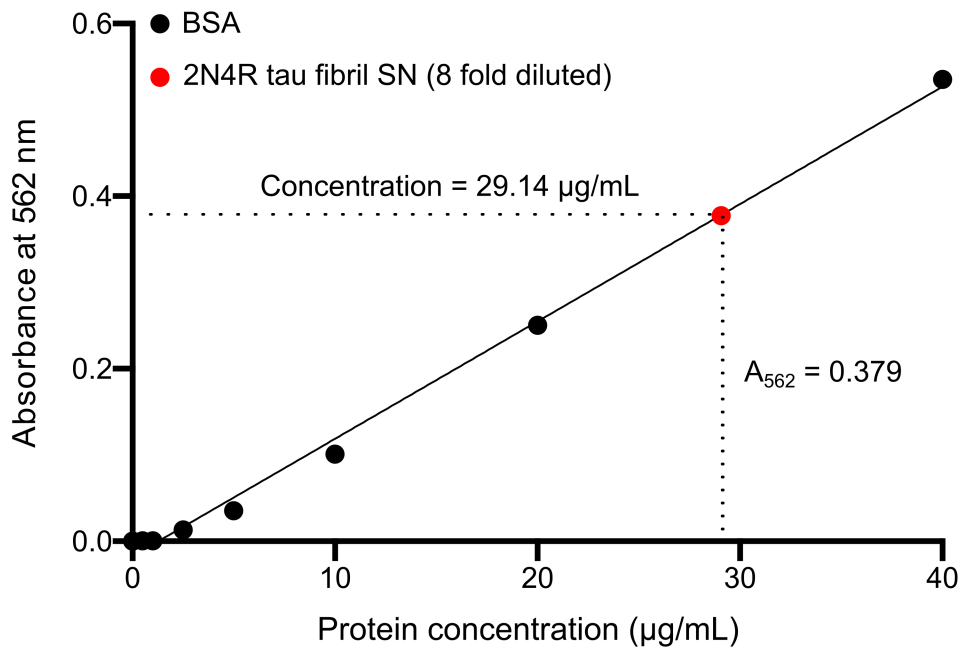


**Supplementary information for**

**Co-factor-free aggregation of tau into seeding-competent RNA-sequestering amyloid fibrils**

Pijush Chakraborty, Gwladys Rivière, Shu Liu, Alain Ibáñez de Opakua, Riza Dervişoğlu, Alina Hebestreit, Loren B. Andreas, Ina M. Vorberg and Markus Zweckstetter



Concentration of 2N4R tau before aggregation = 1.146 mg/mL

Concentration of 2N4R tau in SN =  $(29.14 \times 8) \mu\text{g/mL} = 0.233 \text{ mg/mL}$

~ 80 % 2N4R tau aggregated

**Supplementary Fig. 1** | Determination of the concentration of 2N4R tau present in the supernatant (SN; red dot; 8-fold dilution) after aggregation in the absence of heparin. Concentrations were determined using the BCA assay. Absorbance values, which were detected for BSA standard solutions with known concentrations, are shown in black.

Sequence	Observed m/z	Absolute mass (Da)	Charge	TIC	Sequence	Observed m/z	Absolute mass (Da)	Charge	TIC
LQTAPVPMPLK	663.3665	1,324.72	2	1085120	KDLSNVQSK	566.325	1,130.64	2	688486
LQTAPVPMPLK	663.3668	1,324.72	2	488941	KDLSNVQSK	377.8875	1,130.64	3	1543860
IGSTENLK	431.2404	860.4663	2	8.25E+07	KDLSNVQSK	566.3237	1,130.63	2	1.02E+07
IGSTENLK	431.2383	860.4621	2	4.33E+07	KDLSNVQSK	377.8856	1,130.63	3	1.48E+07
IGSTENLK	431.2396	860.4646	2	3.77E+07	KDLSNVQSK	377.8873	1,130.64	3	1270980
IGSTENLK	431.2394	860.4643	2	7759340	KDLSNVQSK	377.8866	1,130.64	3	2090420
IGSTENLK	431.2393	860.464	2	1.84E+07	KDLSNVQSK	377.8857	1,130.64	3	8531420
IGSTENLK	431.2398	860.4651	2	5.58E+07	KDLSNVQSK	566.3245	1,130.63	2	564889
IGSTENLK	431.2397	860.4648	2	370087	KDLSNVQSK	377.887	1,130.64	3	1.03E+08
IGSTENLK	431.2382	860.4618	2	899899	KDLSNVQSK	566.3256	1,130.64	2	424992
IGSTENLK	431.2389	860.4632	2	623917	KDLSNVQSK	377.8865	1,130.64	3	1810990
IGSTENLK	431.2384	860.4623	2	451482	KDLSNVQSK	377.8867	1,130.64	3	5202380
IGSTENLK	431.2404	860.4663	2	380415	KDLSNVQSK	566.3248	1,130.64	2	389941
IGSTENLK	431.2386	860.4627	2	846224	KDLSNVQSK	566.3263	1,130.64	2	2100210
IGSTENLKHQPGGGK	508.2739	1,521.80	3	259432	KDLSNVQSK	377.8846	1,130.63	3	3.19E+07
IGSTENLKHQPGGGK	381.4566	1,521.80	4	184486	KDLSNVQSK	377.8861	1,130.64	3	2383330
HQPGGGKQIINK	459.2663	1,374.78	3	4335000	KDLSNVQSK	377.8848	1,130.63	3	1.29E+08
HQPGGGKQIINK	459.2674	1,374.78	3	887532	KDLSNVQSK	377.8873	1,130.64	3	3748570
HQPGGGKQIINK	459.2677	1,374.78	3	490639	KDLSNVQSK	377.8847	1,130.63	3	2.92E+07
HQPGGGKQIINK	459.2662	1,374.78	3	712940	KDLSNVQSK	566.3254	1,130.64	2	398022
HQPGGGKQIINK	459.2674	1,374.78	3	397008	KDLSNVQSK	377.8855	1,130.63	3	8.94E+07
VQIINK	421.7779	841.5412	2	1974720	KDLSNVQSK	377.8847	1,130.63	3	787084
VQIINK	421.7784	841.5423	2	2.86E+07	KDLSNVQSK	377.885	1,130.63	3	3073160
VQIINK	421.7774	841.5402	2	5661730	KDLSNVQSK	377.8874	1,130.64	3	4185900
VQIINK	421.7774	841.5402	2	2975050	KDLSNVQSK	377.8843	1,130.63	3	9419510
VQIINK	421.7784	841.5423	2	517547	KDLSNVQSK	377.8844	1,130.63	3	8.83E+07
