

A **uPA mRNA 3' UTR**

NM_002658

**B****Site 1**

5' UGCCCAGACCCUCU-GGCCAGGAU 3' hsa plau
 | III |IIII|
 3' UGACCCUGAAACAU-CCGGUCA 5' hsa-miR-193a

5' UGCCCAGACCCUCU-GGCCAGGAU 3' hsa plau
 5' UGCCCAGACCCUCU-GGCCAGGAU 3' ptr plau
 5' cauCCAGACCcucCc-GaCCAGGAU 3' mmu plau
 5' cauCCAGACCaucCU-GaCCAGGAU 3' rno plau
 5' UGCCCAGAuCuCUugaGGCCAGGAU 3' cfa plau

Site 2

5' AGAUGUCCCUUUCUUGGCCAGUU 3' hsa plau
 |III| |IIIIII|
 3' UGACCCUGAAACAUCCGGUCA 5' hsa-miR-193a

5' AGAUGUCCCUUUCUUGGCCAGUU 3' hsa plau
 5' AGAUGUCCCUUUCUUGGCCAGUU 3' ptr plau
 5' AGAUGUCCacUUCUUGGCCAGUU 3' mmu plau
 5' AGAUGUCCCcUUCcUGGCCAGUU 3' rno plau
 5' AGAUGUUC-UUUCUUGGCCAGUU 3' cfa plau