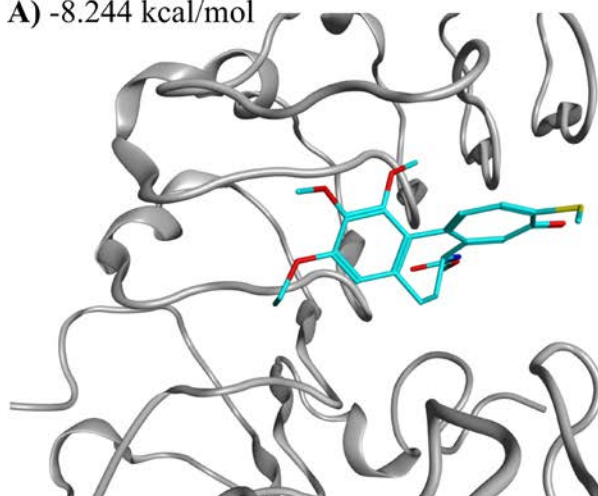
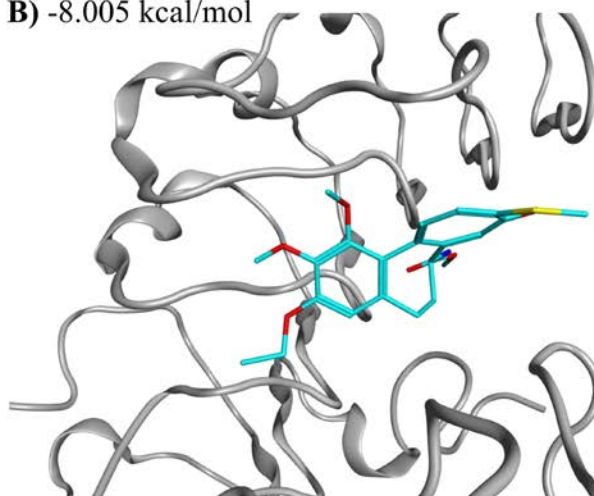


Supplementary Materials

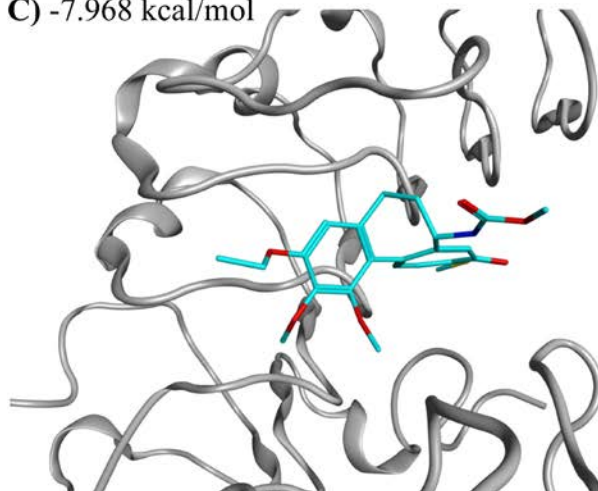
A) -8.244 kcal/mol



B) -8.005 kcal/mol



C) -7.968 kcal/mol



D) -7.912 kcal/mol

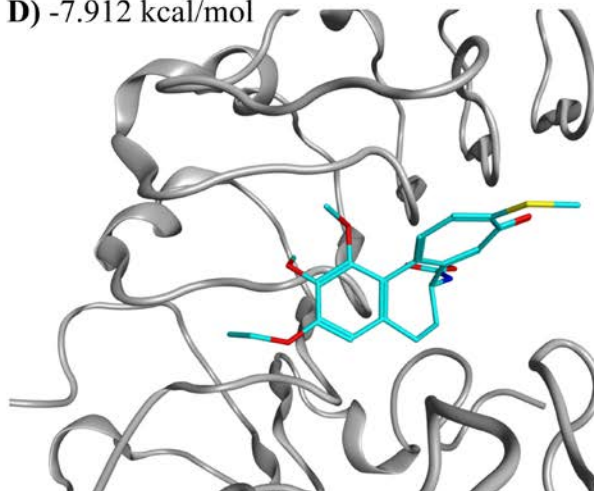


Figure S1: The additional four poses of CR42-24 obtained with molecular docking to the R-837 binding site of TLR7. Computed binding energies are also reported.

