

Summary of integrative structure determination of Integrative structure and functional anatomy of eight spokes of a nuclear pore complex (PDB ID: 8ZZC, PDB-Dev ID: PDBDEV_0000012)

1. Model Composition	
	<ul style="list-style-type: none"> - Nup57: Chain LW (200 residues) - Nup159: Chain GV (600 residues) - Nup116: Chain RC (1113 residues) - Pom34: Chain OJ (299 residues) - Sec13: Chain CE (297 residues) - Nsp1: Chain JF (823 residues) - Nup60: Chain SL (100 residues) - Nsp1: Chain HG (823 residues) - Nsp1: Chain KM (823 residues) - Nup42: Chain RD (430 residues) - Nup49: Chain JX (200 residues) - Nup42: Chain BI (430 residues) - Nup57: Chain JB (200 residues) - Pom34: Chain AX (299 residues) - Nsp1: Chain JW (600 residues) - Sec13: Chain G (297 residues) - Sec13: Chain DN (297 residues) - Nup192: Chain NK (1683 residues) - Nic96: Chain IO (839 residues) - Nup49: Chain KO (472 residues) - Sec13: Chain DU (297 residues) - Nup100: Chain RY (540 residues) - Nsp1: Chain AJ (600 residues) - Dyn2: Chain GB (92 residues) - Seh1: Chain CK (349 residues) - Nup145: Chain TJ (200 residues) - Nup85: Chain FF (744 residues) - Nup159: Chain GG (1460 residues) - Ndc1: Chain PH (655 residues) - Nup85: Chain DI (744 residues) - Nup145: Chain SP (1317 residues) - Ndc1: Chain PC (655 residues) - Nup188: Chain MD (1655 residues) - Nup159: Chain FQ (1460 residues) - Nup192: Chain MI (1683 residues) - Nup116: Chain QV (1113 residues) - Nup57: Chain JS (200 residues) - Nup49: Chain IZ (200 residues) - Nup57: Chain ID (200 residues) - Nup100: Chain SD (959 residues) - Nup60: Chain TP (539 residues) - Mlp2: Chain TV (1679 residues) - Mlp2: Chain UB (1679 residues) - Nup192: Chain NW (1683 residues) - Nsp1: Chain AF (600 residues) - Nup133: Chain FH (1157 residues) - Nup60: Chain BN (100 residues) - Nup100: Chain BE (540 residues) - Nup192: Chain AP (1683 residues) - Dyn2: Chain O (92 residues) - Nup57: Chain LO (200 residues) - Nup145: Chain TN (1317 residues) - Nup133: Chain FA (1157 residues) - Nup170: Chain AN (1502 residues) - Nup49: Chain JI (472 residues)

- Nup57: Chain IT (200 residues)
- Ndc1: Chain OD (655 residues)
- Nup116: Chain RT (1113 residues)
- Nup120: Chain CH (1037 residues)
- Seh1: Chain CR (349 residues)
- Nup59: Chain OM (528 residues)
- Nup120: Chain EZ (1037 residues)
- Nup192: Chain MA (1683 residues)
- Ndc1: Chain PR (655 residues)
- Nup116: Chain RH (1113 residues)
- Nup57: Chain JZ (200 residues)
- Nsp1: Chain LR (600 residues)
- Nic96: Chain LH (839 residues)
- Nup188: Chain MP (1655 residues)
- Nup170: Chain NU (1502 residues)
- Nup84: Chain EJ (726 residues)
- Nup157: Chain AQ (1391 residues)
- Sec13: Chain BX (297 residues)
- Nic96: Chain JU (839 residues)
- Nup82: Chain GL (713 residues)
- Nsp1: Chain LK (600 residues)
- Nsp1: Chain JO (600 residues)
- Nup100: Chain RF (540 residues)
- Nup159: Chain FP (1460 residues)
- Nup59: Chain PG (528 residues)
- Nup49: Chain IJ (200 residues)
- Nup82: Chain FV (713 residues)
- Nsp1: Chain GY (600 residues)
- Nup57: Chain KA (200 residues)
- Nup170: Chain NQ (1502 residues)
- Sec13: Chain CS (297 residues)
- Nup57: Chain KH (200 residues)
- Nup145: Chain TI (200 residues)
- Nup49: Chain IB (200 residues)
- Nup53: Chain OB (475 residues)
- Nup85: Chain CN (744 residues)
- Ndc1: Chain QL (655 residues)
- Nsp1: Chain IQ (823 residues)
- Nup60: Chain BN (539 residues)
- Nsp1: Chain AB (600 residues)
- Nup57: Chain KY (541 residues)
- Nup145c: Chain BV (712 residues)
- Nup116: Chain SA (1113 residues)
- Nup116: Chain RC (740 residues)
- Nup192: Chain NO (1683 residues)
- Nup57: Chain LN (541 residues)
- Dyn2: Chain GJ (92 residues)
- Nsp1: Chain LK (823 residues)
- Nsp1: Chain FZ (823 residues)
- Nup59: Chain OW (528 residues)
- Nsp1: Chain HO (823 residues)
- Nup145: Chain TJ (1317 residues)
- Seh1: Chain BW (349 residues)
- Nup57: Chain IU (200 residues)
- Nup116: Chain RZ (740 residues)
- Nup188: Chain MH (1655 residues)
- Dyn2: Chain HA (92 residues)
- Nsp1: Chain LS (823 residues)
- Sec13: Chain EP (297 residues)
- Nup120: Chain C (1037 residues)
- Pom152: Chain QI (1337 residues)
- Nsp1: Chain LJ (600 residues)
- Nsp1: Chain U (600 residues)
- Nup133: Chain CW (1157 residues)

