

Supplementary Information

Structure-Based in Silico Screening Identifies a Potent Ebolavirus Inhibitor from a Traditional Chinese Medicine Library

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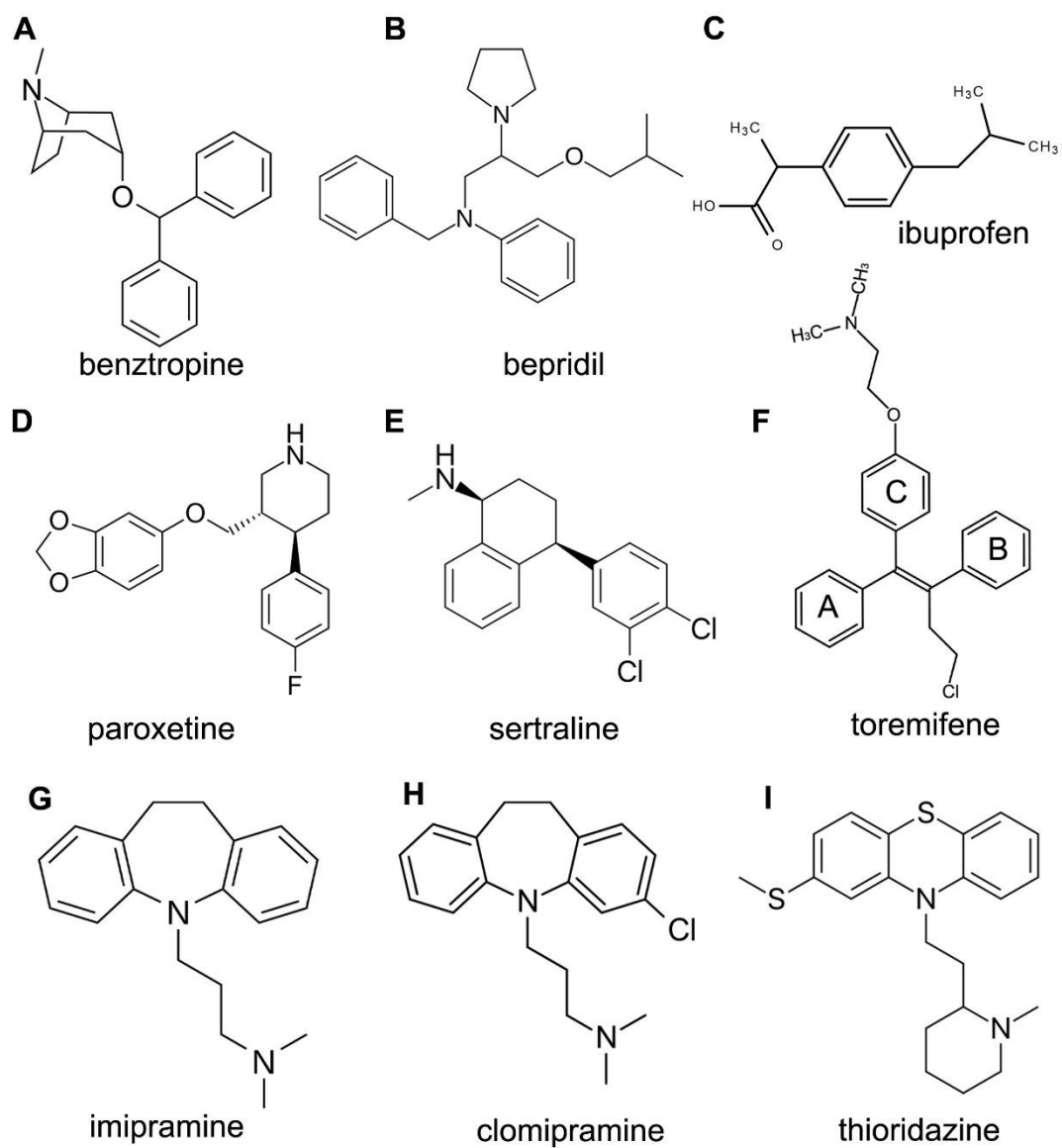


Figure S1. Chemical structures of approved drugs known to bind EBOV GP. Crystal structures of these nine drugs in complexes with EBOV GP have been previously reported.

