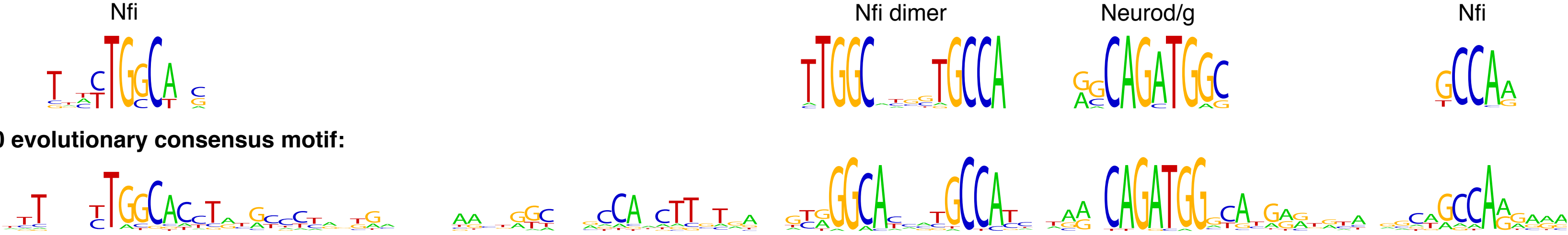
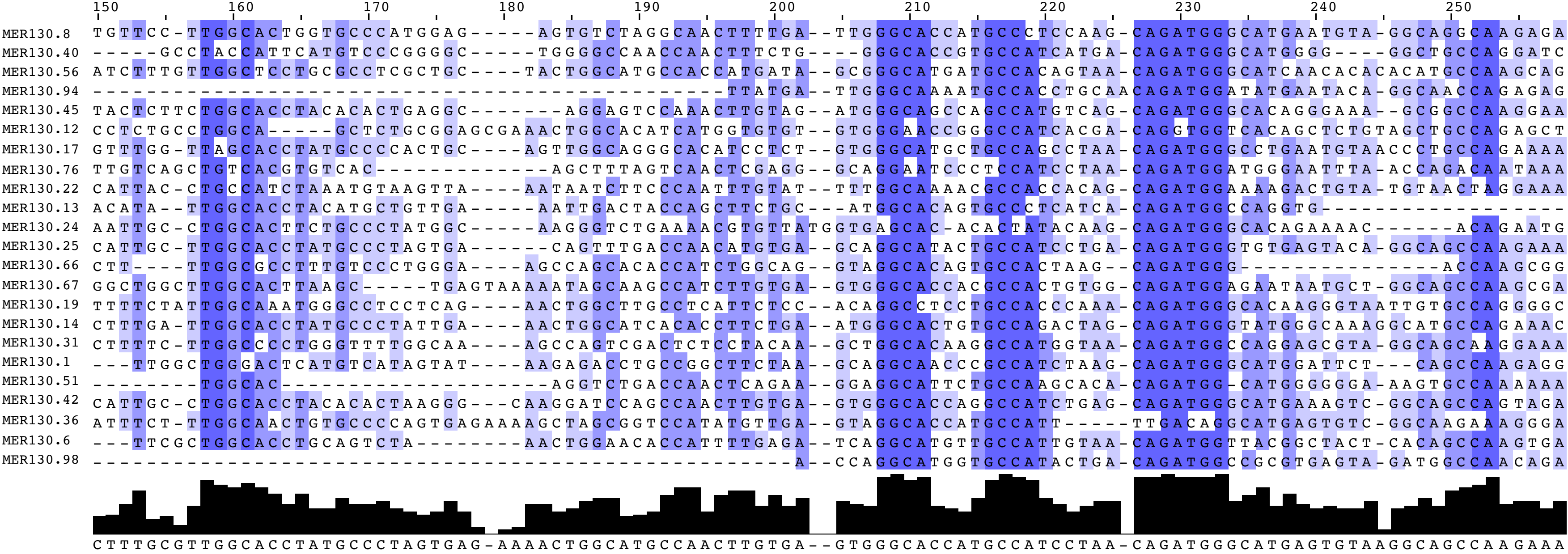


A Transcription factor biophysical binding preferences:



B Multiple alignment:



C MER130.31 transfection mutations:



MER130.45 transfection mutations:



Mut 1

Mut 2

Mut 3

Mut 4

Mut 5