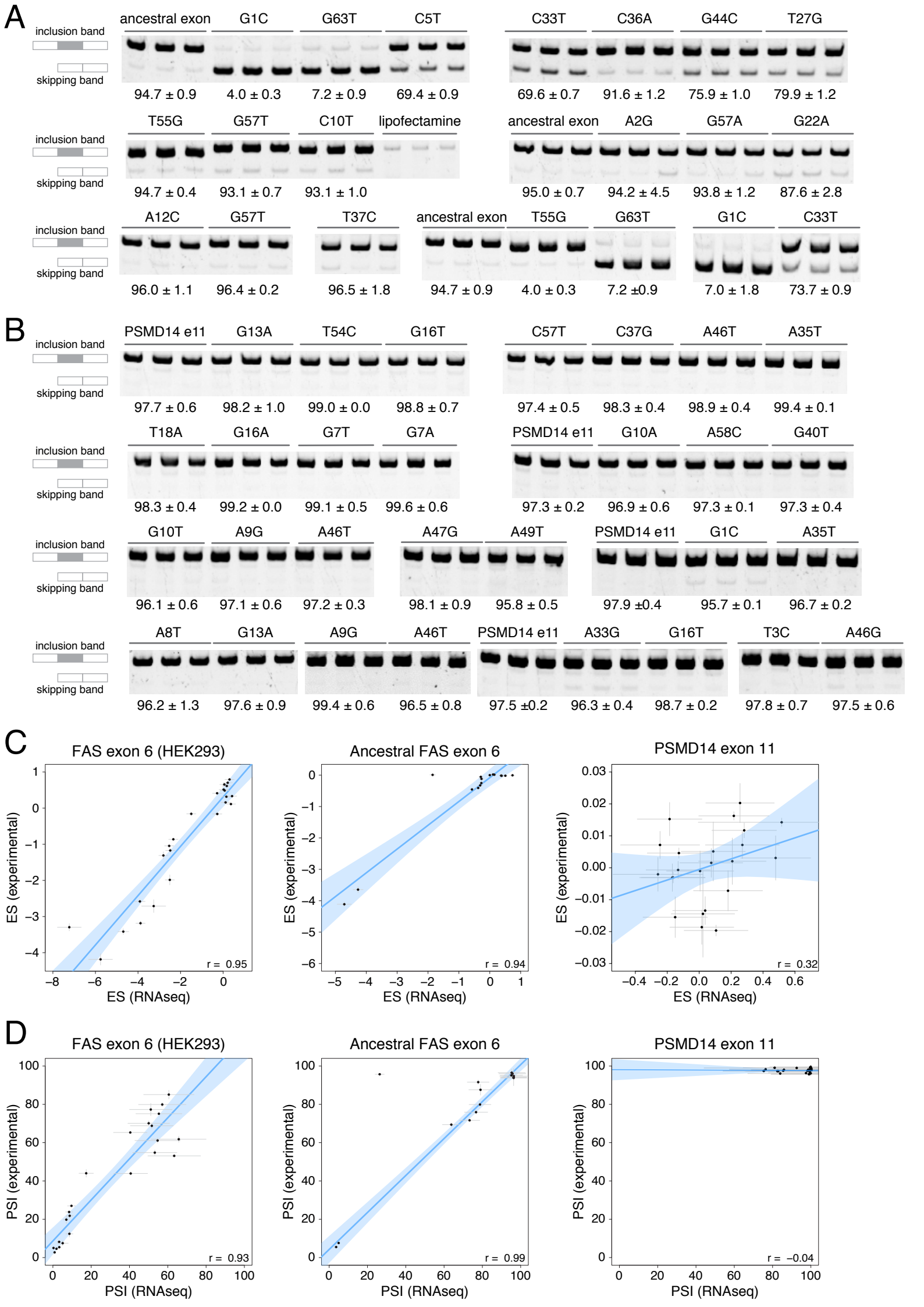
A
inclusion band
skipping band
ancestral exon
G1C
G63T
C5T
94.7 ± 0.9
4.0 ± 0.3
7.2 ± 0.9
69.4 ± 0.9
C33T
C36A
G44C
T27G
69.6 ± 0.7
91.6 ± 1.2
75.9 ± 1.0
79.9 ± 1.2
T55G
G57T
C10T
lipofectamine
94.7 ± 0.4
93.1 ± 0.7
93.1 ± 1.0
ancestral exon
A2G
G57A
G22A
95.0 ± 0.7
94.2 ± 4.5
93.8 ± 1.2
87.6 ± 2.8
A12C
G57T
T37C
96.0 ± 1.1
96.4 ± 0.2
96.5 ± 1.8
ancestral exon
T55G
G63T
G1C
C33T
94.7 ± 0.9
4.0 ± 0.3
7.2 ±0.9
7.0 ± 1.8
73.7 ± 0.9
B
PSMD14 e11
G13A
T54C
G16T
97.7 ± 0.6
98.2 ± 1.0
99.0 ± 0.0
98.8 ± 0.7
C57T
C37G
A46T
A35T
97.4 ± 0.5
98.3 ± 0.4
98.9 ± 0.4
99.4 ± 0.1
T18A
G16A
G7T
G7A
98.3 ± 0.4
99.2 ± 0.0
99.1 ± 0.5
99.6 ± 0.6
PSMD14 e11
G10A
A58C
G40T
97.3 ± 0.2
96.9 ± 0.6
97.3 ± 0.1
97.3 ± 0.4
G10T
A9G
A46T
96.1 ± 0.6
97.1 ± 0.6
97.2 ± 0.3
A47G
A49T
98.1 ± 0.9
95.8 ± 0.5
PSMD14 e11
G1C
A35T
97.9 ±0.4
95.7 ± 0.1
96.7 ± 0.2
A8T
G13A
A9G
A46T
96.2 ± 1.3
97.6 ± 0.9
99.4 ± 0.6
96.5 ± 0.8
PSMD14 e11
A33G
G16T
97.5 ±0.2
96.3 ± 0.4
98.7 ± 0.2
T3C
A46G
97.8 ± 0.7
97.5 ± 0.6
C
FAS exon 6 (HEK293)
ES (experimental)
ES (RNAseq)
r = 0.95
Ancestral FAS exon 6
ES (experimental)
ES (RNAseq)
r = 0.94
PSMD14 exon 11
ES (experimental)
ES (RNAseq)
r = 0.32
D
FAS exon 6 (HEK293)
PSI (experimental)
PSI (RNAseq)
r = 0.93
Ancestral FAS exon 6
PSI (experimental)
PSI (RNAseq)
r = 0.99
PSMD14 exon 11
PSI (experimental)
PSI (RNAseq)
r = −0.04