

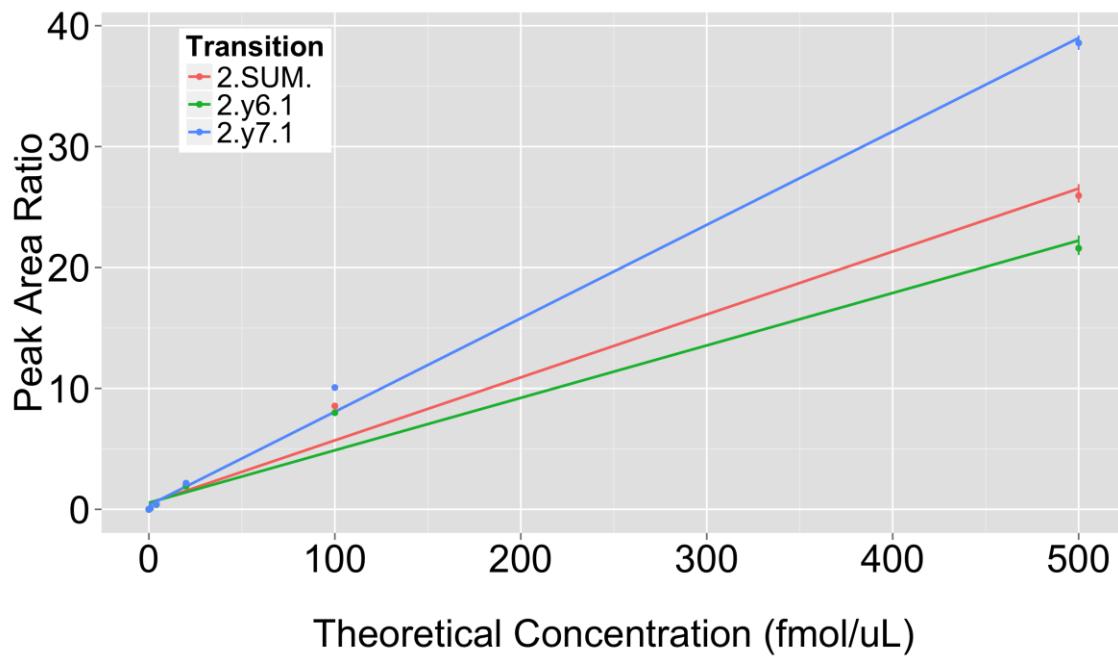
Supplemental Figure 2. Immuno-MRM response curves plotted in linear and log space. The heavy-to-light peak area ratios for the peptide's transitions are plotted vs. the heavy peptide's theoretical concentration. The abbreviations (e.g. 2.y6.1) refer to the precursor ion's charge state (+2), followed by the transition ion and the transition ion's charge state (+1); 2.SUM. refers to the response curve for which the peak areas of all transitions were summed before taking the peak area ratios; the error bars indicate the minimum and maximum (i.e., range) of the peak area ratios of the three capture and LC-MRM-MS replicates.

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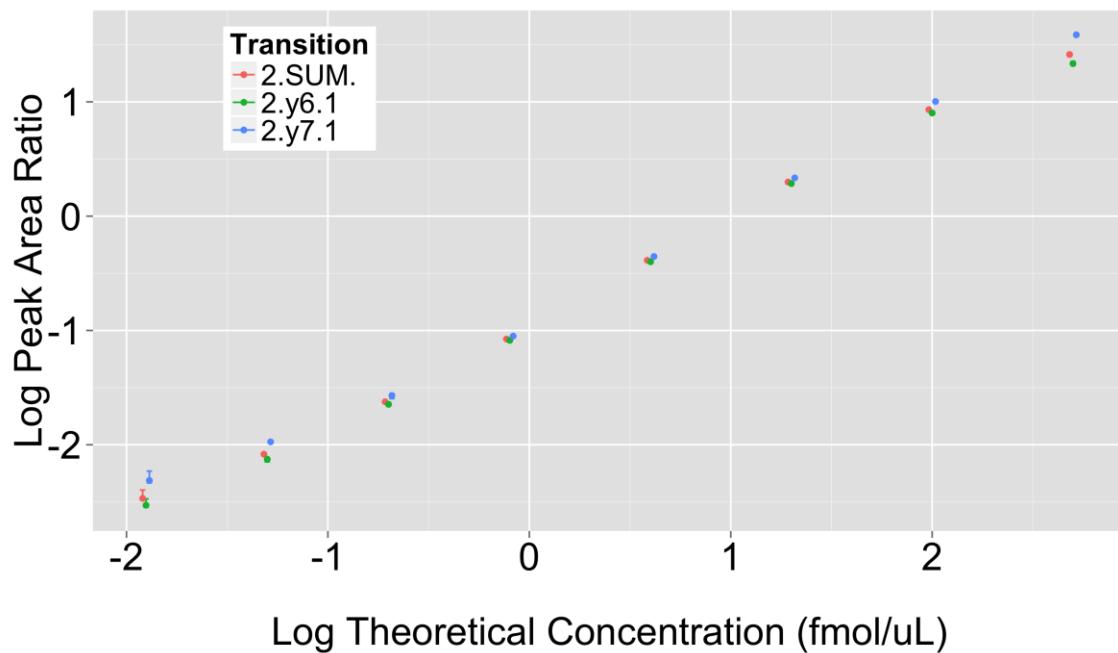
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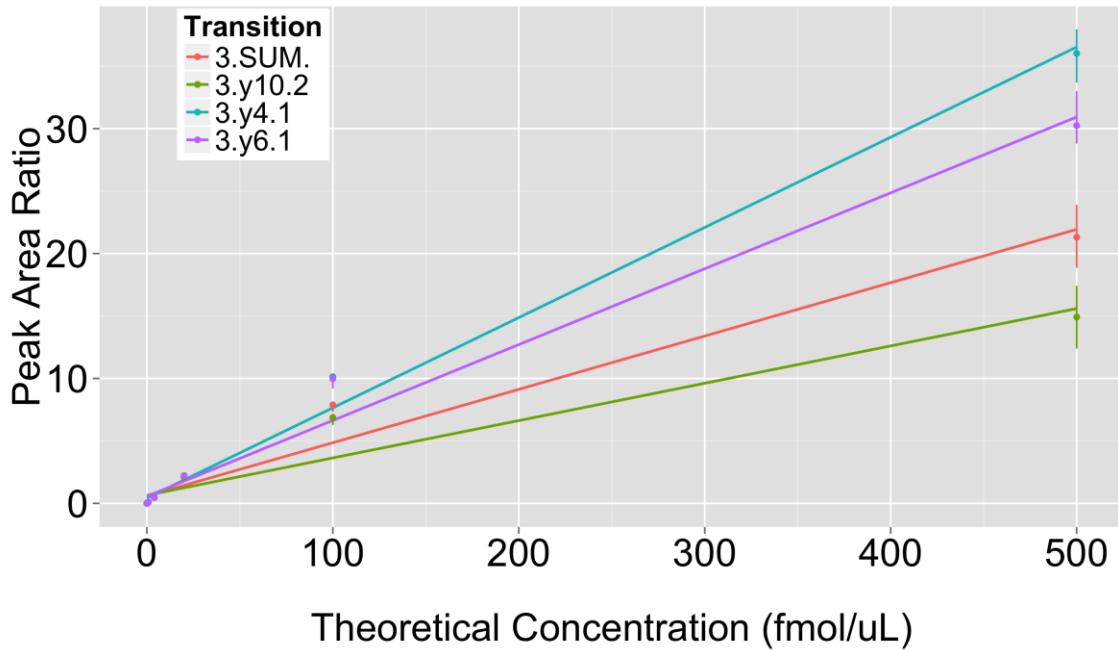
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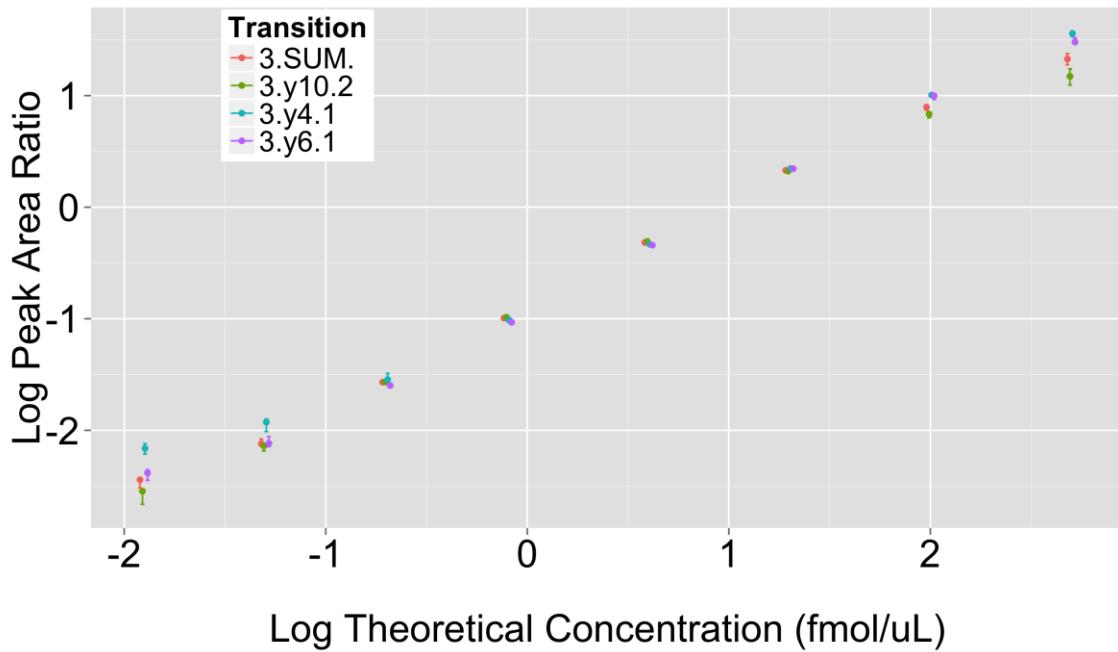
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Analyte: SAIC02A.GLQSLPTHDP SPLQR

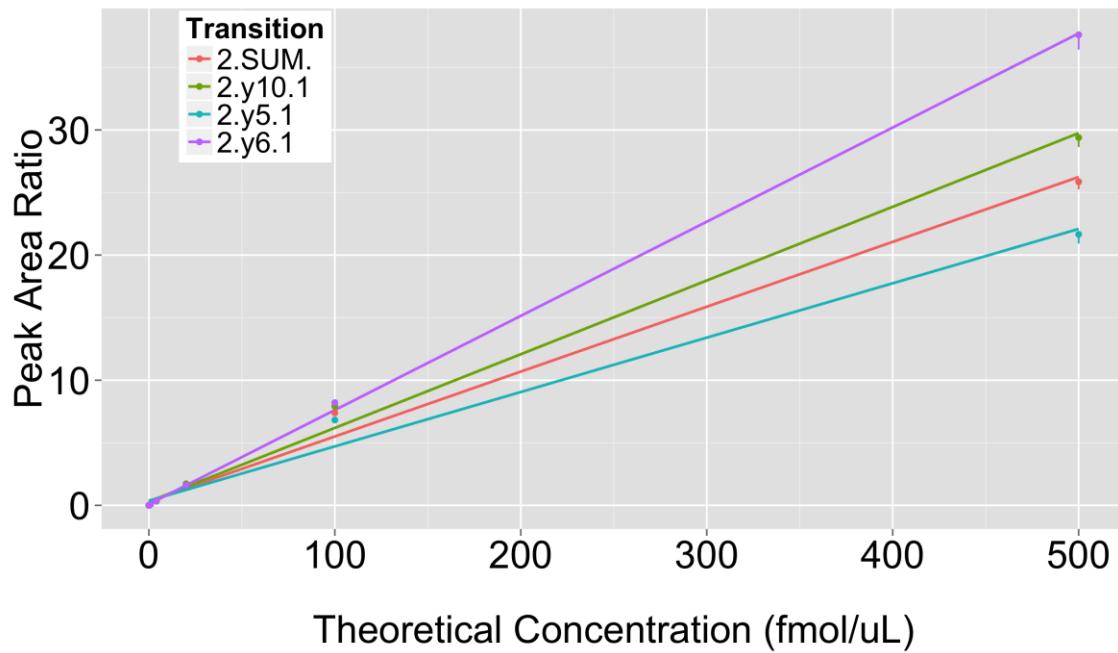


Analyte: SAIC02A.GLQSLPTHDP SPLQR

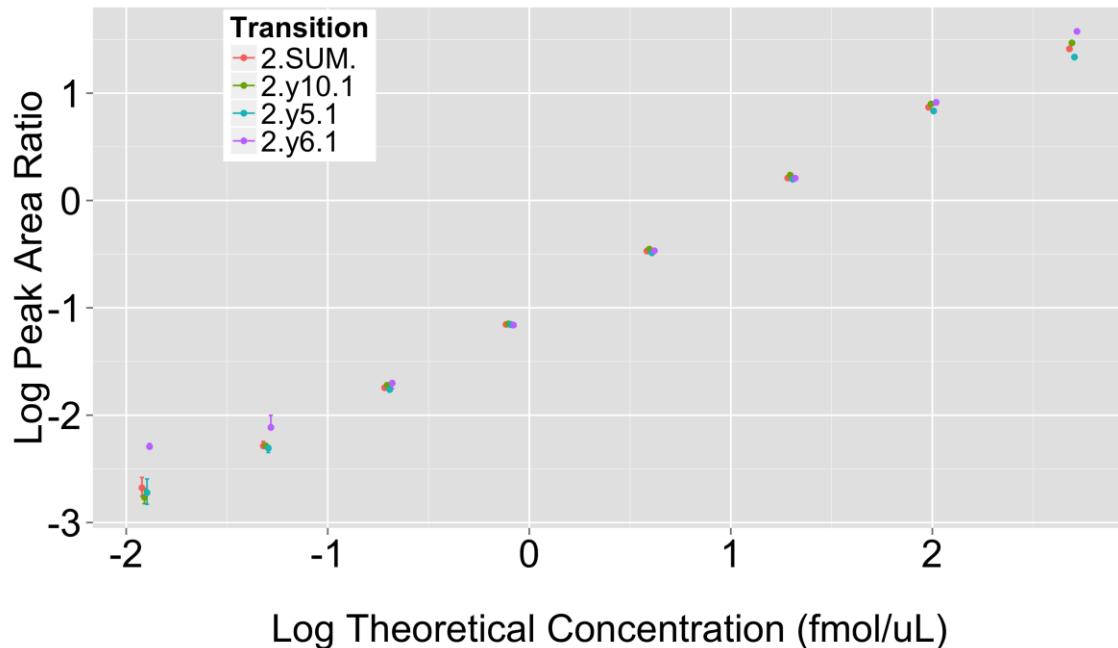


SAIC-07B, XPNPEP1, GSLTFEPLTLVPIQTK.

Analyte: SAIC07B.GSLTFEPLTLVPIQTK

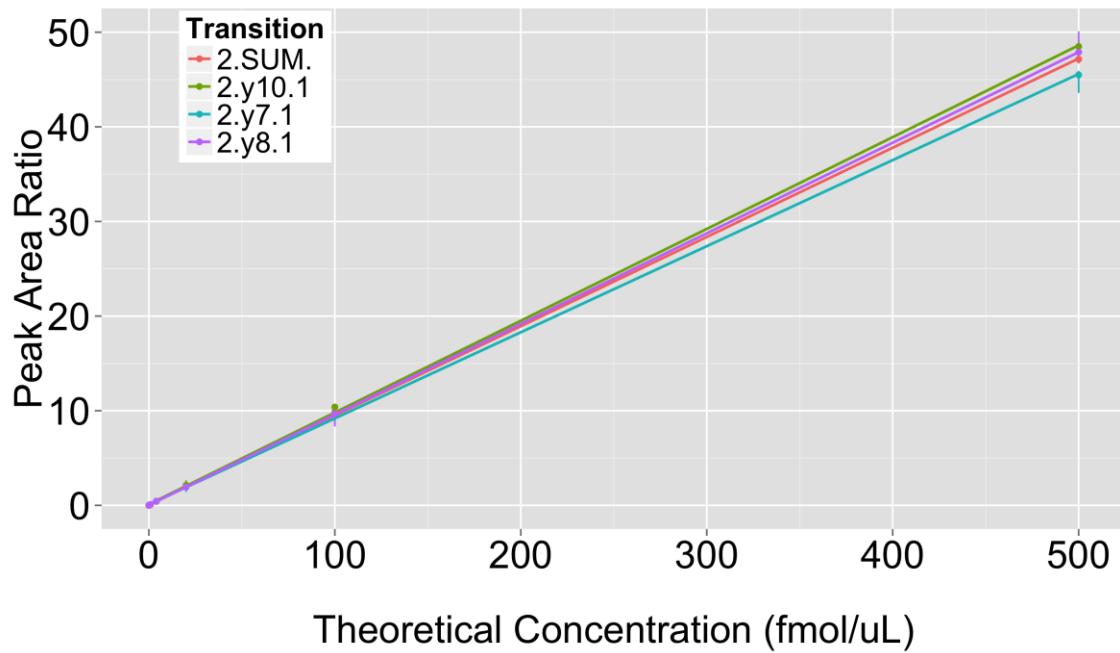


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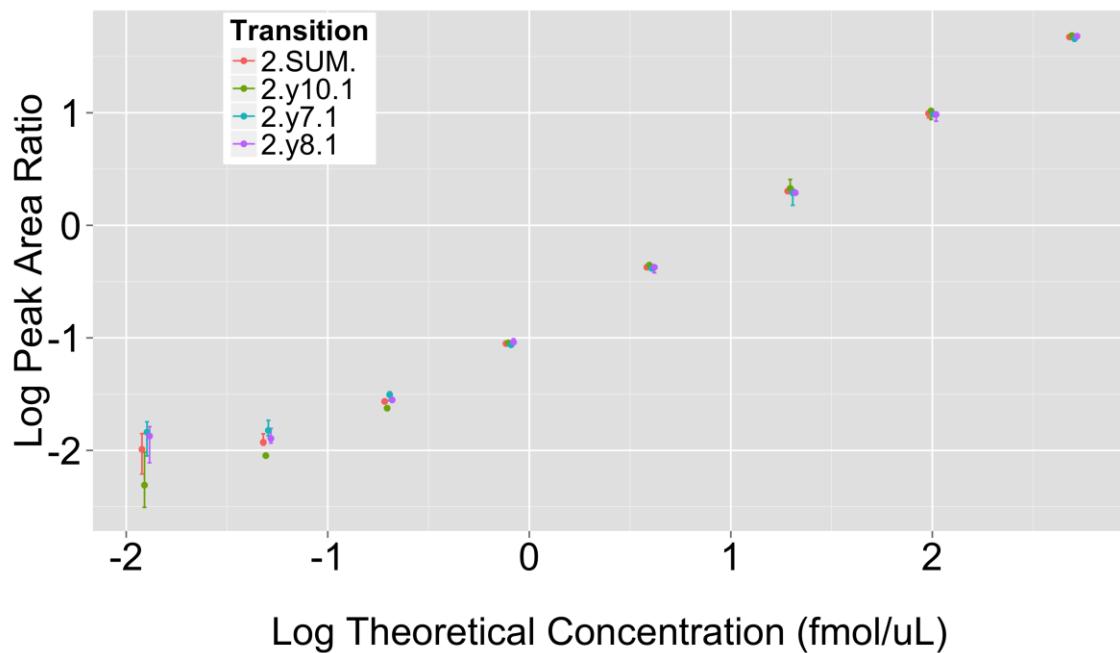


SAIC-07E, XPNPEP1, TLSLDEVYLIDSGAQYK.

Analyte: SAIC07E.TLSLDEVYLIDSGAQYK

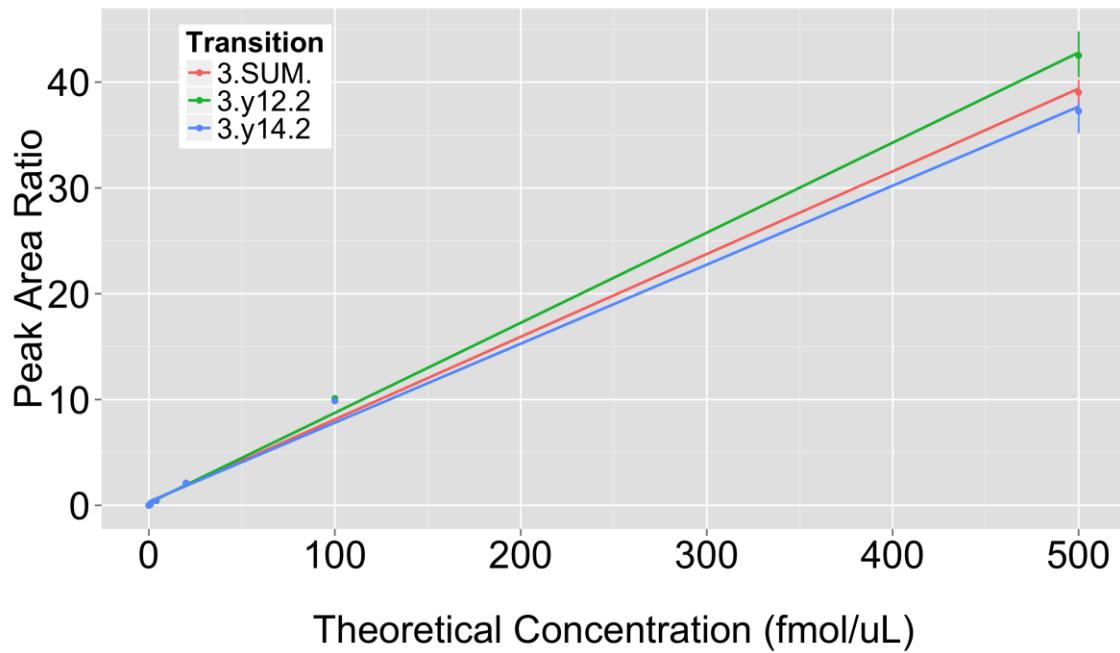


Analyte: SAIC07E.TLSLDEVYLIDSGAQYK

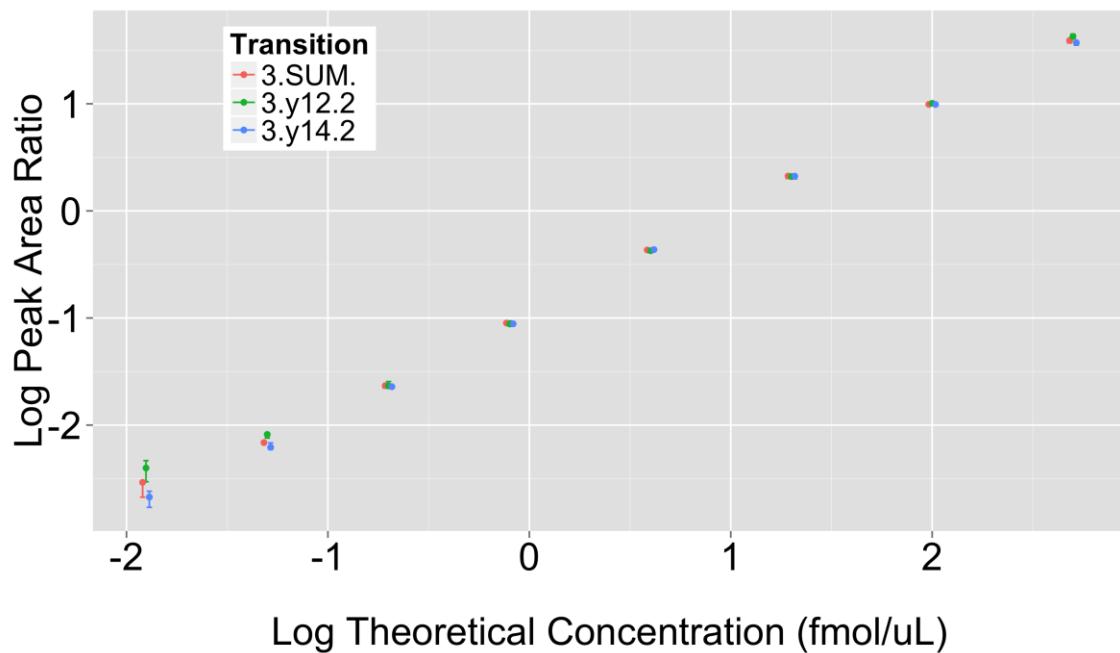


SAIC-13B, ANXA1, AAYLQETGKPLDETLK.