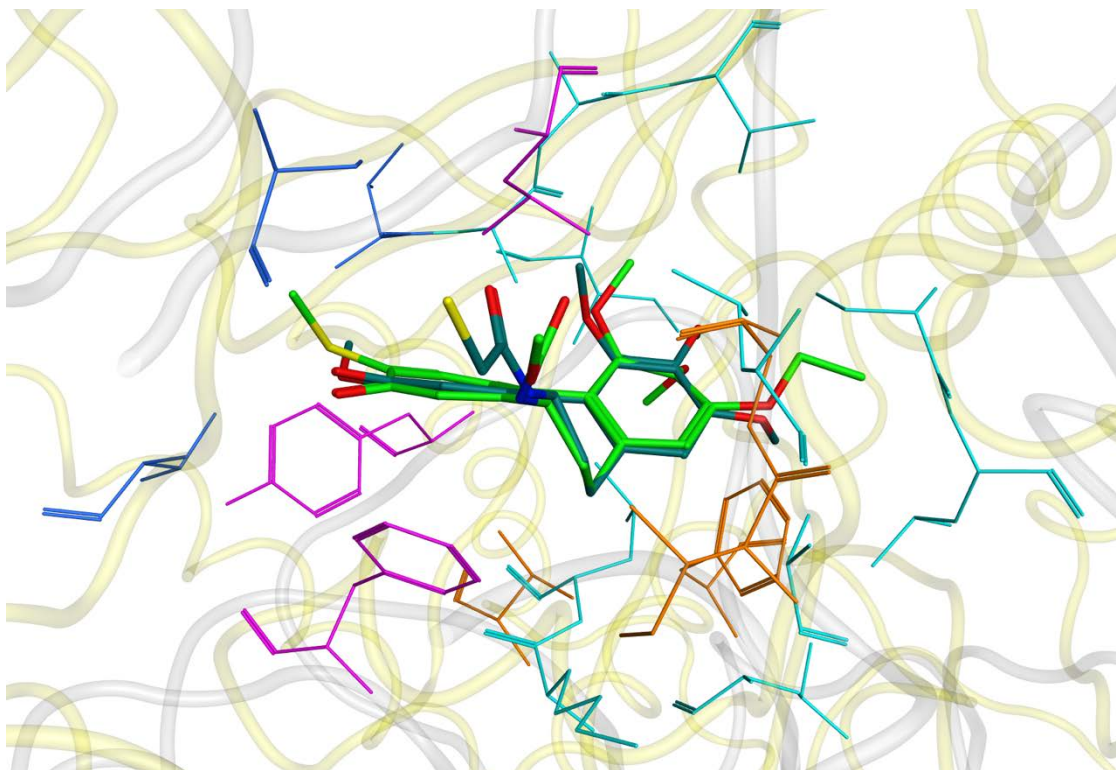


Figure S2: Superposition of the crystallographic pose of DAMA-colchicine (dark green carbons) in tubulin from pdb ID 1SA0 and the CR42-24 docking pose in TLR7 (carbons in light green). The first hydrophobic cleft of the tubulin binding site, interacting with the C ring of the DAMA-colchicine, is reported in blue sticks. The corresponding first hydrophobic zone of TLR7 is reported in purple sticks. The second hydrophobic zone is reported in cyan sticks for tubulin and orange sticks for TLR7. Tubulin backbone is reported in yellow ribbons, TLR7 backbone in grey ribbons.

