



Expression array: A siPOOL combines 30 siRNAs, designed with proprietary algorithms for maximal gene knock-down. As the working concentration of each siRNA in a siPOOL is in the picomolar range, off-target effects are greatly reduced while on-target silencing is cooperatively maintained.

Phenotypic correlation: If reliable and specific, two reagents that target the same gene should produce similar phenotypes. The highly-specific siPOOLs clearly outperformed individual siRNAs, giving rise to reproducible and reliable phenotypes.

