT	

Table S2. Sequences of primers used in this study.

Primer		Sequence (5' – 3')	Vendor
RPLP0	forward	TCTACAACCCTGAAGTGCTTGAT	Eurofins Genomics, Ebersberg, Germany
	reverse	CAATCTGCAGACAGACTGG	
GAPDH	forward	GGTGTGAACCATGAGAAGTATGA	Bioscience, Gera, Germany
	reverse	GAGTCCTTCCACGATACCAAAG	
Luc3	forward	TTACACCCGAGGGGGATGAT	Eurofins Genomics, Ebersberg, Germany
	reverse	TTCACACACAGTTCGCCTC	