

Figure S1. EMASAs gels for  $\lambda$ -CI with A) 50 nM LOL1, B) 50 nM LOL2 and C) 50 nM LOL3. All the binding data were measured using ImageJ and curve fitted using python.

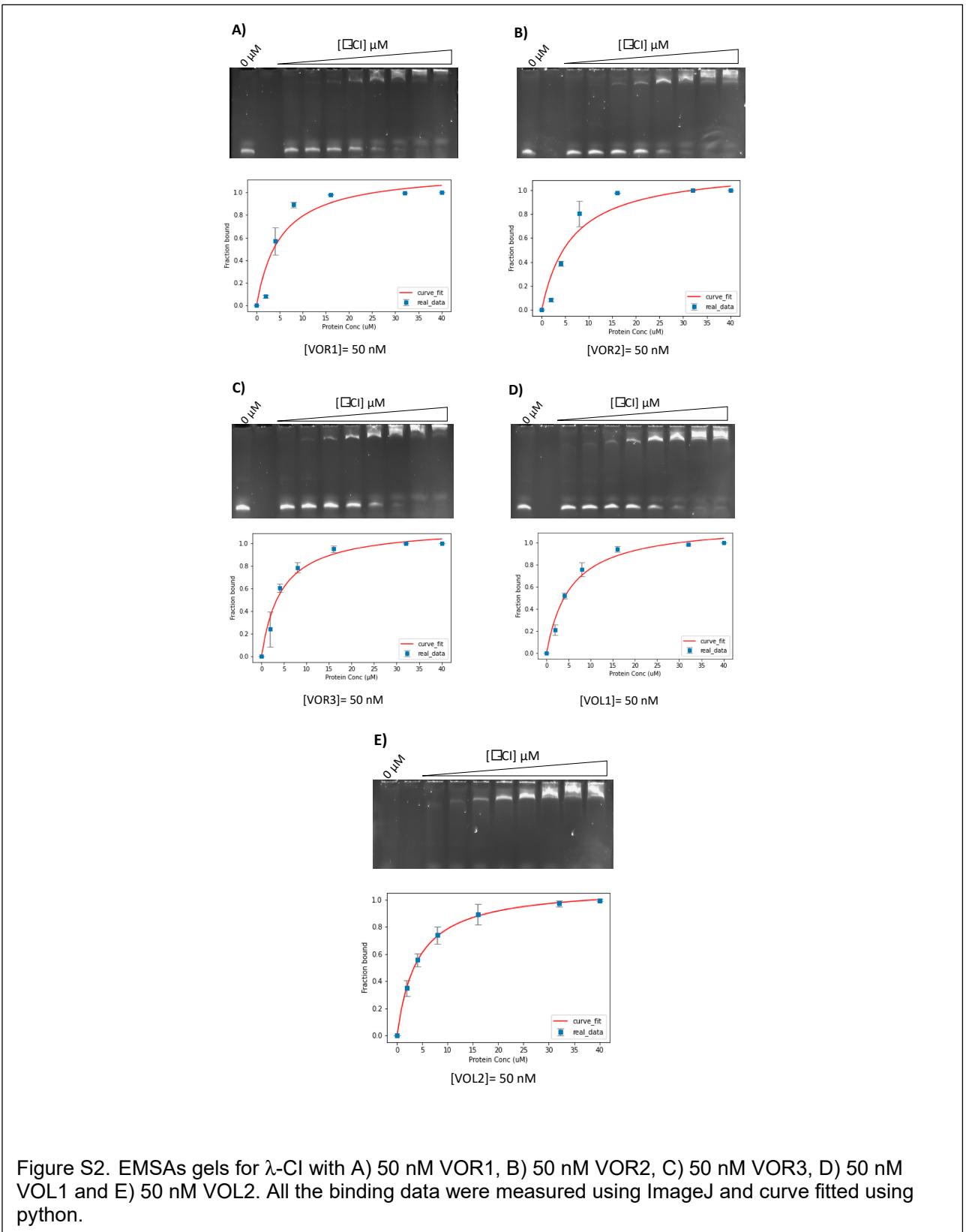


Figure S2. EMASAs gels for  $\lambda$ -Cl with A) 50 nM VOR1, B) 50 nM VOR2, C) 50 nM VOR3, D) 50 nM VOL1 and E) 50 nM VOL2. All the binding data were measured using ImageJ and curve fitted using python.