VQIINK	421.779	841.5434	2	2.54E+07	KDLSNVQSK	377.8846	1,130.63	3	1.60E+07
VQIINK	421.7782	841.5419	2	1.11E+07	KDLSNVQSK	377.8857	1,130.64	3	1774280
VQIINK	421.7787	841.5428	2	2.52E+07	KDLSNVQSK	377.886	1,130.64	3	1.05E+07
VQIINK	421.7789	841.5432	2	881426	KDLSNVQSK	377.8846	1,130.63	3	1374120
VQIINK	421.7775	841.5404	2	2254640	KDLSNVQSK	566.3251	1,130.64	2	357209
VQIINK	421.7777	841.5408	2	2.51E+07	KDLSNVQSK	566.3255	1,130.64	2	245725
VQIINK	421.7781	841.5415	2	2.88E+07	KDLSNVQSK	566.3232	1,130.63	2	389543
VQIINK	421.7788	841.5431	2	1766970	KDLSNVQSK	377.8852	1,130.63	3	1125230
VQIINK	421.778	841.5414	2	1.80E+07	KDLSNVQSK	566.3254	1,130.64	2	234322
VQIINK	421.7773	841.5401	2	5624820	KDLSNVQSK	566.3273	1,130.64	2	442197
VQIINK	421.7789	841.5432	2	3.11E+07	KDLSNVQSK	566.3274	1,130.64	2	240791
VQIINK	421.778	841.5415	2	1221890	LDLSNVQSK	502.2765	1,002.54	2	1.10E+08
VQIINK	421.7782	841.5418	2	423916	LDLSNVQSK	502.2768	1,002.54	2	2.53E+07
VQIINK	421.7779	841.5412	2	447356	LDLSNVQSK	502.2781	1,002.54	2	2.82E+07
VQIINK	421.7775	841.5404	2	459569	LDLSNVQSK	502.2783	1,002.54	2	1.07E+07
VQIINK	421.7788	841.5431	2	536818	LDLSNVQSK	502.2776	1,002.54	2	534743
VQIINKLDLSNVQSK	457.5269	1,826.08	4	623597	LDLSNVQSK	502.2746	1,002.53	2	3000260
QIINK	372.2438	742.4731	2	1779580	LDLSNVQSK	502.2748	1,002.54	2	3.65E+07
QIINK	372.2432	742.4719	2	1993940	LDLSNVQSK	502.2784	1,002.54	2	5150750
KDLSNVQ	458.7625	915.5105	2	1086550	LDLSNVQSK	502.2776	1,002.54	2	708110
KDLSNVQSK	566.3255	1,130.64	2	1.44E+07	LDLSNVQSK	502.2777	1,002.54	2	1712560
KDLSNVQSK	566.3273	1,130.64	2	3897200	LDLSNVQSK	502.2783	1,002.54	2	2509220
KDLSNVQSK	566.326	1,130.64	2	7390680	LDLSNVQSK	502.2787	1,002.54	2	777637
KDLSNVQSK	566.3244	1,130.63	2	9.73E+07	LDLSNVQSK	502.2791	1,002.54	2	639433
KDLSNVQSK	566.3249	1,130.64	2	1.12E+08	LDLSNVQSK	502.2792	1,002.54	2	1122120
KDLSNVQSK	566.3258	1,130.64	2	1482140	LDLSNVQSK	502.2781	1,002.54	2	328470
KDLSNVQSK	566.3266	1,130.64	2	1472020	LDLSNVQSK	502.2778	1,002.54	2	345601
KDLSNVQSK	566.323	1,130.63	2	4895010	LDLSNVQSK	502.2767	1,002.54	2	951925
KDLSNVQSK	566.3239	1,130.63	2	1.53E+07	LDLSNVQSK	502.2763	1,002.54	2	407844
KDLSNVQSK	566.3268	1,130.64	2	996572	CGSKDNIK	461.229	920.4434	2	1325970
KDLSNVQSK	566.3272	1,130.64	2	1049920	CGSKDNIK	461.2292	920.4438	2	1558090
KDLSNVQSK	566.3276	1,130.64	2	5839370	CGSKDNIK	461.2301	920.4455	2	1260580
KDLSNVQSK	566.324	1,130.63	2	3.11E+07	CGSKDNIK	461.2299	920.4452	2	1359890
KDLSNVQSK	566.3257	1,130.64	2	1630260	CGSKDNIK	461.229	920.4434	2	928951
KDLSNVQSK	566.3244	1,130.63	2	1.17E+07	DNKHVPGGGSVQIVYKPVDSLK	613.3406	2,449.33	4	402953
KDLSNVQSK	566.327	1,130.64	2	621179	HVPGGGSVQ	419.2169	836.4192	2	5175300
KDLSNVQSK	566.3265	1,130.64	2	1641550	HVPGGGSVQIVY	606.8246	1,211.63	2	5196730
KDLSNVQSK	566.3248	1,130.64	2	7.11E+07	HVPGGGSVQIVY	606.8227	1,211.63	2	1927530
KDLSNVQSK	566.3248	1,130.64	2	3.45E+07	HVPGGGSVQIVY	606.8281	1,211.64	2	607544
KDLSNVQSK	566.3266	1,130.64	2	2668430	HVPGGGSVQIVY	606.8248	1,211.64	2	658417

