Supporting Information for

Animal toxicity of hairpin pyrrole-imidazole polyamides varies with the turn unit

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Table S2: MALDI-ToF characterization of polyamides 1-5.

Delverside	5'- TTGC TGTTCT GCAA -3		
Polyamide	T _m /°C	$\Delta T_{\rm m}$ / °C	
_	61.8 (±0.5)	_	
1	74.1 (±0.3)	12.3	
2	75.1 (±0.4)	13.3	
3	70.1 (±0.2)	8.3	
4	74.9 (±0.2)	13.2	

Fig. S1. DNA thermal stabilization analysis of compounds 1-4.

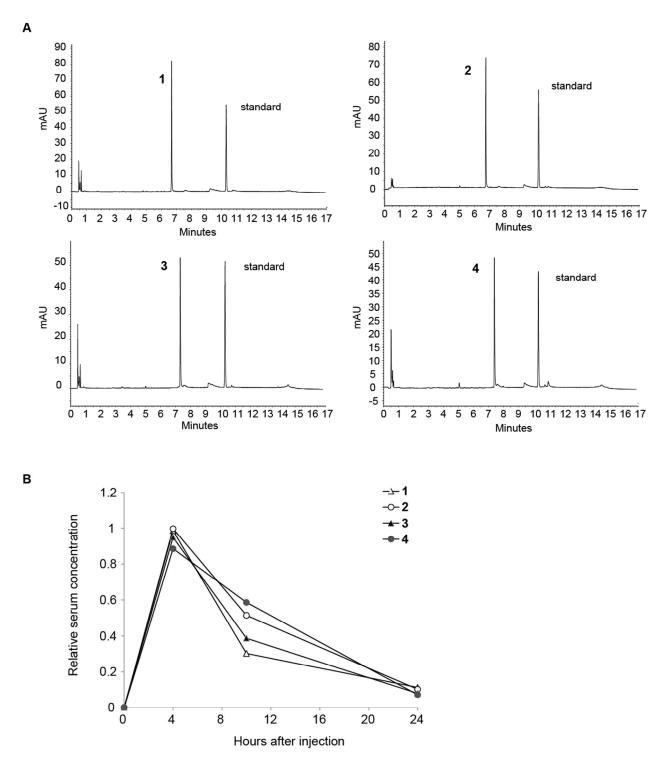


Fig S2. (A) Analytical HPLC traces of compounds **1-4** in the serum 4 hr after injection. **(B)** Relative serum levels of compounds **1-4** at 4 hr, 10 hr, and 24 hr after a single subcutaneous injection of each compound at 10 mg/kg.

