

1 **Reversible Lysine Derivatization Enables Improved Arg-C (iArg-C)**
2 **Digestion, a Highly Specific Arg-C Digestion Using Trypsin**

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10 **Supplemental Information**

11 iArg-C Digestion Protocol S-2, S-3

12 Table S-1. The identification results using different citraconylation and
13 decitraconylation treatments.

14 Table S-2. The identification results using iArg-C and Arg-C digestion.

15 Table S-3. The identification results using Lys-C and trypsin digestion.

iArg-C Digestion Protocol

Sample preparation TIMING 3 h

- 1) Cell lysate. Add 20 volumes of lysis buffer (4 M guanidine hydrochloride, 100 mM TEAB) to the collected cells after washing three times with PBS buffer, and sonicate for 6 min (2 s sonication with 5 s intervals).
- 2) Collection. Collect supernatant via centrifugation at 20,000g for 20 min at 4 °C.
- 3) Protein determination. Determine the protein concentration using Bradford assay.
- 4) Reduction. Add 10 mM DTT and then incubate at 37 °C for 45 min.
- 5) Alkylation. Add 100 mM acrylamide and incubate for 1 h at room temperature.
- 6) Quenching. Add DTT to a final concentration of 50 mM to quench unreacted acrylamide.
- 7) Dilute the proteins to 1 mg/mL with lysis buffer.

Citraconylation TIMING 2 h

- 8) Add 5 µL of 2M citraconic anhydride to 200 µL protein sample, and then immediately add 5 µL 4M NaOH. After short vortex, incubate the sample for 10 min at room temperature.
- 9) Repeat step 8 for another four times to reach a final concentration of 200 mM citraconic anhydride.
- 10) Incubate the solution at room temperature for 1 h.

Digestion TIMING 15-18 h

- 11) Buffer displacement. Transfer samples from Eppendorf tubes to Microcon YM-10 filters and centrifuged at 13,800 g for three-time buffer displacement with 100 mM TEAB (pH 8.0). This step takes 3 to 6 h depending on sample

1 volume and protein concentration.

2 12) Digestion. Add digestion buffer (100 mM TEAB, pH 8.0) to reach the protein
3 concentration of 1 mg/mL. Add trypsin at a ratio of enzyme/protein as 1:50
4 and incubate at 37 °C for 12 h.

5 **Sample collection** TIMING 3-5 h

6 13) After digestion, collect the filtrate via centrifugation at 13,800g. To minimize
7 sample loss, wash the filter twice with 10% ACN and collect the filtrates via
8 centrifugation at 13,800g. Pool the three filtrates together.

9 14) Remove the remaining ACN and concentrate sample to about 1 mg/mL by a
10 Speedvac.

11 **Decitraconylation** TIMING 2 h

12 15) Decitraconylation. Add TFA to a final concentration of 1% (v/v) and incubate
13 at room temperature for 2 h.

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