Unraveling the molecular basis of subunit specificity in
spectrometry

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#	Article	IF	Citations
1	Structural Determinants of Polymerization Reactivity of the P pilus Adaptor Subunit PapF. Structure, 2008, 16, 1724-1731.	1.6	22
3	Structural biology of the chaperoneâ€“usher pathway of pilus biogenesis. Nature Reviews Microbiology, 2009, 7, 765-774.	13.6	298
4	Use of a combined cryo-EM and X-ray crystallography approach to reveal molecular details of bacterial pilus assembly by the chaperone/usher pathway. Current Opinion in Microbiology, 2009, 12, 326-332.	2.3	10
5	Biological â€˜glueâ€™ and â€˜Velcroâ€™: molecular tools for adhesion and biofilm formation in the hairy and gluey bug <i>Pseudomonas aeruginosa</i>. Environmental Microbiology Reports, 2010, 2, 343-358.	1.0	23
6	The differential affinity of the usher for chaperoneâ€“subunit complexes is required for assembly of complete pili. Molecular Microbiology, 2010, 76, 159-172.	1.2	25
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8	Histidine Kinase-Mediated Production and Autoassembly of <i>Porphyromonas gingivalis</i> Fimbriae. Journal of Bacteriology, 2010, 192, 1975-1987.	1.0	34
9	The Outer Membrane Usher Guarantees the Formation of Functional Pili by Selectively Catalyzing Donor-Strand Exchange between Subunits That Are Adjacent in the Mature Pilus. Journal of Molecular Biology, 2010, 396, 1-8.	2.0	22
10	Two-step and one-step secretion mechanisms in Gram-negative bacteria: contrasting the typeÂIV secretion system and the chaperone-usher pathway of pilus biogenesis. Biochemical Journal, 2010, 425, 475-488.	1.7	55
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13	Site-specific methionine oxidation in calmodulin affects structural integrity and interaction with Ca2+/calmodulin-dependent protein kinase II. Journal of Structural Biology, 2011, 174, 187-195.	1.3	33
14	The PprAâ€“PprB twoâ€component system activates CupE, the first nonâ€archetypal <i>Pseudomonas aeruginosa</i> chaperoneâ€“usher pathway system assembling fimbriae. Environmental Microbiology, 2011, 13, 666-683.	1.8	73
15	Function of the usher Nâ€terminus in catalysing pilus assembly. Molecular Microbiology, 2011, 79, 954-967.	1.2	28
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17	Quantitative Analysis of the Interaction Strength and Dynamics of Human IgG4 Half Molecules by Native Mass Spectrometry. Structure, 2011, 19, 1274-1282.	1.6	82
18	Mass spectrometry: come of age for structural and dynamical biology. Current Opinion in Structural Biology, 2011, 21, 641-649.	2.6	240
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22	Chaperoneâ€“usher pathways: diversity and pilus assembly mechanism. Philosophical Transactions of the Royal Society B: Biological Sciences, 2012, 367, 1112-1122.	1.8	118
23	The Structure of the PapD-PapGII Pilin Complex Reveals an Open and Flexible P5 Pocket. Journal of Bacteriology, 2012, 194, 6390-6397.	1.0	18
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31	Enterotoxigenic Escherichia coli CS1 Pilus: Not One Structure but Several. Journal of Bacteriology, 2013, 195, 1357-1359.	1.0	0
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