Analyte: SAIC13B.AAYLQETGKPLDETLK

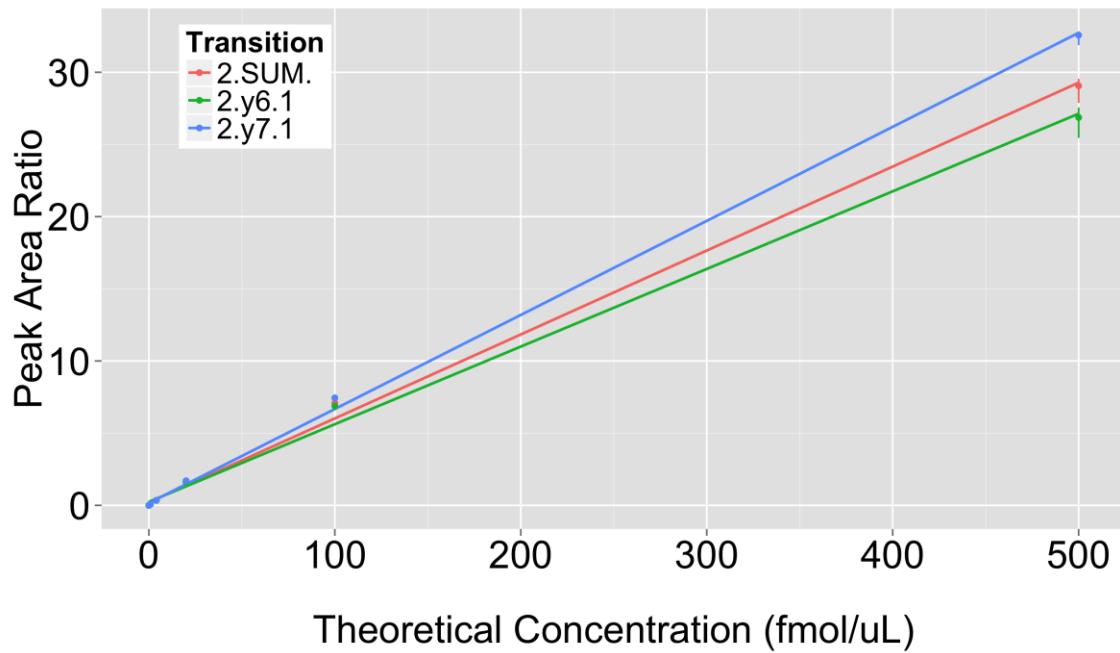


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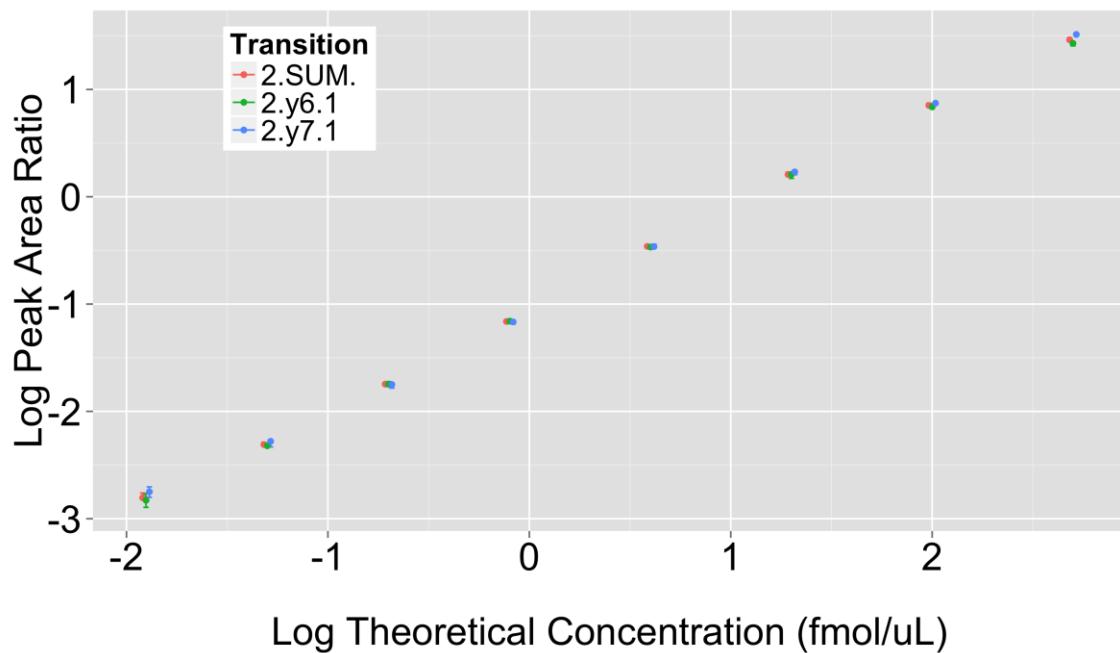


SAIC-13C, ANXA1, GVDEATIIDILTK.

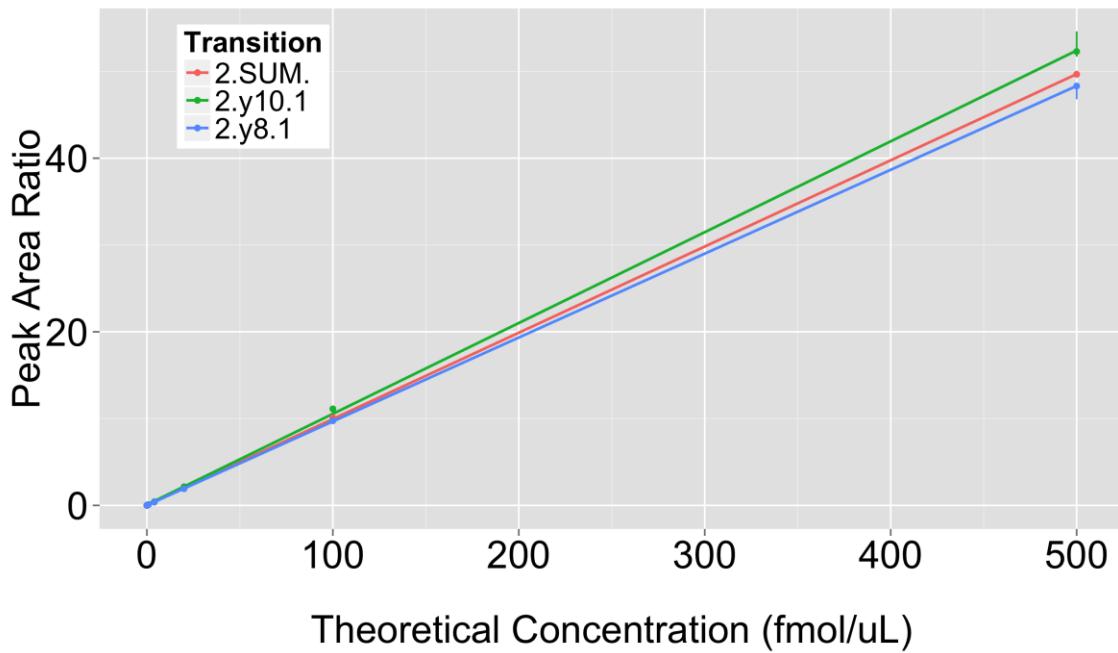
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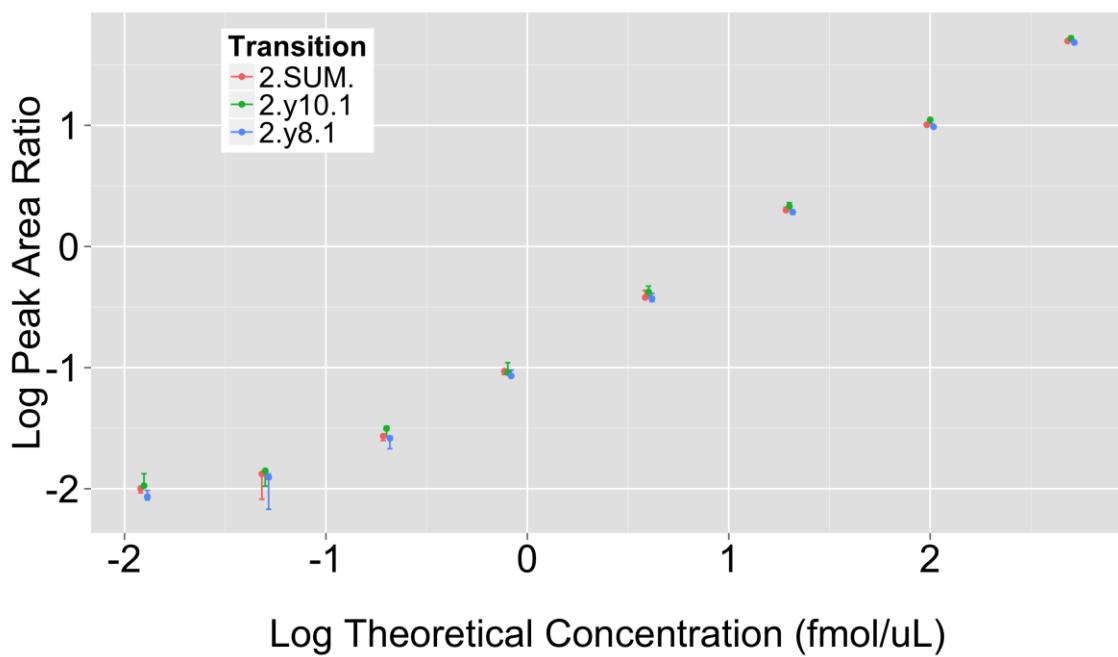
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Analyte: SAIC14C.DEGNYLDDALVR

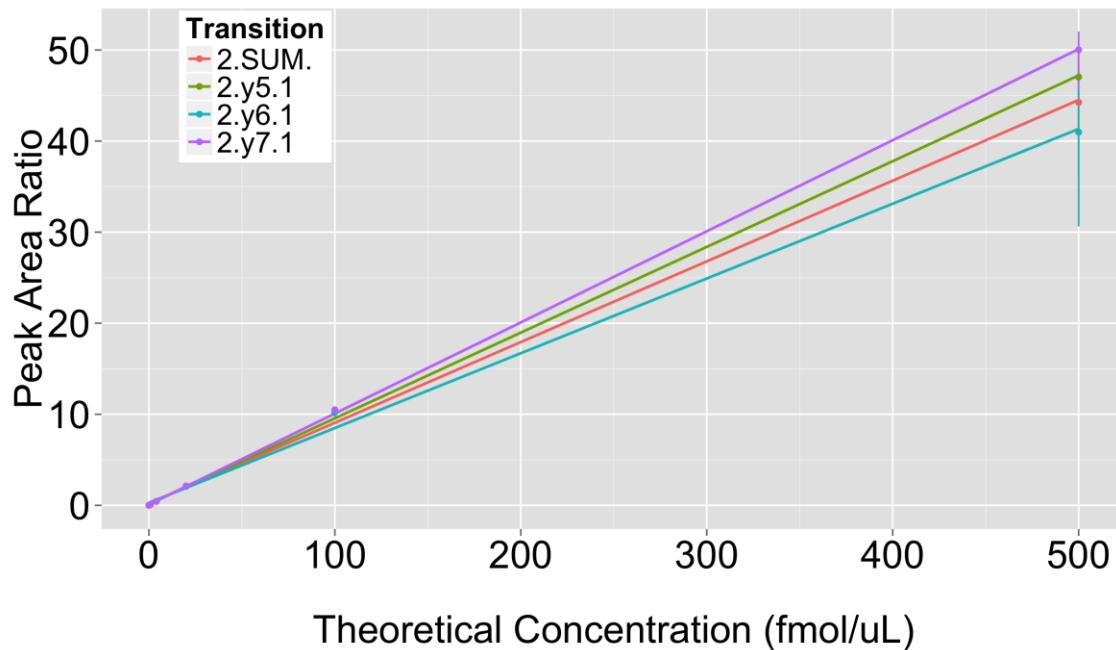


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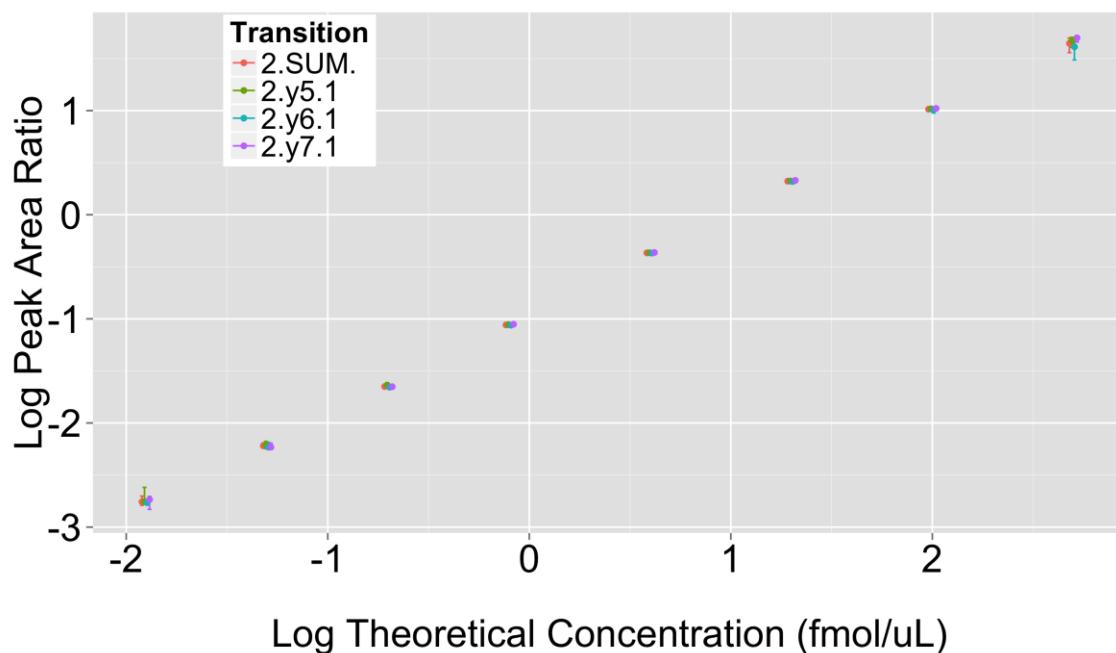


SAIC-16D, CALR, GLQTSQDAR.