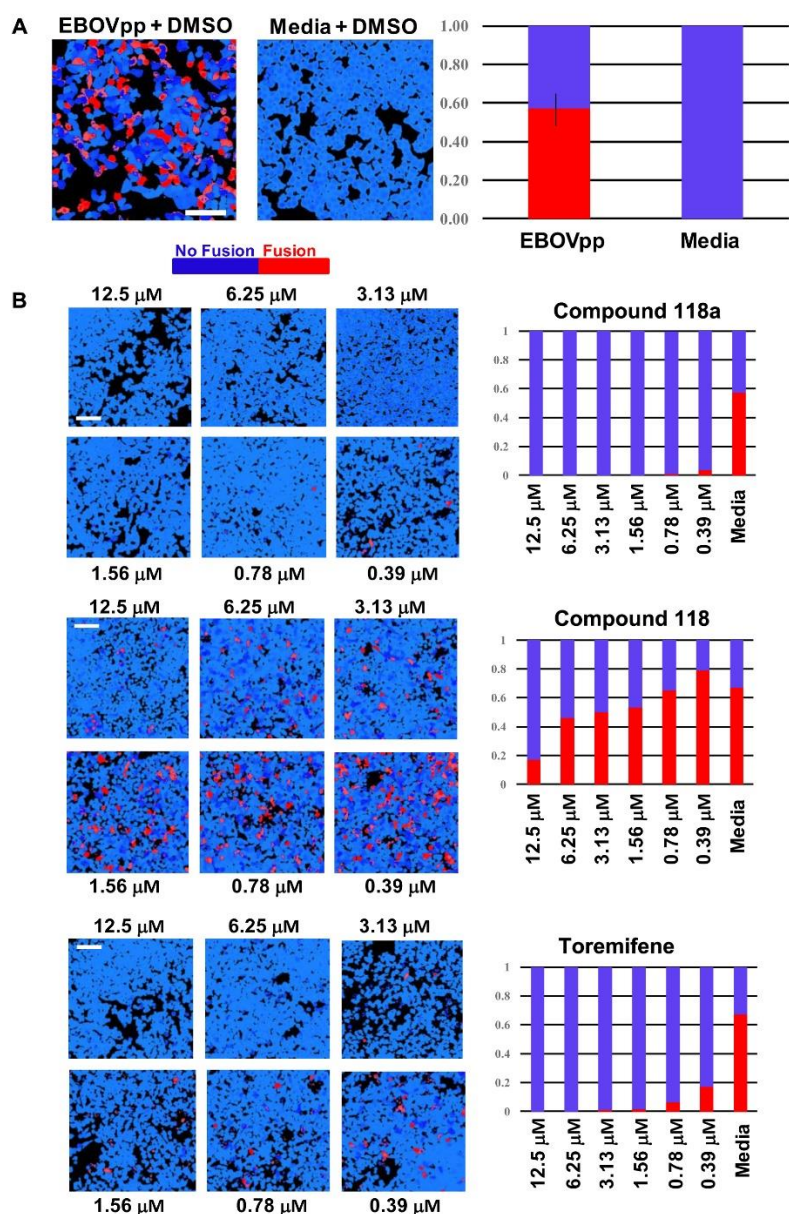


Figure S2. Compound **118a** is a potent and specific inhibitor of EBOVpp fusion in live cells. Fusion assays were carried out using a β -lactamase based reporter assay with different dilutions of **118a**, **118** and toremifene. (A) TZM-bl cells exposed to EBOVpp and media are shown, images were pseudocolored with red (fusion positive) and blue (fusion negative), error bar 10 μ M. The bar diagram show the total proportion of fusion positive cells (57% \pm 14) for TZM-bl cells exposed to EBOVpp (0.5 MOI) and Media (0%). (B) TZM-bl cells exposed to EBOVpp and **118a**, **118** and toremifene. Error bars showing the extent of fusion for each concentration of the corresponding compounds, error bar 20 μ M.

