C13 Arg – A (21.00%)

>sp|Q6UX06|OLFM4_HUMAN Olfactomedin-4 OS=Homo sapiens GN=OLFM4 PE=1 SV=1

MRPGLSFLLALLFFLGQAAGDLGDVGPPIPSPGFSSFPGVDSSSSFSSSSRSGSSSSRSL
GSGGSVSQLFSNFTGSVDDRGTCQCSVSLPDTTFPVDRVER**LEFTAHVLSQK**FEKELSK**V**
REYVQLISVYEKKLLNLTVRIDIMEKDTISYTELDFELIKVEVKEMEKLVIQLKESFGGS
SEIVDQLEVEIRNMTLLVEK**LETLDKNNVLAIR**REIVALKTKLKECEASKDQNTPVVHPP
PTPGSCGHGGVVNISKPSVVQLNWR**GFSYLYGAWGR**DYSPQHPNK**GLYWVAPLNTDGR**LL
EYYR**LYNTLDDLLLYINAR**ELRITYGQGSGTAVYNNNMYVNMYNTGNIARVNLTTNTIAV
TQTLPNAAYNNRFSYANVAWQDIDFAVDENGLWVIYSTEASTGNMVISKLNDTTLQVLNT
WYTKQYKPSASNAFMVCGVLYATRTMNTR**TEEIFYYYDTNTGK**EGKLDIVMHKMQEK**VQS**
INYNPFDQKLYVYNDGYLLNYDLSVLQKPQ

C13 Arg – B (20.50%)

>sp|Q6UX06|OLFM4_HUMAN Olfactomedin-4 OS=Homo sapiens GN=OLFM4 PE=1 SV=1

MRPGLSFLLALLFFLGQAAGDLGDVGPPIPSPGFSSFPGVDSSSSFSSSSRSGSSSSRSL
GSGGSVSQLFSNFTGSVDDRGTCQCSVSLPDTTFPVDRVER**LEFTAHVLSQK**FEKELSKV
R**EYVQLISVYEK**KLLNLTVRIDIMEKDTISYTELDFELIKVEVKEMEKLVIQLKESFGGS
SEIVDQLEVEIRNMTLLVEK**LETLDKNNVLAIR**REIVALKTKLKECEASKDQNTPVVHPP
PTPGSCGHGGVVNISKPSVVQLNWR**GFSYLYGAWGR**DYSPQHPNK**GLYWVAPLNTDGR**LL
EYYR**LYNTLDDLLLYINAR**ELRITYGQGSGTAVYNNNMYVNMYNTGNIARVNLTTNTIAV
TQTLPNAAYNNRFSYANVAWQDIDFAVDENGLWVIYSTEASTGNMVISKLNDTTLQVLNT
WYTKQYKPSASNAFMVCGVLYATRTMNTR**TEEIFYYYDTNTGK**EGKLDIVMHKMQEK**VQS**
INYNPFDQKLYVYNDGYLLNYDLSVLQKPQ

Supplementary Figure 6f: Amino acid sequence of OLFM4 and its identified peptides.

Figure to show the percentage coverage of the full amino acid sequence of OLFM4 that is represented by the peptide sequences identified in all four SILAC experiments. (The amino acids highlighted in red bold represent identified peptides).