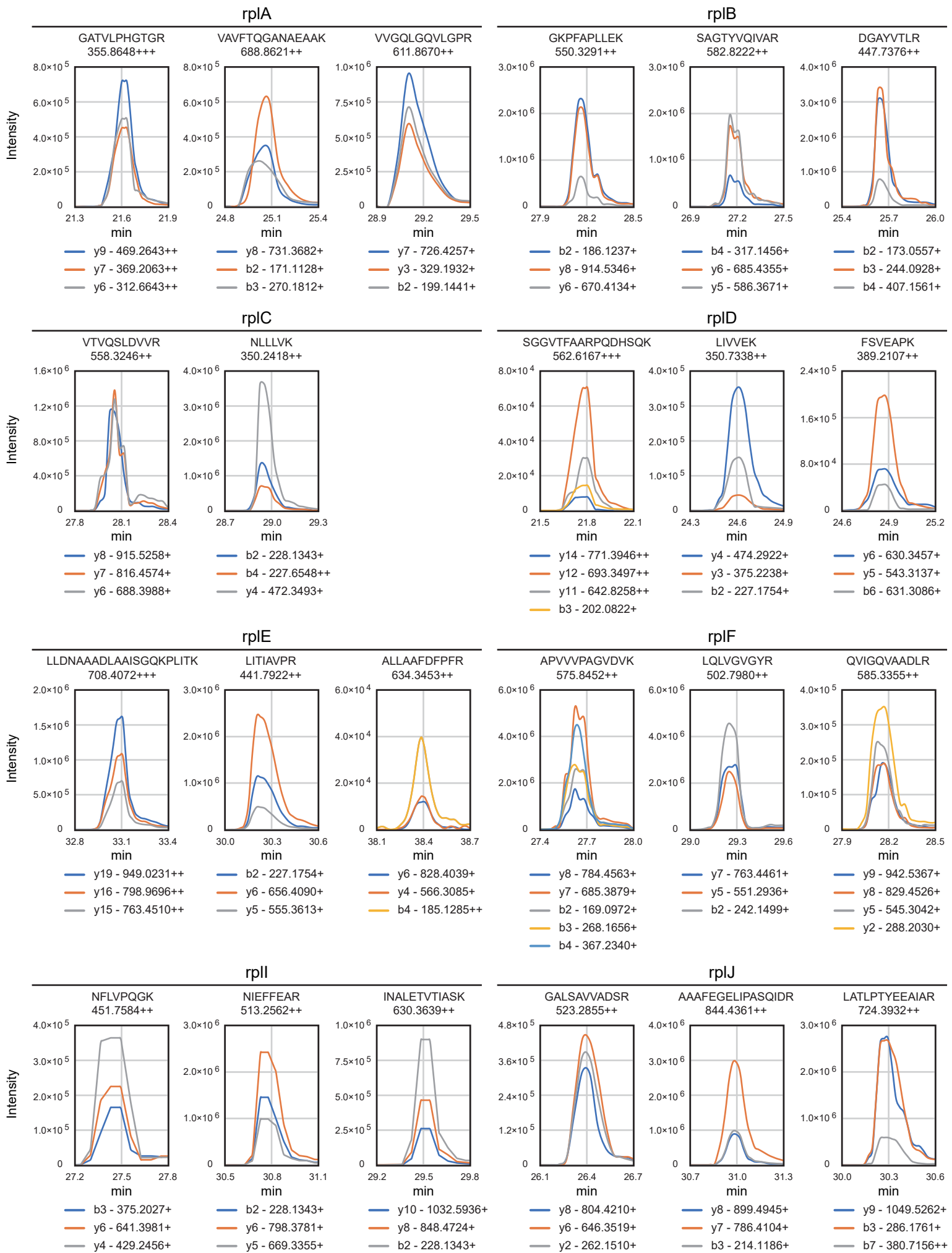
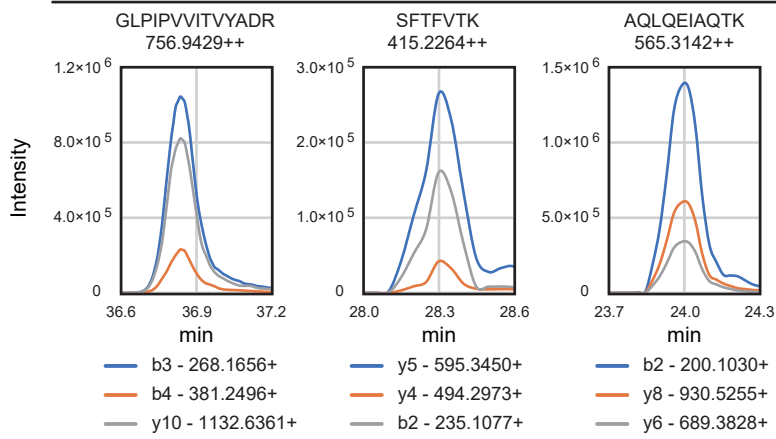


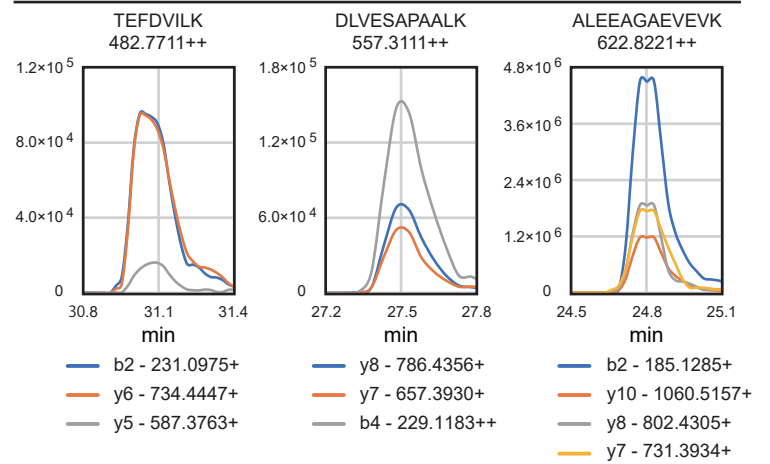
S2 Fig. Ion chromatograms corresponding to the selected transitions associated with peptides from ribosomal proteins. For each peptide, several transitions with intense peaks were selected based on the quantification of r-proteins from purified ribosomes. The calibration curves of all transitions are described in Table S2.