- Nup57: Chain IE (541 residues)
- Pom152: Chain OZ (1337 residues)
- Nup49: Chain IB (472 residues)
- Pom152: Chain PO (1337 residues)
- Nup1: Chain BM (1076 residues)
- Nup188: Chain NJ (1655 residues)
- Nup49: Chain Y (472 residues)
- Nup49: Chain KW (472 residues)
- Nup159: Chain S (1460 residues)
- Nup100: Chain RF (959 residues)
- Nic96: Chain KR (839 residues)
- Nsp1: Chain JN (600 residues)
- Nup170: Chain NA (1502 residues)
- Nup188: Chain MT (1655 residues)
- Nup57: Chain KX (200 residues)
- Nup49: Chain HT (472 residues)
- Nup49: Chain JA (200 residues)
- Nup49: Chain LU (472 residues)
- Nup85: Chain DB (744 residues)
- Nup133: Chain DK (1157 residues)
- Nup1: Chain TR (1076 residues)
- Nup84: Chain DV (726 residues)
- Nup145: Chain ST (1317 residues)
- Nic96: Chain AA (839 residues)
- Nsp1: Chain IX (823 residues)
- Ndc1: Chain OI (655 residues)
- Nup120: Chain DX (1037 residues)
- Nup120: Chain FG (1037 residues)
- Nup49: Chain IK (200 residues)
- Nup82: Chain GU (713 residues)
- Nsp1: Chain GH (823 residues)
- Nup170: Chain NY (1502 residues)
- Nup49: Chain KF (200 residues)
- Nup192: Chain MM (1683 residues)
- Nup85: Chain DP (744 residues)
- Nup120: Chain CA (1037 residues)
- Nsp1: Chain AF (823 residues)
- Seh1: Chain DT (349 residues)
- Dyn2: Chain HI (92 residues)
- Nsp1: Chain X (823 residues)
- Nic96: Chain LP (839 residues)
- Nup57: Chain JJ (541 residues)
- Nup59: Chain PL (528 residues)
- Nup42: Chain RJ (430 residues)
- Nup159: Chain HE (600 residues)
- Nup57: Chain AH (200 residues)
- Gle1: Chain RQ (538 residues)
- Nup159: Chain GO (600 residues)
- Nup85: Chain ER (744 residues)
- Nup49: Chain JH (200 residues)
- Nup49: Chain AC (472 residues)
- Nup116: Chain RU (1113 residues)
- Nup192: Chain AT (1683 residues)
- Nup57: Chain AD (541 residues)
- Nup133: Chain EF (1157 residues)
- Mlp2: Chain TT (1679 residues)
- Nsp1: Chain HS (600 residues)
- Nup49: Chain KO (200 residues)
- Dyn2: Chain FM (92 residues)
- Ndc1: Chain PM (655 residues)
- Nup49: Chain JI (200 residues)
- Nup170: Chain MW (1502 residues)
- Seh1: Chain CY (349 residues)
- Nup49: Chain IJ (472 residues)

- Nsp1: Chain KD (600 residues)
- Nup60: Chain TG (539 residues)
- Nup49: Chain AG (200 residues)
- Nup159: Chain GF (1460 residues)
- Nup84: Chain CM (726 residues)
- Pom152: Chain PJ (1337 residues)
- Nup116: Chain SG (740 residues)
- Nsp1: Chain V (600 residues)
- Nup49: Chain IK (472 residues)
- Ndc1: Chain OX (655 residues)
- Seh1: Chain CD (349 residues)
- Nic96: Chain HX (839 residues)
- Nsp1: Chain V (823 residues)
- Nup133: Chain BU (1157 residues)
- Nup53: Chain OL (475 residues)
- Nup57: Chain JS (541 residues)
- Nup82: Chain GE (713 residues)
- Nup85: Chain BZ (744 residues)
- Nup57: Chain IE (200 residues)
- Nup60: Chain SW (539 residues)
- Nup145c: Chain DE (712 residues)
- Nup159: Chain HD (1460 residues)
- Nup57: Chain KQ (541 residues)
- Mlp1: Chain UC (1875 residues)
- Nsp1: Chain IQ (600 residues)
- Nup133: Chain CI (1157 residues)
- Nup60: Chain TB (100 residues)
- Nup49: Chain AC (200 residues)
- Nup42: Chain QX (430 residues)
- Nup57: Chain JK (200 residues)
- Nsp1: Chain JV (600 residues)
- Nup157: Chain NT (1391 residues)
- Nup84: Chain DH (726 residues)
- Nup49: Chain JH (472 residues)
- Nic96: Chain HQ (839 residues)
- Nup57: Chain JK (541 residues)
- Nup192: Chain MY (1683 residues)
- Nic96: Chain IG (839 residues)
- Pom34: Chain OY (299 residues)
- Nup100: Chain BF (540 residues)
- Nup85: Chain DW (744 residues)
- Nup49: Chain JP (200 residues)
- Nsp1: Chain LB (823 residues)
- Nup57: Chain AD (200 residues)
- Nup145: Chain SY (1317 residues)
- Dyn2: Chain GC (92 residues)
- Nup57: Chain LF (541 residues)
- Nup100: Chain QZ (540 residues)
- Nup145: Chain SK (200 residues)
- Mlp1: Chain TU (1875 residues)
- Nup170: Chain MC (1502 residues)
- Nup53: Chain QJ (475 residues)
- Nup49: Chain KN (200 residues)
- Nup116: Chain BG (1113 residues)
- Nup60: Chain SQ (100 residues)
- Nup53: Chain PP (475 residues)
- Nup84: Chain DA (726 residues)
- Nup85: Chain CG (744 residues)
- Nup57: Chain Z (541 residues)
- Nup120: Chain EL (1037 residues)
- Nsp1: Chain IY (823 residues)
- Nup116: Chain RI (740 residues)
- Nup100: Chain RL (959 residues)
- Nup1: Chain SN (698 residues)

