

Isolation of the Dot/Icm Type IV Secretion System Core Complex from *Legionella pneumophila* for Negative Stain Electron Microscopy Studies

Tomoko Kubori* and Hiroki Nagai

Research Institute for Microbial Diseases, Osaka University, Osaka, Japan

*For correspondence: tkubori@biken.osaka-u.ac.jp

[Abstract] *Legionella* possesses a pivotal secretion machinery to deliver virulence factors to eukaryotic host cells. In this protocol, we describe the procedure for isolation of the native core complex of the Dot/Icm type IV secretion system from *L. pneumophila* aiming to perform biochemical and transmission electron microscopy analyses.

Keywords: *Legionella*, Type IV secretion, Core complex, Isolation, Electron microscopy, Bacteria, Nanomachine, Structure

[Background] *Legionella pneumophila* is a Gram-negative bacterial pathogen that causes lung infection known as Legionnaires' disease (Fields *et al.*, 2002). *L. pneumophila* utilizes a type IV secretion system (T4SS) encoded by the *dot/icm* genes to transport about 300 bacterial proteins into the cytosol of their eukaryotic host to hijack cellular processes (Hubber and Roy, 2010). Composed of more than 20 proteins, the T4SS is a nanomachine built on the bacterial inner and outer membranes (Nagai and Kubori 2011; Kubori and Nagai 2016). The core complex of Dot/Icm T4SS is a biochemically stable part of the system and forms a transport conduit bridging the inner and outer membrane (Kubori *et al.* 2014). The core complex is composed of at least five proteins; three outer membrane-associated proteins, DotC, DotD and DotH, and two inner membrane proteins, DotF and DotG (Vincent *et al.*, 2006). Based on the procedure for biochemical isolation of another bacterial nanomachine, the type III secretion system, from *Salmonella typhimurium* (Kubori *et al.*, 1998; Marlovits *et al.*, 2004), we modified the protocol to adapt it to the purification of the T4SS of *L. pneumophila*. In this protocol, we present the procedure to isolate the native core complex of the T4SS from detergent lysed wild-type *L. pneumophila* based on separation by ultracentrifugation. T4SS isolated using this procedure can be used to perform biochemical and transmission electron microscopy analyses described previously (Kubori *et al.*, 2014).

Materials and Reagents

1. Sterile swabs
2. Sterile cell scrapers (IWAKI, catalog number: 9000-220)
3. Sterile conical tubes (50 ml and 15 ml)
4. Sterile Petri dishes (100 mm in diameter)
5. Cuvettes for spectrophotometer (1.5 ml) (BOECO, catalog number: BRA 759017)
6. Millex-GP filter units (EMD Millipore, catalog number: SLGP033RS)

7. Sterile 10 ml syringe (Terumo, catalog number: SS-10LZ)
8. Ultrafree MC filters (EMD Millipore, catalog number: UFC30GV00)
9. Sterile pipets (10 ml) (Thermo Fisher Scientific, Thermo Scientific™, catalog number: 170356)
10. *L. pneumophila* Lp01 strain (Philadelphia-1 *rpsL hsdR*) (Berger and Isberg, 1993)
11. cOmplete™ protease inhibitor cocktail (Roche Diagnostics, catalog number: 11697498001)
12. Sodium chloride (NaCl) (Nacalai Tesque, catalog number: 31320-05)
13. 12.5% precast polyacrylamide gels (ATTO, catalog number: e-PAGEL E-R12.5L; or equivalent)
14. Glow-discharged carbon grids (Nisshin EM, catalog number: 649)
15. Coomassie brilliant blue (CBB) stain One (Nacalai Tesque, catalog number: 04543-51)
16. ACES (Sigma-Aldrich, catalog number: A3594)
17. Bacto™ yeast extract (BD, Bacto™, catalog number: 212750)
18. MilliQ water
19. Activated charcoal (Sigma-Aldrich, catalog number: C5510)
20. Bacto™ agar (BD, Bacto™, catalog number: 214010)
21. L-cysteine hydrochloride monohydrate (Nacalai Tesque, catalog number: 10313-55)
22. Iron(III) nitrate enneahydrate, Fe(NO₃)₃·9H₂O (Nacalai Tesque, catalog number: 19514-55)
23. Tris(hydroxymethyl)aminomethane (Tris) (Nacalai Tesque, catalog number: 35434-21)
24. Hydrochloric acid (HCl) (Nacalai Tesque, catalog number: 18321-05)
25. Sucrose (Nacalai Tesque, catalog number: 30404-45)
26. Phenylmethylsulfonyl fluoride (PMSF) (Nacalai Tesque, catalog number: 27327-94)
27. Isopropanol (Sigma-Aldrich, catalog number: 190764)
28. EDTA·2Na (Nacalai Tesque, catalog number: 15130-95)
29. Sodium hydroxide (NaOH) (Nacalai Tesque, catalog number: 31511-05)
30. Lysozyme (Wako Pure Chemical Industries, catalog number: 120-02674)
31. Triton X-100 (Nacalai Tesque, catalog number: 35501-15)
32. AG501-X8 Resin (Bio-Rad Laboratories, catalog number: 143-7425)
33. Magnesium sulfate heptahydrate (MgSO₄·7H₂O) (Nacalai Tesque, catalog number: 21003-75)
34. DNase I (Sigma-Aldrich, catalog number: DN25)
35. Potassium hydroxide (KOH) (Nacalai Tesque, catalog number: 28616-45)
36. Uranyl acetate (UA) (Merck, catalog number: 8473)
37. Phosphotungstic acid (PTA) (TAAB, catalog number: p013)
38. CYE plate (see Recipes)
39. AYE medium (see Recipes)
40. Tris-Cl solution (pH 8.0) (see Recipes)
41. Sucrose solution (see Recipes)
42. PMSF stock solution (see Recipes)
43. EDTA stock solution (see Recipes)
44. Lysozyme solution (see Recipes)
45. Triton X-100 stock solution (see Recipes)