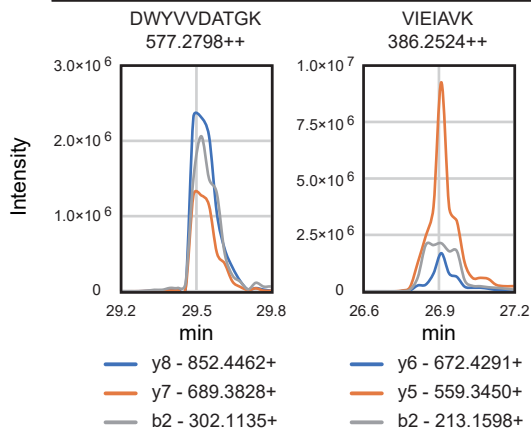
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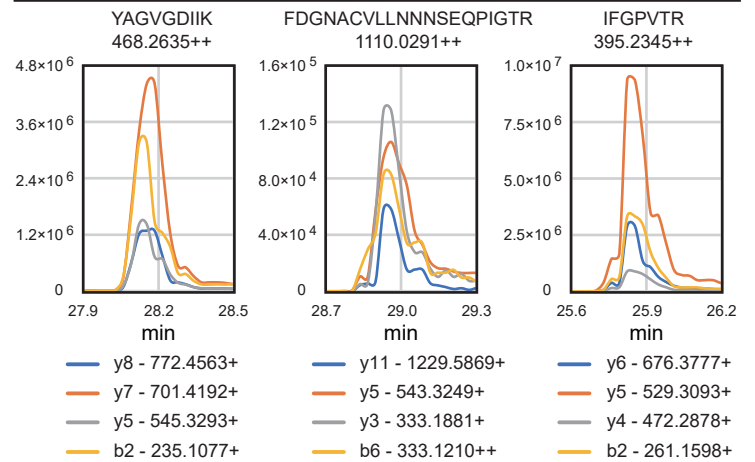
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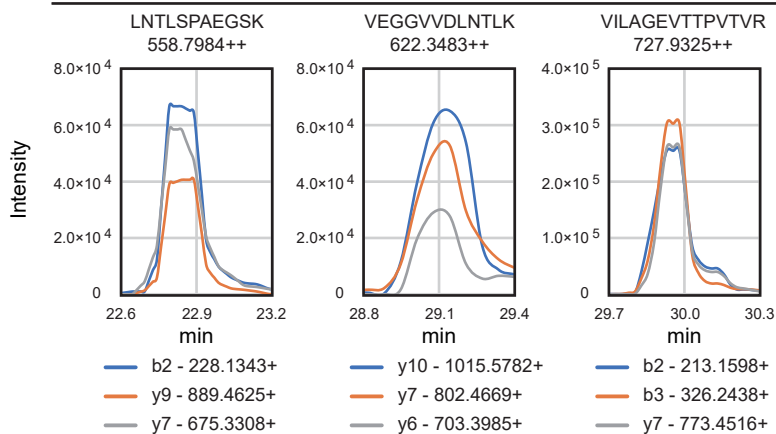
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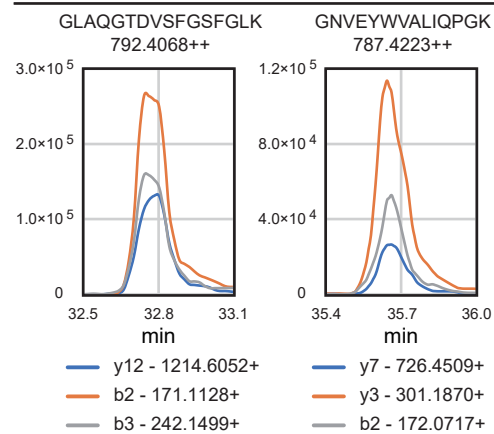
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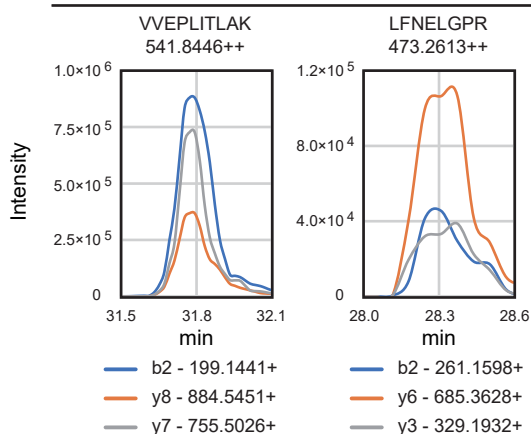
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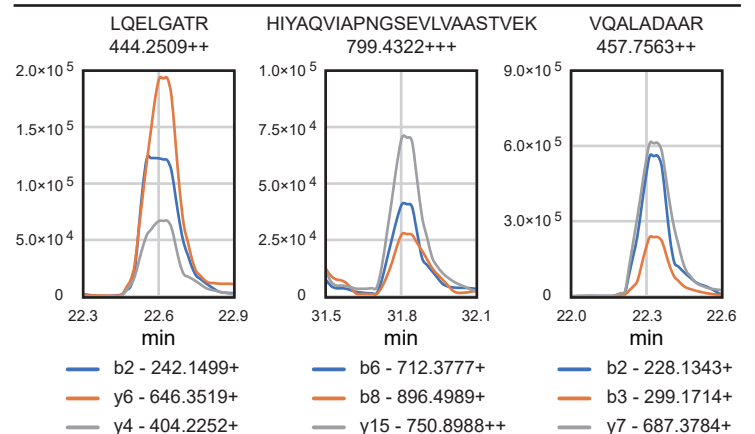
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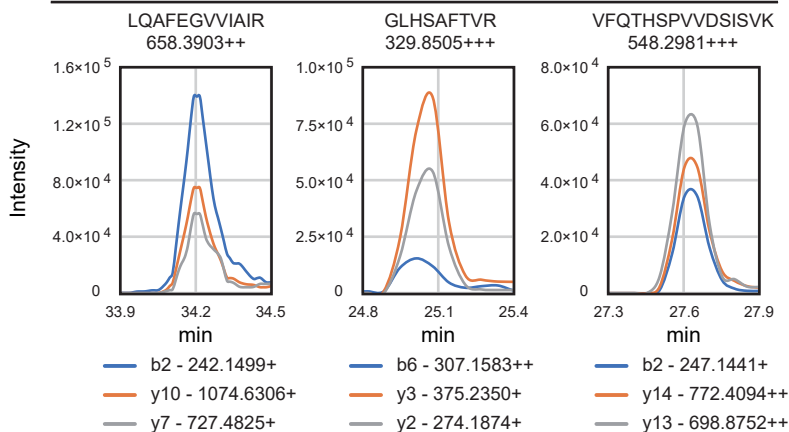