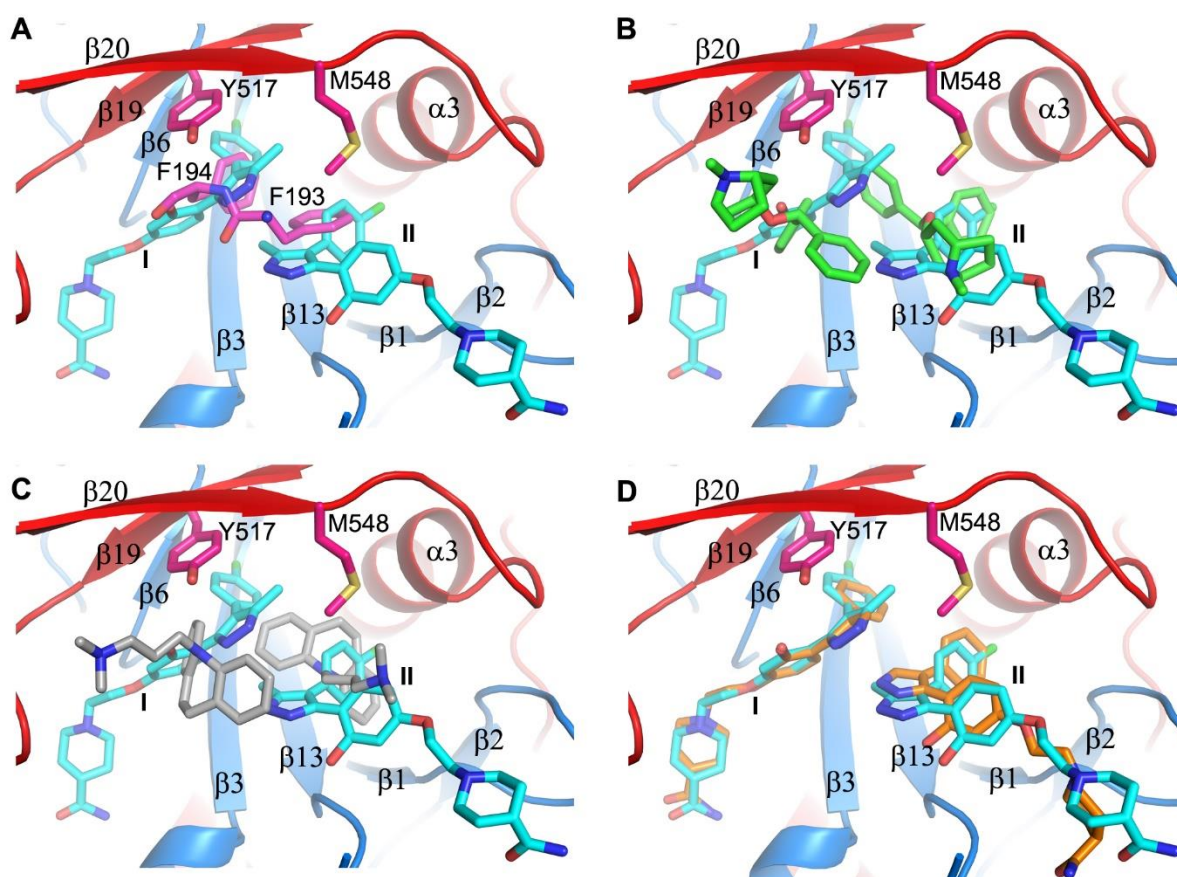


Figure S3. Comparison of binding modes of inhibitors that have two molecules bound in the GP cavity (these are labelled I and II, with I being the most internal). (A) The positions and orientations of the bound **118a** molecules are compared with F193 and F194 (magenta sticks) in the apo GP structure. (B to D) Overlays of bound benztropine (green sticks in (B)), imipramine (grey sticks in (C)) and compound **118** (orange sticks in (D)) molecules with **118a**. The superimpositions were done using protein C α atoms only. In all panels protein structure shown is GP-**118a** with main-chain as ribbons (GP1 in blue and GP2 in red). For clarity, only side-chains of Y517 and M548 (red sticks) are shown to mark the relative position of the bound inhibitors.

Table S1. Glide and QPLD docking scores of known EBOV GP binders. pIC₅₀ values are converted from the IC₅₀s obtained from virus infection assays in vero E6 cells (reference 10).

Drug	Glide Emodel (kcal/mol)	QPLD Emodel (kcal/mol)	pIC₅₀ (-log₁₀ M)
Toremifene	-67.733	-77.125	6.79
Bepriidil	-50.37	-55.746	5.29
Benztropine	-47.379	-53.416	5.09
Paroxetine	-45.973	-51.792	5.13
Imipramine	-61.331	-44.892	4.86
Clomipramine	-44.13	-44.209	4.94
Thioridazine	-38.835	-44.026	5.20
Sertraline	-28.265	-31.91	5.50

Table S2. Glide XP and QPLD docking scores of top 16 natural compounds from *In silico* screening. About 2.5 million compounds in the ZINC natural compound library from the Traditional Chinese Medicine database@Taiwan have been screened (reference 27). All the top 16 hits have passed PAINS-Remover (reference 28). The docking results for toremifene are included for comparison. Compounds selected for *in vitro* experiment are marked with an asterisk.