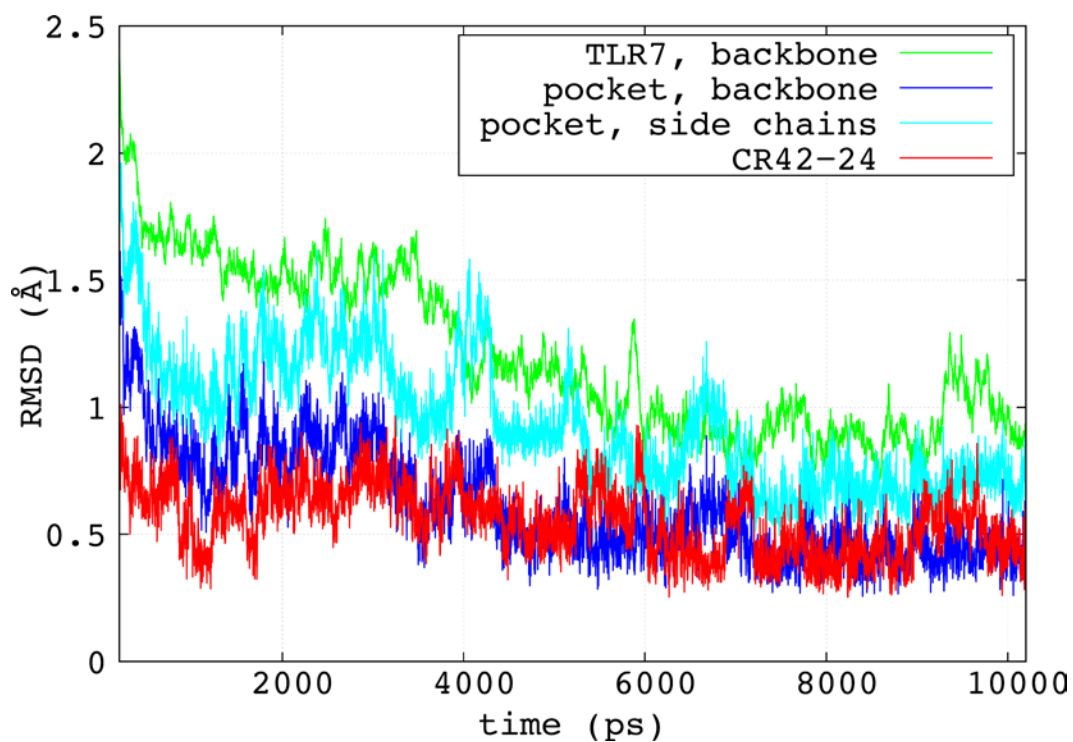


Figure S3: RMSD trends of the MD production simulation for the TLR7 backbone atoms (green), the backbone atoms of the binding pocket (blue), the side chain atoms of the binding pocket (cyan) and the heavy atoms of CR42-24 (red), calculated using the average structure of the last 5 ns as reference. The highest fluctuations are observed for the side chain atoms of the binding pocket, as they adapted to the bound conformation upon ligand binding.