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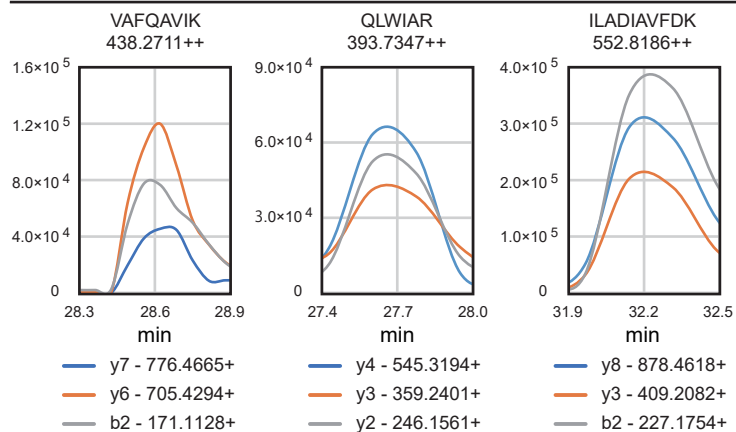
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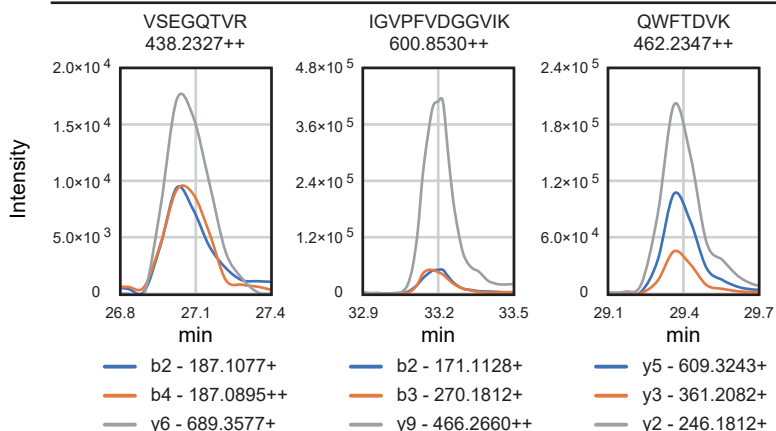
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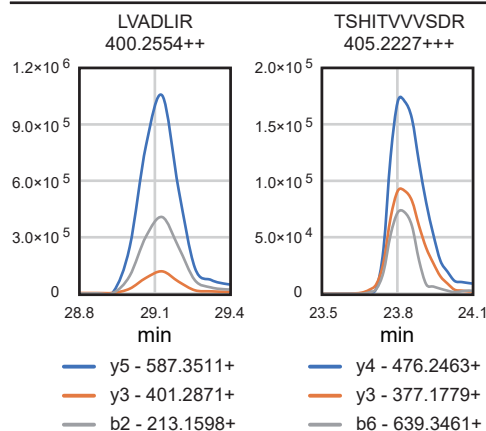
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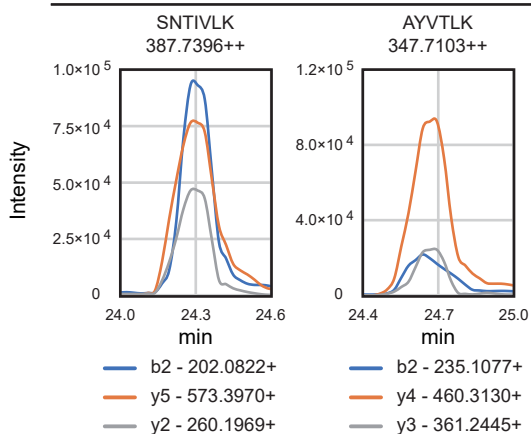
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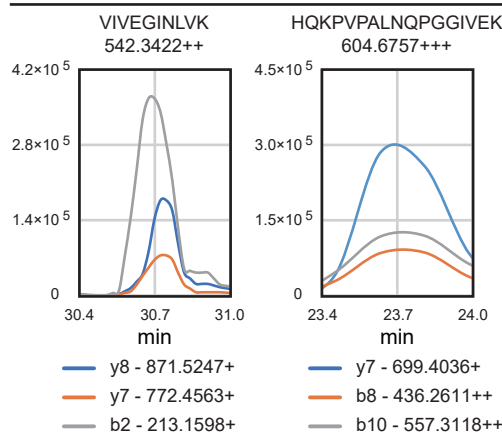
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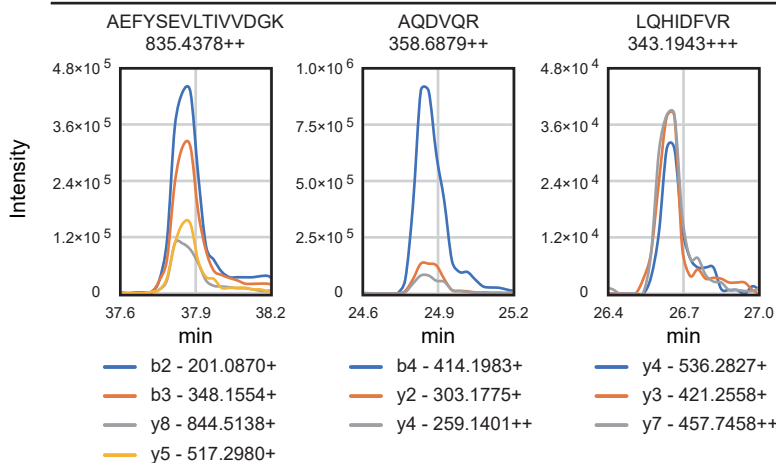
