

**Table S4: Unperturbed CSN6<sup>ΔC</sup> residues upon addition of CSN5<sup>ΔC</sup> as evaluated by NMR HSQC.** Residues were considered as unperturbed those with Euclidian distances less than 0.077 ppm (1). Residues marked with an asterisk (\*) are disordered in the human CSN5<sup>ΔC</sup> crystal structure.

C33*	V171
G34*	I172
V35*	D173
T36*	I174
G37*	I175
S40	N176
G78	G177
R79	A179
N80	T180
M84	L182
L90	L186
K97	T187
E109	T189
L119	E196
H139	G199
K140	V200
F151	H202
K159	T209*

## Reference.

1. Williamson, M. P. (2013) Using chemical shift perturbation to characterise ligand binding. *Prog Nucl Magn Reson Spectrosc* **73**, 1-16