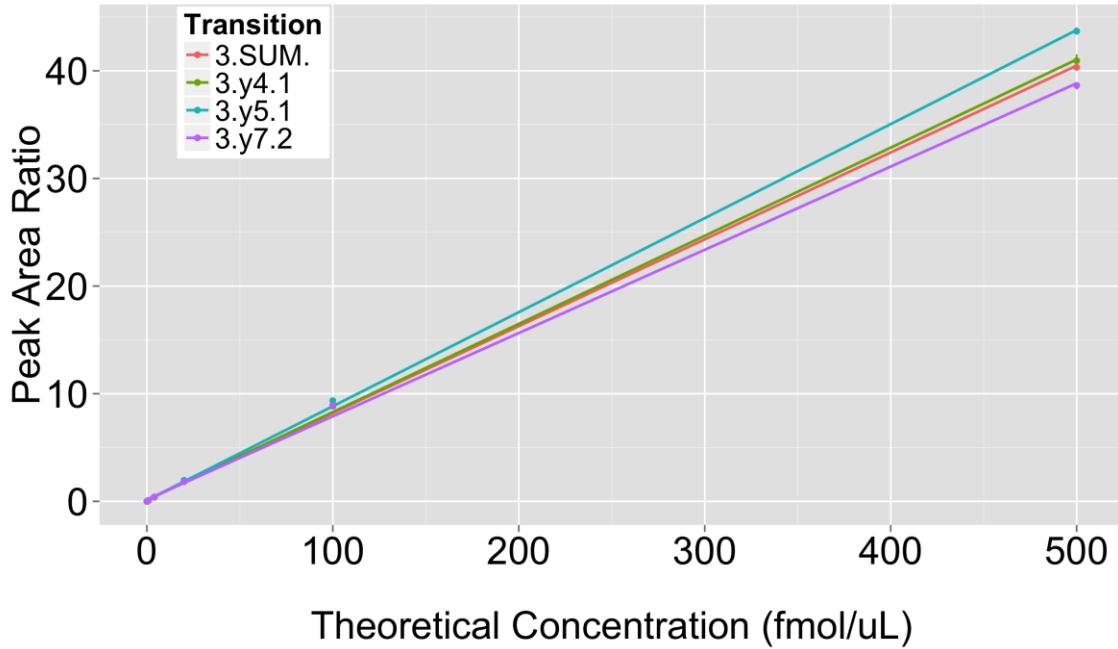
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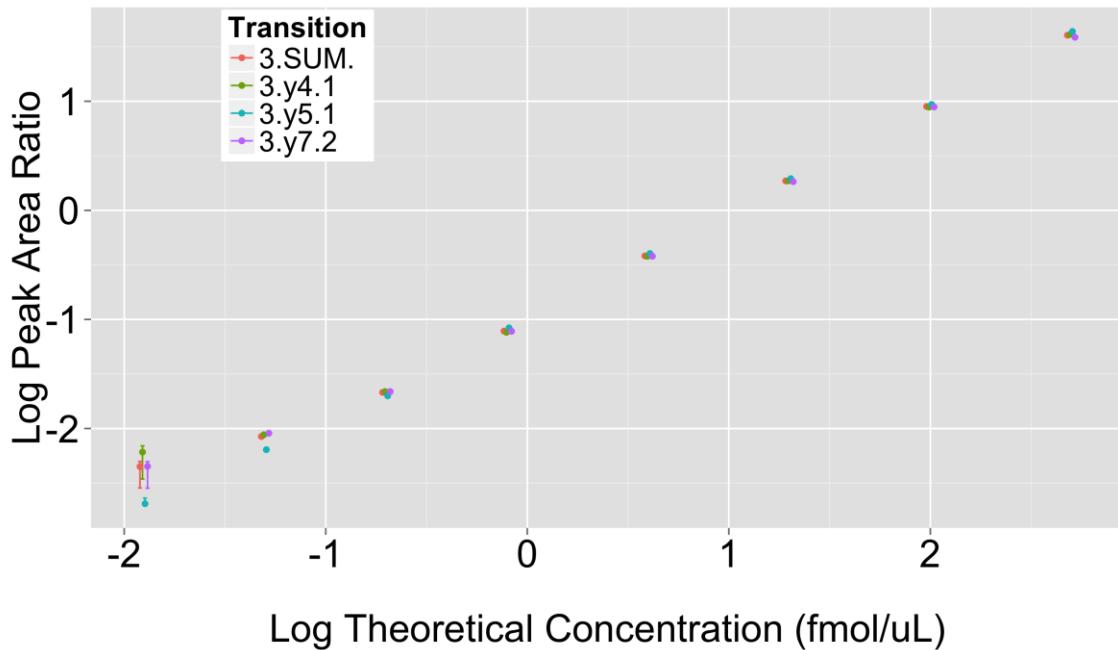
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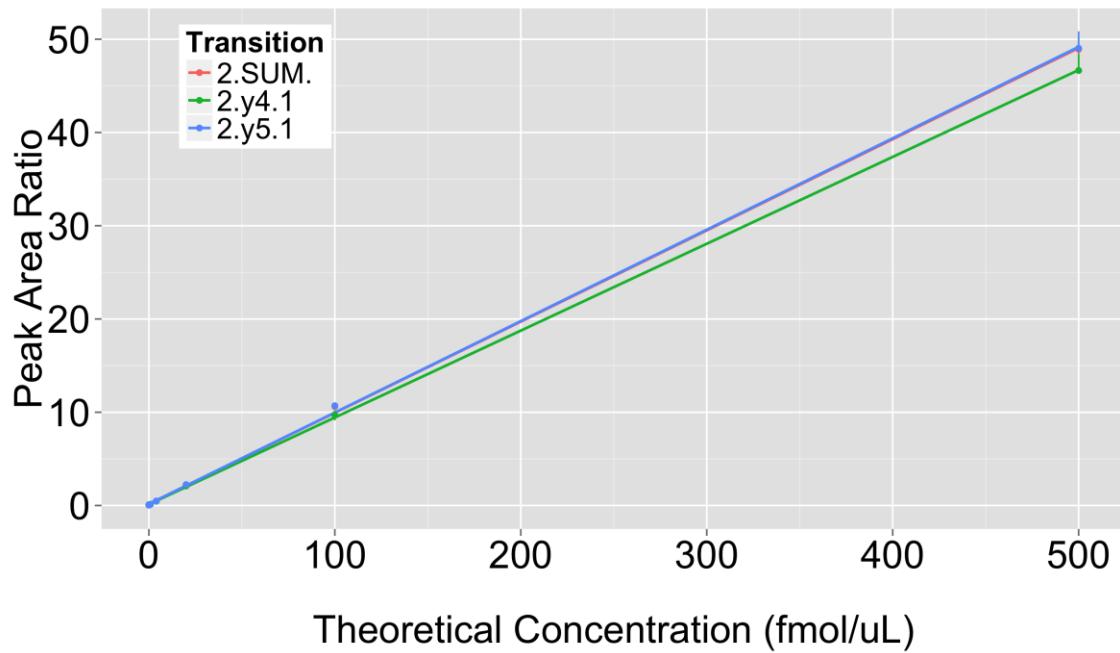
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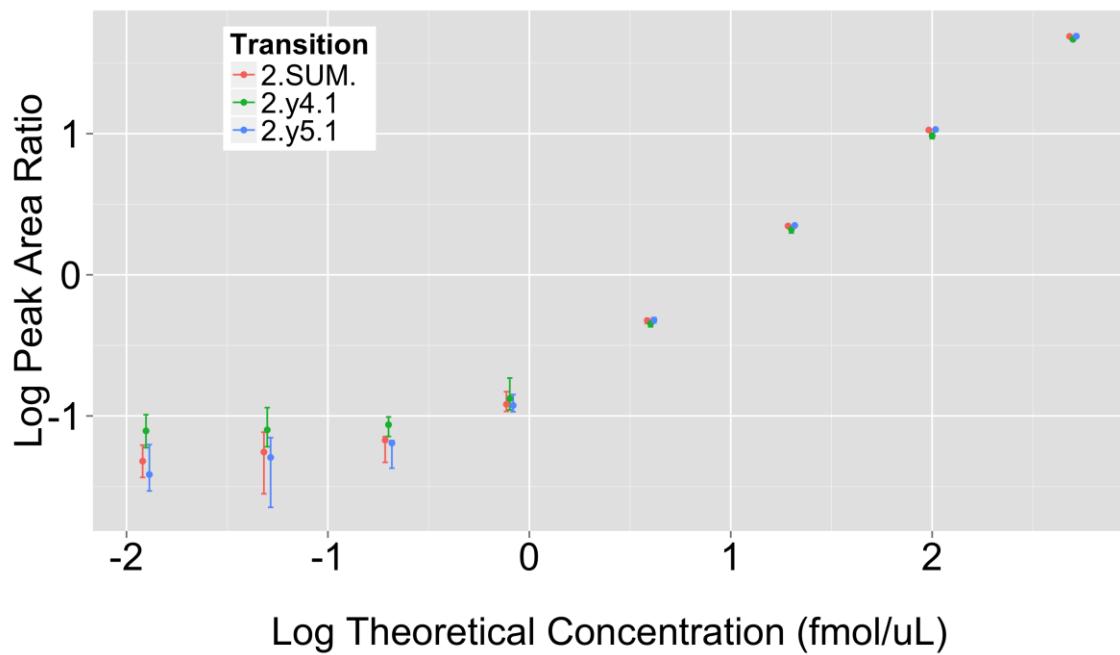
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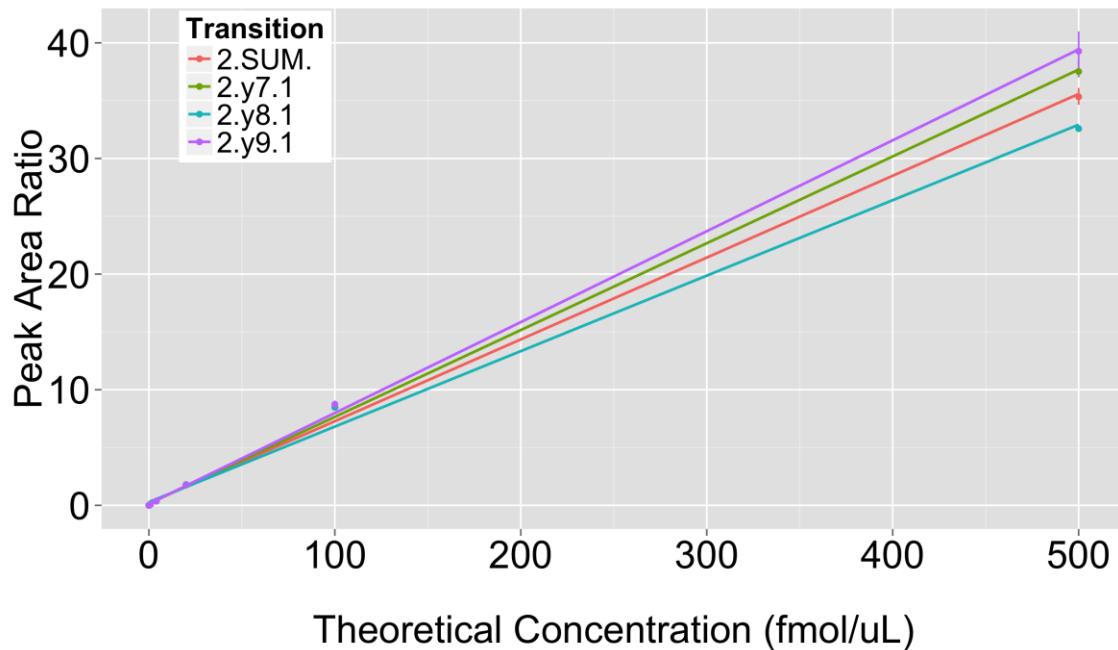
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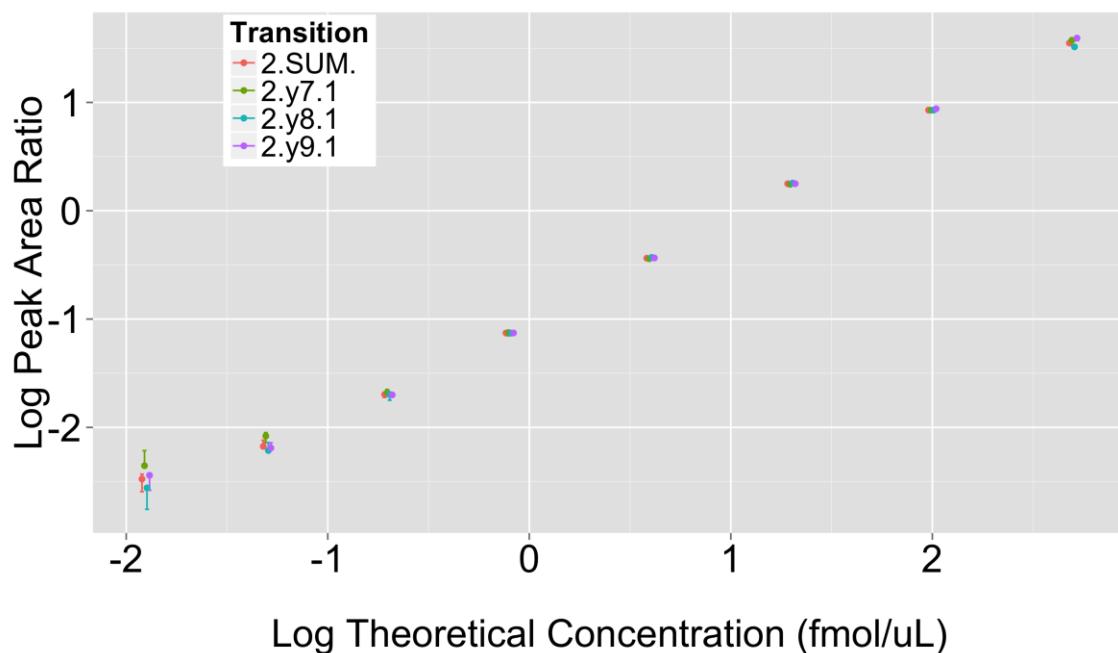
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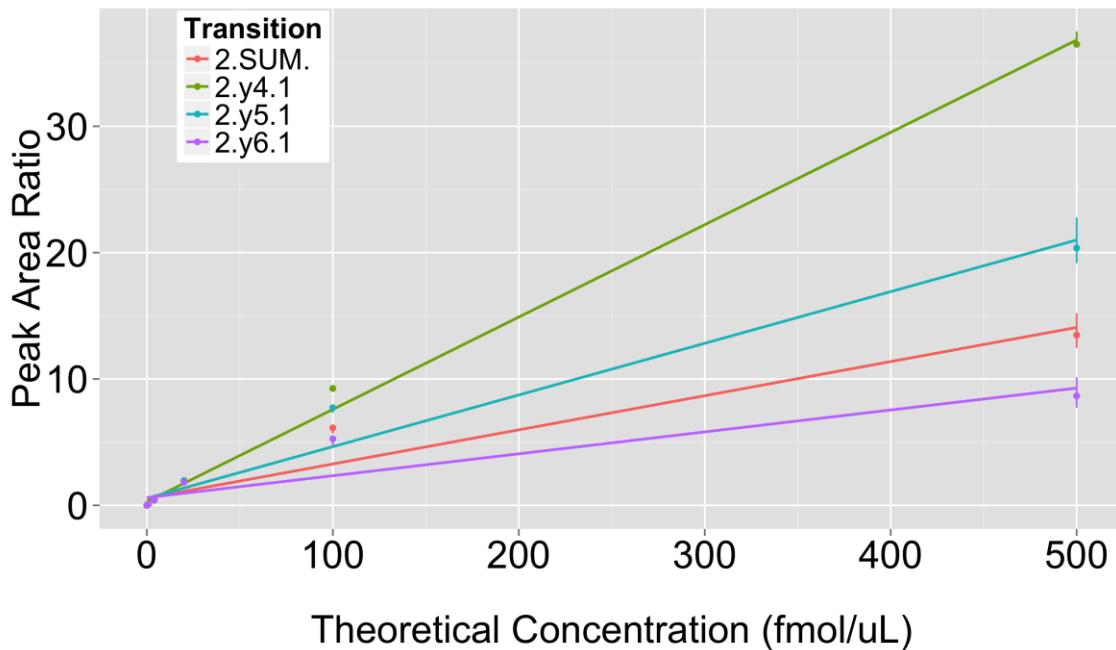
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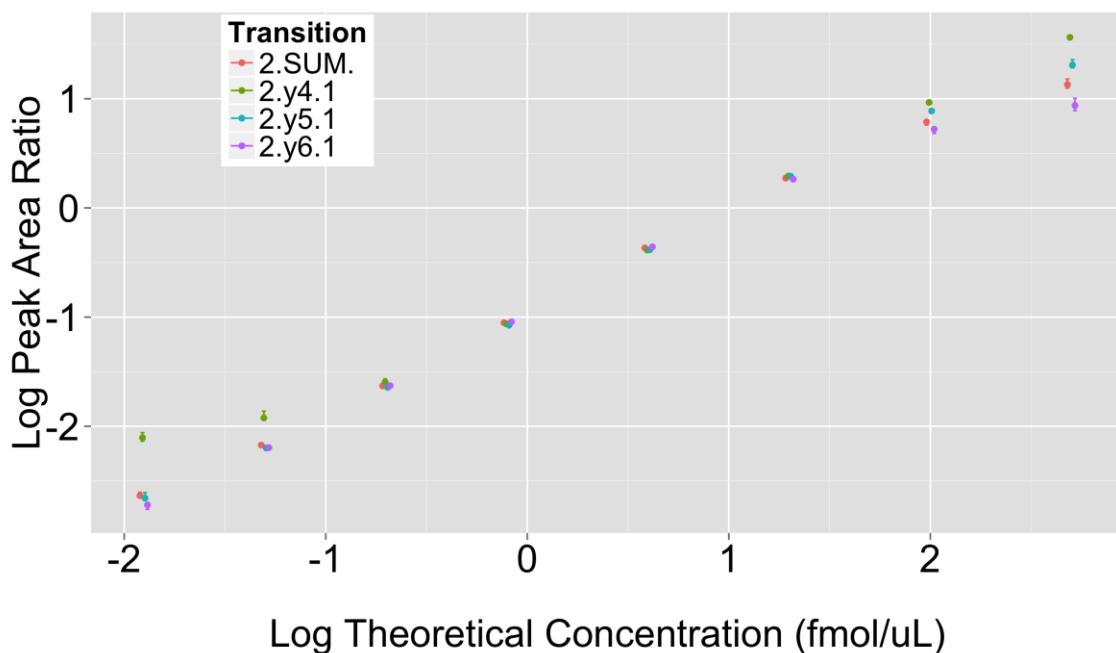
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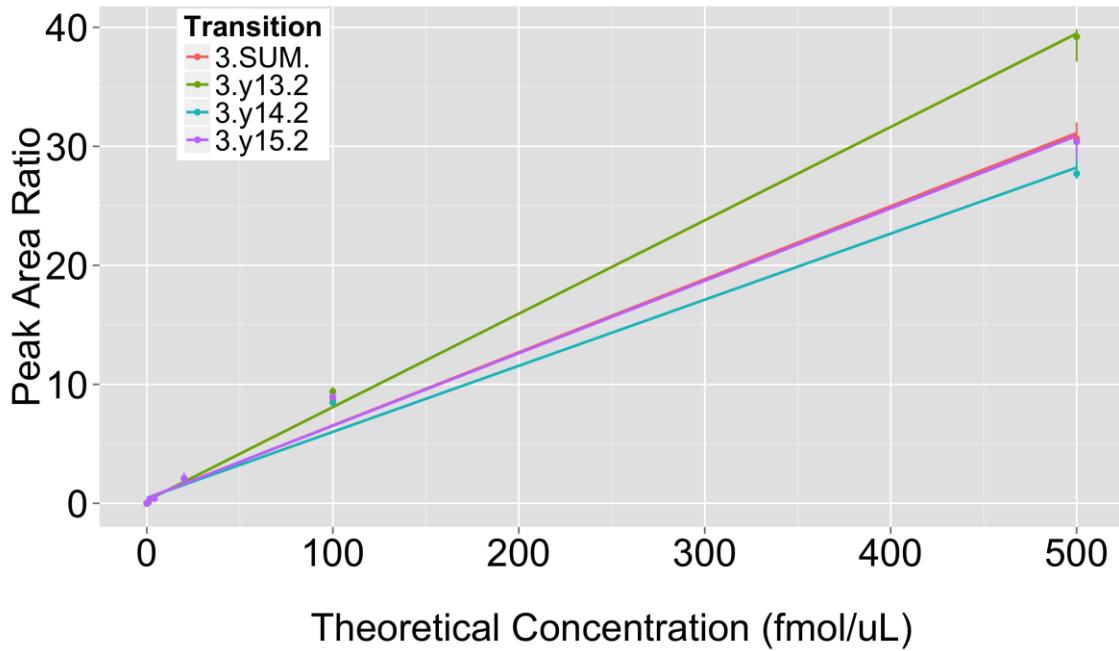
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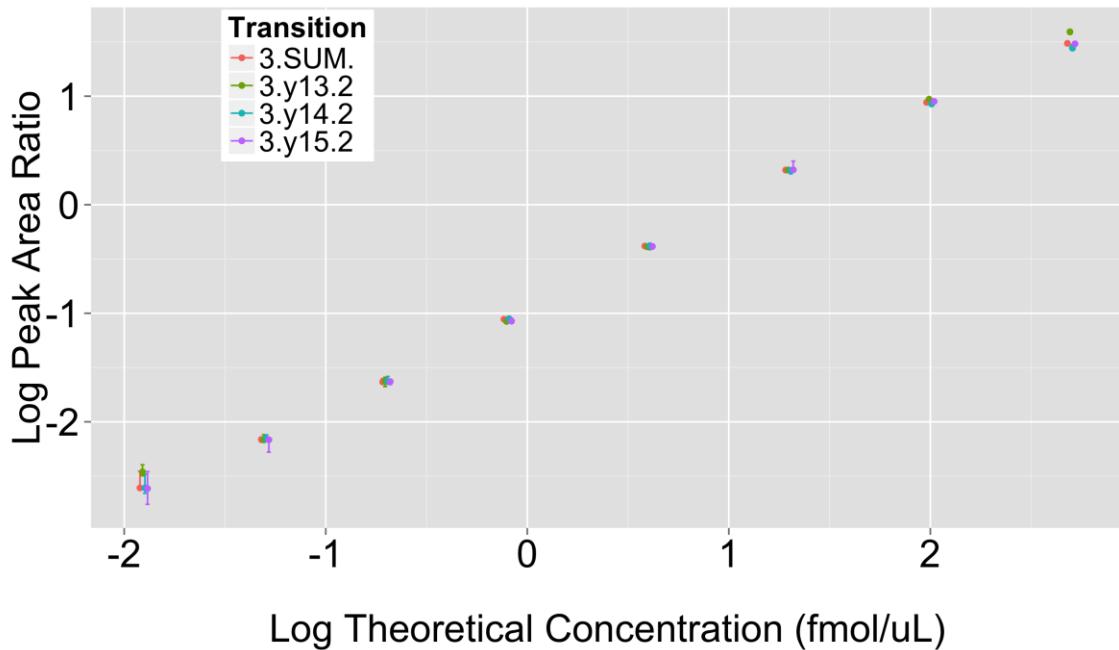
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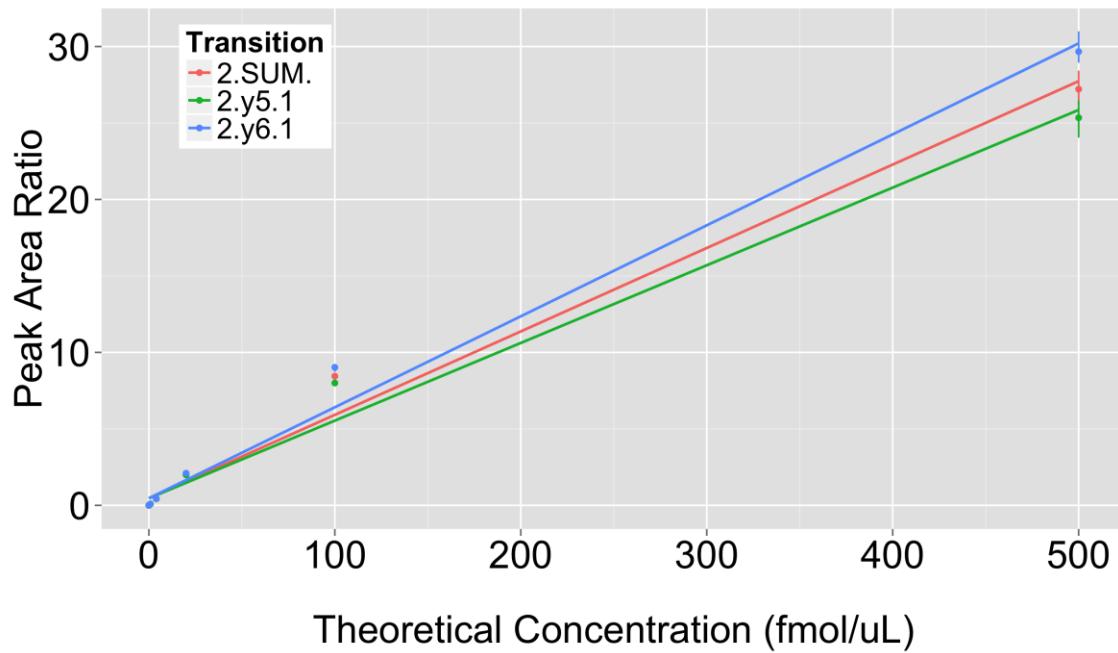
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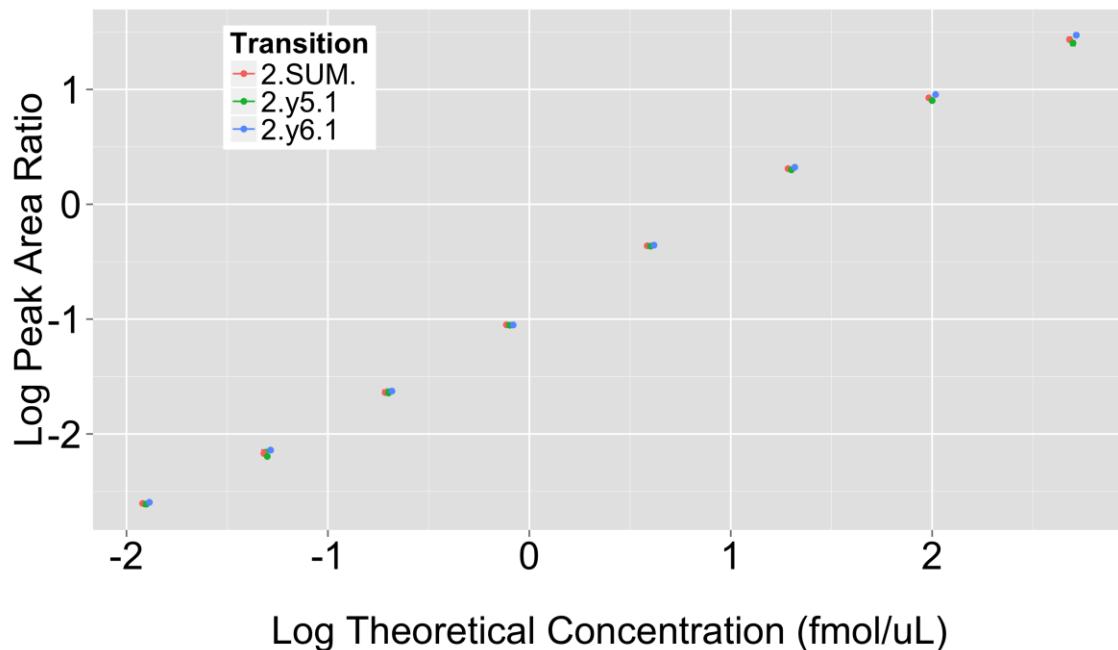
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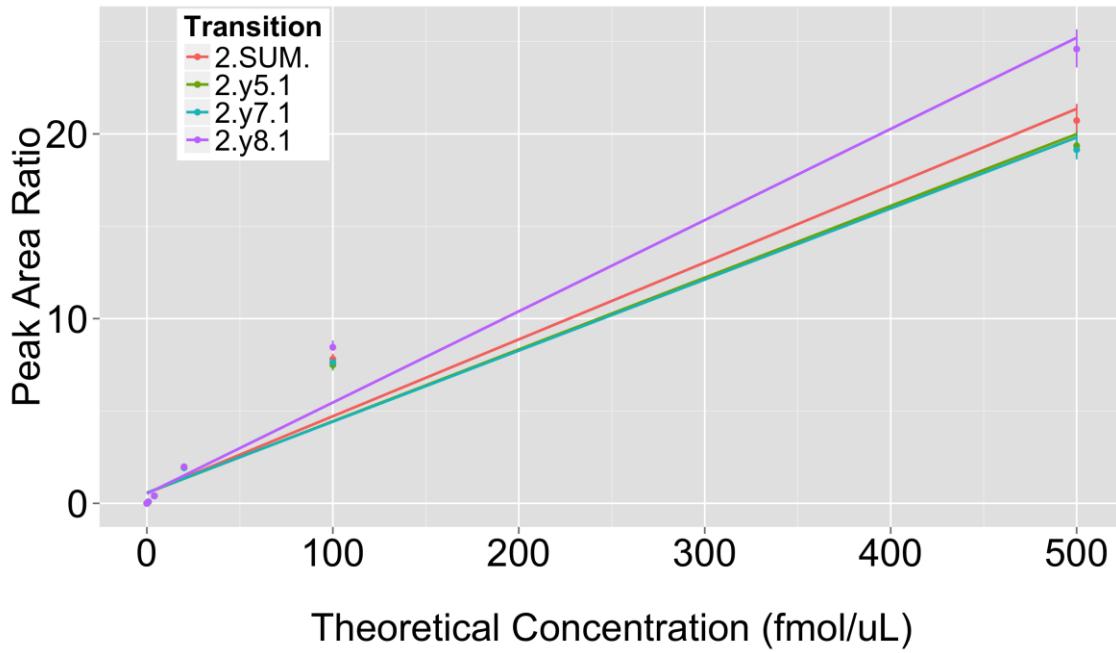
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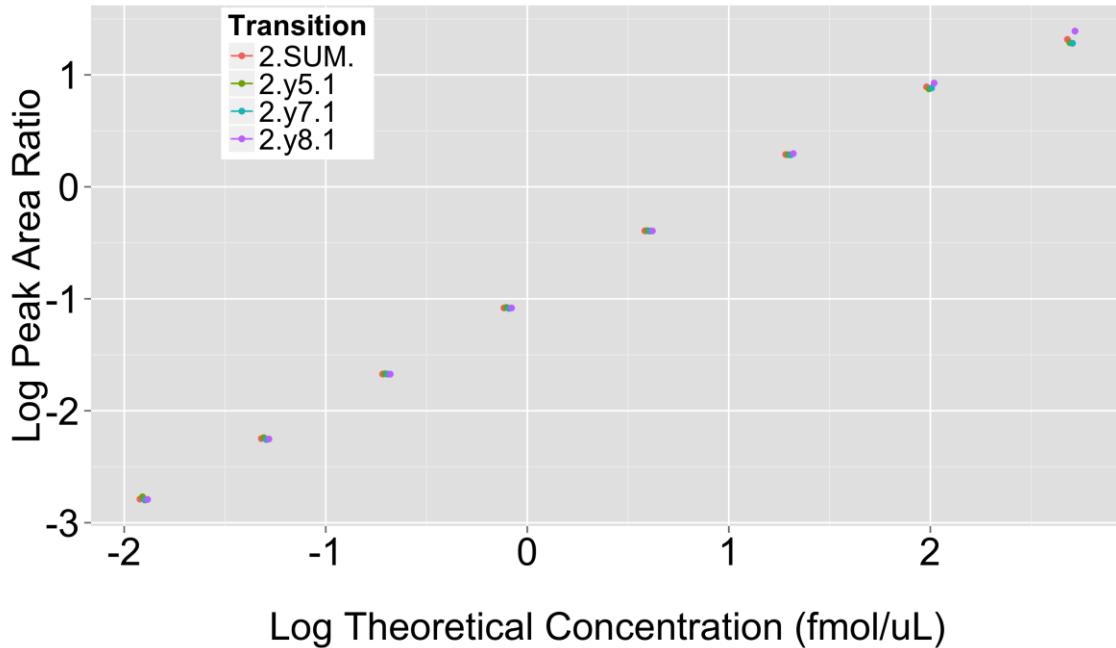
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Analyte: SAIC24B.ISTLSC[+57]ENK