46. MgSO₄ stock solution (see Recipes)
47. DNase I stock solution (see Recipes)
48. KOH solution (see Recipes)
49. NaOH stock solution (see Recipes)
50. TET solution (see Recipes)
51. PTA solution (see Recipes)
52. Uranyl acetate solution (see Recipes)

Equipment

1. Glass flasks (2 L)
2. 37 °C shaking incubator
3. 37 °C incubator
4. Spectrophotometer
5. Refrigerated centrifuge (KUBOTA, model: 7780; or equivalent models)
6. Rotor for centrifugation (KUBOTA, models: AG-5006, AG-6512C)
7. Sterile centrifuge tubes (500 ml capacity, polypropylene or polycarbonate)
8. Sterile centrifuge tubes (50 ml capacity, polyallomer or polycarbonate)
9. Clean glass beakers (200 ml)
10. Magnetic stir bars
11. Magnetic stirrer
12. pH meter
13. Ultracentrifugation (Beckman Coulter, model: Optima™ L-100 XP; or equivalent models)
14. Rotor for ultracentrifugation (Beckman Coulter, model: Type 70Ti)
15. Tubes for ultracentrifugation (Beckman Coulter, catalog number: 355631)
16. Microfuge (Eppendorf, model: 5415 R)
17. Glass beakers (1 L)
18. Autoclavable flasks (2 L)
19. Autoclave
20. Electrophoresis apparatus (ATTO, model: AE-6530; or equivalent model)
21. Electron microscope (JEOL, model: JEM-1011)
22. ÄKTA purifier (GE Healthcare)
23. Superose 6 10/300 GL column (GE Healthcare, catalog number: 17517201)

Procedure

1. Grow *L. pneumophila* Lp01 strain on a charcoal-yeast extract (CYE) plate (Recipe No. 1) from a glycerol frozen stock for 3 days at 37 °C.
2. Use a sterile swab to streak a *L. pneumophila* colony on the entire surface of CYE plates and

cultivate for 2 days at 37 °C.

Note: Two plates covered with a bacterial lawn are needed to obtain sufficient starting material to prepare 1 L culture.

3. Take the whole bacteria using sterile cell scrapers and suspend it with ~5 ml ACES-buffered yeast extract (AYE) medium (Recipe No. 2) in a sterile tube. Measure OD₆₀₀. Add the bacterial solution into 1 L AYE medium in a 2 L flask to make the suspension of OD₆₀₀ 0.2.
4. Grow the bacteria for 12 h at 37 °C with rotary shaking (250 rpm).
5. Chill the flask in icy water. Measure OD₆₀₀. (It should be 2.0-3.0, indicating the late log phase of growth). Transfer the culture solution to chilled centrifuge tubes (2-3 tubes of 500 ml capacity).
6. Harvest the bacteria by centrifugation at 12,000 x g (8,000 rpm) for 15 min at 4 °C (Refrigerated centrifuge: Kubota, model: 7780 or equivalent models; Rotor: Kubota, model: AG-5006).
7. Resuspend the bacterial pellet with 140 ml of cold sucrose solution (Recipe No. 4) containing 1x cOmplete™ protease inhibitor cocktail.

