

Supplementary information

Biocatalytic application and structural elucidation of robust bacterial protein nanocages

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Table S1. Sequences of *E. coli* codon optimized encapsulin and cargo enzyme encoding genes.

Name of the gene	Sequence '3 - '5
CyanoPOX-cargo	ATGGGTAGCAGCCACCACCATCAACCACTCAAGCGGGCTGTCCCCCTGGCTCACATGCTCAGAACCGTACTATTGATGGGA CCGACAACAACCTGTACATAATGTTTGGGGGTCAACAAACCAACATTTAGATCGCGCAGGTCTGCTGCATACGCAGACGGCAT GTCAATGCCAGCCGGAGGTTTCGCGCCCTCCGCACGCGCAGTCAGTAACGGGATTGCCGCGCAGACGGGTTCTATGTTGAATGA CCGTATGTTGAGTGATTGGGTATGGCAGTGGGGTCAATTCCTTGATCATGACCTGGACTTGACGGATGCACGCAGTCCCGCGGA GAGCTTCCCAATCCAGTTCCAATGGGTGACCTTTCTCGACCCCTTCAATACGGGAACCCAGACCATCGGGCTTAGCCGTTACAG CCTATGATCCTGCTACCGGATCTGTTGATGCCGCCAGCAGATGAACCAAATCACAAGTTGGATCGATGCGTCGAATGTATACGG GTCTGATATGACTCGCGCTAACGCTCTTCGCACTATGAGCGGAGGTGTTTACTACTTCAGCTGGGGATCTGCTTCCCTTCAATA CAGGGGGGCTGCCAATGCAGGTGGGACTTCGCCAAGTCTGTTCTTGGCCGAGACGTGCGTTGCAACGAACAGAGTGGCCTTG CTGCCGTACATACTTATTTGTACGCGAGCATAACCGTCTGGCGGACCAAAATTGCTGCAGCGAACCCCGTATGGGAGATGAGGA CATCTACCAGCAAGCTCGCAAGATCGTGGGAGCCAGATGCAAATCATTACGTATAATGAATTTTGGCGCTTATTGGGTTACAG CCGCTCCTTCGCTATGTCGATCGGCTATGATGACTCAATCAATCCAACATCATGAATGAATTTGCGAACGCGTGTACCGCGTC GGTCACACCATGCTGTCGCGACCATTTTGGCTTAGATAACGCTGGGAACGTAATTCACCGGTAAATTTAGCACTGCAGGACG CGTTTTTAAACCAATCGTATCATTAAATGAAGGGGTATCGCCCCATTTTAAAGGCGCTGGCCTCTCAGGCCATGCAGGAGATC GACAACAAGATCGTCGACGACGTTCTGAATTTCTGTTTGGGCCACAGGATCTGGCGGCTTGATCTGGCGAGCTTGAATATTC AACGTGGTCTGACCATGTTTACCGGACTACAATTCGACGCGTGAATGATGGGCTTGACCTCTGTATCTAGCTTCGCTGACATT TCTTCTGACCTGCAGTCCAAGCCGCTTAATGTATTATGGAACCGTCAATGATATTGATCTGTGGTCTGGGGCGCTTGGCG AGGACCACCTGGCTGGCTCAAGCGTAGGCGAGCTTATGACGCTGTCTTGGCGAGCAGTTACTCGTCTGCTGATGGAGACC GTTATTGGTATGAGCGCGACGATTTTTCTGTTAACAACCCATCGTTACTGGCAGAGCTTCAAGCAACTCGTTAAGTGACATTATT CGTCGTAATAGTGACATACGAATATTCAGGACAATGTTTTCTTATCCCTGAACCAGCCACCTGGGGTATTGATGTTCCGGGGC TGCTTTTTGCGTAACGTCGCTCTCCACTCCCTTCCGACAGTGAGCCGGACCGTGAGATCCCGGCCGATGACGGCTCCTTGG GCATCGGTTCTTGAAGGTACACGCTCA