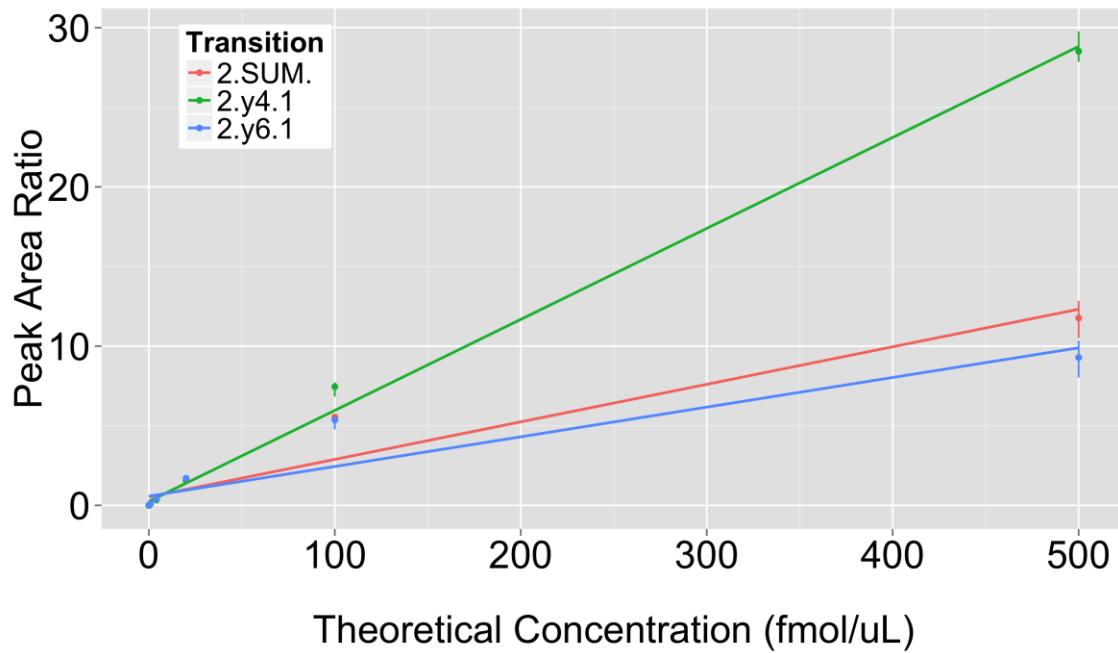


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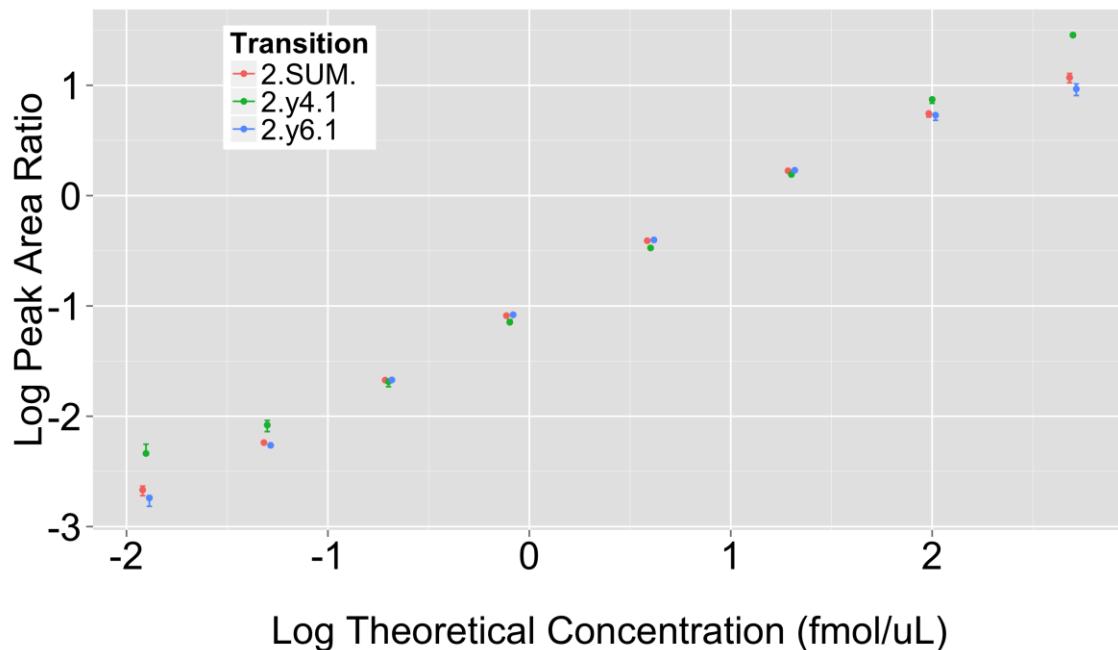


SAIC-25B, LGALS1, SFVNLGK.

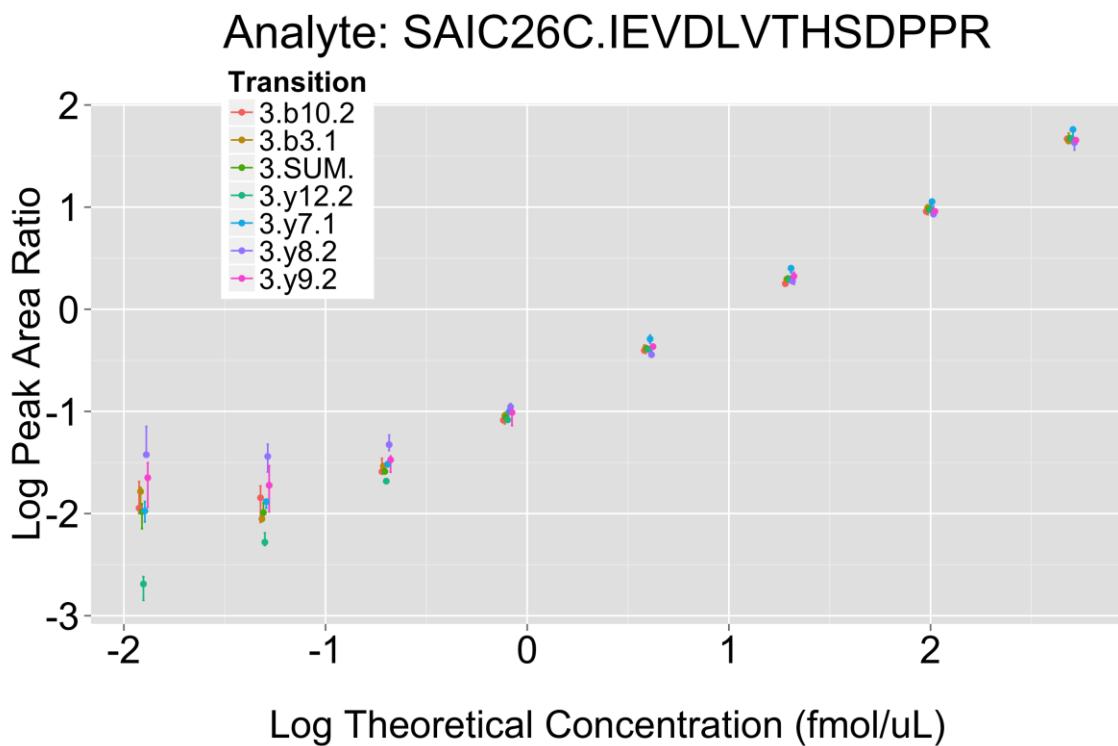
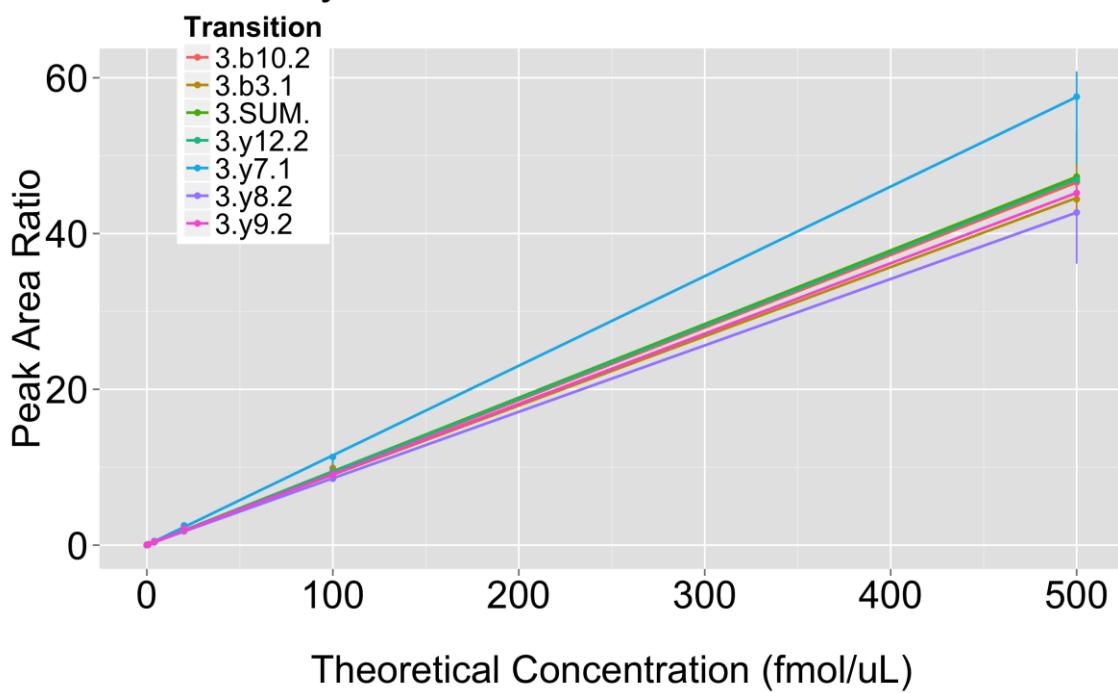
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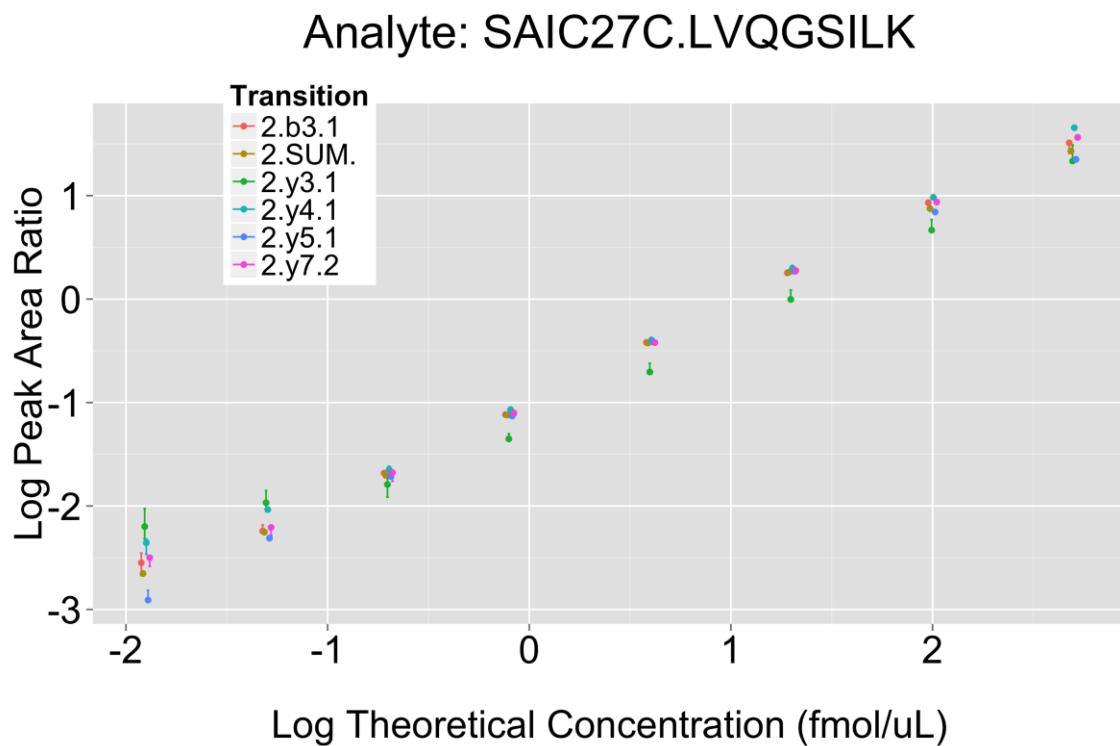
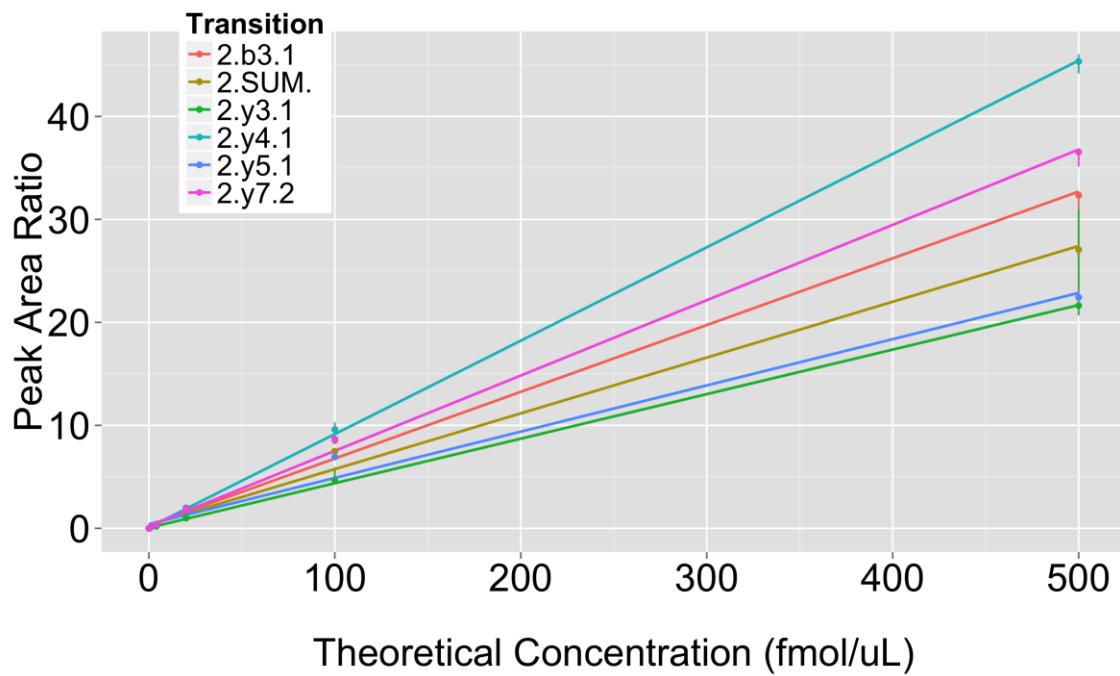
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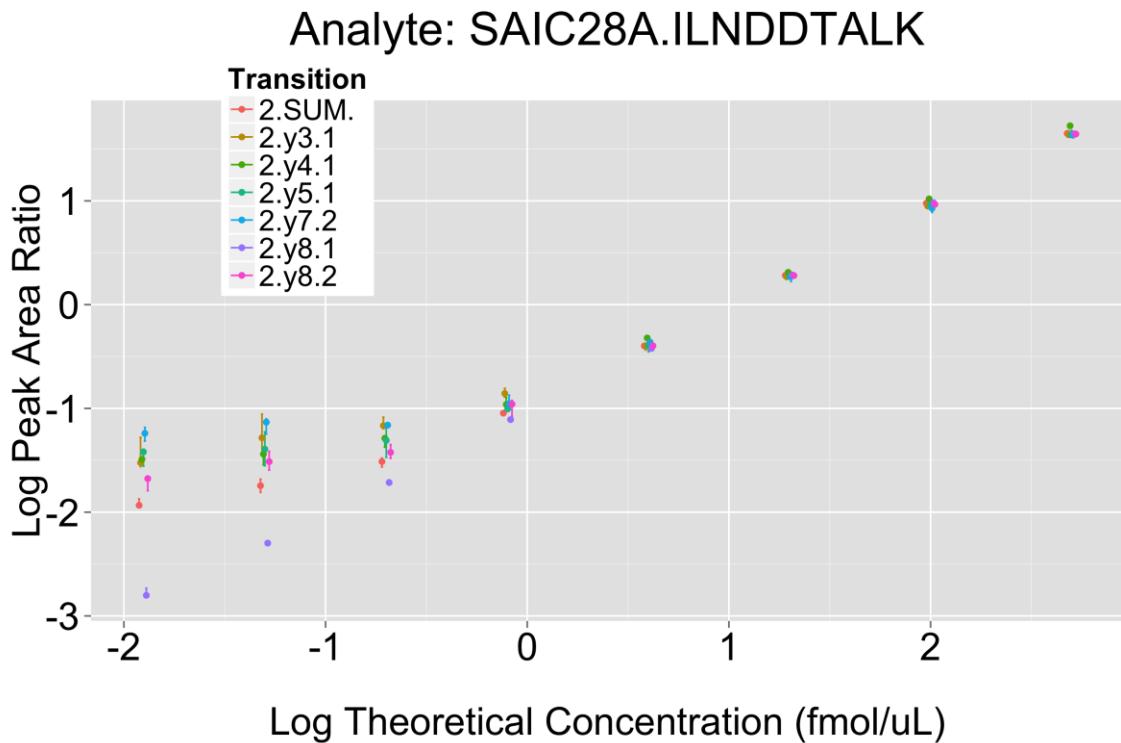
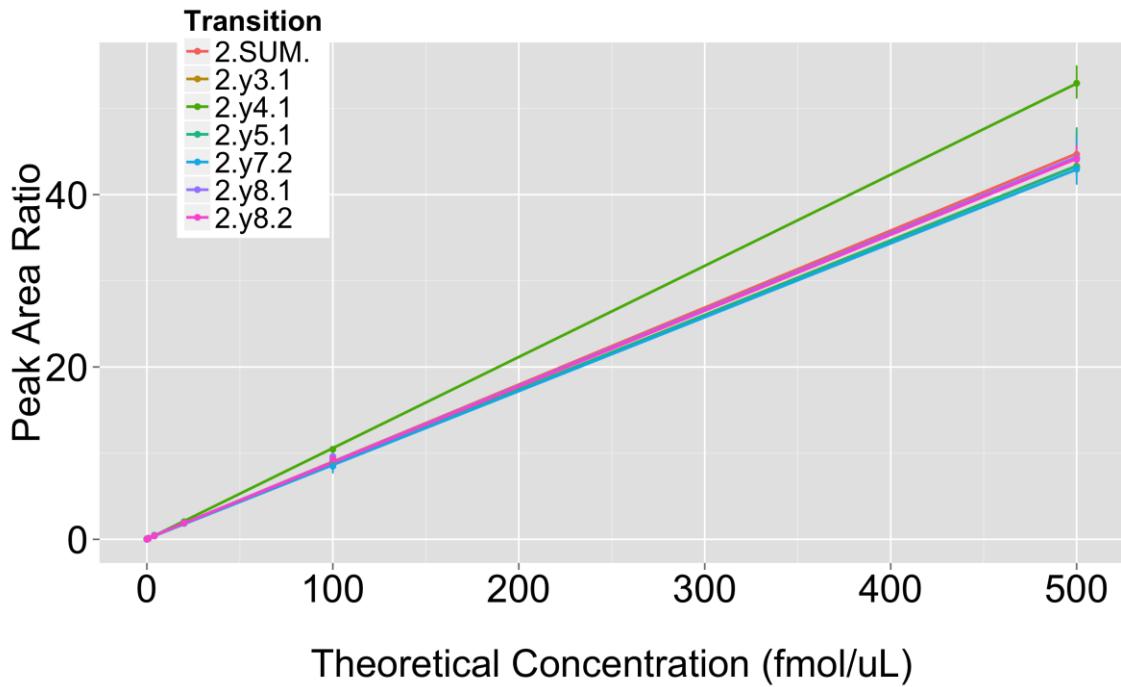
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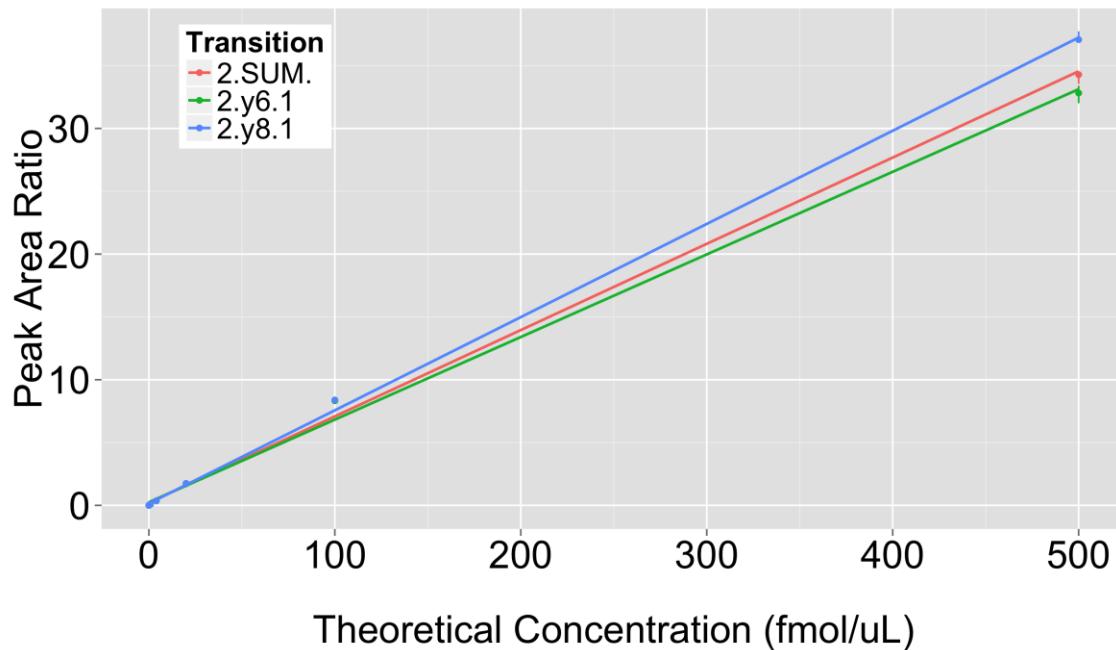
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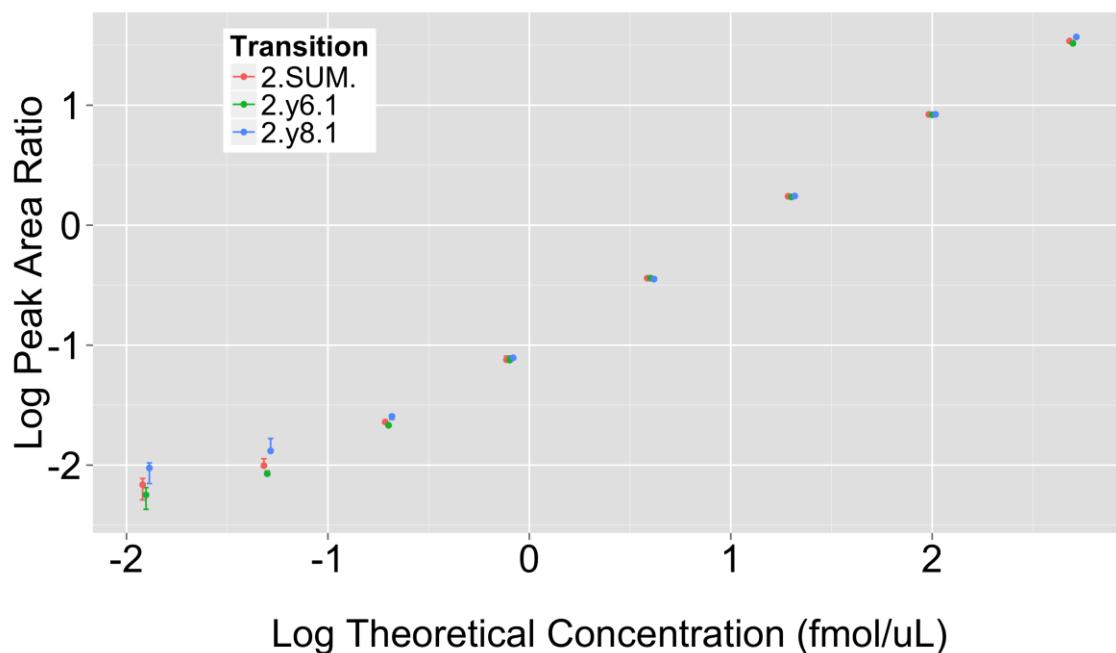
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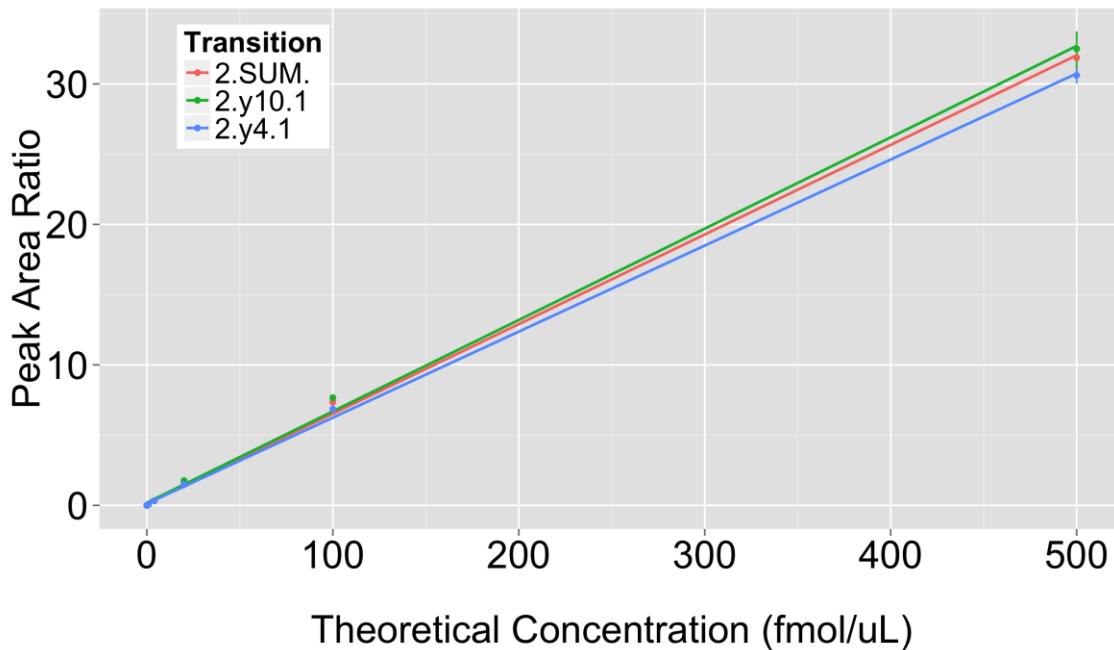
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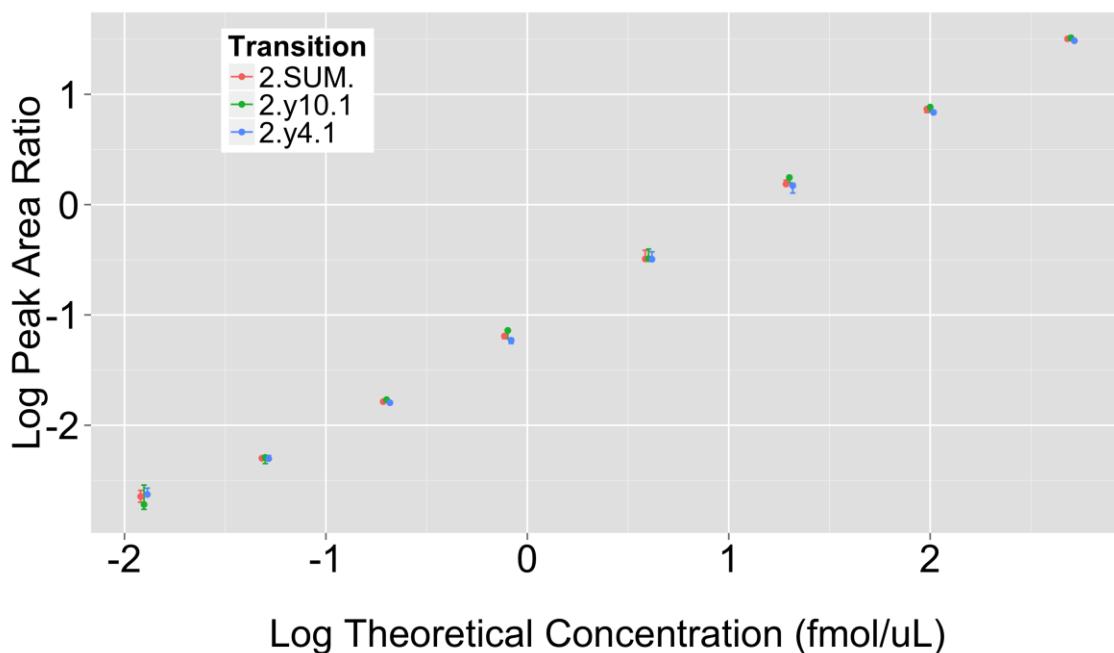
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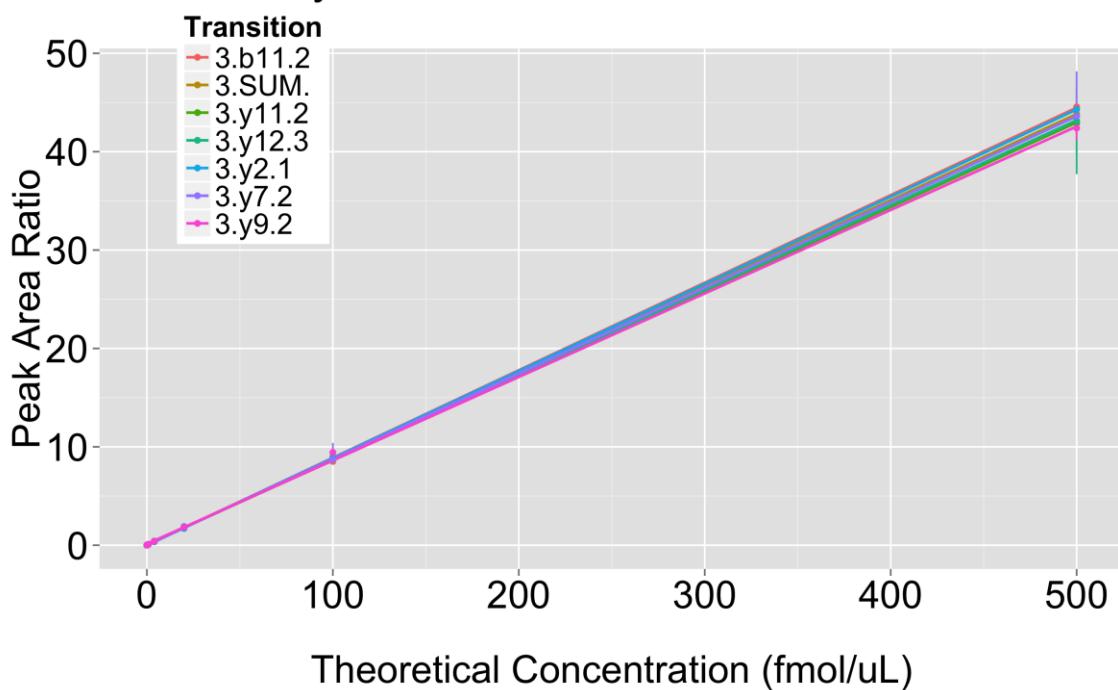
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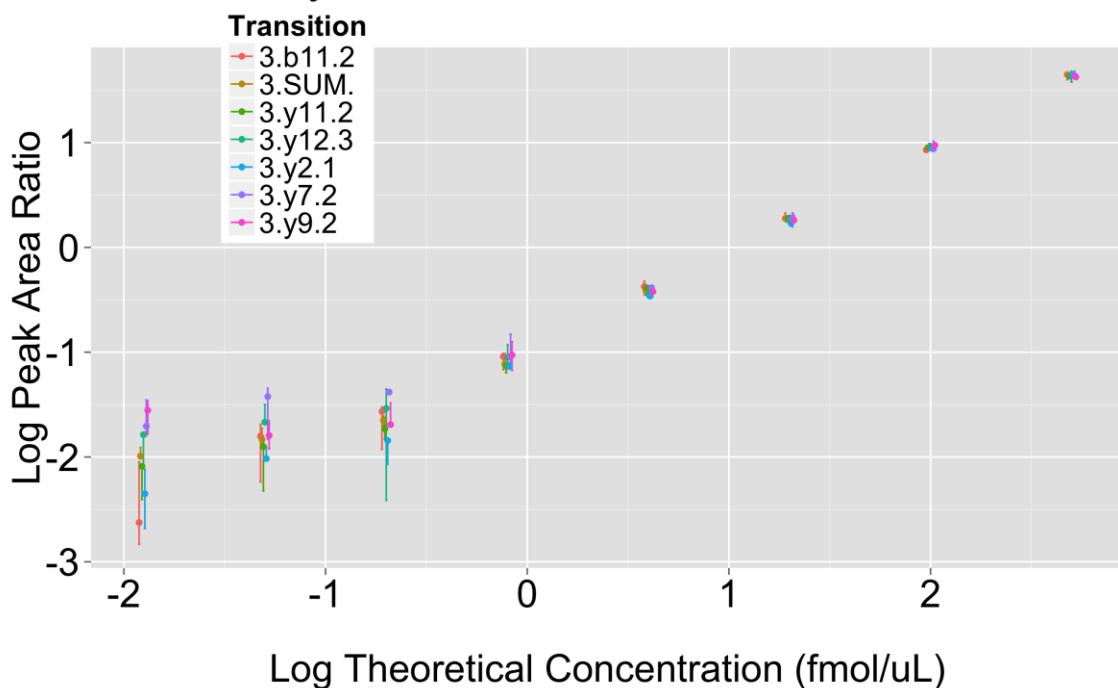
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Analyte: SAIC30E.DIQHWESLKPEER