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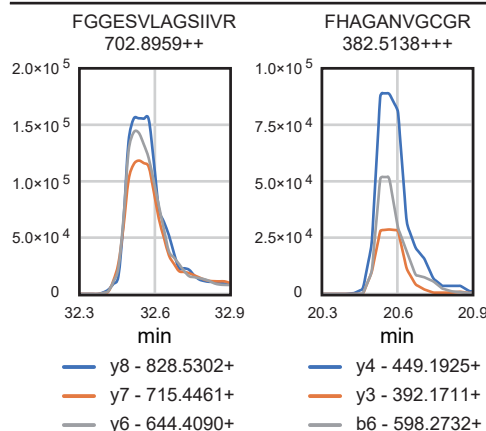
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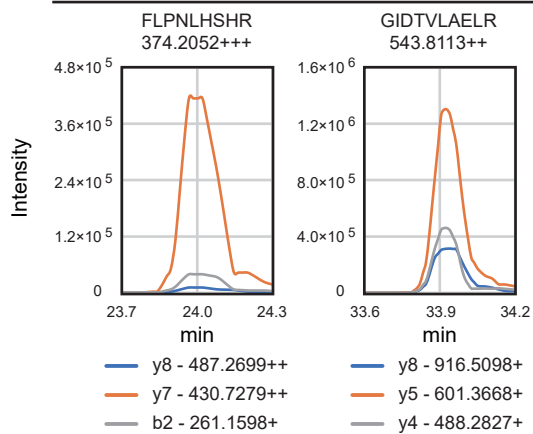
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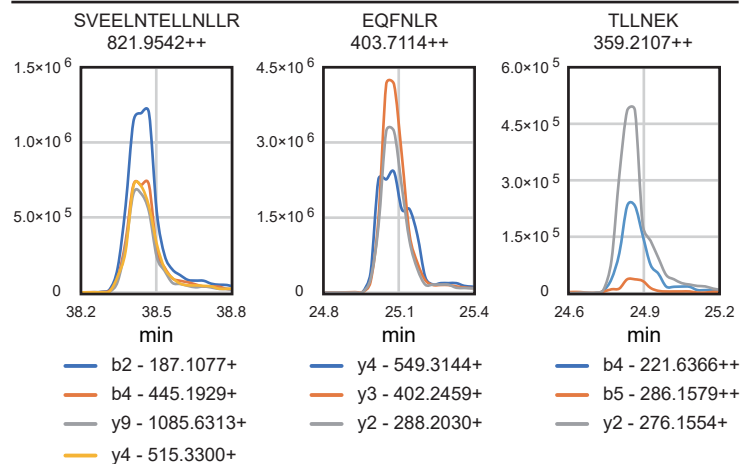
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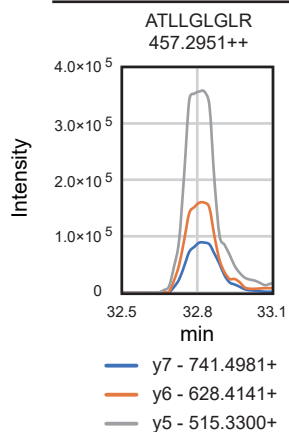
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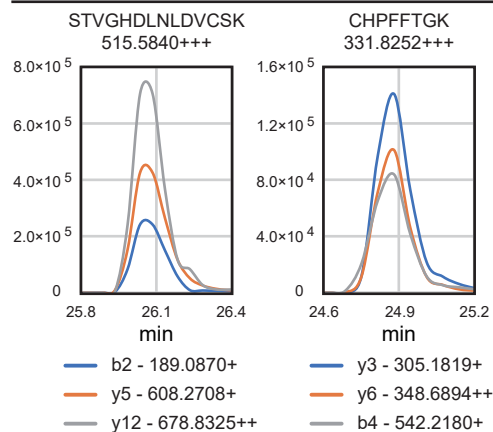
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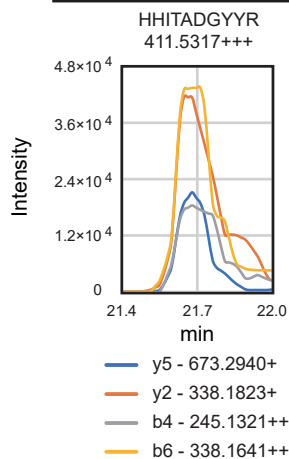
