Supporting Information for

Oligonucleotide-Palladacycle Conjugates as Splice Correcting Agents

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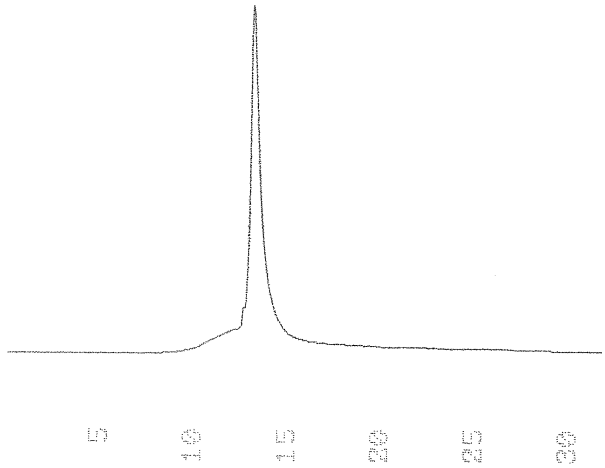

Figure S1. HPLC trace of oligonucleotide ON1b; Hypersil ODS C18 column (250 × 4.6 mm, 5 μm); flow rate = 1.0 mL min^{-1}; linear gradient (0 to 50% over 30 min) of MeCN in 50 mM aqueous triethylammonium acetate.

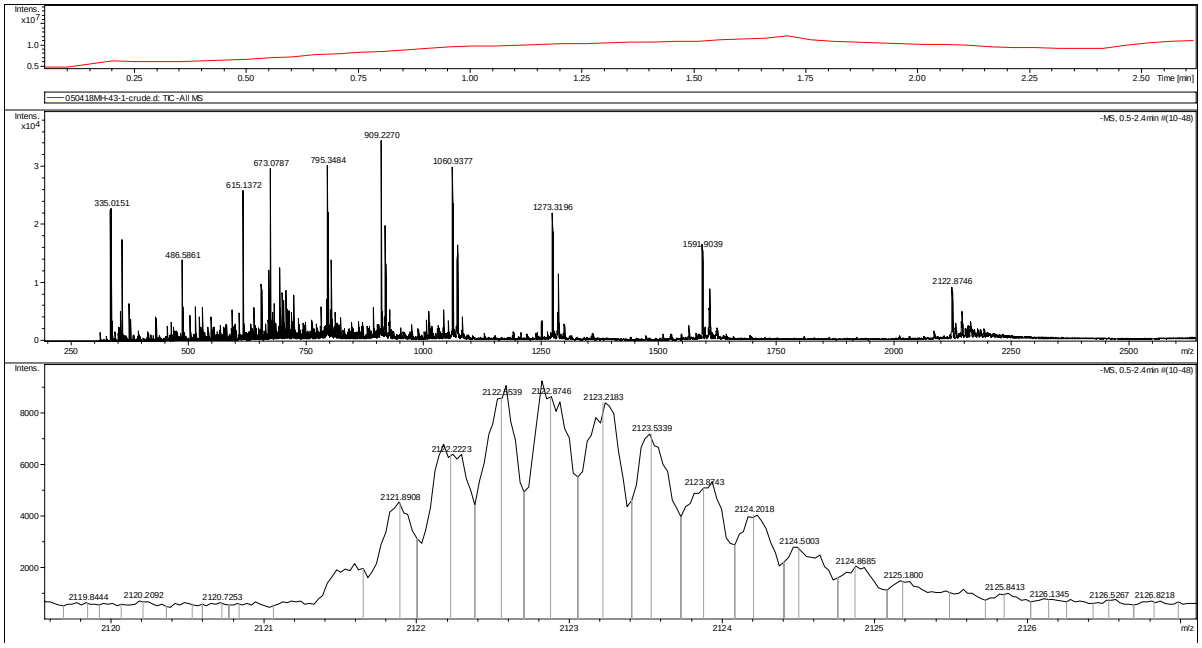

Figure S2. Mass spectrum of oligonucleotide ON1b; m/z calcd for $C_{196}H_{260}N_{59}O_{112}P_{18}S_{18}$: 2122.2265; found: 2122.2223 $[M - 3H]^{3-}$.