PTDH-mFMO-cargo	ATGGGAAGTTCACATCACCATCATCACACAGTTCGGGCTTGGTCCCTCGTGGGTCTCACATGTTGCCCAAATTAGTAATCACGCA CCGTGTTACGAGGAAATTTCTCAGCTGCTGGCTCCTCACTGCGAATTGATACGAACCAAGACTGATTCAACCTTAACACGTGAGG AGATTTTCGTCGCTGTCGTGATGCTCAGGCTATGATGGCTTCATGCCGATCGCGTTGACGCCGATTTTCTCAAGCTTGTCCTC GAATTACGCGTTATTGGATGCGCACTGAAAGGCTTTGACAATTCGACGTAGATGCCTGCACAGCGCGTGGAGTTTGGCTGACTT TTGTACCGGACCTGTTAACTGTCCTACAGCAGAGCTTGCGATCGGCTGGCGGTGGGATTGGGGCGTCATTTGCGTGGCGCG ACGCTTTGTCGCTCAGGTAAATTTTCGCGCTGGCAGCGCGCTTTTACGGTACGGGATTAGACAACGCCACAGTCGGGTTTCT TGGAATGGGTGCCATTGGCTTAGCTATGGCCGATCGTTTGCAAGGGTGGGGGGCAACTTTGCAATACCATGCCCGCAAGGCATT GGACACGCAAACTGAGCAACGTTTAGGGCTTCGTCAAGTAGCGTGCTCCGAGCTTTTCGCGAGTAGTGATTTTATTCTTCTGCTC TTCGCTTAATGCAACACGCTGCACCTTGTAACGCGGAACGCTTGCTCTTGACGTCCAGGCGCTTACTGGTCAACCCGTCG CGTGGCAGCGTCTGATGAGGCGGCTGTGCTGGTCTGCTGGAACGCGGACAACCTGGGTGGCTATGACGCGGATGTTTTGAG ATGGAGGATTGGGCCGTCGCGACCGTCTCAACAGATTGACCCGGCACTTCTGCACACCCGAACACGTTGTTTACTCCACAT CGGGAGTGCCTCCGCGAGTTCGCTTAGAGATCGAACGCTGCGCCGCGCAAAACATTCTGACGGCCTGGCTGGGGAACGCC GATTAATGCTGTGAATCGCCTTCAAAAGCAACCCGGCAGCGGATTACGTTCTGCTGCTGGCATGGCGACGCGTATTGCAATCT TGGGTGCTGGCCATCAGGGATGGCTCAATTGCGCGCAATTCGAATCAGCGCAGGAGAAGGGGGCAGAAATCCAGAACTGGTA TGTTTTGAAAAACAGGCGGATTGGGGAGGTCAAGTGAACACTTACCTTGGCGACGGGTTTAGACGAGAACGGCGAGCCGTACA CTCTCCATGTACCGCTATCTGGTCAAATGGCCAAAGGAGTGTCTTGAGTTTCCGATTACACATTGATGAGCACTTTGGCA AACCAGTTCCTCGTATCCGCCACGCGAGGTACTGTGGGATTATACAAAGGGCGTGTAGAGAAAGCCGAGTCCGTAATATA TTCGTTTCAATACAGCGGTGCGCATGTGCAATCAATGAAGACTCCAGACCTTTACTGTACGGTGCAAGATCATACGACAGAC ACTATCTATAGCGAGGAGTTTGATTATGTGGTCTGCTGTACCGGACATTTTCGACGCCATATGTTCTGAATTTGAAGGGTTCGA GAAGTTCGAGGCCGCATCTTGATGCAACGACTTTCTGACGCTTTGGAATTCAAAGACAAACCGTGCTGTTAGTTGGCAGT TCCTACTCAGCTGAGGATATCGGAAGCCAGTCTATAAGTATGGTGCCAAGAAATTAATTTTATGTTATCGTACGGCGCTATGG GATATAAGTGGCCGAAACTGGGATGAGCGCCCACTTAGTCCGTGTCGATACTGAAATGCATATTTGCTGACGGCAGCTC GGAGAAAGTCGACGCCATCATCTTGTGACGCGGTATATTATCATCTTCCATTTCTTAACGATGATCTGCTTTAGTAACAAATA ATCGCTTTGGCCGCTTAACCTGTACAAAGGGGTGTATGGGAAGACAATCCCAATTTCTACATTGGAATGCAAGACCAATG GTATTCATTTAATGTTGATGCCAGGCTGGTACGCGCGCAGCTTATTATGGGCGTTTACCTTGGCTCAAAAGAAAGAG ATGAAGGCCGACTCTATGGCTTGGCGGAAAAAGAGCTTACTTTGGTCACTGCTGAGGAGATGTATACATATCAAGGGGACTAT ATCCAAATTTAATTGATATGACTGACTACCCAGTTTGTATTTCCGGCCACGAATAAGACCTTCTTAGAGTGGAAGCATCATAA GAAAGAAAATATCATGACATTCGCTGACCATAGTTACCGTCTCCCTATGACCGGTACTATGGCACCAACATCATACACCTGGA TTGATGCTTTAGACGACTCATTAGAAGCGTACTTATCGGATAAGTCGGAATTCGGTGGCAAGGAAGCCCAACCCCTTACC CGACTCGGAGCCGACCGTGAGATTCGGCAGATGACGGTTCTCTGGGAATTGGGTGTTAAAGGGACCCGTTCT