Note: Thoroughly suspending bacterial pellets enhances the efficiency of the detergent lysis in step 11.

8. Transfer the suspension to a 200 ml beaker with a magnetic stir bar on ice.
9. Set the beaker on a magnetic stirrer at room temperature, and stir mildly until uniform suspension is achieved.
10. Add PMSF (final 1 mM, Recipe No. 5), EDTA (final 1 mM, Recipe No. 6) and lysozyme (final 0.1 mg/ml, Recipe No. 7) in this order.

Note: This is the process of spheroplast formation.

Prepare a 20 mg/ml 200x stock solution. Add 0.7 ml of this stock solution to the 140 ml sucrose solution, yielding a final lysozyme concentration of approximately 0.1 mg/ml in the bacterial resuspension. Keep stirring for another 30 min at room temperature.

11. Add slowly (drop by drop) 7 ml of 20% (w/v) Triton X-100 stock solution (final 1% w/v, Recipe No. 8) with stirring.

Note: This is the most important step for detergent lysis of bacterial membranes. Carefully monitor the change of color and viscosity.

12. Keep stirring until the solution becomes very clear (approximately for 30 min).
13. Add 420 µl of 1 M MgSO₄ stock solution (final 3 mM, Recipe No. 9) and 70 µl of 10 mg/ml DNase I stock solution (final 5 µg/ml, Recipe No. 10) in this order.

Note: This step is required to digest DNA.

14. Stir for 10 min.
15. Add EDTA (final 10 mM).
16. Monitoring pH using a pH meter, adjust pH to 10.0 by adding 1 N NaOH (Recipe No. 12).

Note: During this step, large membrane vesicles and fragments that remain are disrupted. The core complex is stable at this high pH, while loosely associated contaminant proteins can be detached from the core complex. For the purpose of analyzing the loosely associated components of the T4SS, pH value can be modified to maintain these proteins in the complex.

17. Move the beaker on ice.

Note: Procedure below should be done at 4 °C.

18. Transfer the lysate to chilled 50 ml centrifuge tubes (divide into 3 tubes).

19. Centrifuge the lysate at 12,000 $\times g$ for 20 min at 4 °C to remove non-lysed materials. (Refrigerated centrifuge: Kubota, model: 7780 or equivalent models; Rotor: Kubota, model: AG-6512C).

20. Recover supernatant and transfer to ultracentrifuge tubes (Beckman Coulter).

21. Apply ultracentrifugation at 100,000 $\times g$ for 30 min at 4 °C to precipitate protein complexes. (Ultracentrifuge: Beckman Coulter, model: Optima L-100 XP or equivalent models; Rotor: Beckman Coulter, model: Type 70Ti)

22. Discard supernatant. Soak the pellet with 0.5 ml of cold TET solution (Recipe No. 13) including 1 mM PMSF per tube. To dissolve completely, leave the pellet in TET overnight at 4 °C.

23. Merge the completely dissolved suspensions and transfer them in a new conical or centrifuge tube (total ~5 ml). Use extra 2-3 ml of cold TET solution to completely recover the dissolved proteins from the tubes and merge them into the new tubes.

24. Centrifuge the suspension at 14,000 $\times g$ for 15 min at 4 °C to remove precipitate. (Refrigerated centrifuge: Kubota, model: 7780 or equivalent models; Rotor: Kubota, model: AG-6512C).

25. Submit the supernatant to a second round of ultracentrifugation at 100,000 $\times g$ for 30 min at 4 °C. (Ultracentrifuge: Beckman Coulter, model: Optima L-100 XP or equivalent models; Rotor: Beckman Coulter, model: Type 70Ti)

26. Resuspend the pellet in ~300 μ l of cold TET. The sample is ready for biochemical and electron microscopic analyses (Figures 1A and 1B).

27. (Optional) Further separation by Superose 6 10/300 column chromatography equilibrated with TET plus 50 mM NaCl can be used to remove large aggregates either made of T4SS or contaminant proteins. Collect the fractions and identify the fractions containing the core complex by gel electrophoresis and transmission electron microscopy (as shown in steps 28 and 29) (Figures 1C and 1D).

Note: As TET solution contains Triton X-100 that disturbs UV monitoring, the UV peaks do not always accord with the presence of the protein complexes. Alternative approaches like sucrose density gradient ultracentrifugation can be applicable for further purification.