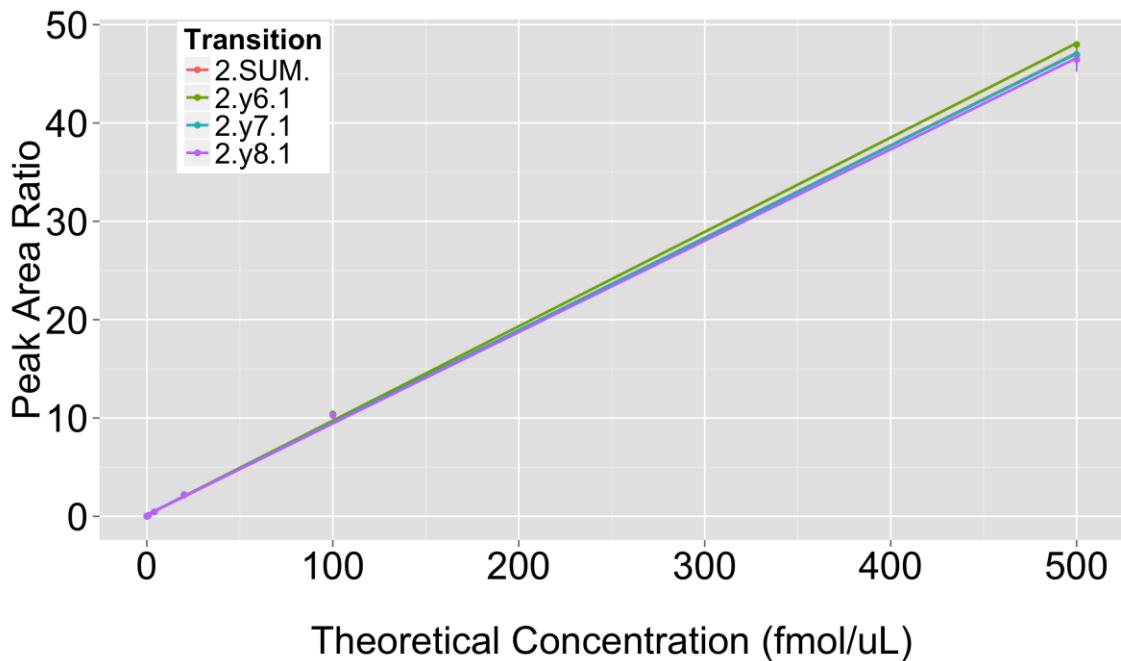


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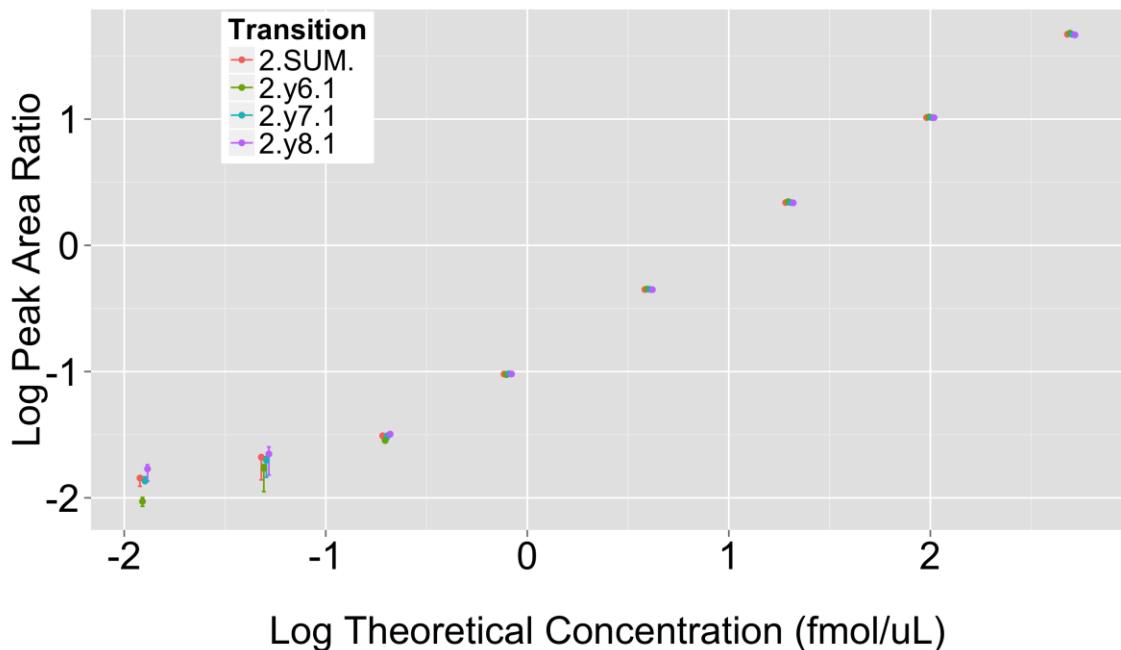


SAIC-31A, SNCG, TVEEAENIAVTSGVVR.

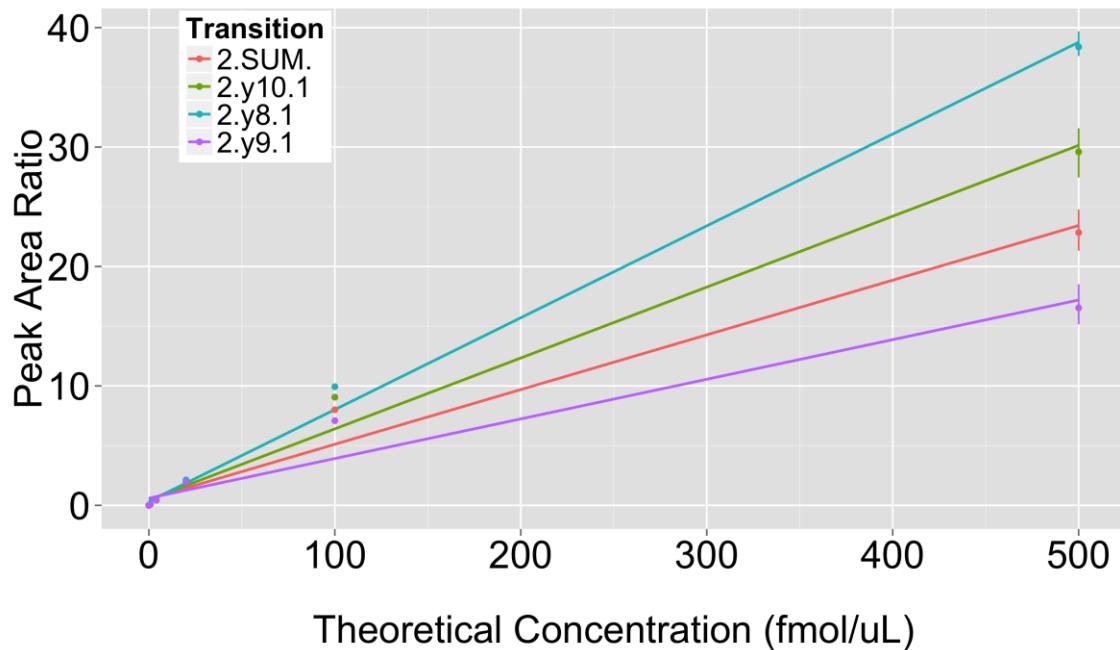
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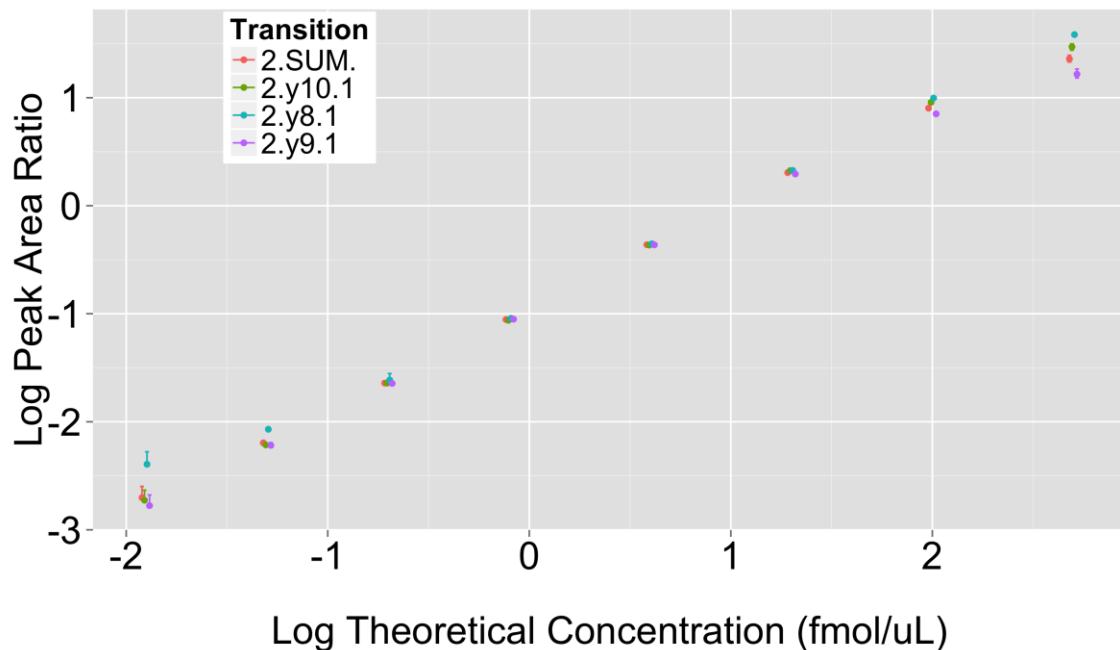
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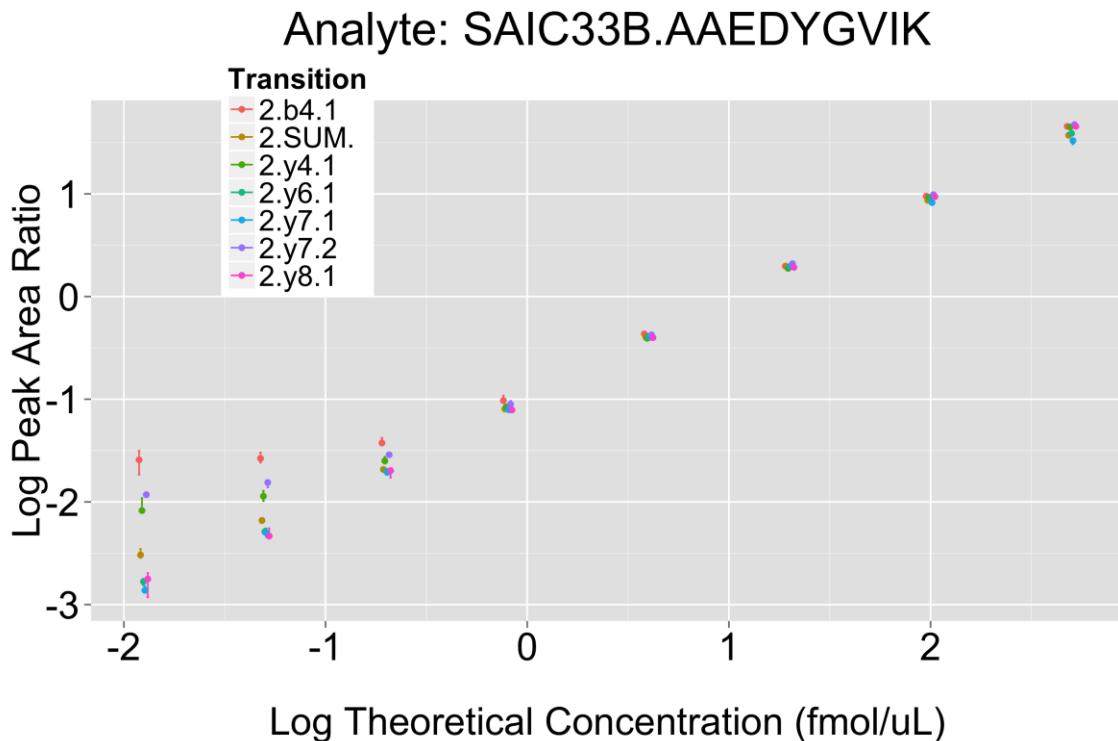
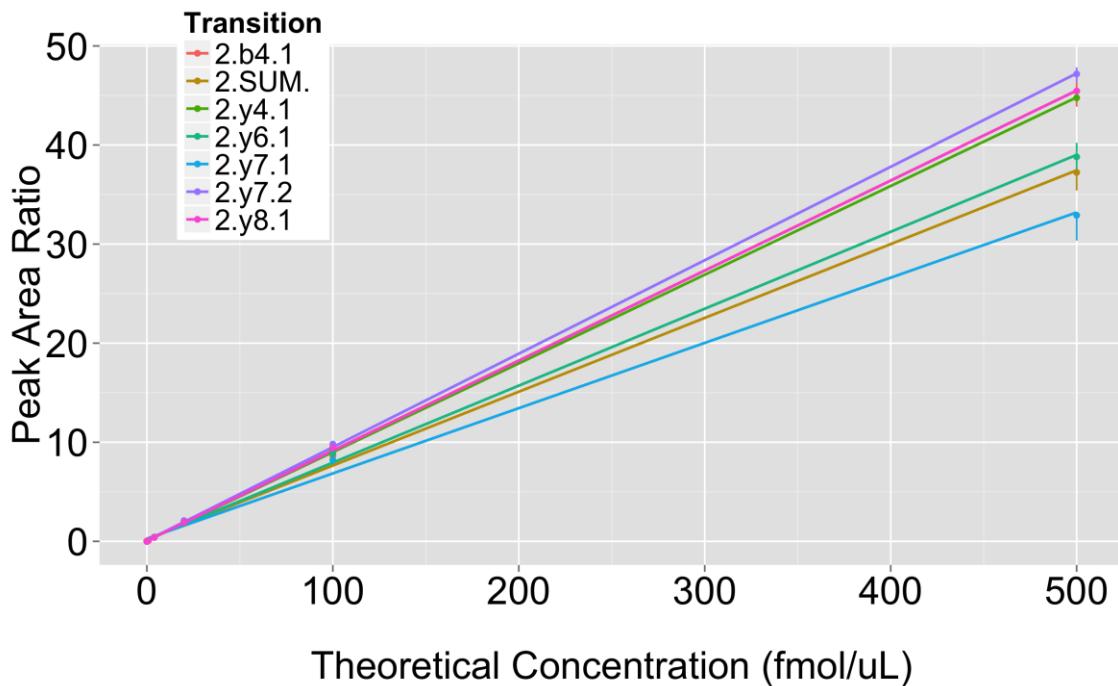
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Analyte: SAIC32C.YLAPSGPSGTLK

