

## Supplementary Figure S1

P62988 (UBIQ\_HUMAN) MW = 8560, Ubiquitin

MQIFVKTLTG KTITLEVEPS DTIENVKAKI QDKEGIPPDO QRLIFAGKQL  
EDGRTLSDYN IQESTLHLV LRLRGG

Query	Observed	Mr(expt)	Mr(calc)	ppm	Miss	Score	Expect	Rank	Peptide
<u>6</u>	<u>391.1400</u>	<u>780.2655</u>	<u>780.4204</u>	<u>-198.47</u>	<u>0</u>	<u>33</u>	<u>0.027</u>	<u>1</u>	<u>-.MQIFVK.T</u>
<u>138</u>	<u>534.2090</u>	<u>1066.4035</u>	<u>1066.6135</u>	<u>-196.88</u>	<u>0</u>	<u>49</u>	<u>0.00052</u>	<u>1</u>	<u>K.ESTLHLVLR.L</u>
<u>142</u>	<u>541.1877</u>	<u>1080.3609</u>	<u>1080.5451</u>	<u>-170.48</u>	<u>0</u>	<u>34</u>	<u>0.019</u>	<u>1</u>	<u>R.TLSDynIQK.E</u>
<u>286</u>	<u>508.5109</u>	<u>1522.5109</u>	<u>1522.7740</u>	<u>-172.76</u>	<u>1</u>	<u>62</u>	<u>2.1e-005</u>	<u>1</u>	<u>K.IQDKEGIPPDQQR.L</u>
<u>361</u>	<u>894.3104</u>	<u>1786.6063</u>	<u>1786.9200</u>	<u>-175.53</u>	<u>0</u>	<u>59</u>	<u>4.3e-005</u>	<u>1</u>	<u>K.TITLEVEPSDTIENVK.A</u>
<u>409</u>	<u>662.9076</u>	<u>1985.7011</u>	<u>1986.0521</u>	<u>-176.72</u>	<u>1</u>	<u>79</u>	<u>3.5e-007</u>	<u>1</u>	<u>K.TITLEVEPSDTIENVKAK.I</u>

P25815 (S100P\_HUMAN) MW= 10400, S100P

MTELETAMGM IIDVFSRYSG SEGSTQTLTK GELKVLMEKE LPGFLQSGKD KDAVDKLLKD  
LDANGDAQVD FSEFIVFVAA ITSACHKYFE KAGLK

Query	Observed	Mr(expt)	Mr(calc)	ppm	Miss	Score	Expect	Rank	Unique	Peptide
<u>199</u>	<u>538.2917</u>	<u>1074.5688</u>	<u>1074.5710</u>	<u>-1.96</u>	<u>0</u>	<u>29</u>	<u>1.9</u>	<u>1</u>	<u>U</u>	<u>K.ELPGFLQSGK.D</u>
<u>373</u>	<u>679.8246</u>	<u>1357.6346</u>	<u>1357.6361</u>	<u>-1.10</u>	<u>0</u>	<u>85</u>	<u>3.3e-006</u>	<u>1</u>	<u>U</u>	<u>R.YSGSEGSTQTLTK.G</u>

Figure S1. Tryptic peptides (underlined in the full sequence) used to identify ubiquitin and S100P by MASCOT searching.