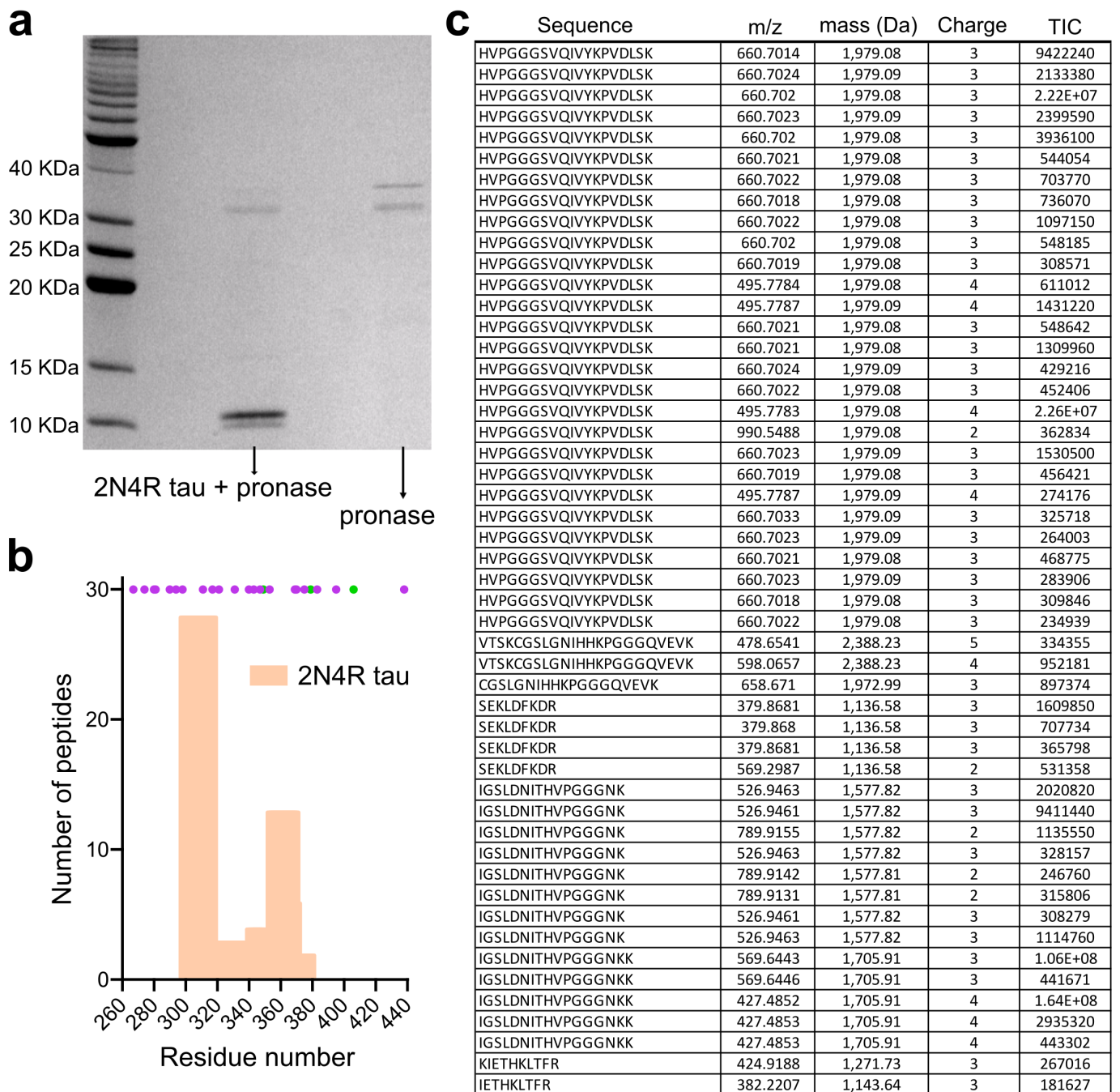
Sequence	Observed m/z	Absolute mass (Da)	Charge	TIC	Sequence	Observed m/z	Absolute mass (Da)	Charge	TIC
HVPGGGSVQIVYK	447.5846	1,339.73	3	3309080	CGSLGNIHHK	374.8553	1,121.54	3	889273
HVPGGGSVQIVYK	447.5861	1,339.74	3	1744260	CGSLGNIHHKPGGGQVEVK	658.672	1,972.99	3	1137040
HVPGGGSVQIVYK	447.5854	1,339.73	3	2346670	CGSLGNIHHKPGGGQVEVK	658.6739	1,973.00	3	1327920
HVPGGGSVQIVYK	670.8753	1,339.74	2	2555930	CGSLGNIHHKPGGGQVEVK	658.6737	1,973.00	3	992953
HVPGGGSVQIVYK	670.8753	1,339.74	2	1447930	CGSLGNIHHKPGGGQVEVK	494.2573	1,973.00	4	3370610
HVPGGGSVQIVYK	670.873	1,339.73	2	2009850	CGSLGNIHHKPGGGQVEVK	658.6732	1,973.00	3	1395720
HVPGGGSVQIVYKPVDL	588.9972	1,763.97	3	1244210	CGSLGNIHHKPGGGQVEVK	658.6652	1,972.97	3	1190500
HVPGGGSVQIVYKPVDL	588.9971	1,763.97	3	1857990	CGSLGNIHHKPGGGQVEVK	658.6742	1,973.00	3	1297690
HVPGGGSVQIVYKPVDL	660.7059	1,979.10	3	4.06E+07	CGSLGNIHHKPGGGQVEVK	658.6736	1,973.00	3	1622510
HVPGGGSVQIVYKPVDL	990.5541	1,979.09	2	3740060	CGSLGNIHHKPGGGQVEVK	658.6757	1,973.01	3	269325
HVPGGGSVQIVYKPVDL	495.7817	1,979.10	4	9258010	CGSLGNIHHKPGGGQVEVK	494.2581	1,973.00	4	2476020
HVPGGGSVQIVYKPVDL	990.557	1,979.10	2	6676660	CGSLGNIHHKPGGGQVEVK	658.6744	1,973.00	3	374600
HVPGGGSVQIVYKPVDL	495.7825	1,979.10	4	3428870	CGSLGNIHHKPGGGQVEVK	658.6737	1,973.00	3	932126
HVPGGGSVQIVYKPVDL	495.7829	1,979.10	4	2320600	CGSLGNIHHKPGGGQVEVK	494.2572	1,973.00	4	4194500
HVPGGGSVQIVYKPVDL	495.783	1,979.10	4	2371390	CGSLGNIHHKPGGGQVEVK	494.257	1,973.00	4	3819660
HVPGGGSVQIVYKPVDL	660.7048	1,979.09	3	3.59E+08	CGSLGNIHHKPGGGQVEVK	658.6749	1,973.00	3	498095
HVPGGGSVQIVYKPVDL	495.7805	1,979.09	4	2914550	CGSLGNIHHKPGGGQVEVK	658.6703	1,972.99	3	1691620
HVPGGGSVQIVYKPVDL	660.7063	1,979.10	3	9921910	CGSLGNIHHKPGGGQVEVK	494.2565	1,973.00	4	2067020
HVPGGGSVQIVYKPVDL	495.783	1,979.10	4	5478000	CGSLGNIHHKPGGGQVEVK	494.2576	1,973.00	4	1971720
HVPGGGSVQIVYKPVDL	495.7815	1,979.10	4	3192610	CGSLGNIHHKPGGGQVEVK	494.2567	1,973.00	4	631635
HVPGGGSVQIVYKPVDL	495.7822	1,979.10	4	4334490	CGSLGNIHHKPGGGQVEVK	494.2575	1,973.00	4	3016150
HVPGGGSVQIVYKPVDL	495.7813	1,979.10	4	2.57E+07	CGSLGNIHHKPGGGQVEVK	494.2559	1,972.99	4	2382460
HVPGGGSVQIVYKPVDL	660.7051	1,979.09	3	1.48E+08	CGSLGNIHHKPGGGQVEVK	494.2588	1,973.01	4	1049420
HVPGGGSVQIVYKPVDL	495.7816	1,979.10	4	1.11E+07	CGSLGNIHHKPGGGQVEVK	494.2595	1,973.01	4	1020780
HVPGGGSVQIVYKPVDL	495.7821	1,979.10	4	2.84E+07	CGSLGNIHHKPGGGQVEVK	658.6698	1,972.99	3	499499
HVPGGGSVQIVYKPVDL	495.7801	1,979.09	4	7.49E+07	CGSLGNIHHKPGGGQVEVK	494.2555	1,972.99	4	843446
HVPGGGSVQIVYKPVDL	660.7036	1,979.09	3	4.47E+08	HKPGGGQVEVK	379.2153	1,134.62	3	788203
HVPGGGSVQIVYKPVDL	495.78	1,979.09	4	2.57E+08	HKPGGGQVEVK	379.213	1,134.62	3	943892
HVPGGGSVQIVYKPVDL	495.7807	1,979.09	4	6.33E+07	KPGGGQVEVK	499.788	997.5615	2	320523