Chemical ID	Glide XP Score (kcal/mol)	Glide XP Emodel (kcal/mol)	QPLD Score (kcal/mol)	QPLD Emodel (kcal/mol)	Ligand efficiency score (kcal/mol)
NP-005129	-12.582	-79.021	-12.53	-75.31	-0.307
NP-016821	-12.465	-77.247	-11.67	-83.967	-0.296
NP-019105	-11.006	-58.284	-9.407	-64.866	-0.297
*ZINC32540717 (118)	-10.851	-70.036	-11.029	-73.801	-0.346
*ZINC00167626	-10.894	-37.646	-9.143	-47.781	-0.545
ZINC 32540716	-10.781	-71.098	-10.645	-84.044	-0.333
*ZINC09410451 (118a)	-9.210	-58.141	-9.210	-78.738	-0.288
NP-001471	-10.398	-54.738	-12.19	-70.721	-0.334
554127	-10.348	-37.327	-11.222	-59.189	-0.470
ZINC35464211	-10.127	-64.476	-9.887	-60.507	-0.316
*ZINC00407254	-9.869	-44.638	-9.65	-57.035	-0.539
*ZINC12893941	-9.791	-39.059	-9.187	-51.477	-0.484
*ZINC04772639	-9.341	-49.893	-9.283	-55.833	-0.404
ZINC05071941	-9.275	-41.034	-9.323	-49.881	-0.459
*ZINC11865143	-9.351	-40.886	-9.292	-53.402	-0.457
*ZINC00056827	-8.166	-42.016	-9.694	-42.819	-0.606
Toremifene	-8.413	-67.733	-8.670	-77.125	-0.299

Table S3. Coordinates of compound **118** docked with the EBOV GP (PDB ID 5JQ7).