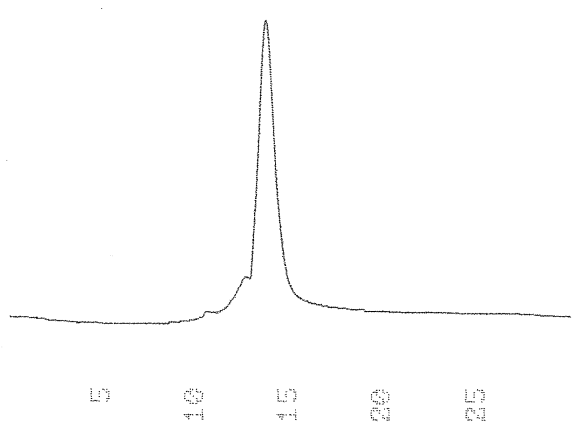

Figure S3. HPLC trace of oligonucleotide ON2b; Hypersil ODS C18 column (250 × 4.6 mm, 5 μm); flow rate = 1.0 mL min^{-1}; linear gradient (0 to 50% over 30 min) of MeCN in 50 mM aqueous triethylammonium acetate.

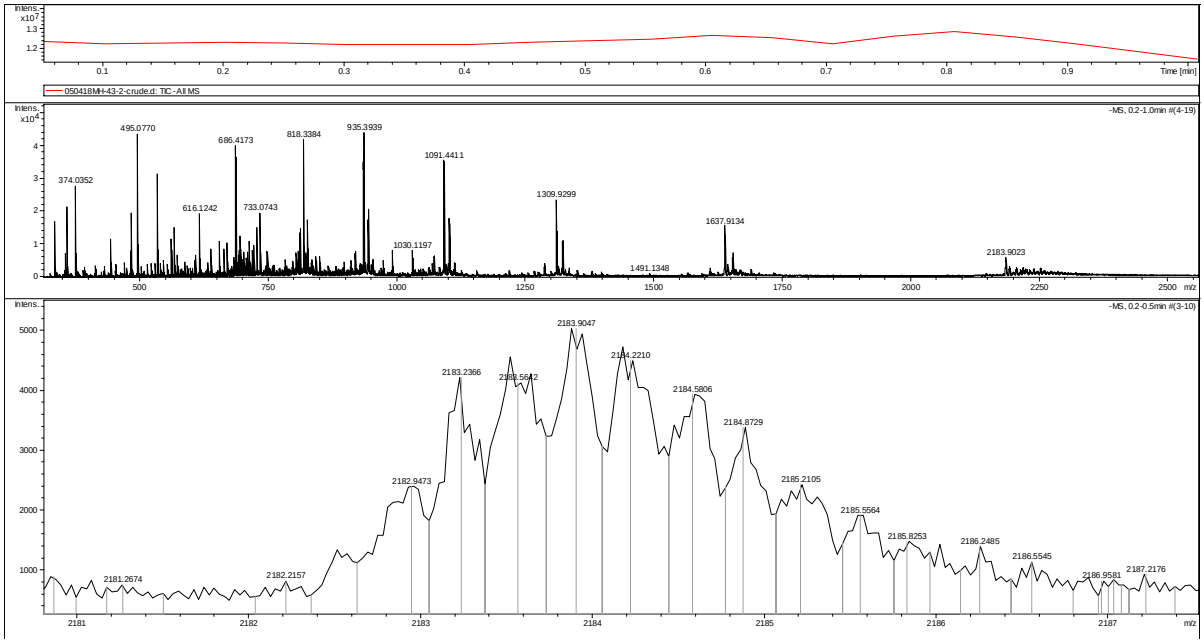

Figure S4. Mass spectrum of oligonucleotide ON2b; m/z calcd for $C_{201}H_{261}N_{70}O_{110}P_{18}S_{18}$: 2183.2438; found: 2183.2366 $[M - 3H]^{3-}$.