- Nup145c: Chain CC (712 residues)
- Nup100: Chain RM (959 residues)
- Nup57: Chain LG (200 residues)
- Nup100: Chain RX (959 residues)
- Nup57: Chain KA (541 residues)
- Nup42: Chain RV (430 residues)
- Nic96: Chain LQ (839 residues)
- Pom34: Chain QR (299 residues)
- Nup100: Chain RA (959 residues)
- Nup57: Chain AL (541 residues)
- Nup157: Chain NL (1391 residues)
- Nup49: Chain IS (472 residues)
- Nup100: Chain RS (540 residues)
- Nsp1: Chain FR (823 residues)
- Nup84: Chain H (726 residues)
- Nup120: Chain BT (1037 residues)
- Nup100: Chain RR (540 residues)
- Nup49: Chain LE (472 residues)
- Nup1: Chain TH (698 residues)
- Gle1: Chain RK (538 residues)
- Nsp1: Chain FR (600 residues)
- Nup82: Chain GT (713 residues)
- Nup145: Chain SJ (1317 residues)
- Nup82: Chain FO (713 residues)
- Nup157: Chain LX (1391 residues)
- Mlp2: Chain TX (1679 residues)
- Nup120: Chain CO (1037 residues)
- Nup116: Chain RH (740 residues)
- Nup100: Chain SD (540 residues)
- Nup145: Chain SY (200 residues)
- Nsp1: Chain IA (823 residues)
- Sec13: Chain EB (297 residues)
- Nup60: Chain SR (100 residues)
- Nup133: Chain EM (1157 residues)
- Ndc1: Chain BB (655 residues)
- Nsp1: Chain LR (823 residues)
- Pom34: Chain QH (299 residues)
- Nup59: Chain QF (528 residues)
- Ndc1: Chain QG (655 residues)
- Nup60: Chain TA (539 residues)
- Nup116: Chain RN (740 residues)
- Nsp1: Chain KL (823 residues)
- Nup57: Chain IL (541 residues)
- Nup159: Chain HD (600 residues)
- Nup116: Chain QV (740 residues)
- Nup170: Chain AR (1502 residues)
- Nup42: Chain SH (430 residues)
- Nup49: Chain HU (472 residues)
- Gle1: Chain QY (538 residues)
- Nup145c: Chain FB (712 residues)
- Nup60: Chain TF (100 residues)
- Nsp1: Chain KD (823 residues)
- Nup116: Chain RI (1113 residues)
- Nsp1: Chain HG (600 residues)
- Nup57: Chain KX (541 residues)
- Nup159: Chain GW (1460 residues)
- Nup59: Chain PQ (528 residues)
- Seh1: Chain M (349 residues)
- Nup57: Chain LG (541 residues)
- Nup192: Chain OA (1683 residues)
- Nsp1: Chain HO (600 residues)
- Sec13: Chain N (297 residues)
- Nup159: Chain HM (600 residues)
- Nup49: Chain JX (472 residues)

- Nup57: Chain IL (200 residues)
- Nup145: Chain TE (1317 residues)
- Nup57: Chain ID (541 residues)
- Nup57: Chain IM (200 residues)
- Nsp1: Chain IP (823 residues)
- Nup145c: Chain DL (712 residues)
- Nup100: Chain QZ (959 residues)
- Nup188: Chain AO (1655 residues)
- Nup159: Chain HL (600 residues)
- Nic96: Chain IV (839 residues)
- Mlp2: Chain BQ (1679 residues)
- Nup57: Chain KP (541 residues)
- Nup157: Chain NP (1391 residues)
- Nup188: Chain ML (1655 residues)
- Nup60: Chain TB (539 residues)
- Nsp1: Chain JO (823 residues)
- Nup49: Chain LL (200 residues)
- Nup116: Chain SA (740 residues)
- Nup120: Chain DJ (1037 residues)
- Nup157: Chain MR (1391 residues)
- Nup49: Chain AG (472 residues)
- Pom152: Chain QS (1337 residues)
- Nup159: Chain GV (1460 residues)
- Mlp1: Chain BP (1875 residues)
- Nup145: Chain BK (1317 residues)
- Nup133: Chain DD (1157 residues)
- Nsp1: Chain JW (823 residues)
- Nup157: Chain MV (1391 residues)
- Nup188: Chain NZ (1655 residues)
- Nup188: Chain NF (1655 residues)
- Nsp1: Chain JV (823 residues)
- Nup49: Chain IR (472 residues)
- Nup60: Chain TF (539 residues)
- Nup57: Chain KH (541 residues)
- Nsp1: Chain FS (600 residues)
- Nup57: Chain LW (541 residues)
- Nsp1: Chain HF (600 residues)
- Nsp1: Chain KM (600 residues)
- Nup159: Chain GO (1460 residues)
- Nup157: Chain NX (1391 residues)
- Nup60: Chain TK (100 residues)
- Sec13: Chain CZ (297 residues)
- Nsp1: Chain GH (600 residues)
- Nup159: Chain GN (600 residues)
- Gle1: Chain SC (538 residues)
- Nup188: Chain MX (1655 residues)
- Nup57: Chain JR (200 residues)
- Nsp1: Chain IH (823 residues)
- Nup57: Chain HW (541 residues)
- Ndc1: Chain PW (655 residues)
- Nup159: Chain GG (600 residues)
- Nup57: Chain KI (200 residues)
- Nup145: Chain TN (200 residues)
- Nup159: Chain HL (1460 residues)
- Nup49: Chain LU (200 residues)
- Pom34: Chain QM (299 residues)
- Nup145: Chain SJ (200 residues)
- Nup82: Chain HB (713 residues)
- Nup49: Chain HU (200 residues)
- Nup82: Chain GD (713 residues)
- Dyn2: Chain GK (92 residues)
- Nsp1: Chain GX (823 residues)
- Pom34: Chain PI (299 residues)
- Nup57: Chain LV (541 residues)