HETATM	1	C	UNK	900	-47.581	11.604	-11.973	0.00	0.00	C
HETATM	2	C	UNK	900	-48.422	11.198	-10.946	0.00	0.00	C
HETATM	3	C	UNK	900	-48.109	11.514	-9.626	0.00	0.00	C
HETATM	4	C	UNK	900	-46.925	12.190	-9.316	0.00	0.00	C
HETATM	5	C	UNK	900	-46.088	12.603	-10.356	0.00	0.00	C
HETATM	6	C	UNK	900	-46.432	12.331	-11.680	0.00	0.00	C
HETATM	7	C	UNK	900	-46.599	12.453	-7.882	0.00	0.00	C
HETATM	8	C	UNK	900	-46.853	11.617	-6.802	0.00	0.00	C
HETATM	9	N	UNK	900	-46.388	12.274	-5.698	0.00	0.00	N
HETATM	10	H	UNK	900	-46.425	11.919	-4.752	0.00	0.00	H
HETATM	11	N	UNK	900	-45.860	13.464	-5.990	0.00	0.00	N
HETATM	12	C	UNK	900	-45.976	13.580	-7.297	0.00	0.00	C
HETATM	13	C	UNK	900	-45.498	14.811	-7.976	0.00	0.00	C
HETATM	14	C	UNK	900	-46.441	15.669	-8.542	0.00	0.00	C
HETATM	15	C	UNK	900	-46.024	16.798	-9.241	0.00	0.00	C
HETATM	16	C	UNK	900	-44.659	17.102	-9.352	0.00	0.00	C
HETATM	17	C	UNK	900	-43.725	16.241	-8.782	0.00	0.00	C
HETATM	18	C	UNK	900	-44.135	15.097	-8.106	0.00	0.00	C
HETATM	19	O	UNK	900	-43.182	14.258	-7.600	0.00	0.00	O
HETATM	20	O	UNK	900	-44.156	18.178	-10.051	0.00	0.00	O
HETATM	21	C	UNK	900	-44.909	19.385	-10.134	0.00	0.00	C
HETATM	22	C	UNK	900	-44.392	20.350	-9.065	0.00	0.00	C
HETATM	23	N	UNK	900	-45.309	21.593	-8.926	0.00	0.00	N1+
HETATM	24	C	UNK	900	-44.909	22.324	-7.626	0.00	0.00	C
HETATM	25	C	UNK	900	-45.863	23.502	-7.411	0.00	0.00	C
HETATM	26	C	UNK	900	-45.783	24.502	-8.585	0.00	0.00	C
HETATM	27	C	UNK	900	-46.079	23.784	-9.928	0.00	0.00	C
HETATM	28	C	UNK	900	-45.193	22.539	-10.150	0.00	0.00	C
HETATM	29	C	UNK	900	-46.719	25.713	-8.337	0.00	0.00	C
HETATM	30	O	UNK	900	-47.616	26.017	-9.117	0.00	0.00	O
HETATM	31	N	UNK	900	-46.509	26.453	-7.251	0.00	0.00	N
HETATM	32	H	UNK	900	-47.828	11.370	-12.999	0.00	0.00	H
HETATM	33	H	UNK	900	-49.309	10.629	-11.170	0.00	0.00	H
HETATM	34	H	UNK	900	-48.786	11.199	-8.851	0.00	0.00	H
HETATM	35	H	UNK	900	-45.171	13.134	-10.145	0.00	0.00	H
HETATM	36	H	UNK	900	-45.795	12.663	-12.487	0.00	0.00	H
HETATM	37	H	UNK	900	-47.309	10.640	-6.721	0.00	0.00	H
HETATM	38	H	UNK	900	-47.497	15.452	-8.457	0.00	0.00	H
HETATM	39	H	UNK	900	-46.783	17.408	-9.703	0.00	0.00	H
HETATM	40	H	UNK	900	-42.674	16.468	-8.887	0.00	0.00	H
HETATM	41	H	UNK	900	-42.295	14.518	-7.856	0.00	0.00	H
HETATM	42	H	UNK	900	-45.988	19.265	-10.061	0.00	0.00	H
HETATM	43	H	UNK	900	-44.728	19.809	-11.122	0.00	0.00	H
HETATM	44	H	UNK	900	-43.372	20.690	-9.265	0.00	0.00	H
HETATM	45	H	UNK	900	-44.399	19.850	-8.096	0.00	0.00	H
HETATM	46	H	UNK	900	-43.869	22.643	-7.731	0.00	0.00	H
HETATM	47	H	UNK	900	-44.974	21.602	-6.814	0.00	0.00	H
HETATM	48	H	UNK	900	-46.882	23.125	-7.290	0.00	0.00	H
HETATM	49	H	UNK	900	-45.598	23.986	-6.472	0.00	0.00	H
HETATM	50	H	UNK	900	-44.784	24.942	-8.621	0.00	0.00	H
HETATM	51	H	UNK	900	-47.129	23.485	-9.961	0.00	0.00	H
HETATM	52	H	UNK	900	-45.943	24.476	-10.762	0.00	0.00	H
HETATM	53	H	UNK	900	-44.138	22.800	-10.255	0.00	0.00	H
HETATM	54	H	UNK	900	-45.516	21.983	-11.030	0.00	0.00	H
HETATM	55	H	UNK	900	-47.111	27.244	-7.095	0.00	0.00	H
HETATM	56	H	UNK	900	-45.812	26.192	-6.555	0.00	0.00	H
HETATM	57	H	UNK	900	-46.283	21.294	-8.851	0.00	0.00	H

Table S4. Coordinates of compound **118a** docked with the EBOV GP (PDB ID 5JQ7).