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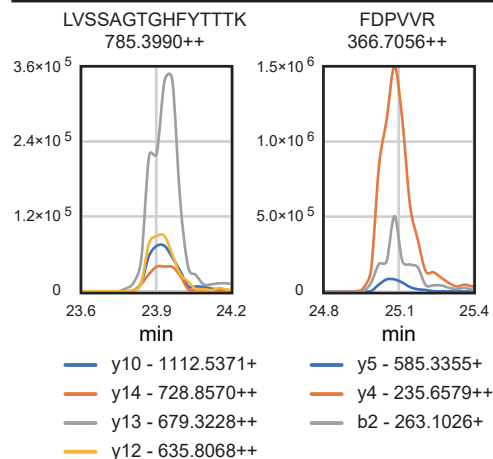
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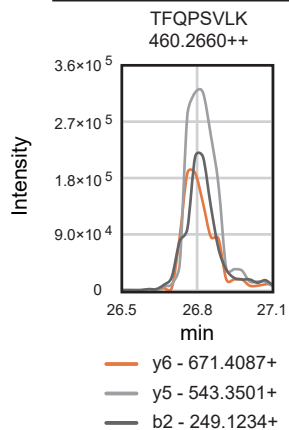
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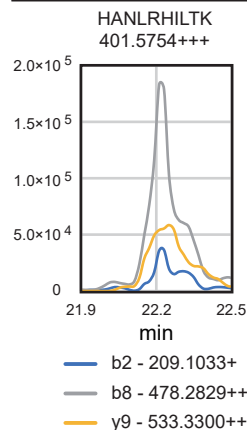
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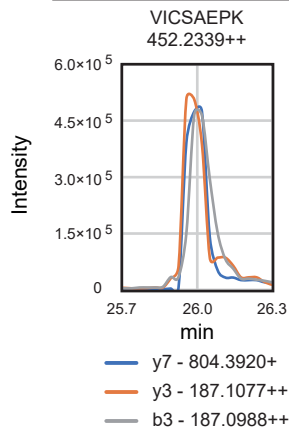
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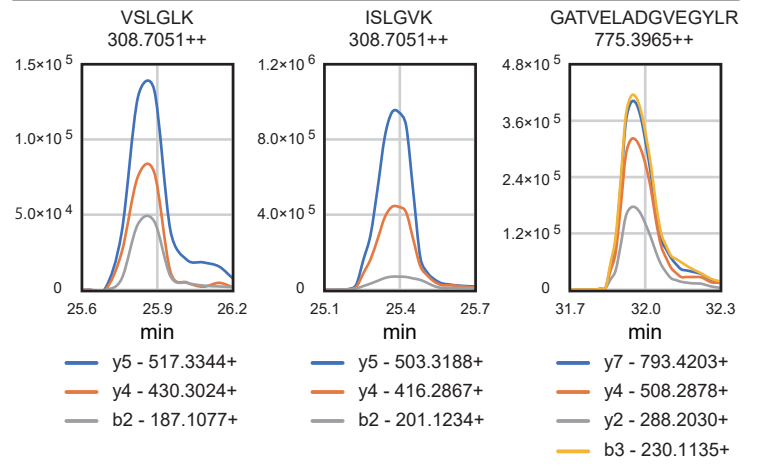
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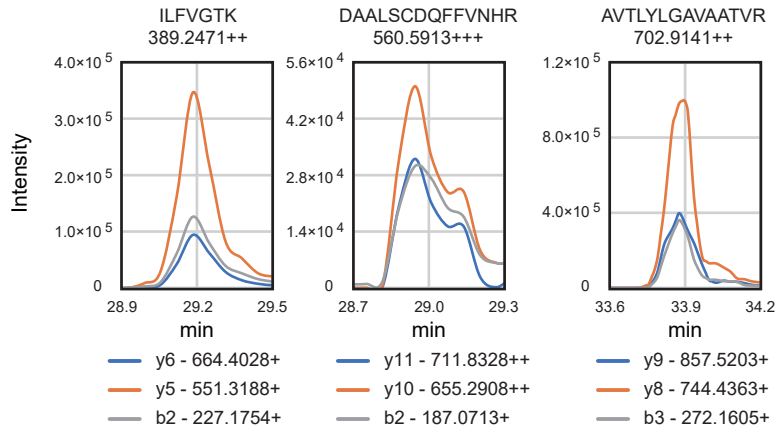