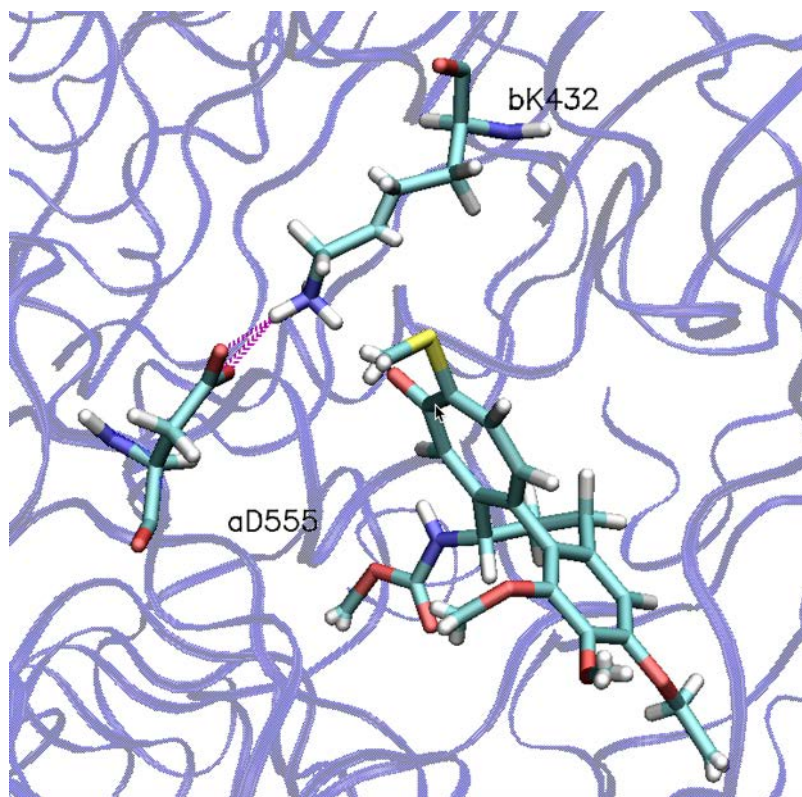


Figure S4: Salt bridge (in purple) established between aD555 and bK432, keeping the former far from the amine of the acetamide group of CR42-24.

Table S1: Cell lines in the Oncolines™ panel used in this study and their genetic status. Indicated in the second column are the genes that are either mutated, part of a translocation or that have altered copy numbers (italic) in that cell line.

Cell line	Cancer genes
5637	ERBB2, RB1, TP53
769-P	No coding alterations found in 38 cancer driver genes
786-O	<i>CDKN2A</i> , PTEN, TP53
A-172	<i>CDKN2A</i> , <i>PTEN</i>
A-204	ABL-driven
A-427	<i>CDKN2A</i> , CTNNB1, KRAS
A-498	<i>CDKN2A</i> , SETD2
A-549	<i>CDKN2A</i> , KRAS, SMARCA4, STK11
A-704	PBRM1, TP53
A375	BRAF, <i>CDKN2A</i> , <i>CDKN2A</i>
A388	<i>EGFR</i> , NOTCH1, PIK3CA, TP53
ACHN	<i>CDKN2A</i> , PBRM1
AN3 CA	CCND1, FBXW7, NSD1, PIK3R1, PTEN, SETD2, TLR7, TP53, ZFHX3
AsPC-1	<i>CDKN2A</i> , FBXW7, KRAS, TP53
AU-565	<i>ERBB2</i> , MYC, SMAD4, TLR7, TP53