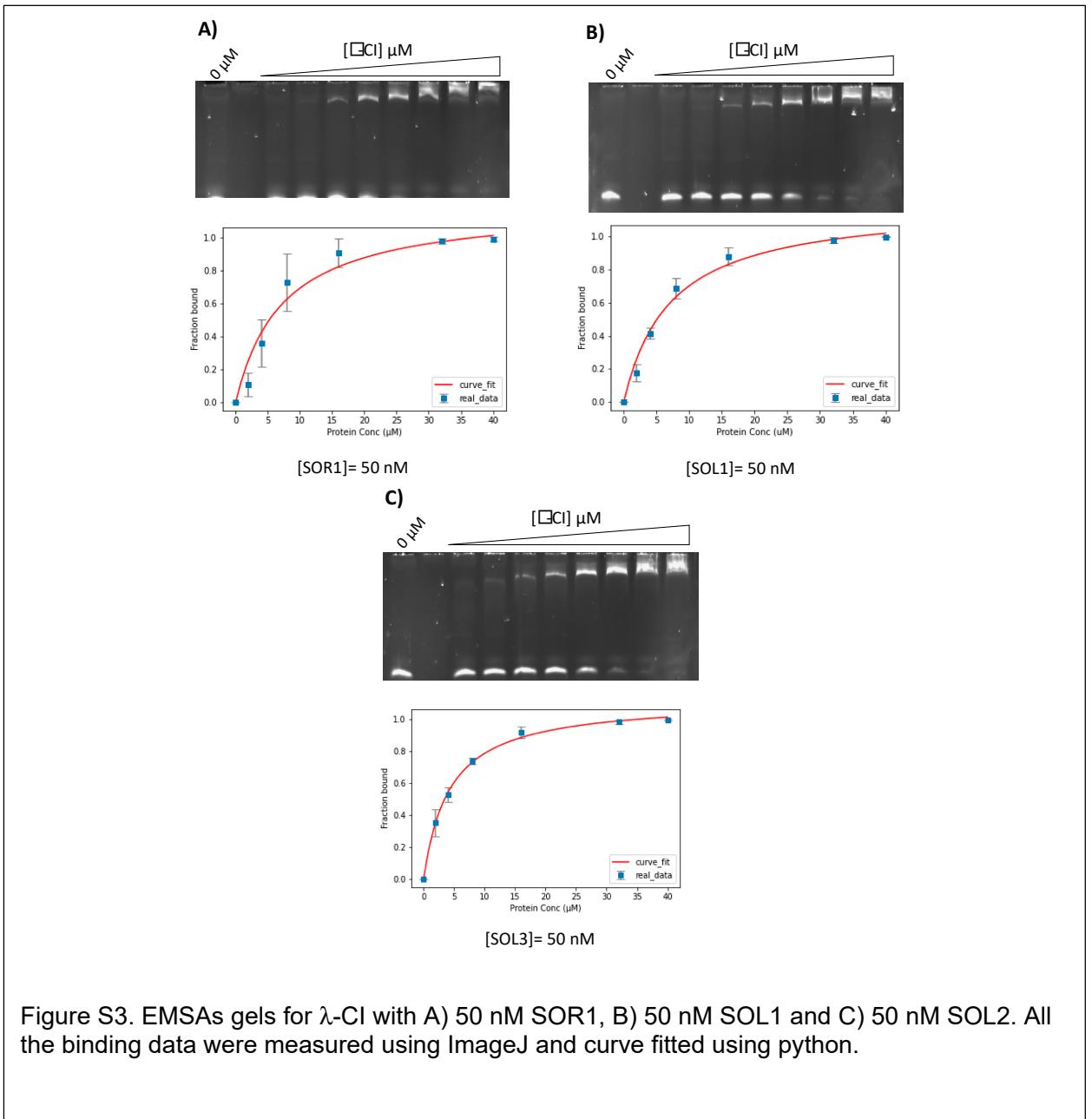


Table S1. The DNA sequences of the three phages' binding sites in black. Overhang sequences are in green.