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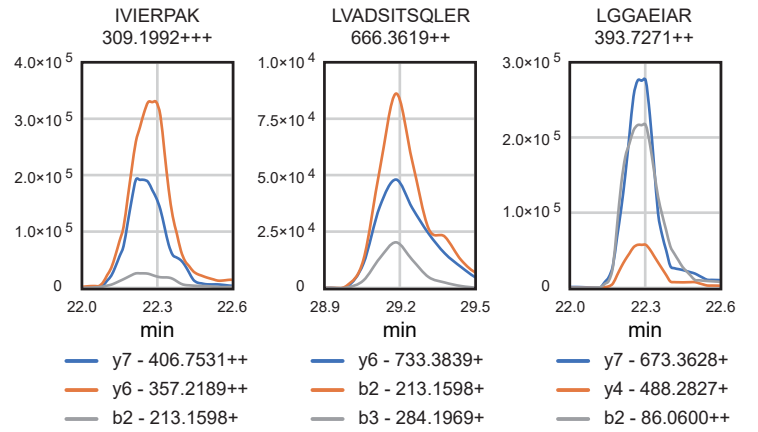
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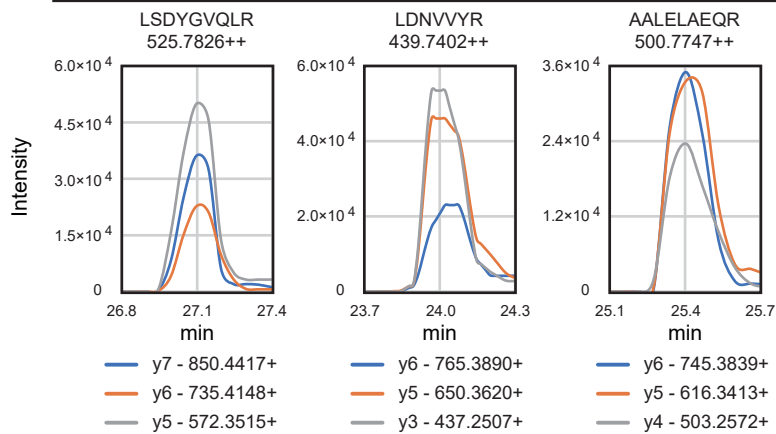
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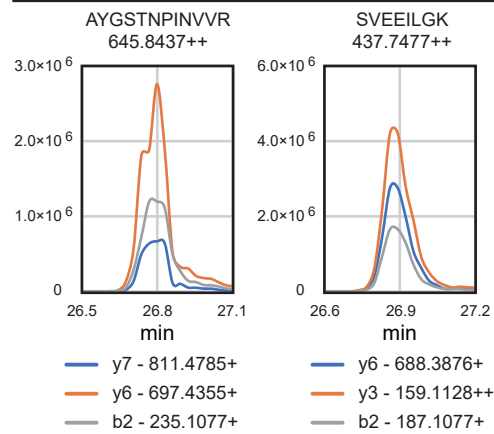
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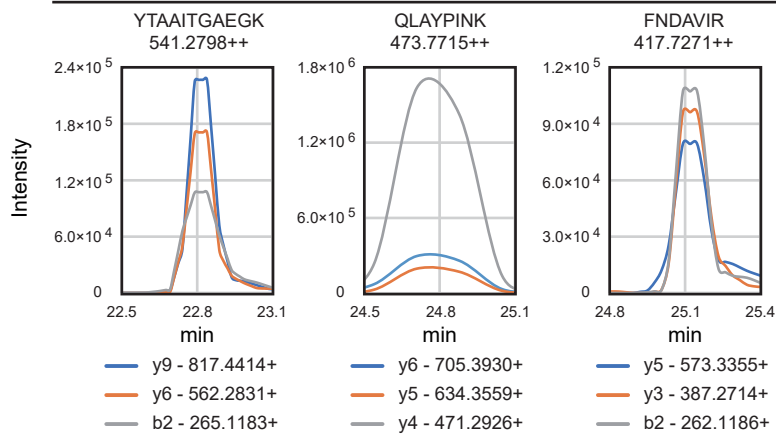
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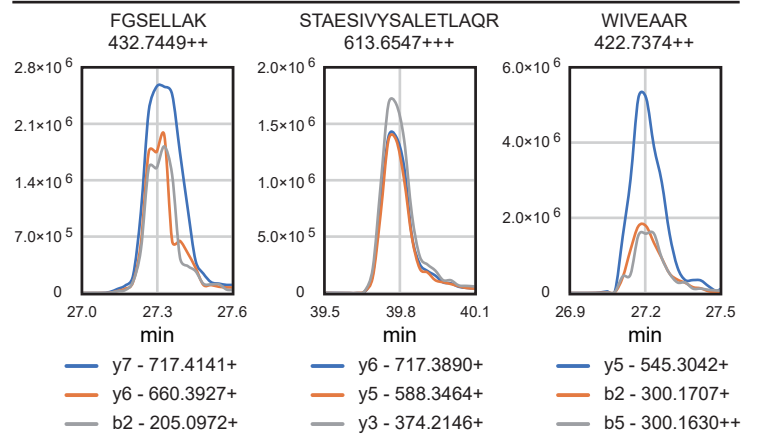
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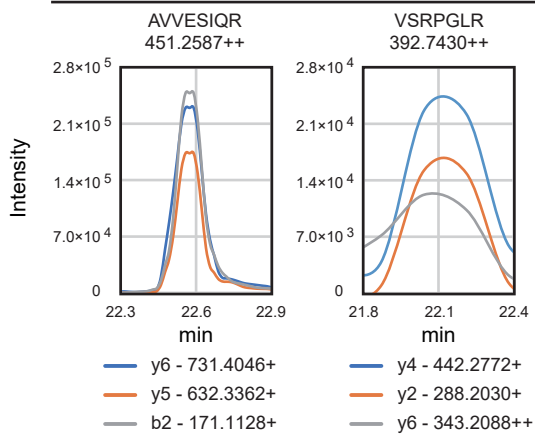