HVPGGGSVQIVYKPVDL	660.7083	1,979.10	3	1.23E+07	KPGGGQVEVK	499.7853	997.5561	2	93,917.00
HVPGGGSVQIVYKPVDL	660.7052	1,979.09	3	1.43E+08	PGGGQVEVK	435.7387	869.4629	2	675391
HVPGGGSVQIVYKPVDL	495.7815	1,979.10	4	2583980	PGGGQVEVK	435.7404	869.4663	2	1.46E+07
HVPGGGSVQIVYKPVDL	990.5507	1,979.09	2	2008380	PGGGQVEVK	435.7382	869.4619	2	1262160
HVPGGGSVQIVYKPVDL	495.7827	1,979.10	4	2159700	PGGGQVEVK	435.7381	869.4617	2	2111660
HVPGGGSVQIVYKPVDL	495.7801	1,979.09	4	3.08E+08	PGGGQVEVK	435.7405	869.4665	2	2.70E+07
HVPGGGSVQIVYKPVDL	990.5532	1,979.09	2	7801890	PGGGQVEVK	435.7384	869.4622	2	4518720
HVPGGGSVQIVYKPVDL	660.7071	1,979.10	3	1.17E+07	PGGGQVEVK	435.739	869.4635	2	1.56E+07
HVPGGGSVQIVYKPVDL	990.5562	1,979.10	2	381200	PGGGQVEVK	435.7386	869.4627	2	835934
HVPGGGSVQIVYKPVDL	660.7081	1,979.10	3	1.13E+07	PGGGQVEVK	435.7368	869.459	2	9522350
HVPGGGSVQIVYKPVDL	660.7067	1,979.10	3	1.17E+07	PGGGQVEVK	435.742	869.4694	2	528237
HVPGGGSVQIVYKPVDL	660.7082	1,979.10	3	3.74E+07	GGGQVEVK	387.2131	772.4116	2	322668
HVPGGGSVQIVYKPVDL	495.7822	1,979.10	4	2446550	GGQVEVK	358.7026	715.3906	2	595542
HVPGGGSVQIVYKPVDL	660.7058	1,979.10	3	1.38E+07	SEKLDK	433.7355	865.4564	2	2037840
HVPGGGSVQIVYKPVDL	660.7045	1,979.09	3	2311750	SEKLDK	433.7361	865.4577	2	1644280
GGGSVQIVYKPVDL	549.6429	1,645.91	3	945519	SEKLDK	433.736	865.4574	2	2002940
GGSVQIVYKPVDL	530.6381	1,588.89	3	1860150	SEKLDK	433.7367	865.4588	2	782481
GGSVQIVYKPVDL	530.6366	1,588.89	3	2202040	SEKLDK	433.7353	865.456	2	1552130
GGSVQIVYKPVDL	511.6294	1,531.87	3	3717640	SEKLDK	433.7364	865.4582	2	713546
GGSVQIVYKPVDL	511.6291	1,531.87	3	1347090	SEKLDK	433.7361	865.4576	2	1239020
SVQIVYKPVDL	492.6227	1,474.85	3	1216430	SEKLDK	433.7363	865.4581	2	370496
SVQIVYKPVDL	738.4289	1,474.84	2	815127	SEKLDK	433.7362	865.4579	2	499564
SVQIVYKPVDL	738.4314	1,474.85	2	979506	LDKDR	397.2141	792.4137	2	1977410
SVQIVYKPVDL	492.6235	1,474.85	3	1558740	LDKDR	397.2155	792.4164	2	1013110
IVYKPVDL	387.9027	1,160.69	3	840122	LDKDR	397.2157	792.4168	2	836590
IVYKPVDL	387.903	1,160.69	3	922994	LDKDR	397.2175	792.4204	2	1887930
KPVDL	393.7404	785.4662	2	555221	LDKDR	397.2141	792.4136	2	917648
KPVDL	393.7412	785.4678	2	387968	IGSLDNITHVPGGGNK	526.9511	1,577.83	3	1608920
KPVDL	393.7415	785.4685	2	568540	IGSLDNITHVPGGGNK	526.9508	1,577.83	3	1536100
KPVDL	393.74	785.4654	2	591744	IGSLDNITHVPGGGNK	526.951	1,577.83	3	489459
KPVDL	393.7413	785.468	2	430025	IGSLDNITHVPGGGNK	526.9509	1,577.83	3	689049
KPVDL	393.7404	785.4663	2	579873	IGSLDNITHVPGGGNKK	569.6486	1,705.92	3	440431
KPVDL	393.7396	785.4647	2	364953	IGSLDNITHVPGGGNKK	569.6498	1,705.93	3	420993
KPVDL	393.7407	785.4668	2	466148	KIETHK	378.2239	754.4333	2	130320
KPVDL	393.7422	785.4699	2	339201	KIETHK	378.2263	754.4381	2	125368
KPVDL	393.7424	785.4702	2	816433	KIETHK	378.2264	754.4382	2	693732
CGSLGNIHHK	374.8554	1,121.54	3	738854					