28. For biochemical analysis, apply samples on 12.5% SDS-polyacrylamide gel (PAGE) and stain with ready-made Coomassie Brilliant Blue stain solution, CBB stain One (Nacalai).

29. For transmission electron microscopy analysis, apply samples on glow-discharged carbon grids and negatively stained with 2% (w/v) PTA pH 7.0 (Recipe No. 14) or 2% (w/v) uranyl acetate (Recipe No. 15). Take micrographs at an accelerating voltage of 80 kV.

Data analysis

The second round of ultracentrifugation enhances the purity of the isolated complex, and the following column chromatography further removes the background contaminations (Figure 1).

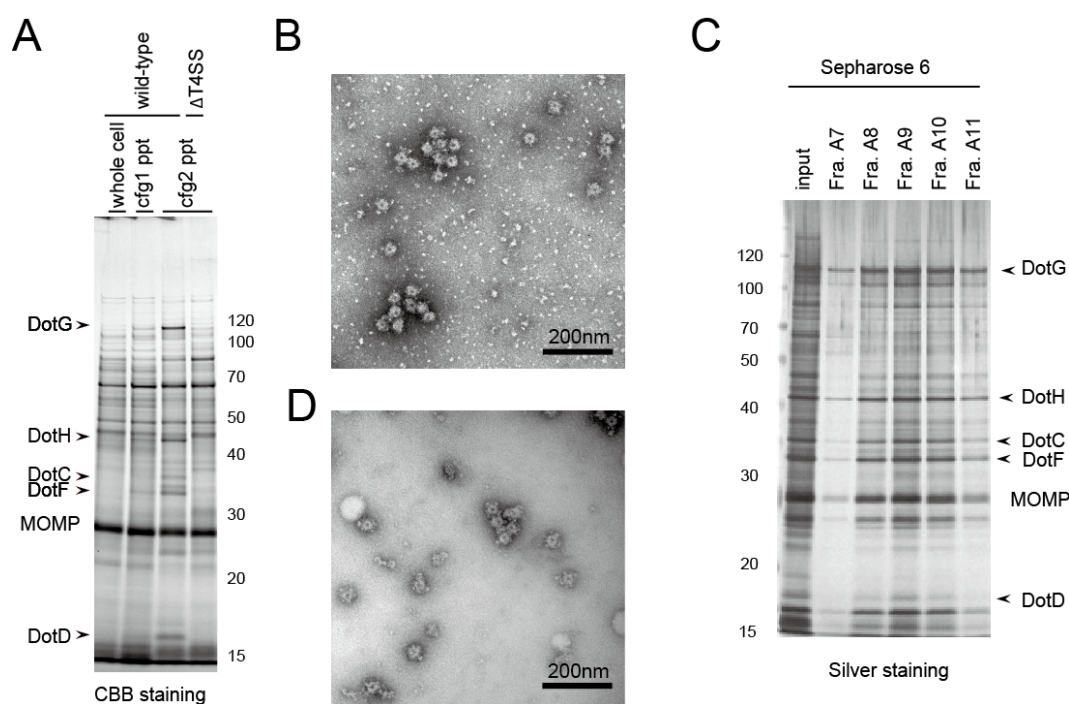


Figure 1. Example of isolated Dot/Icm T4SS core complex. A. SDS-PAGE analysis of the isolated complex in comparison between 1st and 2nd rounds of ultracentrifugation (steps 21 and 25, respectively). Whole cell: Whole cell lysate (step 20); cfg1 ppt: 1st ultracentrifugation pellet (step 22); cfg2 ppt: 2nd ultracentrifugation pellet (step 26). As a negative control, a *L. pneumophila* strain lacking all *dot/icm* genes ($\Delta T4SS$) was also submitted to the isolation procedure (the rightmost lane). B. The electron micrograph of the fraction obtained by the 2nd ultracentrifugation; C. SDS-PAGE analysis of the fractions obtained by a size exclusion column chromatography (step 27); D. The electron micrograph of the fraction A10 of (C). Molecular weight markers are shown in kDa. MOMP: Major Outer membrane Protein. The images are adapted from Kubori *et al.* (2014).