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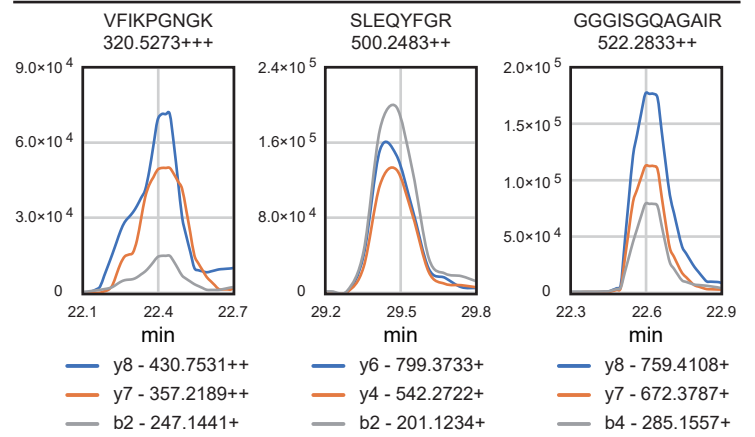
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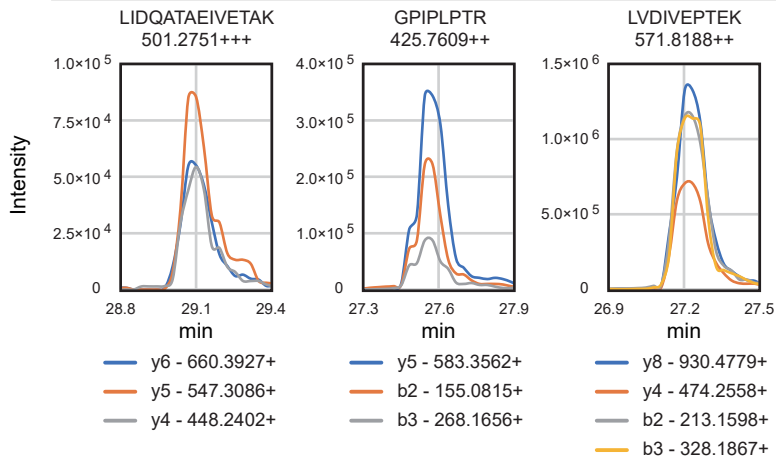
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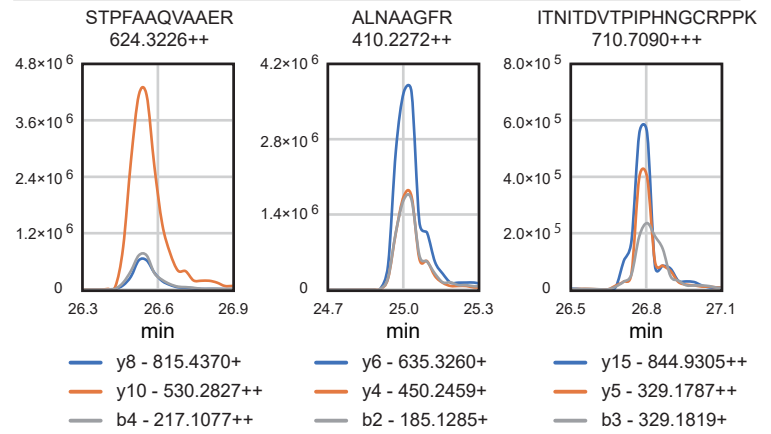
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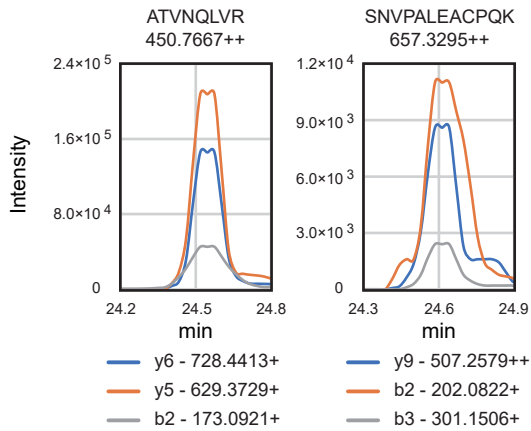
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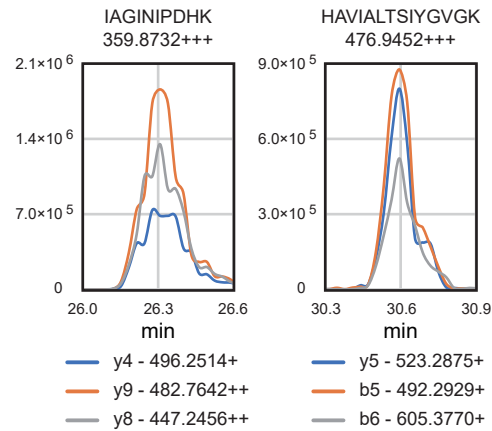
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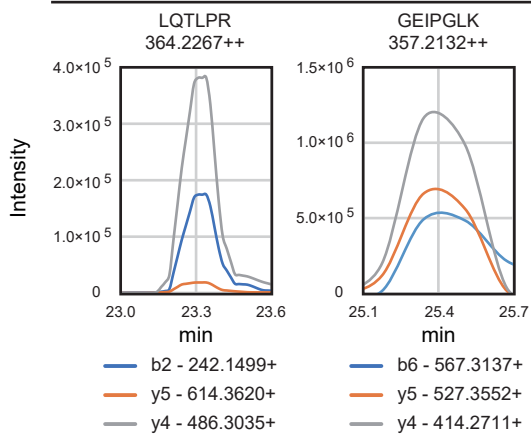
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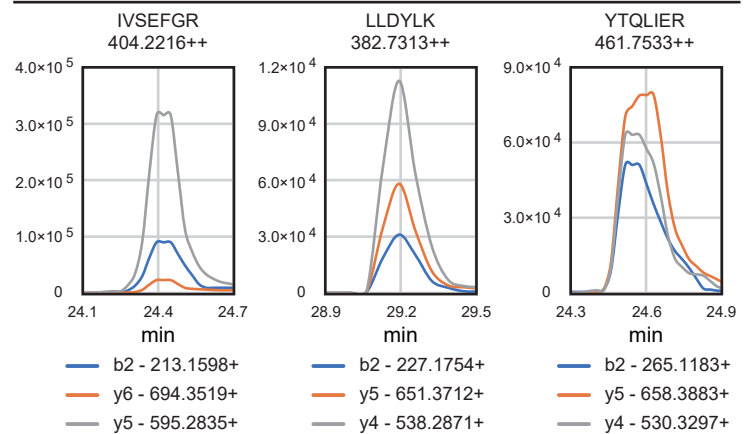
