C13 Arg – A (15.20%)

>sp|P04083|ANXA1_HUMAN Annexin A1 OS=Homo sapiens GN=ANXA1 PE=1 SV=2

MAMVSEFLKQAWFIENEEQEYVQTVKSSKGGPGSAVSPYPTFNPSSDVAALHKAIMVKGV
DEATIIDILTKRNNAQRQQIKAAYLQETGKPLDETLKK**ALTGHLEEVVLALLK**TPAQFDA
DELRAAMK**GLGTDEDTLIEILASR**TNKEIRDINRVYREELKRDLAKDITSDTSGDFRNAL
LSLAKGDRSEDFGVNEDLADSDARALYEAGERRKGTDVNVFNTILTTRSYPQLRRVFQKY
TKYSKHDMNKVLDLELKGDIEKCLTAIVKCATSKPAFFAEKLHQAMKGVGTRHKALIRIM
VSRSEIDMNDIKAFYQKMYGISLCQAILDETKGDYEKILVALCGGN

C13 Arg – B (33.30%)

>sp|P04083|ANXA1_HUMAN Annexin A1 OS=Homo sapiens GN=ANXA1 PE=1 SV=2

MAMVSEFLKQAWFIENEEQEYVQTVKSSK**GGPGSAVSPYPTFNPSSDVAALHK**AIMVK**GV**
DEATIIDILTKRNNAQRQQIKAAYLQETGKPLDETLKK**ALTGHLEEVVLALLK**TPAQFDA
DELRAAMK**GLGTDEDTLIEILASR**TNKEIRDINRVYREELKRDLAKDITSDTSGDFRNAL
LSLAKGDRSEDFGVNEDLADSDARALYEAGERRKGTDVNVFNTILTTRSYPQLRRVFQKY
TKYSKHDMNKVLDLELKGDIEKCLTAIVKCATSKPAFFAEKLHQAMKGVGTRHKALIRIM
VSRSEIDMNDIKAFYQKMYGISLCQAILDETKGDYEKILVALCGGN

C13 Lys – A (28.40%)

>sp|P04083|ANXA1_HUMAN Annexin A1 OS=Homo sapiens GN=ANXA1 PE=1 SV=2

MAMVSEFLKQAWFIENEEQEYVQTVKSSK**GGPGSAVSPYPTFNPSSDVAALHK**AIMVKGV
DEATIIDILTKRNNAQRQQIK**AAYLQETGKPLDETLKK**ALTGHLEEVVLALLK**TPAQFDA**
DELRAAMKGLGTDEDTLIEILASRTNKEIRDINRVYREELKRDLAKDITSDTSGDFRNAL
LSLAKGDR**SEDFGVNEDLADSDAR**ALYEAGERRK**GTDVNVFNTILTTR**SYPQLRRVFQKY
TKYSKHDMNKVLDLELKGDIEKCLTAIVKCATSKPAFFAEKLHQAMKGVGTRHKALIRIM
VSRSEIDMNDIKAFYQKMYGISLCQAILDETKGDYEKILVALCGGN

C13 Lys – B (35.70%)

>sp|P04083|ANXA1_HUMAN Annexin A1 OS=Homo sapiens GN=ANXA1 PE=1 SV=2

MAMVSEFLKQAWFIENEEQEYVQTVKSSK**GGPGSAVSPYPTFNPSSDVAALHK**AIMVK**GV**
DEATIIDILTKRNNAQRQQIK**AAYLQETGKPLDETLKK**ALTGHLEEVVLALLK**TPAQFDA**
DELRAAMKGLGTDEDTLIEILASRTNKEIRDINRVYREELKRDLAKDITSDTSGDFRNAL
LSLAKGDR**SEDFGVNEDLADSDAR**ALYEAGERR**KGTDVNVFNTILTTR**SYPQLRRVFQKY
TKYSKHDMNKVLDLELKGDIEKCLTAIVKCATSKPAFFAEKLHQAMKGVGTRHKALIRIM
VSRSEIDMNDIKAFYQKMYGISLCQAILDETKGDYEKILVALCGGN

Supplementary Figure 6c: Amino acid sequence of Annexin A1 and its identified peptides.

Figure to show the percentage coverage of the full amino acid sequence of Annexin A1 that is represented by the peptide sequences identified in all four SILAC experiments. (The amino acids highlighted in red bold represent identified peptides).