Recipes

1. CYE plate (for 24 plates, 1 L)
 - 10 g ACES
 - 10 g yeast extract
 - Dissolve in ~0.9 L MilliQ water in a 1 L glass beaker, adjust pH by adding 1 N KOH (see Recipe No. 11; ~40 ml) and monitoring with pH meter
 - Bring the volume up to 1 L

Add to a 2 L autoclavable flask containing a magnetic stirrer bar and 2 g activated charcoal and 15 g agar

Mix briefly by stirring

Autoclave the media at 121 °C for 30 min

Cool down at room temperature until the temperature reaches ~60 °C with gentle stirring. During the time, prepare L-cysteine (0.4 g solved in 10 ml sterile MilliQ water) and Fe(NO₃)₃ (0.135 g solved in 10 ml sterile MilliQ water) solutions in sterile 15 ml conical tubes, and filter them with Millex-GP filter unit with 10 ml syringe

Add L-cysteine and Fe(NO₃)₃ solutions (10 ml each) to 1 L medium with stirring

Keep stirring for another 5 min to mix homogeneously

Pour ~40 ml per Petri dish and cool down to solidify

2. AYE medium (1 L)

Same as CYE except that agar is omitted

Note: AYE is not good for use on the same day of preparation. For good result, AYE should be made one day before the Legionella culture. Old media (more than a week after preparation) is not recommended.

3. Tris-Cl solution (pH 8.0)

1 M solution

Adjust pH to 8.0 with HCl

Autoclave at 121 °C for 20 min

4. Sucrose solution

0.5 M sucrose

150 mM Tris-Cl (pH 8.0)

5. PMSF stock solution

100 mM in isopropanol

Storage at -20 °C

6. EDTA stock solution

0.5 M EDTA

Adjust pH to 8.0 with NaOH

Autoclave at 121 °C for 20 min

7. Lysozyme solution

20 mg/ml in sucrose solution

Prepare just before use

8. Triton X-100 stock solution

20% (w/v) solution containing ~2 g of AG501-X8 Resin for deionizing

9. MgSO₄ stock solution

1 M solution

Autoclave at 121 °C for 20 min

10. DNase I stock solution

Dissolve the powder in sterile water to give 10 mg/ml solution

Storage at -20 °C

11. KOH solution

1 N solution

12. NaOH stock solution

1 N solution

13. TET solution

10 mM Tris-Cl (pH 8.0)

1 mM EDTA

0.1% Triton X-100

14. PTA solution

2% (w/v) phosphotungstic acid. Adjust to pH 7.0

15. Uranyl acetate solution

2% (w/v) uranyl acetate

Filter with 0.22 µm ultra-free MC

Acknowledgments

This work has been financially supported by MEXT/JSPS KAKENHI Grants 15H01322 (to TK). The protocol presented here has been adapted from Kubori *et al.* (2014).

References

1. Berger, K. H. and Isberg, R. R. (1993). [Two distinct defects in intracellular growth complemented by a single genetic locus in *Legionella pneumophila*](#). *Mol Microbiol* 7(1): 7-19.
2. Fields, B. S., Benson, R. F. and Besser, R. E. (2002). [Legionella and Legionnaires' disease: 25 years of investigation](#). *Clin Microbiol Rev* 15(3): 506-526.
3. Hubber, A. and Roy, C. R. (2010). [Modulation of host cell function by *Legionella pneumophila* type IV effectors](#). *Annu Rev Cell Dev Biol* 26: 261-283.
4. Kubori, T., Koike, M., Bui, X. T., Higaki, S., Aizawa, S. and Nagai, H. (2014). [Native structure of a type IV secretion system core complex essential for *Legionella* pathogenesis](#). *Proc Natl Acad Sci U S A* 111(32): 11804-11809.
5. Kubori, T., Matsushima, Y., Nakamura, D., Uralil, J., Lara-Tejero, M., Sukhan, A., Galan, J. E. and Aizawa, S. I. (1998). [Supramolecular structure of the *Salmonella typhimurium* type III protein secretion system](#). *Science* 280(5363): 602-605.
6. Kubori, T. and Nagai, H. (2016). [The Type IVB secretion system: an enigmatic chimera](#). *Curr Opin Microbiol* 29: 22-29.
7. Marlovits, T. C., Kubori, T., Sukhan, A., Thomas, D. R., Galan, J. E. and Unger, V. M. (2004).

[Structural insights into the assembly of the type III secretion needle complex.](#) *Science* 306(5698): 1040-1042.

8. Nagai, H. and Kubori, T. (2011). [Type IVB secretion systems of legionella and other Gram-negative bacteria.](#) *Front Microbiol* 2: 136.
9. Vincent, C. D., Friedman, J. R., Jeong, K. C., Buford, E. C., Miller, J. L. and Vogel, J. P. (2006). [Identification of the core transmembrane complex of the *Legionella* Dot/Icm type IV secretion system.](#) *Mol Microbiol* 62(5): 1278-1291.