[Entry composition](#)

- Nsp1: Chain LB (600 residues)
- Nup60: Chain SV (100 residues)
- Nup49: Chain IC (472 residues)
- Nup159: Chain GW (600 residues)
- Nup49: Chain JQ (200 residues)
- Mlp2: Chain TZ (1679 residues)
- Nup82: Chain GM (713 residues)
- Nup100: Chain RY (959 residues)
- Nup159: Chain HE (1460 residues)
- Nup192: Chain NC (1683 residues)
- Nsp1: Chain GQ (600 residues)
- Nic96: Chain IW (839 residues)
- Nup116: Chain RO (740 residues)
- Nup49: Chain JY (472 residues)
- Nsp1: Chain IP (600 residues)
- Nup157: Chain MN (1391 residues)
- Nup57: Chain HW (200 residues)
- Nup1: Chain SS (698 residues)
- Nup192: Chain MQ (1683 residues)
- Nup49: Chain LM (472 residues)
- Nup192: Chain MU (1683 residues)
- Nup116: Chain QW (1113 residues)
- Nup53: Chain OV (475 residues)
- Nup116: Chain SF (1113 residues)
- Nup53: Chain PK (475 residues)
- Nup188: Chain LZ (1655 residues)
- Nup116: Chain QW (740 residues)
- Nsp1: Chain II (600 residues)
- Nup59: Chain PB (528 residues)
- Nsp1: Chain LC (600 residues)
- Nsp1: Chain II (823 residues)
- Mlp2: Chain UD (1679 residues)
- Ndc1: Chain ON (655 residues)
- Seh1: Chain FJ (349 residues)
- Nup159: Chain T (600 residues)
- Nup133: Chain ET (1157 residues)
- Nic96: Chain W (839 residues)
- Nup60: Chain SM (539 residues)
- Nup49: Chain KV (200 residues)
- Nic96: Chain JE (839 residues)
- Nup59: Chain QK (528 residues)
- Pom152: Chain AY (1337 residues)
- Ndc1: Chain QB (655 residues)
- Nsp1: Chain KT (823 residues)
- Nup116: Chain RB (1113 residues)
- Ndc1: Chain AW (655 residues)
- Nup57: Chain JB (541 residues)
- Nup49: Chain LT (200 residues)
- Dyn2: Chain GS (92 residues)
- Nup100: Chain SE (540 residues)
- Nic96: Chain KZ (839 residues)
- Nup159: Chain S (600 residues)
- Nic96: Chain LI (839 residues)
- Pom152: Chain PE (1337 residues)
- Nup49: Chain KG (200 residues)
- Nup53: Chain OG (475 residues)
- Nic96: Chain JD (839 residues)
- Gle1: Chain BJ (538 residues)
- Nup100: Chain RM (540 residues)
- Nup59: Chain BA (528 residues)
- Nup145: Chain SZ (200 residues)
- Nup100: Chain RX (540 residues)
- Pom34: Chain OO (299 residues)
- Nup57: Chain Z (200 residues)

- Sec13: Chain EI (297 residues)
- Nup57: Chain LN (200 residues)
- Nsp1: Chain KE (823 residues)
- Nup116: Chain RN (1113 residues)
- Nup1: Chain TR (698 residues)
- Nup170: Chain MK (1502 residues)
- Nup170: Chain MG (1502 residues)
- Nup100: Chain QT (959 residues)
- Nup49: Chain LD (472 residues)
- Nup133: Chain DY (1157 residues)
- Nup84: Chain BR (726 residues)
- Nup145: Chain BL (1317 residues)
- Nup60: Chain SV (539 residues)
- Nup60: Chain TQ (100 residues)
- Sec13: Chain EW (297 residues)
- Nsp1: Chain KE (600 residues)
- Nup42: Chain SB (430 residues)
- Nup145: Chain SU (1317 residues)
- Dyn2: Chain FL (92 residues)
- Nsp1: Chain HS (823 residues)
- Nup145: Chain ST (200 residues)
- Nup1: Chain SX (698 residues)
- Nup49: Chain LT (472 residues)
- Nup170: Chain NM (1502 residues)
- Nic96: Chain IF (839 residues)
- Nup49: Chain IR (200 residues)
- Nup49: Chain LD (200 residues)
- Dyn2: Chain FT (92 residues)
- Nup60: Chain SR (539 residues)
- Nsp1: Chain KL (600 residues)
- Nup60: Chain TL (100 residues)
- Sec13: Chain DG (297 residues)
- Mlp2: Chain UF (1679 residues)
- Nsp1: Chain AJ (823 residues)
- Nup1: Chain TC (1076 residues)
- Nup49: Chain AK (472 residues)
- Seh1: Chain EA (349 residues)
- Nic96: Chain JM (839 residues)
- Nup60: Chain TK (539 residues)
- Nsp1: Chain KU (823 residues)
- Nup120: Chain ES (1037 residues)
- Nup60: Chain SL (539 residues)
- Nup59: Chain QA (528 residues)
- Nup57: Chain JC (200 residues)
- Nup192: Chain NG (1683 residues)
- Nic96: Chain HY (839 residues)
- Mlp1: Chain TW (1875 residues)
- Nup100: Chain BE (959 residues)
- Nup49: Chain LE (200 residues)
- Pom34: Chain PX (299 residues)
- Nup82: Chain R (713 residues)
- Nsp1: Chain LS (600 residues)
- Seh1: Chain F (349 residues)
- Nup133: Chain DR (1157 residues)
- Nic96: Chain LA (839 residues)
- Nup133: Chain CP (1157 residues)
- Nup120: Chain J (1037 residues)
- Nup84: Chain DO (726 residues)
- Nup100: Chain SE (959 residues)
- Nsp1: Chain JG (823 residues)
- Nup157: Chain ND (1391 residues)
- Nup57: Chain LO (541 residues)
- Nup49: Chain KW (200 residues)
- Nup53: Chain AZ (475 residues)