BT-20	<i>CDKN2A</i> , <i>EGFR</i> , <i>PIK3CA</i> , <i>RB1</i> , <i>SPEN</i> , <i>TP53</i>
BT-549	<i>PTEN</i> , <i>RB1</i>
BxPC-3	<i>BRAF</i> , <i>CDKN2A</i> , <i>EP300</i> , <i>SMAD4</i> , <i>TP53</i>
C-33 A	<i>ARID1A</i> , <i>FBXW7</i> , <i>PIK3CA</i> , <i>PTEN</i> , <i>SMARCA4</i> , <i>TP53</i> , <i>ZFHX3</i>
CAL 27	<i>CDKN2A</i> , <i>SMAD4</i> , <i>TP53</i>
CCF-STTG1	<i>PTEN</i>
CCRF-CEM	<i>CCND1</i> , <i>CDKN2A</i> , <i>FBXW7</i> , <i>KRAS</i> , <i>NRAS</i> , <i>NSD1</i> , <i>PTEN</i> , <i>TLR7</i> , <i>TP53</i>
COLO 205	<i>APC</i> , <i>BRAF</i> , <i>SMAD4</i>
COLO 829	<i>BRAF</i> , <i>CDKN2A</i> , <i>PTEN</i>
Daoy	<i>CDKN2A</i> , <i>NF1</i> , <i>PIK3R1</i> , <i>TP53</i>
DB	<i>TP53</i>
DLD-1	<i>APC</i> , <i>EP300</i> , <i>KRAS</i> , <i>NCOR1</i> , <i>PIK3CA</i> , <i>SMARCA4</i> , <i>TP53</i>
DoTc2 4510	<i>BRCA2</i> , <i>FAT1</i> , <i>NSD1</i>
DU 145	<i>CDKN2A</i> , <i>CREBBP</i> , <i>FAT1</i> , <i>KRAS</i> , <i>RB1</i> , <i>STK11</i> , <i>XIRP2</i>
DU4475	<i>APC</i> , <i>BRAF</i> , <i>RB1</i>
ES-2	<i>BRAF</i> , <i>CCND1</i> , <i>TP53</i>
FaDu	<i>CCND1</i> , <i>FAT1</i> , <i>SMAD4</i> , <i>TP53</i>
G-361	<i>BRAF</i> , <i>CDKN2A</i> , <i>STK11</i>
HCT-15	<i>APC</i> , <i>BRCA2</i> , <i>EP300</i> , <i>KRAS</i> , <i>NCOR1</i> , <i>PIK3CA</i> , <i>TP53</i> , <i>ZFHX3</i>
HCT 116	<i>BRCA2</i> , <i>CDKN2A</i> , <i>CHD4</i> , <i>CTNNB1</i> , <i>EP300</i> , <i>KRAS</i> , <i>NCOR1</i> , <i>NF1</i> , <i>PIK3CA</i> , <i>SMARCA4</i> , <i>XIRP2</i>
HL-60	<i>CDKN2A</i> , <i>MYC</i> , <i>NRAS</i>
Hs 578T	<i>MYC</i> , <i>PIK3R1</i> , <i>TP53</i>
Hs 746T	<i>TP53</i>
Hs 766T	<i>ARID1A</i> , <i>KRAS</i> , <i>LRP1B</i> , <i>SMAD4</i>
HT	<i>EP300</i> , <i>SPEN</i> , <i>TP53</i>
HT-1080	<i>CDKN2A</i> , <i>NRAS</i>
HuTu 80	<i>CTNNB1</i> , <i>SMARCA4</i>
J82	<i>ERBB2</i> , <i>PTEN</i>
JAR	No coding alterations found in 38 cancer driver genes
Jurkat E6.1	<i>APC</i> , <i>ARID1A</i> , <i>CDKN2A</i> , <i>CREBBP</i> , <i>FAT1</i> , <i>FBXW7</i> , <i>PTEN</i> , <i>SMARCA4</i> , <i>TP53</i>
K-562	<i>ABL</i> -driven, <i>CDKN2A</i> , <i>TP53</i>
KATO III	<i>NOTCH1</i>
KG-1	No coding alterations found in 38 cancer driver genes
KLE	<i>CCNE1</i> , <i>FBXW7</i> , <i>KRAS</i> , <i>TP53</i>
KU812	<i>ABL</i> -driven, <i>TP53</i>
LNCaP FGC	<i>ATM</i> , <i>BRCA2</i> , <i>PIK3R1</i> , <i>PTEN</i> , <i>SETD2</i> , <i>XIRP2</i>
LoVo	<i>APC</i> , <i>ARID1A</i> , <i>FBXW7</i> , <i>KRAS</i> , <i>SPEN</i>
LS 174T	<i>ARID1A</i> , <i>CTNNB1</i> , <i>KRAS</i> , <i>PBRM1</i> , <i>PIK3CA</i> , <i>TLR7</i> , <i>ZFHX3</i>
LS411N	<i>APC</i> , <i>ARID1A</i> , <i>BRAF</i> , <i>EP300</i> , <i>FBXW7</i> , <i>PTEN</i> , <i>TP53</i> , <i>XIRP2</i>
MCF7	<i>CDKN2A</i> , <i>EP300</i> , <i>PIK3CA</i>
MeWo	<i>CDKN2A</i> , <i>NF1</i> , <i>TP53</i> , <i>XIRP2</i>