Phage Strain	Binding site	DNA Sequence		
Lambda ( $\lambda$ )	OR1	GGTTATTATGG	TACCTCTGGCGGTGATA	TGCAAGTGC
	OR2	GGTTATTATGG	TAACACCGTGCCTGTTG	TGCAAGTGC
	OR3	GGTTATTATGG	TATCACCGCAAGGGATA	TGCAAGTGC
	OL1	GGTTATTATGG	TATCACCGCCAGTGGTA	TGCAAGTGC
	OL2	GGTTATTATGG	CAACACCGCCAGAGATA	TGCAAGTGC
	OL3	GGTTATTATGG	TATCACCGCAGATGGTT	TGCAAGTGC
VT2-SA	OR1	GGTTATTATGG	AAGTACATATTGTATTT	TGCAAGTGC
	OR2	GGTTATTATGG	ATGCACAATGTGTATTT	TGCAAGTGC
	OR3	GGTTATTATGG	TATTACGCTCTGTACTG	TGCAAGTGC
	OL1	GGTTATTATGG	TAGTACGATATGTACTT	TGCAAGTGC
	OL2	GGTTATTATGG	CAATACAATTGTTCTA	TGCAAGTGC
	OL3	GGTTATTATGG	TACAACGCTTGTTGTATTT	TGCAAGTGC
Stx2I	OR1	GGTTATTATGG	TTTCAGTTCAACCATAA	TGCAAGTGC
	OR2	GGTTATTATGG	ATGAGTACGATACTAAA	TGCAAGTGC
	OR3	GGTTATTATGG	GGCGAATTCTTGCGC	TGCAAGTGC
	OL1	GGTTATTATGG	CAATGATGATTCATGC	TGCAAGTGC
	OL2	GGTTATTATGG	TGTAATATTTCTTACA	TGCAAGTGC
	OL3	GGTTATTATGG	GCTTCTTGAATAACAGC	TGCAAGTGC

**Table S2. The binding data for  $\lambda$ -CI with LOR1**

CI conc (uM)	trials				average	SD
	1	2	3	4		
0	0	0	0	0	0	0
0.4	0.64215724	0.63714968	0.36051448		0.54660713	0.16118041
0.8	0.81082533	0.80156855	0.62118743	0.51932969	0.68822775	0.14247487
2	0.95705854	0.93095137	0.90621801	0.86012149	0.91358735	0.04124786
4	0.97754402	0.9889962	0.96971053	0.96144871	0.97442487	0.01172829
8	0.98305379	0.99586596	0.99014985	0.98752654	0.98914903	0.00535111
16	0.99659658		1	0.99738435	0.99077755	0.99618962
32		1	1	1	1	1
40		1	1	1	1	0

**Table S3. The binding data for  $\lambda$ -CI with LOR2**

CI conc (uM)	trials						average	SD
	1	2	3	4	5	6		
0	0	0	0	0	0	0	0	0
0.4	0.10279775	0.08526245	0.14515904	0.12326493	0.13748501	0.11015927	0.11735474	0.02239122
0.8	0.32027149	0.41100334	0.20627587	0.19479572	0.47697192	0.19163987	0.3001597	0.12300053
2	0.73205124	0.73126619	0.56563037	0.71177839	0.65448723	0.67827907	0.67891541	0.06336345
4	0.90328677	0.93873148	0.88492019	0.78289039	0.6949698	0.88846965	0.84887805	0.09158886
8	0.9662105	0.99088871	0.94249541	0.9235707	0.98185285	0.97730478	0.96372049	0.02576873
16	0.99545487	0.99892665	0.95906681	0.97708132	0.99000016	0.99346508	0.98566582	0.01504986
32		1	1	1	1	1	1	0
40		1	1	1	1	1	1	0

**Table S4. The binding data for  $\lambda$ -Cl with LOR3**

Cl conc ( $\mu$ M)	trials				average	SD
	1	2	3	4		
0	0	0	0	0	0	0
0.4	0.01074645	0.02371539	0.0429321	0.03378716	0.02779528	0.01381224
0.8	0.08518358	0.06505735	0.083499	0.04957068	0.07082765	0.01685043
2	0.37680368	0.30580252	0.31940157	0.35947863	0.3403716	0.03330234
4	0.86521344	0.81709052	0.84190748	0.86737039	0.84789546	0.02355126
8	0.99583305	0.96966002	0.97585521	0.97558424	0.97923313	0.01142988
16	1	1	1	1	1	0
32	1	1	1	1	1	0
40	1	1	1	1	1	0