- Nup116: Chain BH (1113 residues)
- Nsp1: Chain KU (600 residues)
- Nup188: Chain NV (1655 residues)
- Nup120: Chain DQ (1037 residues)
- Nup145c: Chain EG (712 residues)
- Nsp1: Chain FS (823 residues)
- Nup100: Chain RG (959 residues)
- Nup57: Chain LV (200 residues)
- Nup120: Chain CV (1037 residues)
- Nup192: Chain ME (1683 residues)
- Nsp1: Chain U (823 residues)
- Nup100: Chain RL (540 residues)
- Sec13: Chain FD (297 residues)
- Nup157: Chain MF (1391 residues)
- Nup116: Chain RO (1113 residues)
- Nup145c: Chain DZ (712 residues)
- Nup49: Chain KN (472 residues)
- Nup100: Chain BF (959 residues)
- Pom34: Chain PD (299 residues)
- Nup53: Chain PA (475 residues)
- Nup170: Chain MO (1502 residues)
- Seh1: Chain EH (349 residues)
- Nup57: Chain JC (541 residues)
- Nup82: Chain HJ (713 residues)
- Nup60: Chain TA (100 residues)
- Nup49: Chain JQ (472 residues)
- Nic96: Chain AE (839 residues)
- Nup170: Chain LY (1502 residues)
- Nup49: Chain JP (472 residues)
- Nic96: Chain KC (839 residues)
- Nsp1: Chain IY (600 residues)
- Nup145: Chain SP (200 residues)
- Nup53: Chain QO (475 residues)
- Nup1: Chain TM (1076 residues)
- Gle1: Chain RW (538 residues)
- Pom152: Chain PY (1337 residues)
- Nup157: Chain AM (1391 residues)
- Nup159: Chain FY (600 residues)
- Nup159: Chain GF (600 residues)
- Nsp1: Chain HR (600 residues)
- Nup1: Chain TM (698 residues)
- Nup57: Chain LF (200 residues)
- Nup84: Chain BY (726 residues)
- Nup116: Chain RU (740 residues)
- Nup100: Chain RR (959 residues)
- Nsp1: Chain IH (600 residues)
- Nic96: Chain JL (839 residues)
- Nsp1: Chain FZ (600 residues)
- Nup100: Chain RS (959 residues)
- Seh1: Chain EV (349 residues)
- Nsp1: Chain IA (600 residues)
- Nup60: Chain TG (100 residues)
- Nup49: Chain KF (472 residues)
- Sec13: Chain FK (297 residues)
- Nup82: Chain HK (713 residues)
- Nup53: Chain PU (475 residues)
- Nup57: Chain JJ (200 residues)
- Nup1: Chain TH (1076 residues)
- Nup49: Chain KV (472 residues)
- Nup84: Chain EX (726 residues)
- Nsp1: Chain HN (823 residues)
- Ndc1: Chain QQ (655 residues)
- Nup145: Chain SO (200 residues)
- Nup57: Chain KQ (200 residues)

- Nic96: Chain HP (839 residues)
- Nup157: Chain NH (1391 residues)
- Nup133: Chain D (1157 residues)
- Nup145: Chain TI (1317 residues)
- Nup145c: Chain L (712 residues)
- Nup157: Chain MJ (1391 residues)
- Mlp1: Chain UE (1875 residues)
- Nup57: Chain KI (541 residues)
- Nup49: Chain IZ (472 residues)
- Nup145: Chain BK (200 residues)
- Nup145c: Chain EN (712 residues)
- Sec13: Chain CL (297 residues)
- Nup188: Chain NR (1655 residues)
- Nup145: Chain TD (200 residues)
- Nup100: Chain RG (540 residues)
- Nic96: Chain KK (839 residues)
- Nup145: Chain SZ (1317 residues)
- Nup1: Chain SN (1076 residues)
- Nsp1: Chain GA (600 residues)
- Nup170: Chain NE (1502 residues)
- Dyn2: Chain P (92 residues)
- Nup188: Chain AS (1655 residues)
- Nsp1: Chain GP (600 residues)
- Nup49: Chain Y (200 residues)
- Nup57: Chain KY (200 residues)
- Pom152: Chain OK (1337 residues)
- Nup85: Chain B (744 residues)
- Nup84: Chain CT (726 residues)
- Nic96: Chain AI (839 residues)
- Nsp1: Chain HN (600 residues)
- Nup59: Chain QP (528 residues)
- Nsp1: Chain LC (823 residues)
- Nup159: Chain T (1460 residues)
- Nup60: Chain BO (539 residues)
- Nup159: Chain FY (1460 residues)
- Nup116: Chain RT (740 residues)
- Nup145: Chain TE (200 residues)
- Dyn2: Chain GZ (92 residues)
- Pom34: Chain BC (299 residues)
- Dyn2: Chain HH (92 residues)
- Nsp1: Chain HZ (823 residues)
- Nic96: Chain KB (839 residues)
- Nup100: Chain RA (540 residues)
- Pom152: Chain BD (1337 residues)
- Seh1: Chain DM (349 residues)
- Nup57: Chain IM (541 residues)
- Nsp1: Chain LJ (823 residues)
- Nup49: Chain IC (200 residues)
- Nup159: Chain FX (1460 residues)
- Nup1: Chain SS (1076 residues)
- Nup57: Chain AH (541 residues)
- Nsp1: Chain HR (823 residues)
- Nup57: Chain HV (200 residues)
- Nup159: Chain FQ (600 residues)
- Nup133: Chain CB (1157 residues)
- Dyn2: Chain FU (92 residues)
- Nup116: Chain RB (740 residues)
- Nic96: Chain IN (839 residues)
- Nup120: Chain DC (1037 residues)
- Nup84: Chain A (726 residues)
- Nup57: Chain KP (200 residues)
- Nup84: Chain EC (726 residues)
- Nsp1: Chain JN (823 residues)
- Nup145c: Chain CJ (712 residues)