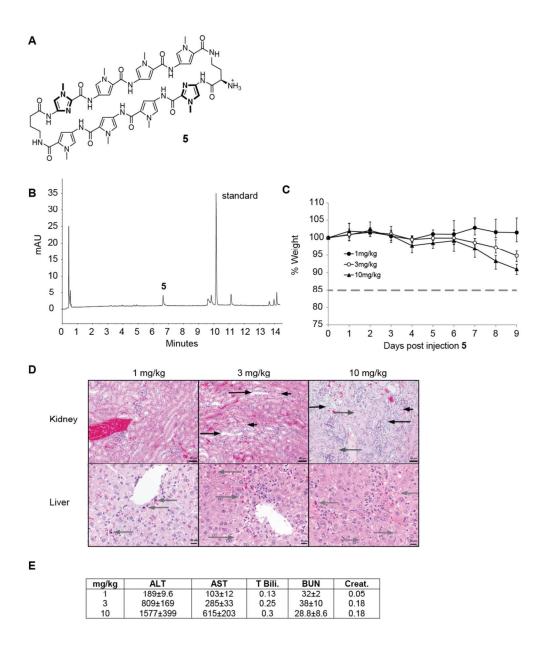


Fig S3. Characterization of a cyclic polyamide targeting the sequence 5'-WGWWCW-3'. (A) Chemical structure of compound 5. (B) Serum circulation of 5 4 hr after SC injection. (C) Changes in animal weights after a single SC injection of 5 at the indicated concentrations. (D) Kidney and liver histology of sacrificed animals after 9 days of monitoring. (E) Serum chemistry of animals treated with 5. Liver: long gray arrow=hepatocellular apoptosis/necrosis, arrowheads=outline area of bridging hepatocellular necrosis/apoptosis. Kidney: short gray arrow=tubular epithelial karyomegaly, long gray arrow=tubular epithelial apoptosis/necrosis, short black arrow=tubular epithelial mitoses, long black arrow=tubular epithelial attenuation

	Test conc (μM)	Test species	Mean remaining parent with NADPH (%)	Mean remaining parent NADPH-free (%)
Verapamil high metabolism control	1	Human	4.2%	100%
	1	Mouse	1.1%	100%
Warfarin low metabolism control	1	Human	100%	100%
	1	Mouse	100%	100%
1	1	Human	96.9%	92.3%
	1	Mouse	95.2%	96.8%
2	1	Human	91.9%	100%
	1	Mouse	92.4%	100%
3	1	Human	95.3%	94.9%
	1	Mouse	97.3%	100%
4	1	Human	3.0%	3.8%
	1	Mouse	4.0%	4.9%

Table S1. Microsomal stability analysis of **1-4** in the presence and absence of NADPH. Samples were incubated for 1 hr at 37 °C with 1 mg/ml of human or mouse microsomes.

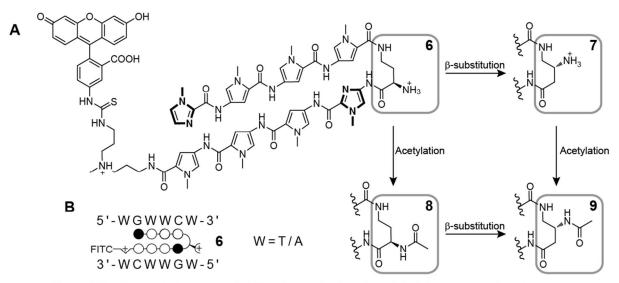


Figure S4. Chemical structures. **(A)** Structures of polyamides **6-9**. The compounds only vary by the amino substitution on the γ -turn unit. **(B)** The prefered DNA binding sequence of the polyamide core. Polyamide **6** is shown bound to the sequence 5'-WGWWCW-3'. Closed circles represent imidazole units and open circles represent pyrrole units.

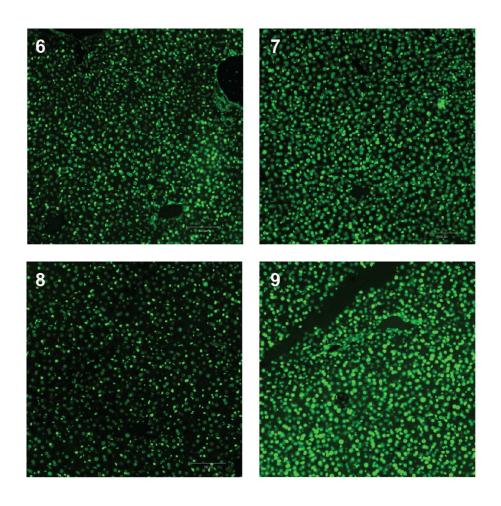


Figure S5. Nuclear localization of compounds 6-9 in the liver 24 hr after SC injection

Compound	Chemical Formula	Calculated Mass		Observed Mass
1	C65H76N22O12	[M+H]+	1357.44	1357.86
2	C65H76N22O12	[M+H]+	1357.44	1357.69
3	C67H78N22O13	[M+H]+	1399.48	1399.91
4	C67H78N22O13	[M+H]+	1399.48	1399.36
5	C54H61N21O10	[M+Na]+	1186.5	1186.6

 Table S2.
 MALDI-ToF analysis of compounds.