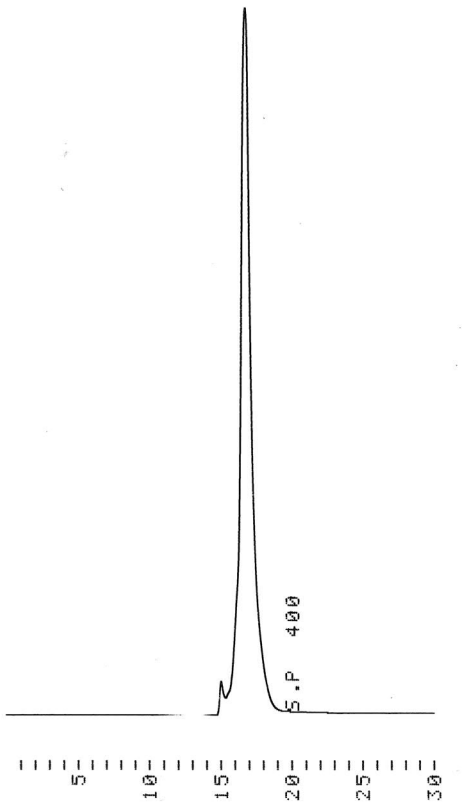

Figure S5. HPLC trace of oligonucleotide ON1b-Pd; Hypersil ODS C18 column (250 × 4.6 mm, 5 μm); flow rate = 1.0 mL min^{-1}; linear gradient (0 to 50% over 30 min) of MeCN in 50 mM aqueous triethylammonium acetate.

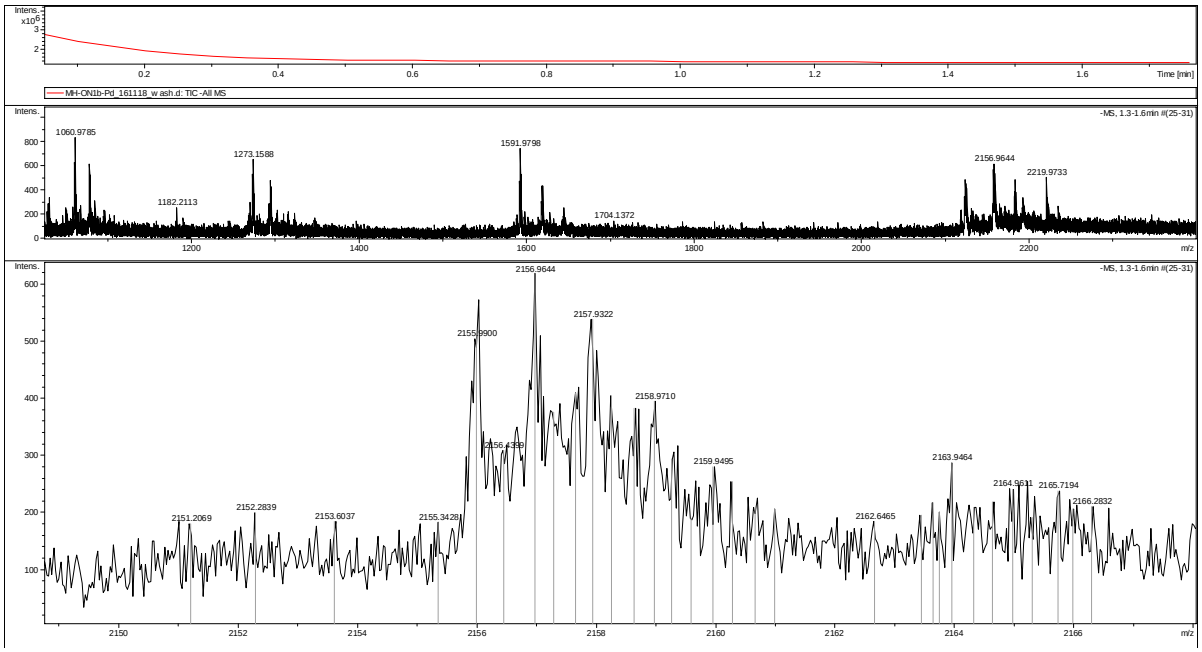

Figure S6. Mass spectrum of oligonucleotide ON1b-Pd; m/z calcd for $C_{196}H_{258}N_{59}O_{112}P_{18}S_{18}Pd$: 2156.8558; found: 2156.9644 $[M - 4H]^{3-}$.