DendroEnc	ATGGACTTTCTTGATCGTAACGCGGCCCTTTAAATGAAGGTGAGTGGCAGCGCATTGACGAGGCAGTCGTCTCGACAGCGCGTC GTACACTTGTGCTCGCCGCATCATTGACGTTTTAGGGCCCTTAGGCTCGGGTGTTTACAGCATCCCATACTCCGTTTTCTCAGGTA AGTCACCTACGGGAATTGACATGGTCGGAGAGAACGAAGATTTGTGGTAGAGGCTTCGCGCCGCGCTACGATCAATCTGCCGA TCTTGTAACAAGGACTTTAAGATTATGTGGCGCGACGTCGAGGCAGACCGTCACCTGGGGCTGCCATTGACGTCAGCACAGCAGC AGTAGCAGGGAACCTTTGTCGCGGTACAGGAAGATCACCTGATCTTCAACGGTAATGTAGAGCTTGGGCATGATGGCCTTTTACG GTGAAAGACCGCCAGACTGTTGCCATTTAGATTGGAACGAAACGGGTGCAGCGCTTGCTGACGTGGTGAAGGCGGTAGGAGC TTTGTCTCAAGCAGGTCACTACGGACCTTACGCTATGGTCGTTTCCCGGTTTTGTTGGGCGTATGATTGCGGTATTTGGGAATA CGGGTATGCTGGAATTAGACCAAGTTAAGGCATTGATTACTGGCGGGGTGCATTATAGCAACGTATTGGAGGAAGTAAAGCTG TTGTCGTCGCCACCGGGTCACAGAACCTTAATCTGGCATTAGGACAAGACATGGTCACAGCATACATGGGACCGACTTCTATGAA CCACGTTTTTTCGCGTCTTAGAGACCGCCGATTATTAGTGCGCCGCCGACGCCATCTGCACCATCGAA
ArthroEnc	ATGAATAATCTGTACCGCGAATTAGCACCAGTCACAGACGAGGCCTGGGGTGAAATTGAACAAGAAGCTACCCGCACTTTCAAG CGTCACATTGCAGGCCGCCGCGTGGTAGACTGTACGGGACCTACTGGTGCCACTACCGCTGCAGTATCTACGGGTATCTTCTGG ACGTAACATCACCTGGAGATGGGGTTATTGCTCACTTGCAGGATTCTCGCCCCCTTGTCGTTTTGCGTGTCTTTTACCATTACAC GTGATGCTATTGATGATGTAGAGCGCGGGAGCCAAGACTCCGATTGGGATCCGGTTAAGGATGCTGCCAAAAAGTTAGCGTTTG CCGAAGATCGTGCCATCTTCGAGGGGTATGCAGCTGCGAATATTCAAGGGATTTCGCGCAGGTAGTAGCAATCCAGGTCTTGCACT TCCTGACGACCCCCGTGGTTATCCAGACGTTATCGCTCAGGCTCTTAGTGAATTGCGCCTTGCAAGGAGTTGATGGACCCTACAGC GTCCTTCTGAGCGCCGACGCATACACTAAGGTCAAGTGAAGACGACGAGCACGGGTACCCACTGCGCGAACACCTTAATCGCGTA GTTGACGGTGACATTATTTGGGCACCTGCAATCGATGGTGCCTTCGTTCTTTCAACGCGCGGTGGAGACTTCGACTTACAATTAG GTACGGATGTATCTATCGGTTATTCATCTCATGATGCCGACACGGTGCAGTTGTATCTGCAGGAGACATTTACCTTTTTGAATTAC ACAGCTGAAGCAAGTGTGGCGCTGTCTGCA

Table S2. Primers used for DendroEnc N198W mutant generation.