**Supplementary Fig. 2** | Details of the peptides detected (shown in Fig. 2c, black) by enzymatic digestion of the trypsin protected core (Fig. 2b) of heparin-induced 2N4R tau fibrils.

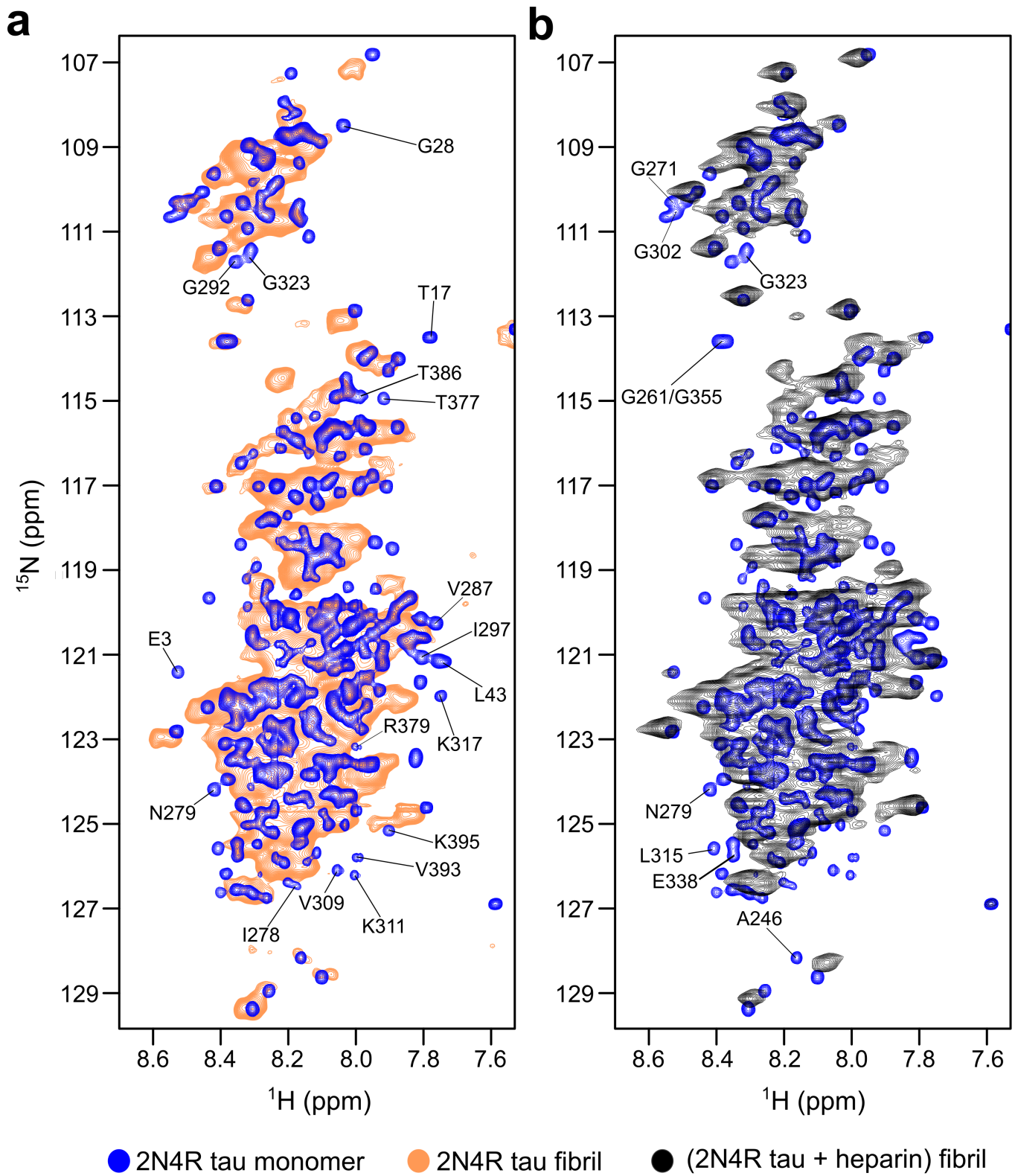
Sequence	Observed m/z	Absolute mass (Da)	Charge	TIC	Sequence	Observed m/z	Absolute mass (Da)	Charge	TIC
AKTDHGAEIVYK	666.352	1,330.69	2	2112160	HVPGGGSVQIVYKPVDSLK	660.7082	1,979.10	3	2811180
AKTDHGAEIVYK	666.3525	1,330.69	2	1441290	HVPGGGSVQIVYKPVDSLK	660.7073	1,979.10	3	1991730
AKTDHGAEIVYK	444.5705	1,330.69	3	2.02E+07	HVPGGGSVQIVYKPVDSLK	660.7075	1,979.10	3	2627530
AKTDHGAEIVYK	444.57	1,330.69	3	5533990	HVPGGGSVQIVYKPVDSLK	660.7018	1,979.08	3	4.21E+09
CGSLGNIHHKPGGGQVEVK	987.5019	1,972.99	2	458225	HVPGGGSVQIVYKPVDSLK	660.7077	1,979.10	3	2581060
CGSLGNIHHKPGGGQVEVK	658.6707	1,972.99	3	2941340	HVPGGGSVQIVYKPVDSLK	495.7808	1,979.09	4	532325
CGSLGNIHHKPGGGQVEVK	494.2547	1,972.99	4	1580130	HVPGGGSVQIVYKPVDSLK	660.7045	1,979.09	3	2597390
DNIKHVPGGGSVQIVYKPVDSLK	817.4449	2,449.31	3	2528420	HVPGGGSVQIVYKPVDSLK	495.7801	1,979.09	4	636190
DNIKHVPGGGSVQIVYKPVDSLK	613.3411	2,449.34	4	7021630	HVPGGGSVQIVYKPVDSLK	495.781	1,979.09	4	532791
DNIKHVPGGGSVQIVYKPVDSLK	817.4438	2,449.31	3	1155600	HVPGGGSVQIVYKPVDSLK	495.7811	1,979.10	4	714262
DNIKHVPGGGSVQIVYKPVDSLK	613.3351	2,449.31	4	5540500	HVPGGGSVQIVYKPVDSLK	495.7824	1,979.10	4	482556
DRVQSKIGSLDNITHVPGGGNK	764.7397	2,291.20	3	3097320	HVPGGGSVQIVYKPVDSLK	660.7021	1,979.08	3	1.16E+09
DRVQSKIGSLDNITHVPGGGNK	459.2466	2,291.20	5	2628490	HVPGGGSVQIVYKPVDSLK	495.7807	1,979.09	4	581133
DRVQSKIGSLDNITHVPGGGNK	484.8668	2,419.30	5	1873910	HVPGGGSVQIVYKPVDSLK	495.7804	1,979.09	4	906451
HQPGGGKVQIINK	459.2658	1,374.78	3	690270	HVPGGGSVQIVYKPVDSLK	990.553	1,979.09	2	501700
HQPGGGKVQIINK	376.7255	1,502.87	4	352294	HVPGGGSVQIVYKPVDSLK	660.7021	1,979.08	3	2.84E+07
HVPGGGSVQIVYKPVDSLK	990.5506	1,979.09	2	1.50E+07	HVPGGGSVQIVYKPVDSLK	495.7809	1,979.09	4	639708
HVPGGGSVQIVYKPVDSLK	660.704	1,979.09	3	8369030	HVPGGGSVQIVYKPVDSLK	495.7788	1,979.09	4	2486670
HVPGGGSVQIVYKPVDSLK	660.7055	1,979.09	3	5284610	HVPGGGSVQIVYKPVDSLKVTSK	599.5907	2,394.33	4	2.31E+07
HVPGGGSVQIVYKPVDSLK	660.7059	1,979.10	3	7011870	HVPGGGSVQIVYKPVDSLKVTSK	799.1186	2,394.33	3	4130730
HVPGGGSVQIVYKPVDSLK	660.7037	1,979.09	3	4.69E+07	IETHKLTFRENAK	793.9355	1,585.86	2	2190220
HVPGGGSVQIVYKPVDSLK	495.7793	1,979.09	4	949283	IETHKLTFRENAK	529.6262	1,585.86	3	1.15E+07
HVPGGGSVQIVYKPVDSLK	660.7061	1,979.10	3	2224980	IETHKLTFRENAKAK	596.0042	1,784.99	3	1762750
HVPGGGSVQIVYKPVDSLK	660.7016	1,979.08	3	1.06E+07	IGSLDNITHVPGGGNK	789.9158	1,577.82	2	3.73E+08
HVPGGGSVQIVYKPVDSLK	660.7066	1,979.10	3	3102850	IGSLDNITHVPGGGNK	789.9163	1,577.82	2	3.21E+07
HVPGGGSVQIVYKPVDSLK	495.7789	1,979.09	4	1561800	IGSLDNITHVPGGGNK	789.9152	1,577.82	2	9.96E+08
HVPGGGSVQIVYKPVDSLK	660.7066	1,979.10	3	2940350	IGSLDNITHVPGGGNK	789.9168	1,577.82	2	810409
HVPGGGSVQIVYKPVDSLK	660.7041	1,979.09	3	1.25E+07	IGSLDNITHVPGGGNK	526.9493	1,577.83	3	1651830
HVPGGGSVQIVYKPVDSLK	495.7804	1,979.09	4	722816	IGSLDNITHVPGGGNK	789.9173	1,577.82	2	2651900
HVPGGGSVQIVYKPVDSLK	660.7077	1,979.10	3	2105500	IGSLDNITHVPGGGNK	526.9488	1,577.82	3	2066100
HVPGGGSVQIVYKPVDSLK	660.7056	1,979.09	3	3327480	IGSLDNITHVPGGGNK	526.9498	1,577.83	3	1975390
HVPGGGSVQIVYKPVDSLK	660.7067	1,979.10	3	1646850	IGSLDNITHVPGGGNK	789.9175	1,577.82	2	5733840
HVPGGGSVQIVYKPVDSLK	495.7806	1,979.09	4	1388860	IGSLDNITHVPGGGNK	526.9476	1,577.82	3	1.48E+07
HVPGGGSVQIVYKPVDSLK	660.7051	1,979.09	3	4058520	IGSLDNITHVPGGGNK	526.9492	1,577.83	3	1.53E+08