MG-63	<i>CDKN2A</i> , <i>MYC</i>
MIA PaCa-2	<i>CDKN2A</i> , <i>KRAS</i> , <i>TP53</i>
MOLT-4	<i>CDKN2A</i> , <i>CHD4</i> , <i>EP300</i> , <i>NOTCH1</i> , <i>NRAS</i> , <i>PTEN</i> , <i>SMARCA4</i> , <i>TP53</i>
NCCIT	<i>PTEN</i> , <i>TP53</i>
NCI-H460	<i>ARID1A</i> , <i>CDKN2A</i> , <i>KRAS</i> , <i>LRP1B</i> , <i>MYC</i> , <i>PIK3CA</i> , <i>STK11</i>
NCI-H661	<i>CCNE1</i> , <i>TP53</i>
NCI-H82	<i>MYC</i>
OVCAR-3	<i>CCNE1</i> , <i>LRP1B</i> , <i>TP53</i>
PA-1	<i>NRAS</i>
PC-3	<i>MYC</i> , <i>PTEN</i> , <i>TP53</i>
PFSK-1	<i>TP53</i>
RD	<i>NF1</i> , <i>NRAS</i> , <i>TP53</i>
RKO	<i>ARID1A</i> , <i>BRAF</i> , <i>BRCA2</i> , <i>EP300</i> , <i>FAT1</i> , <i>NCOR1</i> , <i>NF1</i> , <i>NSD1</i> , <i>PIK3CA</i> , <i>ZFH3</i>
RL	<i>EP300</i>
RL95-2	<i>ARID1A</i> , <i>ATM</i> , <i>BRCA2</i> , <i>EP300</i> , <i>PIK3R1</i> , <i>PTEN</i> , <i>SMARCA4</i> , <i>SPEN</i> , <i>TP53</i> , <i>ZFH3</i>
RPMI-7951	<i>BRAF</i> , <i>PTEN</i> , <i>TP53</i>
RS4-11	<i>CDKN2A</i> , <i>SMARCA4</i>
RT4	<i>CDKN2A</i> , <i>SMAD4</i>
SHP-77	<i>KRAS</i> , <i>TP53</i>
SJCRH30	<i>TP53</i>
SK-N-AS	<i>NRAS</i> , <i>TP53</i> , <i>XIRP2</i>
SK-N-FI	<i>NF1</i> , <i>TP53</i>
SNU-5	<i>ARID1A</i> , <i>CDKN2A</i>
SNU-C2B	<i>ERBB2</i> , <i>KRAS</i> , <i>SMARCA4</i> , <i>TP53</i>
SR	<i>CDKN2A</i> , <i>ZFH3</i>
SU-DHL-1	<i>CREBBP</i> , <i>TP53</i>
SU-DHL-6	<i>CREBBP</i> , <i>EP300</i> , <i>TP53</i>
SUP-T1	<i>CDKN2A</i> , <i>CHD4</i> , <i>PBRM1</i> , <i>PIK3CA</i> , <i>TP53</i>
SW48	<i>BRCA2</i> , <i>CTNNB1</i> , <i>EGFR</i> , <i>EP300</i> , <i>FBXW7</i> , <i>NCOR1</i> , <i>SETD2</i> , <i>ZFH3</i>
SW480	<i>APC</i> , <i>KRAS</i> , <i>TP53</i>
SW620	<i>APC</i> , <i>EP300</i> , <i>KRAS</i> , <i>KRAS</i> , <i>MYC</i> , <i>TP53</i>
SW626	<i>APC</i> , <i>KRAS</i> , <i>SMAD4</i>
SW837	<i>APC</i> , <i>FBXW7</i> , <i>KRAS</i> , <i>TP53</i>
SW872	<i>BRAF</i> , <i>CDKN2A</i> , <i>PTEN</i>
SW900	<i>CDKN2A</i> , <i>FAT1</i> , <i>KRAS</i> , <i>KRAS</i> , <i>NF1</i> , <i>NSD1</i> , <i>TP53</i>
SW948	<i>APC</i> , <i>ATM</i> , <i>KRAS</i> , <i>PIK3CA</i> , <i>SMAD4</i> , <i>TP53</i>
SW982	<i>BRAF</i> , <i>CDKN2A</i>
T24	<i>EP300</i> , <i>FAT1</i> , <i>TP53</i>
T98G	<i>CDKN2A</i> , <i>TP53</i>
TCCSUP	<i>ARID1A</i> , <i>PIK3CA</i> , <i>RB1</i> , <i>TP53</i>
THP-1	<i>ARID1A</i> , <i>CDKN2A</i> , <i>NRAS</i> , <i>PTEN</i> , <i>TP53</i>

TT	TP53
U-118 MG	<i>CDNK2A</i> , TP53
U-2 OS	<i>LRP1B</i>
U-87 MG	<i>CDNK2A</i> , NF1
VA-ES-BJ	<i>CDNK2A</i>