**Table S5. The binding data for  $\lambda$ -Cl with LOL1**

Cl conc ( $\mu$ M)	trials				average	SD
	1	2	3	4		
0	0	0	0	0	0	0
0.4	0.74275585	0.73421081	0.66133934	0.64591473	0.69605518	0.04951787
0.8	0.81615543	0.8250863	0.68089326	0.70580733	0.75698558	0.07426985
2	0.8781074	0.91176577	0.83248591	0.82143733	0.8609491	0.04182533
4	0.97661281	0.93991034	0.88264283	0.92777798	0.93173599	0.03875802
8	0.98323567	0.98496397	0.96620679	0.9741487	0.97713878	0.00869587
16	1	1	1	0.99544462	0.99886115	0.00227769
32	1	1	1	1	1	0
40	1	1	1	1	1	0

**Table S6. The binding data for  $\lambda$ -Cl with LOL2**

Cl conc (uM)	trials				average	SD
	1	2	3	4		
0	0	0	0	0	0	0
0.4	0.23799133	0.22091393	0.16967566	0.11503125	0.18590304	0.05545295
0.8	0.33608317	0.33083302	0.3667847	0.24084449	0.31863635	0.05423095
2	0.57557648	0.69539164	0.58713738	0.7317327	0.64745955	0.07790034
4	0.76388376	0.8240741	0.86713628	0.81002573	0.81627997	0.04255033
8	0.9615783	0.92528135	0.93196017	0.96090744	0.94493182	0.01903269
16	0.99231078	0.9777523	0.98439195	1	0.98861376	0.00964554
32	1	1	1	1	1	0
40	1	1	1	1	1	0

**Table S7. The binding data for  $\lambda$ -Cl with LOL3**

Cl conc (uM)	trials				average	SD
	1	2	3	4		
0	0	0	0	0	0	0
0.4	0.30476998	0.30763117	0.19783118	0.23669997	0.26173307	0.05375537
0.8	0.41190544	0.43094454	0.20352838	0.45163638	0.37450368	0.11513249
2	0.64275015	0.72500806	0.67464069	0.71664094	0.68975996	0.03831244
4	0.76551701	0.74636287	0.7682345	0.86030706	0.78510536	0.05107056
8	0.8120679	0.87721704	0.94002501	0.93063476	0.88998618	0.05885151
16	0.94658288	0.96583899	0.98879278	0.97235494	0.9683924	0.0174552
32	1	0.98430365	1	1	0.99607591	0.00784817
40	1	1	1	1	1	0

**Table S8. The binding data for  $\lambda$ -Cl with VOR1**

Cl conc ( $\mu$ M)	trials				average	SD
	1	2	3	4		
0	0	0	0	0	0	0
0.4	0.01402669	0.01181229	0.00889833	0.01088209	0.01140485	0.00212885
0.8	0.02997044	0.03306916	0.03822149	0.02695978	0.03205522	0.00480835
2	0.07618689	0.09965214	0.06991182	0.07678799	0.08063471	0.01305403
4	0.53991689	0.4101596	0.67461383	0.64890473	0.56839876	0.12057322
8	0.92098892	0.86184614	0.89215216	0.89052237	0.8913774	0.02415415
16	0.97992512	0.98719576	0.98117429	0.97780621	0.98152534	0.0040278
32	0.99667168	0.99487729	0.99319429	0.996975	0.99542956	0.0017543
40	1	1	1	1	1	0

**Table S9. The binding data for  $\lambda$ -Cl with VOR2**

Cl conc ( $\mu$ M)	trials				average	SD
	1	2	3	4		
0	0	0	0	0	0	0
0.4	0	0	0	0	0	0
0.8	0.02057431	0.02229398	0.01539598	0.00478789	0.01576304	0.00788226
2	0.08824867	0.06646483	0.09858852	0.0820022	0.08382606	0.01344406
4	0.41114593	0.36050232	0.39504381	0.37904633	0.3864346	0.02169361
8	0.64663339	0.87280696	0.85236506	0.85185881	0.80591605	0.10663584
16	0.97295743	0.96887687	0.98230809	0.97825865	0.97560026	0.00589493
32		0.99704761	0.99495211	0.97481748	0.98893907	0.01227445
40		1	1	0.98318958	0.99439653	0.0097055