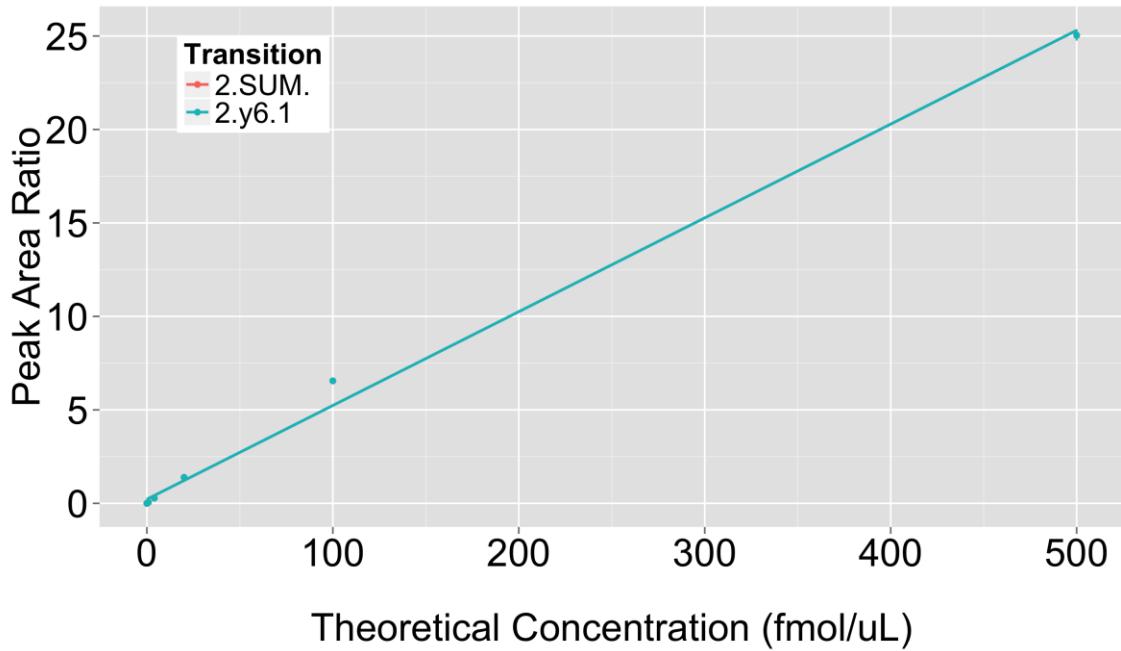


Analyte: SAIC33B.AAEDYGVIK

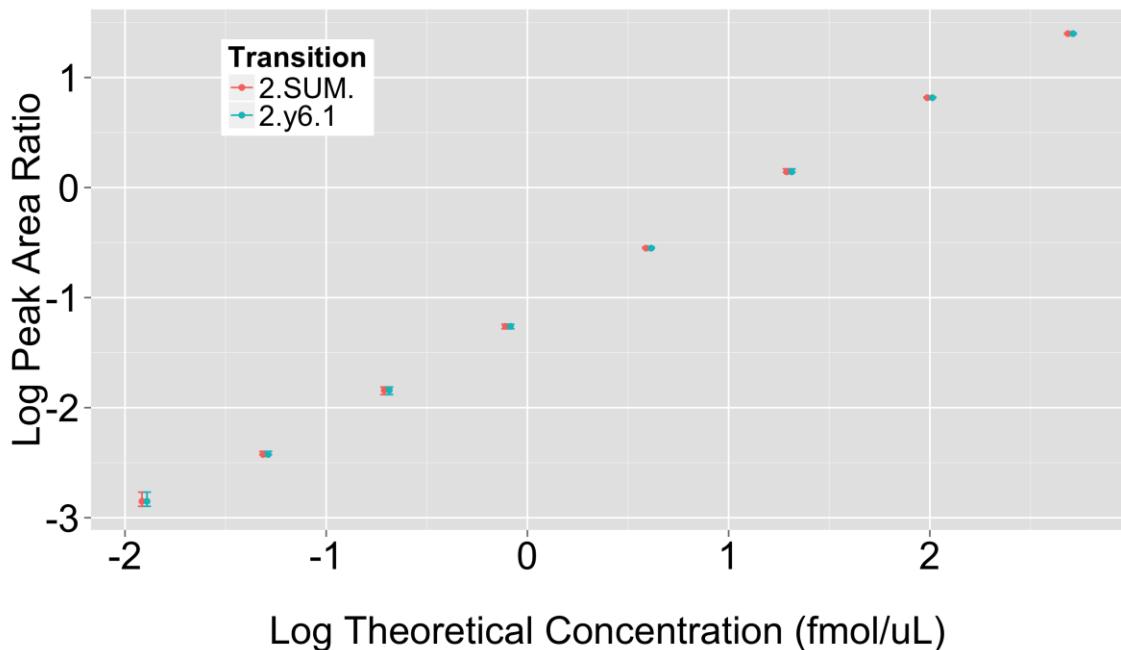


SAIC-34B, PRDX2, LSEDYGVLK.