HETATM	1	C1	UNK	900	-48.141	11.598	-12.609	0.00	0.00	C1
HETATM	2	O	UNK	900	-43.308	18.223	-9.872	0.00	0.00	O
HETATM	3	O	UNK	900	-42.932	24.629	-9.642	0.00	0.00	O
HETATM	4	O	UNK	900	-42.364	15.448	-6.205	0.00	0.00	O
HETATM	5	N	UNK	900	-45.309	21.404	-9.352	0.00	0.00	N1+
HETATM	6	N	UNK	900	-43.309	25.525	-7.651	0.00	0.00	N
HETATM	7	N	UNK	900	-43.853	13.093	-5.902	0.00	0.00	N
HETATM	8	N	UNK	900	-44.757	12.194	-5.259	0.00	0.00	N
HETATM	9	C	UNK	900	-45.008	22.406	-10.487	0.00	0.00	C
HETATM	10	C	UNK	900	-44.712	21.872	-8.002	0.00	0.00	C
HETATM	11	C	UNK	900	-45.162	24.297	-8.714	0.00	0.00	C
HETATM	12	C	UNK	900	-45.612	23.778	-10.109	0.00	0.00	C
HETATM	13	C	UNK	900	-45.361	23.215	-7.623	0.00	0.00	C
HETATM	14	C	UNK	900	-44.909	19.962	-9.749	0.00	0.00	C
HETATM	15	C	UNK	900	-43.706	24.831	-8.712	0.00	0.00	C
HETATM	16	C	UNK	900	-43.470	19.547	-9.383	0.00	0.00	C
HETATM	17	C	UNK	900	-43.661	17.233	-9.020	0.00	0.00	C
HETATM	18	C	UNK	900	-42.806	16.811	-8.063	0.00	0.00	C
HETATM	19	C	UNK	900	-44.908	16.539	-9.251	0.00	0.00	C
HETATM	20	C	UNK	900	-44.256	14.816	-7.610	0.00	0.00	C
HETATM	21	C	UNK	900	-44.496	13.606	-7.023	0.00	0.00	C
HETATM	22	C	UNK	900	-45.478	12.555	-7.300	0.00	0.00	C
HETATM	23	C	UNK	900	-43.074	15.654	-7.188	0.00	0.00	C
HETATM	24	C	UNK	900	-45.174	15.397	-8.603	0.00	0.00	C
HETATM	25	C	UNK	900	-45.655	11.772	-6.218	0.00	0.00	C
HETATM	26	C	UNK	900	-46.167	12.339	-8.602	0.00	0.00	C
HETATM	27	C	UNK	900	-46.504	10.559	-5.918	0.00	0.00	C
HETATM	28	C	UNK	900	-47.468	11.831	-8.672	0.00	0.00	C
HETATM	29	C	UNK	900	-45.492	12.603	-9.799	0.00	0.00	C
HETATM	30	C	UNK	900	-48.078	11.597	-9.902	0.00	0.00	C
HETATM	31	C	UNK	900	-46.109	12.398	-11.029	0.00	0.00	C
HETATM	32	C	UNK	900	-47.394	11.884	-11.081	0.00	0.00	C
HETATM	33	H	UNK	900	-46.323	21.379	-9.251	0.00	0.00	H
HETATM	34	H	UNK	900	-45.467	22.017	-11.397	0.00	0.00	H
HETATM	35	H	UNK	900	-43.924	22.428	-10.612	0.00	0.00	H
HETATM	36	H	UNK	900	-44.909	21.104	-7.255	0.00	0.00	H
HETATM	37	H	UNK	900	-43.633	21.965	-8.137	0.00	0.00	H
HETATM	38	H	UNK	900	-45.751	25.179	-8.458	0.00	0.00	H
HETATM	39	H	UNK	900	-45.382	24.508	-10.887	0.00	0.00	H
HETATM	40	H	UNK	900	-46.701	23.690	-10.114	0.00	0.00	H
HETATM	41	H	UNK	900	-44.982	23.554	-6.657	0.00	0.00	H
HETATM	42	H	UNK	900	-46.431	23.061	-7.469	0.00	0.00	H
HETATM	43	H	UNK	900	-45.657	19.348	-9.252	0.00	0.00	H
HETATM	44	H	UNK	900	-45.090	19.851	-10.818	0.00	0.00	H
HETATM	45	H	UNK	900	-42.744	20.191	-9.886	0.00	0.00	H
HETATM	46	H	UNK	900	-43.250	19.626	-8.314	0.00	0.00	H
HETATM	47	H	UNK	900	-42.367	25.882	-7.646	0.00	0.00	H
HETATM	48	H	UNK	900	-43.891	25.599	-6.818	0.00	0.00	H
HETATM	49	H	UNK	900	-41.864	17.329	-7.962	0.00	0.00	H
HETATM	50	H	UNK	900	-45.640	16.965	-9.920	0.00	0.00	H
HETATM	51	H	UNK	900	-46.150	14.960	-8.765	0.00	0.00	H
HETATM	52	H	UNK	900	-43.507	13.726	-5.190	0.00	0.00	H
HETATM	53	H	UNK	900	-44.186	11.527	-4.758	0.00	0.00	H
HETATM	54	H	UNK	900	-46.556	9.900	-6.785	0.00	0.00	H
HETATM	55	H	UNK	900	-47.513	10.867	-5.645	0.00	0.00	H
HETATM	56	H	UNK	900	-46.099	9.999	-5.079	0.00	0.00	H
HETATM	57	H	UNK	900	-48.030	11.629	-7.775	0.00	0.00	H
HETATM	58	H	UNK	900	-44.477	12.973	-9.783	0.00	0.00	H
HETATM	59	H	UNK	900	-49.074	11.184	-9.941	0.00	0.00	H
HETATM	60	H	UNK	900	-45.594	12.645	-11.943	0.00	0.00	H