Primer	Sequence '3 - '5
DendroEnc N198W Fw	TCGGATGGACTGGCATGCTGGAAGTGGAT
DendroEnc N198W Rv	GCATGCCAGTCCATCCGAAGACGCGAATCATGCG

Table S3. Cryo-EM data collection, refinement, and validation statistics.

	ArthroEnc	WT DendroEnc	N198W DendroEnc
Microscope	Talos Arctica		
Voltage (kV)	200		
Defocus range (μm)	-0.5 to -2.5		
Camera	Gatan K3 Superresolution mode		
Pixel size (Å)	1.04		
Total electron dose (e/Å ²)	45		
Exposure time (s)	3.5		
Frames per movie	50		
Number of movies	719	4498	7874
3D Refinement			
Number of particles	15677	4549	3997
Symmetry applied	Icosahedral (I)		
Final map pixel size	1.04	1.59	1.59
Final resolution (Å)	2.9	3.4	3.6
Atomic model statistics*			
Non-hydrogen atoms	2017	8436	8464
Number of chains	1	4	4
Molprobity score	0.89	1.29	1.46
Rama distribution Z-score	-1.14 ± 0.50	-1.96 ± 0.22	-1.73 ± 0.23
Clashscore	1.51	2.13	3.43
Bond RMSD (Å)	0.003	0.002	0.003
Angle RMSD (°)	0.573	0.608	0.646
Poor rotamers (%)	0	0	0
Favored rotamers (%)	100	99.78	99.78
Ramachandran favored (%)	98.1	95.56	95.29
Ramachandran allowed (%)	1.9	4.44	4.71
Ramachandran outliers (%)	0	0	0

* Statistics given for the deposited PDB model with one asymmetric unit.

Table S4. Specific activity raw data of DendroEnc CyanoPOX constructs in the presence of ABTS and guaiacol.

Enzyme	DendroEnc CyanoPOX	DendroEnc N198W CyanoPOX	DendroEnc CyanoPOX
Substrate	ABTS		guaiacol
Concentration (mM)	10		
mAbs/t (mAbs /min)	714.140	132.675	15.540
	729.740	138.225	15.990
Abs/t (Abs/min)	0.7141	0.1327	0.0155
	0.7297	0.1382	0.0160
$\frac{\Delta Abs}{\Delta t}$ $\varepsilon * l$ (rate of substrate conversion (mM/min))	0.0198	0.0037	0.0006
	0.0203	0.0038	0.0006
U/mg	0.0198	0.0037	0.0006
	0.0203	0.0038	0.0006
average U/mg	0.0201	0.0038	0.0006
standard deviation U/mg	0.00031	0.00011	0.00001

$$\text{Specific activity} \left(\frac{U}{mg} \right) = \frac{\left[\left(\left(\frac{\Delta Abs / \Delta t}{\varepsilon * l} \right) \right) \times \text{reaction volume (mL)} \right]}{\text{protein concentration} \left(\frac{mg}{mL} \right)}$$

Equation S1. Equation used to calculate the specific activity of DendroEnc CyanoPOX constructs in the presence of ABTS and guaiacol. ε (molar extinction coefficient ($\text{mM}^{-1} \text{cm}^{-1}$), l (path length (cm)).

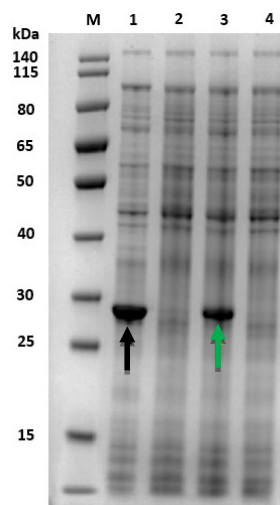


Figure S1. SDS-PAGE analysis of total *E. coli* lysates demonstrating overexpression of encapsulin proteins. Lane M: molecular weight marker. Lane 1: induced expression of ArthroEnc. Lane 2: uninduced ArthroEnc (negative control). Lane 3: induced expression of DendroEnc. Lane 4: uninduced DendroEnc (negative control). The encapsulin bands are indicated by black (ArthroEnc) and green (DendroEnc) arrows.