- Nic96: Chain KS (839 residues)
- Nup57: Chain JZ (541 residues)
- Nup170: Chain NI (1502 residues)
- Nup188: Chain NB (1655 residues)
- Nup145: Chain TO (1317 residues)
- Nsp1: Chain GY (823 residues)
- Pom152: Chain OP (1337 residues)
- Nup170: Chain MS (1502 residues)
- Dyn2: Chain GR (92 residues)
- Nup53: Chain PZ (475 residues)
- Seh1: Chain FC (349 residues)
- Nup59: Chain OH (528 residues)
- Nup145: Chain BL (200 residues)
- Nsp1: Chain GI (600 residues)
- Nup84: Chain CF (726 residues)
- Nup53: Chain PF (475 residues)
- Seh1: Chain EO (349 residues)
- Nup59: Chain PV (528 residues)
- Nup116: Chain SG (1113 residues)
- Nup85: Chain ED (744 residues)
- Nup49: Chain LL (472 residues)
- Nup42: Chain RP (430 residues)
- Nup145: Chain SO (1317 residues)
- Gle1: Chain RE (538 residues)
- Nup57: Chain HV (541 residues)
- Nup60: Chain BO (100 residues)
- Nup57: Chain IU (541 residues)
- Nup57: Chain JR (541 residues)
- Nsp1: Chain GA (823 residues)
- Nup60: Chain SW (100 residues)
- Nup159: Chain GN (1460 residues)
- Ndc1: Chain OS (655 residues)
- Nup116: Chain SF (740 residues)
- Nup59: Chain OC (528 residues)
- Nup120: Chain EE (1037 residues)
- Nup49: Chain JA (472 residues)
- Nup159: Chain FP (600 residues)
- Nup82: Chain Q (713 residues)
- Pom34: Chain PN (299 residues)
- Nup53: Chain AU (475 residues)
- Nup60: Chain TL (539 residues)
- Nsp1: Chain HZ (600 residues)
- Nup49: Chain AK (200 residues)
- Nup100: Chain QU (540 residues)
- Nup60: Chain SM (100 residues)
- Pom34: Chain PS (299 residues)
- Nup1: Chain SX (1076 residues)
- Nsp1: Chain IX (600 residues)
- Nup57: Chain IT (541 residues)
- Nup60: Chain TP (100 residues)
- Nup145: Chain TD (1317 residues)
- Pom34: Chain OE (299 residues)
- Nup59: Chain AV (528 residues)
- Nsp1: Chain GX (600 residues)
- Nup57: Chain AL (200 residues)
- Nsp1: Chain AB (823 residues)
- Nup116: Chain BH (740 residues)
- Nup116: Chain BG (740 residues)
- Mlp1: Chain TS (1875 residues)
- Nup60: Chain TQ (539 residues)
- Nup53: Chain OQ (475 residues)
- Nup85: Chain EY (744 residues)
- Pom152: Chain QN (1337 residues)
- Nup159: Chain HM (1460 residues)

- Nsp1: Chain GI (823 residues)
- Nup145c: Chain CQ (712 residues)
- Nup84: Chain EQ (726 residues)
- Nup82: Chain FW (713 residues)
- Nup84: Chain FE (726 residues)
- Nup85: Chain CU (744 residues)
- Nup1: Chain BM (698 residues)
- Nup145c: Chain E (712 residues)
- Nsp1: Chain JG (600 residues)
- Nup159: Chain FX (600 residues)
- Pom152: Chain OU (1337 residues)
- Mlp1: Chain TY (1875 residues)
- Nsp1: Chain X (600 residues)
- Nup145: Chain SK (1317 residues)
- Nup145c: Chain FI (712 residues)
- Nup49: Chain IS (200 residues)
- Nup82: Chain HC (713 residues)
- Pom34: Chain OT (299 residues)
- Nup192: Chain NS (1683 residues)
- Nup60: Chain SQ (539 residues)
- Nup116: Chain RZ (1113 residues)
- Nup49: Chain HT (200 residues)
- Pom34: Chain QC (299 residues)
- Nup53: Chain QE (475 residues)
- Seh1: Chain DF (349 residues)
- Nup59: Chain OR (528 residues)
- Nup145c: Chain DS (712 residues)
- Nup133: Chain K (1157 residues)
- Nup82: Chain FN (713 residues)
- Nup100: Chain QT (540 residues)
- Nup145c: Chain EU (712 residues)
- Nup145: Chain SU (200 residues)
- Nup145c: Chain CX (712 residues)
- Nup157: Chain MB (1391 residues)
- Nsp1: Chain JF (600 residues)
- Nup85: Chain BS (744 residues)
- Pom152: Chain PT (1337 residues)
- Nup49: Chain JY (200 residues)
- Nsp1: Chain HF (823 residues)
- Nup188: Chain NN (1655 residues)
- Nup85: Chain I (744 residues)
- Pom152: Chain QD (1337 residues)
- Gle1: Chain SI (538 residues)
- Pom152: Chain OF (1337 residues)
- Nup157: Chain MZ (1391 residues)
- Nup100: Chain QU (959 residues)
- Nup1: Chain TC (698 residues)
- Nsp1: Chain KT (600 residues)
- Nup49: Chain KG (472 residues)
- Nsp1: Chain GP (823 residues)
- Nup85: Chain EK (744 residues)
- Nic96: Chain JT (839 residues)
- Mlp1: Chain UA (1875 residues)
- Nup49: Chain LM (200 residues)
- Nsp1: Chain GQ (823 residues)
- Nic96: Chain KJ (839 residues)
- Nup145: Chain TO (200 residues)