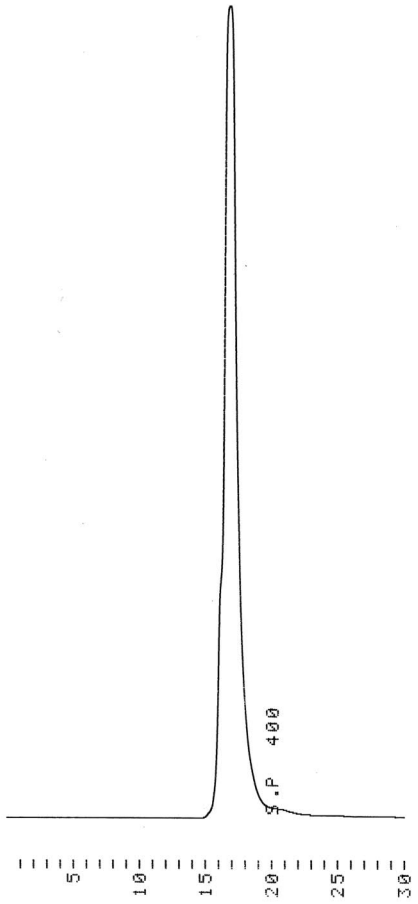

Figure S7. HPLC trace of oligonucleotide ON2b-Pd; Hypersil ODS C18 column (250 × 4.6 mm, 5 μm); flow rate = 1.0 mL min^{-1}; linear gradient (0 to 50% over 30 min) of MeCN in 50 mM aqueous triethylammonium acetate.