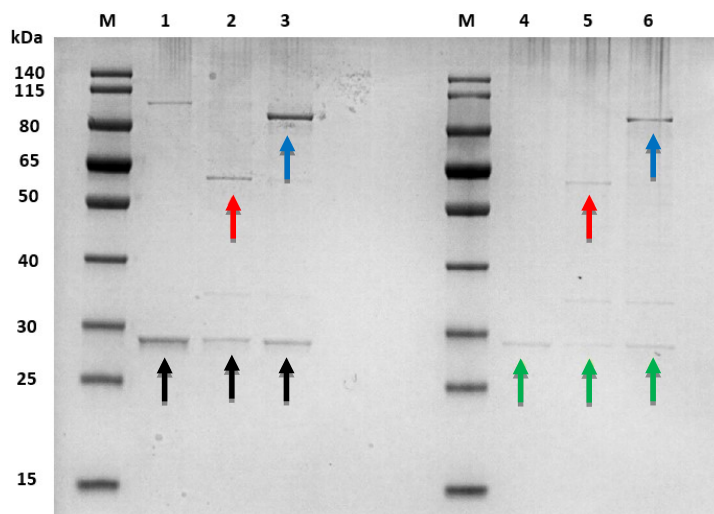


Figure S2. SDS-PAGE gel of purified DendroEnc and ArthroEnc constructs after SEC. Lane M: molecular weight marker. Lane 1: DendroEnc (black arrow). Lane 2: DendroEnc with encapsulated CyanoPOX. CyanoPOX is shown with the red arrow. Lane 3: DendroEnc with encapsulated PTDH-mFMO. PTDH-mFMO is shown by the blue arrow. Lane 4: ArthroEnc (green arrow). Lane 5: ArthroEnc with encapsulated CyanoPOX. CyanoPOX is shown with the red arrow. Lane 6: ArthroEnc with encapsulated PTDH-mFMO. PTDH-mFMO is shown by the blue arrow.

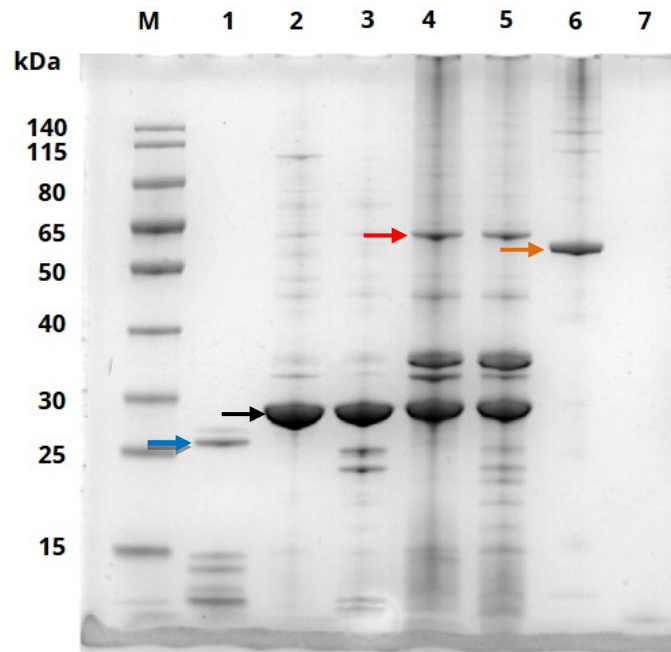


Figure S3. SDS-PAGE gel showing proteolytic stability of empty DendroEnc, CyanoPOX-loaded DendroEnc, and free CyanoPOX after incubation with chymotrypsin. Lane M: molecular weight marker. Lane 1: chymotrypsin, 20x amount compared with the digestions shown in lane 3, 5 and 7. (blue arrow). Lane 2: DendroEnc (black arrow: encapsulin protein). Lane 3: DendroEnc incubated with chymotrypsin. Lane 4: DendroEnc with encapsulated CyanoPOX (red arrow: CyanoPOX with encapsulin targeting sequence). Lane 5: DendroEnc with encapsulated CyanoPOX incubated with chymotrypsin. Lane 6: CyanoPOX (red arrow: CyanoPOX). Lane 7: CyanoPOX incubated with chymotrypsin.