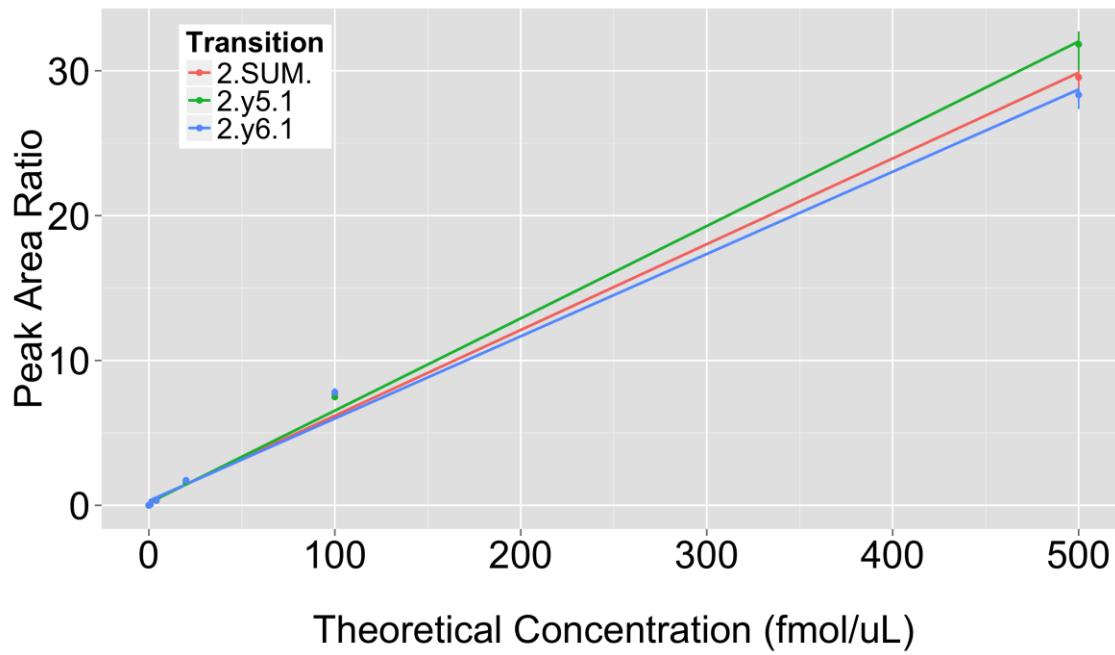
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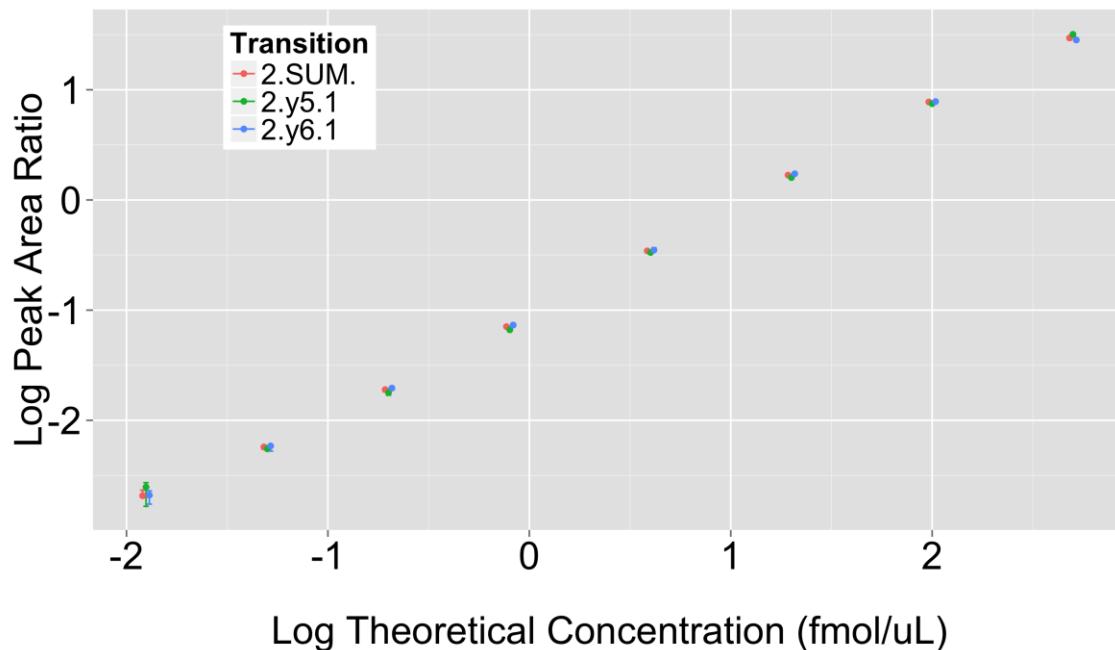
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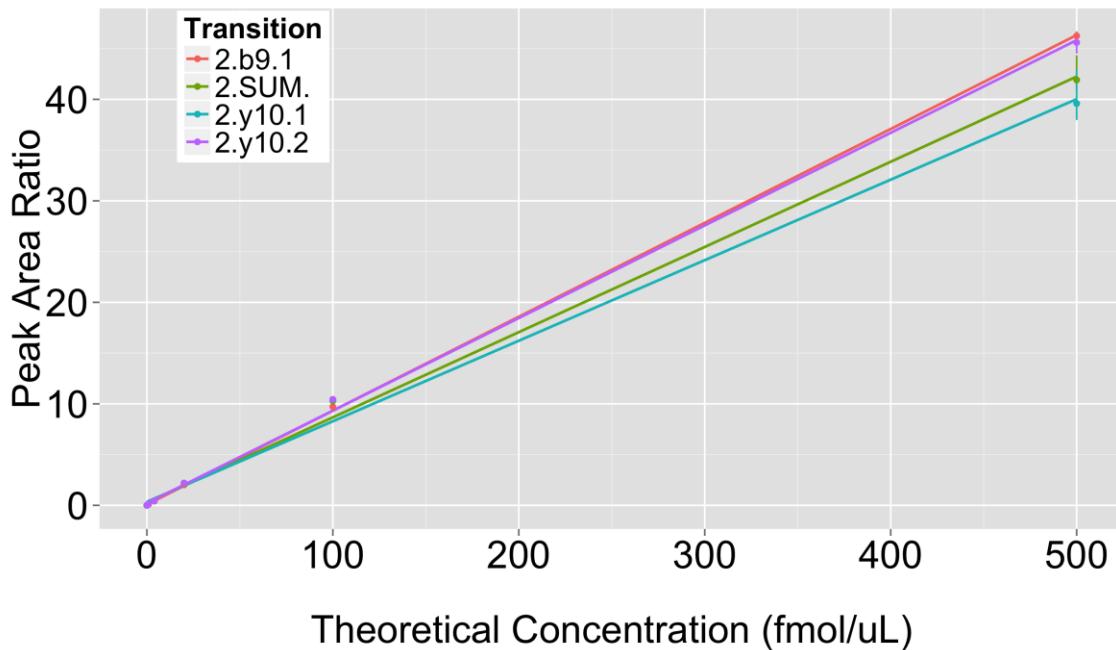
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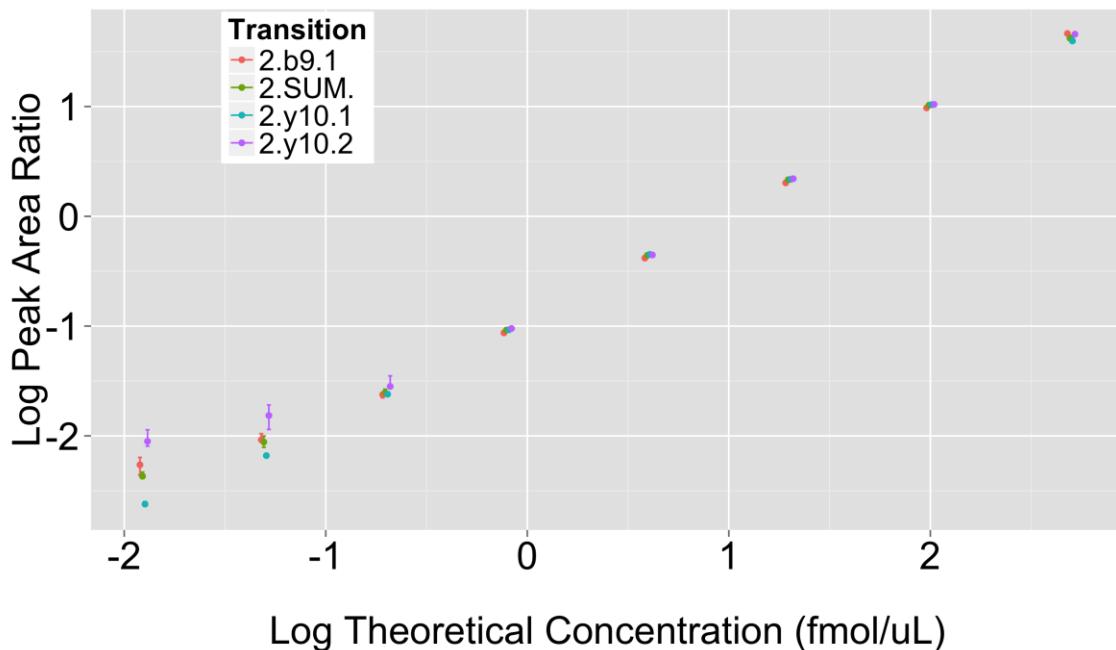
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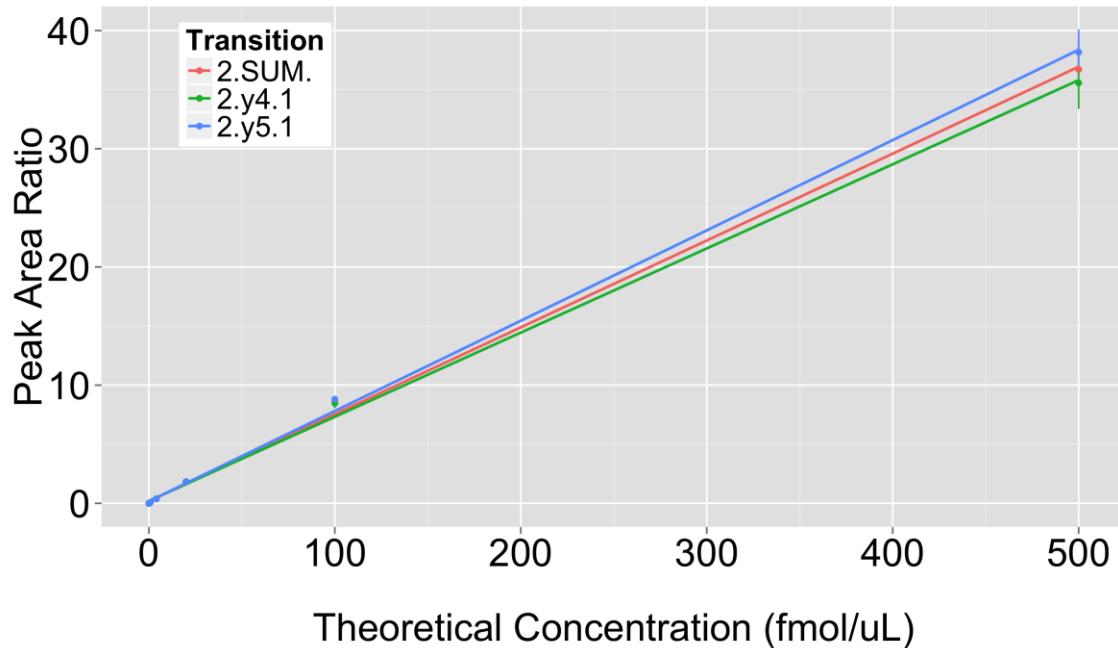
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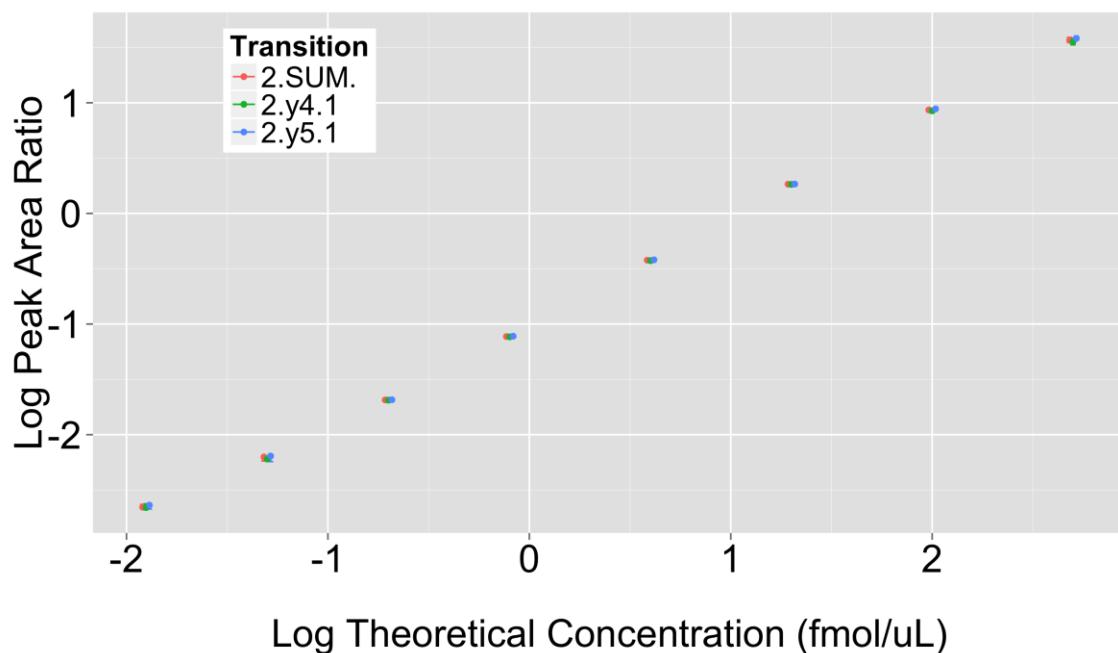
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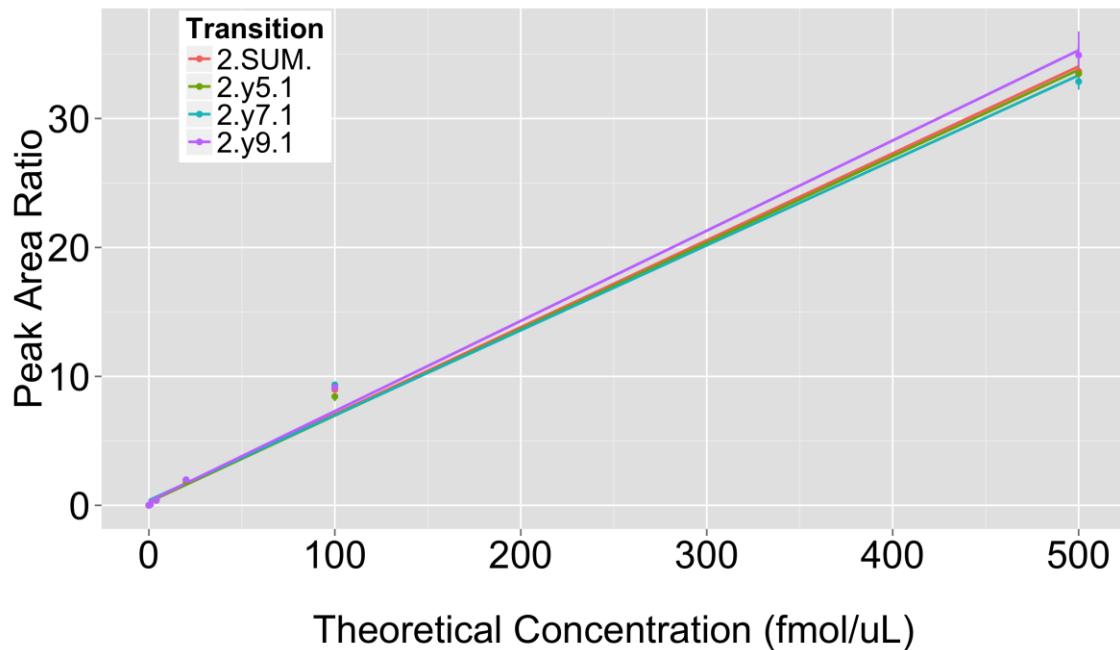
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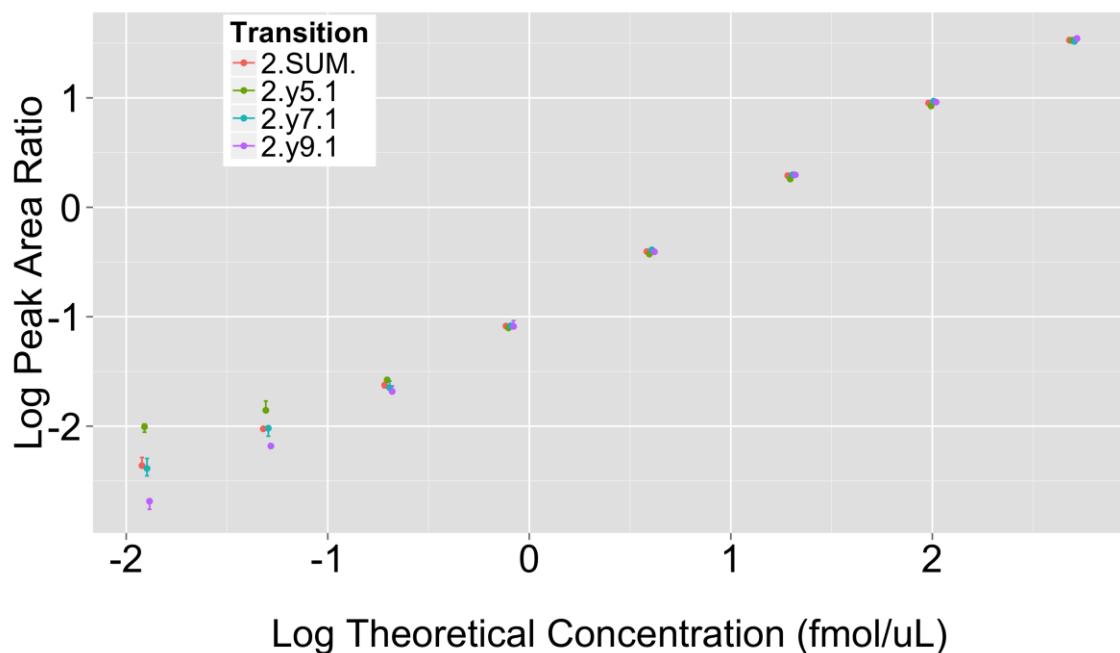
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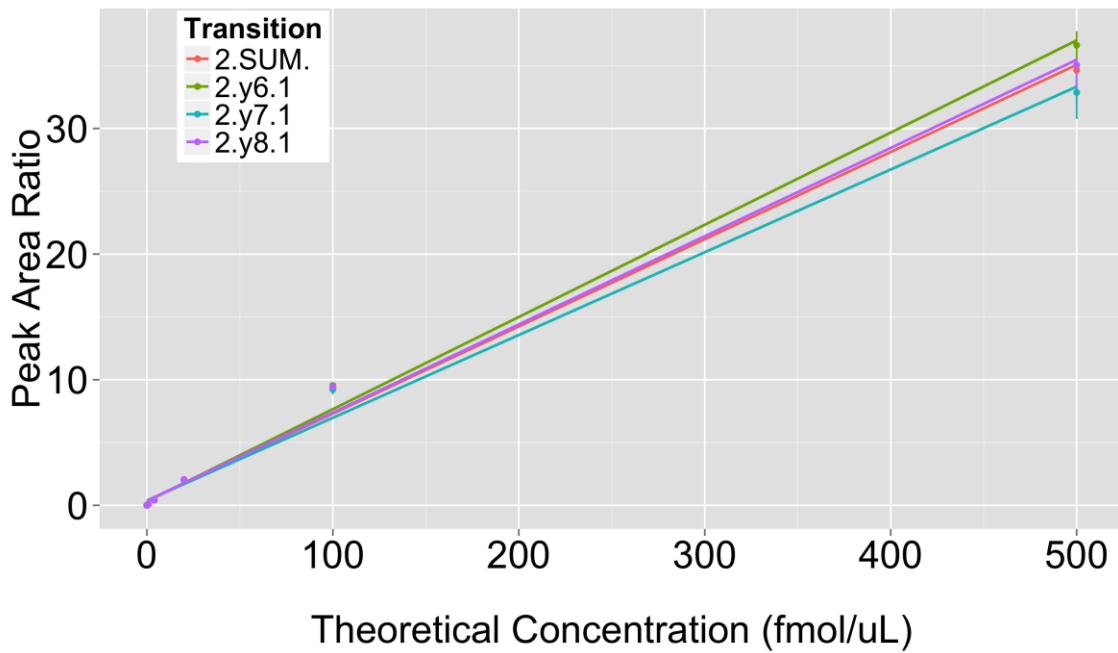
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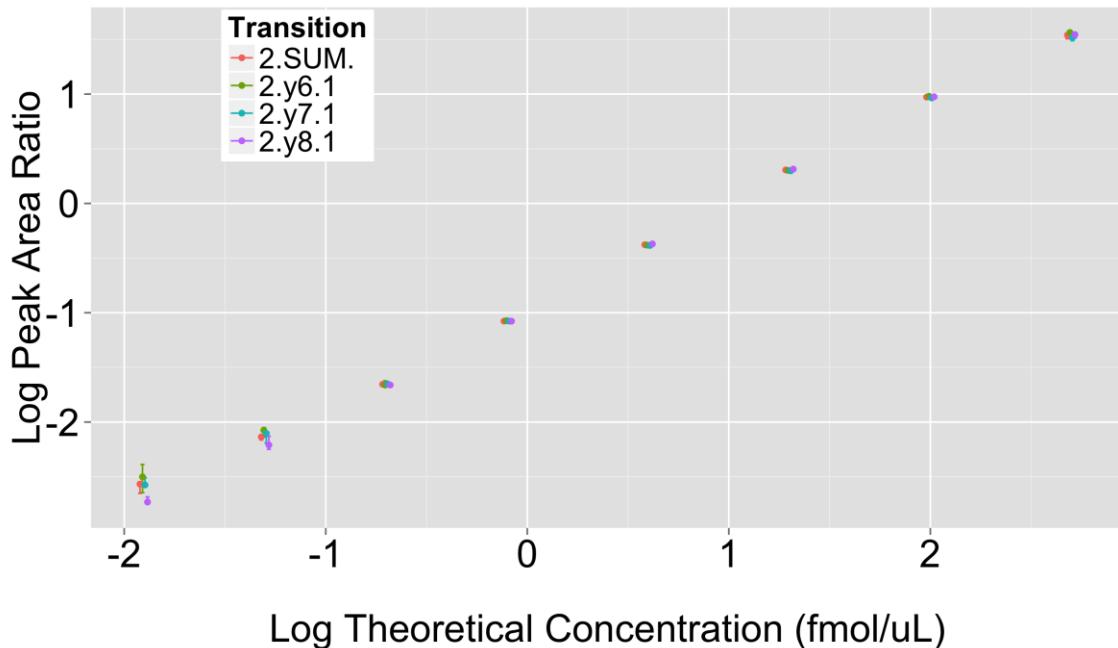
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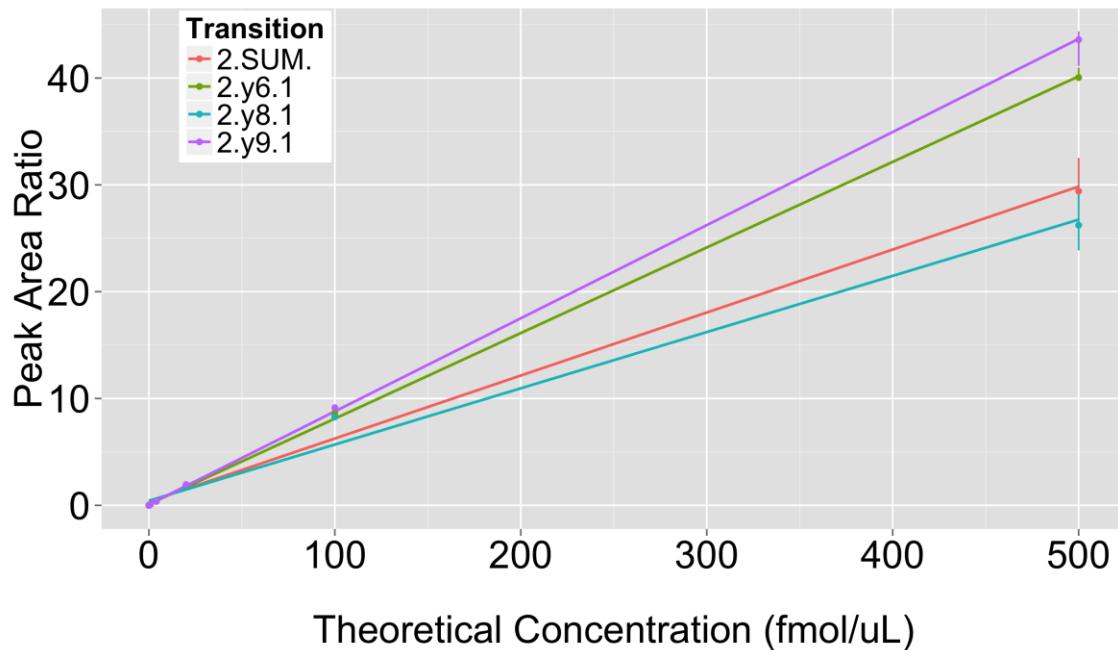
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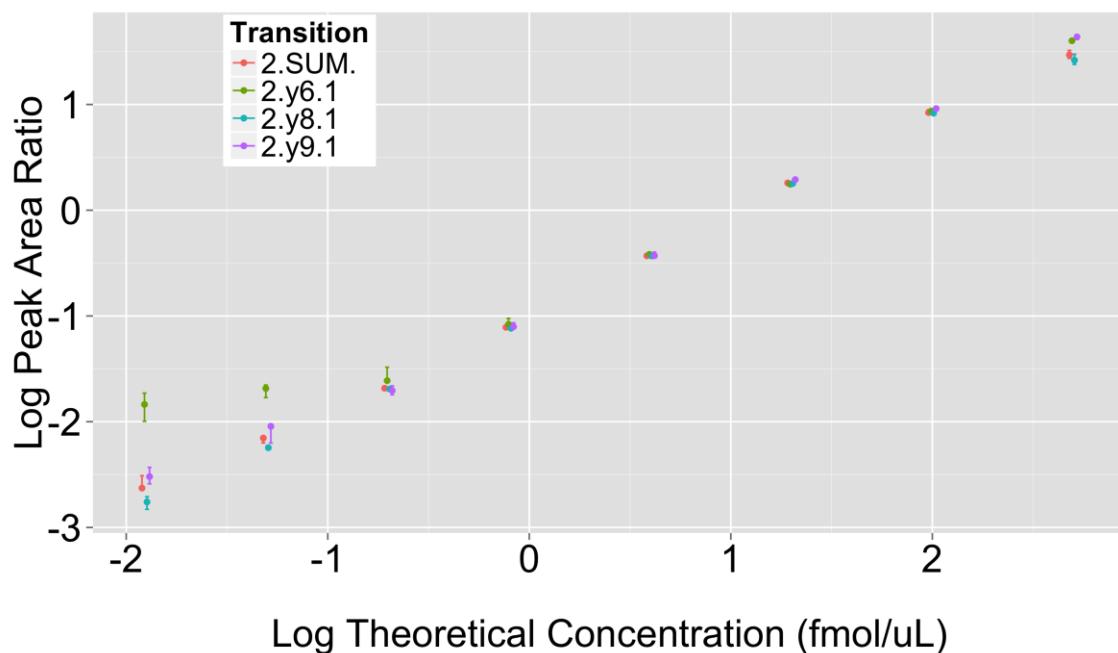
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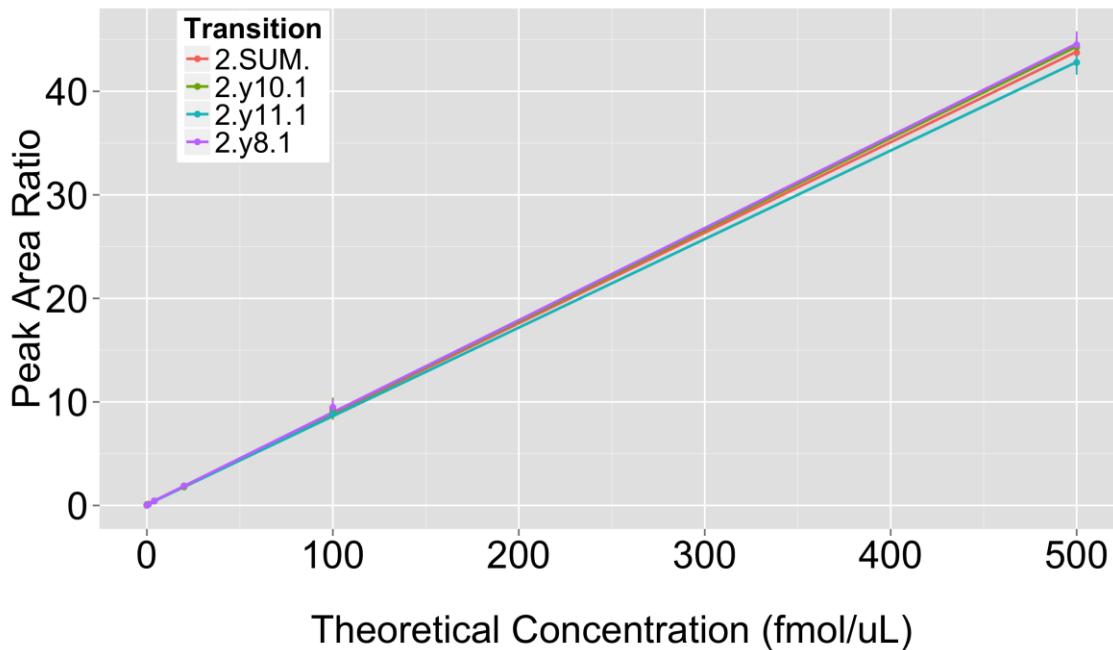
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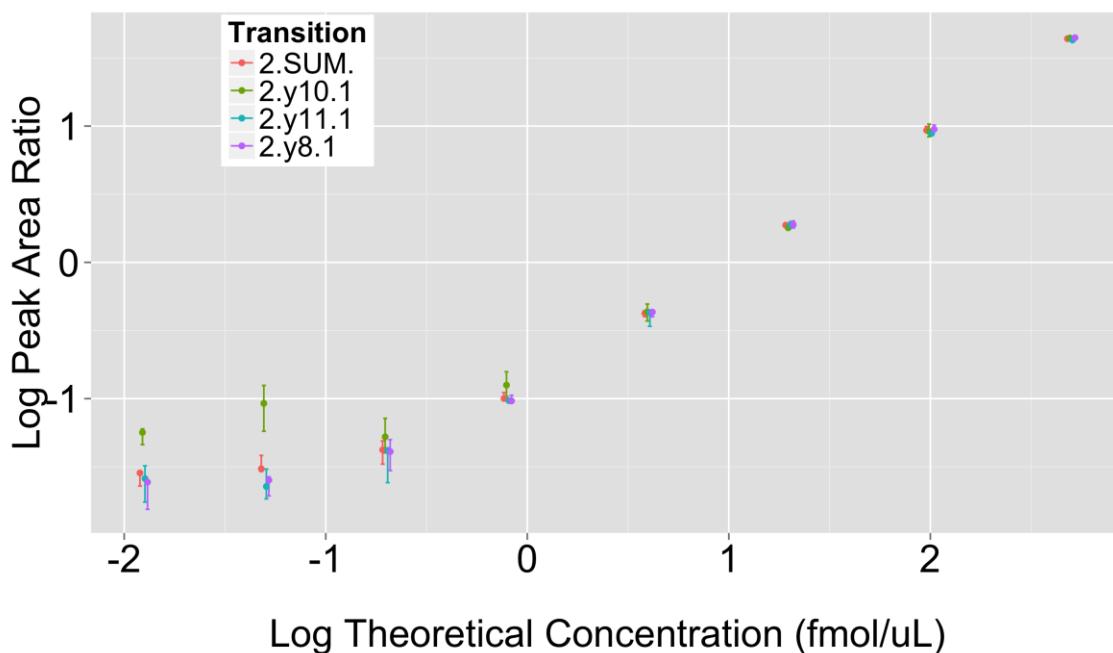
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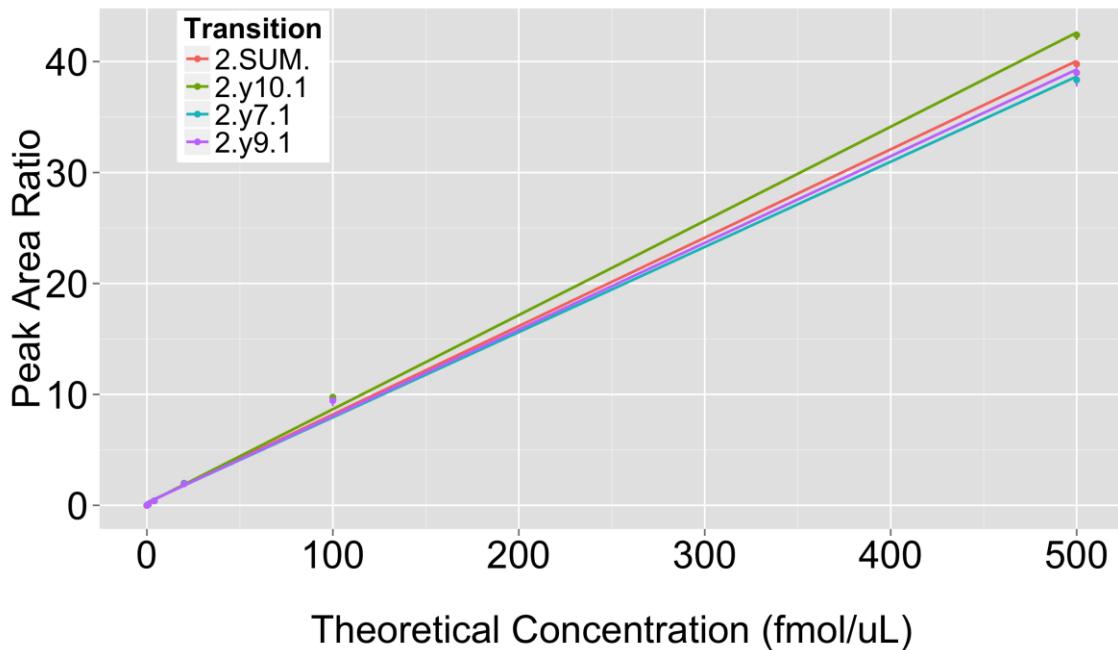
Analyte: SAIC41C.LSLEFPGYPPYNAPTVK



