

Table 1 Peptides and MRM Transitions used for relative quantification

<b>Protein/Peptide</b>	<b>Q1 (charge)</b>	<b>Q3 (type, charge)</b>
<b>P15127_IR</b>		
TIDSVTSAQELR	660.3 (+2)	804.4 (2y <sup>7</sup> , +1)
TVNESASLR	488.8 (+2)	776.4 (2y <sup>7</sup> , +1)
<b>O35567_ATIC</b>		
NLASLGLSLVASGGTAK	779.9 (+2)	890.5 (2y <sup>10</sup> , +1)
<b>D4ABI7_PTPLAD-1</b>		
QVNITVQK	465.3 (+2)	588.4 (2y <sup>5</sup> , +1)
LESEGSPETLTNLK	759.4 (+2)	915.5 (2y <sup>8</sup> , +1)