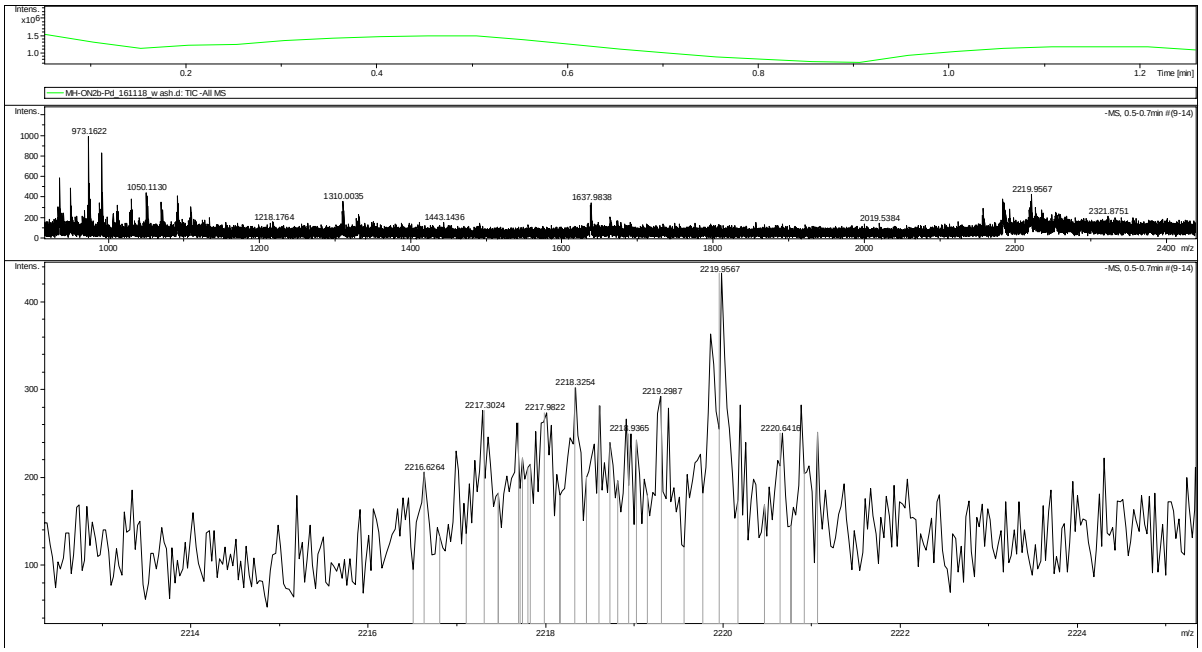

Figure S8. Mass spectrum of oligonucleotide ON2b-Pd; m/z calcd for $C_{201}H_{259}N_{70}O_{110}P_{18}S_{18}Pd$: 2217.8731; found: 2217.9622 $[M - 4H]^{3-}$.

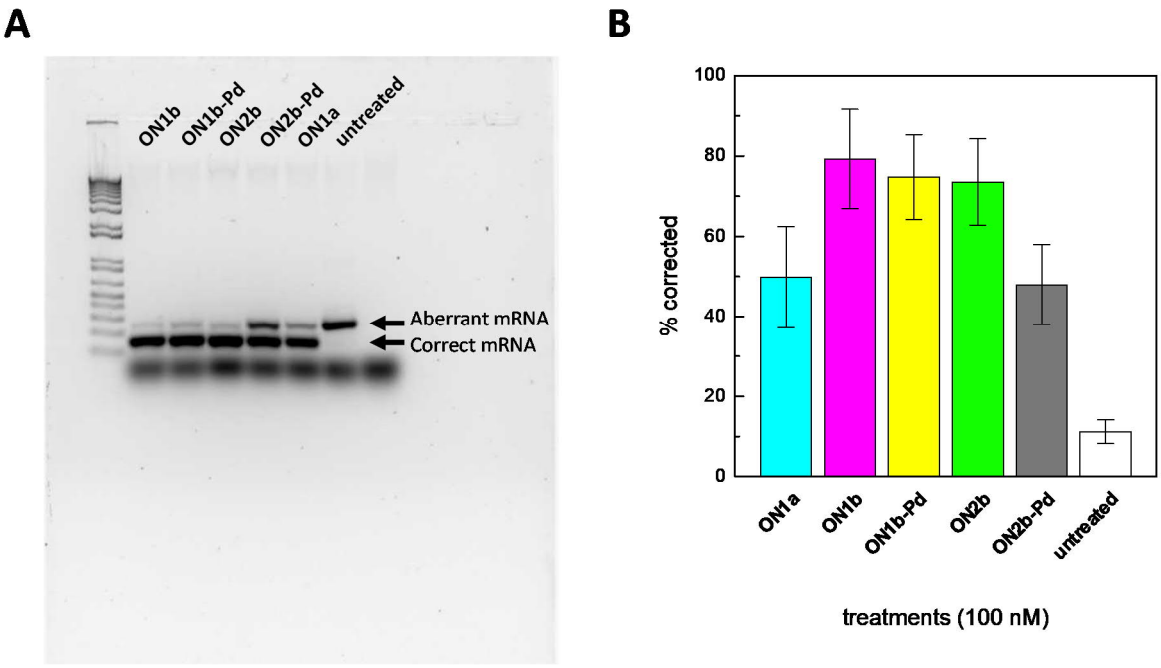

Figure S9. A gel of RT-PCR of luciferase mRNA (A) and efficiency of splice correction (B) in HeLa Luc/705 cells by oligonucleotides ON1a (cyan), ON1b (magenta), ON1b-Pd (yellow), ON2b (green) and ON2b-Pd (grey), as well as without any treatment (white) . The oligonucleotides were delivered by lipofection. Each column represents the mean with the standard error of the mean (SEM) of at least three independent experiments ($n \geq 3$).