**Table S10.** The binding data for  $\lambda$ -Cl with VOR3

Cl conc (uM)	trials				average	SD
	1	2	3	4		
0	0	0	0	0	0	0
0.4	0	0	0	0	0	0
0.8	0.04886266	0.02413568	0.04429557	0.04827195	0.04139147	0.01168126
2	0.11815051	0.11920422	0.44086945	0.28888996	0.24177853	0.15509678
4	0.55060737	0.6220591	0.62070109	0.63122297	0.60614763	0.03732055
8	0.74041737	0.77285962	0.84720425	0.7840791	0.78614008	0.04472143
16	0.96392937	0.91393197	0.97830775	0.96667963	0.95571218	0.02854209
32	0.95861666	0.95324985	0.99652787	0.99295673	0.97533778	0.02256043
40		1	1	1	1	0

**Table S11.** The binding data for  $\lambda$ -Cl with VOL1

Cl conc (uM)	trials				average	SD
	1	2	3	4		
0	0	0	0	0	0	0
0.4	0.03057005	0.05819495	0.02850897	0.02397533	0.03531232	0.01550179
0.8	0.07160596	0.10721945	0.09607109	0.07593393	0.08770761	0.01681807
2	0.17766816	0.26756022	0.22409557	0.17032692	0.20991272	0.04520743
4	0.48212511	0.51637814	0.53282156	0.54807347	0.51984957	0.02828455
8	0.70521508	0.8244549	0.70404005	0.80055965	0.75856742	0.06304552
16	0.94163359	0.97338456	0.8784265	0.93045004	0.93097367	0.03947065
32	0.98056496	0.98508735	0.97074512	0.98298215	0.9798449	0.00634166
40		1	0.98956712	0.98955347	0.9930402	0.00602737

**Table S12.** The binding data for  $\lambda$ -Cl with VOL2

Cl conc (uM)	trials				average	SD
	1	2	3	4		
0	0	0	0	0	0	0
0.4	0.08000414	0.08056843	0.11537065	0.06226504	0.08455207	0.02223398
0.8	0.14845294	0.11181472	0.14178042	0.09284196	0.12372251	0.02603258
2	0.40916511	0.26656848	0.36058305	0.35906424	0.34884522	0.05958231
4	0.52267241	0.51340244	0.61217308	0.58201066	0.55756465	0.04742523
8	0.66298645	0.71520899	0.81242318	0.76559397	0.73905315	0.06440047
16	0.83293489	0.82922647	0.93710878	0.97912333	0.89459837	0.07533807
32	0.94858903	0.95762087	0.9848738	1	0.97277093	0.02382009
40		0.98826482		1	0.99413241	0.00829802

**Table S13.** The binding data for  $\lambda$ -Cl with VOL3

Cl conc (uM)	trials				average	SD
	1	2	3	4		
0	0	0	0	0	0	0
0.4	0.11351357	0.11800529	0.09511538	0.03921621	0.09146261	0.03621136
0.8	0.1928493	0.22432755	0.22305369	0.17858345	0.2047035	0.02269076
2	0.40941596	0.41946489	0.37259066	0.3713141	0.3931964	0.02487662
4	0.5644268	0.55823846	0.56719082	0.54784369	0.55942494	0.00858022
8	0.6946271	0.75325841	0.72598929	0.75732082	0.73279891	0.02900214
16	0.88021251	0.88280774	0.84857208	0.88110808	0.8731751	0.01643729
32	1	0.9538181	0.97302409	0.98232416	0.97729159	0.01923725
40	1	0.97942233	1	0.98440898	0.99095783	0.01063762

**Table S14. The binding data for  $\lambda$ -Cl with SOR1**

Cl conc ( $\mu$ M)	trials				average	SD
	1	2	3	4		
0	0	0	0	0	0	0
0.4	0	0	0	0.01947984	0.00486996	0.00973992
0.8	0.01304342	0.04837218	0.0538849	0.0574276	0.04318203	0.02043495
2	0.0402713	0.08397515	0.20851674	0.10078474	0.10838698	0.07145965
4	0.3027054	0.19663327	0.52325953	0.41840329	0.36025038	0.14146345
8	0.5000063	0.68899767	0.89545658	0.82701548	0.72786901	0.17449739
16	0.82689133	0.83896227	0.98386784	0.98746369	0.90929628	0.08833367
32	0.9617668	0.97551262	0.99444645	0.99496582	0.98167292	0.01606307
40	0.9779765	1	1	1	0.99449412	0.01101175