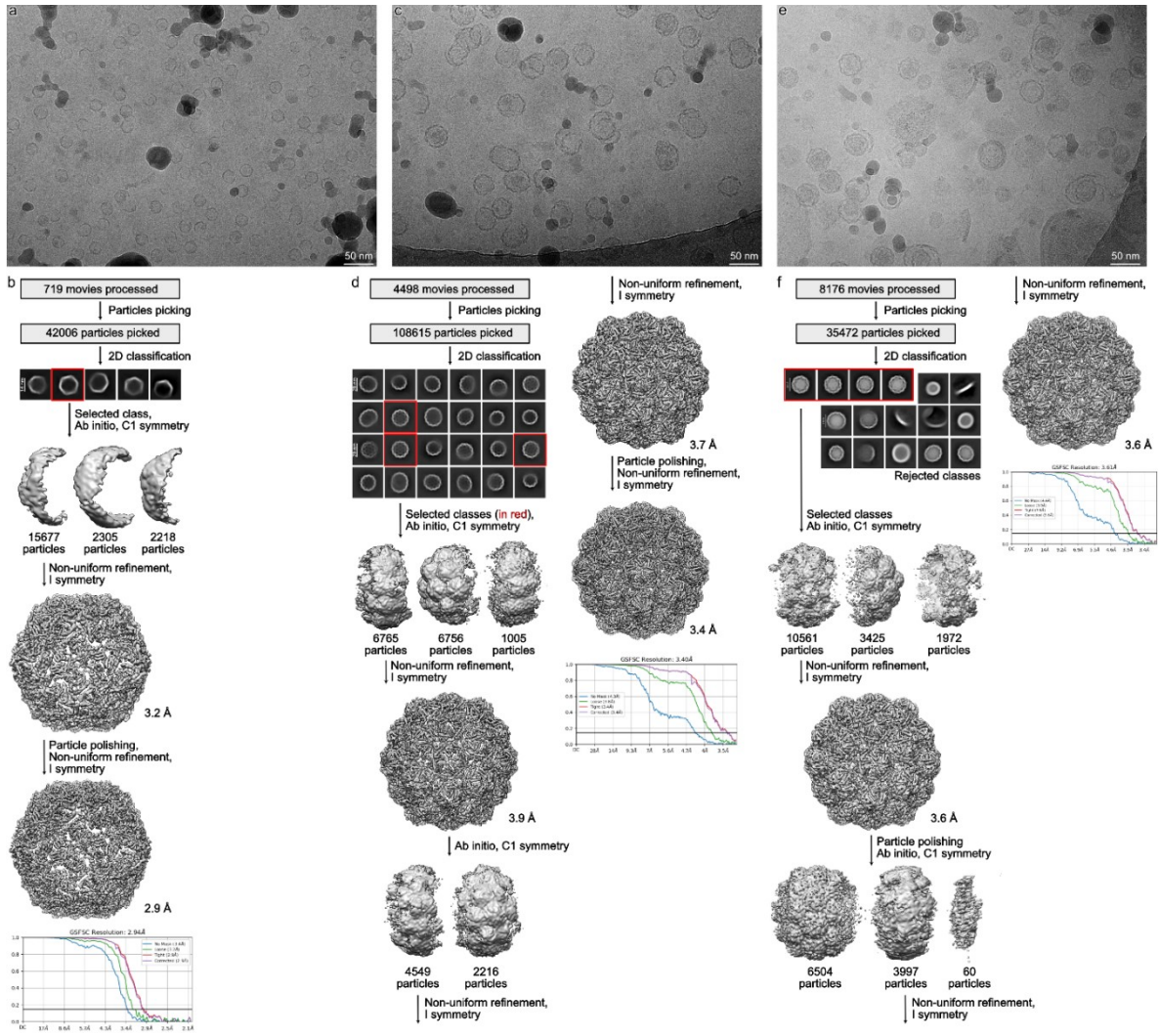


Figure S4. Cryo-EM data processing. (a), (c) and (e) Exemplary cryo-EM micrographs of ArthroEnc, WT DendroEnc and N198W DendroEnc, respectively. (b), (d) and (f) Processing overview and fourier shell correlation (FSC) curves of final reconstructions for ArthroEnc, WT DendroEnc and N198W DendroEnc, respectively.