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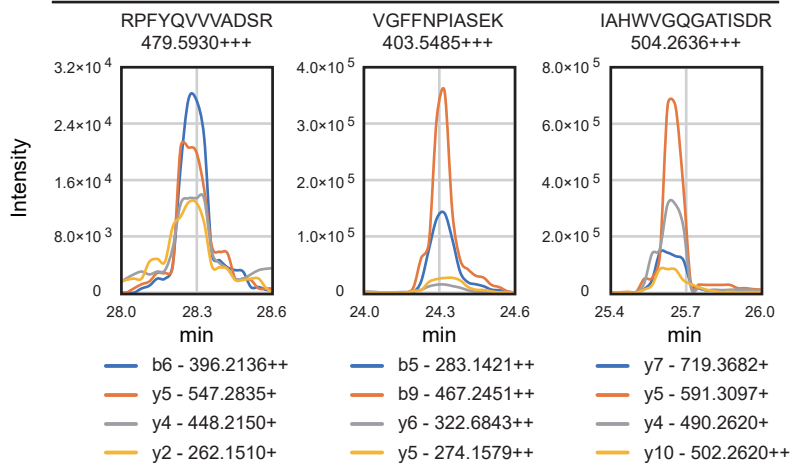
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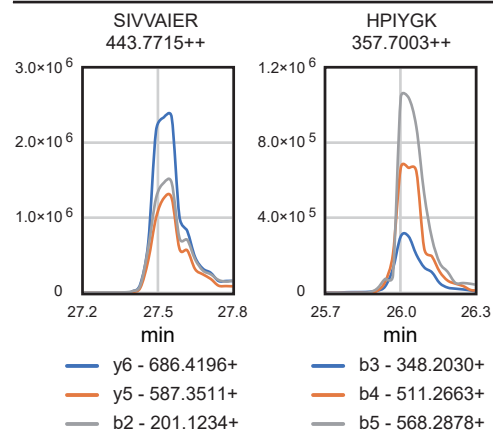
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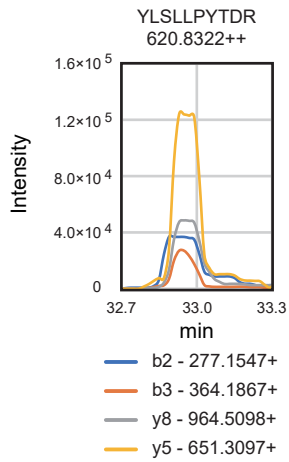
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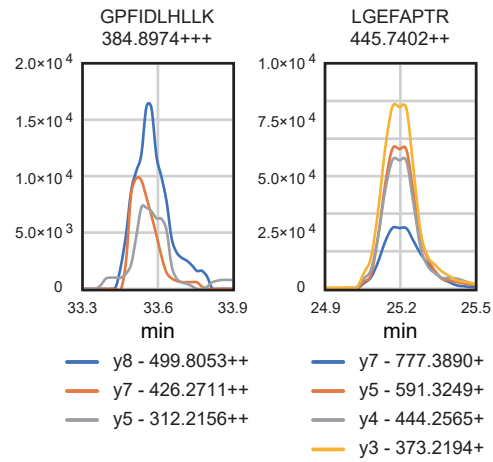
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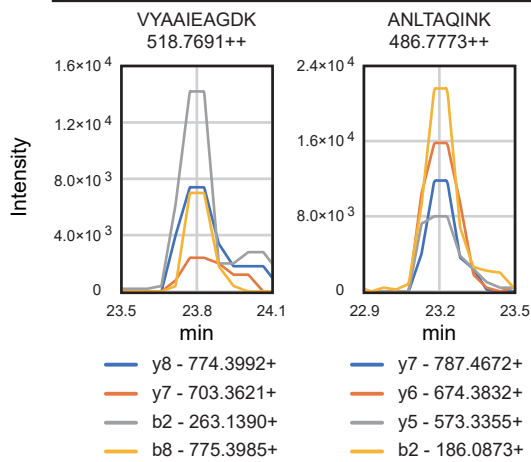
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rpsS



rpsT



rpsU