- Integrative model, File: 10.1016/j.cell.2016.10.028
- Integrative model, File: 10.5281/zenodo.1194547
- Experimental model, PDB ID: 5CWS
- Comparative model, template PDB ID: Not available
- Experimental model, PDB ID: 2QX5

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- CI: -
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- CL: -
- CM: -
- CN: -
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- CP: -
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- DY: -
- DZ: -
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- EO: -

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Rigid bodies

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- HG: -
- HH: -
- HI: -
- HJ: -
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- HM: -
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- HY: -
- HZ: -
- IA: -
- IB: -
- IC: -
- ID: -
- IE: -
- IF: -
- IG: -
- IH: -
- II: -
- IJ: -
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- IM: -
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- IO: -
- IP: -
- IQ: -
- IR: -
- IS: -
- IT: -
- IU: -
- IV: -
- IW: -
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- IY: -
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- JG: -
- JH: -
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- JL: -
- JM: -

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- JP: -
- JQ: -
- JR: -
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- JT: -
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- LP: -
- LQ: -
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- LS: -
- LT: -
- LU: -
- LV: -
- LW: -
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- LY: -

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- MF: -
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- MH: -
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- OJ: -
- OK: -

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 - UF: -

- A: 1-6, 7-20, 21-26, 27-80, 81-95, 96-126, 127-135, 136-364, 365-371, 372-483, 484-505, 506-562, 563-574, 575-726
 - B: 1-46, 47-126, 127-131, 132-230, 231-234, 235-436, 437-450, 451-492, 493-495, 496-544, 545-552, 553-560, 561-566, 567-585, 586-589, 590-597, 598-602, 603-612, 613-615, 616-634, 635-637, 638-655, 656-660, 661-675, 676-684, 685-699, 700-706, 707-719, 720-724, 725-744
 - C: 1-1, 2-29, 30-52, 53-305, 306-310, 311-711, 712-714, 715-726, 727-732, 733-746, 747-753, 754-766, 767-769, 770-781, 782-806, 807-818, 819-820, 821-833, 834-837, 838-853, 854-861, 862-879, 880-883, 884-895, 896-900, 901-913, 914-916, 917-931, 932-942, 943-955, 956-959, 960-971, 972-975, 976-987, 988-993, 994-1008, 1009-1024, 1025-1036, 1037-1037
 - D: 1-55, 56-78, 79-85, 86-125, 126-132, 133-144, 145-161, 162-184, 185-192, 193-200, 201-205, 206-249, 250-257, 258-480, 481-489, 490-763, 764-771, 772-1155, 1156-1157
 - E: 1-91, 92-99, 100-125, 126-144, 145-148, 149-550, 551-553, 554-560, 561-565, 566-576, 577-586, 587-602, 603-611, 612-624, 625-630, 631-645, 646-653, 654-673, 674-680, 681-689, 690-702, 703-712
 - F: 1-248, 249-287, 288-346, 347-349
 - G: 1-9, 10-158, 159-165, 166-296, 297-297
 - H: 1-6, 7-20, 21-26, 27-80, 81-95, 96-126, 127-135, 136-364, 365-371, 372-483, 484-505, 506-562, 563-574, 575-726
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- K: 1-55, 56-78, 79-85, 86-125, 126-132, 133-144, 145-161, 162-184, 185-192, 193-200, 201-205, 206-249, 250-257, 258-480, 481-489, 490-763, 764-771, 772-1155, 1156-1157

- L: 1-91, 92-99, 100-125, 126-144, 145-148, 149-550, 551-553, 554-560, 561-565, 566-576, 577-586, 587-602, 603-611, 612-624, 625-630, 631-645, 646-653, 654-673, 674-680, 681-689, 690-702, 703-712

- M: 1-248, 249-287, 288-346, 347-349

- N: 1-9, 10-158, 159-165, 166-296, 297-297

- O: 1-6, 7-92

- P: 1-6, 7-92

- Q: 1-6, 7-16, 17-22, 23-120, 121-122, 123-452, 453-521, 522-612, 613-624, 625-669, 670-677, 678-713

- R: 1-6, 7-16, 17-22, 23-120, 121-122, 123-452, 453-521, 522-612, 613-624, 625-669, 670-677, 678-713

- S: 1082-1116, 1117-1126, 1127-1210, 1211-1239, 1240-1265, 1266-1321, 1322-1331, 1332-1372, 1373-1381, 1382-1412, 1413-1428, 1429-1456, 1457-1460, 482-1081

- T: 1082-1116, 1117-1126, 1127-1210, 1211-1239, 1240-1265, 1266-1321, 1322-1331, 1332-1372, 1373-1381, 1382-1412, 1413-1428, 1429-1456, 1457-1460, 482-1081