HVPGGGSVQIVYKPVDSLK	660.7028	1,979.09	3	5430780	IGSLDNITHVPGGGNK	526.9499	1,577.83	3	2.83E+09
HVPGGGSVQIVYKPVDSLK	495.7783	1,979.08	4	1.01E+09	IGSLDNITHVPGGGNK	526.9506	1,577.83	3	1707060
HVPGGGSVQIVYKPVDSLK	495.7788	1,979.09	4	847724	IGSLDNITHVPGGGNK	526.949	1,577.83	3	1876050
HVPGGGSVQIVYKPVDSLK	660.7027	1,979.09	3	1.19E+07	IGSLDNITHVPGGGNK	526.9502	1,577.83	3	1906910
HVPGGGSVQIVYKPVDSLK	990.5495	1,979.08	2	1409330	IGSLDNITHVPGGGNK	526.9468	1,577.82	3	3882530
HVPGGGSVQIVYKPVDSLK	495.7795	1,979.09	4	1153680	IGSLDNITHVPGGGNK	526.9481	1,577.82	3	7.70E+08
HVPGGGSVQIVYKPVDSLK	495.7788	1,979.09	4	4.69E+07	IGSLDNITHVPGGGNK	526.9464	1,577.82	3	2791160
HVPGGGSVQIVYKPVDSLK	660.7034	1,979.09	3	1.47E+08	IGSLDNITHVPGGGNK	526.9465	1,577.82	3	3411380
HVPGGGSVQIVYKPVDSLK	660.7051	1,979.09	3	3208480	IGSLDNITHVPGGGNK	526.946	1,577.82	3	1529550
HVPGGGSVQIVYKPVDSLK	660.704	1,979.09	3	2.81E+07	IGSLDNITHVPGGGNK	526.9482	1,577.82	3	1737920
HVPGGGSVQIVYKPVDSLK	495.7811	1,979.10	4	771167	IGSLDNITHVPGGGNK	526.9475	1,577.82	3	2552780
HVPGGGSVQIVYKPVDSLK	660.7043	1,979.09	3	9691810	IGSLDNITHVPGGGNK	526.9481	1,577.82	3	3058160
HVPGGGSVQIVYKPVDSLK	495.7798	1,979.09	4	693595	IGSLDNITHVPGGGNK	526.9468	1,577.82	3	9277720
HVPGGGSVQIVYKPVDSLK	495.7801	1,979.09	4	976561	IGSLDNITHVPGGGNK	526.946	1,577.82	3	1.13E+07
HVPGGGSVQIVYKPVDSLK	660.7021	1,979.08	3	8.28E+07	IGSLDNITHVPGGGNK	526.9472	1,577.82	3	2636520
HVPGGGSVQIVYKPVDSLK	495.7785	1,979.08	4	4.53E+08	IGSLDNITHVPGGGNK	526.9487	1,577.82	3	2985840
HVPGGGSVQIVYKPVDSLK	495.7791	1,979.09	4	1097450	IGSLDNITHVPGGGNK	526.9488	1,577.82	3	1773780
HVPGGGSVQIVYKPVDSLK	660.7044	1,979.09	3	1.00E+07	IGSLDNITHVPGGGNK	526.9481	1,577.82	3	2273800
HVPGGGSVQIVYKPVDSLK	990.5496	1,979.08	2	3.50E+08	IGSLDNITHVPGGGNK	789.9166	1,577.82	2	889692
HVPGGGSVQIVYKPVDSLK	660.707	1,979.10	3	2223410	IGSLDNITHVPGGGNK	526.9468	1,577.82	3	3380620
HVPGGGSVQIVYKPVDSLK	495.7791	1,979.09	4	4140980	IGSLDNITHVPGGGNK	526.9499	1,577.83	3	1899070
HVPGGGSVQIVYKPVDSLK	660.7019	1,979.08	3	1.89E+07	IGSLDNITHVPGGGNK	526.9486	1,577.82	3	2689600
HVPGGGSVQIVYKPVDSLK	495.7818	1,979.10	4	693787	IGSLDNITHVPGGGNK	789.9147	1,577.81	2	2997430
HVPGGGSVQIVYKPVDSLK	660.7065	1,979.10	3	2884480	IGSLDNITHVPGGGNK	526.9478	1,577.82	3	4.87E+07
HVPGGGSVQIVYKPVDSLK	660.7054	1,979.09	3	4776200	IGSLDNITHVPGGGNK	526.9464	1,577.82	3	3526230
HVPGGGSVQIVYKPVDSLK	495.7782	1,979.08	4	2.81E+07	IGSLDNITHVPGGGNK	526.9468	1,577.82	3	6623530
HVPGGGSVQIVYKPVDSLK	495.781	1,979.09	4	1361110	IGSLDNITHVPGGGNK	526.948	1,577.82	3	2208650
HVPGGGSVQIVYKPVDSLK	660.706	1,979.10	3	2744780	IGSLDNITHVPGGGNK	526.9481	1,577.82	3	5383120
HVPGGGSVQIVYKPVDSLK	660.7045	1,979.09	3	9240940	IGSLDNITHVPGGGNK	526.9467	1,577.82	3	4841190
HVPGGGSVQIVYKPVDSLK	495.7788	1,979.09	4	793429	IGSLDNITHVPGGGNK	526.9462	1,577.82	3	2.41E+07
HVPGGGSVQIVYKPVDSLK	660.7033	1,979.09	3	1.32E+07	IGSLDNITHVPGGGNK	526.948	1,577.82	3	1.15E+08
HVPGGGSVQIVYKPVDSLK	495.7794	1,979.09	4	1.22E+07	IGSLDNITHVPGGGNK	526.9489	1,577.82	3	1.03E+07
HVPGGGSVQIVYKPVDSLK	495.7797	1,979.09	4	1596920	IGSLDNITHVPGGGNK	526.9501	1,577.83	3	1713510
HVPGGGSVQIVYKPVDSLK	660.7037	1,979.09	3	4333350	IGSLDNITHVPGGGNK	526.9457	1,577.82	3	4650700
HVPGGGSVQIVYKPVDSLK	990.5502	1,979.09	2	406341	IGSLDNITHVPGGGNK	526.9487	1,577.82	3	1656710
HVPGGGSVQIVYKPVDSLK	660.7068	1,979.10	3	4017330	IGSLDNITHVPGGGNK	789.9152	1,577.82	2	2125410
HVPGGGSVQIVYKPVDSLK	495.779	1,979.09	4	8845590	IGSLDNITHVPGGGNK	526.9503	1,577.83	3	1577080
HVPGGGSVQIVYKPVDSLK	495.7822	1,979.10	4	594417	IGSLDNITHVPGGGNK	526.9464	1,577.82	3	9456110
HVPGGGSVQIVYKPVDSLK	495.7813	1,979.10	4	658746	IGSLDNITHVPGGGNK	526.95	1,577.83	3	1738300

