

## Supplementary Data

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# Mass Spectrometric Characterization of Amyloid- $\beta$ Species in the 7PA2 Cell Model of Alzheimer's Disease

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Supplementary Table 1  
(Continued)

Peptide	Theoretical mass [Da]	$\Delta m$ [ppm] <sup>a</sup>	MS/MS <sup>b</sup>	Effect of BACE1 inhibition <sup>c</sup>
A $\beta$ <sub>1-17</sub>	2066.956	-0.1	Y	↓
A $\beta$ <sub>1-18</sub>	2166.024	+0.3	Y	↓
A $\beta$ <sub>1-19</sub>	2313.093	-1.9	Y	↓
A $\beta$ <sub>1-20</sub>	2460.161	-1.2	Y	↓
A $\beta$ <sub>1-27</sub>	3132.433	-0.7	Y	↓
A $\beta$ <sub>1-28</sub>	3260.528	-1.0		→
A $\beta$ <sub>1-29</sub>	3317.549	-0.6	Y	→
A $\beta$ <sub>1-30</sub>	3388.586	-1.1		↓
A $\beta$ <sub>1-31</sub>	3501.670	-1.6		↓
A $\beta$ <sub>1-32</sub>	3614.754	+0.7	Y	↓
A $\beta$ <sub>1-33</sub>	3671.776	-0.4	Y	↓
A $\beta$ <sub>1-34</sub>	3784.860	+1.0	Y	↓
A $\beta$ <sub>1-37</sub>	4071.990	+0.9	Y	→
A $\beta$ <sub>1-38</sub>	4129.012	+2.1	Y	↓
A $\beta$ <sub>1-39</sub>	4228.080	+0.1	Y	→
A $\beta$ <sub>1-40</sub>	4327.148	+1.7	Y	→
A $\beta$ <sub>1-41</sub>	4440.232	+2.5	Y	↓
A $\beta$ <sub>1-42</sub>	4511.270	+1.8	Y	→
A $\beta$ <sub>3-17</sub>	1880.892	-1.0		↓
A $\beta$ <sub>3-20</sub>	2274.097	-0.5	Y	↓
A $\beta$ <sub>5-29</sub>	2855.374	-0.9	Y	↑
A $\beta$ <sub>5-33</sub>	3209.601	-0.5	Y	↑
A $\beta$ <sub>5-38</sub>	3666.837	+0.9		↑
A $\beta$ <sub>5-39</sub>	3765.905	+0.4		↑
A $\beta$ <sub>5-40</sub>	3864.973	+1.5	Y	↑
A $\beta$ <sub>5-41</sub>	3978.057	+2.6		↑
A $\beta$ <sub>5-42</sub>	4049.095	+4.4	Y	↑
A $\beta$ <sub>11-34</sub>	2607.381	+0.9	Y	↓
A $\beta$ <sub>11-38</sub>	2951.533	+0.4		↓
A $\beta$ <sub>11-39</sub>	3050.601	+2.8		↓
A $\beta$ <sub>11-40</sub>	3149.670	+2.3	Y	↓
A $\beta$ <sub>11-42</sub>	3333.791	+1.3		↓
A $\beta$ <sub>17-38</sub>	2193.150	-0.5		↑
A $\beta$ <sub>17-39</sub>	2292.219	+3.5	Y	↑
A $\beta$ <sub>17-40</sub>	2391.287	+1.9		↑

<sup>a</sup>  $\Delta m$  denotes the relative experimental mass deviation,  $\Delta m = (m_{\text{exp}} - m_{\text{theory}})/m_{\text{theory}}$ ; <sup>b</sup> Y indicates identification by MS/MS; <sup>c</sup> Denotes the relative change in BACE1 inhibited cells compared to untreated cells. NB: this is a label free determination using the peak area ratios from LC-MS/MS analyses. An up or down arrow denotes a relative change of at least 3 times, while a right arrow indicates a relative change of less than 3 times.