- U: 601-636, 637-727, 728-741, 742-778, 779-787, 788-823, 1-600

- V: 601-636, 637-727, 728-741, 742-778, 779-787, 788-823, 1-600

- W: 1-19, 57-204, 205-360, 361-365, 366-374, 375-404, 405-444, 445-454, 455-515, 516-532, 533-747, 748-752, 753-835, 836-839

- X: 601-636, 728-741, 779-787, 1-600

- Y: 201-269, 360-368, 408-432, 1-200

- Z: 201-286, 424-432, 477-504, 541-541, 1-200

- AA: 1-19, 57-204, 205-360, 361-365, 366-374, 375-404, 405-444, 445-454, 455-515, 516-532, 533-747, 748-752, 753-835, 836-839

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- AH: 201-286, 287-423, 424-432, 433-476, 477-504, 505-540, 541-541, 1-200

- AI: 1-19, 20-56, 57-204, 205-360, 361-365, 366-374, 375-404, 405-444, 445-454, 455-515, 516-532, 533-747, 748-752, 753-835, 836-839

- AJ: 601-636, 637-727, 728-741, 742-778, 779-787, 788-823, 1-600

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- AN: 1-97, 98-299, 300-310, 311-319, 320-352, 353-471, 472-504, 505-537, 538-573, 574-717, 718-764, 765-791, 792-830, 831-862, 863-883, 884-916, 917-918, 919-930, 931-935, 936-992, 993-999, 1000-1016, 1017-1020, 1021-1033, 1034-1043, 1044-1116, 1117-1140, 1141-1191, 1192-1194, 1195-1243, 1244-1256, 1257-1502

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- OB: 1-247, 248-284, 285-303, 304-360, 361-475

- OC: 1-265, 266-302, 303-345, 346-402, 403-528

- OD: 1-655

- OE: 1-299

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- OJ: 1-299
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- PW: 1-655

- PX: 1-299
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- PZ: 1-247, 248-284, 285-303, 304-360, 361-475
- QA: 1-265, 266-302, 303-345, 346-402, 403-528
- QB: 1-655
- QC: 1-299
- QD: 1-378, 379-472, 473-519, 520-611, 612-615, 616-714, 715-721, 722-818, 819-823, 824-918, 919-930, 931-1026, 1027-1035, 1036-1141, 1142-1149, 1150-1229, 1230-1243, 1244-1337
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- QF: 1-265, 266-302, 303-345, 346-402, 403-528
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- QI: 1-378, 379-472, 473-519, 520-611, 612-615, 616-714, 715-721, 722-818, 819-823, 824-918, 919-930, 931-1026, 1027-1035, 1036-1141, 1142-1149, 1150-1229, 1230-1243, 1244-1337
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- QQ: 1-655
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- QU: 551-815, 959-959, 11-550
- QV: 751-965, 966-1111, 1112-1113, 11-750
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- QX: -
- QY: 1-120
- QZ: 551-815, 959-959, 11-550
- RA: 551-815, 959-959, 11-550
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- TY: 238-716
- TZ: 215-690
- UA: 238-716
- UB: 215-690
- UC: 238-716
- UD: 215-690
- UE: 238-716
- UF: 215-690

Structural coverage (rigid bodies)	0%
3. Restraints	
Physical principles	Information about physical principles was not provided
Experimental data	<ul style="list-style-type: none"> - 1 unique CrossLinkRestraint: Gaussian mixture models - 1 unique SASRestraint: Assembly name: SAXS subassembly Fitting method: FoXS Multi-state: False - 1 unique EM2DRestraint: DSS, 508 cross-links - 1 unique GeometricRestraint: Number of micrographs: 800, Image resolution: 35.0 - 1 unique EM3DRestraint: DSS, 509 cross-links
4. Validation	
Number of ensembles	2
Number of models in ensembles	5, 1000
Number of deposited models	2
Model precision (uncertainty of models)	1.0, Å, 1.0, Å
Data quality	<ul style="list-style-type: none"> - SASDBV9: Rg from Guinier is 1.77nm and Rg from p(r) is 1.82nm - SASDBW9: Rg from Guinier is 2.71nm and Rg from p(r) is 2.79nm - SASDBZ9: Rg from Guinier is 4.34nm and Rg from p(r) is 4.63nm - SASDBX9: Rg from Guinier is 2.78nm and Rg from p(r) is 2.64nm - SASDBY9: Rg from Guinier is 2.95nm and Rg from p(r) is 2.98nm
Model quality: assessment of excluded volume	Satisfaction: 100.00-100.00%
Fit to data used for modeling	Fit of model to information used to compute it has not been determined
Fit to data used for validation	Fit of model to information not used to compute it has not been determined
5. Methodology and Software	
1. Method	Sampling
Name	Replica exchange monte carlo
Number of computed models	500
2. Method	Sampling

<i>Name</i>	Replica exchange monte carlo
<i>Number of computed models</i>	3000
3. <i>Method</i>	Sampling
<i>Name</i>	Replica exchange monte carlo
<i>Number of computed models</i>	1000
4. <i>Method</i>	Sampling
<i>Name</i>	Brownian dynamics
<i>Number of computed models</i>	1000
<i>Software</i>	<ul style="list-style-type: none"> - Integrative Modeling Platform (IMP) (version develop-0a5706e202) - IMP PMI module (version 67456c0) - HHpred (version 2.0.16) - PSIPRED (version 4.0) - DISOPRED (version 3) - DomPred (version Not available) - COILS/PCOILS (version Not available) - EMAN2 (version 2.2) - RELION (version 1.4) - SGD (version Not available) - HeliQuest (version Not available) - MODELLER (version 9.15) - MODELLER (version 9.13)