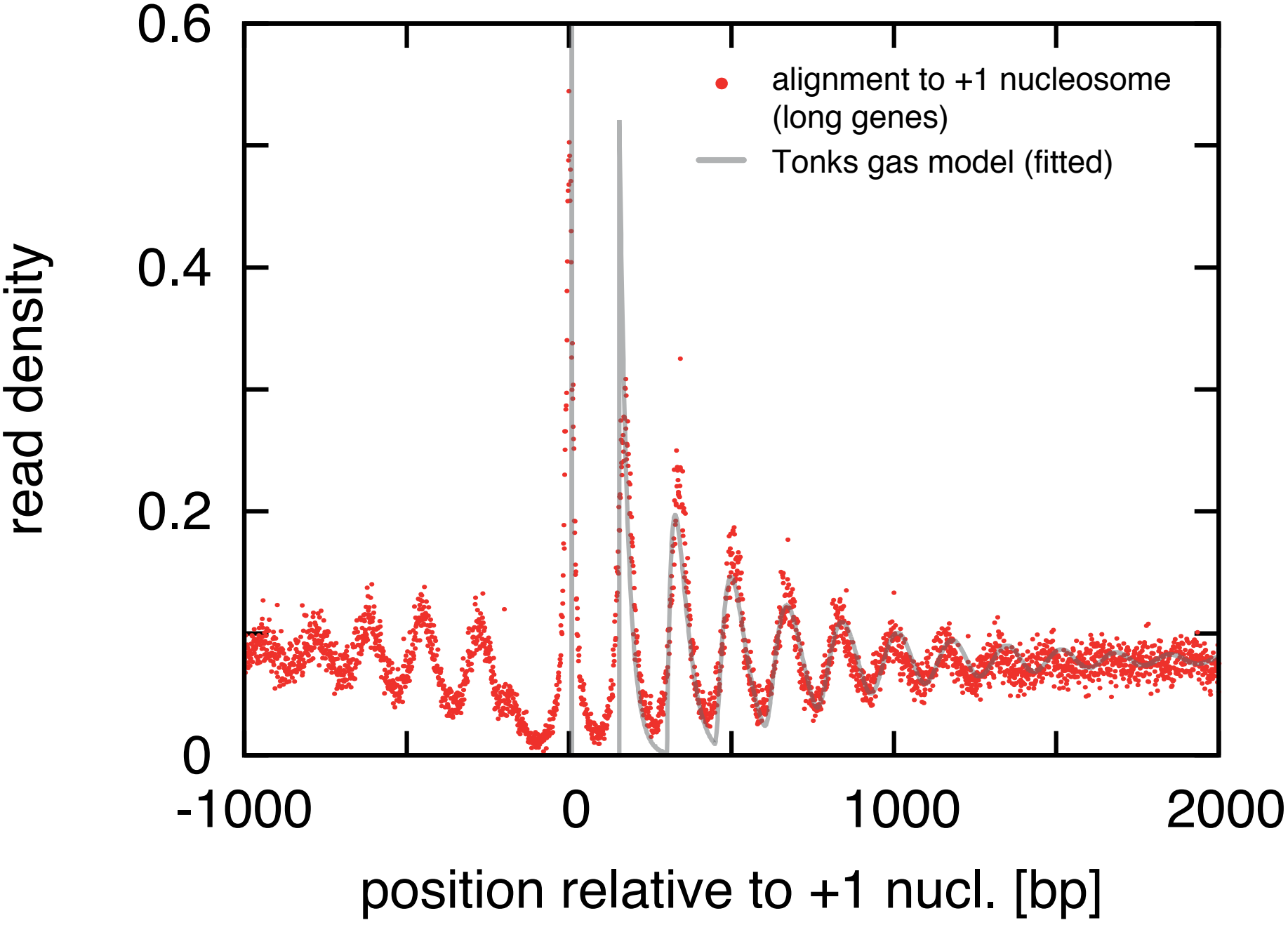
0.6
0.4
0.2
0
read density
-1000
0
1000
2000
position relative to +1 nucl. [bp]
alignment to +1 nucleosome
(long genes)
Tonks gas model (fitted)