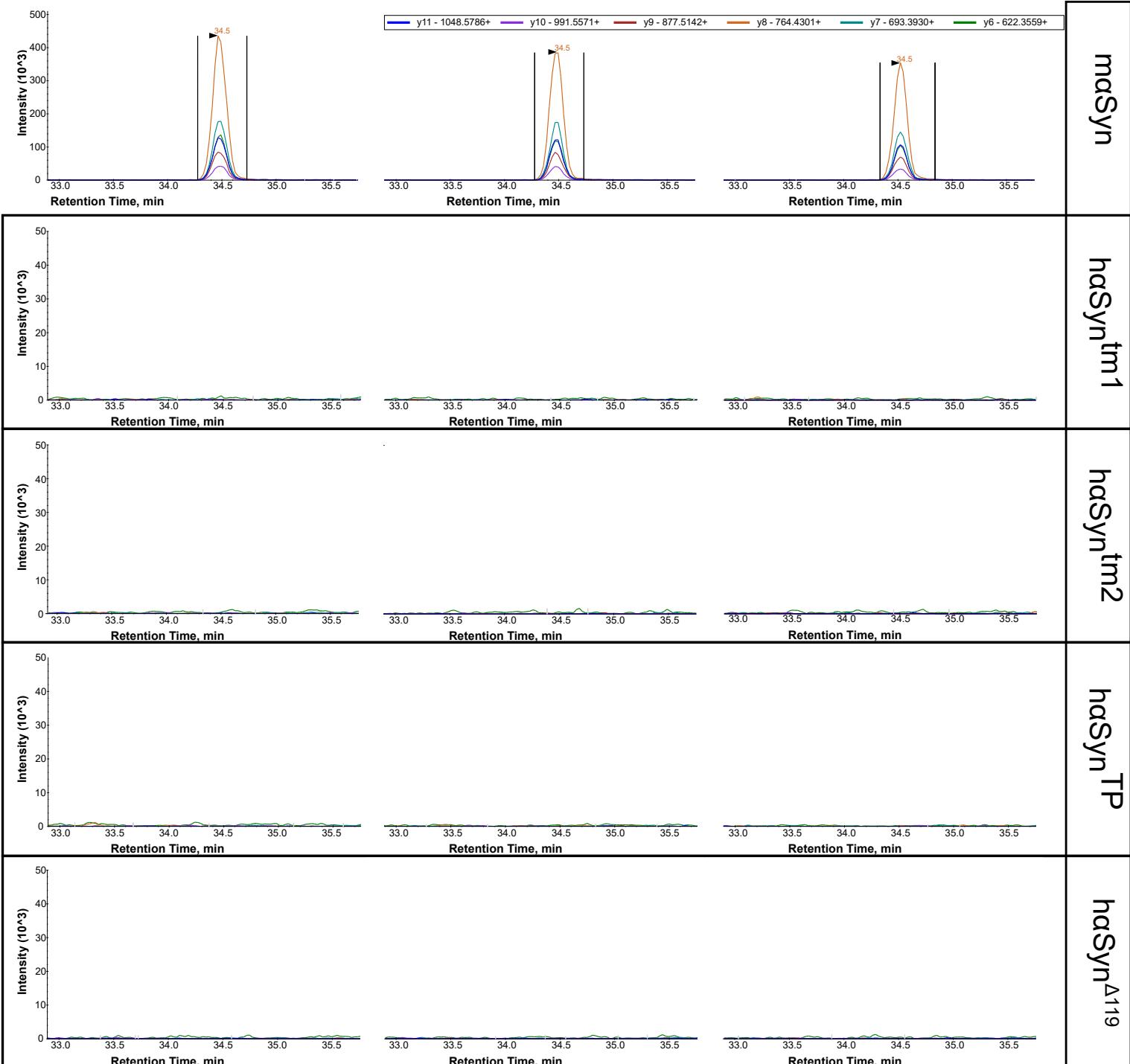


Supp. Fig. 3

81 TVEGAGNIAATGFVK 96 (maSyn)



N-terminus (1 - 60)

maSyn 1 MDVFMKGLSKAEGVAAAEEKTKQGVAEAAGKTKEGVLYVGSKTKEGVVHGVTVAEKTK

NAC domain (61 - 95)

C-terminus (96 - 140)

maSyn 61 EQVTNVGGA~~V~~TGVTAVAQKTVEGAGNIAATGFVKKDQMGKGEEGYPQEGLIEDMPVDP
81 96

maSyn 121 GSEAYEMPSEEGYQDYEPEA 140

Supp. Fig. 3. Elution profiles of the six most-intense fragment ions of an m α Syn-specific peptide 81 – TVEGAGNI-AAATGFVK – 96. Each fragment ion was named as: “y” fragments containing C-terminus, or “b” fragments containing N-terminus, followed by the number of amino acids that the fragment ion consisted of and its corresponding mass (single-charged). We detected this peptide only in wild type mice, confirming that no partial m α Syn was produced in any of the transgenic lines. Note that this peptide is located in the NAC-domain of the m α Syn protein (bottom diagram).