**Table S15. The binding data for  $\lambda$ -Cl with SOR2**

Cl conc ( $\mu$ M)	trials				average	SD
	1	2	3	4		
0	0	0	0	0	0	0
0.4	0	0	0	0	0	0
0.8	0.00592222	0.02254033	0	0	0.00711564	0.01065536
2	0.07986793	0.11951658	0.05727472	0.16368442	0.10508591	0.04677565
4	0.26374299	0.38601725	0.40590062	0.22666082	0.32058042	0.08871847
8	0.46515866	0.75354023	0.76322201	0.74266345	0.68114609	0.14423629
16	0.90565095	0.92694548	0.93200518	0.92711663	0.92292956	0.01175552
32	0.98106216	0.99225687	1	0.98680554	0.99003114	0.00806596
40	1	1	1	0.9950606	0.99876515	0.0024697

**Table S16. The binding data for  $\lambda$ -Cl with SOR3**

Cl conc ( $\mu$ M)	trials				average	SD
	1	2	3	4		
0	0	0	0	0	0	0
0.4	0	0	0	0	0	0
0.8	0.0263119	0.0322789	0.02467223	0.02234692	0.02640249	0.00424193
2	0.07515644	0.10437324	0.11292587	0.20621888	0.12466861	0.05672065
4	0.25803062	0.22253041	0.28552586	0.26466536	0.25768806	0.02620308
8	0.71389634	0.6372837	0.74045782	0.80179675	0.72335865	0.06817466
16	0.95229108	0.94763626	0.87673028	0.91823087	0.92372212	0.03476808
32	1	0.98218216	0.98313518	0.98713873	0.98811402	0.00820982
40	1	1	1	1	1	0

**Table S17. The binding data for  $\lambda$ -Cl with SOL1**

Cl conc ( $\mu$ M)	trials				average	SD
	1	2	3	4		
0	0	0	0	0	0	0
0.4	0.01463321	0	0	0	0.0036583	0.0073166
0.8	0.05317782	0.03911125	0.02728853	0.02268064	0.03556456	0.01362955
2	0.1501219	0.20956129	0.22729639	0.11753331	0.17612823	0.05113917
4	0.36805867	0.44948844	0.42042621	0.41620982	0.41354578	0.03374112
8	0.66252366	0.61674738	0.76790415	0.69628315	0.68586459	0.06366847
16	0.86342997	0.81569019	0.88523374	0.94945947	0.87845334	0.05553605
32	1	0.95291522	0.97682745	0.97735973	0.9767756	0.01922702
40	1	1	0.99311855	0.99574557	0.99721603	0.00338883

**Table S18.** The binding data for  $\lambda$ -Cl with SOL2

Cl conc (uM)	trials				average	SD
	1	2	3	4		
0	0	0	0	0	0	0
0.4	0.0906104	0.03082151	0.05562639	0.02286552	0.04998096	0.03046831
0.8	0.0891321	0.0933386	0.08992544	0.06878079	0.08529423	0.0111592
2	0.39913809	0.22492358	0.4390636	0.33564389	0.34969229	0.09344644
4	0.61427868	0.50909633	0.63514091	0.57753222	0.58401204	0.05533107
8	0.86265941	0.72272238	0.83910715	0.83233576	0.81420618	0.06235848
16	0.90159999	0.96373558	0.99163682	0.97925797	0.95905759	0.03996966
32	0.98467916	0.99040399	1	0.99443241	0.99237889	0.00646747
40	1	1	1	1	1	0

**Table S19.** The binding data for  $\lambda$ -Cl with SOL3

Cl conc (uM)	trials				average	SD
	1	2	3	4		
0	0	0	0	0	0	0
0.4	0.12759012	0.09792431	0.0715258	0.03365832	0.08267464	0.03990342
0.8	0.20679682	0.13688742	0.11548178		0.15305534	0.04775626
2	0.43725049	0.41679413	0.2863562	0.27681588	0.35430418	0.08447179
4	0.54367471	0.58354003	0.47767705	0.50614501	0.5277592	0.04597173
8	0.74873527	0.7451234	0.75414504	0.70663891	0.73866065	0.02166736
16	0.87716312	0.95709068	0.92910767	0.91189007	0.91881289	0.03343551
32	1	0.98414138	0.98647421	0.96934697	0.98499064	0.01255558
40	1	0.98994859	0.9926234	1	0.995643	0.00514818