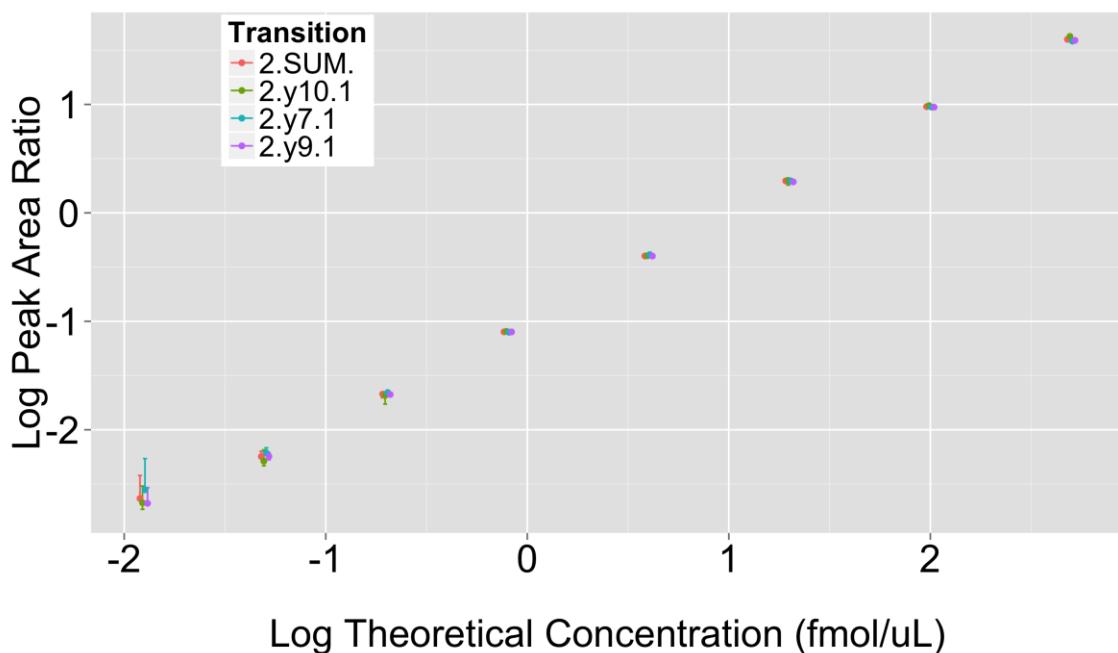
Analyte: SAIC41C.LSLEFPGYPPYNAPTVK



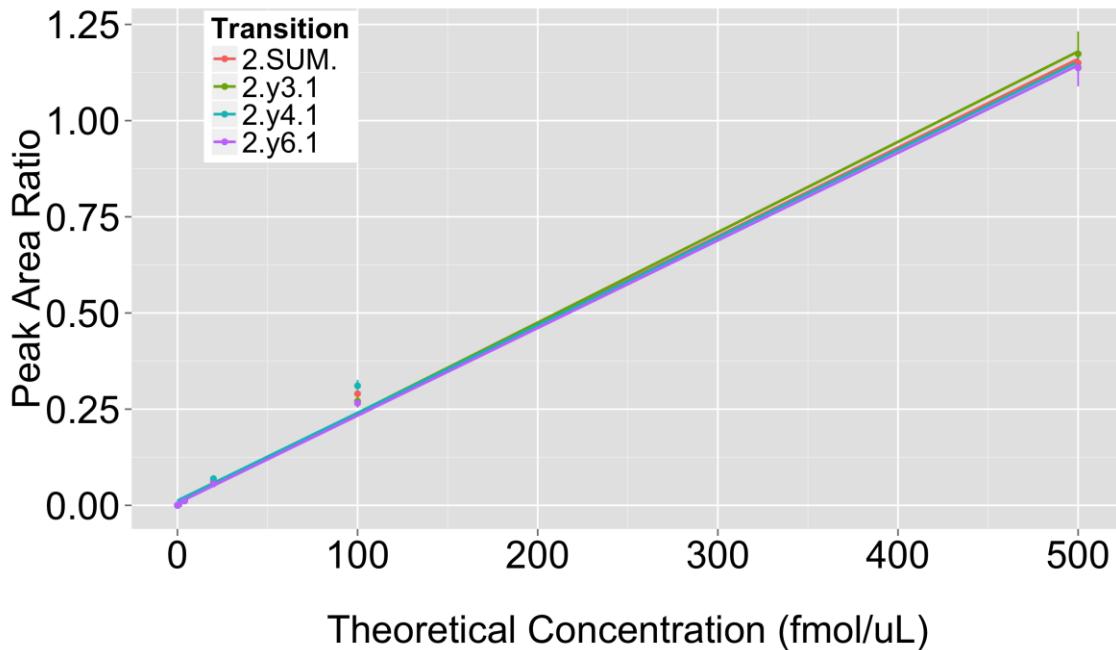
Analyte: SAIC42B.AQLLQGLGFNLTER



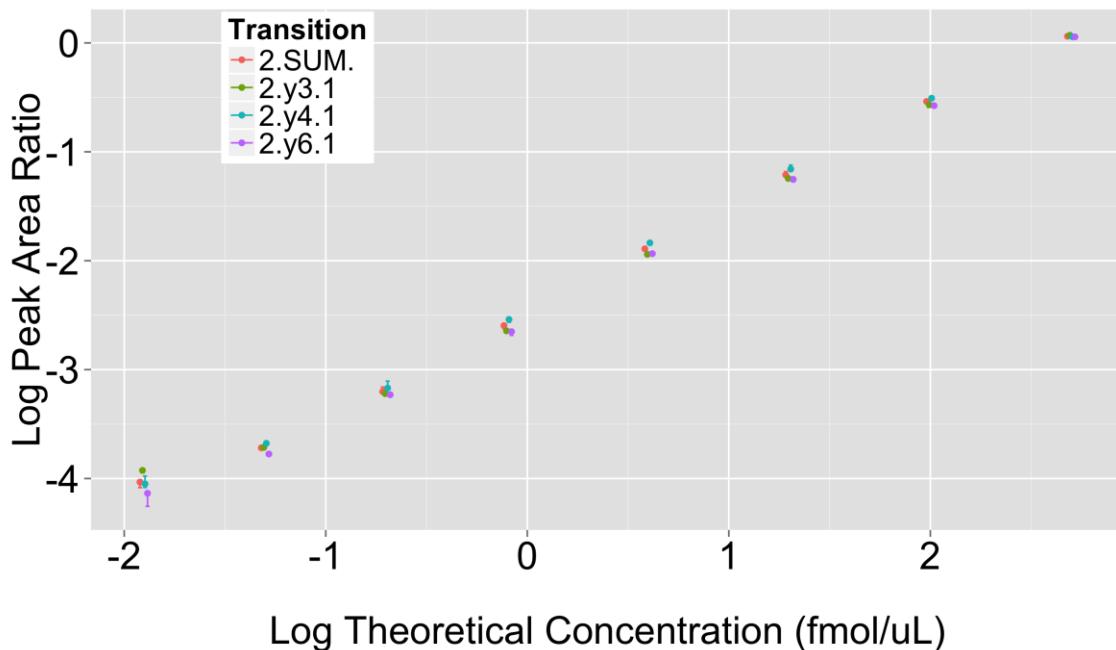
Analyte: SAIC42B.AQLLQGLGFNLTER



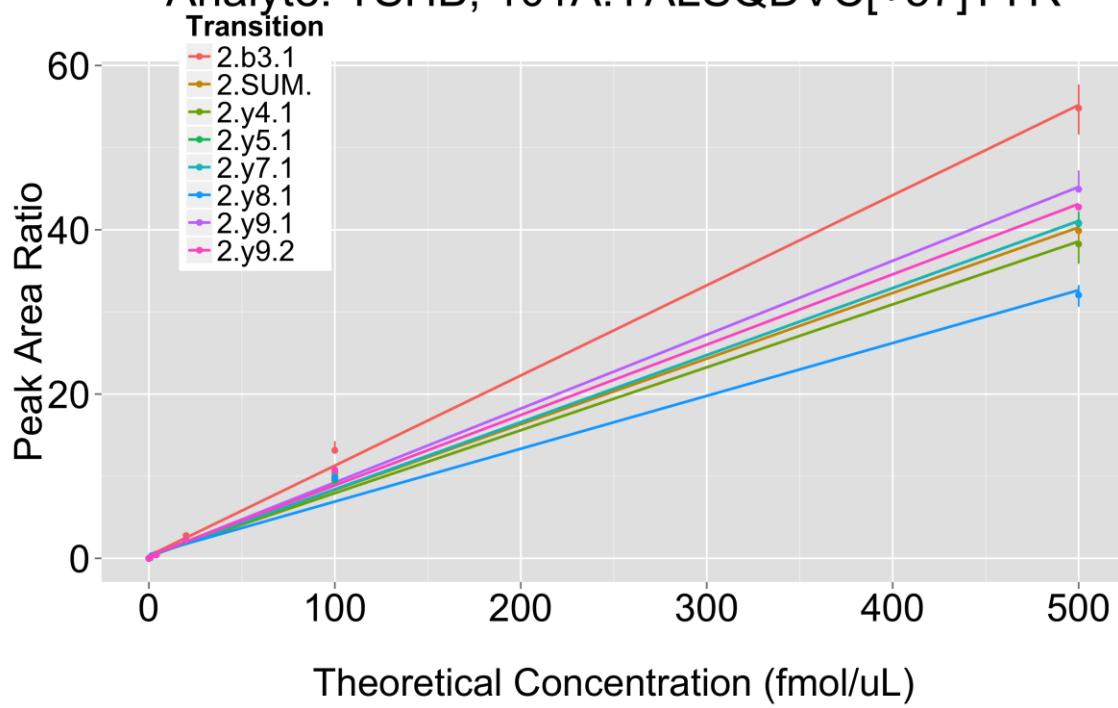
Analyte: SAIC43B.ASSIIDELFQDR



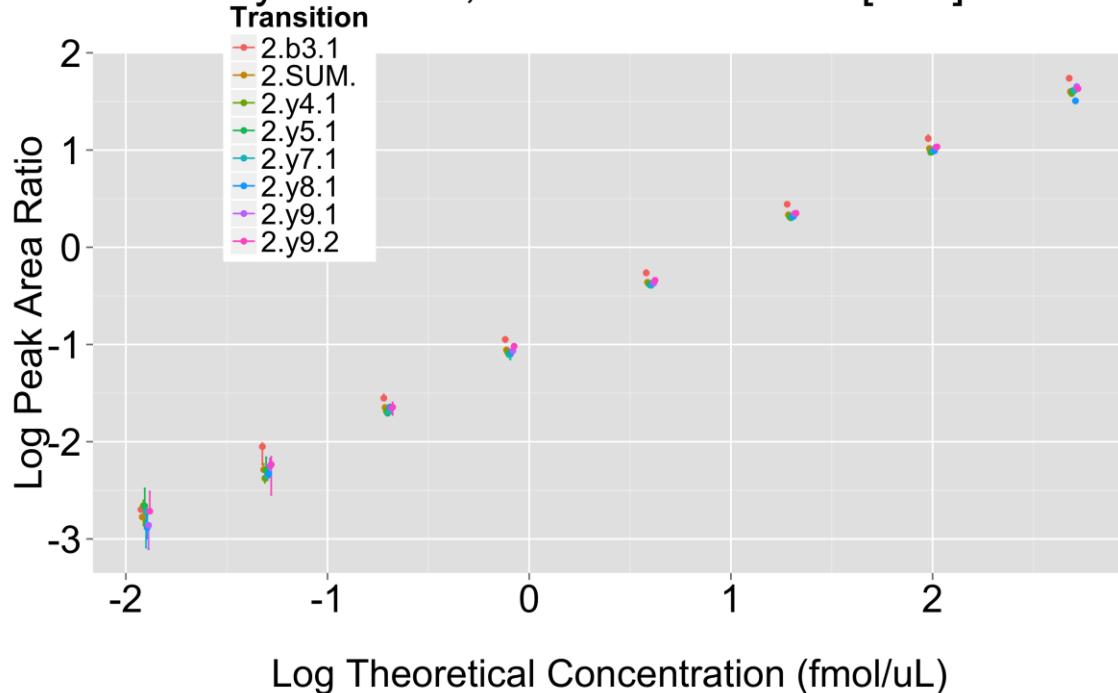
Analyte: SAIC43B.ASSIIDELFQDR

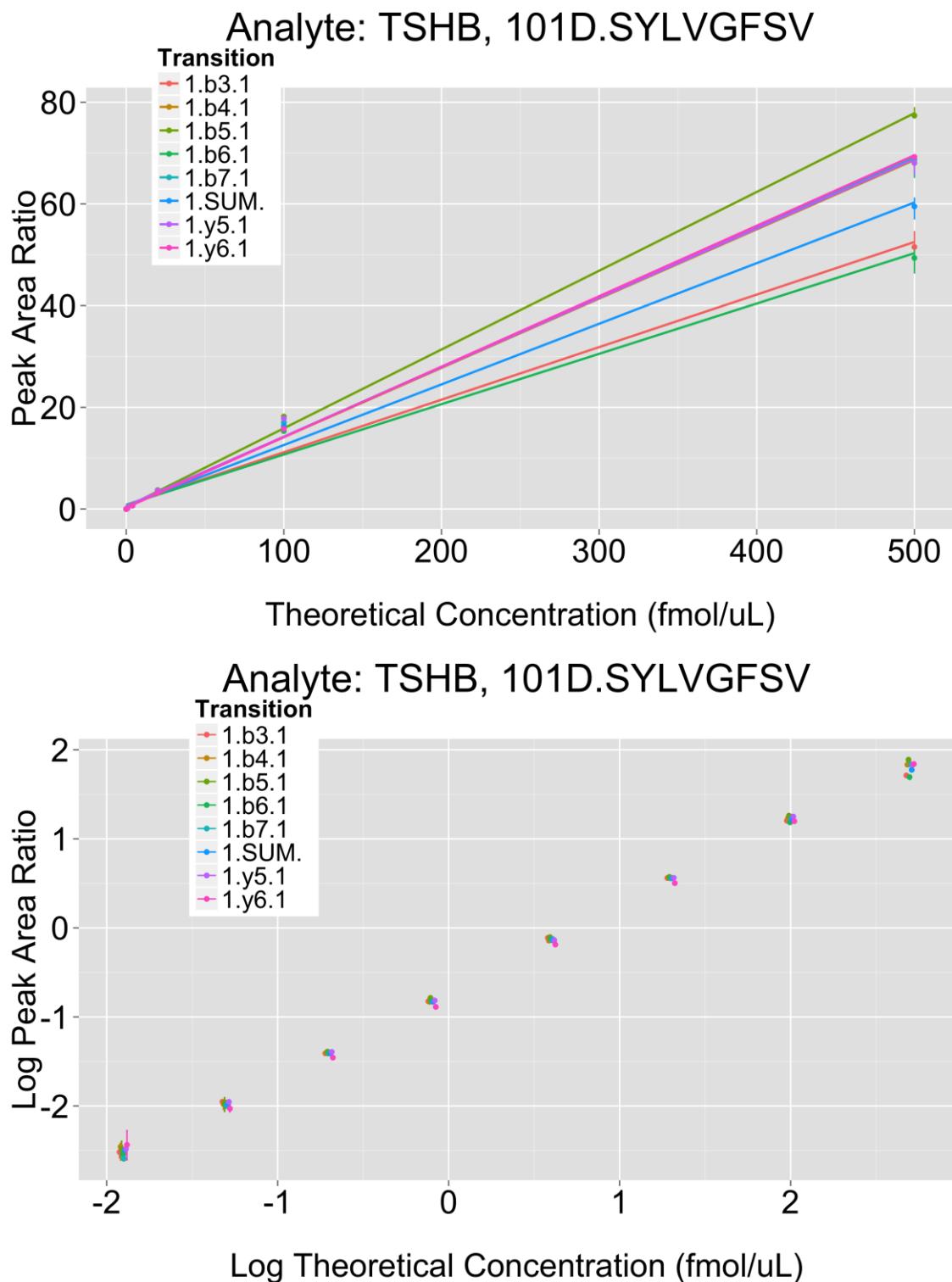


Analyte: TSHB, 101A.YALSQDV[+57]TYR

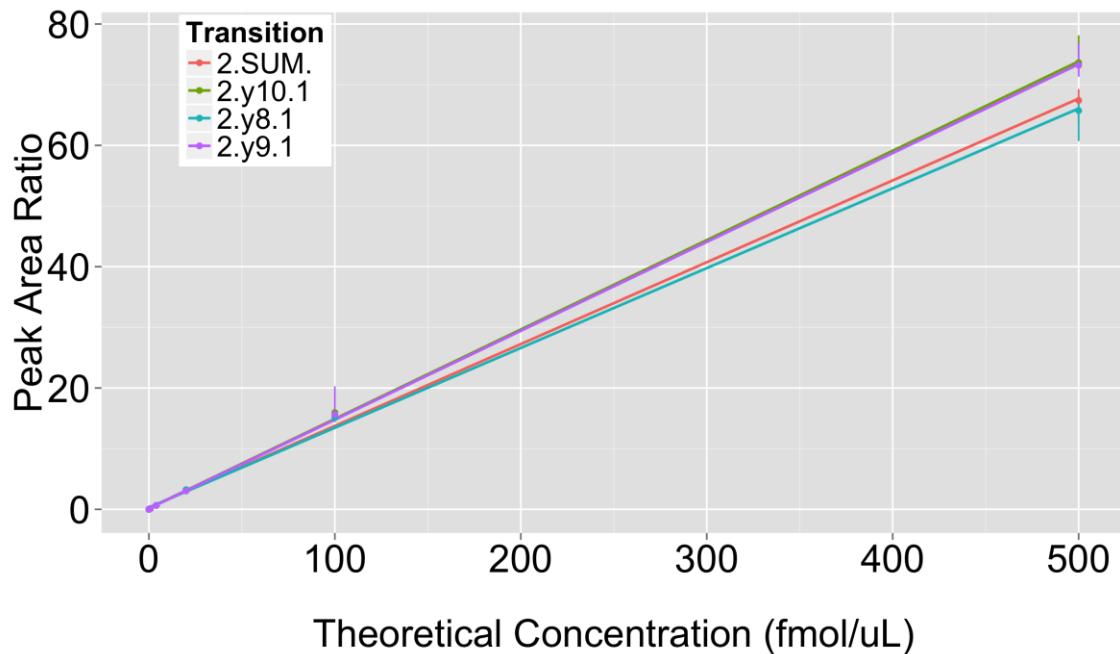


Analyte: TSHB, 101A.YALSQDV[+57]TYR

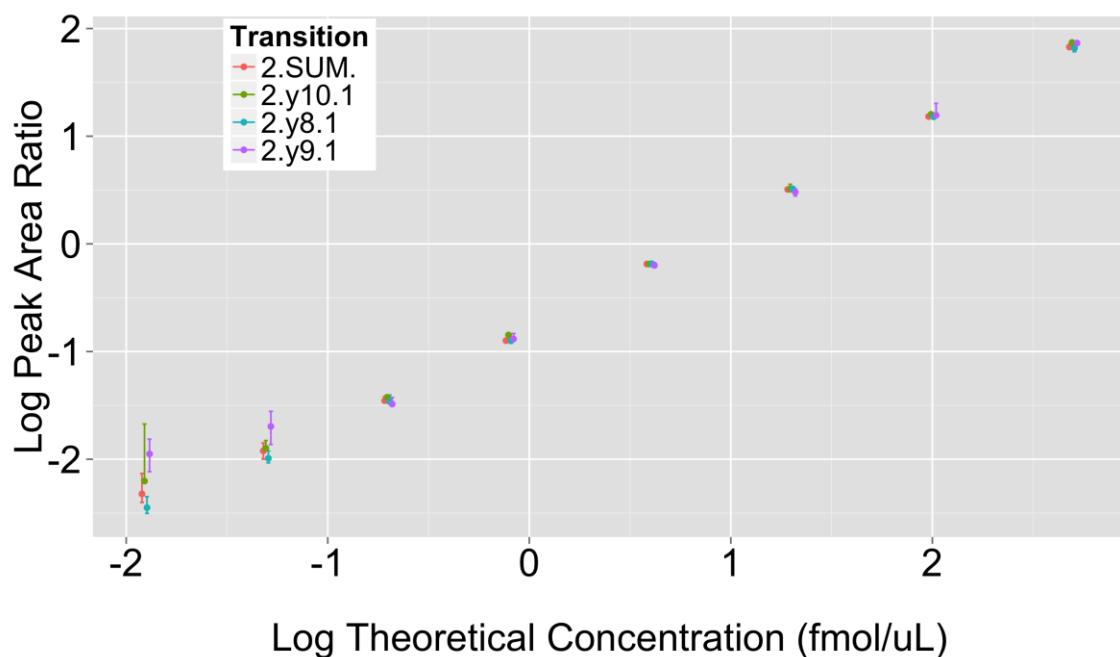




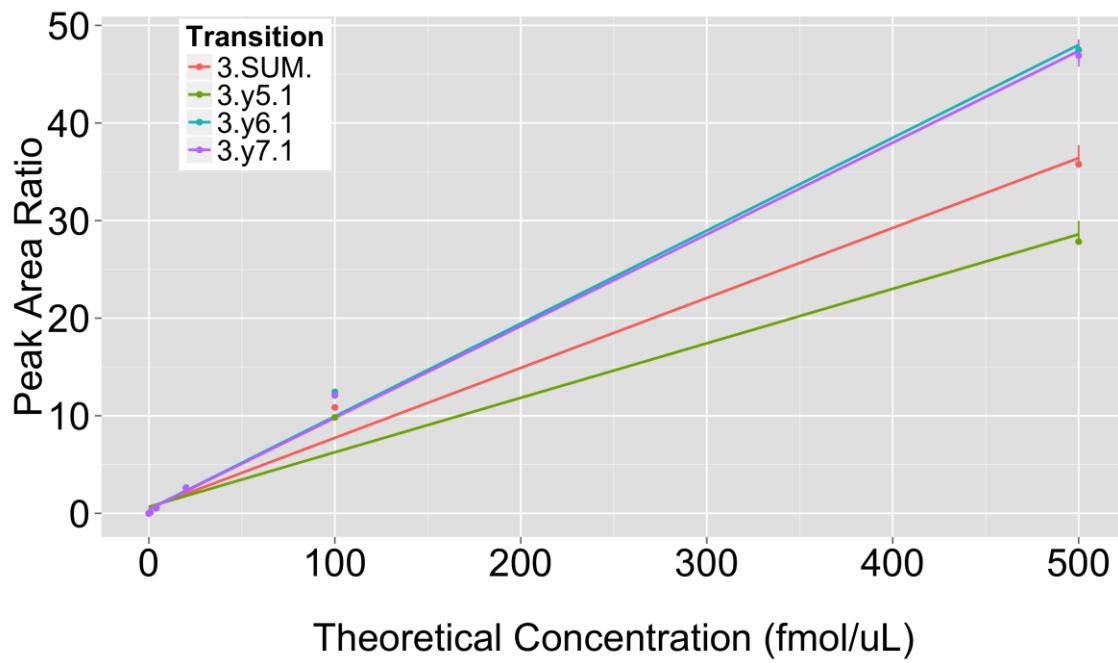
Analyte: CGB*, 102A.VLQGVLPALPQVVC[+57]NYR



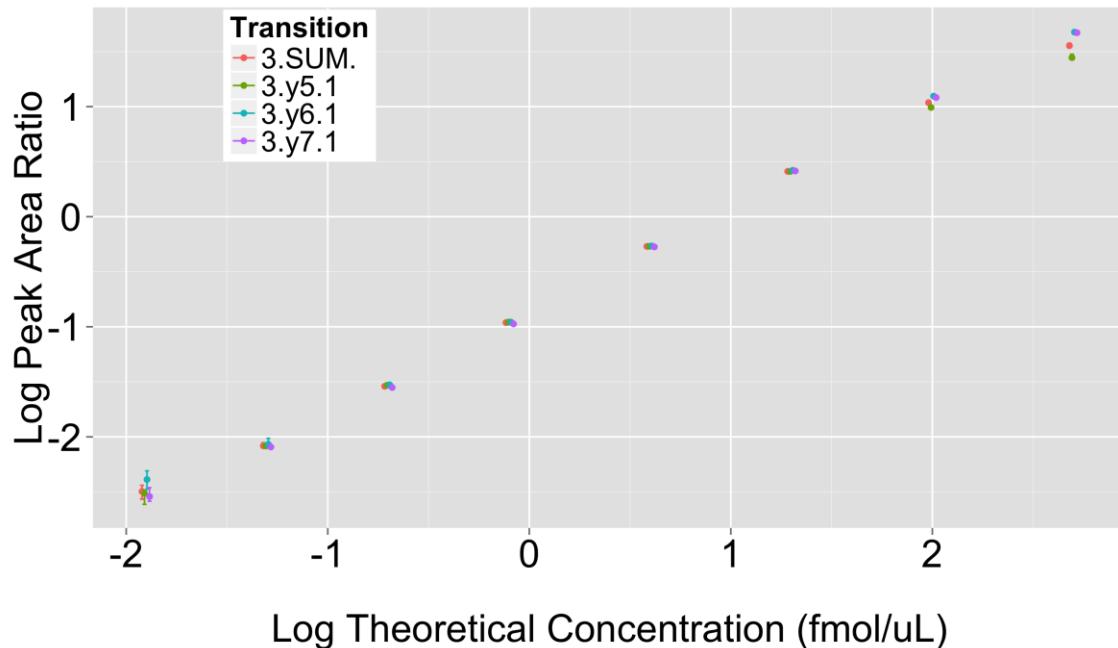
Analyte: CGB*, 102A.VLQGVLPALPQVVC[+57]NYR



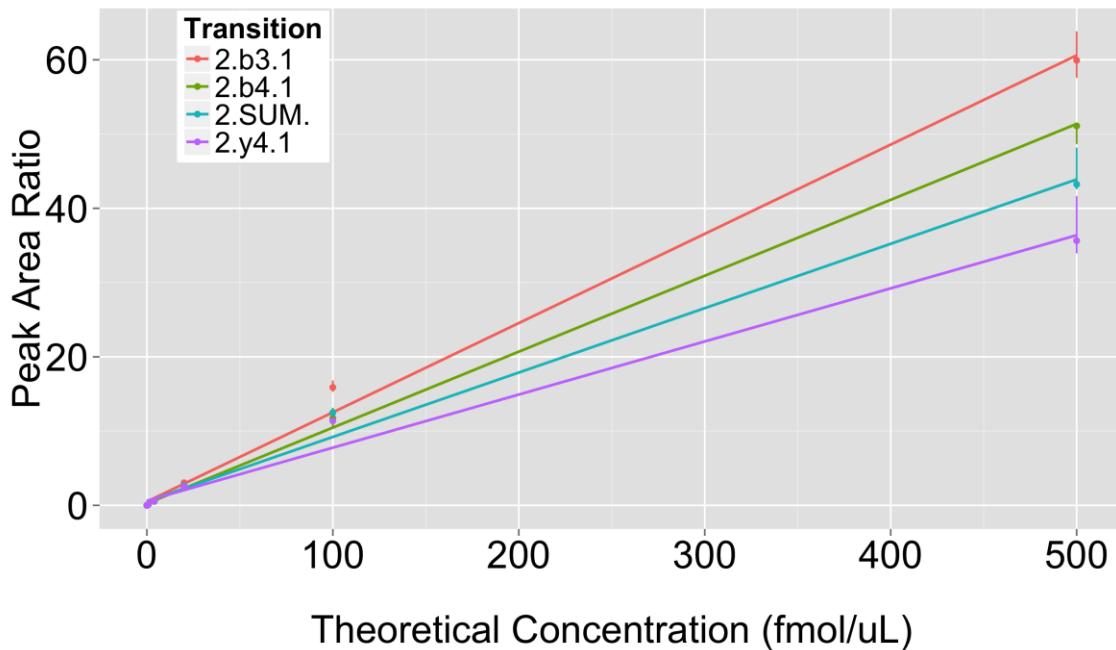
Analyte: MUC1, 103A.EGTINVHDVETQFNQYK



Analyte: MUC1, 103A.EGTINVHDVETQFNQYK



Analyte: MUC1, 103b.NYGQLDIFPAR



Analyte: MUC1, 103b.NYGQLDIFPAR