Sequence	Observed m/z	Absolute mass (Da)	Charge	TIC	Sequence	Observed m/z	Absolute mass (Da)	Charge	TIC
IGSLDNITHVPGGGNK	526.9491	1,577.83	3	1155100	SEKLDKDRVQSK	395.7166	1,578.84	4	1045490
IGSLDNITHVPGGGNK	526.9484	1,577.82	3	221418	SPVVSVDTSRPHLSNVSSSTGSDIMVDSPLATLADEVASLAK	1,448.05	4,341.14	3	5789350
IGSLDNITHVPGGGNK	526.9493	1,577.83	3	286334	SPVVSVDTSRPHLSNVSSSTGSDIMVDSPLATLADEVASLAK	1,086.30	4,341.14	4	3540890
IGSLDNITHVPGGGNK	526.9454	1,577.81	3	5477080	SPVVSVDTSRPHLSNVSSSTGSDIMVDSPLATLADEVASLAK	869.2347	4,341.14	5	7381630
IGSLDNITHVPGGGNK	526.9497	1,577.83	3	2062070	TDHGAEIVYK	566.7853	1,131.56	2	2.79E+08
IGSLDNITHVPGGGNK	526.9471	1,577.82	3	2340520	TDHGAEIVYK	566.7865	1,131.56	2	5572430
IGSLDNITHVPGGGNK	526.9483	1,577.82	3	2040250	TDHGAEIVYK	378.1934	1,131.56	3	2524120
IGSLDNITHVPGGGNK	789.9179	1,577.82	2	1253410	TDHGAEIVYK	378.1937	1,131.56	3	1684140
IGSLDNITHVPGGGNK	789.9111	1,577.81	2	1343070	TDHGAEIVYK	378.1937	1,131.56	3	1289780
IGSLDNITHVPGGGNK	526.9494	1,577.83	3	166880	TDHGAEIVYK	378.1925	1,131.56	3	4498190
IGSLDNITHVPGGGNK	526.9495	1,577.83	3	321867	TDHGAEIVYK	378.1926	1,131.56	3	1391590
IGSLDNITHVPGGGNK	853.964	1,705.91	2	1.66E+07	TDHGAEIVYK	378.1935	1,131.56	3	1948310
IGSLDNITHVPGGGNK	853.9636	1,705.91	2	3424590	TDHGAEIVYK	378.1924	1,131.56	3	672651
IGSLDNITHVPGGGNK	569.6451	1,705.91	3	1.05E+08	TDHGAEIVYK	566.7853	1,131.56	2	6059200
IGSLDNITHVPGGGNK	569.6451	1,705.91	3	8.15E+08	TDHGAEIVYK	566.7852	1,131.56	2	2.88E+07
IGSLDNITHVPGGGNK	853.9622	1,705.91	2	1052510	TDHGAEIVYK	378.1931	1,131.56	3	3423510
IGSLDNITHVPGGGNK	427.487	1,705.92	4	3.26E+07	TDHGAEIVYK	566.7856	1,131.56	2	3038160
IGSLDNITHVPGGGNK	427.4857	1,705.91	4	4857820	TDHGAEIVYK	378.1936	1,131.56	3	1510270
IGSLDNITHVPGGGNK	427.4857	1,705.91	4	897717	TDHGAEIVYK	378.1924	1,131.56	3	5707550
IGSLDNITHVPGGGNK	427.486	1,705.91	4	987470	TDHGAEIVYK	566.7869	1,131.56	2	1789910
IGSLDNITHVPGGGNK	427.4863	1,705.92	4	7266090	TDHGAEIVYK	566.7855	1,131.56	2	1.23E+09
IGSTENLK	431.239	860.4635	2	4936660	TDHGAEIVYK	378.1933	1,131.56	3	2052700
IGSTENLKHQPGGGK	381.4567	1,521.80	4	428848	TDHGAEIVYK	566.7878	1,131.56	2	606715
KIETHKLTFRNAK	572.3261	1,713.96	3	2728970	TDHGAEIVYK	378.1934	1,131.56	3	1.32E+07
KIETHKLTFRNAK	429.4954	1,713.95	4	1568200	TDHGAEIVYK	378.1928	1,131.56	3	796246
KIETHKLTFRNAK	429.4954	1,713.95	4	1179910	TDHGAEIVYK	378.194	1,131.56	3	1.24E+09
KLDLSNVQSK	566.3237	1,130.63	2	6777390	TDHGAEIVYK	378.1933	1,131.56	3	816200
KLDLSNVQSK	377.8845	1,130.63	3	6797930	TDHGAEIVYK	378.1932	1,131.56	3	1392660
LDKDRVQSK	618.3409	1,234.67	2	2709780	TDHGAEIVYK	378.1932	1,131.56	3	899653
LDKDRVQSK	618.3421	1,234.67	2	4709350	TDHGAEIVYK	566.7892	1,131.56	2	276842
LDKDRVQSK	618.3406	1,234.67	2	3.65E+07	TDHGAEIVYK	566.7862	1,131.56	2	1995230
LDKDRVQSK	618.3407	1,234.67	2	2.44E+08	TDHGAEIVYK	378.1941	1,131.56	3	725129
LDKDRVQSK	618.3415	1,234.67	2	2048270	TDHGAEIVYK	378.1931	1,131.56	3	1147450
LDKDRVQSK	618.3406	1,234.67	2	590390	TDHGAEIVYK	378.1928	1,131.56	3	793357
LDKDRVQSK	412.563	1,234.67	3	4055280	TDHGAEIVYK	378.1926	1,131.56	3	4495100
LDKDRVQSK	412.565	1,234.67	3	1648240	TDHGAEIVYK	566.7894	1,131.56	2	643479
LDKDRVQSK	412.5636	1,234.67	3	2329710	TDHGAEIVYK	566.7849	1,131.56	2	2295400
LDKDRVQSK	412.5623	1,234.67	3	6494730	TDHGAEIVYK	378.1934	1,131.56	3	2985250
LDKDRVQSK	412.5643	1,234.67	3	621133	TDHGAEIVYK	566.7894	1,131.56	2	619166
LDKDRVQSK	412.5646	1,234.67	3	631690	TDHGAEIVYK	566.7863	1,131.56	2	1401040
LDKDRVQSK	412.5642	1,234.67	3	605167	TDHGAEIVYK	378.1931	1,131.56	3	1147240
LDKDRVQSK	412.5644	1,234.67	3	1714350	TDHGAEIVYK	378.1939	1,131.56	3	1067540
LDKDRVQSK	412.5627	1,234.67	3	8990510	TDHGAEIVYK	566.787	1,131.56	2	1621820
LDKDRVQSK	412.5637	1,234.67	3	1405350	TDHGAEIVYK	378.1934	1,131.56	3	828802
LDKDRVQSKIGSLDNITHVPGGGNK	699.6224	2,794.46	4	2945180	TDHGAEIVYK	378.193	1,131.56	3	791990
LDKDRVQSKIGSLDNITHVPGGGNK	699.6266	2,794.48	4	1.04E+07	TDHGAEIVYK	378.1919	1,131.56	3	873625
LDLSNVQSK	502.2763	1,002.54	2	1.49E+08	TDHGAEIVYK	378.1946	1,131.56	3	856211
LDLSNVQSK	502.2764	1,002.54	2	3602480	TDHGAEIVYK	378.1934	1,131.56	3	937579
LDLSNVQSK	502.2762	1,002.54	2	2210260	TDHGAEIVYK	566.788	1,131.56	2	1283420
LTFRENAK	489.7738	977.533	2	1.10E+09	TDHGAEIVYK	378.1931	1,131.56	3	1210650
LTFRENAK	489.7716	977.5286	2	1.52E+07	TDHGAEIVYK	566.7864	1,131.56	2	222469
LTFRENAK	489.7725	977.5305	2	2.51E+08	TDHGAEIVYK	378.1927	1,131.56	3	656860
LTFRENAK	489.7747	977.5348	2	2271510	TDHGAEIVYK	378.1929	1,131.56	3	674648
LTFRENAK	489.7745	977.5345	2	925989	TDHGAEIVYK	378.1915	1,131.56	3	646219
LTFRENAK	489.7739	977.5332	2	484169	TDHGAEIVYK	378.1924	1,131.56	3	868838
LTFRENAK	489.7726	977.5306	2	286652	TDHGAEIVYK	378.1935	1,131.56	3	652849
LTFRENAK	489.7739	977.5332	2	467440	TDHGAEIVYK	378.193	1,131.56	3	1234380
LTFRENAK	489.7727	977.5308	2	428820	TDHGAEIVYK	378.1936	1,131.56	3	1078030
LTFRENAK	489.7749	977.5352	2	396975	TDHGAEIVYK	378.1939	1,131.56	3	846087
LTFRENAK	489.7744	977.5343	2	313402	TDHGAEIVYK	378.1937	1,131.56	3	788023
LTFRENAK	489.7719	977.5292	2	4153020	TDHGAEIVYK	378.1934	1,131.56	3	820444
LTFRENAK	489.7743	977.5341	2	556436	TDHGAEIVYK	378.1934	1,131.56	3	4.24E+08
LTFRENAK	489.7764	977.5382	2	343102	TDHGAEIVYK	378.1934	1,131.56	3	7208060
LTFRENAK	489.7726	977.5306	2	325833	TDHGAEIVYK	566.786	1,131.56	2	1689280
LTFRENAK	489.7721	977.5296	2	1304710	TDHGAEIVYK	378.1951	1,131.56	3	1187870
LTFRENAK	489.7716	977.5287	2	1926160	TDHGAEIVYK	378.1952	1,131.56	3	700429
LTFRENAK	489.7726	977.5306	2	494865	TDHGAEIVYK	378.1929	1,131.56	3	805639
LTFRENAK	489.7729	977.5312	2	668740	TDHGAEIVYK	378.1937	1,131.56	3	994250
LTFRENAKAK	589.3387	1,176.66	2	1.17E+07	TDHGAEIVYK	378.1927	1,131.56	3	648024
LTFRENAKAKTDHGAEIVYK	573.5607	2,290.21	4	7300470	TDHGAEIVYK	566.7864	1,131.56	2	795048
SEKLDKDRVQSK	790.4258	1,578.84	2	862065	TDHGAEIVYK	378.1926	1,131.56	3	675877
SEKLDKDRVQSK	527.2865	1,578.84	3	4558580	TDHGAEIVYK	566.7859	1,131.56	2	355248
SEKLDKDRVQSK	527.2872	1,578.84	3	5798240	TDHGAEIVYKSPVVSVDTSRPHLSNVSSSTGSDIMVDSPLATLADEVASLAK	1,091.95	5,454.70	5	4.77E+07
					TDHGAEIVYKSPVVSVDTSRPHLSNVSSSTGSDIMVDSPLATLADEVASLAK	907.4586	5,438.71	6	5919710
					TDHGAEIVYKSPVVSVDTSRPHLSNVSSSTGSDIMVDSPLATLADEVASLAK	1,088.75	5,438.72	5	600274

**Supplementary Fig. 3** | Details of the peptides detected (shown in Fig. 2c, orange) by enzymatic digestion of the trypsin protected core (Fig. 2b) of heparin-free 2N4R tau fibrils.

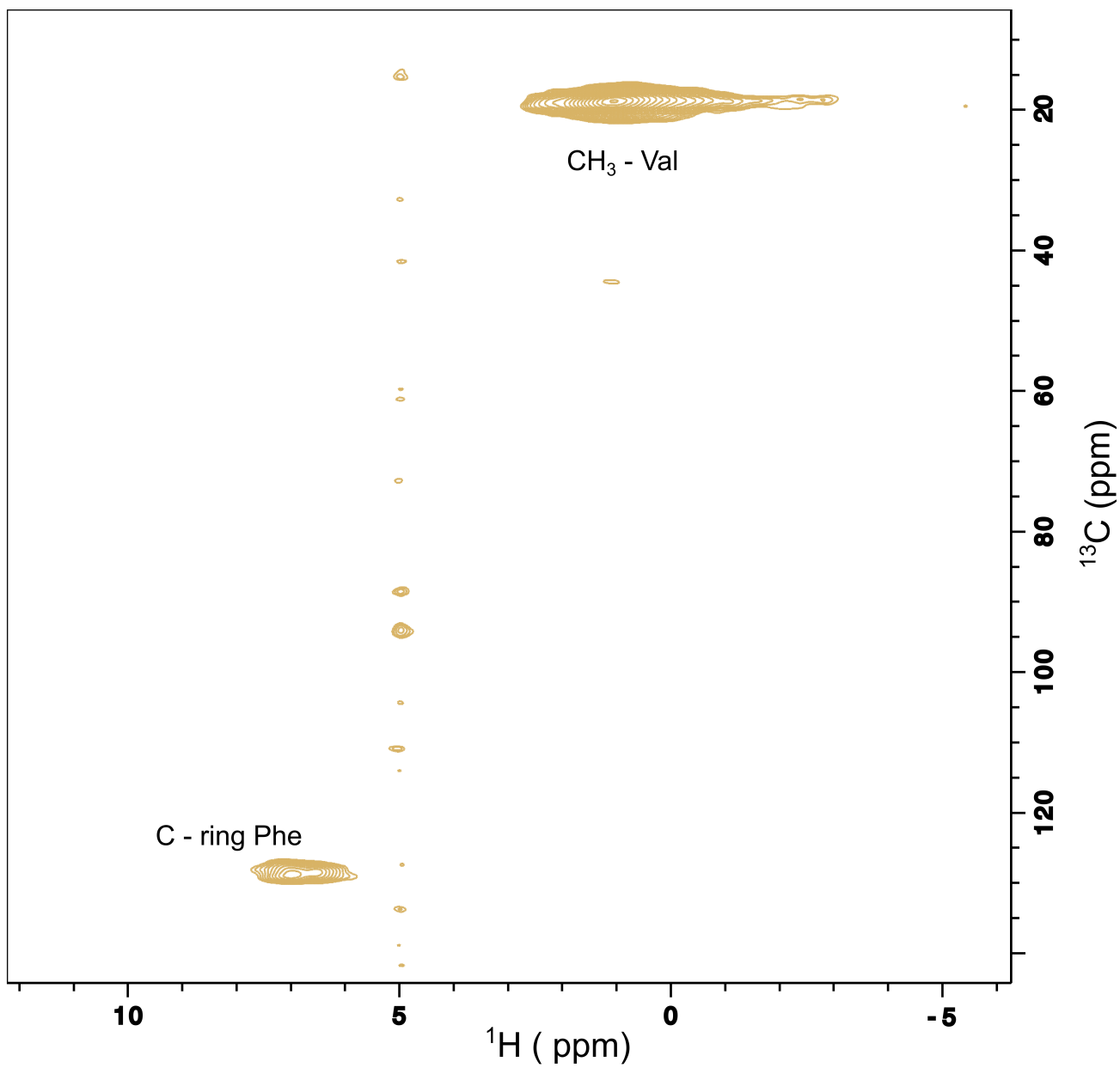


**Supplementary Fig. 4 | Pronase-resistant core of heparin-free 2N4R tau fibrils.** **a**, SDS-PAGE gel of pronase-digested tau fibrils formed in the absence of heparin. The pronase band is indicated in a separate lane. The result was reproducible for three independently performed experiments. **b**, Number of peptides detected from the enzymatic digestion of the tau band observed in SDS-PAGE in (a). The position of lysine and arginine residues in 2N4R tau are marked with purple and green dots, respectively. **c**, Details of the peptides detected as shown in (b).

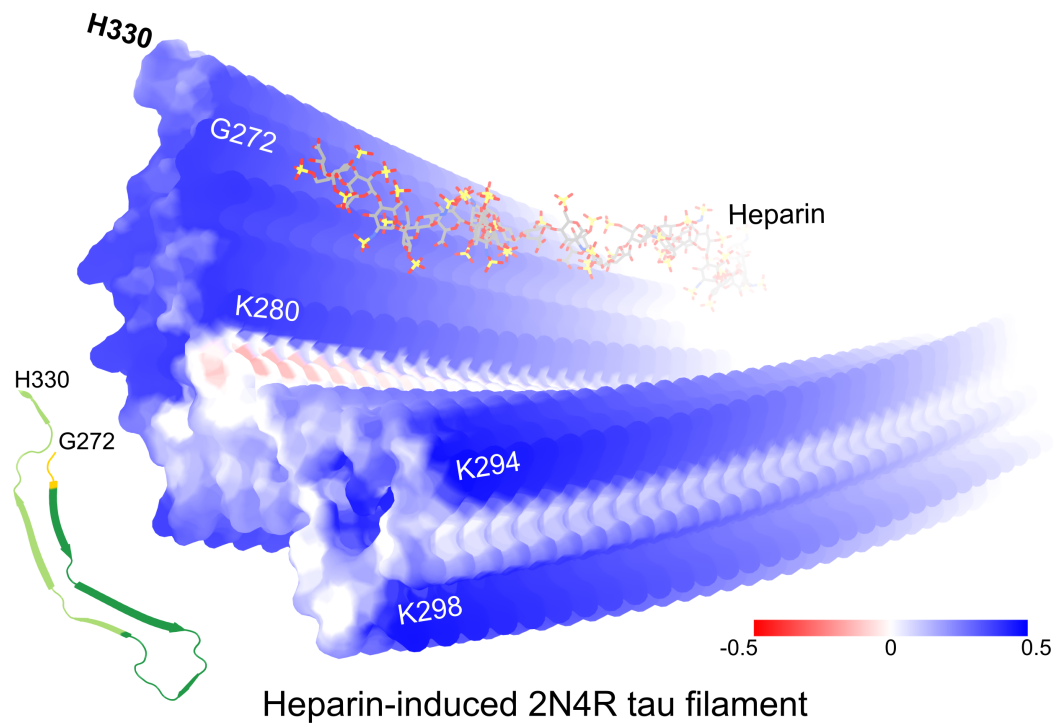


**Supplementary Fig. 5** | Superposition of the  $^1\text{H}$ - $^{15}\text{N}$  HSQC spectrum of monomeric 2N4R tau (blue) with  $^1\text{H}$ - $^{15}\text{N}$  INEPT spectra of heparin-free 2N4R fibrils (orange; left) and heparin-induced fibrils of 2N4R tau (black; right).





**Supplementary Fig. 6** | 2D hCH spectra of heparin-free fibrils of  $^{13}\text{C}_\gamma$  valine,  $^{13}\text{C}$ -ring phenylalanine-labeled 2N4R tau.



**Supplementary Fig. 7** | Electrostatic surface potential of the heparin-induced 2N4R tau fibrils (snake form; PDB code: 6QJH). Positively and negatively charged residues are shown in blue and red, respectively. A molecule of heparin is displayed to illustrate the presence of heparin on the surface of the fibril.